

HMGB1 基因调控胶质瘤恶性生物学行为研究进展

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【摘要】 胶质瘤因其复杂的生物学特性以及胶质瘤干细胞的干细胞特性,标准治疗后易复发。高迁移率族蛋白 1(HMGB1)作为癌基因通过多种不同方式参与调控胶质瘤恶性进展。本文对 HMGB1 基因与胶质瘤预后、肿瘤病理分级、干细胞相关性以及 HMGB1 基因调控胶质瘤方式如非编码 RNA 途径、细胞自噬途径、肿瘤微环境等进行综述,为胶质瘤的治疗提供参考。

【关键词】 神经胶质瘤; HMGB1 蛋白质; RNA,未翻译; 自噬; 肿瘤微环境; 综述

Progress on the regulation of malignant biological behavior of glioma by HMGB1 gene

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【Abstract】 Glioma is often recurred after standard treatment due to its complex biological characteristics and the presence of glioma stem cells (GSCs). High-mobility group box 1 (HMGB1) can regulate the malignant progression of glioma through a variety of ways as an oncogene. This article reviews the relationship between HMGB1 gene and the prognosis of glioma patients, tumor pathological grading, GSCs, and the ways including non coding RNA pathway, autophagy pathway, tumor microenvironment, etc., in which HMGB1 gene regulates glioma, so as to provide the reference for the treatment of glioma.

【Key words】 Glioma; HMGB1 protein; RNA, untranslated; Autophagy; Tumor microenvironment; Review

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胶质瘤是临床最常见且最致命的原发性脑肿瘤,

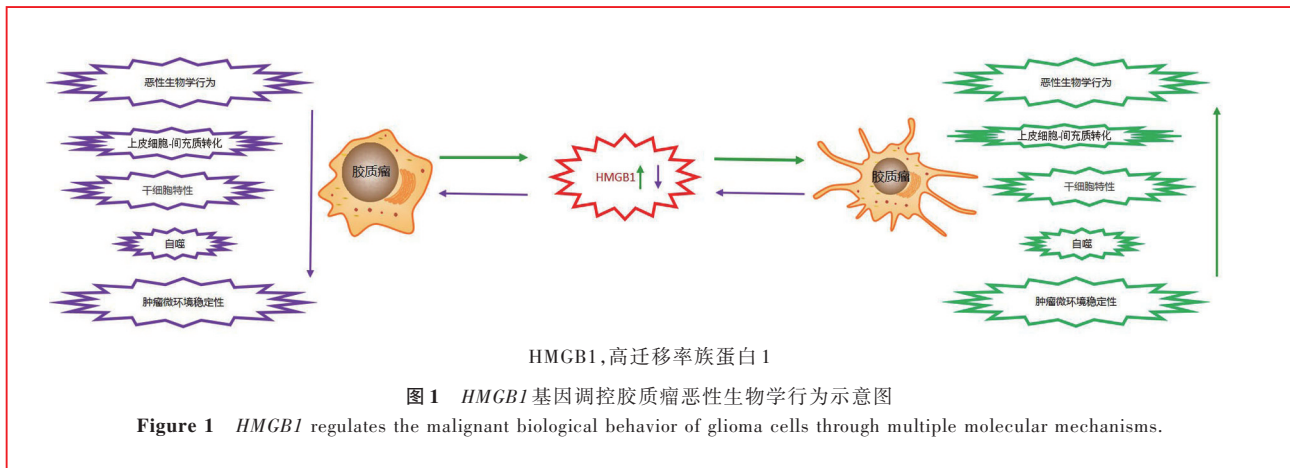
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予以手术、放疗和替莫唑胺化疗的标准治疗后仍不可避免复发,严重危及生命^[1-3]。胶质瘤的确切发病机制尚未阐明,随着分子病理学的发展,其恶性进展除增殖、侵袭、耐药、胶质瘤干细胞(GSCs)及微血管生成外,还与肿瘤微环境(TME)中独特的信息传递和不同类型的表观遗传学征象密切相关,如组蛋白修饰、DNA 甲基化、染色质重塑等^[4-7],成为胶质瘤治疗的新靶点。高迁移率族蛋白 1(HMGB1)是一种高度保守的染色质结合蛋白,可作为 DNA 伴侣发挥作用,亦是一种分泌蛋白,参与调控细胞损伤,在炎症反应和自身免疫性疾病的发生发展中发挥重要作用^[8-10]。近年有研究显示, HMGB1 基因在



多种肿瘤细胞增殖、浸润中起关键作用^[11-14],并经多种途径参与调控胶质瘤恶性生物学行为及进展(图1)^[15-17]。本文拟综述 *HMGB1* 基因在胶质瘤恶性进展中的作用,以为胶质瘤的治疗提供参考。

一、*HMGB1* 基因与胶质瘤预后

HMGB1 基因表达变化与胶质母细胞瘤患者生存和预后呈负相关。Kluckova 等^[18]检测胶质母细胞瘤患者血清 *HMGB1*、 $CD14^+TREM-1^+$ 和 $CD14^+TREM-2^+$ 细胞水平, Cox 回归显示, $CD14^+TREM-2^+$ 细胞比例越高、患者生存率越高,血清 *HMGB1* 水平与 $CD14^+TREM-1^+$ 细胞比例以及 $TREM-1/TREM-2$ 比值呈负相关。杨如意等^[19]对比分析 87 例胶质瘤患者术后肿瘤标本与 20 例颅内减压术后脑组织标本 *HMGB1* 及其受体 Toll 样受体 4 (TLR4) 表达变化,发现胶质瘤标本 *HMGB1* 和 TLR4 阳性表达率均高于颅内减压脑组织,且 *HMGB1* 水平与 TLR4 水平呈正相关,二者均随肿瘤级别的增加而升高,而患者生存期则随肿瘤级别的增加而缩短,预后较差。郭丹等^[20]的 Meta 分析亦得出上述结论。周少龙等^[21]检测胶质瘤周围组织 *HMGB1* 和 TLR4 表达量,其表达水平越高、患者患癫痫并发症的风险越高、预后越差。然而,*HMGB1* 与胶质瘤患者预后的相关性并未在胶质瘤标准诊断条件下进一步验证。

二、*HMGB1* 基因与胶质瘤恶性行为

外源性干预 *HMGB1* 基因可以调控胶质瘤恶性进展^[22-23]。李壮等^[24]选取肿瘤基因组学图谱计划 (TCGA) 数据库中 319 例样本的 *HMGB1* 基因测序数据进行差异表达及生存分析,发现 *HMGB1* mRNA 在肿瘤组织中显著表达上调,且随肿瘤分级 (WHO 分级) 的增加而升高,并与患者预后呈负相关;沉默

HMGB1 基因的胶质母细胞瘤细胞系增殖和侵袭能力下降,且对替莫唑胺的反应更敏感。黄俊强等^[25]将 *HMGB1* 基因小干扰 RNA (siRNA) 质粒在脂质体介导下转染 U373 和 U87 胶质瘤细胞,显示位于 S 期的细胞比例明显减少,提示 *HMGB1* 基因表达下调后细胞周期发生 S 期阻滞,导致胶质瘤细胞增殖、侵袭受到抑制。Zhang 等^[26]通过糖尿病小鼠构建胶质瘤模型,无论肿瘤植入何部位,高血糖小鼠 GL261 胶质瘤细胞生长体积均增加,并发现其作用机制为通过上调晚期糖基化终末产物受体 (RAGE) 及其配体 *HMGB1* 表达以及抑制抗肿瘤免疫反应,促进胶质瘤生长。王新军等^[27]的裸鼠成瘤实验采用 RAGE 抑制剂 FPS-ZM1 干预 LN229 胶质瘤细胞,发现肿瘤细胞 *HMGB1* 表达受到明显抑制,成瘤速度明显减慢。Li 等^[28]采用重组人 *HMGB1* 干预胶质母细胞瘤细胞系和原代细胞,发现 *HMGB1* 可通过糖原合成酶激酶-3 β (GSK3 β)/Snail 信号转导通路促进细胞上皮间质转化 (EMT),后者是胶质瘤恶性进展的重要因素,并通过激活 TLR4/P38/核因子 E2 相关因子 2 (Nrf2) 信号转导通路介导 P62 蛋白的过表达, *p62* 基因敲低则反向抑制 *HMGB1* 诱导的上皮间质转化。李卫玲等^[29]采用钾离子通道阻断剂 4-氨基吡啶、四乙胺和 ATP 敏感性钾离子通道阻断剂格列苯脲干预 U87 和 U251 胶质瘤细胞,发现肿瘤细胞迁移、侵袭均受到抑制,且 *HMGB1* 水平降低。李文涛等^[30]构建胶质瘤大鼠模型并进行体外研究,通过感染外源性病毒上调促凋亡基因 *TRAIL* 的表达,大鼠肿瘤细胞和原代肿瘤细胞 *HMGB1* 释放增加,他们还发现,即将凋亡的胶质瘤细胞诱发巨噬细胞释放 *HMGB1*,并通过 TLR4 刺激成熟树突状细胞 (DC) 免疫应答肿瘤抗原,形成一定的自我保护,而下调 *HMGB1* 表达可诱

发抗肿瘤免疫反应,使大鼠存活期缩短。早期生长反应蛋白1(EGR1)作为胶质母细胞瘤预后的生物标志物,与HMGB1协同作用可以更准确地预测预后,亦可通过诱导HMGB1表达以促进肿瘤细胞侵袭^[31]。肿瘤浸润中性粒细胞(TINs)产生的中性粒细胞胞外诱捕网(NETs)通过调节HMGB1/RAGE/白细胞介素-8(IL-8)信号转导通路以介导胶质瘤细胞与肿瘤微环境间的信号转导,靶向抑制NETs生成或IL-8分泌可能是阻止胶质瘤进展的有效途径^[32]。HMGB1亦可通过甲基化下调SASH1基因的表达,降低整合素 $\beta 8$ 水平,从而抑制胶质瘤细胞黏附,促进细胞迁移^[33]。

三、HMGB1基因与胶质瘤干细胞

GSCs的干细胞特性可导致胶质瘤手术后易复发^[34]。缺氧环境下,HMGB1在GSCs表达上调,沉默HMGB1基因可导致GSCs标志物丢失和自我更新能力减弱;此外,HMGB1基因敲除还可抑制RAGE依赖性细胞外信号调节激酶1/2(ERK1/2)信号转导通路激活,抑制GSCs细胞周期改变^[35]。GSCs被认为是胶质瘤对替莫唑胺耐药的主要原因之一,即替莫唑胺可以促进GSCs生成,同时HMGB1通过TLR2/NEAT1/Wnt信号转导通路促GSCs生成,HMGB1表达下调时替莫唑胺使GSCs生成受限^[36]。Zhang等^[37]对比分析放疗后GSCs和胶质瘤细胞凋亡率以及 γ -H2AX阳性细胞率,结果发现,经6 Gy放疗后GSCs仍有良好的成球能力,肿瘤细胞凋亡率和 γ -H2AX阳性细胞率均较低,而PCAT1在GSCs中表达水平升高,敲低PCAT1基因可抑制GSCs成球,加速肿瘤细胞凋亡和DNA损伤;此外,沉默PCAT1基因还可通过上调微小RNA-129-5p(miRNA-129-5p)、下调HMGB1抑制肿瘤细胞增殖。Zang等^[38]通过腺病毒感染A172和T98G胶质瘤细胞和人原代胶质瘤细胞,结果显示,腺病毒感染使干细胞性标志物水平升高,使被感染肿瘤球的自我更新能力和多谱系分化能力增强;腺病毒感染的肿瘤球在免疫缺陷小鼠中具有更强的异种移植瘤形成潜力;腺病毒通过诱导TLR9基因表达上调促进GSCs生成,TLR9基因表达下调则使GSCs成球能力下降;此外,髓样分化因子88(MyD88)、总信号转导与转录激活因子3(STAT3)、磷酸化STAT3和长链非编码RNA(lncRNA)NEAT1在腺病毒诱导的GSCs中亦表达上调,敲除MYD88基因或抑制STAT3、NEAT1基因可以减弱腺病毒诱导的GSCs干细胞特性。

四、HMGB1基因与胶质瘤非编码RNA

非编码RNA(ncRNA)是不参与蛋白质编码的RNA,主要以癌基因或抑癌基因方式在转录、RNA加工、翻译、表观遗传修饰和翻译后修饰中作为关键调控因子发挥作用^[39-40]。目前,胶质瘤领域研究最广泛的有lncRNA、miRNA和环状RNA(circRNA),其中lncRNA占比最高(>80%),较miRNA具有更大的变异性,且二者之间存在双向调控机制;circRNA高度保守,表达模式具有组织特异性和肿瘤分期依赖性,通常经海绵化miRNA调节基因转录和蛋白质合成^[41-42]。

1.长链非编码RNA lncRNA异常表达与胶质瘤发生发展密切相关。Chen等^[43]发现,LINC00665基因在胶质瘤中表达上调,其表达变化与肿瘤直径、WHO病理分级、MRI信号强化、瘤周水肿呈正相关,通过吸附miRNA-129-5p间接升高HMGB1的表达水平,促进胶质瘤恶性进展。Wu等^[44]发现LINC00662基因可直接与miRNA-107基因相互作用,而HMGB1基因又是miRNA-107基因的已知作用靶点;LINC00662基因在胶质瘤中过表达与晚期恶病质和不良预后相关,沉默LINC00662基因可抑制胶质瘤细胞增殖、侵袭及其在裸鼠体内的生长,HMGB1基因过表达(或miRNA-107基因抑制)可以逆转敲低LINC00662基因所引起的胶质瘤生长抑制。Tian等^[45]在U251和LN229胶质瘤细胞中敲除或过表达HMGB1基因,采用逆转录-聚合酶链反应(RT-PCR)和lncRNA芯片筛选发现,LINC00320基因变化>10倍提示LINC00320受HMGB1的调控;细胞核定位的LINC00320在体内外均可抑制胶质瘤细胞增殖,新鲜胶质瘤组织中LINC00320表达下调提示预后不良。Zhang等^[46]发现,lncRNA TP73-AS1(又称TP73-AS1)在胶质瘤及其细胞系中特异性表达上调,而miRNA-142表达下调且与预后不良相关,他们还通过体外研究证实HMGB1/RAGE信号转导通路参与TP73-AS1/miRNA-142对胶质瘤细胞增殖、侵袭的调控,这一过程中TP73-AS1与HMGB1竞争miRNA-142的结合位点,即TP73-AS1通过海绵化miRNA-142上调HMGB1的表达,从而促进胶质瘤细胞的增殖和侵袭。

2.微小RNA Zheng等^[47]采用RT-PCR法对胶质母细胞瘤和瘤旁组织进行检测发现,肿瘤组织miRNA-339-5p表达量低于瘤旁组织,而瘤旁组织PTP4A1表达量升高;体外培养U251胶质瘤细胞,外

源性上调 miRNA-339-5p 表达,肿瘤细胞 PTP4A1 和 HMGB1 降低,血管样结构减少,提示 miRNA-339-5p 可通过抑制 PTP4A1/HMGB1 信号转导通路以抑制 U251 细胞的血管生成、迁移和侵袭。Cheng 等^[48]认为,miRNA-505-3p 低表达与胶质母细胞瘤患者预后不良相关,过表达则可抑制胶质瘤细胞增殖、迁移和侵袭;体外研究证实, HMGB1 是 miRNA-505-3p 的直接靶点,后者通过靶向抑制 HMGB1 和蛋白激酶 B (PKB) 以阻止胶质母细胞瘤进展。Li 等^[49]报告, miRNA-10b 在胶质瘤组织和 U87、U251 细胞中特异性表达上调,转染 miRNA-10b-5p 抑制剂可使细胞划痕愈合能力和侵袭性下降,转染 miRNA-10b-5p 拟似物则呈相反效果,肿瘤细胞侵袭相关蛋白 HMGB1、RhoC 和基质金属蛋白酶 2(MMP-2) 水平升高。体外研究结果显示, miRNA-218-5p 通过与 *HMGB1* mRNA 3' 非翻译区(3'UTR) 的位点结合下调 HMGB1 在胶质瘤细胞中的表达,从而抑制胶质瘤细胞增殖、侵袭和迁移,并诱导其凋亡^[50]。动物实验显示, miRNA-218-5p 通过下调 HMGB1 表达抑制裸鼠胶质瘤生长; miRNA-142-3p 通过与 HMGB1 3'UTR 结合抑制 Wnt/ β -catenin 信号转导通路关键分子,激活 Caspase-3,从而抑制胶质瘤细胞增殖,并诱导其凋亡^[51]。Gu 等^[52]发现, miRNA-218 在 U251 和 U87 胶质瘤细胞中表达下调,而 HMGB1 表达上调;转染 miRNA-218 拟似物可以显著降低肿瘤细胞活性和集落形成,促进细胞凋亡,减少细胞侵袭和迁移,并将细胞阻滞在 G0/G1 期; miRNA-218 可下调 HMGB1、RAGE、细胞周期蛋白 D1 和 MMP-9 表达,上调 Caspase-9 表达,沉默 *HMGB1* 基因后肿瘤细胞 RAGE、细胞周期蛋白 D1 和 MMP-9 水平升高,而 Caspase-9 水平下降; HMGB1 和 miRNA-218 共同过表达时胶质瘤生长抑制明显改善,但细胞凋亡增加,而 HMGB1 和 miRNA-218 共同抑制时上述作用消除, HMGB1 过表达而 miRNA-218 抑制时肿瘤细胞侵袭性增强。Yang 等^[53]发现, miRNA-129-2 在胶质瘤中表达下调,其主要通过靶向抑制 HMGB1 而发挥抑制肿瘤生长作用。

3. 环状 RNA circRNA 在胶质瘤领域的研究较少,但其功能不可忽视。Tao 等^[54]采用 RT-PCR 法检测胶质瘤 circFANCL、miRNA-337-3p 和 HMGB1 的表达变化,结果显示, circFANCL 在胶质瘤组织和细胞中显著升高,且与预后呈负相关,下调其表达可抑制肿瘤细胞增殖、诱导细胞周期阻滞、促进细胞

凋亡,进而抑制胶质瘤生长;同时, circFANCL 还可作为 miRNA-337-3p 的海绵对其进行调控, HMGB1 又是 miRNA-337-3p 的作用靶点,即 circFANCL 通过调控 miRNA-337-3p/HMGB1 信号转导通路促进胶质瘤恶性进展。

4. *HMGB1* 基因与胶质瘤细胞自噬 自噬是细胞基于细胞质内容物的循环和通过溶酶体降解清除受损分子,从而面对各种刺激维持内环境稳态的关键过程^[55]。Lei 等^[56]采用碳离子束照射 U251 胶质瘤细胞发现其可诱导细胞凋亡,1 Gy 照射 24 小时后自噬显著增强,随后呈时间和剂量依赖性下降, HMGB1 则以时间和剂量依赖性方式释放至细胞外,提高自噬和细胞外 HMGB1 水平,从而在一定程度上保护自身免受程序性死亡的调控。释放入细胞质的 HMGB1 定位于内质网或线粒体,与 Beclin1 或 P53 结合,内质网蛋白 Sigma1-R 可调控 HMGB1 表达,与线粒体结合的 HMGB1 使线粒体变长,导致“巨型”线粒体形成、线粒体 DNA(mtDNA) 耗竭和线粒体蛋白质破坏,进而调控自噬^[57]。由于胶质母细胞瘤微环境氧含量和营养缺乏,故其具有较高的自噬活性,自噬在胶质母细胞瘤的进展中发挥重要作用^[58]。YAP 在胶质瘤中过表达,并与 HMGB1 水平呈正相关,体外研究显示, YAP 过表达可增强基础和诱导条件下 U251 和 U87 胶质瘤细胞的自噬,以氯喹阻断自噬可消除 YAP 的促肿瘤生长作用,下调 HMGB1 表达亦可消除 YAP 对自噬和胶质瘤生长的促进作用,提示 YAP 通过增强 HMGB1 介导的自噬以促进胶质瘤进展,抑制 YAP/HMGB1 信号转导通路是胶质母细胞瘤可行治疗靶点^[59]。

5. *HMGB1* 基因与胶质瘤微环境 胶质瘤细胞间通讯依赖肿瘤微环境的稳态,不同的信息传递方式表现出不同功能^[60-61]。外泌体是一种源自细胞并介导细胞间通讯的细胞外囊泡,不同类型细胞各有其独特的外泌体进行信息传递^[62-63]。Ma 等^[64]发现,星形胶质细胞外泌体 *HMGB1* 表达量高于 C6 胶质瘤细胞外泌体,且从星形胶质细胞中提取的外泌体可升高 C6 细胞 SASH1 水平,而自 *HMGB1* 耗尽的星形胶质细胞中提取的外泌体则无此功能,但在细胞外直接加入重组 HMGB1 后, SASH1 水平下降,提示 HMGB1 作为细胞外蛋白可下调 SASH1 的表达,但作为外泌体则可上调 SASH1 的表达,从而在不同条件下使胶质瘤细胞的病理活动向相反方向发展。Hong 等^[65]以溶瘤病毒感染胶质瘤小鼠模型和体外

培养的U87胶质瘤细胞,结果发现,感染后肿瘤微环境HMGB1分泌增多,小鼠瘤周脑组织血管渗出和水肿增加,抑制HMGB1表达可提高小鼠存活率。

综上所述,HMGB1基因参与基因复制、表达、DNA损伤修复,并积极维持基因组稳定性,通过促进并维持细胞免疫应答在非感染性疾病、炎症、自身免疫性疾病、缺血再灌注、肿瘤微环境免疫调控中发挥重要作用,特别是在胶质瘤中作为癌基因以不同方式参与调控肿瘤恶性进展^[66-68]。深入探讨HMGB1基因在胶质瘤中的异质性以及研发其特异性药物进行抗肿瘤研究可能是未来研究方向。

利益冲突 无

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· 小词典 ·

中英文对照名词词汇(六)

中性粒细胞/淋巴细胞比值	peripheral nerve hyperexcitability syndromes(PNHS)
neutrophil-to-lymphocyte ratio(NLR)	主要组织相容性复合物
肿瘤基因组学图谱计划	major histocompatibility complex(MHC)
The Cancer Genome Atlas(TCGA)	转化生长因子- β transforming growth factor- β (TGF- β)
肿瘤浸润中性粒细胞 tumor-infiltrating neutrophils(TINs)	自然杀伤细胞 natural killer lymphocyte(NK)
肿瘤微环境 tumor microenvironment(TME)	自身免疫性脑炎 autoimmune encephalitis(AE)
重症肌无力 myasthenia gravis(MG)	Miller Fisher综合征 Miller Fisher syndrome(MFS)
重症肌无力定量评分	阻塞性睡眠呼吸暂停 obstructive sleep apnea(OSA)
Quantitative Myasthenia Gravis Score(QMGS)	阻塞性睡眠呼吸暂停综合征
周期性肢体运动障碍	obstructive sleep apnea syndrome(OSAS)
periodic limb movement disorder(PLMD)	最小临床表现 minimal manifestation(MM)
周围神经过度兴奋综合征	