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**EFEITOS DOS PRINCIPAIS ÁCIDOS ORGÂNICOS ACUMULADOS NA
ACIDÚRIA 3-HIDROXI-3-METILGLUTÁRICA SOBRE A HOMEOSTASE
REDOX, A RESPOSTA INFLAMATÓRIA E A FOSFORILAÇÃO DE
PROTEÍNAS DO CITOESQUELETO EM CÓRTEX CEREBRAL E ESTRIADO
DE RATOS JOVENS**

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“Uma vida não questionada não merece ser vivida.”

(Platão)

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SUMÁRIO

PARTE I – Introdução e Objetivos	1
RESUMO	2
ABSTRACT	4
Lista de Abreviaturas	6
Lista de Figuras	8
Lista de Tabelas	9
I INTRODUÇÃO	10
I.1 Erros Inatos do Metabolismo (EIM)	10
I.1.1 Conceito e classificação dos EIM	10
I.1.2 Manifestações clínicas	12
I.2 Acidúrias orgânicas	15
I.2.1 Acidúria 3-hidróxi-3-metilglutárica	16
I.2.1.1 Achados clínicos e neuropatológicos	18
I.2.1.2 Diagnóstico	19
I.2.1.3 Tratamento	19
I.2.1.4 Fisiopatologia	20
I.3 Radicais livres	21
I.3.1 Mecanismos de dano celular induzido por radicais livres	23
I.3.1.1 Lipoperoxidação	24
I.4 Defesas antioxidantes	26
I.4.1 Defesas antioxidantes não enzimáticas	27
I.4.2 Defesas antioxidantes enzimáticas	27
I.5 Estresse oxidativo	28
I.6 Citoesqueleto	32
I.6.1 Filamentos intermediários	33
I.6.1.1 Neurofilamentos	36
I.6.1.2 Proteína glial fibrilar ácida	37
I.6.1.3 Vimentina	38
I.7 Fosforilação	39
II OBJETIVOS	45
II.1 Objetivo geral	45
II.2 Objetivos específicos	45

PARTE II – Artigos Científicos	47
Capítulo 1	48
Capítulo 2	59
Capítulo 3	88
PARTE III – Discussão e Conclusões	136
III DISCUSSÃO	137
IV CONCLUSÕES	151
V PERSPECTIVAS	153
Referências bibliográficas	154

PARTE I

Introdução e Objetivos

RESUMO

A deficiência da atividade da enzima 3-hidroxi-3-metilglutaril-CoA liase (HL) (OMIM 246450), também conhecida por acidúria 3-hidroxi-3-metilglutárica (HMGA), é um distúrbio genético caracterizado bioquimicamente pelo acúmulo predominante dos ácidos 3-hidroxi-3-metilglutárico (HMG), 3-metilglutárico (MGA) e 3-metilglutacônico (MGT), bem como em menor grau do ácido 3-hidroxiisovalérico em tecidos e líquidos biológicos de indivíduos afetados. Os pacientes apresentam sintomas neurológicos graves e anormalidades no córtex cerebral e gânglios da base cuja fisiopatogenia é pouco conhecida. No presente estudo investigamos os efeitos dos principais metabólitos acumulados na HMGA sobre importantes parâmetros de estresse oxidativo em estriado de ratos injetados com esses compostos e em astrócitos cultivados de córtex cerebral, bem como sobre a fosforilação de proteínas do citoesqueleto de estriado e córtex cerebral de ratos com 30 dias de idade. Assim, estudamos inicialmente os efeitos da administração intraestriatal de HMG e MGA sobre importantes parâmetros de estresse oxidativo em ratos em desenvolvimento. Nossos resultados demonstram que o HMG e o MGA induziram dano oxidativo em lipídios e proteínas. Os dois ácidos também aumentaram a oxidação da 2'-7'-diclorofluorescina (DCFH), ao passo que apenas o HMG induziu a produção de óxido nítrico. Em relação às defesas antioxidantes o HMG e o MGA provocaram uma diminuição das concentrações de glutatona reduzida e das atividades da superóxido dismutase e glutatona redutase, bem como um aumento da atividade da glutatona peroxidase. Já o HMG também causou um aumento de atividade da catalase e uma diminuição da atividade da glicose-6-fosfato desidrogenase. Finalmente observou-se que antioxidantes preveniram completamente ou atenuaram as alterações dos parâmetros de estresse oxidativo causadas pelo HMG, reforçando a participação de espécies reativas nesses efeitos. Observou-se também que MK-801, um antagonista não-competitivo do receptor glutamatérgico do tipo N-metil-D-aspartato (NMDA), preveniu alguns dos efeitos provocados pelo HMG, indicando o envolvimento do receptor NMDA no desequilíbrio redox causado por esse metabólito. Tendo em vista que os astrócitos são importantes para proteção neuronal e são susceptíveis a danos por neurotoxinas, o passo seguinte de nossa investigação foi determinar os efeitos do HMG e MGA sobre parâmetros importantes da homeostase redox e produção de citocinas em astrócitos corticais cultivados. Ambos os ácidos orgânicos reduziram a função mitocondrial astrocitária sem alterarem a viabilidade celular e também diminuíram as concentrações de glutatona reduzida. Em contrapartida, ambos metabólitos aumentaram a formação de espécies reativas (oxidação da DCFH) e também provocaram um aumento na liberação das IL-1 β , IL-6 e TNF α através da via de sinalização de Erk. Finalmente investigamos os efeitos do HMG, MGA e MGT sobre a fosforilação da GFAP e subunidades NFL, NFM e NFH dos neurofilamentos (NF) em fatias de estriado e de córtex cerebral. Nossos resultados demonstraram que o HMG, o MGA e o MGT provocaram uma hipofosforilação da GFAP e todas as subunidades dos NF nas duas estruturas cerebrais. Verificamos ainda que a hipofosforilação induzida por HMG nas proteínas do citoesqueleto foi mediada pela inibição das proteínas cinases, sem alterações de proteínas fosfatases. Assim, demonstramos que uma inibição da PKA estaria envolvida na hipofosforilação da Ser55 na região aminoterminal da NFL,

bem como uma inibição da JNK, resultaria na hipofosforilação das repetições do tipo KSP na região carboxiterminal das subunidades NFM e NFH. Observou-se também que a hipofosforilação do citoesqueleto era dependente dos receptores glutamatérgicos sinápticos e extrasinápticos (subunidade NR2B) do tipo NMDA e também do íon Ca^{2+} . Além disso, o inibidor da óxido nítrico sintase, L-NAME, e o antioxidante TROLOX (análogo hidrossolúvel da vitamina E) preveniram completamente a hipofosforilação e a inibição das atividades da PKA e da JNK causada pelo HMG. Podemos então presumir que, os presentes dados fornecem evidências sólidas de que o estresse oxidativo induzido pelo HMG e MGA em estriados de ratos e em astrócitos corticais cultivados, além de uma resposta inflamatória evidenciada nessas células neurais e uma hipofosforilação em proteínas do citoesqueleto possam representar mecanismos patológicos importantes que contribuem, ao menos em parte, com as alterações cerebrais observadas nos pacientes afetados pela deficiência da HL.

ABSTRACT

3-Hydroxy-3-methylglutaryl-CoA lyase (HL) deficiency, also known as 3-hydroxy-3-methylglutaric aciduria (HMGA) is a genetic disorder biochemically characterized by predominant accumulation of 3-hydroxy-3-methylglutaric (HMG), 3-methylglutaric (MGA) and 3-methylglutaconic (MGT), as well as lesser amounts of 3-methylglutaconic and 3-hydroxyisovaleric acids in tissues and biological fluids of affected individuals. Clinically, the patients present neurological symptoms and basal ganglia injury, whose pathomechanisms are partially understood. In the present study, we investigated the effects of the main metabolites accumulating in HMGA on important parameters of oxidative stress in intrastrially injected rats, in cortical cultured astrocytes, as well as on the phosphorylation of striatal and cortical cytoskeletal proteins of 30 day old rats. Thus, we first investigated the effects of the intrastriatal administration of HMG and MGA on important parameters of oxidative stress in developing rats. Our results demonstrate that HMG and MGA induced lipid and protein oxidative damage. HMG and MGA also increased 2',7'-dichlorofluorescein oxidation, whereas only HMG elicited nitric oxide production. Regarding the antioxidant defenses, both organic acids decreased reduced glutathione concentrations and the activities of superoxide dismutase and glutathione reductase and increased glutathione peroxidase activity. HMG also provoked an increase of catalase activity and a diminution of glucose-6-phosphate dehydrogenase activity. We finally observed that antioxidants fully prevented or attenuated HMG-induced alterations of the oxidative stress parameters, further indicating the participation of reactive species in these effects. We also observed that MK-801, a non-competitive antagonist of the N-methyl-D-aspartate (NMDA) receptor, prevented some of these effects, indicating the involvement of the NMDA receptor in the redox imbalance provoked HMG effects. Considering that astrocytes are important for neuronal protection and are susceptible to damage by neurotoxins the next step of this work was investigated the effects of HMG and MGA on important parameters of redox homeostasis and cytokine production in cortical cultured astrocytes. Both organic acids decreased astrocytic mitochondrial function without altering cell viability, and also decreased the concentrations of reduced glutathione. In contrast, they increased reactive species formation (DCFH oxidation) and also provoked a significant increase of IL-1 β , IL-6 and TNF α release through the Erk signaling pathway. Finally, we investigated the effects of HMG, MGA and MGT on the phosphorylation of GFAP and the neurofilaments (NF) subunits NFL, NFM and NFH in striatal and cortical slices. Our results demonstrated that HMG, MGA and MGT caused hypophosphorylation of GFAP and of all NF subunits in both cerebral structures. We also found that the HMG-induced hypophosphorylation on the cytoskeletal proteins was mediated by the inhibition of protein kinases without altering protein phosphatases. Thus, we demonstrated that PKA inhibition was involved in the hypophosphorylation of Ser55 in the amino terminal region of NFL, as well as JNK inhibition resulting in the hypophosphorylation of KSP repeats in the carboxyl terminal region of NFM and NFH subunits. It was also observed that the cytoskeletal hypophosphorylation was dependent on synaptic and extra synaptic (NR2B subunit) NMDA glutamatergic receptors and also on Ca²⁺. Furthermore, the nitric oxide synthase inhibitor, L-NAME, and the antioxidant TROLOX (polar analog of

vitamin E) fully prevented the hypophosphorylation and the inhibition of PKA and JNK activities caused by HMG. We can then presume that all these data provide solid evidence that oxidative stress induced by HMG and MGA in rat striatum and cortical cultured astrocytes, beyond of a inflammatory response verified in this neural cell type and an hypophosphorylation of cytoskeletal proteins could represent important pathomechanisms that contribute, at least in part, the cerebral alterations observed in the patients affected by the HL deficiency.

Lista de abreviaturas

AMPc – adenosina 3',5'-monofosfato cíclico
ATP – adenosina trifosfato
CAT – catalase
CDK5 – proteína cinase dependente de ciclina 5
CREB – elemento de ligação em resposta ao AMPc
C-terminal – extremidade carboxiterminal
DCF – 2',7'-diclorofluoresceína
DCFH – 2',7'- diclorofluorescin
DCFH-DA – 2',7'-diclorofluorescin diacetato
DNA – ácido desoxiribonucléico
EIM – erros inatos do metabolismo
ERN – espécies reativas de nitrogênio
ERO – espécies reativas de oxigênio
ER – espécies reativas
FI – filamentos intermediários
G6PD – glicose-6-fodfato desidrogenase
GFAP – proteína glial fibrilar ácida
GPx – glutaciona peroxidase
GR – glutaciona redutase
GSH – glutaciona reduzida
HL – 3-hidróxi-3-metilglutaril-CoA liase
HMG – ácido -3-hidróxi-3-metilglutárico
HMGA – acidúria 3-hidróxi-3-metilglutárica
HNE – 4-hidroxinonenal
IL-1 β – interleucina 1 β
IL-6 – interleucina 6
JNK – cinase c-Jun N-terminal
KSP – serina-lisina-prolina
LDL – lipoproteína de baixa densidade
L-NAME – L-NG-nitroarginina metil ester
MAPK – proteínas cinases ativadas por mitógeno
MDA – malondialdeído

MF – microfilamentos
MGA – ácido 3-metilglutárico
MGT – ácido 3-metilglutacônico
MK-801 – dizocilpina
MT – microtúbulo
MTT – 3-(4,5-dimetiltiazol-2-il)-2,5-difeniltetrazólio
NF - neurofilamentos
NFH – neurofilamento de alto peso molecular
NFL – neurofilamento de baixo peso molecular
NFM – neurofilamento de médio peso molecular
NMDA – N-metil-D-aspartato
NOS – óxido nítrico sintase
NR2B – subunidade 2B do receptor NMDA
N-terminal – extremidade aminoterminal
8-OHdGA – 8-hidroxi-2'-deoxiguanosina
OHIVA – ácido 3 hidróxiisovalérico
OMIM – online mendelian inheritance in man
PKA – proteína cinase A
PKC – proteína cinase C
PKCamII – proteína cinase dependente de cálcio e calmodulina
PP1 – proteína fosfatase 1
PP2A – proteína fosfatase 2A
PP2B – proteína fosfatase 2B ou calcineurina
PUFA – ácidos graxos de cadeia lateral polinsaturada
RNA – ácido ribonucléico
SNC – sistema nervoso central
SNP – sistema nervoso periférico
SOD – superóxido dismutase
TNF α – fator de necrose tumoral α
TROLOX – ácido 6-hidróxi-2,5,7,8-tetrametilcromano-2-carboxílico

Lista de Figuras

Figura 1. Citoesqueleto das células eucarióticas.	33
Figura 2. Os FI são integradores da citoarquitetura.	35
Figura 3. Regulação das proteínas celulares por fosforilação.	40

Lista de Tabelas

Tabela I. Algumas espécies reativas.

22

I INTRODUÇÃO

I.1 Erros Inatos do Metabolismo (EIM)

I.1.1 Conceito e classificação dos EIM

O termo erros inatos do metabolismo (EIM) foi utilizado pela primeira vez por Archibald Garrod em 1908 durante estudos realizados com pacientes com alcaptonúria, doença em que os afetados excretam grandes quantidades de ácido homogentísico na urina. O pesquisador observou que, frequentemente, um ou mais indivíduos da mesma família eram portadores da doença sem que seus pais fossem afetados. Baseado também na observação da maior incidência de consanguinidade entre os pais dos pacientes e nas leis de Mendel, Garrod propôs um modelo de herança autossômica recessiva para este distúrbio. Através da determinação do ácido homogentísico na urina de pacientes com alcaptonúria e da observação de que esta substância era um metabólito normal da degradação da tirosina, relacionou este acúmulo a um bloqueio na conversão do ácido homogentísico até fumarato e acetoacetato. Verificou-se mais tarde que tais alterações resultavam da síntese qualitativa ou quantitativamente anormal de uma proteína enzimática (SCRIVER *et al.*, 2001).

Atualmente, mais de 500 EIM foram bioquimicamente caracterizados, sendo que a maioria envolve processos de síntese, degradação, transporte e armazenamento de moléculas no organismo (SCRIVER *et al.*, 2001). Os pacientes portadores de algum tipo de EIM apresentam sintomatologia variada, embora haja uma predominância de sintomas neurológicos e anormalidades cerebrais. Portanto, as manifestações clínicas mais comuns nos pacientes

afetados por EIM estão relacionadas ao acometimento do SNC, com retardo neuropsicomotor, crises encefalopáticas, neurodegeneração progressiva e outras alterações neurológicas (BURTON, 1987).

A maioria dos EIM deve-se à deficiência grave na atividade de uma enzima. A ausência ou deficiência severa da atividade enzimática leva a um bloqueio da rota metabólica com acúmulo dos substratos e seus derivados que potencialmente podem provocar sintomatologia grave e mesmo morte (MITCHELL e FUKAO, 2001).

Sinclair (1982) classificou os EIM em quatro grandes grupos conforme o tipo de função exercida pela proteína deficiente e o tecido envolvido, considerando ainda aspectos clínicos, bioquímicos, patológicos e terapêuticos:

- a) Distúrbios de transporte: envolvem proteínas transportadoras de moléculas orgânicas ou inorgânicas. Exemplos: deficiência de dissacaridases e defeito no transporte de magnésio.
- b) Distúrbios de armazenamento, degradação e secreção: envolvem proteínas relacionadas com o aparelho de Golgi ou lisossomos na sua grande maioria, provocando acúmulo de macromoléculas em tecidos específicos. Exemplos: doenças lisossômicas de depósito, glicogenoses e cistinose;
- c) Distúrbios de síntese: envolvem a síntese de proteínas com funções hormonais, de defesa imunológica, estrutural, etc. Exemplos: hiperplasia adrenal congênita por deficiência na atividade da enzima de 21-hidroxilase, hipogamaglobulinemia;
- d) Distúrbios do metabolismo intermediário: caracterizam-se por

deficiências enzimáticas das rotas do metabolismo intermediário de moléculas pequenas comprometendo importantes rotas, tais como o ciclo do ácido tricarboxílico, o ciclo da uréia ou outras rotas. Assim, o substrato da enzima deficiente se acumula a menos que haja uma rota alternativa para metabolizá-lo. Por outro lado, o produto final da reação não será formado, podendo haver uma deficiência do mesmo no organismo. Os mecanismos responsáveis pelos danos teciduais podem ocorrer, desta forma pela: 1. ação do substrato acumulado que pode ser liberado na circulação e transportado para todo o organismo causando efeitos tóxicos devido a alterações bioquímicas e danos em vários tecidos e particularmente no SNC; 2. pela falta de substâncias essenciais ao desenvolvimento do organismo causada pelo bloqueio metabólico. Os mais frequentes EIM constituem os grupos das acidúrias orgânicas, as aminoacidopatias, entre outros.

A identificação de um EIM permite a introdução de um tratamento eficaz quando estiver disponível. No entanto, não há tratamento efetivo para a maioria dos EIM no momento. Além disso, a ocorrência de outros afetados no mesmo grupo familiar pode ser evitada pela identificação de casais em risco, aos quais pode ser oferecido o aconselhamento genético e, em muitos casos, o diagnóstico pré-natal (SCRIVER *et al.*, 2001).

I.1.2 Manifestações clínicas

Como mencionado anteriormente, pacientes portadores de EIM apresentam sintomas muito variados e inespecíficos, inclusive aqueles que são afetados pelo mesmo distúrbio. Essa variação fenotípica deve-se a diferentes

graus de deficiência enzimática, área do metabolismo envolvida e tecidos afetados. Algumas manifestações clínicas aparecem com maior frequência no período neonatal, tais como deficiência no crescimento, vômitos, diarreia, convulsões, letargia ou coma, odor peculiar na pele ou urina e dificuldade alimentar, enquanto outras (atraso no desenvolvimento psicomotor, neurodegeneração progressiva) podem aparecer mais tarde (BURTON, 1987). Uma gama de sintomas e sinais clínicos deve alertar o médico para esse grupo de doenças. A forma de aparecimento dos sintomas é um fator importante na distinção entre os dois grandes grupos de EIM: erros inatos devido a alterações do metabolismo intermediário e erros inatos devidos ao depósito celular de macromoléculas. Os defeitos do metabolismo intermediário que levam ao acúmulo de moléculas pequenas (ex.: aminoácidos e ácidos orgânicos) geralmente têm uma apresentação clínica súbita e a evolução se caracteriza por episódios de agudização recorrentes geralmente precedidos por infecções, ingestão alimentar exagerada de alimentos específicos, cirurgia, jejum ou outras condições de catabolismo elevado, pois nestas situações ocorre degradação de proteínas ou lipídeos que vão originar os metabólitos tóxicos (ex.: defeitos do ciclo da uréia e acidemias orgânicas). São consideradas doenças de intoxicação e/ou de comprometimento do metabolismo energético, pois várias dessas entidades patológicas caracterizam-se por prejuízo deste metabolismo.

Nos intervalos entre as crises, os pacientes podem estar clinicamente normais e a concentração dos compostos acumulados em algumas dessas doenças próxima ao normal. Para muitas destas doenças é, portanto, essencial que as amostras para análise laboratorial sejam coletadas nos momentos de

crise metabólica. O exame físico geralmente é inespecífico, assim como os exames histopatológicos dos órgãos mais afetados. O tratamento agudo com restrição alimentar específica (proteínas, lipídios ou glicídios) associada ou não à suplementação de vitaminas (que ajudam as reações enzimáticas) mostra resultados extraordinários para algumas dessas doenças, retirando o paciente da crise em poucas horas ou dias.

Outros defeitos do metabolismo intermediário mostram uma evolução crônica desde o nascimento ou nos primeiros meses ou anos de vida. Nesses casos, a intoxicação é crônica (ex.: fenilcetonúria) e os afetados apresentam um atraso na aquisição das habilidades motoras (ex.: caminhar), não adquirindo as habilidades cognitivas normais.

Já os EIM de moléculas complexas ou de organelas (doenças lisossômicas de depósito e peroxissomais) manifestam-se na sua quase totalidade de uma forma crônica e progressiva, atingindo tecidos e órgãos (fígado, baço, medula óssea e encéfalo) em que os substratos (glicogênio, lipídeos complexos, mucopolissacarídeos) que não podem ser degradados ali se depositam. É comum encontrar nesses pacientes distorções e sinais clínicos específicos (hepatomegalia, esplenomegalia, leucodistrofia, etc.). A doença geralmente se manifesta após os primeiros meses ou anos de vida e é comum o afetado perder as habilidades motoras e cognitivas que já tinham sido adquiridas. Os exames histopatológicos nos afetados por várias dessas doenças são bastante informativos (presença de glicogênio no fígado nas glicogenoses, de lipídeos específicos em vários tecidos nas lipidoses, etc.). O tratamento dessas doenças, no entanto, é ainda experimental para a maioria

das doenças deste grupo, sendo que a reposição enzimática tem tido recentemente sucesso em várias delas (SCRIVER *et al.*, 2001; HUGHES *et al.*, 2007; SIMON *et al.*, 2008; CHOI *et al.*, 2008; ORTIZ *et al.*, 2008; VEDDER *et al.*, 2008; GRABOWSKI, 2008; van DUSSEN *et al.*, 2014; ITIER *et al.*, 2014; ANDERSON *et al.*, 2014a,b; HENLEY *et al.*, 2014).

I.2 Acidúrias orgânicas

As acidemias ou acidúrias orgânicas constituem um grupo de EIM caracterizados pelo acúmulo de um ou mais ácidos orgânicos nos líquidos biológicos e tecidos dos pacientes afetados devido à deficiência da atividade de enzimas do metabolismo de aminoácidos, lipídeos ou carboidratos (CHALMERS e LAWSON, 1982). A frequência destas doenças na população em geral é pouco conhecida, o que pode ser creditado à falta de laboratórios especializados para o seu diagnóstico e ao desconhecimento médico sobre essas enfermidades. Na Holanda, país considerado referência para o diagnóstico de EIM, a incidência dessas doenças é estimada em 1:2.200 recém-nascidos, enquanto, na Alemanha, Israel e Inglaterra é de aproximadamente 1:6.000 - 1:9.000 recém-nascidos (HOFFMANN *et al.*, 2004). Na Arábia Saudita, onde a consanguinidade é elevada, a frequência é de 1:740 nascidos vivos (RASHED *et al.*, 1994). Chalmers e colaboradores (1980) demonstraram que as acidemias orgânicas eram os EIM mais frequentes em crianças hospitalizadas motivando diversos estudos clínicos, laboratoriais e epidemiológicos nos anos seguintes.

Clinicamente os pacientes afetados por acidemias orgânicas apresentam predominantemente disfunção neurológica em suas mais diversas formas de

expressão: regressão neurológica, convulsões, coma, ataxia, hipotonia, hipertonia, irritabilidade, tremores, movimentos coreatéticos, tetraparesia espástica, atraso no desenvolvimento psicomotor, retardo mental e outras manifestações. As mais frequentes manifestações laboratoriais são cetose, cetonúria, neutropenia, trombocitopenia, acidose metabólica, baixos níveis de bicarbonato, hiperglicinemia, hiperamonemia, hipo / hiperglicemia, acidose láctica, aumento dos níveis séricos de ácidos graxos livres e outros (SCRIVER *et al.*, 2001). Recentemente, com o uso da tomografia computadorizada e ressonância magnética, foram encontradas na maioria dos pacientes afetados por essas doenças alterações de substância branca (hipomielização e / ou desmielização), atrofia cerebral generalizada ou dos gânglios da base (necrose ou calcificação), megaencefalia, atrofia frontotemporal e atrofia cerebelar (MAYATEPEK *et al.*, 1996).

1.2.1 Acidúria 3-hidroxi-3-metilglutárica

A acidúria 3-hidroxi-3-metilglutárica (HMGA), causada pela deficiência da enzima 3-hidroxi-3-metilglutaril-CoA liase (HL; EC 4.1.3.4), é uma acidúria orgânica de herança autossômica recessiva inicialmente descrita por Faull e colaboradores (1976a,b). A HL catalisa a clivagem do 3-hidroxi-3-metilglutaril-CoA (HMG-CoA) a acetoacetato e acetil-CoA, a qual é a última reação da síntese de corpos cetônicos e do catabolismo da leucina. Estudos demonstram a existência de 90-100 casos de HMGA no mundo, sendo que, com exceção da Arábia Saudita, Península Ibérica (Portugal e Espanha) e mais recentemente no Brasil a HMGA é rara em países europeus e no Japão (VARGAS *et al.*, 2007; MENAO *et al.*, 2009).

Duas isoformas da HL são encontradas, uma localizada na mitocôndria e a outra nos peroxissomos. A HL mitocondrial é um homodímero e o gene da enzima está localizado no cromossomo 1 (1p35.1-36.1), sendo composto por 9 éxons e 8 íntrons. A isoforma mitocondrial é composta por 298 resíduos de aminoácidos e contém uma seqüência peptídica de 27 resíduos de aminoácidos na extremidade N-terminal, que sinaliza a entrada da enzima na mitocôndria. Dentro da mitocôndria, o peptídeo é removido, formando a enzima madura com 298 resíduos de aminoácidos. Por outro lado, a isoforma peroxissomal possui 325 resíduos de aminoácidos. Existe uma grande diversidade e heterogeneidade de mutações na deficiência da HL; porém, dentro de comunidades específicas o padrão pode ser mais homogêneo. As mutações c.122G>A e c.109G>A são prevalentes na Arábia Saudita e Península Ibérica, respectivamente. Apesar do conhecimento de diferentes mutações, não há correlação entre o genótipo, a atividade enzimática e o prognóstico dos pacientes (SWEETMAN e WILLIAMS, 2001; AL-SAYED *et al.*, 2006; PIÉ *et al.*, 2007).

Devido ao defeito da HL, o principal metabólito acumulado na HMGA é o ácido 3-hidroxi-3-metilglutárico (HMG), derivado da hidrólise do HMG-CoA. Na urina de pacientes em tratamento, as concentrações de HMG variam entre 200-4.000 mmol / mol de creatinina (indivíduos normais nos primeiros meses de vida: 50-90 mmol / mol de creatinina). Contudo, durante crises de descompensação metabólica, esse metabólito pode alcançar níveis de 1.500-19.000 mmol / mol de creatinina. Com a reversibilidade da reação catalisada pela enzima 3-metilglutaconil-CoA hidratase, o ácido 3-metilglutacônico (MGT) também se encontra acumulado nos tecidos e líquidos biológicos dos

pacientes. Além disso, o MGT pode ser hidrogenado a ácido 3-metilglutárico (MGA). Também devido à reversibilidade da enzima 3-metilcrotonil-CoA carboxilase, altas concentrações de 3-metilcrotonil-CoA são encontradas e este metabólito pode ainda ser hidratado, gerando ácido 3-hidroxiisovalérico (OHIVA). Além disso, durante as crises de descompensação metabólica são detectadas outros ácidos, tais como o glutárico e o adípico (BONAFÉ *et al.*, 2000; SWEETMAN e WILLIAMS, 2001).

1.2.1.1 Achados clínicos e neuropatológicos

As manifestações clínicas da HMGA se apresentam usualmente no período neonatal (SWEETMAN e WILLIAMS, 2001). Durante episódios agudos de descompensação metabólica (jejum prolongado ou situações hipercatabólicas), os pacientes apresentam vômitos, diarreia, desidratação, hipotonia, hipotermia, letargia e apnéia que pode progredir ao coma (SWEETMAN e WILLIAMS, 2001; FUNGHINI *et al.*, 2001; POSPISILOVA *et al.*, 2003). Outros sintomas incluem hepatomegalia com hiperamonemia, macrocefalia, microcefalia, pancreatite aguda, retardo no desenvolvimento e cardiomiopatia. Alguns pacientes podem evoluir até a idade adulta (LEUNG *et al.*, 2009; REIMÃO *et al.*, 2009). Enfatiza-se que a deficiência da HL é fatal em aproximadamente 20 % dos casos. Os pacientes também apresentam acidose metabólica com hipoglicemia hipocetótica e elevação de transaminases no plasma. Os achados de ressonância magnética evidenciam lesões na substância branca cerebral que, em alguns casos, podem ser reversíveis com o tratamento baseado na restrição alimentar de leucina (YALÇINKAYA *et al.*, 1999; YYLMAZ *et al.*, 2006; ZAIFEIRIOU *et al.*, 2007). Além disso, foi

verificado o envolvimento dos núcleos caudato, dentado, globus pallidus, bem como do trato corticoespinhal (YALÇINKAYA *et al.*, 1999; YYLMAZ *et al.*, 2006).

I.2.1.2 Diagnóstico

A deficiência da HL deve ser investigada em pacientes que apresentam hipoglicemia hipocetótica e acidose metabólica. Geralmente a HMGA é diagnosticada através do aumento dos metabólitos acumulados HMG, MGT, MGA, 3-metilcrotonilglicina e OHIVA nos tecidos e líquidos biológicos dos pacientes (especialmente na urina) através da cromatografia gasosa acoplada à espectrometria de massa. A determinação da atividade da HL também é importante, podendo ser medida espectrofotometricamente em leucócitos, fibroblastos e plaquetas. A análise de DNA também é útil para o diagnóstico pré-natal (SWEETMAN e WILLIAMS, 2001).

I.2.1.3 Tratamento

A HMGA tem tratamento disponível, baseado em dieta hipoproteica bem como a administração de L-carnitina ($100 \text{ mg} \cdot \text{kg}^{-1} \cdot \text{dia}^{-1}$). Os pacientes devem evitar situações de jejum e ingesta excessiva de gordura. Além disso, a administração de glicose deve ser feita em vigência de episódios de hipoglicemia que ocorrem geralmente durante as crises. A restrição de leucina (proteína) e de gordura também é corriqueiramente utilizada (DASOUKI *et al.*, 1987).

I.2.1.4 Fisiopatologia

A patogênese da acidúria 3-hidróxi-3-metilglutárica não está bem estabelecida, no entanto, é possível que os metabólitos acumulados em tecidos e líquidos biológicos dos pacientes afetados sejam tóxicos e responsáveis por pelo menos parte dos sintomas apresentados. Os mecanismos pelos quais esses ácidos orgânicos levam aos sintomas característicos dessa doença ainda não foram totalmente desvendados, porém evidências *in vitro* e *in vivo* indicam alterações no metabolismo energético e nas oxidações biológicas.

Acredita-se que a deficiência dos corpos cetônicos (ácido acetoacético e 3-OH-butírico) durante o jejum ou durante fases agudas da doença que normalmente levam à hipoglicemia, possa explicar ao menos em parte o quadro clínico dos pacientes, já que os corpos cetônicos são uma alternativa energética importante para o cérebro.

Também foi verificado o envolvimento de espécies reativas na fisiopatologia da HMGA. Foi demonstrado que o HMG e o MGA induzem dano oxidativo lipídico e protéico e reduzem as defesas antioxidantes *in vitro* em córtex cerebral, estriado e fígado de ratos jovens (LEIPNITZ *et al.*, 2008a,b; 2009). Além disso, MGA induz disfunção mitocondrial e diminui a atividade da Na⁺,K⁺-ATPase em sinaptossomas preparados de cérebro de ratos (RIBEIRO *et al.*, 2011).

Um modelo genético nocaute de HMGA foi recentemente desenvolvido em camundongos, mas infelizmente os animais morrem ainda *in utero*,

impossibilitando, desta forma, a utilização desses animais para estudos de patogênese da doença (WANG *et al.*, 1998).

I.3 Espécies reativas e radicais livres

“Espécies reativas” (ER) é um termo usado coletivamente para designar tanto espécies “radicais” quanto “não radicais”, agentes oxidantes que facilmente são convertidos a radicais, tais como o ácido hipocloroso (HOCl), o ácido hipobromoso (HOBr), o ozônio (O₃), o peroxinitrito (ONOO⁻), o oxigênio *singlet* (¹O₂) e o peróxido de hidrogênio (H₂O₂). Sendo assim pode-se dizer que todo radical livre é uma espécie reativa, entretanto, nem toda espécie reativa é considerada um radical livre. E dentro dessa definição enquadram-se uma ampla gama de espécies e algumas delas são listadas na Tabela I (HALLIWELL, 2006).

Radicais livres são moléculas ou átomos que contenham um ou mais elétrons desemparelhados e possuam a capacidade de existir de forma independente (SOUTHORN e POWIS, 1988; HALLIWELL e GUTTERIDGE, 1999). O desemparelhamento de elétrons, situação energeticamente instável, é o que confere alta reatividade a essas espécies. Os radicais livres podem ser formados pela perda de um elétron de um não-radical ou pelo ganho de um elétron por um não-radical. Radicais podem também ser formados em um processo de fissão homolítica, no qual uma ligação covalente é quebrada e cada elétron do par compartilhado permanece com cada um dos átomos envolvidos (HALLIWELL e GUTTERIDGE, 1999). Quando um radical livre reage com um composto não-radical, outro radical livre pode ser formado; assim, a presença de um único radical pode iniciar uma seqüência de reações

em cadeia de transferência de elétrons (redox) (MAXWELL, 1995). Nas reações em cadeia induzidas pelos radicais livres, um radical reativo leva à formação de um produto que também é um radical livre e que, por sua vez, reage produzindo um terceiro radical.

Tabela I – Algumas espécies reativas

Radicais Livres	Não radicais
Espécies reativas de oxigênio (ERO)	
Superóxido, $O_2^{\bullet-}$	Peróxido de hidrogênio, H_2O_2
Hidroxila, OH^\bullet	Ozônio, O_3
Hidroperoxila, OH^\bullet_2 (superóxido protonado)	<i>Singlet</i> , 1O_2
Carbonato, $CO_3^{\bullet-}$	Peróxidos orgânicos, ROOH
Peroxila, RO_2^\bullet	Peroxomonocarbonato, $HOOCO_2^-$
Alcoxila, RO^\bullet	Peróxicarbonato nitroso, $ONOOCO_2^-$
Radical dióxido de carbono, $CO_2^{\bullet-}$	
Espécies reativas de cloro	
Cloro atômico, Cl^\bullet	Ácido hipocloroso, HOCl
	Cloreto de nitrila, NO_2Cl
	Cloraminas
	Gás cloro, Cl_2
	Dióxido de cloro, ClO_2
Espécies reativas de bromo	
Bromo atômico, Br^\bullet	Ácido hipobromoso, HOBr
	Gás bromo, Br_2

Cloreto de bromo, HOBr

Espécies reativas de nitrogênio (ERN)

Óxido nítrico, NO[•]

Ácido nitroso, HNO₂

Dióxido de nitrogênio, NO[•]₂

Cátion nitrosil, NO⁺

Nitrato, NO[•]₃

Ânion nitroxila, NO⁻

Tetróxido dinitrogenado, N₂O₄

Trióxido nitrogenado, N₂O₃

Peroxinitrito, ONOO⁻

Peroxinitrato, O₂NOO⁻

Ácido peroxinitroso, ONOOH

Cátion nitronico, NO₂⁺

Peroxinitritos alquila, ROONO

Peroxinitratos de alquila, RO₂ONO

Cloreto de nitrila, NO₂Cl

Peróxiacetil nitrato, CH₃C(O)OONO₂

Fonte: adaptado de Halliwell (2006)

I.3.1 Mecanismos de dano celular induzido por espécies reativas

As espécies reativas de oxigênio (ERO) e de nitrogênio (ERN) ocorrem tanto em processos fisiológicos quanto patológicos do organismo. Fisiologicamente essas espécies reativas apresentam diversas funções (BERGENDI *et al.*, 1999). Assim, um aumento da liberação local de radicais livres pode ser benéfico, como é o caso da liberação de espécies tóxicas oxidantes pelos neutrófilos, que podem atuar na defesa do hospedeiro contra uma infecção (DELANTY e DICHTER, 1998). Participam ainda de processos

de sinalização celular e também estão envolvidos na síntese e regulação de algumas proteínas (WARD e PETERS, 1995).

Por outro lado, quando formadas em excesso, essas espécies altamente reativas têm o potencial de oxidar moléculas biológicas incluindo proteínas, lipídios e DNA (MAXWELL, 1995). Com relação aos efeitos prejudiciais das reações oxidantes ao organismo, os radicais livres podem promover lipoperoxidação; podem causar a oxidação de lipoproteínas de baixa densidade (LDL); podem reagir com proteínas, levando à sua inativação e conseqüente alteração de sua função; e podem também reagir com o DNA e RNA, levando a mutações somáticas e a distúrbios de transcrição (DELANTY e DICHTER, 1998), entre outros efeitos.

1.3.1.1 Lipoperoxidação

Antes de ser vista como uma consequência de dano oxidativo, a lipoperoxidação deve ser considerada como um processo fisiológico contínuo que ocorre normalmente nas membranas celulares. Além de ser um fator de renovação da membrana, este processo é essencial na síntese de prostaglandinas e leucotrienos, bem como na fagocitose e pinocitose. No entanto, por serem formadas em grande parte por lipídios insaturados e proteínas, as membranas são particularmente vulneráveis ao ataque oxidativo. Assim, quando a produção de espécies reativas aumentar além da capacidade de detoxificação, esse processo será exacerbado, e com isso, a lipoperoxidação poderá acarretar profundas alterações na estrutura e na permeabilidade das membranas celulares. Isso irá causar perda de seletividade na troca iônica, liberação do conteúdo de organelas e formação de produtos

citotóxicos como o malondialdeído (MDA) e o 4-hidroxinonenal (HNE), entre outros eventos (FERREIRA e MATSUBARA, 1997).

Sob pH fisiológico, MDA ataca proteínas e bases do DNA, causando lesões mutagênicas (DRAPER e HADLEY, 1990). HNE pode provocar dano mitocondrial, inibição de síntese protéica e de DNA, e induz atividade quimiotáxica ou lise celular e morte (ESTERBAUER *et al.*, 1991). HNE parece ser particularmente citotóxico a células neurais (KARLHUBER *et al.*, 1997; KELLER *et al.*, 1999; MARK *et al.*, 1997). Nesse contexto, formação de HNE precede morte neuronal (ONG *et al.*, 2000) e concentrações desse produto lipídico final aumentadas são encontradas em tecido cerebral de pacientes afetados por diversas doenças neurodegenerativas (JENNER, 2003). Por lesar proteínas de membranas, esses produtos de lipoperoxidação podem também inibir a atividade de importantes enzimas de membranas, como Ca^{2+} -ATPase, glicerol-3-fosfato aciltransferase, glicose-6-fosfatase e Na^+ , K^+ -ATPase, e também canais de potássio (DUPRAT *et al.*, 1995; McCONNELL *et al.*, 1999).

A lipoperoxidação ocorre pelas etapas de iniciação, propagação e terminação. A iniciação é causada pelo ataque a um lipídio de membrana por parte de qualquer espécie que tenha reatividade suficiente para abstrair um átomo de hidrogênio de um grupo metileno ($-\text{CH}_2-$). Nesse contexto, os radicais hidroxila podem prontamente iniciar a lipoperoxidação. Já que o átomo de hidrogênio tem apenas um elétron, a abstração de H^\bullet de um grupo metileno deixa um elétron desemparelhado no carbono ($-\text{CH}^\bullet-$). Esse radical formado é geralmente estabilizado por um rearranjo molecular, formando um dieno conjugado. O destino mais provável desse radical é reagir com o O_2 , formando

um radical peroxila (ROO^*). Os radicais peroxila, por sua vez, são capazes de abstrair um próton de outra molécula lipídica, sendo esta fase conhecida como a fase de propagação. O radical de carbono formado pode reagir com o O_2 para formar outro radical peroxila, e assim sucessivamente. A abstração de um hidrogênio de outro lipídio por parte do radical peroxila gerará um hidroperóxido lipídico (LOOH). O término da reação poderá ocorrer quando dois radicais produzidos nas etapas anteriores reagirem entre si, formando um produto estável (HALLIWELL e GUTTERIDGE, 1999).

I.4 Defesas antioxidantes

Antioxidantes são substâncias endógenas ou exógenas que reduzem a formação de radicais livres ou reagem com os mesmos, neutralizando-os. A célula pode se proteger contra o dano oxidativo através de antioxidantes não enzimáticos e enzimáticos.

Embora diferindo na composição, as defesas antioxidantes estão amplamente distribuídas no organismo e compreendem agentes que removem cataliticamente os radicais livres, como as enzimas superóxido dismutase, catalase, glutathione peroxidase, entre outras; proteínas que minimizam a disponibilidade de pró-oxidantes (íons de ferro e cobre, por exemplo), ao se ligarem aos mesmos como as transferrinas; proteínas que protegem biomoléculas de danos (incluindo dano oxidativo) por outros mecanismos; agentes de baixo peso molecular que aprisionam espécies reativas de oxigênio e nitrogênio, como glutathione, α -tocoferol, ácido ascórbico e a bilirrubina.

I.4.1 Defesas antioxidantes não enzimáticas

As defesas antioxidantes não enzimáticas podem ser separadas em dois grandes grupos: agentes hidrossolúveis e lipossolúveis. O primeiro grupo inclui glutathiona reduzida (GSH), ácido ascórbico, ácido úrico, melatonina, os cofatores selênio e coenzima Q10 e proteínas plasmáticas. O tripeptídeo GSH é um dos mais efetivos e abundantes antioxidantes contra ERO, particularmente no cérebro onde as concentrações alcançam 1-10mmol/L (BAST, 1993). GSH mantém o equilíbrio redox na célula e inativa ERO. Ácido ascórbico também parece ser muito importante no cérebro porque células neurais têm um sistema de captação altamente eficiente e os níveis de ácido ascórbico no líquor são muito mais altos do que no plasma (LONNROT *et al.*, 1996).

I.4.2 Defesas antioxidantes enzimáticas

A atividade de antioxidantes enzimáticos também é importante para a detoxificação de radicais livres. Exemplos desses são as izoenzimas da superóxido dismutase contendo cobre e zinco (Cu, Zn-SOD) ou magnésio (Mn-SOD), a catalase (CAT), a glutathiona peroxidase (GPx) e a glutathiona redutase (GR). Qualquer desequilíbrio nesse sistema pode resultar em um aumento de radicais livres derivados do oxigênio. As isoformas Cu, Zn-SOD e Mn-SOD estão presentes no citosol e mitocôndria cerebrais, e juntos com a GSH, representam a linha de defesa mais importante contra a toxicidade do $O_2^{\bullet-}$. Isoformas da SOD geram H_2O_2 , o qual é removido pela atividade da GPx combinada com GSH ou GR, ou pela CAT (CHANCE *et al.*, 1979; HALLIWELL

e GUTTERIDGE, 1999b), ou adicionalmente por outras peroxidases (CHAE *et al.*, 1999).

I.5 Estresse oxidativo

Espécies reativas são necessárias para a função normal da célula, servindo como moléculas de sinalização para importantes papéis fisiológicos. Elas são continuamente produzidas e neutralizadas por sistemas de defesa antioxidante. No entanto, quando produzidos em altas concentrações ou quando defesas antioxidantes estão deficientes, elas podem causar dano celular. Se o aumento de espécies reativas é relativamente pequeno, a resposta antioxidante será suficiente para compensar esse aumento. No entanto, sob certas condições patológicas, a produção de espécies reativas está muito mais aumentada, e as defesas antioxidantes podem ser insuficientes para restabelecer a homeostase redox.

O rompimento entre o equilíbrio pro-oxidante e antioxidante é descrito como estresse oxidativo, e pode representar um mecanismo fundamental de doenças humanas (HALLIWELL e GUTTERIDGE, 2007; SIES, 1985). Assim, o termo “Estresse Oxidativo” é usado para se referir à situação na qual a geração de espécies reativas ultrapassa a capacidade das defesas antioxidantes disponíveis. Pode resultar tanto de uma diminuição das defesas antioxidantes quanto de uma produção aumentada de oxidantes, bem como da liberação de metais de transição ou a combinação de quaisquer desses fatores (HALLIWELL, 2001).

O estresse oxidativo pode promover adaptação, dano ou morte celular. Quando ocorre adaptação as células podem geralmente tolerar um estresse oxidativo moderado, que geralmente resulta em *up-regulation* da síntese de sistemas de defesa antioxidante a fim de restaurar o balanço oxidante/antioxidante. Apesar disso, nem sempre o estresse oxidativo precisa envolver defesas antioxidantes aumentadas.

No dano celular, o estresse oxidativo pode danificar todos os alvos moleculares (DNA, proteínas, carboidratos e lipídios) (HALLIWELL e GUTTERIDGE, 2007). A resposta ao dano pode ser reversível: a célula entra em *steady state* alterado temporário ou prolongado que não leva à morte celular.

A morte celular pode ocorrer tanto por necrose quanto por apoptose. Na morte celular por necrose, a célula incha e se rompe, liberando seu conteúdo para o meio extracelular. Pode haver a liberação de antioxidantes, como a catalase e a glutatona, e também de pró-oxidantes, como os íons cobre e ferro e proteínas do grupo HEME, agentes esses que podem afetar as células adjacentes, podendo até mesmo impor a elas um estresse oxidativo. Já na apoptose, o mecanismo intrínseco de suicídio celular é ativado, e não há a liberação do conteúdo celular. A morte celular por apoptose pode ser acelerada em certas doenças, como as doenças neurodegenerativas, havendo envolvimento do estresse oxidativo (HALLIWELL e GUTTERIDGE, 2007).

Além da indução de necrose e apoptose, o estresse oxidativo pode levar a um aumento da lipoperoxidação, cujos produtos (MDA e HNE, entre outros) são altamente neurotóxicos, e a um dano oxidativo tanto às proteínas, inibindo

a atividade de diversas enzimas e alterando a função celular, quanto ao DNA, causando alteração de bases púricas e pirimídicas (HALLIWELL e GUTTERIDGE, 2007).

O cérebro é altamente dependente de energia para o seu funcionamento normal e a mitocôndria é a estrutura intracelular responsável pela produção dessa energia. Para a produção eficiente de energia na forma de ATP, a mitocôndria possui uma alta demanda por oxigênio, já que utiliza uma grande quantidade de O_2 em uma massa de tecido relativamente pequena, o que torna esse tecido altamente susceptível à ação de espécies reativas.

Além disso, a presença de membranas celulares ricas em ácidos graxos de cadeia lateral poliinsaturada (PUFA), os quais são especialmente sensíveis ao ataque de radicais livres e conseqüente oxidação de lipídeos pode levar ao dano por radicais livres no tecido cerebral. Alto conteúdo de ferro, o qual favorece a lipoperoxidação e autooxidação de neurotransmissores através da OH^\bullet formado na reação de Fenton (ZALESKA e FLOYD, 1985), a considerável quantidade de microglia, macrófagos residentes do sistema nervoso que podem produzir $O_2^{\bullet-}$ e H_2O_2 , sua modesta defesa antioxidante, sendo os níveis de catalase particularmente baixos em muitas regiões cerebrais (HALLIWELL e GUTTERIDGE, 1996; HALLIWELL e GUTTERIDGE, 2007), aumentam a suscetibilidade do cérebro ao dano causado por radicais livres.

Além disso, uma disfunção mitocondrial pode ocorrer por diminuição da atividade dos complexos da cadeia respiratória com um conseqüente prejuízo no transporte de elétrons, o que leva a uma dispersão dos elétrons na forma de radicais livres potencialmente danosos à célula.

Numerosas evidências sugerem que os radicais livres e o estresse oxidativo possam estar envolvidos na patogênese dos danos neurológicos de várias doenças neurodegenerativas, como doença de Alzheimer, doença de Parkinson e esclerose lateral amiotrófica. Como medida de parâmetros de estresse oxidativo, existe um considerável número de casos onde se relatou aumento nos níveis de MDA e HNE no cérebro de pacientes, além de produtos da oxidação protéica, como grupos carbonil e 3-nitrotirosina, e também produtos que resultam da oxidação de DNA, bem como concentrações reduzidas dos antioxidantes não enzimáticos GSH e ácido ascórbico e diminuição da atividade das enzimas antioxidantes CAT e GPx (JENNER e OLANOW 1996; LIU *et al.*, 1999; PERRY *et al.*, 2003).

Estudos demonstraram uma diminuição na atividade do complexo I da cadeia respiratória em cérebros *postmortem* de pacientes portadores de doença de Parkinson. Essa inibição do complexo I pode acarretar na geração de espécies reativas, tais como ânion superóxido, radicais hidroxila e peroxinitrito, as quais poderiam causar um prejuízo ainda maior na cadeia transportadora de elétrons. Dessa forma, é possível que o estresse oxidativo e as disfunções mitocondriais formem um ciclo vicioso na doença de Parkinson (GU *et al.*, 1996; JANETSKY *et al.*, 1994; SCHAPIRA *et al.*, 1989, 1990 a, b).

Na doença de Alzheimer, a mais comum dentre as doenças neurodegenerativas, é possível que o estresse oxidativo tenha um papel chave na morte neuronal. Tem sido proposto que o peptídeo β -amilóide, o formador das chamadas placas senis, tenha a capacidade de gerar radicais livres espontaneamente. Estudos *in vivo* também evidenciaram um dano oxidativo

em cérebros humanos *postmortem* com doença de Alzheimer através da observação de aumento de 8-hidroxi-2'-deoxiguanosina (8-OHdGA), produtos de oxidação de outras bases e de RNA, carbonilas de proteínas, nitrotirosina e marcadores de peroxidação lipídica (SMITH *et al.*, 1991; MARKESBERY *et al.*, 1999; NOUROOZ-ZADEH *et al.*, 1999; LOVELL *et al.*, 2000).

I.6 Citoesqueleto

Para que as células funcionem adequadamente elas devem se organizar no espaço, interagir mecanicamente com o ambiente ao seu redor, apresentar uma conformação correta, ser fisicamente robustas e estar estruturadas de forma adequada internamente. Muitas células devem também ser capazes de modificar sua forma e migrar para outros locais. Além disso, todas as células devem ser capazes de reorganizar seus componentes internos como decorrência dos processos de crescimento, divisão e/ou adaptação à mudança no ambiente. Todas essas funções estruturais e mecânicas apresentam-se muito desenvolvidas em células eucarióticas, sendo dependentes de um sistema de filamentos denominado citoesqueleto (ALBERTS *et al.*, 2008).

As diferentes atividades do citoesqueleto dependem de três tipos de filamentos protéicos: filamentos de actina ou microfilamentos (MF), microtúbulos (MT) e filamentos intermediários (FI). Cada tipo é formado pela polimerização de monômeros específicos (CARRAWAY, 2000; ALBERTS *et al.*, 2008). Os três tipos de filamentos do citoesqueleto são conectados entre si e suas funções são coordenadas, permitindo a participação em inúmeras atividades celulares em conjunto com diversas proteínas associadas (BEAR *et al.*, 2002; ALBERTS *et al.*, 2008) (Figura 1).

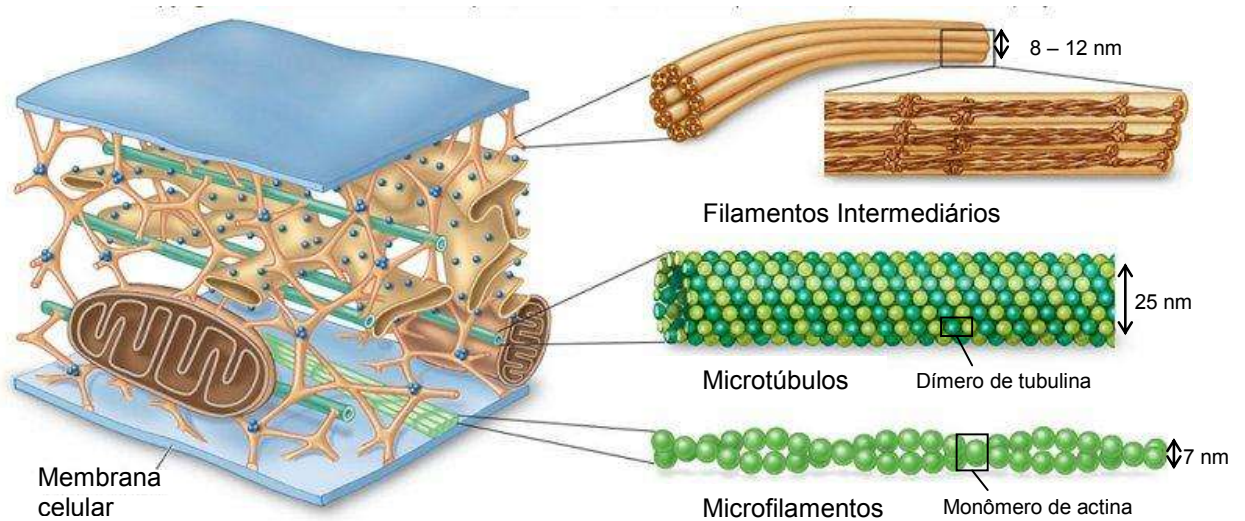


Figura 1 – Citoesqueleto das células eucarióticas. O citoesqueleto é constituído por microtúbulos, filamentos intermediários e microfilamentos. Os microfilamentos são formados por uma dupla fita e são responsáveis pela contração muscular, forma da célula e movimentos citoplasmáticos. Os filamentos intermediários são formados por subunidades que formam polímeros e ajudam na manutenção da forma celular, suportam a extensão das células nervosas e mantém as células unidas. Os microtúbulos são formados por dímeros de α e β -tubulina e estão associados com o transporte de organelas e cromossomos (adaptado de <<http://www.gceadvancelevel.com/cytoskeleton/>>).

I.6.1 Filamentos intermediários

Os filamentos intermediários são formados pela associação de subunidades fibrosas, constituindo fibras resistentes encontradas na maioria das células animais. Eles foram denominados intermediários por causa do seu diâmetro (8 – 10 nm), que está entre o diâmetro dos MF (7 – 8 nm) e o diâmetro dos MT (25 nm) (ALBERTS *et al.*, 2008). Os FI contituem a família mais diversificada de proteínas do citoesqueleto e são codificadas por cerca de 70 genes no genoma humano (FUCHS, 1994), possuindo um alto grau de especificidade celular e sendo frequentemente considerados marcadores de diferenciação celular (ALBERTS *et al.*, 2008, ERIKSSON *et al.*, 2009). As subunidades dos FI são constituídos por um domínio central altamente conservado em α -hélice e domínios N- e C-terminal variáveis sendo

classificados em seis sub-grupos, de acordo com o tipo celular e o padrão de desenvolvimento (FUCHS, 1994).

Inicialmente os FI eram considerados estruturas bastante estáticas e rígidas com funções relacionadas à integridade estrutural de células e tecidos (CHOU *et al.*, 1997; COULOMBE *et al.*, 2000; FUCHS e CLEVELAND, 1998; OMARY e KU, 1997). De fato, o papel dos FI em conferir resistência mecânica é muito relevante e a consequente perda de função pode estar relacionada com doenças envolvendo a fragilidade celular (FUCHS e CLEVELAND, 1998; OMARY *et al.*, 2004). Entretanto, nos últimos anos as funções atribuídas aos FI tem ampliado significativamente, pois tem sido relacionados a uma variedade de eventos celulares. Os FI desempenham um papel ativo no transporte de sinais da periferia celular para o núcleo (CHANG e GOLDMAN, 2004; PARAMIO e JORCANO, 2002), estão envolvidos na motilidade (ECKES *et al.*, 1998; HELFAND *et al.*, 2003), na adesão e migração celular (IVASKA *et al.*, 2005) e na modificação de vários processos celulares devido a sua habilidade de regular moléculas de sinalização (PALLARI e ERIKSSON, 2006). Outras funções atribuídas aos FI são a capacidade de agir como proteínas adaptadoras em vias de sinalização, bem como estarem envolvidos no crescimento e regeneração das células (KIM e COULOMBE, 2007) (Figura 2). Além disso, diversos trabalhos descrevem alterações dos FI em resposta a danos e nos processos de reparo tecidual, câncer e outras doenças (DePIANTO e COULOMBE, 2004; KUCHMA *et al.*, 2012).

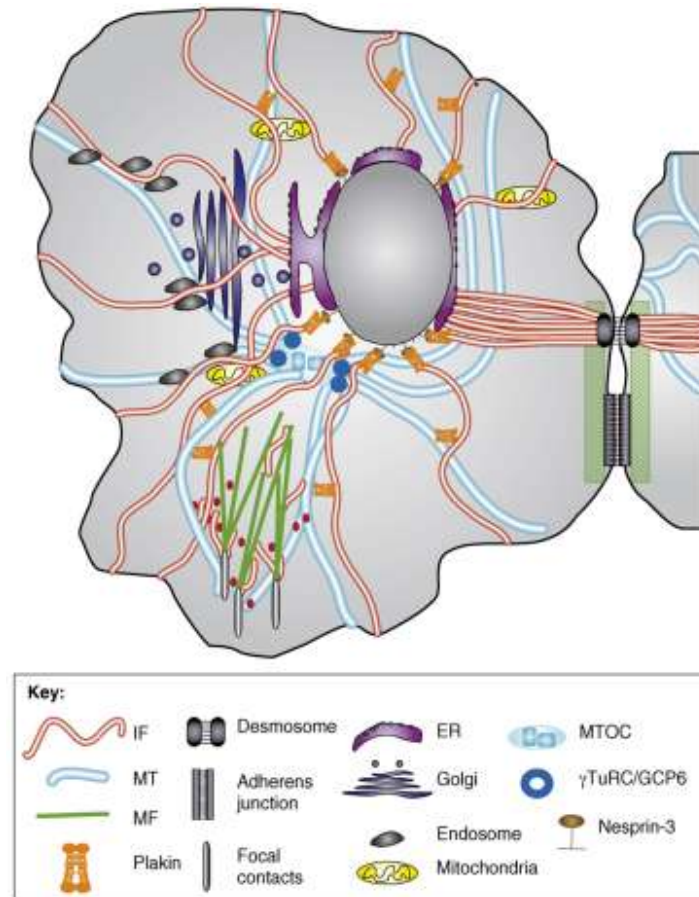


Figura 2 – Os FI são integradores da citoarquitetura. Eles estão ancorados no núcleo e se estendem para o citoplasma formando pontos de contato na membrana plasmática associando-se com a mitocôndria, às membranas do complexo de Golgi e ao sistema endolisossomal. Os FI posicionam os centros de organização dos microtúbulos (MTOC), através de interações com o complexo γ TuRC/GCP6, influenciando a organização dos MT e as propriedades de suas proteínas transportadoras, bem como, da distribuição das proteínas de membrana (adaptado de Godsel e colaboradores, 2007).

No sistema nervoso central (SNC) os FI tem papel fundamental na criação e manutenção da forma de neurônios e astrócitos, na manutenção do calibre axonal, bem como no transporte de organelas e substâncias envolvidas na transmissão sináptica (KIRKPATRICK e BRADY, 1999; ACKERLEY *et al.*, 2000). Essa rede protéica é um importante alvo para mecanismos de transmissão de sinais a partir de receptores de membrana plasmática, levando a uma resposta celular apropriada. Além disso, está envolvida na organização

e reorganização dos receptores de membrana, sendo essencial para os mecanismos de reconhecimento celular (CARRAWAY, 2000), bem como para a modulação da viabilidade celular através do processo de apoptose (NDOZANGUE-TOURIGUINE *et al.*, 2008).

1.6.1.1 Neurofilamentos

Os neurofilamentos (NF) são FI importantes na fisiologia neuronal, sendo os principais componentes do citoesqueleto dos neurônios maduros. São formados pela polimerização de três subunidades: os NF de baixo (NFL; 68 KDa), médio (NFM; 150 KDa) e alto peso molecular (NFH; 200 KDa) (ALBERTS *et al.*, 2008). A extremidade amino-terminal juntamente com a região em α -hélice da subunidade NFL interagem lateralmente e longitudinalmente formando a estrutura propriamente dita do NF (GEISLER e WEBER, 1981), enquanto as regiões carboxi-terminais das subunidades NFM e NFH são responsáveis pelas projeções laterais, que permitem a interação dos NFs entre si e com os demais constituintes do citoesqueleto (HISANAGA e HIROKAWA, 1988; GOTOW *et al.*, 1992; KIRKPATRICK e BRADY, 1999). Originalmente se assumia que os NF eram compostos apenas pelas subunidades NFL, NFM e NFH, porém estudos recentes indicam que outras proteínas como a α -internexina, no SNC, e a periferina, no sistema nervoso periférico (SNP), também se associam com os NF participando da formação do FI neuronal (BEAULIEU *et al.*, 1999; YUAN *et al.*, 2006).

A principal função atribuída aos NF é a manutenção do calibre axonal e conseqüentemente o aumento da velocidade de condução do impulso elétrico, processo esse regulado por fosforilação. Além disso, os NF também

contribuem para as propriedades dinâmicas do citoesqueleto axonal durante a diferenciação neuronal e para o crescimento, a regeneração e a orientação dos axônios (NIXON e SHEA, 1992). Eles são formados dentro do corpo celular e posteriormente transportados para o axônio, sendo a fosforilação do domínio carboxi-terminal um importante mecanismo regulatório do transporte axonal (JUNG *et al.*, 2000).

Acúmulo anormal de NF é descrito em várias doenças neurodegenerativas, tais como esclerose lateral amiotrófica, doença de Parkinson, doença de Alzheimer (AL-CHALABI e MILLER, 2003; LIU *et al.*, 2004) e em neuropatias tóxicas (ZHU *et al.*, 1998) sendo considerado, em muitos casos, um marcador nessas disfunções neuronais (SU *et al.*, 2012). Ainda não é compreendido como esse acúmulo de NF contribui para o processo neurodegenerativo nessas doenças, mas sugere-se que o transporte dos NF esteja interrompido nos neurônios afetados (ACKERLEY *et al.*, 2000; LARIVIERE e JULIEN, 2004).

1.6.1.2 Proteína glial fibrilar ácida

A proteína glial fibrilar ácida (GFAP) é uma proteína estrutural de 50 KDa sintetizada nos astrócitos e em algumas células de Schwann (GUO-ROSS *et al.*, 1999; KANEKO *et al.*, 1995; KOSAKO *et al.*, 1997). A GFAP é considerada marcador de astrócitos e é importante na modulação da motilidade e forma celular por fornecer estabilidade estrutural aos astrócitos. Além disso, esse FI é importante para a interação astrócito-neurônio (McCALL *et al.*, 1996; ELIASSON *et al.*, 1999) participando do sofisticado sistema de comunicação

intercelular recíproco que pode regular a liberação de neurotransmissores, a excitabilidade neuronal e a transmissão sináptica (CARMIGNOTO, 2000).

No SNC, após um dano, os astrócitos tornam-se reativos e respondem de uma maneira típica denominada astrogliose, que é caracterizada por uma rápida síntese de GFAP (EDDLESTON e MUCKE, 1993), sendo essa proteína, em muitos casos, um marcador de neurotoxicidade. Mudanças na expressão de GFAP podem alterar a morfologia dos astrócitos o qual pode afetar indiretamente outros tipos celulares e a estrutura do cérebro (MIDDELDORP e HOL, 2011). Uma intensiva gliose reativa é verificada em doenças como esclerose múltipla (HALLPIKE *et al.*, 1983), adrenoleucodistrofia (SCHAUMBURG *et al.*, 1975) e doença de Alexander (ENG *et al.*, 1998).

I.6.1.3 Vimentina

Entre as proteínas dos filamentos intermediários a vimentina é a mais amplamente distribuída ocorrendo em muitas células de origem mesenquimal. Além disso, é expressa de forma transitória durante o desenvolvimento em muitos tecidos (ALBERTS *et al.*, 2008). No SNC é expressa na glia radial, astrócitos imaturos e também na glia de Bergmann no cerebelo. Durante a diferenciação dos astrócitos é substituída progressivamente por GFAP (MENET *et al.*, 2001). Essa proteína, juntamente com a GFAP, participa do processo de gliose reativa após dano no SNC (MENET *et al.*, 2001; PEKKNY *et al.*, 1999).

A vimentina desempenha uma função importante na integridade estrutural de células e tecidos (NIEMINEM *et al.*, 2006), na adesão, na migração (GONZALES *et al.*, 2001; HOMAN *et al.*, 1998; TSURUTA e JONES,

2003; KREIS *et al.*, 2005), na transdução de sinal (JANOSCH *et al.*, 2000; IVASKA *et al.*, 2005; PERLSON *et al.*, 2005; KUMAR *et al.*, 2007) e em processos apoptóticos (YANG *et al.*, 2005).

Os FI sofrem alterações dinâmicas na sua organização citoplasmática durante diferentes estágios do ciclo celular ou em resposta a sinais celulares (STEINERT e ROOP, 1988; HELFAND *et al.*, 2005) sendo a fosforilação um mecanismo central na regulação da organização da rede citoplasmática de FI (ERIKSSON e GOLDMAN, 1993).

I.7 Fosforilação

A fosforilação protéica é uma modificação pós-traducional importante para as vias de transdução de sinal que controla o metabolismo, crescimento, divisão e diferenciação celular, motilidade, tráfego de organelas, transporte de membrana, contração muscular, imunidade, memória e aprendizado (MANNING *et al.*, 2002a,b).

Essa modificação covalente tem a propriedade de regular a função de proteínas em resposta a estímulos extracelulares (NESTLER e GREENGARD, 1999). Segundos mensageiros intracelulares, tais como o AMPc e o cálcio regulam funções neuronais através de alteração no estado de fosforilação de proteínas intracelulares. A fosforilação altera a função proteica de uma maneira rápida e reversível. O grau de fosforilação de uma proteína alvo reflete um balanço entre as ações contrárias de proteínas cinases e fosfatases, integrando um conjunto de rotas de sinalização celular (Figura 3). Entre o substrato de cinases e fosfatases estão enzimas, receptores de

neurotransmissores, canais iônicos e proteínas estruturais (PURVES *et al.*, 2005).

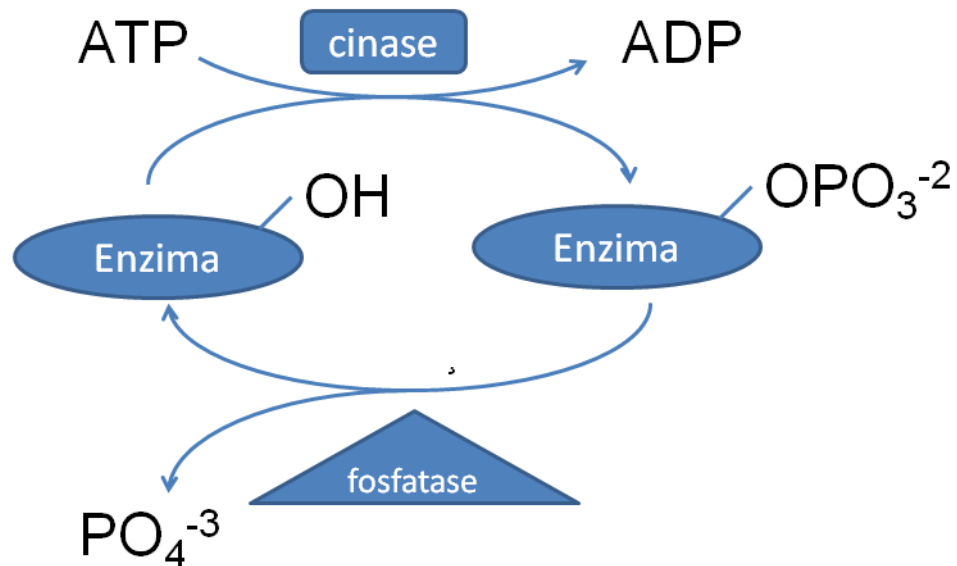


Figura 3 – Regulação das proteínas celulares por fosforilação. Proteínas cinases transferem grupos fosfato do ATP para resíduos de serina, treonina ou tirosina em um substrato protéico. A remoção dos grupos fosfato é catalisada por proteínas fosfatases (adaptado de Purves e colaboradores, 2005).

Os componentes do citoesqueleto estão entre as principais proteínas alvo modificadas por fosforilação em resposta a sinais extracelulares que podem determinar a morfologia neuronal (SANCHEZ *et al.*, 2000). Nos FI a fosforilação é o principal mecanismo responsável pela sua modulação, contribuindo na sua organização e função de maneira célula e tecido específica (GRANT e PANT, 2000; OMARY *et al.*, 2006).

Os FI podem ser fosforilados nas regiões aminoterminal e/ou carboxiterminal. A fosforilação do domínio aminoterminal está relacionada com a capacidade de associação e desassociação dos FI, sendo controlada por

cinases dependentes de segundo mensageiros (SIHAG *et al.*, 2007). Já no domínio carboxiterminal a fosforilação é realizada por cinases independentes de segundo mensageiros e tem implicações na interação dos FI com outras estruturas do citoesqueleto (CHOU *et al.*, 1997; GRANT e PANT, 2000).

Os FI são substratos de inúmeras serina/treonina cinases incluindo a proteína cinase C (PKC), a proteína cinase dependente de cálcio e calmodulina (PKCaMII), a proteína cinase dependente de AMPc (PKA), a família das proteínas cinases ativadas por mitógeno (MAPK), a proteína cinase dependente de ciclina 5 (CDK5), entre outras. Estas cinases possuem um papel significativo na regulação da estrutura e associação dos FI, bem como na interação deles com outras proteínas (ERIKSSON *et al.*, 2004; SIHAG *et al.*, 2007). Além disso, a ação dessas cinases nos FI é sítio específica, ou seja, elas fosforilam um resíduo específico dentro da estrutura protéica (SIHAG *et al.*, 2007). Proteínas serina/treonina fosfatases como a proteína fosfatase 1 (PP1) e as proteínas fosfatases 2A (PP2A) e 2B (PP2B ou calcineurina) também desempenham um papel importante na regulação do citoesqueleto (GRANT e PANT, 2000).

O estado de fosforilação dos NF tem um papel importante no controle da integridade do citoesqueleto, no transporte e no diâmetro axonal (MOTIL *et al.*, 2006; SIHAG *et al.*, 2007; STRACK *et al.*, 1997), além de promover a interação desses FI com as mitocôndrias (WAGNER *et al.*, 2003). Também pode modular a função de organelas (TOIVOLA *et al.*, 2005) e proteger os NF da proteólise (GRANT e PANT, 2000). Os NF são altamente fosforilados *in vivo*, porém o grau de fosforilação do NFM e do NFH e, especialmente do NFL, é

diferente em cada compartimento neuronal sendo altamente fosforilados no axônio, mas pouco fosforilados no corpo celular e nos dendritos (GOTOW e TANAKA, 1994; GOTOW *et al.*, 1994).

A região amino-terminal dos NF é fosforilada em sítios distintos na NFL e NFM. Os sítios mais frequentes de fosforilação dessas subunidades são os resíduos Ser55 na NFL fosforilados pela PKA e pela PKC e PKA nos resíduos Ser23 da NFM (SIHAG e NIXON, 1990; SIHAG *et al.*, 1999). Além disso, estudos *in vitro* mostraram que os resíduos Ser57 e Ser51 da NFL são substratos da PKCaMII e PKC, respectivamente (GONDA *et al.*, 1990; HASHIMOTO *et al.*, 2000).

A fosforilação da região carboxi-terminal das subunidades NFM e NFH dos FI ocorre em uma sequência de repetições de lisina, serina e prolina (repetições do tipo KSP) (GEISLER *et al.*, 1987; XU *et al.*, 1992; LEE *et al.*, 1988). A fosforilação desses sítios pode ser regulada por cascatas de transdução de sinal disparadas por fatores de crescimento, influxo de cálcio ou neurotoxinas, sendo realizadas por cinases como as MAPK e a CDK5 que são direcionadas por prolina para fosforilar a serina (LI *et al.*, 1999 a,b; PIEROZAN *et al.*, 2012).

Assim como nos NF, a polimerização de GFAP e da vimentina também é regulada pela fosforilação/desfosforilação do domínio aminoterminal que contém muitos sítios fosforiláveis (INAGAKI *et al.*, 1996; GOHARA *et al.*, 2001; TAKEMURA *et al.*, 2002). Para a GFAP foram identificados seis diferentes sítios de fosforilação Thr7, Ser8, Ser13, Ser17 e Ser34 na região aminoterminal e Ser389 na região carboxiterminal (INAGAKI *et al.*, 1990; NAKAMURA *et al.*,

1992; TSUJIMURA *et al.*, 1994; SEKIMATA *et al.*, 1996), entretanto pouco se sabe a respeito do papel de cada um desses sítios na associação, organização estrutural e disposição dos FI. A fosforilação/desfosforilação da região amino-terminal da GFAP em aminoácidos específicos regula a associação dessa proteína e é importante para a sua distribuição durante o ciclo celular (RALTON *et al.*, 1994; YASUI *et al.*, 1998; KAWAJIRI *et al.*, 2003). Além disso, sugere-se que a fosforilação da GFAP regula a plasticidade estrutural dos filamentos gliais e eventualmente as funções dos astrócitos e protege a GFAP da degradação (TAKEMURA *et al.*, 2002; KOROLAINEN *et al.*, 2005). Já a vimentina apresenta vários sítios específicos de fosforilação na região amino-terminal. Essa modificação pós-traducional nessa proteína parece ter um papel importante na migração celular (IVASKA *et al.*, 2005).

A fosforilação/desfosforilação é o principal mecanismo regulatório dos FI e de outras proteínas do citoesqueleto e alterações nos níveis fisiológicos de fosforilação/desfosforilação dessas proteínas são consideradas eventos críticos em patologias do SNC podendo levar a disfunção neurológica e morte celular (MILLER *et al.*, 2002). Várias doenças neurodegenerativas, tais como a doença de Alzheimer, a doença de Parkinson e a doença de Huntington são caracterizadas por acúmulo de agregados de filamentos insolúveis no citoplasma das células (GOEDERT, 1998; JULIEN, 1999; DiPROSPERO *et al.*, 2004), os quais estão relacionados à desregulação na atividade de cinases e fosfatases aos FI, constituindo-se em um mecanismo na neurodegeneração (PETZOLD, 2005).

Assim, no presente trabalho investigamos os efeitos dos principais ácidos orgânicos acumulados na HMGA sobre a homeostase redox em estriado de ratos jovens, sobre a resposta inflamatória e a homeostase redox em astrócitos de córtex cerebral de ratos neonatos e sobre a fosforilação de proteínas do citoesqueleto em estriado e córtex cerebral ratos jovens. Dividimos esse trabalho em três capítulos que correspondem a três artigos científicos, o primeiro já aceito e publicado e outros dois submetidos.

II OBJETIVOS

II.1 Objetivo geral

O objetivo geral desse trabalho foi o de explorar alguns dos mecanismos patogênicos dos principais ácidos orgânicos acumulados na acidúria 3-hidroxi-3-metilglutárica que possam explicar ao menos em parte o dano cortical e estriatal característico dos pacientes afetados por essa doença, focando na homeostase redox, resposta inflamatória e fosforilação de proteínas do citoesqueleto de ratos jovens.

II.2 Objetivos específicos

- Investigar os efeitos *ex vivo* da administração intraestriatal dos ácidos HMG e MGA sobre o dano oxidativo lipídico (níveis de substâncias reativas ao ácido tiobarbitúrico), dano oxidativo proteico (formação de carbonilas), defesas antioxidantes não-enzimáticas (concentrações de glutathione reduzida), formação de espécies reativas (oxidação da 2',7'-diclorofluorescina e determinação de nitritos e nitratos) e enzimáticas (atividades das enzimas antioxidantes glutathione peroxidase, glutathione redutase, superóxido dismutase, catalase e glicose-6-fosfato desidrogenase), em estriado de ratos de 30 dias de vida.

- Investigar os efeitos *in vitro* dos ácidos HMG e MGA sobre a formação de espécies reativas (oxidação da 2',7'-diclorofluorescina), as defesas antioxidantes não-enzimáticas (concentrações de glutathione reduzida), a viabilidade celular (formação do formazan e incorporação do iodeto de propídeo), bem como sobre a resposta inflamatória (concentrações de IL-1 β ,

IL-6, TNF α e NF κ B) e a indução de gliose reativa (quantificação de GFAP) em astrócitos de córtex cerebral de ratos neonatos cultivados (1 a 2 dias de vida).

- Investigar os efeitos *in vitro* dos ácidos HMG, MGA e MGT sobre o sistema de fosforilação dos FI (incorporação de [³²P]-ortofosfato de sódio, proteínas cinases e fosfatases) de astrócitos (GFAP) e neurônios (NFL, MFM e NFH), bem como, investigar os mecanismos de ação dos efeitos desencadeados pelos metabólitos testados em estriado e córtex cerebral de ratos de 30 dias de vida.

PARTE II

Artigos Científicos

Capítulo 1

In vivo experimental evidence that the major metabolites accumulating in 3-hydroxy-3-methylglutaryl-CoA lyase deficiency induce oxidative stress in striatum of developing rats: A potential pathophysiological mechanism of striatal damage in this disorder

Carolina Gonçalves Fernandes, Mateus Struecker da Rosa, Bianca Seminotti, Paula Pierozan, Rafael Wolter Martell, Valeska Lizzi Lagranha, Estela Natacha Brandt Busanello, Guilhian Leipnitz, Moacir Wajner

Artigo Científico publicado no periódico

Molecular Genetics and Metabolism



In vivo experimental evidence that the major metabolites accumulating in 3-hydroxy-3-methylglutaryl-CoA lyase deficiency induce oxidative stress in striatum of developing rats: A potential pathophysiological mechanism of striatal damage in this disorder



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Striatum

ABSTRACT

3-Hydroxy-3-methylglutaryl-CoA lyase (HL) deficiency is a genetic disorder biochemically characterized by predominant accumulation of 3-hydroxy-3-methylglutaric (HMG) and 3-methylglutaric (MGA) acids in tissues and biological fluids of affected individuals. Clinically, the patients present neurological symptoms and basal ganglia injury, whose pathomechanisms are partially understood. In the present study, we investigated the *ex vivo* effects of intra-striatal administration of HMG and MGA on important parameters of oxidative stress in striatum of developing rats. Our results demonstrate that HMG and MGA induce lipid and protein oxidative damage. HMG and MGA also increased 2',7'-dichlorofluorescein oxidation, whereas only HMG elicited nitric oxide production, indicating a role for reactive oxygen (HMG and MGA) and nitrogen (HMG) species in these effects. Regarding the enzymatic antioxidant defenses, both organic acids decreased reduced glutathione concentrations and the activities of superoxide dismutase and glutathione reductase and increased glutathione peroxidase activity. HMG also provoked an increase of catalase activity and a diminution of glucose-6-phosphate dehydrogenase activity. We finally observed that antioxidants fully prevented or attenuated HMG-induced alterations of the oxidative stress parameters, further indicating the participation of reactive species in these effects. We also observed that MK-801, a non-competitive antagonist of the N-methyl-D-aspartate (NMDA) receptor, prevented some of these effects, indicating the involvement of the NMDA receptor in HMG effects. The present data provide solid evidence that oxidative stress is induced *in vivo* by HMG and MGA in rat striatum and it is presumed that this pathomechanism may explain, at least in part, the cerebral alterations observed in HL deficiency.

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Abbreviations: CAT, catalase; DCF, 2',7'-dichlorofluorescein; DCFH, 2',7'-dichlorofluorescein; DCFH-DA, 2',7'-dichlorofluorescein diacetate; DNPH, 2,4-dinitrophenylhydrazine; GC/MS, gas chromatography/mass spectrometry; G6PD, glucose-6-phosphate dehydrogenase; GPx, glutathione peroxidase; GR, glutathione reductase; GSH, reduced glutathione; GSSG, oxidized glutathione; HL, 3-hydroxy-3-methylglutaryl-CoA lyase; HMG, 3-hydroxy-3-methylglutaric acid; HMGA, 3-hydroxy-3-methylglutaric aciduria; L-NAME, N^ω-nitro-L-arginine methyl ester; MEL, melatonin; MGA, 3-methylglutaric acid; NAC, N-acetylcysteine; NMDA, N-methyl-D-aspartate; RNS, reactive nitrogen species; ROS, reactive oxygen species; SOD, superoxide dismutase; SPSS, statistical package for the social sciences; TBA-RS, thiobarbituric acid-reactive substances; TCA, trichloroacetic acid.

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1. Introduction

3-Hydroxy-3-methylglutaryl-CoA lyase (HL) deficiency (OMIM 246450) is an autosomal recessive genetic disorder biochemically characterized by predominant tissue accumulation and high urinary excretion of large quantities of 3-hydroxy-3-methylglutarate (HMG) and 3-methylglutarate (MGA), as well as 3-methylglutaconate, 3-hydroxyisovalerate and 3-methylcrotonylglycine to lesser amounts [1–3].

Clinical presentation usually occurs in the first year of life. Acute episodes are common and are characterized by vomiting, diarrhea, dehydration, hypotonia, hypothermia, lethargy, coma and apnea, as well as by metabolic acidosis and hypoketotic hypoglycemia. Other signs include macrocephalia, developmental delay, hepatomegalia with liver test alterations, acute pancreatitis and dilated cardiomyopathy [2,4–7]. This disease is fatal in approximately 20% of cases, although

the symptoms tend to be milder after childhood [8,9]. Cerebral magnetic resonance neuroimaging shows predominantly abnormalities in the caudate nucleus and dentate nucleus and multiple coalescent lesions in periventricular subcortical white matter and arcuate fibers [10–12].

The mechanisms underlying the pathogenesis of the brain damage in HL deficiency are poorly known. However, it has been suggested that hypoglycemia associated with deficit of ketone bodies, secondary carnitine deficiency and intracellular accumulation of toxic organic acids may be involved in the pathophysiology of this disease [13–15]. In this context, previous *in vitro* studies demonstrated that HMG and MGA induce lipid and protein oxidative damage and reduce the non-enzymatic antioxidant defenses in cerebral cortex, striatum and liver of young rats [16–18]. It was also shown that MGA induces mitochondrial dysfunction and inhibits Na^+ , K^+ -ATPase activity in synaptosomes prepared from rat brain [19]. However, to the best of our knowledge, there is no study on the *in vivo* effects of the major metabolites accumulating in HL deficiency on redox cellular homeostasis.

Therefore, in the present work we investigated whether high striatum concentrations of HMG and MGA achieved by an acute intrastriatal injection of these organic acids to young rats could affect redox homeostasis in the hope to better clarify the pathomechanisms of striatal abnormalities observed in HL deficiency. For this purpose, important parameters of oxidative stress were determined, including malondialdehyde (MDA) concentrations (lipid peroxidation), carbonyl formation (protein oxidative damage), 2',7'-dichlorofluorescein (DCFH) oxidation, nitric oxide production, reduced glutathione (GSH) concentrations (non-enzymatic antioxidant defenses) and the activities of the antioxidant enzymes superoxide dismutase (SOD), glutathione reductase (GR), glucose-6-phosphate dehydrogenase (G6PD), glutathione peroxidase (GPx) and catalase (CAT).

2. Materials and methods

2.1. Reagents

All reagents used were of analytical grade and purchased from Sigma Co. (St. Louis, MO, USA). The solutions of HMG (4 M), MGA (4 M) and NaCl were prepared in water and the pH was adjusted to 7.4 with NaOH. N-acetylcysteine (NAC), melatonin (MEL), N^{G} -nitro-L-arginine methyl ester (L-NAME), dizocilpine maleate (MK-801) and the combination of α -tocopherol (vitamin E) plus ascorbic acid (vitamin C) were dissolved in water, and the pH was similarly adjusted to 7.4.

2.2. Animals

Thirty-day-old male Wistar rats, obtained from the Central Animal House of the Department of Biochemistry, ICBS, Universidade Federal do Rio Grande do Sul, Porto Alegre, RS, Brazil, were used. The animals were maintained on a 12:12 h light/dark cycle (lights on 07.00–19.00 h) in air conditioned constant temperature ($22 \pm 1^\circ\text{C}$) colony room, with free access to water and 20% (w/w) protein commercial chow (SUPRA, Porto Alegre, RS, Brazil). The experimental protocol was approved by the Ethics Committee for Animal Research of the Universidade Federal do Rio Grande do Sul, Porto Alegre, Brazil and followed the NIH Guide for the Care and Use of Laboratory Animals (NIH publication 85-23, revised 1996). All efforts were made to minimize the number of animals used and their suffering.

2.3. Administration of 3-hydroxy-3-methylglutaric (HMG) and 3-methylglutaric acids (MGA)

Rats were anesthetized with an intraperitoneal (i.p.) injection of equitiesine (3.33 mL/Kg), which consists of a mixture of 0.25 M chloral hydrate, 88 mM magnesium sulfate heptahydrate, 10 mg/mL

sodium thiopental, 5.8 M propylene glycol and 1.97 M ethanol. They were thereafter placed on a stereotaxic apparatus. Two small holes were drilled in the skull and 1 μL (4 μmol) of a HMG, MGA or NaCl solution (control group) was slowly injected into each striatum over 3 min via a needle connected by a polyethylene tube to a 10 μL Hamilton syringe. The needle was left in place for another 1 min before being softly removed, so that the total procedure lasted 4 min. The coordinates for injections were as follows: 0.6 mm posterior to the bregma, 2.6 mm lateral to the midline and 4.5 mm ventral from dura [20]. In some experiments, animals were pre-treated i.p. with MEL (100 mg/kg), NAC (150 mg/kg) or vitamin E (40 mg/kg) plus vitamin C (100 mg/kg) for 3 days, after which they received an acute injection of HMG. In other experiments, MK-801 (6 nmol) or L-NAME (0.2 nmol) was injected intrastrially 30 min before the administration of HMG.

Some animals aged 7 and 30 days of life were injected with a single dose of HMG or MGA (10 $\mu\text{mol/g}$ body weight of each organic acid) and sacrificed 30 min after the administration. The concentrations of the organic acids were measured in striatum from these animals 30 min after injection.

2.4. Striatum preparation

The rats were sacrificed by decapitation without anesthesia 30 min after the intrastriatal injection of the organic acid or NaCl. We sacrificed the animals and collected the samples 30 min after the injection because we aimed to investigate short-lived effects (acute evaluation) of the major metabolites accumulating in 3-hydroxy-3-methylglutaric aciduria. The brain was rapidly excised on a Petri dish placed on an ice plate. The olfactory bulb, pons, medulla, cerebral cortex and cerebellum were discarded and the striatum was dissected, weighed, and homogenized in 4 (1:4, w/v) (measurement of nitric oxide production) or 10 volumes (1:10, w/v) (evaluation of the other oxidative stress parameters) of 20 mM sodium phosphate buffer, pH 7.4, containing 140 mM KCl. Homogenates were centrifuged at 750 g for 10 min at 4°C to discard nuclei and cell debris [21]. The pellet was discarded and the supernatant, a suspension of mixed and preserved organelles, including mitochondria, was separated and used to measure oxidative stress parameters. Slices (400 μm) of striatum were also prepared and used for DCFH oxidation.

2.5. Organic acid analysis

HMG and MGA quantification in brain homogenates was performed by gas chromatography/mass spectrometry (GC/MS) according to Sweetman [22] with some modifications [23]. For this analysis, we used striatum homogenates (1:10 w/v in 20 mM sodium phosphate buffer, pH 7.4, containing 140 mM KCl) prepared from rats that received a single intrastriatal administration of HMG (4 μmol) or MGA (4 μmol), as described previously. Brain homogenates previously incubated for 1 h at 37°C with HMG (20 mM) or MGA (20 mM) to evaluate the metabolism of these organic acids in the brain. We also utilized striatum from 7- and 30-day-old rats injected i.p. with 10 $\mu\text{mol/g}$ body weight of HMG or MGA, which was homogenized (1:10, w/v) in 20 mM sodium phosphate buffer, pH 7.4, containing 140 mM KCl. An aliquot of 500 μL of each sample (brain homogenates) was first acidified to pH 1.5–2.0 and mixed with 100 μL hexadecane (internal standard). Standard solutions of HMG and MGA at the concentration of 1 mM were also submitted to the same procedure.

2.6. Oxidative stress parameters

2.6.1. Malondialdehyde (MDA) concentrations

MDA concentrations were estimated by measuring thiobarbituric acid-reactive substances (TBA-RS) [24], with slight modifications. Briefly, 200 μL of 10% trichloroacetic acid (TCA) and 300 μL of 0.67%

thiobarbituric acid in 7.1% sodium sulfate were added to tissue supernatants containing 0.3 mg of protein and incubated for 2 h in a boiling water bath. The resulting pink-stained complex was extracted with 400 μ L of butanol. Fluorescence of the organic phase was read at 515 nm and 553 nm as excitation and emission wavelengths, respectively. Calibration curve was performed using 1,1,3,3-tetramethoxypropane. MDA concentrations were calculated as nmol MDA/mg of protein.

2.6.2. Protein carbonyl formation

Protein carbonyl formation was measured spectrophotometrically according to Reznick and Packer [25]. Two hundred microliters of striatum supernatants containing 0.3 mg of protein were treated with 400 μ L of 10 mM 2,4-dinitrophenylhydrazine (DNPH) dissolved in 2.5 N HCl or with 2.5 N HCl (blank) and left in the dark for 1 h. Samples were then precipitated with 600 μ L of 20% TCA and centrifuged for 5 min at 10,000 g. The pellet was then washed with 1 mL ethanol:ethyl acetate (1:1, v/v) and suspended in 550 μ L of 6 M guanidine prepared in 2.5 N HCl at 37 °C for 5 min. The absorbance was read at 365 nm. The difference between the DNPH-treated and HCl-treated samples (blank) was used to calculate the carbonyl content determined at 365 nm. The results were calculated as nmol of carbonyl groups/mg of protein, using the extinction coefficient of $22,000 \times 10^6$ nmol/mL for aliphatic hydrazones.

2.6.3. 2',7'-Dichlorofluorescein (DCFH) oxidation

Reactive oxygen species (ROS) production was assessed by determining DCFH oxidation according to LeBel [26]. 2',7'-Dichlorofluorescein diacetate (DCFH-DA) was prepared in 20 mM sodium phosphate buffer, pH 7.4, containing 140 mM KCl and incubated with tissue slices (30 mg) during 30 min at 37 °C. DCFH-DA is enzymatically hydrolyzed by intracellular esterases to form non-fluorescent DCFH, which is then rapidly oxidized to form highly fluorescent 2',7'-dichlorofluorescein (DCF) in the presence of ROS. The DCF fluorescence intensity parallels the amount of ROS formed. Fluorescence was measured using excitation and emission wavelengths of 480 and 535 nm, respectively. Calibration curve was performed with standard DCF (0.1–1 μ M) and the levels of ROS were calculated as pmol DCF formed/mg protein.

2.6.4. Nitrate and nitrite content

Nitrate and nitrite concentrations were evaluated according to Navarro-González [27] with some modifications. One hundred and fifty microliters of tissue supernatants (containing approximately 1.2 mg of protein) were deproteinized by adding 125 μ L of 75 mM ZnSO₄ solution, followed by centrifugation at 9000 g for 2 min at 25 °C. The supernatant obtained was neutralized with 55 mM NaOH solution and diluted in 5 volumes of glycine buffer solution, pH 9.7. Copper-coated cadmium granules (600–1000 mg) were added to the supernatants to convert all nitrate into nitrite in the biological samples. Aliquots of 200 μ L were then treated with 200 μ L of Griess reagent (2% sulfanilamide in 5% HCl and 0.1% N-1-(naphthyl)ethylenediamine in H₂O) and incubated at room temperature by 10 min. The absorbance was read at 505 nm. A calibration curve was prepared with NaNO₂ at concentrations ranging from 1 to 125 μ M. The final results were expressed in μ mol of nitrate and nitrite/mg of protein.

2.6.5. Reduced glutathione (GSH) concentrations

GSH concentrations were measured according to Browne and Armstrong [28] with slight modifications. One hundred and eighty-five microliters of 100 mM sodium phosphate buffer, pH 8.0, containing 5 mM EDTA, and 15 μ L of o-phthalaldehyde (1 mg/mL) were added to 30 μ L of sample (0.3–0.5 mg of protein) previously deproteinized with metaphosphoric acid. This mixture was incubated at room temperature in a dark room for 15 min. Fluorescence was measured using excitation and emission wavelengths of 350 nm and 420 nm,

respectively. Calibration curve was prepared with standard GSH (0.001–1 mM) and the concentrations were calculated as nmol/mg of protein.

2.6.7. Superoxide dismutase (SOD) activity

SOD activity was assayed according to Marklund [29]. The reaction medium contained 50 mM Tris buffer/1 mM ethylenediaminetetraacetic acid, pH 8.2, 80 U/mL catalase, 0.38 mM pyrogallol and approximately 1 μ g of protein. The inhibition of autoxidation of pyrogallol, a process dependent on superoxide, occurs in the presence of SOD, whose activity can be then indirectly assayed spectrophotometrically at 420 nm. The results are reported as U/mg of protein.

2.6.8. Glutathione reductase (GR) activity

GR activity was measured according to Carlberg and Mannervik [30]. The enzyme activity was determined by monitoring the NADPH decrease at 340 nm in a medium with 200 mM sodium phosphate buffer, pH 7.5, containing 6.3 mM ethylenediaminetetraacetic acid, 1 mM GSSG, 0.1 mM NADPH and tissue supernatants (approximately 3 μ g of protein). The specific activity was calculated and expressed as U/mg of protein.

2.6.9. Glucose-6-phosphate dehydrogenase (G6PD) activity

G6PD activity was measured by the method of Leong and Clark [31], in which the reaction mixture contained 100 mM Tris-HCl, pH 7.5, 10 mM MgCl₂, 0.5 mM NADP⁺ and approximately 3 μ g of protein. The reaction was started by the addition of 1 mM glucose-6-phosphate and was followed in a spectrophotometer at 340 nm. The specific activity is represented as U/mg of protein.

2.6.10. Glutathione peroxidase (GPx) activity

GPx activity was measured according to Wendel [32] using tert-butylhydroperoxide as substrate. The enzyme activity was determined by monitoring the NADPH decrease at 340 nm in a medium containing 100 mM potassium phosphate buffer/1 mM EDTA, pH 7.7, 2 mM GSH, 0.1 U/mL glutathione reductase, 0.4 mM azide, 0.5 mM tert-butyl-hydroperoxide, 0.1 mM NADPH and approximately 3 μ g of protein. The specific activity was calculated as U/mg of protein.

2.6.11. Catalase (CAT) activity

CAT activity was assayed according to Aebi [33] by measuring the absorbance decrease at 240 nm in a reaction medium containing 20 mM H₂O₂, 0.1% Triton X-100, 10 mM potassium phosphate buffer, pH 7.0, and approximately 1 μ g of protein. The specific activity was calculated as U/mg of protein.

2.7. Protein content

The protein content was determined by the method of Lowry et al. [34], using bovine albumin as standard.

2.8. Statistical analysis

Data were expressed as means \pm SD for absolute values. Assays were performed in duplicate or triplicate and the mean was used for statistical analysis. Data were analyzed using one-way analysis of variance (ANOVA) followed by the Duncan multiple range test when the F value was significant. Differences between groups were rated significant at $P < 0.05$. All analyses were carried out in an IBM-compatible PC computer using the Statistical Package for the Social Sciences (SPSS) software.

3. Results

3.1. HMG and MGA concentrations achieved in rat striatum

We first determined the concentrations of HMG and MGA achieved in the striatum after a single injection of each of the organic acids. We verified that mean HMG and MGA concentrations achieved in the striatum were respectively 140 and 200 μM .

Next we tested *in vitro* whether 1) HMG could be converted to MGA by measuring MGA formation from HMG in brain homogenates and 2) MGA could be metabolized in the brain. We observed under our conditions that MGA could not be detected when HMG was incubated for 1 h with brain homogenates and that MGA was not metabolized in the brain since the amount of MGA remained the same when brain was exposed for 1 h with this organic acid (results not shown). These data suggest that the brain does not contain the enzymes necessary for HMG and MGA metabolism as it was previously reported [35] and indicate that our results were due to true effects of HMG and MGA injected into the striatum.

We also investigated whether HMG and MGA are able to cross the blood–brain barrier (BBB) by *i.p.* injection of each organic acid (10 $\mu\text{mol/g}$ body weight) to 7- and 30-day-old (young) rats. We barely detected both organic acids in the striatum of 30-day-old rats (concentrations around 20 μM). However, the same dose of these metabolites injected *i.p.* in 7-day-old rats gave rise to high striatum accumulation

of these compounds (HMG: 348 μM ; MGA; 480 μM), implying that the immature BBB is much more permeable to these organic acids.

3.2. HMG and MGA intrastratial administration induces lipid and protein oxidative damage in rat striatum

Fig. 1A shows that HMG and MGA [$F_{(2,13)} = 12.78$; $P < 0.001$] significantly increased MDA levels in rat striatum, indicating that these organic acids induce lipid oxidative damage.

We also evaluated the effect of intrastratial administration of HMG and MGA on carbonyl formation, a product of protein oxidation, in rat striatum. We observed that HMG and MGA [$F_{(2,10)} = 8.13$; $P < 0.01$] significantly increased carbonyl formation (Fig. 1B), indicating that both organic acids provoke protein oxidative damage.

3.3. HMG and MGA intrastratial administration increases reactive species production in rat striatum

We assessed whether reactive species were involved in HMG and MGA-induced pro-oxidant effects by investigating the influence of these organic acids on DCFH oxidation and nitrate plus nitrite production. We verified that HMG and MGA significantly increased DCFH oxidation [$F_{(2,13)} = 16.4$; $P < 0.001$] (Fig. 1C), whereas only HMG increased nitrate and nitrite production [$F_{(2,11)} = 2.97$; $P < 0.05$] (Fig. 1D).

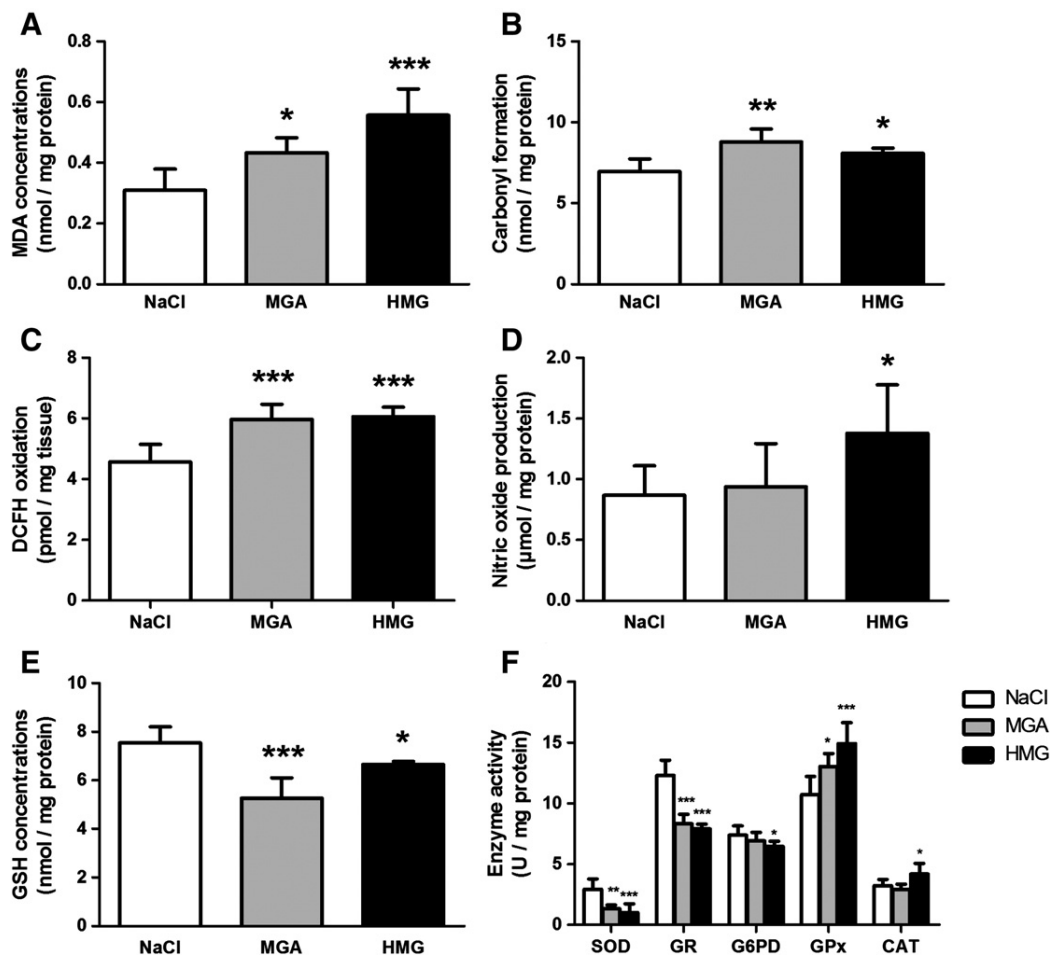


Fig. 1. Effect of intrastratial administration of 3-hydroxy-3-methylglutaric acid (HMG, 4 μmol) or 3-methylglutaric acid (MGA, 4 μmol) on malondialdehyde (MDA) levels (A), carbonyl formation (B), 2',7'-dichlorofluorescein (DCFH) oxidation (C), nitric oxide production (D), GSH concentrations (E), and the activities of superoxide dismutase (SOD), glutathione reductase (GR), glucose-6-phosphate dehydrogenase (G6PD), glutathione peroxidase (GPx) and catalase (CAT) (F) in rat striatum 30 min after injection. Data are expressed as mean \pm SD of 5 to 6 independent experiments (animals) performed in triplicate. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, compared to rats that received intrastratial NaCl injection (Duncan multiple range test).

3.4. HMG and MGA intrastratial administration reduces GSH concentrations in rat striatum

The next step of this work was to investigate the effects of HMG and MGA on GSH concentrations. Fig. 1E depicts that HMG and MGA [$F_{(2,15)} = 20.29$; $P < 0.001$] decreased GSH concentrations in rat striatum, indicating a decrease of striatal non-enzymatic antioxidant defenses.

3.5. HMG and MGA intrastratial administration modulates antioxidant enzyme activities in rat striatum

We then determined the effects of intrastratial administration of HMG and MGA on the activities of the antioxidant enzymes SOD, GR, G6PD, GPx and CAT. It can be observed in Fig. 1F that HMG and MGA significantly decreased SOD [$F_{(2,15)} = 13.78$; $P < 0.001$] and GR [$F_{(2,15)} = 44.97$; $P < 0.001$] activities, whereas HMG decreased the activity of G6PD [$F_{(2,15)} = 2.88$; $P < 0.05$] in rat striatum. On the other hand, GPx activity was significantly increased by HMG and MGA

intrastratial administration [$F_{(2,15)} = 12.58$; $P < 0.01$], whereas HMG increased CAT activity [$F_{(2,15)} = 6.44$; $P < 0.05$] (Fig. 1F).

3.6. Antioxidants and MK-801 prevented HMG-induced alterations of the parameters of oxidative stress

We also investigated the effects of i.p. pre-treatment of MEL (Fig. 2), NAC (Fig. 3) and the combination of vitamin E plus vitamin C (Fig. 4) on HMG-induced lipid oxidative damage and alterations of the antioxidant defenses in rat striatum. The figures demonstrate that these antioxidants fully prevented or attenuated the effects of HMG on MDA levels (Figs. 2A, 3A and 4A), GSH concentrations (Figs. 2B, 3B and 4B) and the activities of SOD, GR, G6PD, GPx and CAT (Figs. 2C, 3C and 4C).

We then evaluated the influence of a single intrastratial injection of the non-competitive NMDA receptor antagonist MK-801 and of the nitric oxide synthase (NOS) inhibitor L-NAME on the effects elicited by HMG (Fig. 5). We observed that MK-801 fully prevented lipid peroxidation induced by HMG (Fig. 5A), whereas L-NAME totally prevented and MK-801 attenuated GSH decrease caused by HMG (Fig. 5B).

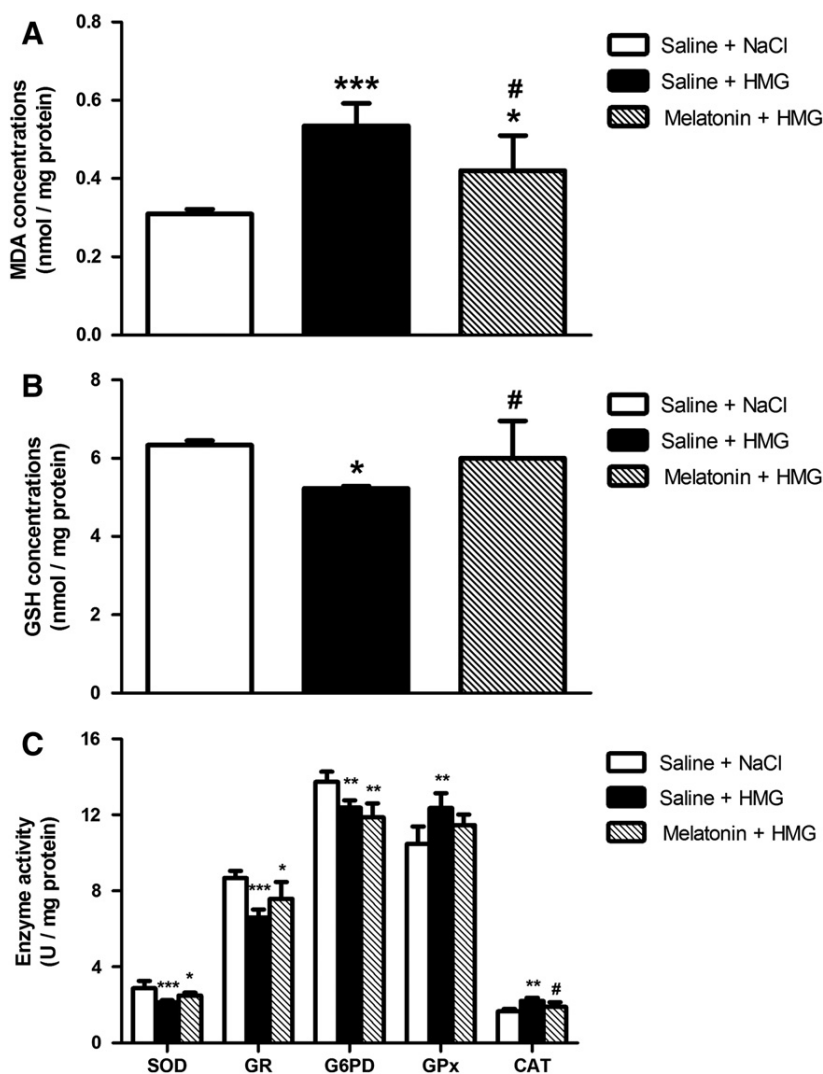


Fig. 2. Effect of melatonin (MEL; 100 mg/kg) on 3-hydroxy-3-methylglutaric acid (HMG, 4 μ mol)-induced alterations on malondialdehyde (MDA) levels (A), glutathione (GSH) concentrations (B) and activities of antioxidant enzymes (C). Values are means \pm standard deviation of five to six independent experiments (animals) performed in triplicate. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, compared to control (NaCl-injected rats); # $P < 0.05$, compared to HMG (Duncan multiple range test).

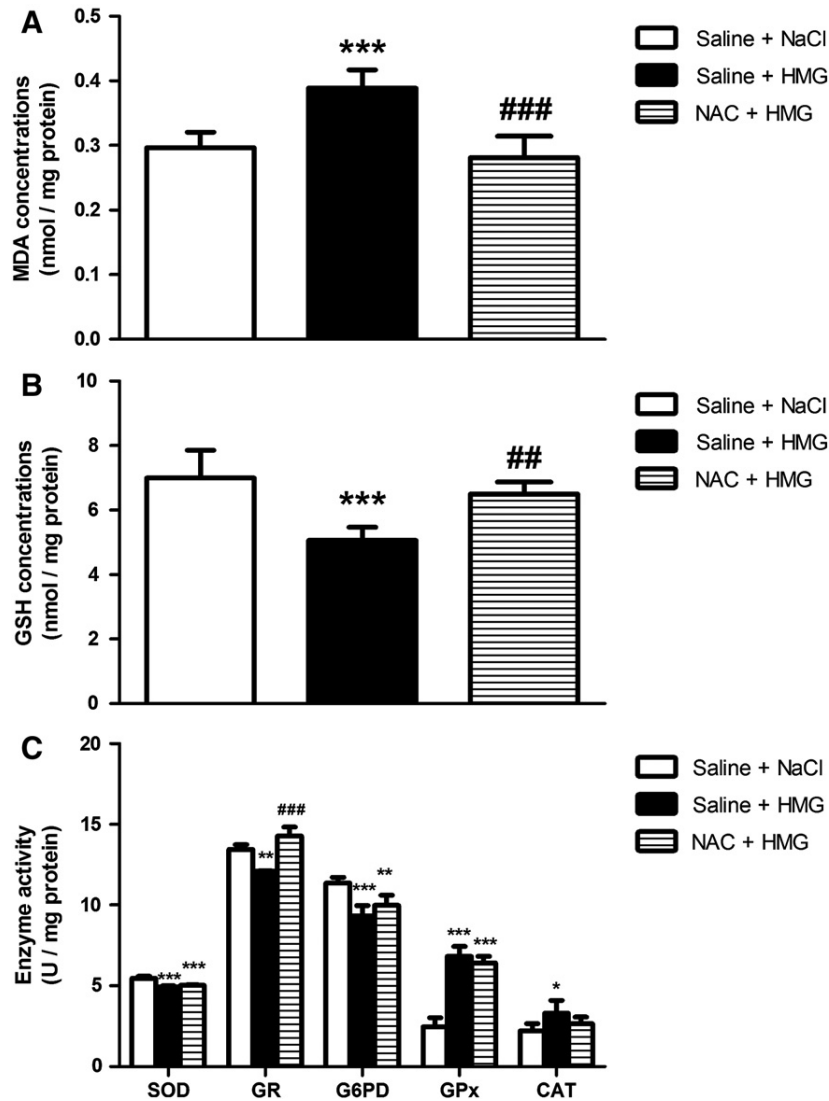


Fig. 3. Effect of N-acetylcysteine (NAC; 150 mg/kg) on 3-hydroxy-3-methylglutaric acid (HMG, 4 μ mol)-induced alterations on malondialdehyde (MDA) levels (A), glutathione (GSH) concentrations (B) and activities of antioxidant enzymes (C). Values are means \pm standard deviation of five to six independent experiments (animals) performed in triplicate. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, compared to control (NaCl-injected rats); ## $p < 0.01$, ### $p < 0.001$, compared to HMG (Duncan multiple range test).

Regarding the enzymatic antioxidant defenses, L-NAME and MK-801 prevented the HMG-elicited decrease of SOD activity (Fig. 5C), whereas MK-801 avoided the decrease of the activity of GR and L-NAME barred the decrease of G6PD provoked by HMG (Fig. 5C).

4. Discussion

The mechanisms of brain injury in HL deficiency are poorly known, although it is conceivable that the hypoglycemia and/or hyperammonemia occurring during crises may be involved in the neurological dysfunction presented by the patients [2,7,35]. It could be also presumed that chronic accumulation of HMG and MGA may represent an important pathomechanism contributing to the cerebral symptoms and abnormalities observed in these patients, particularly during episodes of metabolic decompensation, in which the concentrations of these metabolites dramatically increase. Therefore, it seems relevant to investigate the *in vivo* role of the major metabolites accumulating in HL deficiency on the cell redox homeostasis in the brain in order to clarify the relationship between the clinical features and the biochemical abnormalities in this disorder.

In the present study we evaluated the effects of an acute intrastriatal administration of HMG and MGA on important parameters of oxidative stress in rat striatum, a cerebral structure mainly altered in the brain of the affected patients [10–12]. We found that intrastriatal administration of HMG and MGA induced lipid peroxidation in striatum, corroborating our previous *in vitro* findings showing that both organic acids increased TBA-RS levels in rat brain [18]. Since TBA-RS measurement reflects the amount of MDA generated, a final product formed by the oxidation of unsaturated fatty acids of lipid constituents of the tissues, these results indicate that HMG and MGA induced lipid oxidative damage *in vivo*. We also showed that the antioxidants MEL, NAC, the combination of vitamins C and E prevented or attenuated MDA concentrations increase elicited by HMG, implying that this effect was secondary to elevated reactive species generation, mainly hydroxyl and peroxy radicals, which are scavenged by these antioxidants [36–38].

HMG and MGA also increased carbonyl formation in the striatum. In this context, it should be noted that carbonyl groups are generated in proteins by the oxidation of amino acid side chain residues (Pro, Arg, Lys, and Thr) mediated by reactive species [39]. We cannot also exclude the possibility that aldehydes resulting from lipid peroxidation, which were increased by HMG, may have secondarily induced carbonyl

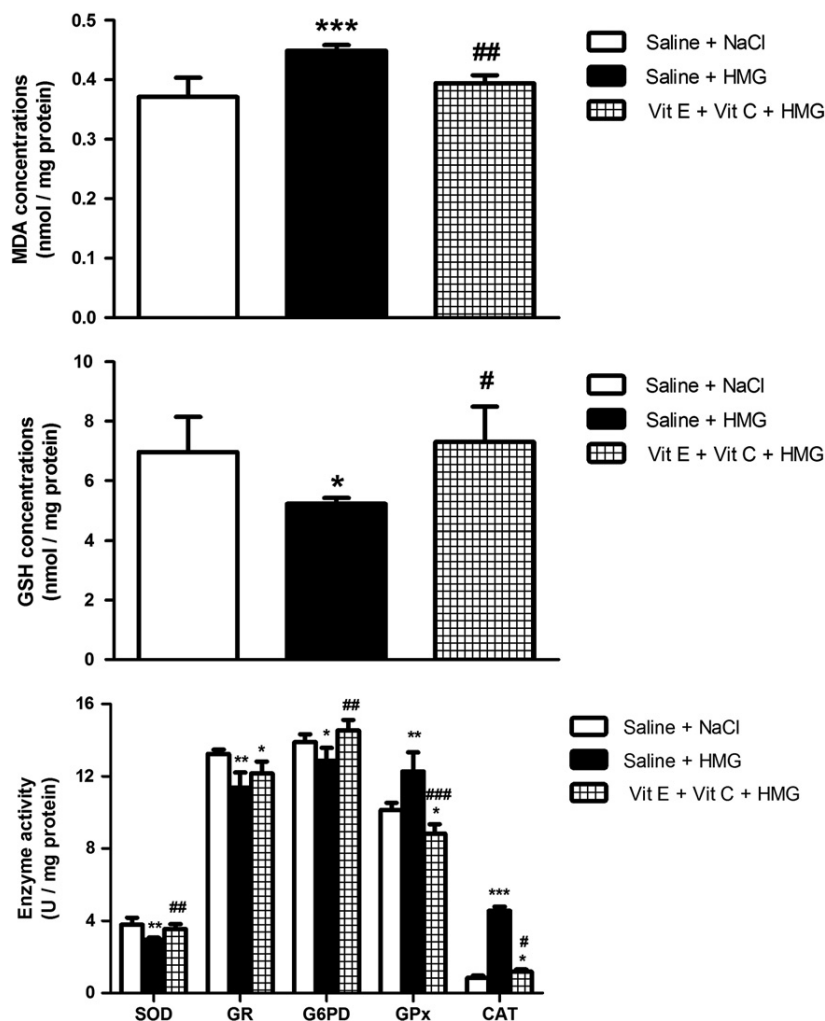


Fig. 4. Effect of the combination of α -tocopherol (vitamin E, Vit E; 40 mg/kg) and ascorbic acid (vitamin C, Vit C; 100 mg/kg) on 3-hydroxy-3-methylglutaric acid (HMG, 4 μ mol)-induced alterations on malondialdehyde (MDA) levels (A), glutathione (GSH) concentrations (B) and activities of antioxidant enzymes (C). Values are means \pm standard deviation of five to six independent experiments (animals) performed in triplicate. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, compared to control (NaCl-injected rats); # $P < 0.05$, ## $P < 0.01$, ### $P < 0.001$ compared to HMG (Duncan multiple range test).

generation [39]. Since carbonyl group generation is currently used as a marker of free radical-mediated protein oxidation [40], it is presumed that these organic acids provoked protein oxidative damage in the striatum.

We also verified that HMG and MGA increased DCFH oxidation, suggesting the involvement of ROS in HMG and MGA pro-oxidant effects [41,42]. HMG also increased nitric oxide production, as determined by the augmented formation of nitrates and nitrites provoked by this metabolite. These data, allied to our findings showing that NAC, MEL, the combination of vitamins C and E and L-NAME (NOS inhibitor) prevented lipid peroxidation and also the alterations of antioxidant defenses elicited by HMG, reinforce the presumption that the oxidative damage was due to increased generation of reactive oxygen and nitrogen species. On the other hand, it is possible that the increase of reactive nitrogen species (RNS) induced by HMG was partly due to overstimulation of NMDA receptors since MK-801, the antagonist of these receptors, also prevented HMG-induced pro-oxidant effects [43].

Regarding the antioxidant system, we found that HMG and MGA significantly reduced GSH concentrations, and this is in line with our previous *in vitro* findings [18]. Considering that endogenous GSH is the main naturally-occurring antioxidant in the brain and is used to evaluate the capacity of a tissue to prevent the damage associated to free radical processes [38], it is presumed that the rat

striatum non-enzymatic antioxidant defenses were compromised by HMG and MGA *in vivo*. Moreover, since GSH is considered an important defense against lipid oxidative damage scavenging reactive species responsible for the initiation of this process, it is feasible that lipid peroxidation elicited by HMG and MGA could be secondary to GSH reduction. It is emphasized that GSH reduction could be due to the increase of reactive species elicited by HMG and MGA, which may have consumed this important brain antioxidant, reducing its final concentrations.

HMG and MGA also significantly decreased SOD and GR activities and increased GPx activity in rat striatum. In addition, HMG caused a significant increase of CAT activity and a decrease of G6PD activity. Although we cannot at present explain the mechanisms by which SOD activity was reduced, it has been suggested that a decrease in the activities of antioxidant enzymes, such as SOD, may be due to ROS causing a site-specific amino acid modification [43,44]. We should emphasize that a reduction of SOD activity may lead to superoxide anions excess that usually generates other forms of carbon-, nitrogen- and oxygen-centered radicals that could contribute to the lipid and protein oxidative damage induced by HMG and MGA in the brain. Furthermore, the increase of GPx activity, DCFH oxidation and lipid peroxidation caused by HMG and MGA administration suggest that hydrogen peroxide and lipid peroxide are probably involved in HMG- and MGA-induced

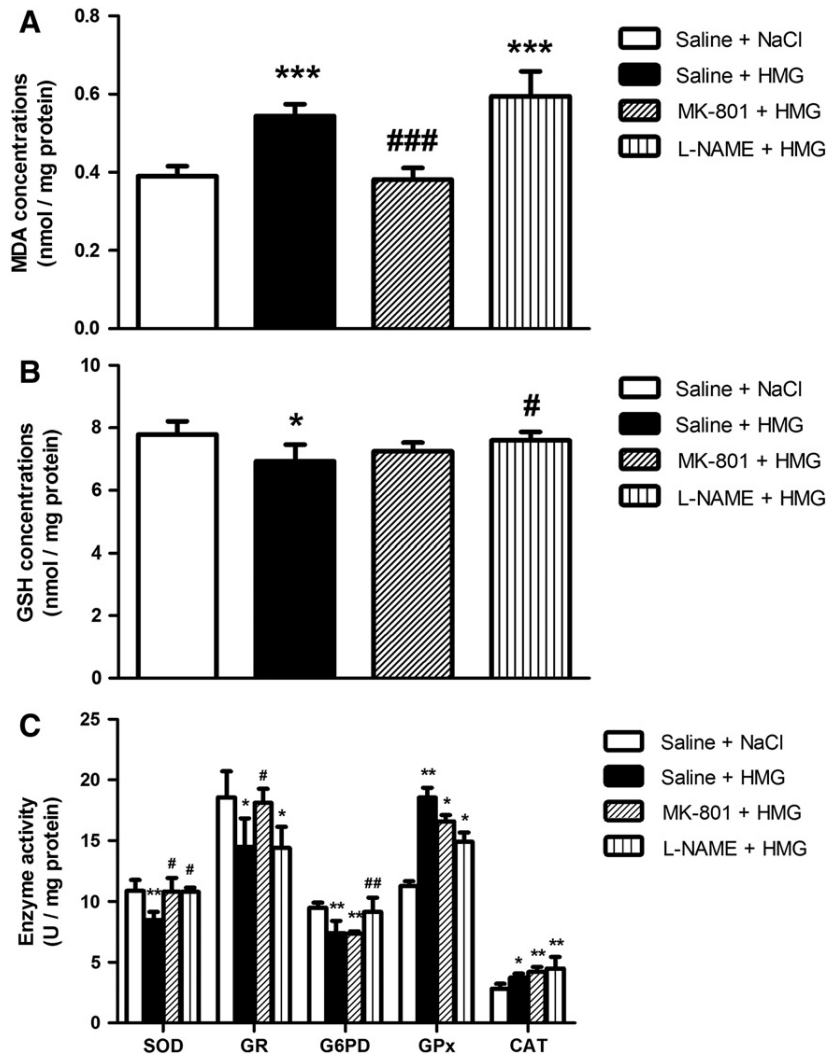


Fig. 5. Effect of dizolciline maleate (MK-801; 6 nmol) or N^o-nitro-L-arginine methyl ester (L-NAME; 0.2 nmol) on 3-hydroxy-3-methylglutaric acid (HMG, 4 μ mol)-induced alterations on malondialdehyde (MDA) levels (A), glutathione (GSH) concentrations (B) and activities of antioxidant enzymes (C). Values are means \pm standard deviation of five to six independent experiments (animals) performed in triplicate. * P < 0.05, ** P < 0.01, *** P < 0.001, compared to control (NaCl-injected rats); # P < 0.05, ## P < 0.01, ### P < 0.001 compared to HMG (Duncan multiple range test).

oxidative damage in rat striatum. As regards the increased CAT activity, induction of the expression of this antioxidant enzyme at the gene level might have taken place as a compensatory mechanism in response to increased formation of hydrogen peroxide [45–47] induced by HMG. Finally, it is feasible that HMG-induced diminution of G6PD activity may have contributed to the decreased GR activity due to the lack of NADPH for GSH recycling through this enzyme.

We also found that the non-competitive NMDA receptor antagonist MK-801 prevented the lipid oxidation and the alterations of antioxidant defenses caused by HMG. These data highlight a role for NMDA receptor in these effects. The fact that NMDA receptors may be involved in HMG-induced effects is not surprising because of the structural similarity between HMG and glutamate. Therefore, it may be of interest to study whether these organic acids and especially HMG interact with NMDA receptors. On the other hand, we cannot exclude the possibility that induction of NOS activity leading to generation of RNS was at least partially due to overstimulation of NMDA glutamate receptors by this organic acid [38,43,48].

Since oxidative stress results from an imbalance between the total antioxidant defenses and the reactive species generated in a tissue,

our present data strongly indicate that HMG and MGA induce oxidative stress *in vivo* in rat striatum. It must be emphasized here that ROS are capable to oxidize different molecules, including lipids, proteins, sugars and DNA, representing a key event in the pathogenic cascade leading to necrotic or apoptotic cell death [49–51]. Furthermore, oxidative stress is a very deleterious condition especially to the brain compared with other tissues [38]. This is supported by the fact that the brain has a high rate of oxidative metabolism coupled to ROS production, lower activity of antioxidant enzymes, reduced content of non-enzymatic antioxidants and higher peroxidation potential because of its high content of polyunsaturated fatty acids [38].

Currently, we cannot determine the pathophysiological relevance of the present data since to our knowledge brain concentrations of HMG and MGA were not yet established in HL deficiency. However, it should be stressed that the significant alterations of the biochemical parameters were obtained at micromolar concentrations in our present *in vivo* model. On the other hand, we found that HMG and MGA were shown to rapidly cross the BBB in 7-day-old rats in the striatum, although only traces of these compounds were detected in this cerebral structure from older animals. These data indicate that the

immature barrier is much more permeable to these organic acids and that treatment for HL deficient patients should be better controlled in early stages of development.

In summary, our present data clearly demonstrate that HMG and MGA disturb cell redox homeostasis in the striatum. We cannot attribute our findings to derivatives of HMG and MGA since their concentrations did not change when these organic acids were incubated with brain slices or homogenates. This is in accordance with recent findings showing that HL is not expressed in the mitochondrial fraction of the brain [35], reinforcing our results and indicating that HMG and MGA are not produced *in situ* in the brain.

In conclusion, to the best of our knowledge this is the first report showing that HMG and MGA induce *in vivo* oxidative stress in the striatum at least partially mediated by NMDA receptors. The present *in vivo* data allied to previous *in vitro* studies [16–19] indicate that this pathomechanism may potentially contribute to the neurologic manifestations found in HL deficiency. In this regard, antioxidants directed towards the mitochondria or specific antagonists of NMDA glutamate subunits focused to selectively block the toxic effects of NMDA stimulation without interfering with its physiological function in neurotransmission may possibly serve in the future as adjuvant therapies in HL deficiency, especially during crises [52,53]. In this context, it was recently shown that the NMDA receptor antagonist ketamine significantly reduced seizures and ameliorated the clinical status of a patient with nonketotic hyperglycinemia [53]. It is also possible that memantine, an open-ion channel antagonist with rapid turnover, that was shown to protect against excitotoxicity in Alzheimer's disease patients, could be used to prevent brain abnormalities in this disorder [54].

Acknowledgments

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Capítulo 2

Induction of a proinflammatory response in cortical astrocytes by the major metabolites accumulating in HMG-CoA lyase deficiency: the role of ERK signaling pathway in cytokine release

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Induction of a proinflammatory response in cortical astrocytes by the major metabolites accumulating in HMG-CoA lyase deficiency: the role of ERK signaling pathway in cytokine release

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Keywords:	3-Hydroxy-3-methylglutaric aciduria; oxidative stress; proinflammatory response; astrocytes; cytokine production; ERK pathway.
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Abstract:	<p>3-Hydroxy-3-methylglutaric aciduria (HMGA) is an inherited metabolic disorder caused by 3-hydroxy-3-methylglutaryl-CoA lyase deficiency. It is biochemically characterized by predominant tissue accumulation and high urinary excretion of 3-hydroxy-3-methylglutarate (HMG) and 3-methylglutarate (MGA), as well as 3-methylglutaconate and 3-hydroxyisovalerate. Affected patients commonly present acute symptoms during metabolic decompensation, including vomiting, seizures and lethargy/coma accompanied by metabolic acidosis and hypoketotic hypoglycaemia. Although neurological manifestations are common, the pathogenesis of brain injury in this disease is poorly known. Astrocytes are important for neuronal protection and are susceptible to damage by neurotoxins. In the present study, we investigated the effects of HMG and MGA, the major metabolites accumulating in HMGA, on important parameters of redox homeostasis and cytokine production in cortical cultured astrocytes. The role of the metabolites on astrocyte mitochondrial function (thiazolyl blue tetrazolium bromide - MTT reduction) and viability (propidium iodide incorporation) was also studied. Both organic acids decreased astrocytic mitochondrial function without altering cell viability, and the concentrations of reduced glutathione. In contrast, they increased reactive species formation (2'-7'-dichlorofluorescein diacetate - DCFHDA oxidation). They also provoked a significant increase of IL-1, IL-6 and TNFα release through the ERK signaling pathway. Taken together, the data indicate that the principal compounds accumulating in HMGA induce a proinflammatory response in cultured astrocytes that may possibly be involved in the neuropathology of this disease.</p>

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January 8, 2015

Dr. Nicolas G. Bazan,
Editor-in-Chief,
Molecular Neurobiology

Dear Dr. Bazan,

I am sending you our manuscript entitled “**Induction of a proinflammatory response in cortical astrocytes by the major metabolites accumulating in HMG-CoA lyase deficiency: the role of ERK signaling pathway in cytokine ‘release’**”, which we would like to submit for publication in Molecular Neurobiology.

This study was performed in strict accordance with the Guide for the Care and Use of Laboratory Animals (Eighth edition, 2011) and was approved by the Ethical Committee of the Federal University of Rio Grande do Sul, Porto Alegre, Brazil. All efforts were also made to use the minimal number of animals necessary to produce reliable scientific data. The manuscript does not contain clinical studies or patient data.

I also inform you that all authors have contributed significantly to the manuscript and agree with the submission of the paper at its present version. Moreover, the whole manuscript, or parts of it, will not be submitted elsewhere for publication. Finally, there is no conflict of interest in the conduct and reporting of research.

This manuscript presents novel findings showing that 3-hydroxy-3-methylglutarate and 3-methylglutarate that accumulate in 3-hydroxy-3-methylglutaric aciduria induce a proinflammatory response in cultured astrocytes that may possibly be involved in the neuropathology of this disease.

I look forward to hearing from you in the near future.

Yours sincerely,

M. Wajner, MD, PhD

Induction of a proinflammatory response in cortical astrocytes by the major metabolites accumulating in HMG-CoA lyase deficiency: the role of ERK signaling pathway in cytokine release

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Running title: The major metabolites accumulating in HMG-CoA lyase deficiency induce proinflammatory response in astrocytes

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Abstract

3-Hydroxy-3-methylglutaric aciduria (HMGA) is an inherited metabolic disorder caused by 3-hydroxy-3-methylglutaryl-CoA lyase deficiency. It is biochemically characterized by predominant tissue accumulation and high urinary excretion of 3-hydroxy-3-methylglutarate (HMG) and 3-methylglutarate (MGA), as well as 3-methylglutaconate and 3-hydroxyisovalerate. Affected patients commonly present acute symptoms during metabolic decompensation, including vomiting, seizures and lethargy/coma accompanied by metabolic acidosis and hypoketotic hypoglycaemia. Although neurological manifestations are common, the pathogenesis of brain injury in this disease is poorly known. Astrocytes are important for neuronal protection and are susceptible to damage by neurotoxins. In the present study, we investigated the effects of HMG and MGA, the major metabolites accumulating in HMGA, on important parameters of redox homeostasis and cytokine production in cortical cultured astrocytes. The role of the metabolites on astrocyte mitochondrial function (thiazolyl blue tetrazolium bromide – MTT reduction) and viability (propidium iodide incorporation) was also studied. Both organic acids decreased astrocytic mitochondrial function without altering cell viability, and the concentrations of reduced glutathione. In contrast, they increased reactive species formation (2',7'-dichlorofluorescein diacetate - DCFHDA oxidation). They also provoked a significant increase of IL-1 β , IL-6 and TNF α release through the ERK signaling pathway. Taken together, the data indicate that the principal compounds accumulating in HMGA induce a proinflammatory response in cultured astrocytes that may possibly be involved in the neuropathology of this disease.

Kew words: 3-Hydroxy-3-methylglutaric aciduria; oxidative stress; proinflammatory response; astrocytes; cytokine production; ERK pathway.

Introduction

3-Hydroxy-3-methylglutaryl-CoA lyase (EC 4.1.3.4) catalyses the cleavage of 3-hydroxy-3-methylglutaryl-CoA to form acetyl-CoA and acetoacetate. This reaction is a key step in ketogenesis and is also the last step in the leucine catabolic pathway [1-2]. Deficiency in this enzyme activity is the cause of the autosomal recessive disorder called 3-hydroxy-3-methylglutaric aciduria (HMGA, MIM 246450). Biochemically, the disease is characterized by predominant tissue accumulation and high urinary excretion of 3-hydroxy-3-methylglutarate (HMG) and 3-methylglutarate (MGA), as well as 3-methylglutaconate, 3-hydroxyisovalerate and 3-methylcrotonylglycine to a lesser degree [3-8].

Clinical presentation usually occurs in the first year of life during fasting or intercurrent illnesses, mainly infections. Acute symptoms include vomiting, seizures and lethargy/coma, accompanied by metabolic acidosis, hypoketotic hypoglycemia and mild hyperammonemia [1, 10-11, 19]. Chronically progressive brain injury and mild hepatic alterations during crises are frequently present in this disorder. Pancreas and heart can also be involved in some patients [8, 12-17]. Magnetic resonance neuroimaging frequently shows multiple and marked coalescent lesions in periventricular subcortical white matter and arcuate fibers, most prominent in frontal or periauricular regions, apart from alterations in the basal ganglia attesting the significance of the neurological involvement [17-20].

Astrocytes, the more versatile cells in the mammalian brain, provide neuronal support and stability, being therefore important modulators of normal brain functioning [21-22]. Moreover, these cells have crucial role in neurodegenerative disorders, playing decisive roles in damaging and stress responses by synthesizing cytokines and chemokines [23]. This process usually lead to reactive astrogliosis and may be beneficial or deleterious because reactive astrocytes may underlie pro- or anti-inflammatory effects, being the proinflammatory response involved in pathogenesis of neurodegenerative processes [24]. Additionally, astrocytes modulate the biosynthesis and release of antioxidant defenses like glutathione (GSH) in the CNS [25].

The pathophysiology of 3-hydroxy-3-methylglutaryl-CoA lyase (HMG-CoA lyase) deficiency is only partially understood. It may be explained by energy deficit (hypoketotic hypoglycaemia), reduction of fatty acid oxidation due to carnitine deficiency or alternatively due to the intracellular accumulation of ammonia and/or the toxic organic acids accumulating secondary to the leucine catabolic pathway blockage. Previous experimental evidence supports neurotoxic effects of the major compounds that accumulate intramitochondrially in HMGA. Oxidative stress has been proposed as one of the primary causes of neurodegeneration, not only through the structural and functional alterations that reactive oxygen species (ROS) produce to cell biomolecules, but also because they are potential mediators of cell death by either necrosis or apoptosis [26]. In this context, previous *in vitro* and *ex vivo* studies demonstrated that HMG and MGA induce lipid and protein oxidative damage and reduce the non-enzymatic antioxidant defenses in cerebral cortex, striatum and liver of young rats [27-30]. It was also shown that MGA reduces mitochondrial redox potential (mitochondrial dysfunction) and inhibits Na⁺,K⁺-ATPase activity in synaptosomes prepared from rat brain [31].

However, to the best of our knowledge, nothing has been reported on the influence of these metabolites on astrocytes to establish whether these cells are vulnerable to the major metabolites accumulating in HMGA. Since cultured astrocytes have made important contributions to the understanding of

neurodegenerative pathologies because of their critical role on the neuronal-astrocytic interactions [32-33], in the present work we investigated the effects of HMG and MGA on oxidative and inflammatory parameters in cortical primary astrocyte cell cultures. To further clarify the pathogenesis of the brain damage in HMGA, we explored the putative mechanisms involved in inflammatory response induced by these organic acids in astrocytes

Material and Methods

1. Materials

Dulbecco's Modified Eagle's Medium (DMEM) and other materials for cell cultures were purchased from Gibco. DNase and polyclonal anti-GFAP was purchased from Dako, whereas thiazolyl blue tetrazolium bromide (MTT), propidium iodide (Pi) and ELISA for NF κ B, TNF- α , IL-1 β and IL-6 were purchased from Invitrogen, PeproTech and eBioscience, respectively. All other chemicals were obtained from Sigma–Aldrich.

2. Animals

Male Wistar rats (1 to 3 days old) were obtained from our breeding colony (Department of Biochemistry, UFRGS, Brazil) and maintained in a controlled environment (12 h light/12 h dark cycle; 22 ± 1 °C; ad libitum access to food and water). The experimental protocol was approved by the Ethics Committee for animal research of the Federal University of Rio Grande do Sul, Porto Alegre, Brazil and followed the “Principles of Laboratory Animal Care (NIH publication 85-23, revised 1985). All efforts were made to minimize the number of animals used and their suffering.

3. Primary astrocyte cultures

Animals had their cerebral cortex aseptically removed from cerebral hemispheres. All meninges were removed. During the dissection, the cerebral cortex was kept in HBSS (Hank's Balanced Salt Solution) containing 0.05% trypsin and 0.003% DNase at 37 °C for 8 min. The tissue was then mechanically dissociated for 7 min using a Pasteur pipette and centrifuged at 100 g for 5 min. The cells were resuspended in HBSS containing DNase (0.003%) and left for decantation during 20 min. The supernatant was collected and centrifuged for 400 g for 7 min. The cells from the supernatant were resuspended in DMEM (10% fetal bovine serum (FBS), 15 mM HEPES, 14.3 mM NaHCO₃, 1% fungizone and 0.04% gentamicin) and plated in 24-well plates pre-coated with poly-L-lysine at a density of $3\text{--}5 \times 10^5$ cells/cm². The cells were cultured at 37 °C in a 95% air / 5% CO₂ incubator. The first medium exchange occurred 24 h after obtaining the culture.

4. Organic acids treatment

After cells reached confluence, the culture medium was removed and the cells were incubated in the absence or presence of HMG or MGA (0.2 to 5 mM) for 24 h at 37 °C in a 95% air / 5% CO₂ incubator in DMEM with 1% FBS.

5. Intracellular ROS levels

Intracellular ROS production was assessed as previously described [34] using the nonfluorescent cell permeating compound 2',7'-dichlorofluorescein diacetate (DCFHDA) on cells under basal conditions or treated with HMG or MGA. DCFHDA is hydrolyzed by intracellular esterases to dichlorofluorescein (DCFH), which is trapped within the cell. This nonfluorescent molecule is then oxidized to fluorescent dichlorofluorescein (DCF) by the action of cellular oxidants. Astrocytes were treated with DCFHDA (10 mM) for 30 min at 37°C. Following DCFHDA exposure, the cells were scraped into PBS supplemented with 0.2% Triton X-100. Fluorescence was measured using a plate reader (Spectra Max M5, Molecular Devices) at excitation and emission wavelengths of 485 nm and 520 nm, respectively.

6. Mitochondrial function

Mitochondrial function was accessed by thiazolyl blue tetrazolium bromide (MTT) reduction assay, in which formazan is produced (activity of mitochondrial dehydrogenases). Briefly, 0.05 mg / ml of MTT was added to the incubation medium. After 3 hours of incubation, the medium from each well was gently removed by aspiration and dimethylsulfoxide (DMSO) was added to each well followed by incubation and shaking for 5 min. The formazan product generated during the incubation was solubilized in DMSO and measured at 560 and 650 nm. Only functional cells are able to reduce MTT.

7. Reduced glutathione (GSH) content

GSH levels were assessed as previously described [35]. Astrocyte homogenates (50 mg) were diluted in 100 mM sodium phosphate buffer (pH 8.0) containing 5 mM EDTA, and the protein was precipitated with 1.7% meta-phosphoric acid. The supernatant was assayed with o-phthalaldehyde (1 mg/ml methanol) at room temperature for 15 min. Fluorescence was measured using excitation and emission wavelengths of 350 and 420 nm, respectively. A calibration curve was performed with standard GSH solutions (0–500 mM). GSH concentrations were calculated as nmol/mg protein.

8. IL-1 β , IL-6 and TNF α measurement

IL-1 β and IL-6 was carried out in an extracellular medium, using a rat IL-1 β or IL-6 ELISA from eBioscience (USA), whereas the TNF α assay was carried out in the same medium, using a rat TNF- α ELISA from Peprotech (USA). The results are expressed as the percentage of control levels. To test whether HO1 and ERK signaling pathways were involved in the production of these classical cytokines, we used their inhibitors zinc proporphyrin IX (ZnPP IX, 10 μ M) and PD98059 (5 μ M), respectively, for 0.5 h before the treatment with HMG and MGA.

9. Measurement of NF κ B levels

The levels of NF κ B p65 in the nuclear fraction, which had been isolated from lysed cells by centrifugation, were measured using an ELISA commercial kit from Invitrogen (USA). The results are expressed as percentages relative to the control levels.

10. Measurement of astrocyte viability

Cell viability was measured by propidium iodide cell incorporation using phase contrast optics. Membrane integrity was assessed by fluorescent image analysis (Nikon inverted microscope using a TE-FM Epi-Fluorescence accessory) of propidium iodide (Pi) uptake (7.5 mM) [36] at 37 °C in an atmosphere of 5% CO₂/ 95% air in DMEM supplemented with 5% FBS.

11. GFAP quantification by western blot analysis

To characterize the astrocyte cultures we measured the glial fibrillar astrocyte protein (GFAP) levels by western blot in the presence or absence of 5 mM HMG or 5 mM MGA. Cells were solubilized with a lysis solution containing 4 % sodium dodecyl sulfate (SDS), 2 mM EDTA and 5 mM Tris-HCl at pH 6.8. Equal amounts of protein from each sample were boiled in a sample buffer (62.5 mM Tris-HCl, pH 6.8, 2 % (w/v) SDS, 5 % β-mercaptoethanol, 10 % (v/v) glycerol, 0.002 % (w/v) bromophenol blue) and submitted to electrophoresis in a 10 % (w/v) SDS-polyacrilamide gel. The separated proteins were transferred to a nitrocellulose membrane. Equal loading of each sample was confirmed with Ponceau S staining (Sigma). The membrane was incubated with polyclonal antibodies targeting GFAP (1:3000). β-actin was used as a loading control. After incubating overnight with the primary antibody at 22 °C, the membrane was washed and incubated with peroxidase-conjugated anti-rabbit immunoglobulin (IgG) at a dilution of 1 : 1000 for 1h. The chemiluminescence signal was detected using enhanced chemiluminescence; the resulting films were scanned and the bands were quantified using the Scion Image software (Scion Corp., Frederick, MD, USA).

12. Protein assay

Protein content was measured using Lowry's method, with bovine serum albumin as a standard [37].

13. Statistical analyses

Differences among groups were statistically analyzed using one-way analysis of variance (ANOVA), followed by the Duncan's multiple range test and in some experiments the Student *t* test for unpaired samples was used. Only significant *F* and *t* values were shown. All analyses were performed using the Statistical Package for Social Sciences software, version 18.0 (SPSS Inc., Chicago, IL, USA).

Results

1. *3-Hydroxy-3-methylglutaric (HMG) and 3-methylglutaric (MGA) acids decrease mitochondrial function in cultured astrocytes*

We first evaluated the mitochondrial function by the MTT reduction assay. Cultured astrocytes exposed to HMG and MGA at 5 mM concentration for 24 h significantly decreased formazan formation (HMG: Fig. 1A [$F_{(2,6)} = 22.6$; $P < 0.001$] and MGA: Fig. 1B [$F_{(2,6)} = 60.1$; $P < 0.001$], respectively). However, astrocyte viability, measured by propidium iodide incorporation, was not significantly changed by HMG and MGA for 24 h (results not shown).

2. *3-Hydroxy-3-methylglutaric (HMG) and 3-methylglutaric (MGA) acids increase reactive oxygen species (ROS) production and reduce the antioxidant defenses in cultured astrocytes*

Next, we investigated whether HMG and MGA could alter ROS production in primary astrocyte cultures from neonatal rat cerebral cortex. Fig. 2A and B show that HMG [$F_{(4,37)} = 40.9$; $P < 0.001$] and MGA [$F_{(4,37)} = 27.5$; $P < 0.001$] significantly increased DCFH oxidation in cultured astrocytes in a dose dependent manner (HMG: $\beta = 0.88$; $P < 0.001$; MGA: $\beta = 0.81$; $P < 0.001$), indicating that these organic acids induce ROS production.

We also evaluated the effect of HMG and MGA on GSH levels, the major naturally occurring brain antioxidant. Fig. 3A and B show that HMG [$F_{(4,22)} = 16.1$; $P < 0.001$] and MGA [$F_{(4,22)} = 10.8$; $P < 0.001$] significantly decreased GSH concentrations in astrocytes in a dose dependent manner (HMG: $\beta = -0.62$; $P < 0.01$; MGA: $\beta = -0.66$; $P < 0.001$), indicating a decrease of non-enzymatic antioxidant defenses.

3. *3-Hydroxy-3-methylglutaric (HMG) and 3-methylglutaric (MGA) acids induce cytokine production in cultured astrocytes*

We then investigated the inflammatory response in cultured astrocytes exposed to HMG or MGA by measuring interleukin production in the extracellular medium. Our results show that both HMG and MGA induced significant increases of the levels of interleukin 1 β (IL-1 β , Fig. 4A [$F_{(2,9)} = 28.7$; $P < 0.001$] and Fig. 4B [$F_{(2,9)} = 29.5$; $P < 0.001$]), interleukin 6 (IL-6, Fig. 4C [$F_{(2,9)} = 35.3$; $P < 0.001$] and Fig. 4D [$F_{(2,9)} = 42.6$; $P < 0.001$]) and tumor necrosis factor α (TNF α , Fig. 5E [$F_{(2,9)} = 40.9$; $P < 0.001$] and Fig. 4F [$F_{(2,9)} = 10.4$; $P < 0.001$]) release, indicating a proinflammatory response caused by these organic acids. It can be also seen that all effects were dose dependent (IL-1 β : HMG: $\beta = 0.86$; $P < 0.001$; MGA: $\beta = 0.87$; $P < 0.001$; IL-6: HMG: $\beta = 0.77$; $P < 0.01$; MGA: $\beta = 0.92$; $P < 0.001$; TNF α : HMG: $\beta = 0.87$; $P < 0.001$; MGA: $\beta = 0.76$; $P < 0.01$). In a separate set of experiments, we observed no changes in cell integrity (evaluated by PI incorporation, data not shown), implying that the increased levels of cytokines most likely resulted from secretion of these proinflammatory factors.

4. *3-Hydroxy-3-methylglutaric (HMG) and 3-methylglutaric (MGA) acids do not activate NF κ B in cultured astrocytes*

We also observed that NF κ B levels were not altered by exposition of cultured astrocytes to HMG and MGA for 24 h (Fig. 5A and B).

5. *ERK signaling pathway mediates the 3-hydroxy-3-methylglutaric (HMG) and 3-methylglutaric (MGA)-induced proinflammatory cytokines release*

In an attempt to determine the mechanisms underlying the HMG and MGA-induced increase proinflammatory cytokines, we measured cytokine production (IL-1 β , IL-6 and TNF α) in the presence of ERK (PD98059) and HO1 (ZnPP IX) inhibitors. Table 1 shows that ERK inhibitor totally abolished the effects of HMG and MGA on IL-1 β and TNF α levels. However, HO1 inhibitor was not able to prevent the increase in cytokines release provoked by HMG and MGA (Table 2).

6. *3-Hydroxy-3-methylglutaric (HMG) and 3-methylglutaric (MGA) do not induce reactive gliosis or alter astrocyte morphology*

We finally measured the glial fibrillar astrocyte protein (GFAP) levels by western blot in cultured astrocytes exposed to 5 mM HMG or 5 mM MGA. No alterations in GFAP expression levels were detected, as compared to controls, indicating that these organic acids do not induce reactive gliosis (Fig. 6). In addition, no modification of cell morphology could be observed on microscope images (results not shown). These results demonstrated that HMG and MGA did not induce reactive gliosis neither changed cell morphology.

Discussion

Patients affected by HMG-CoA deficiency, also known as 3-hydroxy-3-methylglutaric aciduria (HMGA), accumulate predominantly HMG and MGA in their tissues and body fluids. The concentrations of these metabolites dramatically increase during acute metabolic crises, which are characterized by severe metabolic acidosis, hypoketotic hypoglycaemia, mild hyperammonaemia, vomiting, hypotonia, coma, and seizures resulting in permanent neurological damage [6, 11, 17, 20]. Although the neurological symptoms are common and cerebral MRI reveals marked lesions in periventricular and/or subcortical white matter, the mechanisms underlying the brain injury in this disease are poorly known. However, previous *in vitro* and *ex vivo* studies have shown that the major metabolites accumulating in this disorder disrupt mitochondrial dysfunction and redox homeostasis in rat brain [27-30].

Astrocytes modulate ionic homeostasis, energy metabolism, antioxidant and anti-inflammatory responses, among other important CNS functions [38-43]. Furthermore, since the utilization of primary astrocyte cultures have contributed to the understanding of the role of astrocytes in physiological and pathological conditions [33], in the present study we exposed astrocytes derived from cerebral cortex of neonatal rats to equimolar concentrations of HMG and MGA to mimic the biochemical phenotype of HMGA, and evaluated redox homeostasis, cytokine production, cell mitochondrial function and viability.

Initially we observed that HMG and MGA provoked a dose dependent increase of DCFH oxidation in cultured astrocytes, an useful probe for oxidative study. Since oxidation of DCFH occurs mainly through free reactive oxygen formation and is commonly used as a probe for the total ROS production in biological systems, it is concluded that these compounds elicited reactive species generation in an intact cell system (astrocytes), corroborating with previous findings obtained with subcellular fractions from total brain that includes all neural cells [27-31]. So, the first novel finding of the present investigation is the observation that astrocytes are involved in free radical production induced by the major metabolites accumulating in HMGA.

We also found that these organic acids markedly decreased GSH concentrations in cortical cultured astrocytes. In this context, astrocytes are considered to be a major source of GSH by synthesizing and secreting this antioxidant, thus helping the maintenance of the neuronal redox state [44-45]. GSH is the most abundant antioxidant in the brain, exerting a critical role to scavenge reactive oxygen species (ROS), protect protein sulfhydryl groups in the appropriate redox state, also regulating cell death and survival pathways [46].

Since adequate levels of antioxidants are essential to protect cells against oxidative damage and an imbalance in the pro-oxidant/antioxidant homeostasis induces oxidative stress [47], it is possible that the significant reduction of GSH induced by the metabolites accumulating in HMGA may cause loss of functionality of astrocytes under HMG and MGA exposure, as observed by the significant reduction of MTT. It is emphasized that the MTT assay is commonly used to assess mitochondrial functionality, as an indicator of cell energy and metabolic status, although it is also accepted as a marker of cell viability. Since propidium iodide incorporation was not changed in cultured astrocytes exposed to HMG and MGA under the same conditions, we presume that cell viability was preserved. Noteworthy, astrocytes have been demonstrated to be vulnerable to various toxins and in particular to MPP⁺ and this susceptibility seems to be dependent on a pro-oxidant condition induced by MPP⁺, leading astrocytes to lose functionality [48-49].

We cannot establish at the present whether the reduction of GSH levels occurred due to its oxidation by free radicals because of a high production of reactive species (increased DCFH oxidation) leading to GSH consumption, or whether the reduction of GSH levels led to an imbalance of reactive species formation leading to a pro-oxidant state. It is of note that depletion of GSH takes place during stress conditions giving rise to an inflammatory response with oxidative stress that plays a critical role in astrocytic inflammatory response [50]. In addition, glial GSH depletion induces neuroinflammation and neurotoxicity with significant augment of proinflammatory cytokines [51].

In this particular, another original contribution of our work was the demonstration of elevated levels of the cytokines IL-1 β , IL-6 and TNF α in cultured astrocytes exposed to HMG and MGA. TNF α and IL1- β are synthesized predominantly by microglia and astrocytes, acting in acute inflammatory responses and these cytokines are able to induce IL-6 synthesis. IL-6 is also produced in microglia, astrocytes and neurons and plays a pivotal role in a variety of CNS functions such as induction and modulation of astrocytes reactivation, pathological inflammation and neuroprotection [50-54]. Moreover, the elevation of cytokine levels probably reflects glial reactivity that, in combination with an increase in ROS production and a decrease in antioxidant content in the brain, has been reported as primary features in the pathogenesis and development of various human diseases and of neurodegeneration [55-56]. Another point to be emphasized is that in our present study astrocyte activation was observed without GFAP overexpression [57].

ERK pathway has been implicated in the regulation of glial inflammatory response following an insult/damage and is an upstream signal transduction of NF κ B [51, 58]. NF κ B is considered the major inflammatory mediator in the CNS, although its mechanism of activation and subsequent translocation into the nucleus are not completely understood [59-60]. In this sense, the effects of HMG and MGA on classical proinflammatory cytokines (TNF α and IL1- β) were shown to be dependent on ERK signaling pathway. Concerning to IL-6, this cytokine presents a duality of actions, degenerative and reparative and/or anti- and proinflammatory [61]. We found here that IL-6 augment was not associated with ERK, but may be alternatively due to the increase of TNF α and IL1- β , inductors of IL-6 production [62, 50]. Additionally, we explored the involvement of HO1 cascade, another upstream of NF κ B, on cytokine release induced by HMG and MGA. The ZnPP IX inhibitor did not prevent cytokine generation.

The pathophysiological importance of our present data is difficult to predict because brain concentrations of HMG and MGA are not yet established in HMGA affected patients. However, since neurological symptoms appear or become worse during these catabolic crises, in which the concentrations of the accumulating metabolites dramatically increase, it is conceivable a correlation between accumulation of these metabolites and worsening of clinical features. It should be also noted that the concentrations of organic acids in general, and in particular of dicarboxylic acids (such as HMG and MGA), that accumulate in various organic acidemias are higher in the brain as compared with the blood probably because of the low efflux of these compounds from the brain, which is in accordance with the trapping theory [63-66]. Nevertheless, the main relevance of this study is that it provides solid evidence that astrocytes derived from neonatal rats are vulnerable to the toxicants HMG and MGA.

In summary, we describe for the first time a susceptibility of neonatal rat cortical astrocytes to the toxic insults of HMG and MGA, giving rise to a proinflammatory response with increased cytokine production and oxidative stress, accompanied by mitochondrial dysfunction. Our findings confirm previous results carried out in subcellular fractions and homogenates from whole brain and indicate that disruption of astrocyte redox homeostasis caused by the major organic acids accumulating in HMGA may contribute to the neuropathology of patients affected by this disease. Furthermore, our present data indicate that cultured astrocytes represent an interesting and promising biological tool to study toxicity of compounds accumulating in inherited disorders, although their main limitation lies in the fact that it does not involve the intricate relationships between neurons and glial cells. It is therefore presumed that reactive species and cytokine production contribute to the neurological dysfunction observed in this disorder, particularly during metabolic crises, when metabolite production and brain accumulation is even higher. However, we cannot rule out that hyperammonemia and energy deprivation (hypoketotic hypoglycaemia), may also be involved in the brain injury of these patients since these conditions may result in free radical formation in the CNS [58, 67-68]. Overall, these observations suggest that antioxidants may potentially be useful for therapeutic purposes to protect neural cells against oxidative damaging response in HMGA affected patients.

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Conflict of interest

The authors declare that there are no conflicts of interest.

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Legend to figures

Fig. 1 *In vitro* effects of 3-hydroxy-3-methylglutaric (HMG) and 3-methylglutaric acids (MGA) on thiazolyl blue tetrazolium bromide (MTT) reduction in astrocyte cultures from cerebral cortex of rats. Data are represented as mean \pm SD of 3 independent experiments (wells) performed in triplicate and expressed as percentage of controls ** $P < 0.01$, *** $P < 0.001$, compared to control (Duncan's multiple range test)

Fig. 2 *In vitro* effects of 3-hydroxy-3-methylglutaric (HMG, A) and 3-methylglutaric acids (MGA, B) on 2',7'-dichlorofluorescein diacetate (DCFH-DA) oxidation in astrocyte cultures from cerebral cortex of rats. Data are represented as mean \pm SD of 7 to 10 independent experiments (wells) performed in triplicate and expressed as percentage of controls * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, compared to control and expressed as percentage of controls (Duncan's multiple range test)

Fig. 3 *In vitro* effects of 3-hydroxy-3-methylglutaric (HMG, A) and 3-methylglutaric acids (MGA, B) on reduced glutathione (GSH) levels in astrocyte cultures from cerebral cortex of rats. Data are represented as mean \pm SD of 5 to 6 independent experiments (wells) performed in triplicate and expressed as percentage of controls. ** $P < 0.01$, *** $P < 0.001$, compared to control (Duncan's multiple range test)

Fig. 4 *In vitro* effects of 3-hydroxy-3-methylglutaric (HMG) and 3-methylglutaric acids (MGA) on the release of the cytokines interleukin 1 β (IL-1 β - A, B), interleukin 6 (IL-6 - C, D) and tumor necrosis factor α (TNF α - E, F) in astrocyte cultures from cerebral cortex of rats after 24 hours of exposition. Data are represented as mean \pm SD of 4 independent experiments (wells) performed in triplicate and expressed as percentages of controls. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, compared to control (Duncan's multiple range test)

Fig. 5 *In vitro* effects of 3-hydroxy-3-methylglutaric (HMG) and 3-methylglutaric acids (MGA) on the NF κ B levels in astrocyte cultures from cerebral cortex of rats after 24 hours of exposition. Data are represented as mean \pm SD of 4 independent experiments (wells) performed in triplicate and expressed as percentages of controls. No differences between groups were detected (one-way ANOVA)

Fig. 6 *In vitro* effects of 3-hydroxy-3-methylglutaric (HMG) and 3-methylglutaric acids (MGA) on GFAP expression levels in astrocyte cultures from cerebral cortex of rats after 24 hours of exposition. Data are represented mean \pm SD of 2 independent experiments (wells) performed in triplicate and expressed as percentage of controls. No differences between groups were detected (one-way ANOVA)

Table1

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Table 1 - Effects of the ERK inhibitor PD 98059 (5 μ M) on 3-hydroxy-3-methylglutaric (5 mM HMG) and 3-methylglutaric acids (5 mM MGA)-induced alterations on the release of the cytokines interleukin 1 β (IL-1 β), interleukin 6 (IL-6) and tumor necrosis factor α (TNF α) in astrocyte cultures from cerebral cortex of rats

	IL-1 β	IL-6	TNF α
Control	100	100	100
PD 98059	98.7 \pm 7.20	104 \pm 8.67	98.3 \pm 3.78
HMG	137 \pm 10.8 ***	142 \pm 12.9 ***	139 \pm 5.56 ***
MGA	145 \pm 7.44 ***	162 \pm 14.3 ***	140 \pm 5.65 ***
HMG + PD 98059	102 \pm 9.01 ###	147 \pm 11.5 ***	106 \pm 5.40 ###
MGA + PD 98059	96.3 \pm 14.5 ###	160 \pm 19.8 ***	106 \pm 9.45 ###

Values are mean \pm standard deviation for four to six independent (wells) experiments per group expressed as percentage of control. *** $p < 0.001$ compared to control group; ### $p < 0.001$ compared to HMG and MGA (Duncan's multiple range test).

Table2

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Table 2 - Effects of the heme oxygenase 1 (HO1) inhibitor zinc protoporphyrin IX (10 μ M ZnPP IX) on 3-hydroxy-3-methylglutaric (5 mM HMG) and 3-methylglutaric acids (5 mM MGA)-induced alterations on the release of the cytokines interleukin 1 β (IL-1 β), interleukin 6 (IL-6) and tumor necrosis factor α (TNF α) in astrocyte cultures from cerebral cortex of rats

	IL-1 β	IL-6	TNF α
Control	100 \pm 0	100 \pm 0	100 \pm 0
ZnPP IX	98 \pm 6.29	100 \pm 4.03	103 \pm 2.83
HMG	145 \pm 8.33 ***	160 \pm 8.04 ***	136 \pm 5.51 ***
MGA	149 \pm 16.1 ***	172 \pm 18.9 ***	142 \pm 3.77 ***
HMG + ZnPP IX	140 \pm 11.8 ***	152 \pm 9.05 ***	134 \pm 5.16 ***
MGA + ZnPP IX	155 \pm 6.56 ***	177 \pm 6.76 ***	143 \pm 13.9 ***

Values are mean \pm standard deviation for 4 to 6 independent (wells) experiments per group expressed as percentage of control. *** $p < 0.001$ compared to control group (Duncan's multiple range test).

Figure1
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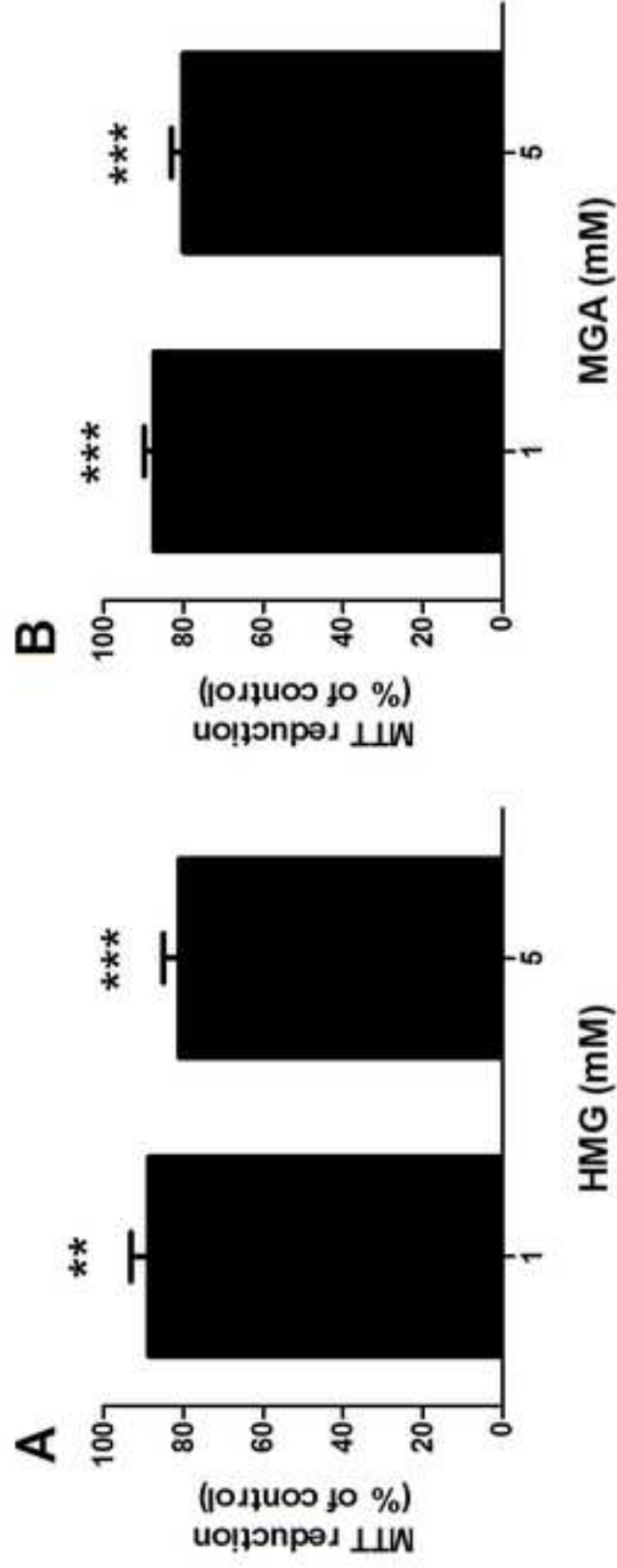
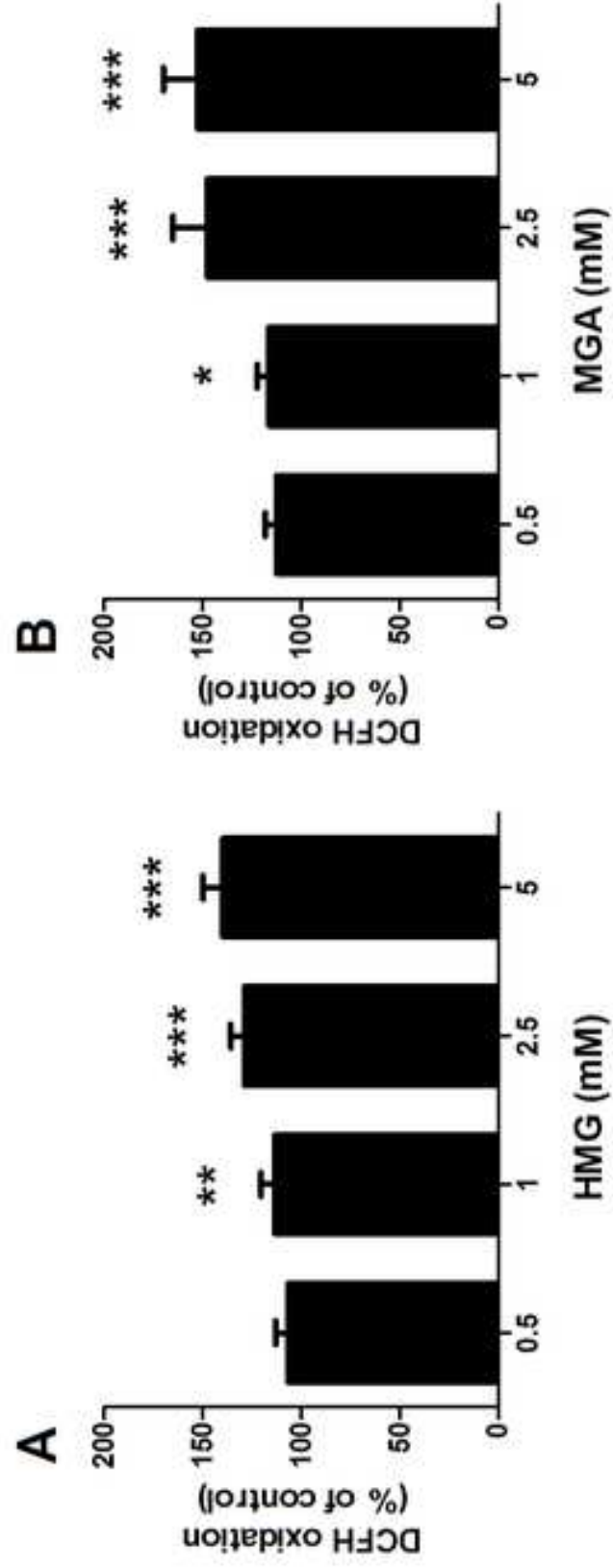


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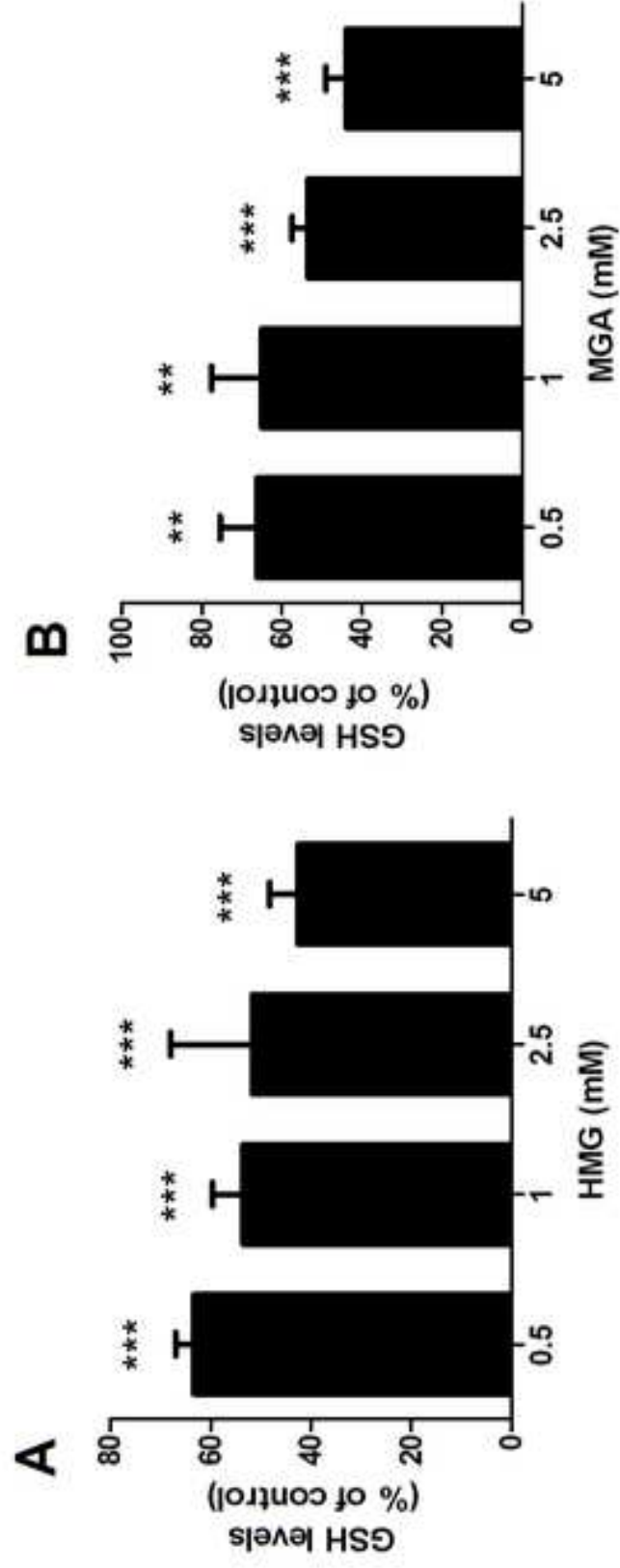


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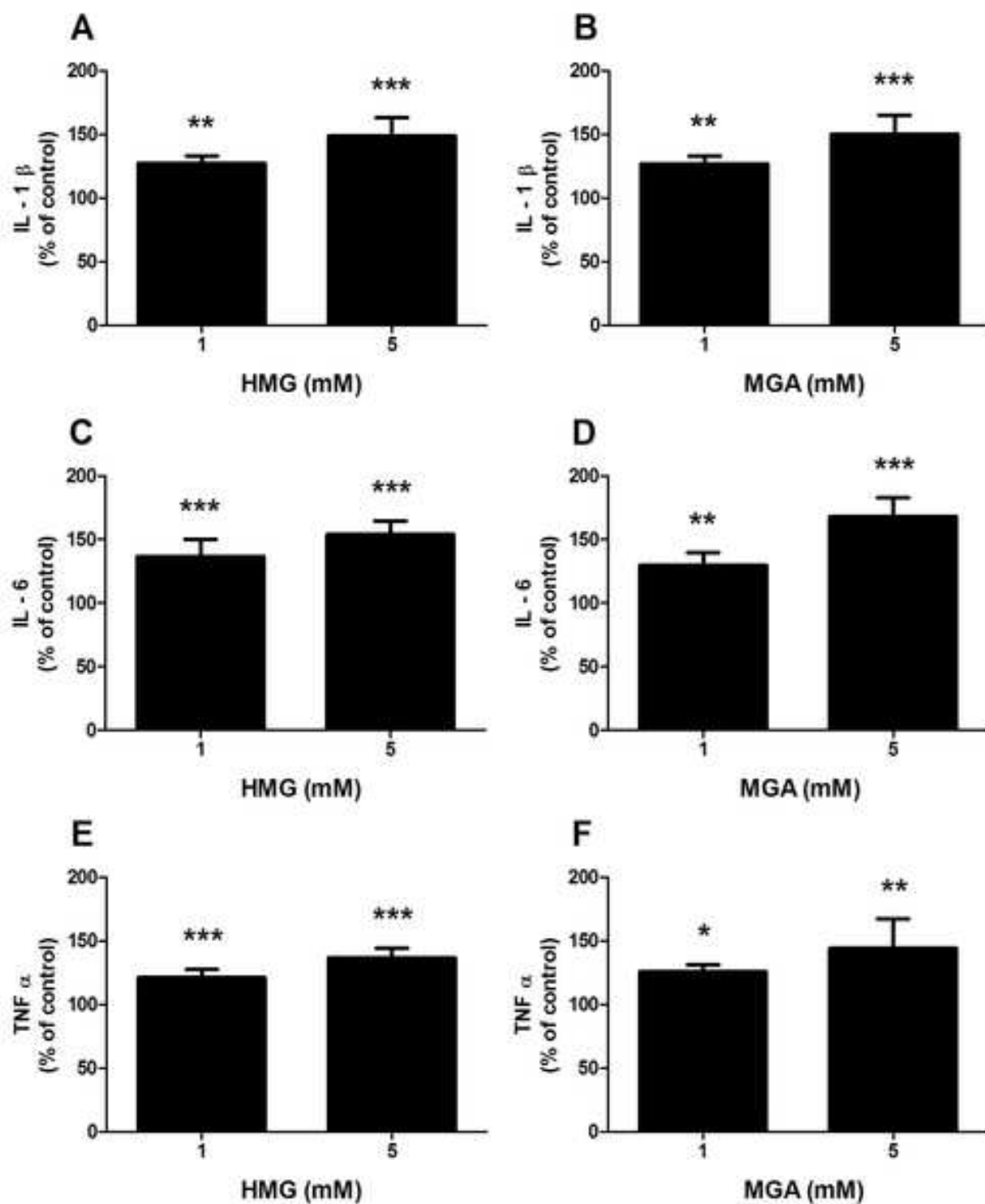


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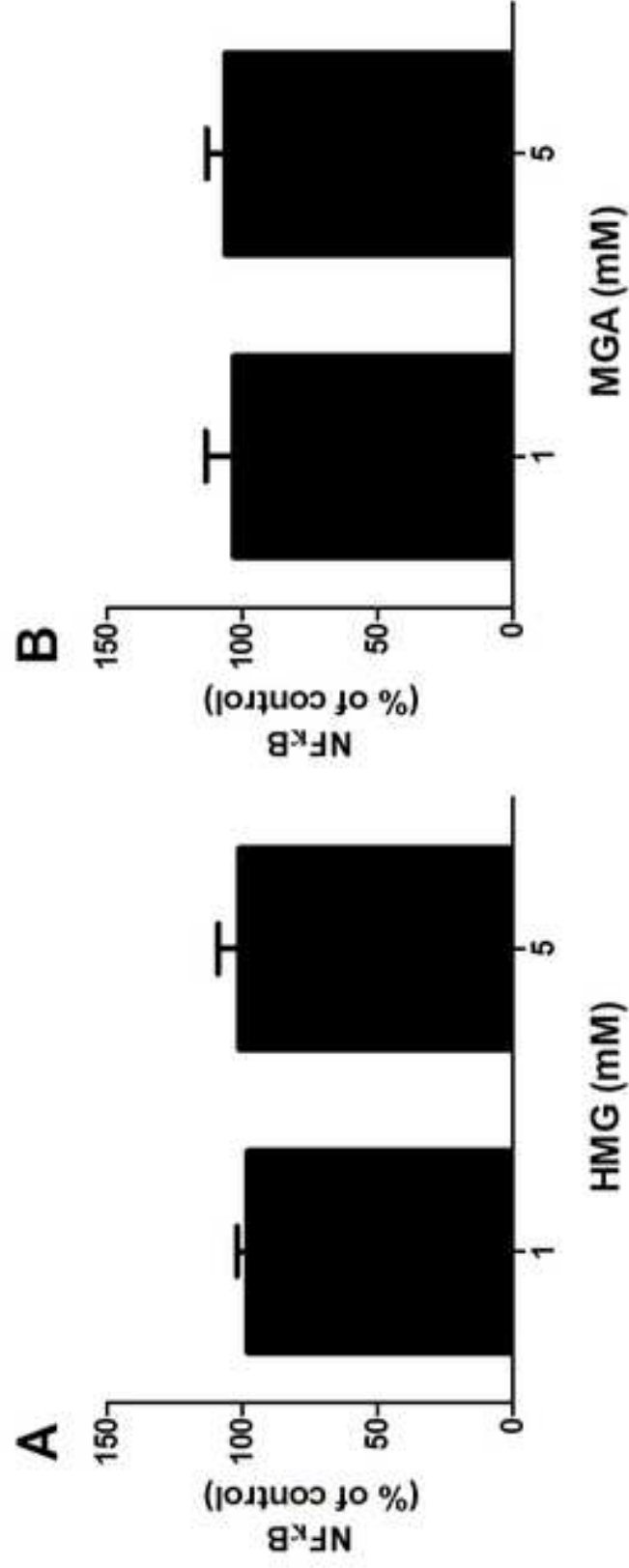
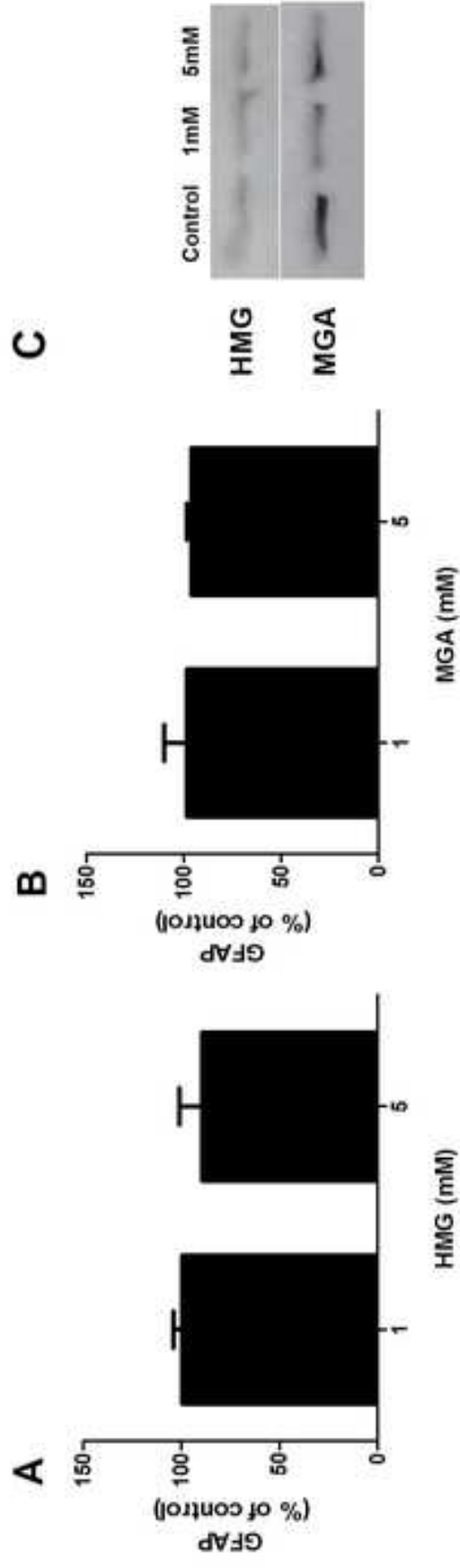


Figure6
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Capítulo 3

Experimental evidence that the major metabolites accumulating in 3-hydroxy-3-methylglutaryl-CoA lyase deficiency induce hypophosphorylation of cytoskeletal proteins of neural cells mediated by protein kinase inhibition and oxidative stress

Carolina Gonçalves Fernandes, Paula Pierozan, Gilberto Machado Soares, Fernanda Ferreira, Ângela Zanatta, Alexandre Umpierrez Amaral, Clarissa Günther Borges, Moacir Wajner and Regina Pessoa-Pureur

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Article Type: Regular Paper

Section/Category: Cellular and Molecular Neuroscience

Keywords: 3-hydroxy-3-methylglutaryl-CoA lyase deficiency; hypophosphorylation; intermediate filaments; cell signaling; glutamatergic receptors

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Abstract: 3-Hydroxy-3-methylglutaryl-CoA lyase (HL) deficiency is an inborn disease biochemically characterized by tissue accumulation and high urinary excretion of 3-hydroxy-3-methylglutaric (HMG), 3-methylglutaric (MGA), 3-methylglutaconic (MGT) acids and also of 3-hydroxyisovaleric acid to a lesser extent. Neurological symptoms and abnormalities are commonly observed in patients with this disease, but their pathogenesis is poorly known. In the present study we found that HMG, MGA and MGT caused hypophosphorylation of glial fibrillary acidic protein and the neurofilament subunits NFL, NFM and NFH in striatal and cortical slices of rats. Energy deprivation was not responsible for these effects since important parameters of mitochondrial energy production were not altered. However, HMG-induced hypophosphorylation of cytoskeletal proteins was mediated by inhibition of protein kinases, with no alterations of protein phosphatases. Thus, we demonstrated that protein cAMP-dependent kinase (PKA) inhibition was involved in the hypophosphorylation of Ser55 of NFL, as well as JNK/MAPK inhibition resulted in the hypophosphorylation of KSP repeats of NFM and NFH subunits. It was also observed that the cytoskeletal hypophosphorylation was dependent on synaptic and extra synaptic (NR2B subunit) NMDA receptors and also on Ca²⁺. Furthermore, the NOS inhibitor, L-NAME, and the antioxidant, TROLOX (polar analog of vitamin E), fully prevented the hypophosphorylation and the inhibition of PKA and JNK activities caused by HMG. These findings indicate that the principal metabolites accumulating in HL deficiency disrupt the cytoskeleton phosphorylating system probably through NMDA receptors, Ca²⁺ and disruption of redox homeostasis. It is presumed that these mechanisms may contribute to the neuropathology of this disease.

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January 26, 2015

Dr. A. Hoke,
Editor-in-Chief,
Experimental Neurology

Dear Dr. Hoke,

I am sending you our manuscript entitled “**Experimental evidence that the major metabolites accumulating in 3-hydroxy-3-methylglutaryl-CoA lyase deficiency induce hypophosphorylation of cytoskeletal proteins of neural cells mediated by protein kinase inhibition and oxidative stress**”, which we would like to submit for publication in **Experimental Neurology**.

This study was performed in strict accordance with the Guide for the Care and Use of Laboratory Animals (Eighth edition, 2011) and was approved by the Ethical Committee of the Federal University of Rio Grande do Sul, Porto Alegre, Brazil. All efforts were also made to use the minimal number of animals necessary to produce reliable scientific data. The manuscript does not contain clinical studies or patient data.

I also inform you that all authors have contributed significantly to the manuscript and agree with the submission of the paper at its present version. Moreover, the whole manuscript, or parts of it, will not be submitted elsewhere for publication. Finally, there is no conflict of interest in the conduct and reporting of research.

This manuscript presents novel findings showing that 3-hydroxy-3-methylglutarate, 3-methylglutarate and 3-methylglutaconate that accumulate in 3-hydroxy-3-methylglutaric aciduria induce a disruption in the cytoskeletal-associated phosphorylating system probably through NMDA receptors, Ca^{2+} and disruption of redox homeostasis that may possibly be involved in the neuropathology of this disease.

I look forward to hearing from you in the near future.

Yours sincerely,

Pessoa-Pureur R. PhD

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Experimental evidence that the major metabolites accumulating in 3-hydroxy-3-methylglutaryl-CoA lyase deficiency induce hypophosphorylation of cytoskeletal proteins of neural cells mediated by protein kinase inhibition and oxidative stress

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Abstract

3-Hydroxy-3-methylglutaryl-CoA lyase (HL) deficiency is an inborn disease biochemically characterized by tissue accumulation and high urinary excretion of 3-hydroxy-3-methylglutaric (HMG), 3-methylglutaric (MGA), 3-methylglutaconic (MGT) acids and also of 3-hydroxyisovaleric acid to a lesser extent. Neurological symptoms and abnormalities are commonly observed in patients with this disease, but their pathogenesis is poorly known. In the present study we found that HMG, MGA and MGT caused hypophosphorylation of glial fibrillary acidic protein and the neurofilament subunits NFL, NFM and NFH in striatal and cortical slices of rats. Energy deprivation was not responsible for these effects since important parameters of mitochondrial energy production were not altered. However, HMG-induced hypophosphorylation of cytoskeletal proteins was mediated by inhibition of protein kinases, with no alterations of protein phosphatases. Thus, we demonstrated that protein cAMP-dependent kinase (PKA) inhibition was involved in the hypophosphorylation of Ser55 of NFL, as well as JNK/MAPK inhibition resulted in the hypophosphorylation of KSP repeats of NFM and NFH subunits. It was also observed that the cytoskeletal hypophosphorylation was dependent on synaptic and extra synaptic (NR2B subunit) NMDA receptors and also on Ca^{2+} . Furthermore, the NOS inhibitor, L-NAME, and the antioxidant, TROLOX (polar analog of vitamin E), fully prevented the hypophosphorylation and the inhibition of PKA and JNK activities caused by HMG. These findings indicate that the principal metabolites accumulating in HL deficiency disrupt the cytoskeleton phosphorylating system probably through NMDA receptors, Ca^{2+} and disruption of redox homeostasis. It is presumed that these mechanisms may contribute to the neuropathology of this disease.

Keywords: 3-hydroxy-3-methylglutaryl-CoA lyase deficiency; hypophosphorylation; intermediate filaments; cell signaling; glutamatergic receptors

Introduction

3-Hydroxy-3-methylglutaryl-coenzyme A (HMG-CoA) lyase (EC 4.1.3.4) is an enzyme found in the mitochondrial matrix and peroxisomes that catalyzes the final step in the metabolism of L-leucine also participating in ketone body synthesis (Gibson et al., 1988; Wysocki and Hähnel, 1986). Deficiency of HMG-CoA lyase (HL), also called 3-hydroxy-3-methylglutaric aciduria (HMGA, OMIM 246450), is an autosomal recessive inborn disease with an estimated prevalence of 1 in 100,000 live births (Pié et al., 2007; Reimão et al., 2009). This enzymatic deficiency results in tissue accumulation and high urinary excretion of 3-hydroxy-3-methylglutaric (HMG), 3-methylglutaric (MGA), 3-methylglutaconic (MGT) and 3-hydroxyisovaleric acids (Faull et al., 1976). The symptoms appearance occurs in the first year of life, normally after a fasting period or other catabolic event. The most prominent findings are vomiting, hepatomegaly with mild liver alterations, lethargy, coma, convulsions, metabolic acidosis and hypoketotic hypoglycaemia (Gibson et al., 1988). Encephalopathy with generalized cerebral edema, structural abnormalities in the white matter (hypomyelination) of cerebral cortex and the basal ganglia are the main findings on MRI (Yalcinkaya et al., 1999; Yilmaz et al., 2006; Zaifeiriou et al., 2007).

Recent *in vivo* studies described oxidative damage and activation of N-methyl-D-aspartate (NMDA) receptors with free radical formation caused by HMG and MGA in the CNS of rats (Fernandes et al., 2013; da Rosa et al., 2013), supporting a role for excitotoxicity and redox imbalance in the brain damage in this disorder. In this context, it must be stressed that these metabolites are structurally similar to glutamate. It is also of note that NMDA receptors and oxidative imbalance are implicated in the misregulation of signaling pathways disrupting the phosphorylating system associated

with the cytoskeleton in various pathological conditions (Loureiro et al., 2010a,b; Cattani et al., 2013; Pierozan et al., 2010).

Intermediate filament (IF) proteins are cell-specific constituents of the cytoskeleton. Mature astrocytes express glial fibrillary acidic protein (GFAP), which is necessary to maintain cell shape, motility/migration, proliferation, glutamate homeostasis and neurite outgrowth, being also important for protection against CNS injury (Middeldorp and Hol, 2011). Neurons express neurofilaments (NF), which are constituted by three distinct subunits classified by their molecular masses: NFH (high molecular weight neurofilament subunit), NFM (middle molecular weight neurofilament subunit) and NFL (light molecular weight neurofilament subunit). They represent the most abundant cytoskeletal elements in mature neurons where their principal function is to control the axonal caliber and nerve conductivity (Perrot and Eyer, 2009).

The physiological role of IF proteins is mainly controlled by phosphorylation, which is a dynamic process mediated by the combined action of several protein kinases and phosphatases in response to extracellular signals. In this context, the phosphorylation sites in IF subunits are located in their amino- and carboxyl-terminal domains and the phosphorylation levels are implicated in the association/disassociation ability as well as in the interaction with other cytoskeletal components (Sihag et al., 2007). *In vivo* and *ex vivo* studies from our group and others demonstrated that the phosphate groups on the N-terminal head domain of GFAP, vimentin, and NFL are added by second messenger-dependent protein kinases, such as cAMP-dependent protein kinase (PKA), Ca²⁺/calmodulin-dependent protein kinase II (PKCaMII), and protein kinase C (PKC) (Sihag et al., 2007; Pierozan et al., 2010, 2012). Phosphorylation sites in the C-terminal domain of NFM and NFH are located in Lys-

Ser-Pro (KSP) repeat regions of these subunits. The KSP repeats are phosphorylated by proline directed kinases such as Cdk5 (Guidato et al., 1996), the mitogen-activated protein kinases (MAPK) Erk1/2, JNK, and p38MAPK (Veeranna et al., 1998; Brownlees et al., 2000) and by the glycogen synthase kinase 3 (GSK3) (Veeranna et al., 2000).

Protein phosphatases are highly concentrated in the mammalian brain (Strack et al., 1997) and the cytoskeleton is a preferential target for the action of Ser-Thr phosphatases 1, 2A and 2B (PP1, PP2A and PP2B) (de Almeida et al., 2003; Funchal et al., 2005a,b; de Mattos-Dutra et al., 1997). In this context, PP1 and PP2B-mediated hypophosphorylation of IF proteins in the cerebral cortex of rats caused by intrastrially injected of quinolinic acid (QUIN) was first reported in our laboratory and proposed to misregulate cytoskeletal homeostasis and be associated with QUIN toxicity (Pierozan et al., 2014).

Thus, considering that the phosphorylating system associated with the cytoskeleton may be related to neural damage and that the pathogenesis of the brain damage in HL deficiency is still poorly elucidated, in the present report we investigated the actions of HMG, MGA and MGT on the phosphorylating system associated with the IF-enriched cytoskeleton in striatal and cortical slices of young rats. We also searched for the role of NMDA receptors and oxidative stress on the alterations of the cytoskeletal phosphorylating system homeostasis provoked by HMG exposure. We hypothesize that disruption of cytoskeletal homeostasis could be implicated in the neural damage induced by the main metabolites accumulating in HMG-CoA lyase deficiency.

Material and Methods

Chemicals

[³²P]Na₂HPO₄ was purchased from Comissão de Energia Nuclear (CNEN), São Paulo, Brazil. 3-Hydroxy-3-methylglutaric acid (HMG), 3-methylglutaconic (MGA) acid, 3-methylglutaconic acid (MGT), acrylamide, antipain, 1,2-bis (2-aminophenoxy) ethane-N-N-N'-N'-tetraacetic acid tetrakis (acetoxymethyl Ester) (BAPTA-AM), benzamidine, bisacrilamide, calyculin A, chymostatin, dizocilpine (MK-801), ethylene glycol tetraacetic acid (EGTA), ifenprodil, leupeptin, NG-nitro-L-arginine methyl ester (L-NAME), N-acetyl-L-cystein (NAC), pepstatin, tacrolimus (FK-506), (±)-6-hydroxy-2,5,7,8-tetramethylchromane-2-carboxylic acid (TROLOX) were obtained from Sigma (St. Louis, MO, USA). Anti-pNFL(Ser55) and anti-p38MAPK and anti-phospho p38MAPK antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). The anti-PKCamII α, anti-JNK (p54), anti-JNK (p46) anti-pJNK (p54), anti-pJNK (p46), anti-Erk (p44), anti-Erk (p42) anti-pErk (p44), anti-pErk (p42) and anti PKAα antibodies were from Cell Signaling (Danvers, MA, USA). The antibody NFH (anti-KSP repeats) was purchased from Chemicon (Temecula, CA, USA). The chemiluminescence ECL kit peroxidase and the conjugated anti-rabbit IgG were obtained from Amersham (Oakville, Ontario, Canada). All chemicals were of analytical grade.

Animals

Thirty-day-old Wistar rats obtained from Central Animal House of the Departamento de Bioquímica da Universidade Federal do Rio Grande do Sul, Porto Alegre, Brazil, were used in the experiments. The animals were maintained on 12:12 h light/dark cycle in an air-conditioned constant temperature (22 °C ± 1 °C) colony room, with food and water

ad libitum. The project followed the experimental protocol “Principles of Laboratory Animal Care” (NHI publication 85-23, revised 1985) and was approved by the Ethics Committee for Animal Research of the Federal University of Rio Grande do Sul.

Phosphorylation assay

Pre-incubation

Rats were sacrificed by decapitation, the striatum and cerebral cortex were dissected onto Petri dishes placed on ice and cut into 400 μm thick slices with a McIlwain chopper. Tissue slices were initially preincubated at 30 °C for 10 min in a Krebs-Hepes medium containing 124 mM NaCl, 4 mM KCl, 1.2 mM MgSO_4 , 25 mM Na-Hepes, 12 mM glucose, 1 mM CaCl_2 , pH 7.4, and the following protease inhibitors: 1 mM benzamide, 0.1 μM leupeptin, 0.7 μM pepstatin and 0.7 μM chymostatin. In some experiments we used 50 μM BAPTA-AM, 0.2 μM caliculyn, 1 mM EGTA, 100 μM FK-506, 10 μM ifenprodil, 1 mM N ω -Nitro-L-arginine methyl ester hydrochloride (L-NAME), 10 μM dizocilpine (MK-801), 1 mM N-acetyl cysteine (NAC), 5 μM 6-hydroxy-2,5,7,8-tetramethylchroman-2-carboxylic acid (TROLOX).

Incubation and extraction of the cytoskeletal fraction

After pre-incubation, the medium was changed by 100 μL of a Krebs-Hepes medium in the absence or presence of HMG, MGA or MGT (0.2 to 5 mM). In some experiments we evaluated the effects of 5 mM HMG using striatum slices in order to find out the mechanisms of action of this organic acid on the cytoskeletal phosphorylating system. 100 μCi of [^{32}P]-orthophosphate was then added to the incubation medium and the labeling reaction was allowed to proceed for 60 min at 30 °C. Reaction was stopped with 1 mM of cold stop buffer (150 mM NaF, 5 mM EDTA, 5 mM EGTA, 50 mM Tris-

HCl, pH6.5, and the protease inhibitors described above). Slices were then washed twice with stop buffer to remove excess radioactivity. These slices were homogenized in 400 μ L of ice-cold high salt buffer containing 5 mM KH_2PO_4 , 600 mM KCl, 10 mM MgCl_2 , 2 mM EGTA, 1 mM EDTA, 1 % triton X-100, pH 7.1 and the protease inhibitors described above. The homogenate was centrifuged at 15,800 x g for 10 min at 4 °C, the supernatant discarded and the pellet washed/resuspended with the same volume of the high salt buffer. The washed/resuspended pellet was centrifuged as described and the supernatant was discarded. The Triton-insoluble IF-enriched pellet, containing NF subunits and GFAP, was dissolved in 1 % SDS and after prepared to electrophoresis with a concentrated sample buffer containing 33.3 % glycerol, 16 % β -mercaptoethanol, 5 % SDS, 0.033 M NaOH and boiled for 3 min (Funchal et al., 2003). Samples were analyzed in 7.5 % SDS-PAGE

Polyacrylamide gel electrophoresis (SDS-PAGE)

Samples of IF-enriched fraction containing equal protein content (50 μ g) were loaded onto a 7.5 or 10 % gel. The gels were exposed to X-ray films (T-mat G/RA) (Kodak) at -70 °C with intensifying screens and after 30 days the autoradiograph was obtained. The [^{32}P]-orthophosphate incorporated to each sample was quantified from each correspondent cytoskeletal protein by scanning the films with a Hewlett-Packard Scanjet 6100C scanner and determining optical densities with an AlphaEaseFC version 6.0.0 software (FluorChem SA). Protein loading was controlled by Comassie blue R staining. All bands measured were within the linear range of the film.

Western blot analysis

Total tissue homogenate was obtained from striatum slices previously incubated for 60 min at 30 °C with or without 5 mM HMG and in some experiments with the addition of

1 mM of L-NAME or 5 μ M of TROLOX. The slices were homogenized in 100 mL of a lysis solution containing 2 mM EDTA, 50 mM Tris-HCl, 4 % SDS, pH 6.8. This was followed by dilution in a solution containing 40 % glycerol, 5 % β -mercaptoethanol, 50 mM Tris-HCl, pH 6.8 and boiled for 3 min. The samples (30 μ g of protein) were loaded onto a 10 % gel by SDS-PAGE according to the discontinuous system of Laemmli (Laemmli, 1970) and transferred to nitrocellulose membranes (Trans-blot SD semi-dry transfer cell, BioRad) for 1 h at 15 V in transfer buffer: 48 mM Trizma, 39 mM glycine, 20 % methanol and 0.25 % SDS. The nitrocellulose membranes were washed for 10 min in Tris buffered saline (TBS: 0.5 M NaCl, 20 mM trizma, pH 7.5), followed by 2 h incubation in blocking solution (TBS plus 5 % deffated dried milk). After incubation, the blots were washed twice for 5 min with TBS plus 0.05 % Tween-20 (T-TBS), and then incubated overnight at 4 °C in blocking solution containing the following monoclonal antibodies: anti-PKCaMII α , anti-p38/MAPK, anti-PKAc α , anti-JNK (p54), anti-JNK (p46) anti-pJNK (p54), anti-pJNK (p46), anti-Erk (p44), anti-Erk (p42) anti-pErk (p44), anti-pErk (p42) diluted 1:1000, anti-pNFL(Ser55) diluted 1:800, anti-NFH/NFM KSP repeats diluted 1:100 anti- β actin diluted 1:1000. The blot was then washed twice for 5 min with T-TBS and incubated for 2 h in TBS containing peroxidase conjugated rabbit anti-mouse IgG or anti-rabbit IgG diluted 1:2000. We also used the following negative controls: antibody negative control (lacking the sample containing the antigen of interest) and sample negative control (lacking the first antibody). The blot was washed twice again for 5 min with T-TBS and twice for 5 in with TBS. The blot was then developed using a chemiluminescence ECL kit and quantified as described above.

Cell viability

Cell viability was evaluated by lactate dehydrogenase (LDH) activity (Whitaker and McKay, 1969) and by MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) reduction (Liu et al., 1997) in striatal and cortical slices. LDH release was determined using a colorimetric commercial kit (from Doles, Brazil) and expressed as percentage of positive control samples after addition of 10 % Triton X-100. Results were expressed relative to 100% LDH activity. For the MTT assay, MTT (0.5 mg/mL) in Krebs–Hepes medium was incubated for 60 min at 30 °C. The tetrazolium ring of MTT is cleaved by various dehydrogenases and then precipitated as a blue formazan product. The medium was then aspirated, the precipitated formazan was solubilized with dimethyl sulfoxide and quantified spectrophotometrically at a wavelength of 550 nm.

Parameters of energy homeostasis

CO₂ production

Cerebral cortex and striatum slices (400 µm) were pre-incubated with or without 5 mM HMG or MGA in Krebs–Ringer bicarbonate buffer (1.17 mM KH₂PO₄, 118 mM NaCl, 4.69 mM KCl, 24.1 mM MgSO₄, 3.33 mM CaCl₂ and 650 mM NaHCO₃, pH 7.4) into small flasks (11 cm³) in a volume of 450 µL at 35 °C for 60 min in a metabolic shaker (90 oscillations / min). After pre-incubation, [U-¹⁴C] glucose (0.055 mCi) plus 5.0 mM of unlabeled glucose was added to the incubation medium. The flasks were gassed with an O₂/CO₂ (95:5) mixture and sealed with rubber stoppers Parafilm M. Glass center wells containing a folded 60 nm / 4 nm piece of Whatman 3 filter paper were hung from the stoppers. After incubation in the metabolic shaker 0.2 mL of 50 % trichloroacetic acid was supplemented to the medium and 0.1 mL of benzethonium hydroxide was

added to the center of the wells with needles introduced through the rubber stopper. The flasks were left to stand for 30 min to complete CO₂ trapping and then opened. The filter paper were removed and added to vials containing scintillation fluid, and radioactivity was counted (Reis de Assis et al., 2004). Results were calculated as nmol CO₂ . h⁻¹ . g tissue⁻¹ and expressed as percentage of control.

Preparation of mitochondrial fractions

Mitochondrial preparations from forebrain were isolated according to Rosenthal and co-workers (1987), with slight modifications (Mirandola et al., 2008). Animals were decapitated, the forebrain was dissected and homogenized with a glass hand-held homogenizer in ice-cold mitochondria isolation buffer containing 225 mM mannitol, 75 mM sucrose, 1 mM EGTA, 0.1 % bovine serum albumin (BSA, free fatty acid) and 10 mM HEPES, pH 7.2. The homogenate was centrifuged at 2,000 x g for 3 min at 4 °C. The pellet was discarded and the supernatant was centrifuged at 12,000 x g for 8 min at 4 °C. The resultant pellet was resuspended in 5 mL of isolation buffer containing 20 µL of 10% digitonin (final concentration of 0.04%), and centrifuged 12,000 x g for 10 min at 4 °C. The supernatant was discarded and the pellet resuspended in 5 mL of isolation buffer without EGTA and centrifuged at 12,000 x g for 10 min at 4 °C. The final pellet was resuspended in isolation buffer without EGTA in an approximate protein concentration of 15-20 mg/mL. Protein concentration was measured by the method of Bradford (1976) using BSA as standard. This preparation results in a mixture of synaptosomal and non-synaptosomal mitochondria similar to the general brain composition. Mitochondria obtained from forebrain were used in the assays immediately after isolation and assays were carried out in the absence or presence of Ca²⁺.

Spectrophotometric analysis of the citric acid cycle enzyme activities

The activities of aconitase and α -ketoglutarate dehydrogenase (α -KGDH) complex enzymes of the citric acid cycle (CAC) were determined using enriched mitochondrial fractions from whole brain, whereas succinate dehydrogenase (SDH) was measured in brain homogenates. HMG or MGA (5 mM) were supplemented to the subcellular preparations and submitted to a pre-incubation at 37 °C for 60 min. The activity of the enzyme aconitase (ACO) was measured according to Morrison (1954), following the reduction of NADP⁺ at wavelengths of excitation and emission of 340 and 466 nm, respectively. The activity of the α -KGDH complex was evaluated according to Lai and Cooper (1986) and Tretter and Adam-Vizi (2004), with slight modifications (Viegas et al., 2009). The reduction of NAD⁺ was recorded in a Hitachi F-4500 spectrofluorometer at wavelengths of excitation and emission of 340 and 466 nm, respectively. The activity of succinate dehydrogenase (SDH) was determined as described by Fischer and colleagues (1985). The activities of these citric acid cycle enzymes were calculated as $\text{nmol} \cdot \text{min}^{-1} \cdot \text{mg protein}^{-1}$; $\text{mmol} \cdot \text{min}^{-1} \cdot \text{mg protein}^{-1}$ or $\text{mmol} \cdot \text{min}^{-1} \cdot \text{mg protein}^{-1}$ and expressed as percentage of control.

Spectrophotometric analysis of the respiratory chain complexes I–IV activities

The activities of succinate–2,6-dichloroindophenol (DCIP)–oxidoreductase (complex II) and succinate:cytochrome *c* oxidoreductase (complexes II–III) were determined in homogenates from striatum or cerebral cortex according to Fischer and co-workers (1985) and that of cytochrome *c* oxidase (complex IV) according to Rustin and co-workers (1994). The methods described to measure these activities were slightly modified, as described in details in a previous report (da Silva et al., 2002). HMG or MGA (5 mM) was added to the reaction medium at the beginning of the assays, while

no metabolite was added to controls. The activities of the respiratory chain complexes were calculated as $\text{nmol} \cdot \text{min}^{-1} \cdot \text{mg protein}^{-1}$ or $\text{mmol} \cdot \text{min}^{-1} \cdot \text{mg protein}^{-1}$ and expressed as percentage of control.

Mitochondrial membrane potential

Mitochondrial membrane potential ($\Delta\Psi_m$) was estimated according to Akerman and Wikstrom (1976) and Kowaltowski and co-workers (2002). The fluorescence of 5 μM cationic dye safranin O, a $\Delta\Psi_m$ indicator, was followed at excitation and emission wavelengths of 495 and 586 nm. CCCP was added in the end of measurements to abolish $\Delta\Psi_m$. Mitochondrial incubations were carried out at 37 °C, with continuous magnetic stirring. The assay was conducted in medium containing 150 mM KCl, 5 mM MgCl_2 , 30 μM EGTA, 0.1 mg / mL BSA, 5 mM HEPES, 2 mM KH_2PO_4 , pH 7.2, 1 μg / mL oligomycin A, using state 4 respiring mitochondria (0.5 mg protein / mL) supported by 2.5 mM glutamate plus 2.5 mM malate or succinate plus rotenone. HMG (5mM), MGA (5mM), CaCl_2 (30 μM) and CCCP (3 μM) were added as indicated by the arrows in the figure. Traces are representative of independent experiments carried out in mitochondrial preparations from forebrain of three animals and were expressed as fluorescence arbitrary units (FAU).

Protein determination

The protein concentration was determined by the methods of Lowry (Lowry et al., 1951) or Bradford (Bradford, 1976) using serum albumin as standard.

Statistical analysis

Data were statistically analyzed by one-way analysis of variance (ANOVA) followed by the Duncan multiple range test when the F-test was significant. A $p < 0.05$ was

considered significant. All analyses were performed using the SPSS 19.0 software program on an IBM-PC compatible computer.

Results

We first observed that cell viability (MTT assay and LDH release) was not altered in striatal and cortical slices exposed to HMG, MGA or MGT for 60 min (results not shown). In contrast, HMG (Figure 1 A, B, G and H) and MGA (Figure 1 C, D, I and J) caused hypophosphorylation of GFAP and NF subunits (NFL, NFM and NFH) in both brain structures, whereas MGT provoked hypophosphorylation of cytoskeletal IF only in the striatum (Figure 1E and K).

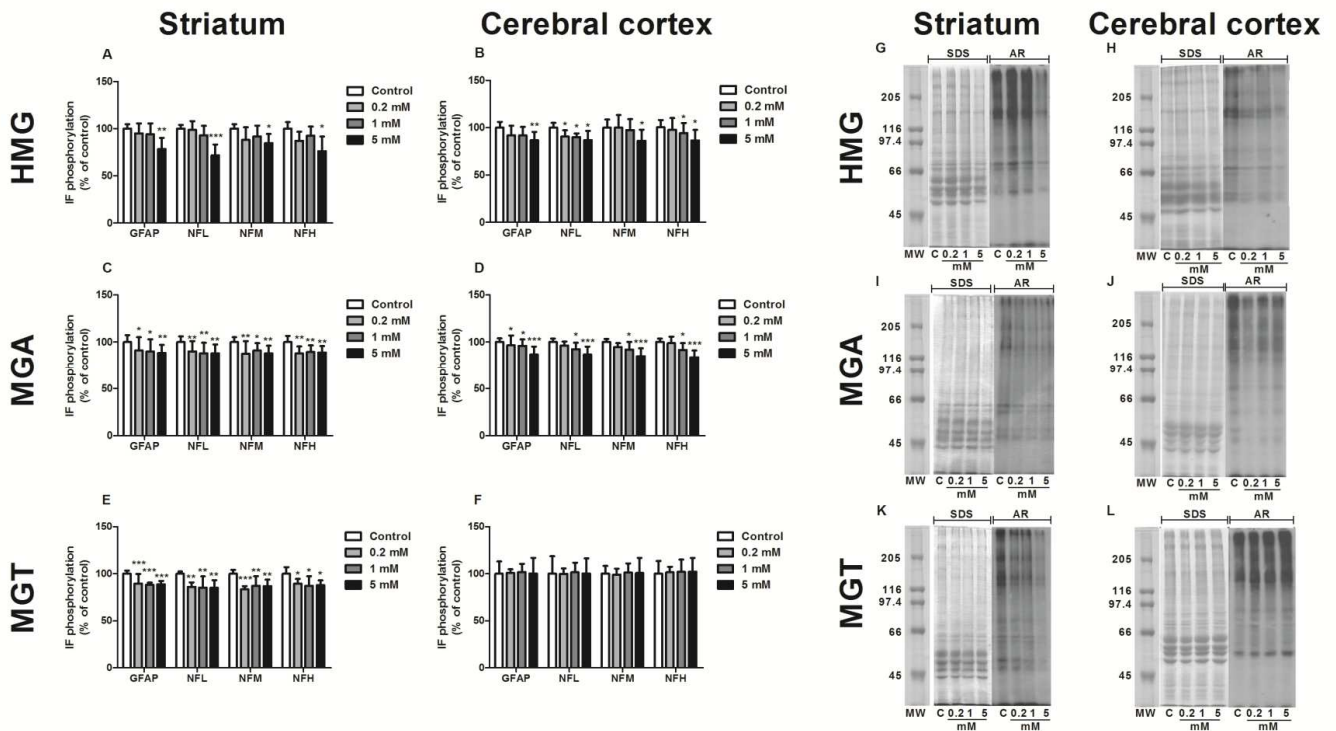


Figure 1 –Effect of HMG, MGA or MGT on the phosphorylation levels of IFs in the cytoskeletal fraction from slices of striatum or cerebral cortex of 30-day-old rats. Tissue slices were incubated during 60 min in the presence or absence (control group) of the metabolites (0.2 – 5 mM). Effects of HMG (A and B), MGA (C and D) and MGT (E and F) on the radioactivity incorporated into IFs in striatal and cortical slices. Data are reported as mean \pm SD of 18 animals in each group and expressed as percentage of controls. Statistically significant from controls: * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ (Duncan’s multiple range test). Comassie blue stained molecular weight standards, representative stained gel and corresponding autoradiography from striatal slices on HMG (G), MGA (I) and MGT effects (K) and from cerebral cortex slices on HMG (H), on MGA (J) and MGT effects (L). AR: autoradiography; C: Control; MW: molecular weight standard proteins; SDS: SDS page.

In order to understand the mechanisms underlying the effects of HMG and MGA on the phosphorylation system associated with the cytoskeleton, we measured some parameters of energy metabolism to test the possibility that decreased ATP availability could be responsible for the disrupted phosphorylating system. We found no alteration of citric acid cycle enzyme activities, CO_2 production, activities of the respiratory chain complexes II, II-III and IV and mitochondrial membrane potential in striatal and cortical preparations exposed to HMG- and MGA, making unlikely that decreased energy

production could explain our results of hypophosphorylation (Tables I, II and III; Figure 2).

Table I – Effect of 3-hydroxy-3-methylglutaric (HMG) or 3-methylglutaric (MGA) acids on the CO₂ production from [U-¹⁴C] glucose in rat brain tissue slices

	Controle	HMG	MGA
Striatum	100 ± 13.1	113 ± 6.15	-
Cerebral cortex	100 ± 26.9	114 ± 40.4	122 ± 39.1

Values are mean ± standard deviation for four to six independent (animals) experiments per group. The CO₂ production was calculated as pmol ¹⁴CO₂ · h⁻¹ · g tissue⁻¹ and expressed as percentage of control. No significant differences between groups were detected (one way ANOVA).

Table II – Effect of 3-hydroxy-3-methylglutaric (HMG) or 3-methylglutaric (MGA) acids on the citric acid cycle (CAC) enzymes in enriched mitochondrial fractions from rat brain

	Control	HMG	MGA
ACO	100 ± 26.3	113 ± 20.9	110 ± 17.7
α-KGDH	100 ± 42.2	98.1 ± 34.6	103 ± 54.8

Values are mean ± standard deviation for four to six independent (animals) experiments per group. The activities of aconitase (ACO) and α-ketoglutarate dehydrogenase (α-KGDH) were calculated as μmol NADPH · min⁻¹ · mg protein⁻¹ and mmol NADH · min⁻¹ · mg protein⁻¹, respectively, and expressed as percentage of control. No significant differences between groups were detected (one way ANOVA).

Table III - Effect of 3-hydroxy-3-methylglutaric (HMG) or 3-methylglutaric (MGA) acids on the activities of respiratory chain complexes and the activity of the CAC enzyme succinate dehydrogenase (SDH) in rat brain homogenates

	Striatum			Cerebral cortex		
	Control	HMG	MGA	Control	HMG	MGA
II	100 ± 14.2	101 ± 13.4	97.8 ± 6.29	100 ± 5.50	95.9 ± 12.7	97.8 ± 7.25
II-III	100 ± 11.8	100 ± 8.46	101 ± 9.13	100 ± 7.33	99.3 ± 3.07	101 ± 6.61
IV	100 ± 10.5	100 ± 17.4	94.4 ± 11.5	100 ± 29.1	99.9 ± 10.4	102 ± 15.2
SDH	100 ± 7.46	100 ± 18.3	98.7 ± 11.3	100 ± 11.3	96.1 ± 4.83	98.3 ± 2.89

Values are mean± standard deviation for four to six independent (animals) experiments per group. The activities of complexes II, II–III, IV and the activity of succinate dehydrogenase (SDH) were calculated as nmol · min⁻¹ · mg protein⁻¹ and nmol DCIP · min⁻¹ · mg protein⁻¹, respectively, and expressed as percentage of control. No significant differences between groups were detected (one way ANOVA).

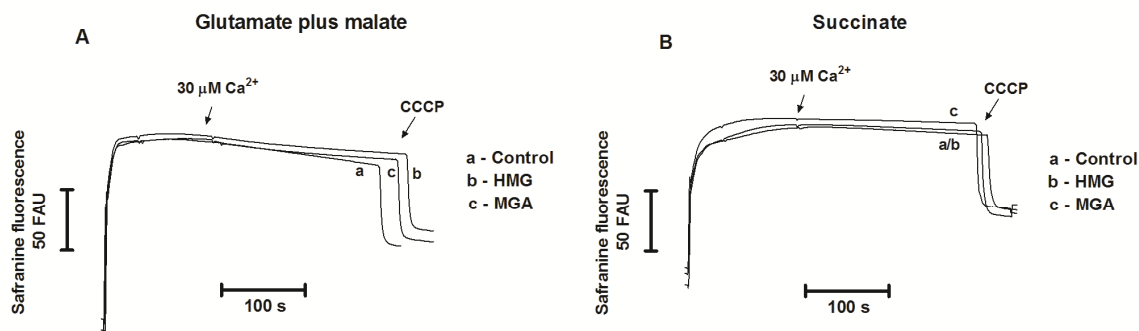


Figure 2 – Effects of HMG or MGA acids on mitochondrial membrane potential in the presence of Ca^{2+} ($30 \mu\text{M}$). All experiments were performed in a reaction media containing mitochondrial preparations ($0.5 \text{ mg protein. mL}^{-1}$) supported by glutamate plus malate (A) or succinate (B). HMG or MGA (5 mM , lines b-c, respectively) were added 50 seconds after the beginning of the assay. Controls (line a) were performed in the absence of the metabolites. The reaction media contained $30 \mu\text{M EGTA}$. CCCP ($3 \mu\text{M}$) was added at the end of the assays. Traces are representative of 3 independent experiments (animals) and were expressed as fluorescence arbitrary units (FAU).

We further investigated the mechanisms underlying the hypophosphorylation of IF induced by HMG, the major accumulating compound, using the striatum that is mainly compromised in patients affected by HL deficiency (Sweetman and Williams, 2001; van der Knaap et al., 1998; Yalcinkaya et al., 1999; Yilmaz et al., 2006).

Since hypophosphorylation can be due to protein phosphatase activation and/or protein kinase inhibition, we first searched for the protein phosphatases involved in the hypophosphorylation of cytoskeletal proteins using specific Ser-Thr phosphatase inhibitors. We observed that the potent PP1 and PP2A inhibitor calyculin A (Sheppek et al., 1997) and the specific PP2B inhibitor FK506 (Baba et al., 2003) were ineffective in preventing HMG-induced decrease of IF phosphorylation (Figure 3).

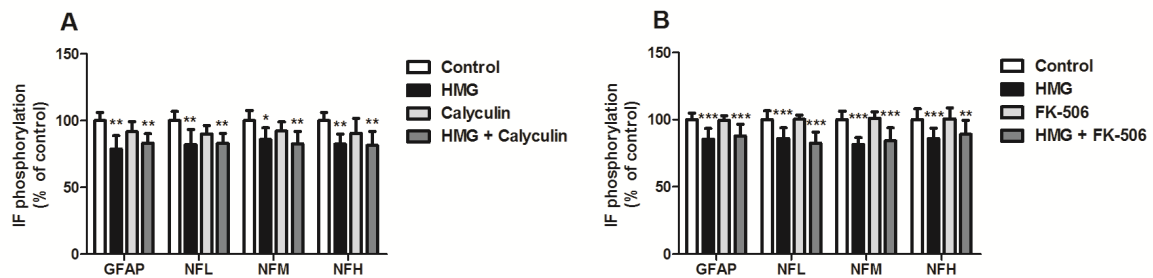


Figure 3 – Effects of PP1, PP2A and PP2B inhibitors on HMG-induced decrease of the phosphorylation levels of IFs in the cytoskeletal fraction from striatum slices of 30-day-old rats. Tissue slices were co-incubated during 60 min in the presence or absence (control group) of the metabolite (5 mM) plus 0.2 μ M calyculin (A; PP1 and PP2A inhibitor) or 100 μ M FK-506 (B; PP2B inhibitor). Data are reported as mean \pm SD of 18 animals in each group and expressed as percentage of controls. Statistically significant differences from controls were determined by one-way ANOVA followed by Duncan's multiple range test. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ compared with control.

We next investigated whether the most common Ser–Thr kinases known to modulate the phosphorylating level of IF cytoskeletal proteins, namely cAMP-dependent protein kinase (PKA), Ca^{2+} /calmodulin-dependent protein kinase (PKCamII) and the mitogen-activated protein kinases (MAPK) Erk, JNK and p38MAPK (Hisanaga et al., 1990; Sihag and Nixon, 1989; Sihag et al., 2007), were involved in HMG hypophosphorylating effect. Western blot analysis using specific antibodies showed unaltered immunocontent of the active PKCamII subunit (Figure 4A), but decreased immunocontent of the active PKA subunit (Figure 4B), evidencing a role for PKA in the hypophosphorylation of the IF proteins induced by this metabolite. We also tested the phosphorylation level of NFLSer55, known to be the preferential phosphorylation site for PKA in the amino-terminal region of NFL subunit (Grant and Pant, 2000). It was observed reduced immunocontent of phosphorylated Ser55, reinforcing the involvement of PKA in HMG hypophosphorylating effects (Figure 4C).

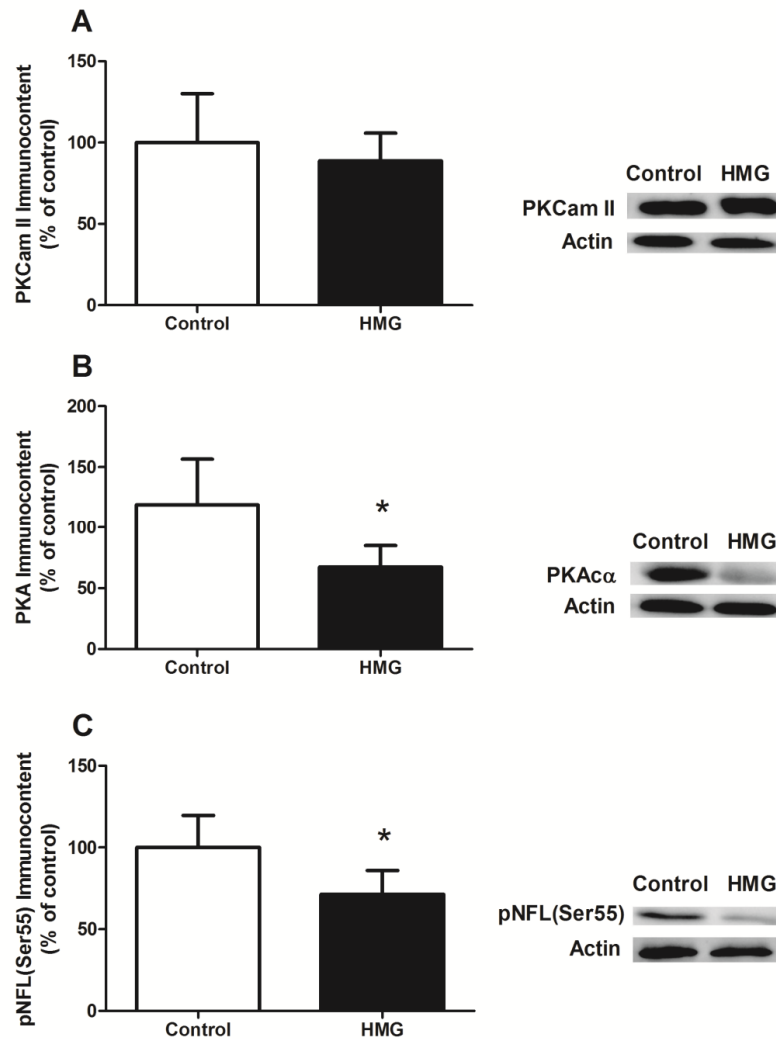


Figure 4 - Effects of HMG on the immunocontent of PKC α (A), PKA (B) and Ser55 phosphorylation site (C) in the total homogenates from striatal slices of 30-day-old rats. Western blot with anti-PKC α , anti-PKA or anti-pNFL(Ser55) antibodies. Tissue slices were incubated during 60 min in the presence or absence (control group) of the metabolite (5 mM). Data are reported as mean \pm SD of 6 animals in each group and expressed as percentage of controls. Statistically significant differences from controls were determined by Student *t* test. * $p < 0.05$ compared with control.

Next, we examined the involvement of the MAPK cascade on HMG-induced IF hypophosphorylation. Western blot analysis with specific monoclonal antibodies showed unaltered immunocontent of Erk1/phospho-Erk1 and Erk2/phospho-Erk2 (Figure 5A), as well as of p38MAPK/phospho-p38MAPK (Figure 5B). In contrast,

phospho-JNK, but not total JNK, immunoccontent was significantly decreased (Figure 5C), implying a reduction of JNK activity/phosphorylation. In an attempt to verify the implications of the downregulated pJNK on the hypophosphorylation of NFM and NFH subunits, we determined the phosphorylation level of the KSP repeats located in the tail domain of these subunits, known to be the preferential phosphorylating sites for MAPKs (Veeranna et al., 2000). Results of Western blot assay with the phosphoKSP repeat antibody showed reduced immunoccontent of these phosphorylated sites in response to HMG (Figure 5D).

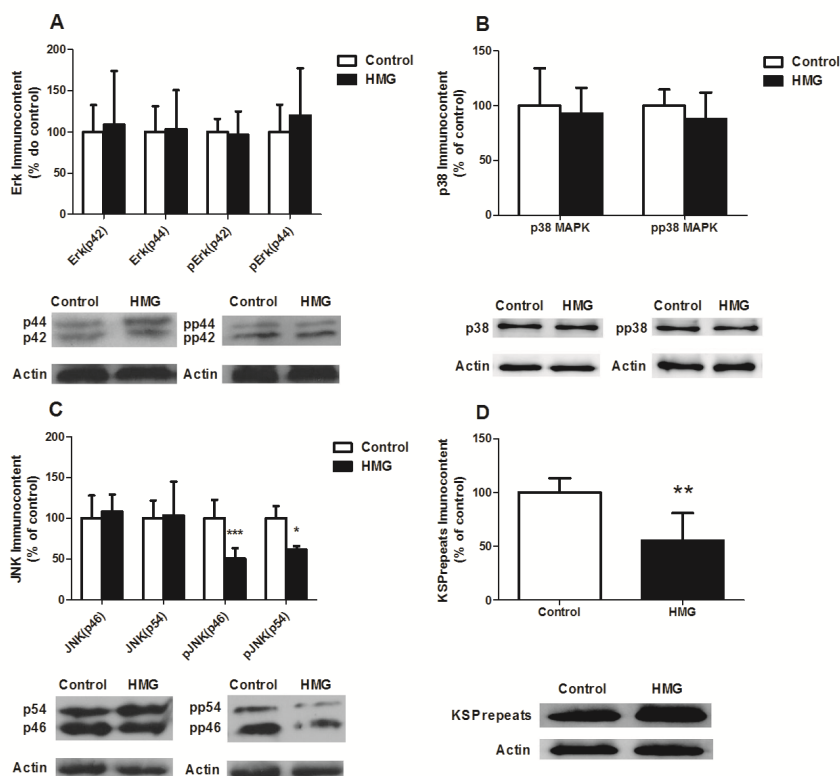


Figure 5 - Effects of HMG on the immunocontent of Erk 1/2 (A), p38MAPK (B), JNK (C) and phosphoKSP repeats (D) in the total homogenates from striatal slices of 30-day-old rats. Western blot with anti-Erk (p42), anti-Erk (p44), anti-pErk (p42), anti pErk (p44), anti-p38/MAPK, anti-phospho-p38/MAPK, anti-JNK (p54), anti-JNK (p46), anti-pJNK (p54), anti-pJNK (p46) or anti-KSP repeat antibodies. Tissue slices were incubated during 60 min in the presence or absence (control group) of the metabolite (5 mM). Data are reported as mean \pm SD of 6 animals in each group and expressed as percentage of controls. Statistically different from controls: * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ (Student's t test).

Since we have shown that NMDA receptors are implicated in oxidative stress caused by intracerebral administration of HMG in rat striatum (Fernandes et al., 2013), we next assessed the involvement of NMDA receptors in the IF hypophosphorylated levels in HMG treated striatal slices. Results showed that 10 μ M MK-801, a non-competitive antagonist of synaptic NMDARs (Pierozan et al., 2010) totally prevented HMG-induced GFAP, NFL, NFM and NFH hypophosphorylation (Figure 6A). Similar results were obtained with 10 μ M ifenprodil, a selective antagonist of extra-synaptic

NMDAR-containing NR2B subunit (Rittase et al., 2014) (Figure 6B). These data indicate the participation of both synaptic and extra-synaptic glutamate receptors in the hypophosphorylating effect of HMG towards IF.

We also found that the IF hypophosphorylation caused by HMG was totally prevented when striatum slices were co-incubated with the intracellular and extracellular Ca^{2+} chelators BAPTA-AM ($50 \mu\text{M}$) and EGTA (1 mM), respectively, in a Ca^{2+} -free medium (Figure 6C). These findings support a role for Ca^{2+} underlying the actions of HMG in the homeostasis of the cytoskeleton associated phosphorylating system.

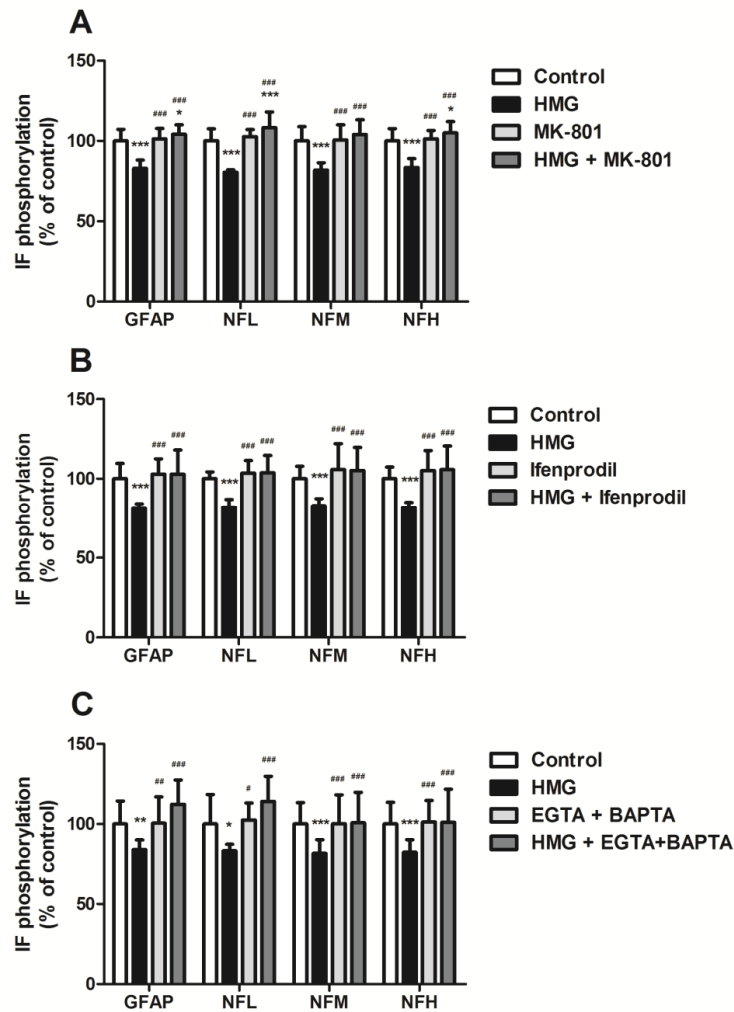


Figure 6 – Effects of a synaptic and extra-synaptic NMDARs and Ca^{2+} levels on the effect of HMG on the phosphorylation level of IF proteins from striatal slices of 30-day-old rats. Tissue slices were co-incubated during 60 min in the presence or absence (control group) of the metabolite (5 mM) plus the synaptic NMDAR antagonist MK-801 (A; 10 μM), the extra-synaptic NMDAR inhibitor (NR2B subunit) ifenprodil (B; 10 μM) or the intra and extra-cellular Ca^{2+} chelators BAPTA-AM, (50 μM) and EGTA, (C; 1 mM), respectively. Data are reported as mean \pm SD of 18 animals in each group and expressed as percentage of controls. Statistically significant differences from controls were determined by one-way ANOVA followed by Duncan's multiple range test. * $p < 0.05$, *** $p < 0.001$ compared with control; ### $p < 0.001$ compared with HMG.

Next, we evaluated the role of reactive species in the hypophosphorylating effects of HMG. We observed that the nitric oxide synthase (NOS) inhibitor L-NAME (1 mM) totally prevented the hypophosphorylation of GFAP and NF subunits (Figure

7A), suggesting a role for reactive nitrogen species in such effect. TROLOX (5 μ M), a soluble polar vitamin E analog, also normalized the phosphorylating level of GFAP and NF subunits (Figure 7B), suggesting a role for peroxy radicals in the HMG action. In contrast, N-acetyl cysteine (NAC) (1 mM), a GSH precursor, failed to prevent HMG-elicited IF hypophosphorylation (Figure 7C).

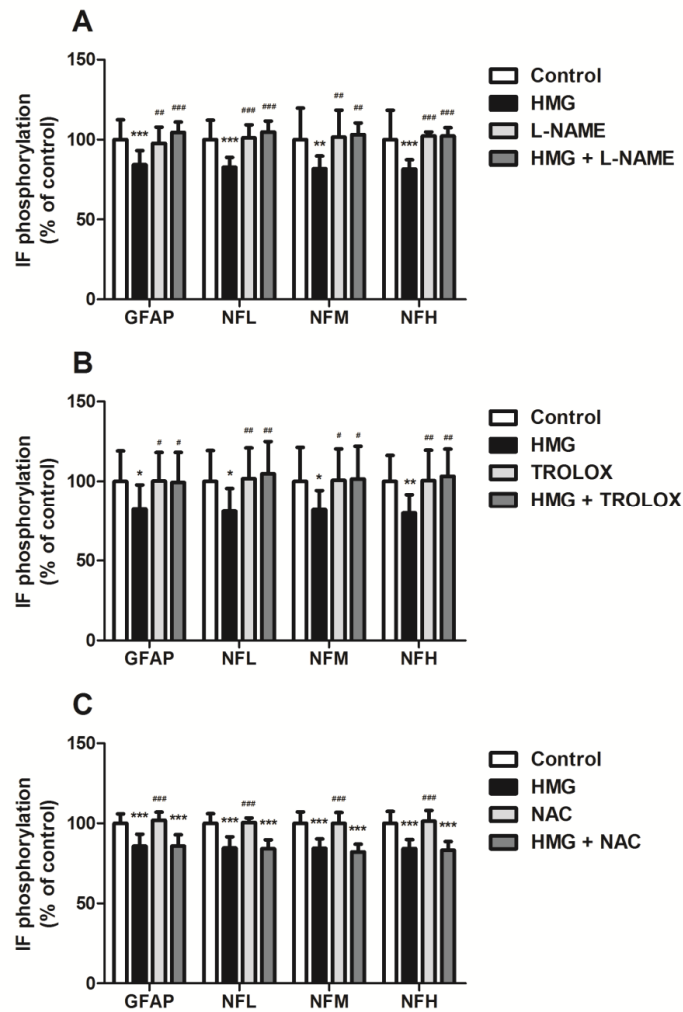


Figure 7 – Effects of L-NAME, TROLOX and NAC on HMG-induced alteration on the phosphorylation levels of intermediate filaments subunits (IFs) in the cytoskeletal fraction from striatal slices of 30-day-old rats. Tissue slices were co-incubated during 60 min in the presence or absence (control group) of the metabolite (5 mM) plus L-NAME, a nitric oxide synthase inhibitor (A;1 mM), TROLOX, a vitamin E hydrosoluble analogue (B;5 μ M) or NAC, a GSH precursor (C;1 mM). Data are reported as mean \pm SD of 18 animals in each group and expressed as percentage of controls. Statistically significant differences were determined by one-way ANOVA followed by Duncan's multiple range test. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ compared with control; # $p < 0.05$, ## $p < 0.01$, ### $p < 0.001$ compared with HMG.

Finally, we searched for whether L-NAME (1 mM) and TROLOX (5 μ M) could prevent the inhibition of PKA and JNK activities provoked by HMG. Thus, striatal slices were co-incubated with HMG and TROLOX or L-NAME and the immunocontent

of the active enzyme subunits was quantified. Western blot analysis showed that TROLOX and L-NAME restored the immunocontent levels of the active form of PKA (Figure 8A and B) and JNK (Figure 8C and D), strengthening a role for oxidative/nitrosative stress on the altered phosphorylating system associated with the IF cytoskeleton of astrocytes and neurons in striatal slices treated with HMG.

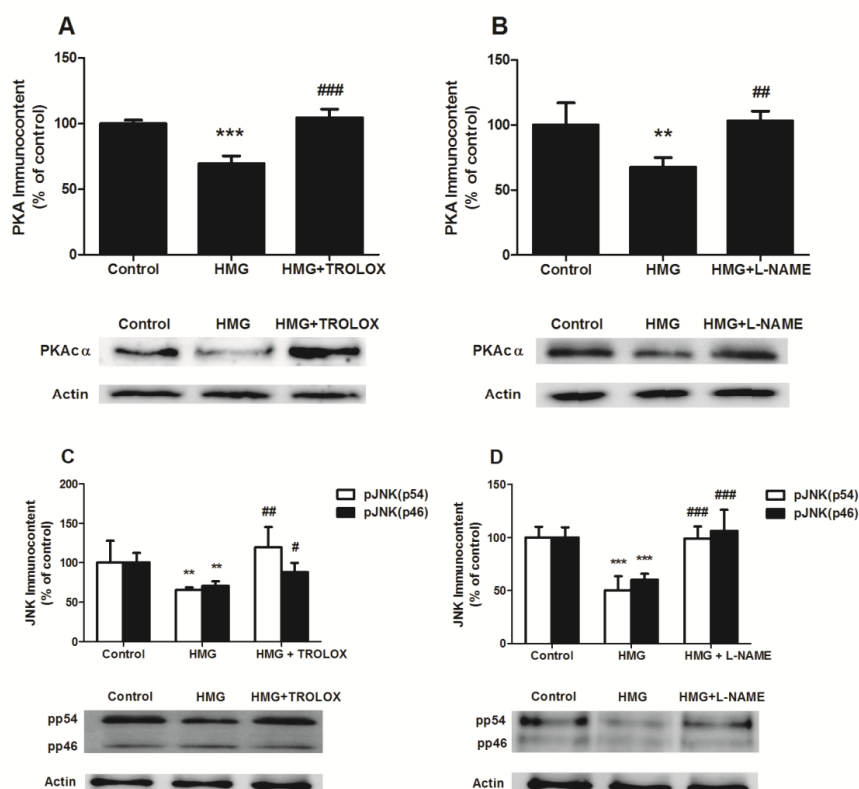


Figure 8 – Effects of TROLOX and L-NAME on PKA α and JNK immunocontent in the total homogenate of striatal slices of 30-day-old rats. Tissue slices were incubated during 60 min in the presence or absence (control group) of the metabolite (5 mM) plus TROLOX (5 μ M) or L-NAME (1 mM). Western blot with anti PKA α (A and B); anti-pJNK (p54), anti-pJNK (p46) (C and D). Data are reported as mean \pm SD of 6 animals in each group and expressed as percentage of controls. Statistically significant differences were determined by one-way ANOVA followed by Duncan's multiple range test. ** $p < 0.01$, *** $p < 0.001$ compared with control; # $p < 0.05$, ## $p < 0.01$, ### $p < 0.001$ compared with HMG.

Discussion

In the present report, we described that HMG, MGA and MGT, at concentrations found to induce mitochondrial redox homeostasis dysfunction (Leipnitz et al, 2008a,b, 2009), were able to disrupt the cytoskeleton phosphorylating system of neural cells in striatal and cortical slices of 30-day-old rats. We initially demonstrated that these organic acids that accumulate in HMGA cause hypophosphorylation of GFAP and NF subunits (NFL, NFM and NFH). The data indicate for the first time that the phosphorylating system associated with the IF-enriched cytoskeleton of astrocytes and neurons is vulnerable to the effects of HMG, MGA and MGT and may contribute to the pathophysiological mechanisms of brain damage in this disease. It was also found that the hypophosphorylation induced by HMG was not due to energy deprivation, since important parameters of mitochondrial energy production were not altered by this compound. Similarly, HMG-elicited decrease of phosphorylated IF could not be attributed to activated protein phosphatases, since specific inhibitors of the most important Ser-Thr phosphatases known to be associated with these cytoskeleton (PP1, PP2A and PP2B) failed to prevent this effect. More interesting, we found that the hypophosphorylation provoked by HMG was mediated by down-regulated PKA and JNK/MAPK activities. The antioxidants TROLOX and the NOS inhibitor L-NAME normalized the phosphorylation levels of IFs and prevented the inhibition of PKA and JNK provoked by HMG in striatum, strongly indicating that oxidative and nitrosative damage of these protein kinases were responsible for the alterations observed. It is also conceivable that synaptic and extra-synaptic NMDA receptors may have also participated in these effects since the non-competitive NMDA antagonists MK-801 and ifenprodil similarly restored the phosphorylated levels of IFs (Hardingham et al., 2002; Zhou et al., 2013a).

It is of note that striatum was used for elucidating the underlying mechanism of HMG, which most accumulates in HMGA, because progressive injury to the basal ganglia and usually found in patients with HMGA, attesting the vulnerability of this brain region. We also emphasize that leucoencephalopathy (cortical atrophy) is also detected in brain imaging of these individuals (van der Knaap et al., 1998; Yalçinkaya et al., 1999; Yılmaz et al., 2006). Furthermore, we have chosen to focus on the phosphorylating system associated with the IF cytoskeleton since these proteins are crucial for cell function, besides participating in the regulation of signaling pathways, axonal caliber and their dynamics is mostly regulated by phosphorylation (Sihag et al., 2007). Besides, several evidences point to a role for the IF cytoskeletal proteins in the cell response to an insult through alterations in the homeostasis of the phosphorylating system of these proteins (hyper and hypophosphorylation) observed in animal models of neurometabolic diseases (Pierozan et al., 2014; Loureiro et al., 2010a,b; Funchal et al., 2005a,b).

It is well known that hypophosphorylated NFs are more susceptible to proteolytic breakdown and accumulation of hypophosphorylated NFs into the axons could disturb the axonal transport of organelles and intracellular compounds (Goldstein et al., 1987; Pant, 1988). In line with this, hypophosphorylated NF proteins have been associated with the neurotoxicity of quinolinic acid and tellurium and the consequent cell damage (Heimfarth et al., 2012a,b, 2013a,b; Pierozan et al., 2014). Also, IF hypophosphorylation in C6 glioma cells exposed to high homocysteine levels has been associated with actin reorganization and altered cell morphology (Loureiro et al., 2010a,b). Therefore, the effects observed in the present work let us suppose that *in vitro* changes on the phosphorylating system associated with the cytoskeleton can be a predictor of the potential *in vivo* neurotoxicity of HMG.

In our findings, decreased PKA activity was reflected by the hypophosphorylation of Ser55 in the amino-terminal head domain of NFL, which is implicated in the regulation of the polymerization/depolymerization dynamic of these proteins (Grant and Pant, 2000). In addition, decreased JNK activity was supported by hypophosphorylated Lys-Ser-Pro (KSP) repeats of NFM and NFH tail domain regions (Giasson and Mushynski, 1997). It is stressed that misregulated phosphorylation levels in NFM and NFH has long been considered to be associated with abnormal axonal transport rate and disrupted stability of mature axons (Sihag et al., 2007).

The relevance of NMDA-mediated actions of HMG on the phosphorylating mechanism associated with the cytoskeleton was also supported by Ca^{2+} buffering, which totally prevented IF hypophosphorylation. Our results corroborate previous evidence that induction of oxidative stress by HMG in intrastrially injected rats is mediated by NMDA receptors (Fernandes et al., 2013). On the other hand, it is important to note that neither PKCaMII nor PP2B (two calcium-dependent enzymes) activities were altered in response to HMG, suggesting that IF hypophosphorylation downstream of NMDA was not dependent exclusively on Ca^{2+} levels on the kinase/phosphatase activities, pointing to more complex mechanisms directed to the phosphorylating system.

Taking into account that activated GluN2BR gives rise to inhibition of cAMP/PKA/CREB signaling pathway (Corcoran et al., 2013) and that ifenprodil, a blocker of extra-synaptic NR2B subunit-containing NMDARs (Zhou et al., 2013a), prevented HMG-induced IF hypophosphorylation, we propose that PKA down-regulation could be, at least partially, ascribed to NR2B-mediated mechanisms of PKA inhibition. Our findings showing that IF proteins can be a target of NR2B actions are in line with the previously described role of NR2B in the regulation of hippocampal actin

cytoskeleton (Akashi et al., 2009). Also, GluN2B-containing NMDARs have been shown to have a role in the deleterious actions of oligomers of amyloid-beta peptide 1-42 in microtubule dynamics in hippocampal cultures, suggesting that NR2B could be implicated in cytoskeletal disruption and neuronal dysfunction (Mota et al., 2012). Interestingly, 3-hydroxyglutaric acid, a structurally HMG similar compound accumulated in glutaric academia type I, was shown to cause neuronal death by overactivation of the NR2B subunit of NMDA glutamate receptors (Kolker et al., 2000a,b), reinforcing the excitotoxic role of NR2B on brain metabolic pathology.

Regarding to the preventive effect of L-NAME on PKA and JNK activities, this is in line with the ability of NO to inhibit JNK via S-nitrosylation (Park et al., 2000). We could also presume that down-regulated JNK/MAPK may occur via co-activation of synaptic and extra-synaptic GluRs with induction of nNOS (Courtney et al., 2014; Zhou et al., 2013b). The observations that TROLOX, an effective scavenger of peroxy radicals, normalized the reduction of protein kinases and the phosphorylating levels of IF provoked by HMG, may be tentatively associated with lipid peroxidation inducing these effects. This is in line with the previously reported modulation of the cytoskeleton in conditions of redox imbalance in different cell types and stress conditions (Loureiro et al., 2010a,b, 2013; Zamoner et al., 2008a,b).

We here demonstrated solid evidence for neurotoxic effects of the principal organic acids accumulating in HMGA disrupting cytoskeletal phosphorylating system with possible consequences for cellular function. We also found that antioxidants prevented these effects directed towards protein kinases, which is in accordance with previous findings showing that these metabolites disturb redox homeostasis (Leipnitz et al., 2008a,b, 2009; Fernandes et al., 2013; da Rosa et al., 2013). Therefore, it is conceivable that, besides hypoketotic hypoglycemia or hyperammonemia, the brain

damage in HMGA may be also a consequence of cytoskeleton dysregulation caused by high toxic levels of these organic acids.

In conclusion, this is the first report demonstrating that the disrupted homeostasis of the phosphorylating system associated with the cytoskeleton is caused by the major metabolites accumulating in HMGA. The evidence of a link between NMDA receptors, redox imbalance, misregulation of cell signaling mechanisms and hypophosphorylation of IF in response to HMG point to a critical role of the signaling pathways regulating the IF phosphorylation in the events of HMG-induced toxicity. Since disorganization of the cytoskeleton may be directly involved in several neurodegenerative disorders, we are tempted to propose that therapeutic strategies to regulate NMDA receptors and oxidative/nitrosative stress might serve as an adjuvant therapy for the protection of cells from the compounds that are found at high concentrations in this disease.

Conflicts of interest

The authors declare that there are no conflicts of interest associated with this manuscript.

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PARTE III

Discussão e Conclusões

III DISCUSSÃO

A HMGA é um distúrbio genético de caráter autossômico recessivo com prevalência estimada de 1 para cada 100.000 nascimentos (PIÉ *et al.*, 2007; REIMÃO *et al.*, 2009). É causada pela deficiência da HL, resultando em um acúmulo tecidual e uma alta excreção urinária dos ácidos 3-hidroxi-3-metilglutárico (HMG), 3-metilglutárico (MGA) e 3-metilglutacônico (MGT), bem como em menor grau dos ácidos 3-hidroxi-isovalérico e 3-metilcrotonilglicina nos indivíduos afetados (BONAFÉ *et al.*, 2000; SWEETAMN e WILLIAMS, 2001; PIÉ *et al.*, 2007). A apresentação clínica geralmente ocorre durante o primeiro ano de vida, usualmente após um período de jejum prolongado ou outro evento catabólico. Os episódios agudos são comuns e são caracterizados por vômitos, diarreia, desidratação, hipotonia, hipotermia, letargia, coma e apnéia, bem como por acidose metabólica e hipoglicemia hipocetótica. Outros sinais incluem macrocefalia, atraso no desenvolvimento, hepatomegalia com alterações no exame de fígado, pancreatite aguda e cardiomiopatia dilatada (SWEETMAN e WILLIAMS, 2001; GIBSON *et al.*, 1994; FUNGHINI *et al.*, 2001; VARGAS *et al.*, 2007; ZAFEIRIOU *et al.*, 2007). Esta doença é fatal em aproximadamente 20% dos casos, embora os sintomas tendem a ser mais leves depois da infância (THOMPSON *et al.*, 1990; PIÉ *et al.*, 2003).

Praticamente todos os pacientes apresentam sintomas neurológicos graves e anormalidades no córtex cerebral e gânglios da base (YALÇINKAYA *et al.*, 1999; YÍLMAZ *et al.*, 2006; ZAFEIRIOU *et al.*, 2007) cuja fisiopatogenia é pouco conhecida. Entretanto, podemos presumir que o acúmulo crônico desses metabólitos pode representar, ao menos em parte, um mecanismo

importante que contribui para os sintomas e anormalidades cerebrais observadas nestes pacientes, especialmente durante episódios de descompensação metabólica, em que as concentrações dos ácidos orgânicos aumentam drasticamente. Portanto, no presente estudo investigamos os efeitos dos principais metabólitos acumulados na HMGA sobre importantes parâmetros de estresse oxidativo, sobre a resposta inflamatória, bem como sobre a fosforilação de proteínas do citoesqueleto no estriado e córtex cerebral de ratos, com o objetivo de esclarecer a relação entre as características clínicas e as alterações bioquímicas nessa doença. É digno de nota que o estriado foi utilizado porque nessa doença ocorre lesão progressiva nos gânglios da base (estriado nos roedores), atestando a vulnerabilidade dessa região do cérebro. Ressaltamos, ainda, que a leucoencefalopatia (atrofia cortical) também é comumente detectada em imagens do cérebro desses indivíduos, justificando o uso do córtex cerebral neste estudo (van der KNAAP *et al.*, 1998; YALÇINKAYA *et al.*, 1999; YÝLMAZ *et al.*, 2006).

Como já mencionado, os mecanismos subjacentes da patogênese das lesões cerebrais na HMGA são pouco conhecidos. No entanto, tem sido sugerido que além do acúmulo intracelular de ácidos orgânicos tóxicos, hipoglicemia associada com déficit de corpos cetônicos, deficiência de carnitina podem também estar envolvidos na patofisiologia da doença (KAHLER *et al.*, 1994; MITCHELL e FUKAO, 2001; LEUNG *et al.*, 2009). Neste contexto, estudos *in vitro* anteriores demonstraram que o HMG e o MGA induzem dano oxidativo lipídico e protéico e reduzem as defesas antioxidantes não enzimáticas no córtex cerebral, no estriado e no fígado de ratos jovens (LEIPNITZ *et al.*, 2008a,b e 2009). Mostrou-se também que o MGA induz

disfunção mitocondrial e inibe a atividade da Na⁺, K⁺-ATPase em sinaptossomas preparados a partir de cérebro de ratos (RIBEIRO *et al.*, 2011).

A partir desses achados iniciamos nossa investigação estudando os efeitos *ex vivo* da administração intrastriatal de HMG e MGA sobre parâmetros de estresse oxidativo em estriado de ratos em desenvolvimento (Capítulo 1). Avaliamos se altas concentrações de HMG e MGA no estriado de ratos jovens obtidas por uma injeção intrastriatal aguda desses ácidos orgânicos poderiam afetar a homeostase redox na esperança de esclarecer melhor os mecanismos patológicos envolvidos nas anormalidades do estriado, uma estrutura cerebral alterada no cérebro dos pacientes afetados (van der KNAAP *et al.*, 1998; YALÇINKAYA *et al.*, 1999; YÝLMAZ *et al.*, 2006). Para isso, foram determinados parâmetros importantes de estresse oxidativo, incluindo as concentrações de MDA (peroxidação lipídica), a formação de carbonilas (dano oxidativo protéico), a produção de espécies reativas (oxidação da DCFH e produção de óxido nítrico), as concentrações de GSH (defesas antioxidantes não enzimáticas) e as atividades das enzimas antioxidantes SOD, GR, G6PD, GPx e CAT.

Nossos resultados demonstram que o HMG e o MGA induziram dano oxidativo a lipídios e a proteínas, bem como aumentaram a produção de espécies reativas de oxigênio, enquanto o HMG também induziu a produção de espécies reativas de nitrogênio. Em relação às defesas antioxidantes, ambos os metabólitos diminuíram os níveis de GSH e causaram alterações em várias atividades enzimáticas, provocando um desequilíbrio nesses sistemas de defesa. Finalmente observou-se que os antioxidantes melatonina, N-

acetilcisteína e a combinação de vitaminas C e E preveniram completamente ou atenuaram as alterações dos parâmetros de estresse oxidativo causadas pelo HMG, reforçando a participação de espécies reativas, mais particularmente os radicais hidroxila e peroxila que são sequestrados por estes antioxidantes (REITER *et al.*, 2001; ANISIMOV, 2006; HALLIWELL e GUTERIDGE 2007). Verificamos também que o MK-801, um antagonista não-competitivo do receptor glutamatérgico do tipo NMDA, impediu alguns dos efeitos provocados pelo HMG, indicando o envolvimento dos receptores NMDA nos efeitos encontrados. O fato de receptores NMDA estarem envolvidos nos efeitos induzidos por HMG não é surpreendente, devido à semelhança estrutural entre a HMG e glutamato. Portanto, pode ser de interesse estudar se os ácidos orgânicos acumulados e, especialmente, o HMG interagem diretamente com receptores NMDA. Por outro lado, não se pode excluir a possibilidade de o aumento na produção das ERN pode ser mediada por uma indução da atividade da óxido nítrico sintase, que conduz à geração dessas espécies foi, pelo menos, parcialmente devido a supraestimulação dos receptores NMDA por este ácido orgânico (HALLIWELL e GUTTERIDGE, 2007; BROWN, 2010; AYATA *et al.*, 1997).

Nossos resultados indicam fortemente que o HMG e o MGA provocam estresse oxidativo *in vivo* em estriado de ratos jovens. Devemos salientar também que as espécies reativas são capazes de oxidar moléculas diferentes, incluindo lípidos, proteínas, açúcares e DNA, o que representa um evento chave na cascata que conduz à patogenicidade da morte celular necrótica ou apoptótica (KROEMER e REED, 2000; NIZUMA *et al.*, 2009; CIRCU e AW, 2010). Além disso, o estresse oxidativo é uma condição muito prejudicial,

especialmente para o cérebro em comparação com outros tecidos, devido ao fato de o cérebro possuir uma alta taxa de metabolismo oxidativo acoplado a uma grande produção de ERO, uma atividade reduzida das enzimas antioxidantes, menores concentrações de antioxidantes e maior potencial de peroxidação, devido ao seu elevado teor de PUFA (HALLIWELL e GUTTERIDGE, 2007).

A partir do conhecimento de que o cérebro é bastante suscetível ao estresse oxidativo causado pelo HMG e pelo MGA fomos investigar se os astrócitos são células alvo desses metabólitos, já que essas células neurais são fundamentais para o funcionamento do sistema nervoso central e susceptíveis a danos por neurotoxinas, além de serem importantes fontes de produção de espécies reativas. Para tal determinamos os efeitos do HMG e MGA sobre importantes parâmetros da homeostase redox, produção de citocinas, bem como sobre a função mitocondrial e a viabilidade de astrócitos cultivados obtidos de córtex cerebral de ratos neonatos (Capítulo 2). Descobrimos que ambos os ácidos orgânicos reduziram a função mitocondrial astrocitária, sem alterar a viabilidade celular, reduzindo também as concentrações de GSH. Em contrapartida, os metabólitos aumentaram a formação de espécies reativas verificada por um aumento na oxidação da DCFH e provocaram um aumento significativo das citocinas pró-inflamatórias IL-1 β , IL-6 e TNF α através da via de sinalização da Erk.

Assim, reforçando nossos resultados encontrados no capítulo 1, verificamos que o HMG e o MGA provocaram um aumento na produção de espécies reativas. Uma vez que a oxidação da DCFH ocorre principalmente por

meio de ERO e é utilizada como uma sonda para a determinação de particularmente ERO em sistemas biológicos, concluiu-se que estes ácidos orgânicos provocaram a geração de espécies reativas em um sistema de células intactas (astrócitos), corroborando com resultados anteriores (LEIPNITZ *et al.*, 2008a,b e 2009; RIBEIRO *et al.*, 2011). Não podemos, no entanto excluir a participação de ERN nesse modelo, visto que a oxidação da DCFH pode também ocorrer através dessas espécies, assim como citado no capítulo 1 deste trabalho. Tais resultados indicam que os astrócitos expostos aos principais metabólitos que se acumulam na HMGA estão envolvidos na produção de radicais livres.

Descobrimos também que estes ácidos orgânicos diminuíram marcadamente as concentrações de GSH. Nesse contexto, os astrócitos são considerados uma das principais fontes de GSH por sintetizar e secretar esse antioxidante, ajudando, assim, a manutenção do estado redox neuronal (DRINGEN, 2000; POPE *et al.*, 2008). GSH é o antioxidante mais abundante no cérebro, exercendo um papel fundamental para eliminar as ERO, proteger os grupamentos sulfidríla das proteínas no estado redox adequado, regulando também a morte celular e as cascatas de sobrevivência celular (SARAFIAN *et al.*, 1996).

Uma vez que níveis adequados de antioxidantes são essenciais para proteger as células contra os danos oxidativos (CASTOLDI *et al.*, 2001), é possível que a redução significativa da GSH induzida pelos metabólitos que se acumulam na HMGA pode causar perda da funcionalidade de astrócitos expostos ao HMG e ao MGA, como observado pela redução significativa de

MTT. Salienta-se que o ensaio MTT é comumente utilizado para avaliar a funcionalidade mitocondrial, como um indicador do estado energético das células refletindo a atividade de desidrogenases, embora também seja aceito como um marcador de viabilidade celular. Uma vez que a incorporação de iodeto de propídio não foi alterado em astrócitos em cultura expostos ao HMG e ao MGA nas mesmas condições, conclui-se que a viabilidade celular foi preservada (TSAI e LEE, 1994; ALARCÓN-AGUILAR *et al.*, 2014).

Não podemos estabelecer no presente se a redução dos níveis de GSH ocorreu devido à sua oxidação por radicais livres devido a uma alta produção de espécies reativas (aumento da oxidação DCFH) causada pelos ácidos orgânicos que leva ao consumo de GSH, ou se a redução dos níveis de GSH levou a um desequilíbrio na formação de espécies reativas levando a um estado pró-oxidante. No entanto, enfatiza-se que a depleção de GSH ocorre durante condições de estresse que podem ser consequência de uma resposta inflamatória com estresse oxidativo, desempenhando portanto um papel crítico na resposta inflamatória astrocítica (TANABE *et al.*, 2010; LEE *et al.*, 2010).

Nesse particular, uma contribuição original de nossa investigação (capítulo 2) foi a demonstração de níveis elevados de citocinas IL-1 β , IL-6 e TNF α em astrócitos cultivados expostos a HMG e MGA. TNF α e IL-1 β são sintetizadas predominantemente por microglia e astrócitos na resposta inflamatória aguda. Essas citocinas também podem induzir a síntese de IL-6. A IL-6 é produzida em microglia, astrócitos e neurônios e desempenha um papel central numa variedade de funções do sistema nervoso central, tais como a indução e a modulação de ativação de astrócitos, inflamação e neuroproteção

(TANABE *et al.*, 2010; LEE *et al.*, 2010; SANTELLO *et al.*, 2011; ERTA *et al.*, 2012; YE *et al.*, 2013). Além disso, a elevação dos níveis de citocinas provavelmente reflete a reatividade da glia que, em combinação com um aumento da produção de ER e um decréscimo no teor de antioxidantes no cérebro, foram descritas como envolvidas na patogênese e no desenvolvimento de várias doenças humanas e na neurodegeneração (DARLINGTON, 2005; KANWAR e NEHRU, 2007).

Assim, verificamos que o HMG e o MGA induziram as citocinas pró-inflamatórias clássicas (TNF α e IL1- β) dependentes da via de sinalização Erk. No que diz respeito ao IL-6, esta citocina apresenta uma dualidade de ações, degenerativas e reparadoras, anti- e pró-inflamatórias (FARINA *et al.*, 2007). Neste particular, é de se notar que a via Erk tem sido implicada na regulação da resposta inflamatória glial após um insulto (dano) e é uma via de transdução de NF κ B (LEE *et al.*, 2010; SANTELLO *et al.*, 2011; ERTA *et al.*, 2012; YE *et al.*, 2013; DARLINGTON, 2005; KANWAR e NEHRU, 2007; BRENNER, 2014; BOBERMIN *et al.*, 2012). NF κ B é considerado o principal mediador inflamatório no SNC, embora o seu mecanismo de ativação e subsequente translocação para o núcleo não são completamente compreendidos (GLOIRE *et al.*, 2006; WAKABAYASHI *et al.*, 2010). Por outro lado, o aumento de IL-6 não foi associado com a via da Erk, mas pode ter sido causado alternativamente pelo aumento de TNF α e IL1- β , indutores de IL-6 (TANABE *et al.*, 2010; TANABE *et al.*, 2011).

Os resultados obtidos no capítulo 2 indicam uma susceptibilidade de astrócitos corticais de ratos neonatos aos insultos tóxicos do HMG e do MGA,

associado a um aumento da resposta pró-inflamatória com aumento da produção de citocinas e de estresse oxidativo acompanhada por uma disfunção mitocondrial. Esses resultados confirmam nossos achados de indução de estresse oxidativo em estriado *in vivo* por esses compostos, bem como achados anteriores *in vitro* realizados em frações subcelulares e homogeneizados de cérebro (LEIPNITZ *et al.*, 2008a,b; 2009) e indicam que a perturbação da homeostase redox astrocitária causada pelos principais ácidos orgânicos que se acumulam na HMGA pode potencialmente contribuir para a neuropatologia de pacientes afetados por esta doença. Além disso, nossos dados indicam que astrócitos em cultura representam uma ferramenta biológica interessante e promissora para estudar a toxicidade dos compostos que se acumulam em outras doenças hereditárias metabólicas, embora a sua principal limitação dos achados presentes reside no fato de que não envolve as relações complexas entre neurônios e células gliais. Presume-se, portanto, que a produção de espécies reativas e citocinas contribuem para a disfunção neurológica observada nesta doença, particularmente durante as crises metabólicas, em que a produção e o acúmulo dos metabólitos no cérebro provavelmente é ainda maior. No entanto, não se pode descartar que a hiperamonemia e a privação de energia (hipoglicemia hipocetótica) também podem estar envolvidas na lesão cerebral desses pacientes uma vez que estas condições também podem resultar na formação de radicais livres no SNC (BOBERMIN *et al.*, 2012; SINGH *et al.*, 2004; SUH *et al.*, 2007).

Finalmente investigamos os efeitos do HMG, MGA e MGT sobre a fosforilação da GFAP e das subunidades dos NF (NFL, NFM e NFH) do citoesqueleto em fatias de estriado e córtex cerebral de ratos jovens (Capítulo

3). Optamos por focar nosso estudo no sistema de fosforilação associada aos FI do citoesqueleto, porque essas proteínas são cruciais para a morfologia celular, além de participarem da regulação de vias de sinalização, na formação e manutenção do calibre axonal e também porque essas dinâmicas são principalmente reguladas por fosforilação (SIHAG *et al.*, 2007) . Além disso, várias evidências apontam para um papel dos FI na resposta a um insulto celular através de alterações na homeostase do sistema de fosforilação dessas proteínas (hiper e hipofosforilação) observada em modelos animais de doenças neurometabólicas (PIEROZAN *et al.*, 2014; LOUREIRO *et al.*, 2010a,b; FUNCHAL *et al.*, 2005a,b).

Resumidamente, nossos resultados obtidos no capítulo 3 foram que o HMG, o MGA e o MGT provocaram uma hipofosforilação da GFAP e das subunidades dos NF nas duas estruturas cerebrais testadas e esse efeito foi mediado pela inibição das proteínas cinases PKA e JNK, sem a participação das proteínas fosfatases. Observou-se também que a hipofosforilação do citoesqueleto era dependente dos receptores do tipo NMDA sinápticos e extrasinápticos, bem como do íon Ca^{2+} . Além disso, os antioxidantes TROLOX e L-NAME preveniram completamente a hipofosforilação e a inibição das atividades da PKA e da JNK causadas pelo HMG.

Esses dados indicam que o sistema de fosforilação do citoesqueleto associado com os FI de astrócitos e neurônios é vulnerável aos efeitos do HMG, do MGA e do MGT, podendo contribuir para explicar os mecanismos patofisiológicos de dano cerebral dos pacientes com a deficiência da HL. Verificou-se também que o hipofosforilação induzida pelo HMG não foi devido

ao déficit energético (o ATP é necessário para que haja a correta fosforilação de proteínas), uma vez que parâmetros importantes da produção de energia não foram alteradas pelo HMG. Também demonstramos que a diminuição da fosforilação dessas proteínas não poderia ser atribuída a ativação de proteínas fosfatases, uma vez que os inibidores específicos Ser-Thr das fosfatases mais importantes associados com o citoesqueleto (PP1, PP2A e PP2B) não conseguiram evitar este efeito. Contudo, descobrimos que a hipofosforilação provocada pelo HMG foi mediada pela inibição das proteínas cinases PKA e JNK. Além disso, observamos que os antioxidantes TROLOX e o inibidor da óxido nítrico sintase (NOS), L-NAME, preveniram totalmente a hipofosforilação dos FI e a inibição da atividade da PKA e da JNK provocadas pelo HMG no estriado, indicando fortemente que ERO e ERN poderiam ter provocado dano oxidativo a essas proteínas cinases e serem responsáveis pelas alterações de fosforilação observadas. Também é concebível que os receptores NMDA sinápticos e extra-sinápticos também podem ter participado nesses efeitos uma vez que os antagonistas de NMDA não competitivo, MK-801, e seletivo da subunidade NR2B, ifenprodil, normalizaram os níveis de fosforilação dos FI (HARDINGHAM *et al.*, 2002; ZHOU *et al.*, 2013a,b).

Para reforçar a participação das cinases nos efeitos causados por HMG verificamos que a diminuição da atividade da PKA foi refletida pela hipofosforilação da Ser55 no domínio aminoterminal da NFL que está implicado na regulação da dinâmica de polimerização / despolimerização dessas proteínas (GRANT e PANT, 2000). Além disso, a diminuição da atividade da JNK refletiu-se na hipofosforilação das repetições do tipo KSP nos domínios carboxiterminal do NFH e do NFN (GIASSON e MUSHYNSKI, 1997). Ressalte-

se que os níveis de fosforilação do NFM e do NFH tem sido associados com a velocidade de transporte axonal e a estabilidade dos axônios maduros (SIHAG *et al.*, 2007).

Os efeitos do HMG sobre o mecanismo de fosforilação associada ao citoesqueleto mediados por NMDA foi reforçada pela importância do Ca^{2+} nesses efeitos, já que o tamponamento e o sequestro desse íon totalmente preveniram essa hipofosforilação. Esses resultados corroboram com os resultados encontrados no capítulo 1 desta tese de que a indução de estresse oxidativo causado por injeção intracerebral de HMG é mediada por receptores glutamatérgicos do tipo NMDA.

Levando-se em conta que a ativação de receptores glutamatérgicos que contém a subunidade NR2B leva à inibição da sinalização CREB via AMPc / PKA (CORCORAN *et al.*, 2013) e que o ifenprodil, um bloqueador seletivo dessa subunidade extrasináptica (ZHOU *et al.*, 2013a,b) preveniu a hipofosforilação dos FI provocada por HMG, é possível que esse efeito tenha sido mediado pela inibição da PKA secundariamente à ativação dos receptores NMDA extrasinápticos. Nossos resultados mostram que proteínas do citoesqueleto podem ser alvo de ações indiretas ao estímulo dos NR2B, o que está de acordo com o papel descrito anteriormente desses receptores na regulação do citoesqueleto de actina no hipocampo (AKASHI *et al.*, 2009). Além disso, os NR2B têm demonstrado possuir um papel prejudicial na dinâmica dos microtúbulos em culturas de hipocampo com consequente disfunção neuronal (MOTA *et al.*, 2012). Curiosamente, o ácido 3-hidroxiglutárico, um composto estruturalmente semelhante ao HMG, que se acumula na acidemia

glutárica tipo I, foi demonstrado provocar morte neuronal por superativação da subunidade NR2B de receptores NMDA (KÖLKER *et al.*, 2000a,b), reforçando portanto o papel excitotóxico dessa subunidade NMDA no cérebro de doenças metabólicas.

Assim, no capítulo 3, demonstramos evidências de que os principais ácidos orgânicos que se acumulam na HMGA alteram o sistema de fosforilação do citoesqueleto com possíveis consequências para a forma e função celular. Descobrimos também que alguns antioxidantes impediram esses efeitos inibitórios em proteínas cinases, o que está de acordo com achados prévios (LEIPNITZ *et al.*, 2008a,b e 2009; da ROSA *et al.*, 2013), além de reforçar os resultados encontrados nos capítulos anteriores desta tese, mostrando que esses metabólitos perturbam a homeostase redox no cérebro. Portanto, é concebível que, além da hipoglicemia-hipocetótica e da hiperamonemia, os danos cerebrais apresentados pelos pacientes acometidos pela HMGA podem ser também uma consequência da desregulação no citoesqueleto causada por níveis tóxicos destes ácidos orgânicos, já que esses podem exercer sinergicamente um papel crítico nas vias de sinalização que regulam a fosforilação dos FI e bem como suas funções celulares.

Finalmente, é difícil de prever a importância fisiopatológica dos dados apresentados nesta investigação, pois as concentrações cerebrais de HMG, MGA e MGT ainda não foram estabelecidas no cérebro dos doentes com HMGA. No entanto, uma vez que os sintomas neurológicos aparecem ou pioram durante as crises catabólicas, em que as concentrações dos metabólitos acumulados aumentam drasticamente durante esses episódios, é

concebível que o acúmulo desses metabolitos possam estar relacionados com a piora clínica. Devemos também enfatizar que as concentrações dos ácidos orgânicos em geral, e em especial de ácidos dicarboxílicos (tais como a HMG, MGA e MGT), que se acumulam em várias acidemias orgânicas são mais elevados no cérebro, em comparação com o sangue, provavelmente devido ao baixo efluxo destes compostos a partir do cérebro, que está de acordo com a teoria de aprisionamento cerebral de ácidos dicarboxílicos (HOFFMANN *et al.*, 1993; HOFFMANN *et al.*, 1994; SAUER *et al.*, 2006; STELLMER *et al.*, 2007). E, uma vez que os resultados obtidos na presente tese podem estar diretamente envolvidos em várias doenças neurodegenerativas, somos tentados a propor que estratégias terapêuticas para regular os receptores NMDA, o estresse oxidativo / nitrosativo e os níveis de citocinas podem potencialmente servir como terapia adjuvante para a proteção do sistema nervoso central dos pacientes afetados por HMGA.

IV CONCLUSÕES

- A injeção intraestriatal dos ácidos HMG e MGA induziram a produção de espécies reativas de oxigênio e nitrogênio, a peroxidação lipídica, a oxidação protéica, diminuíram os níveis de GSH e desestabilizaram as defesas antioxidantes enzimáticas indicando que esses metabólitos induzem estresse oxidativo *ex vivo* em estriado de ratos jovens.
- A exposição de astrócitos cultivados de córtex cerebral de ratos neonatos ao HMG e ao MGA causa aumento na produção de espécies reativas, diminuição dos níveis de GSH, aumento na liberação de citocinas pró-inflamatórias mediado pela cascata da Erk e um desequilíbrio da função mitocondrial, configurando portanto, uma indução de estresse oxidativo e uma indução de resposta inflamatória nessas células neurais importantes para o funcionamento do sistema nervoso central.
- Os ácidos HMG, MGA e MGT provocaram hipofosforilação das proteínas GFAP e das subunidades NFL, NFM e NFH do citoesqueleto de estriado e de córtex cerebral de ratos jovens, mediada pela inibição das proteínas cinases PKA e JNK secundariamente ao estímulo dos receptores glutamatérgicos sinápticos e extrasinápticos bem como de dano oxidativo a essas proteínas.
- Em resumo, os principais ácidos orgânicos acumulados na HMGA provocam um desequilíbrio da homeostase redox e alteração do sistema fosforilante de proteínas do cicloesqueleto com o

envolvimento de receptores NMDA que potencialmente representam mecanismos patogênicos envolvidos no dano estriatal e cortical em pacientes afetados pela deficiência da HL.

V PERSPECTIVAS

- Estudar os efeitos da injeção intraestriatal dos ácidos HMG e MGA em ratos jovens sobre a fosforilação das proteínas do citoesqueleto no estriado.
- Estudar os efeitos dos ácidos HMG e MGA sobre a fosforilação das proteínas do citoesqueleto em culturas de astrócitos e neurônios de córtex cerebral e estriado de ratos.
- Fazer análise histopatológica com hematoxilina/eosina e imunohistoquímica para GFAP e NeuN em estriado de ratos jovens injetados com os ácidos HMG e MGA.
- Estudar a morfologia e a viabilidade celular de neurônios e astrócitos cultivados de córtex cerebral e estriado de ratos expostos aos ácidos HMG e MGA.

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