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p53/miR-34a调控网络在肿瘤中的作用

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摘要: MicroRNA 是重要的调控分子, 长约 22 nt, 属于非编码 RNA, 对肿瘤的发生和发展起着重要的作用。miR-34a 是研究比较清楚的 microRNA, 目前认为是重要的抑癌 microRNA。*p53* 是一个重要的抑癌基因。*p53* 与 miR-34a 形成的正反馈调控网络具有抑制肿瘤细胞生长、转移及抑制肿瘤干细胞的功能。现综述 *p53*/miR-34a 调控网络研究的最新进展, 并探讨其在肿瘤诊断及治疗中的应用。

关键词: miR-34a; *p53*; 信号通路; 肿瘤

中图分类号: Q522; R730.2 **文献标志码:** A

p53/miR-34a signaling networks in tumor

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Abstract: MicroRNAs (miRNAs) are small noncoding RNAs with the length of ~22 nucleotides. They play important roles in tumorigenesis. miR-34a is considered to be a tumor suppressor. *p53* is a tumor suppressor. The positive feedback of *p53*/miR-34a could inhibit tumor cell growth and migration. This review will summarize the progress of *p53*/miR-34a network in tumor, and also discuss its application in diagnosis and treatment of cancer.

Key words: miR-34a; *p53*; signaling pathway; tumor

肿瘤是目前影响人类健康的主要疾病之一。肿瘤的发生涉及多条信号通路的异常。MicroRNA (miRNA) 参与了与肿瘤抑制及发生相关的通路调控。miRNA 是一类广泛存在于真核生物中, 由 18~23 个核苷酸组成的保守、非编码单链 RNA。miRNA 已成为治疗癌症的新途径^[1-4]。其中, miR-34a 是研究最多的 miRNA 之一。它是 miR-34 家族的成员, 与另外两位成员不同, miR-34a 单独位于染色体 1p36, 具有自身转录本。*p53* 基因是人类癌症中最容易发生突变的基因^[5], 也是重要的抑癌基因。*p53* 对 DNA 损伤、细胞应激反应产生应答^[6]。本文将综述 *p53* 所介导的 miR-34a 对肿瘤发生的影响。

1 *p53*与miR-34a

miR-34a 是最常见的受 P53 调控的 miRNA, *p53* 对肿瘤的抑制作用与 *p53* 介导的 miRNA 调控的靶基因有关 (图 1)。当细胞受到某些刺激, 如 DNA 损伤、药物刺激等, 会激活 *p53* 基因的表达以应对

刺激。P53 使得 miR-34a 的 CpG 岛启动子区域去甲基化, 激活 miR-34a 的表达, 进而调控 miR-34a 的靶基因。Hermeking^[7] 的研究表明, P53 蛋白对 miR-34a 的表达有直接靶向调节作用。此外, 也有间接调控因子影响 *p53* 与 miR-34a 的表达, 如沉默信息调节因子相关酶 1 (SIRT1), 它是一种 NAD⁺ 依赖的去乙酰化酶。当它抑制 P53 蛋白的转录后去乙酰化时, P53 蛋白的活性就会下调。miR-34a 通过靶向抑制 SIRT1 的表达进而诱导 *p53* 的活性^[8]。同时, miR-34a 是抑制肿瘤的重要分子, 参与调控多种癌症的发生与发展过程, 包括增殖、凋亡、迁移、侵袭、代谢、上皮间质转化 (epithelial-mesenchymal transition, EMT) 等, 从而抑制肿瘤的生长和转移^[9-11]。

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2 p53/miR-34a在肿瘤中的作用

2.1 对细胞周期的影响

对许多肿瘤的研究表明, miR-34a 异常表达时会抑制 *CDK4* 和 *CDK6* 的表达, 从而促进细胞周期进程^[12-13]。2015年, Kiyonari 等^[14] 研究表明, 诱导 *p53* 表达积累促进 miR-34a 的表达, 使得 E2F 转录因子 3 (E2F transcription factor 3, E2F3) 的表达受到了抑制, 结果促进了细胞周期进程。另外, Ichimura 等^[15] 的研究也表明, miR-34a 能抑制 MEK/ERK 信号通路的重要成员 MAP2K1, 从而达到抑制淋巴瘤

细胞增殖的作用。不仅如此, miR-34a 还可以靶向作用于 NAD^+ 补救途径的关键限速酶烟酰胺磷酸核糖转移酶 (nicotinamide phosphoribosyl-transferase, NAMPT), 从而调控 *SIRT1* 的活性^[16]。Liu 等^[17] 研究揭示, *SIRT1* 和 *c-MYC* 通过一个正反馈的环路相互调控 (图 2)。同时, miR-34a 也可以抑制 *c-MYC/SIRT1* 通路, 从而激活 *p53*, 进而上调 miR-34a 表达^[18-19]。

2.2 对细胞凋亡的影响

miR-34a 在许多癌症中都起到促凋亡的作用, 抗凋亡基因 *c-Myc* 较早被证明是 miR-34a 的靶

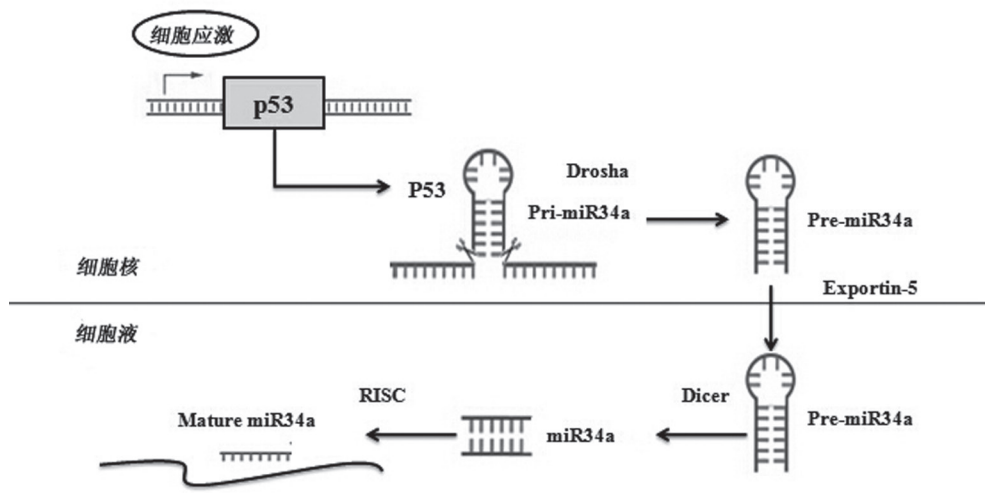
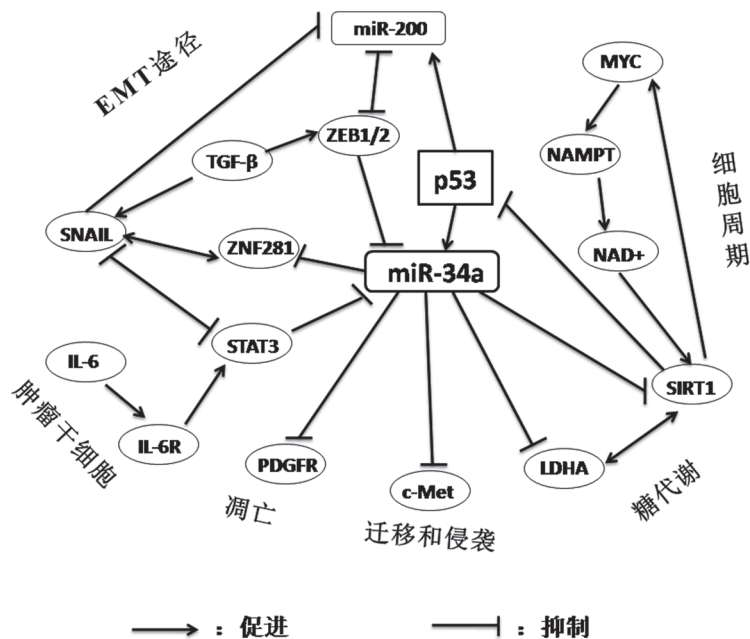


图1 P53调控miR-34a的表达



→ : 促进 —| : 抑制

图2 p53/miR-34a调控网络

基因^[19-20]。miR-34a 靶向调控 *Bcl-2*, 诱导细胞凋亡^[11]。在非小细胞肺癌 (non-small-cell lung carcinoma, NSCLC) 中, *PDGFR* 的表达量与 miR-34a 呈现负相关的关系。过表达 miR-34a 导致 *PDGFR* 的表达下调, 进而促进癌细胞的凋亡^[21]。死亡受体 CD95 (death receptor CD95, CD95) 有促进细胞凋亡的功能。同时, *CD95* 也是 *p53* 的靶基因之一。miR-34a 和 *CD95* 同时应答 *p53*, *CD95* 的表达能影响细胞激活 *p53*, 以及 *p53* 调控 miR-34a 的能力^[22]。miR-34a 与部分确证的靶基因的关系如表 1 所示。

2.3 对上皮间质转化(EMT)的影响

EMT 在肿瘤的发生过程中起到促进作用^[36]。miR-34a 直接靶向抑制 EMT 诱导转录因子 SNAIL, 进而抑制 EMT 过程^[37-38]。P53 蛋白通过靶向作用 miR-34a 和 miR-200 家族来调控 EMT 途径和其相对的 MET 途径, 从而在细胞的可塑性方面起到关键作用。*p53*、miR-34a 和 miR-200 家族与 SNAIL、锌指 E 盒结合同源框 1 (zinc finger E-box binding homeobox 1, ZEB1) 和锌指 E 盒结合同源框 2 (ZEB2) 组成了两条双负反馈回路 (图 2)^[39-40]。ZEB1 和 SNAIL 与 miR-34a 启动子的 E-box 结合, 起到了抑制 miR-34a 表达的作用, 这也进一步增加了 miR-34a/SNAIL 和 miR-200/ZEB 两条回路的关联^[37,41]。

miR-34a 靶基因的产物锌指蛋白 281 (zinc finger protein 281, ZNF281), 受到 SNAIL 和 miR-34a 的共同调节。因此, *ZNF281* 的表达受到 miR-34a 和 SNAIL 相关回路的调控, 一方面 miR-34a 能间接抑制其表达; 另一方面 SNAIL 可以直接诱导 *ZNF281* 的表达^[42]。

2.4 对迁移和侵袭途径的影响

miR-34a 的异常表达影响肿瘤恶化和迁移, miR-34a 低表达的原发性癌症患者体内肿瘤发生转移, 这揭示出 miR-34a 在抑制肿瘤转移方面有着重要作用^[43-45]。在小鼠实验中, 过表达 miR-34a 能够抑制转移瘤形成。体内实验表明, miR-34a 稳定过表达可以明显降低乳腺癌、前列腺癌、肺癌、骨肉瘤细胞的转移^[45-48]。许多研究表明, miR-34a 通过直接调控靶基因, 如 *AXL*、*PDGFR- α* 和 *c-Met*, 从而抑制癌细胞的迁移和侵袭。最近的一项研究表明, 把直肠癌细胞培养在含有白细胞介素 -6 (interleukin-6, IL-6) 的培养基中, 信号转导与转录激活因子 3 (signal transducer and activator of transcription 3, STAT3) 能直接抑制 miR-34a 的表达。*IL-6R/STAT3/miR-34a* 回路机制 (图 2), 即白细胞介素 -6 受体 (interleukin 6 receptor, *IL-6R*)、*STAT3* 和 miR-34a 存在一条正反馈调节的通路, 这条通路的激活与初期直肠肿瘤和

表1 miR-34a与其靶基因的关系

microRNA	靶基因	癌症类型	信号通路	参考文献
miR-34a	<i>Bcl-2</i>	乳腺癌	PI3K-Akt	[23]
	<i>CD24</i>	乳腺癌		[24]
	<i>LMTK3</i>	乳腺癌	AMPK	[25]
	<i>SIRT1</i>	结肠癌		[26]
	<i>MDM4</i>	结肠癌		p53
	<i>c-Kit</i>	结肠癌	PPAR	[28]
	<i>E2F3</i>	子宫颈癌		[29]
	<i>AR</i>	前列腺癌		[30]
	<i>AXL</i>	实体肿瘤	MAPK、Ras、PI3K-Akt	[31]
	<i>YY1</i>	食管鳞状细胞癌		[32]
	<i>PDGFR</i>	肺癌		[33]
	<i>c-Myc</i>	肝细胞性肝癌	MAPK、TGF- β	[34]
	<i>c-Met</i>	恶性间皮瘤	代谢通路	[35]

注: B细胞慢性淋巴细胞白血病/淋巴瘤2 (B-cell CLL/lymphoma 2, *Bcl-2*); CD24分子(CD24 molecule, *CD24*); 狐猴酪氨酸激酶3 (lemur tyrosine kinase 3, *LMTK3*); 沉默信息调节因子2相关酶1 (sirtuin 1, *SIRT1*); *MDM4*, p53调节基因(*MDM4*, p53 regulator, *MDM4*); 受体酪氨酸激酶(v-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog, *c-kit*); 雄激素受体 (androgen receptor, *AR*); *AXL*受体酪氨酸激酶(*AXL* receptor tyrosine kinase, *AXL*); *YY1*转录因子(*YY1* transcription factor, *YY1*); 血小板衍生的生长因子受体(platelet-derived growth factor receptor, *PDGFR*); v-myc禽骨髓细胞瘤病毒癌基因同源物 (v-myc avian myelocytomatosis viral oncogene homolog, *c-Myc*); 上皮间质细胞转换因子(mesenchymal-epithelial transition factor gene, *c-Met*)。

直肠癌的侵袭和迁移有关。

2.5 对糖代谢调控的影响

miR-34a 介导抑制己糖激酶 1 (hexokinase 1, HK1)、己糖激酶 2 (HK2)、葡萄糖-6-磷酸异构酶 (glucose-6-phosphate isomerase, GPI) 和丙酮酸脱氢酶激酶 (pyruvate dehydrogenase kinase, isozyme 1, PDK1), 从而抑制糖酵解和增强线粒体呼吸^[42]。而 p53 则是通过 miR-34a 介导抑制这些酶, 进而调控糖酵解和葡萄糖代谢。乳酸脱氢酶 A (lactate dehydrogenase A, LDHA) 催化丙酮酸转换为乳酸, 在无氧酵解中有着关键作用。Kaller 等^[49]证明了 LDHA 是 miR-34a 作用的靶位点。SIRT1 受到 LDHA 的调控。同时, SIRT1 的表达量受到抑制也会造成 LDHA 转录的下调^[50-51](图 2)。miR-34a 的靶基因 SIRT1, 不仅在细胞周期中起到重要的调控作用, 在 p53/miR-34a 介导的代谢调控中也起到了关键作用。

2.6 对肿瘤干细胞的影响

肿瘤干细胞 (cancer stem cells, CSCs) 具有干细胞的特性。Liu 等^[52]研究表明, 前列腺癌干细胞中 miR-34a 的表达量明显下调, 上调 miR-34a 的表达则抑制 CD44⁺组中肿瘤的再生和转移。miR-34a 在直肠癌干细胞中同样起到了重要作用^[53], 上调或者下调 miR-34a 的表达将影响到 miR-34a 靶基因 Notch1 介导的与 CSCs 有关的信号通路, 从而改变了 CSCs 自我更新与分化之间的平衡^[54]。miR-34a 除了可以调控 Notch 信号通路外, 还可以直接抑制 Notch 的配体 Dll1 (Delta-like ligand 1)^[55]。通过这种机制, miR-34a 可以抑制绒毛癌细胞的增殖、迁移、侵袭以及异种移植肿瘤的生长^[56]。Siemens 等^[28]在研究中发现, *c-kit* 不仅是 miR-34a 的靶基因, 而且 *c-kit* 对 CSC 有负调控作用。在间皮瘤中, p53/miR-34 调控网络 (图 2) 与 CSC 的增殖密切相关, 当下调 p53 或 miR-34a 时 *c-Met* 的表达量上调, 从而促进了 CSC 的增殖。同时, 其迁移和侵袭能力也有所增强^[35]。

3 展望

综上所述, 在 miR-34a 参与调控的信号通路中, p53/miR-34a 调控网络对肿瘤发生与发展起着重要的调控作用。通过这条通路, miR-34a 可以直接或间接地调控不同基因, 从而抑制肿瘤的发生或促进肿瘤细胞凋亡。所以, p53/miR-34a 调控网络与肿瘤的研究日趋深入。毫无疑问的是, 进一步研究这两者之间的关系对于揭示肿瘤发生和发展的机制有

着极其重要的意义。

从目前的研究结果来看, miR-34a 在癌症患者组织中是低表达的; 同时, miR-34a 与肿瘤干细胞、肿瘤的恶性程度、肿瘤大小以及不良预后呈负相关。因此, 检测 miR-34a 表达水平对于癌症患者的预后、诊断有着重要意义。在化疗耐药性方面, miR-34a 能够促进癌细胞对药物的敏感性, 从而增强药物的疗效。例如, miR-34a 可以促进乳腺癌细胞对紫杉醇的敏感性^[57]。有关视网膜母瘤的研究中同样存在这种对药物的敏感性的作用机制^[58]。在美国, miR-34a 的类似物已经进入 I 期临床试验。miR-34a 及其调控网络的研究才仅仅是个开始, 随着研究的深入, 新药物靶点的开发和应用将为期不远, 同时也是肿瘤精准医疗或个性化治疗的目标。

[参 考 文 献]

- [1] He L, Hannon GJ. MicroRNAs: small RNAs with a big role in gene regulation. *Nat Rev Genet*, 2004, 5: 522-31
- [2] Shyu AB, Wilkinson MF, van Hoof A. Messenger RNA regulation: to translate or to degrade. *EMBO J*, 2008, 27: 471-81
- [3] Neely LA, Patel S, Garver J. A single-molecule method for the quantitation of microRNA gene expression. *Nat Methods*, 2006, 3: 41-6
- [4] Berezikov E, Cuppen E, Plasterk RH. Approaches to microRNA discovery. *Nat Genet*, 2006, 38: S2-7
- [5] Kandoth C, McLellan MD, Vandin F, et al. Mutational landscape and significance across 12 major cancer types. *Nature*, 2013, 502: 333-9
- [6] Vogelstein B, Lane D, Levine AJ. Surfing the p53 network. *Nature*. 2000, 408: 307-10
- [7] Hermeking H. p53 enters the microRNA world. *Cancer Cell*, 2007, 12:414-8
- [8] Yamakuchi M, Ferlito M, Lowenstein CJ. miR-34a repression of SIRT1 regulates apoptosis. *Proc Natl Acad Sci USA*, 2008, 105: 13421-6
- [9] Chang TC, Wentzel EA, Kent OA, et al. Transactivation of miR-34a by p53 broadly influences gene expression and promotes apoptosis. *Mol Cell*, 2007, 26: 745-52
- [10] He L, He X, Lim LP, et al. A microRNA component of the p53 tumour suppressor network. *Nature*, 2007, 447: 1130-4
- [11] Bommer GT, Gerin I, Feng Y, et al. p53-mediated activation of miRNA34 candidate tumor-suppressor genes. *Current Biol*, 2007, 17: 1298-307
- [12] Hong JH, Roh KS, Suh SS, et al. The expression of microRNA-34a is inversely correlated with *c-MET* and *CDK6* and has a prognostic significance in lung adenocarcinoma patients. *Tumour Biol*, 2015, 36: 9327-37
- [13] Hargraves KG, He L, Firestone GL. Phytochemical regulation of the tumor suppressive microRNA, miR-34a, by p53-dependent and independent responses in human breast cancer cells. *Mol Carcinogen*, 2016, 55: 486-98

- [14] Kiyonari S, Imori M, Matsuoka K, et al. The 1,2-Diaminocyclohexane carrier ligand in oxaliplatin induces p53-dependent transcriptional repression of factors involved in thymidylate biosynthesis. *Mol Cancer Ther*, 2015, 14: 2332-42
- [15] Ichimura A, Ruike Y, Terasawa K, et al. MicroRNA-34a inhibits cell proliferation by repressing mitogen-activated protein kinase kinase 1 during megakaryocytic differentiation of K562 cells. *Mol Pharmacol*, 2010, 77: 1016-24
- [16] Choi SE, Fu T, Seok S, et al. Elevated microRNA-34a in obesity reduces NAD⁺ levels and SIRT1 activity by directly targeting NAMPT. *Aging Cell*, 2013, 12: 1062-72
- [17] Liu Y, Li X, Zhu S, et al. Ectopic expression of miR-494 inhibited the proliferation, invasion and chemoresistance of pancreatic cancer by regulating SIRT1 and c-Myc. *Gene Ther*, 2015, 22: 729-38
- [18] Menssen A, Hermeking H. c-MYC and SIRT1 locked in a vicious cycle. *Oncotarget*, 2012, 3: 112-3
- [19] Marshall GM, Liu PY, Gherardi S, et al. SIRT1 promotes N-Myc oncogenesis through a positive feedback loop involving the effects of MKP3 and ERK on N-Myc protein stability. *PLoS Genet*, 2011, 7: e1002135
- [20] Long Z, Wang B, Tao D, et al. Hypofractionated radiotherapy induces miR-34a expression and enhances apoptosis in human nasopharyngeal carcinoma cells. *Int J Mol Med*, 2014, 34: 1388-94
- [21] Garofalo M, Jeon YJ, Nuovo GJ, et al. MiR-34a/c-dependent PDGFR- α/β downregulation inhibits tumorigenesis and enhances TRAIL-induced apoptosis in lung cancer. *PLoS One*, 2013, 8: e67581
- [22] Hau A, Ceppi P, Peter ME. CD95 is part of a let-7/p53/miR-34 regulatory network. *PLoS One*, 2012, 7: e49636
- [23] Li L, Yuan L, Luo J, Gao J, et al. MiR-34a inhibits proliferation and migration of breast cancer through down-regulation of Bcl-2 and SIRT1. *Clin Exp Med*, 2013, 13: 109-17
- [24] Muppala S, Mudduluru G, Leupold JH, et al. CD24 induces expression of the oncomir miR-21 via Src, and CD24 and Src are both post-transcriptionally downregulated by the tumor suppressor miR-34a. *PLoS One*, 2013, 8: e59563
- [25] Zhao G, Guo J, Li D, et al. MicroRNA-34a suppresses cell proliferation by targeting LMTK3 in human breast cancer mcf-7 cell line. *DNA Cell Biol*, 2013, 32: 699-707
- [26] Mohan M, Kumar V, Lackner AA, et al. Dysregulated miR-34a-SIRT1-acetyl p65 axis is a potential mediator of immune activation in the colon during chronic simian immunodeficiency virus infection of rhesus macaques. *J Immunol*, 2015, 194: 291-306
- [27] Mandke P, Wyatt N, Fraser J, et al. MicroRNA-34a modulates MDM4 expression via a target site in the open reading frame. *PLoS One*, 2012, 7: e42034
- [28] Siemens H, Jackstadt R, Kaller M, et al. Repression of c-Kit by p53 is mediated by miR-34 and is associated with reduced chemoresistance, migration and stemness. *Oncotarget*, 2013, 4: 1399-415
- [29] Geng D, Song X, Ning F, et al. MiR-34a inhibits viability and invasion of human papillomavirus-positive cervical cancer cells by targeting E2F3 and regulating survivin. *Int J Gynecol Cancer*, 2015, 25: 707-13
- [30] Kashat M, Azzouz L, Sarkar SH, et al. Inactivation of AR and Notch-1 signaling by miR-34a attenuates prostate cancer aggressiveness. *Am J Transl Res*, 2012, 4: 432-42
- [31] Mudduluru G, Ceppi P, Kumarswamy R, et al. Regulation of Axl receptor tyrosine kinase expression by miR-34a and miR-199a/b in solid cancer. *Oncogene*, 2011, 30: 2888-99
- [32] Nie J, Ge X, Geng Y, et al. miR-34a inhibits the migration and invasion of esophageal squamous cell carcinoma by targeting Yin Yang-1. *Oncology Rep*, 2015, 34: 311-7
- [33] Garofalo M, Jeon YJ, Nuovo GJ, et al. Correction: MiR-34a/c-dependent PDGFR- α/β downregulation inhibits tumorigenesis and enhances TRAIL-induced apoptosis in lung cancer. *PLoS One*, 2015, 10: e0131729
- [34] Xu X, Chen W, Miao R, et al. miR-34a induces cellular senescence via modulation of telomerase activity in human hepatocellular carcinoma by targeting FoxM1/c-Myc pathway. *Oncotarget*, 2015, 6: 3988-4004
- [35] Menges CW, Kadariya Y, Altomare D, et al. Tumor suppressor alterations cooperate to drive aggressive mesotheliomas with enriched cancer stem cells via a p53-miR-34a-c-Met axis. *Cancer Res*, 2014, 74: 1261-71
- [36] De Craene B, Berx G. Regulatory networks defining EMT during cancer initiation and progression. *Nat Rev Cancer*, 2013, 13: 97-110
- [37] Siemens H, Jackstadt R, Hunten S, et al. miR-34 and SNAIL form a double-negative feedback loop to regulate epithelial-mesenchymal transitions. *Cell Cycle*, 2011, 10: 4256-71
- [38] Du R, Sun W, Xia L, et al. Hypoxia-induced down-regulation of microRNA-34a promotes EMT by targeting the Notch signaling pathway in tubular epithelial cells. *PLoS One*, 2012, 7: e30771
- [39] Kim T, Veronese A, Pichiorri F, et al. p53 regulates epithelial-mesenchymal transition through microRNAs targeting ZEB1 and ZEB2. *J Exp Med*, 2011, 208: 875-83
- [40] Burk U, Schubert J, Wellner U, et al. A reciprocal repression between ZEB1 and members of the miR-200 family promotes EMT and invasion in cancer cells. *EMBO Rep*, 2008, 9: 582-9
- [41] Ahn YH, Gibbons DL, Chakravarti D, et al. ZEB1 drives prometastatic actin cytoskeletal remodeling by downregulating miR-34a expression. *J Clin Invest* 2012, 122: 3170-83
- [42] Hahn S, Jackstadt R, Siemens H, et al. SNAIL and miR-34a feed-forward regulation of ZNF281/ZBP99 promotes epithelial-mesenchymal transition. *EMBO J*, 2013, 32: 3079-95
- [43] Dang Y, Luo D, Rong M, et al. Underexpression of miR-34a in hepatocellular carcinoma and its contribution towards enhancement of proliferating inhibitory effects of agents targeting c-MET. *PLoS One*, 2013, 8: e61054
- [44] Siemens H, Neumann J, Jackstadt R, et al. Detection of miR-34a promoter methylation in combination with ele-

- vated expression of c-Met and β -catenin predicts distant metastasis of colon cancer. *Clin Cancer Res*, 2013, 19: 710-20
- [45] Yang S, Li Y, Gao J, Zhang T, et al. MicroRNA-34 suppresses breast cancer invasion and metastasis by directly targeting Fra-1. *Oncogene*, 2013, 32: 4294-303
- [46] Yan K, Gao J, Yang T, et al. MicroRNA-34a inhibits the proliferation and metastasis of osteosarcoma cells both *in vitro* and *in vivo*. *PLoS One*, 2012, 7: e33778
- [47] Kong D, Heath E, Chen W, et al. Epigenetic silencing of miR-34a in human prostate cancer cells and tumor tissue specimens can be reversed by BR-DIM treatment. *Am J Transl Res*, 2012, 4: 14-23
- [48] Zhou JY, Chen X, Zhao J, et al. MicroRNA-34a overcomes HGF-mediated gefitinib resistance in EGFR mutant lung cancer cells partly by targeting MET. *Cancer Lett*, 2014, 351: 265-71
- [49] Kaller M, Liffers ST, Oeljeklaus S, et al. Genome-wide characterization of miR-34a induced changes in protein and mRNA expression by a combined pulsed SILAC and microarray analysis. *Mol Cell Proteomics*, 2011, 10: M111 010462
- [50] Mao B, Zhao G, Lv X, et al. Sirt1 deacetylates c-Myc and promotes c-Myc/Max association. *Int J Biochem Cell Biol*, 2011, 43: 1573-81
- [51] Simmons GE Jr, Pruitt WM, Pruitt K. Diverse roles of SIRT1 in cancer biology and lipid metabolism. *Int J Mol Sci*, 2015, 16: 950-65
- [52] Liu C, Kelnar K, Liu B, et al. The microRNA miR-34a inhibits prostate cancer stem cells and metastasis by directly repressing CD44. *Nat Med*, 2011, 17: 211-5
- [53] Bu P, Chen KY, Chen JH, et al. A microRNA miR-34a-regulated bimodal switch targets Notch in colon cancer stem cells. *Cell Stem Cell*, 2013, 12: 602-15
- [54] Winton DJ. miR-34a sets the "sweet spot" for notch in colorectal cancer stem cells. *Cell Stem Cell*, 2013, 12: 499-501
- [55] de Antonellis P, Medaglia C, Cusanelli E, et al. MiR-34a targeting of Notch ligand delta-like 1 impairs CD15⁺/CD133⁺ tumor-propagating cells and supports neural differentiation in medulloblastoma. *PLoS One*, 2011, 6: e24584
- [56] Pang RT, Leung CO, Lee CL, et al. MicroRNA-34a is a tumor suppressor in choriocarcinoma via regulation of Delta-like1. *BMC Cancer*, 2013, 13: 25
- [57] Kang L, Mao J, Tao Y, et al. MicroRNA-34a suppresses the breast cancer stem cell-like characteristics by down-regulating Notch1 pathway. *Cancer Sci*, 2015, 106: 700-8
- [58] Zhao J, Kelnar K, Bader AG. In-depth analysis shows synergy between erlotinib and miR-34a. *PLoS One*, 2014, 9: e89105