

AFP对裸鼠肝癌移植瘤生长的促进作用及对血管形成相关因子的影响

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Title: The effect and possible angiogenesis mechanism of AFP transfection of SMMC-7721 xenografts in nude mice

作者: 魏柏¹; 陈景三¹; 杨盛力²; 占静¹

1.华中科技大学同济医学院附属梨园医院肿瘤科, 湖北 武汉 430077; 2.华中科技大学同济医学院附属协和医院肿瘤中心, 湖北 武汉 430022

Author(s): Wei Bai¹; Chen Jingsan¹; Yang Shengli²; Zhan Jing¹

1. Department of Oncology, Liyuan Hospital, the Affiliated Hospital of Tongji Medical University, Huazhong University of Science and Technology, Hubei Wuhan 430077, China; 2. Cancer Center of Union Hospital, the Affiliated Hospital of Tongji Medical University, Huazhong University of Science and Technology, Hubei Wuhan 430022, China.

关键词: 肝癌; AFP; 肿瘤生长; 血管形成

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摘要: 目的: 探讨AFP对肝癌移植瘤生长及血管生成调节因子表达的影响。方法: 采用pEGFP-N1-APP和pEGFP-N1质粒分别转染SMMC-7721细胞构建SMMC-7721/APP和SMMC-7721/CON细胞, 将裸鼠分为实验组和对照组, 分别于裸鼠皮下接种SMMC-7721/APP和SMMC-7721/CON, 从而建立人肝癌移植瘤模型, 动态观测裸鼠肿瘤体积和质量变化; 取肿瘤组织用实时定量RT-PCR和Western blot检测组织中的HOXA7、eIF4E、VEGFA和FGF2表达。结果: 构建SMMC-7721/APP和SMMC-7721/CON细胞, 并将其接种到裸鼠皮下成功建立肝癌移植瘤模型。与对照组比较, 实验组移植瘤体积与质量显著增高, 成瘤后第12天, 实验组移植瘤体积和质量分别为(307.71±47.63) mm³和(20.243±0.411) g, 差异均具有显著性(P=0.012, P=0.04)。实时定量RT-PCR结果显示APP过表达提高了肝癌移植瘤细胞HOXA7、eIF4E和FGF2 mRNA表达水平, 分别为2.488±1.155、23.828±2.465和4.407±1.164, 与对照组相比增高程度具有显著性差异, 但VEGFA mRNA表达未见显著增强。Western blot结果显示: SMMC-7721/APP较SMMC-7721/CON细胞显著提高HOXA7蛋白表达水平, 但对FGF2和VEGFA无显著影响。结论: APP基因转染可明显促进裸鼠肝癌移植瘤的生长, 该作用可能与其促进肿瘤血管形成因子在mRNA水平的调控有关。

Abstract: Objective: To investigate the effect and possible angiogenesis mechanism of AFP gene transfection of SMMC-7721 xenografts in nude mice. Methods: An expressing vector encoding AFP was recombined and then transfected into cultured SMMC-7721 with lipofectin. SMMC-7721/APP and SMMC-7721/CON were inoculated nude mice to establish nude mice subcutaneous transplantation tumor of hepatocellular carcinoma model. Nude mice tumor volume and quality were observed dynamically HOXA7, eIF4E, VEGFA, FGF2 and MDM2 were measured in tumor tissue by real time RT-PCR and Western blot. Results: SMMC-7721/APP and SMMC-7721/CON were established. Then xenografts in nude mice were successfully established. Compared with untransfected group, APP gene transfection in experimental group promoted tumor volume and quality with significant difference. The volume and quality of the xenografts in nude mice were (307.71±47.63) mm³ and (20.243±0.411) g separately. The mRNA expression of HOXA7, eIF4E, VEGFA, FGF2 and MDM2 were upregulated. RT-PCR showed that APP overexpression improved the mRNA expression of HOXA7, eIF4E and FGF2 mRNA to 2.488±1.155, 23.828±2.465 and 4.407±1.164 with significance. However, VEGFA mRNA expression improved without significant difference. And Western blot results showed that HOXA7 was significantly increased in SMMC-7721/APP compared to that in SMMC-7721/CON. However, the protein level of FGF2 and

VEGFA had no significant alteration. Conclusion: AFP gene transfection can promote the growth of nude mouse transplantation tumor of hepatocellular carcinoma significant, which may be related with tumor angiogenesis.

参考文献/REFERENCES

- [1] Are C,Meyer B,Stack A,et al.Global trends in the burden of liver cancer [J].J Surg Oncol,2017,115(5):591-602.
- [2] Torre LA,Bray F,Siegel RL,et al.Global cancer statistics,2012 [J].CA:A Cancer Journal for Clinicians,2015,65(2):87-108.
- [3] Gong X,Qin S.Study progression of anti-angiogenetic therapy and its combination with other agents for the treatment of advanced hepatocellular carcinoma [J].Hepatobiliary Surgery and Nutrition,2018,7(6):466-474.
- [4] Morse MA,Sun W,Kim R,et al.The role of angiogenesis in hepatocellular carcinoma [J].Clin Cancer Res,2019,25(3):912-920.
- [5] Berretta M,Rinaldi L,Di Benedetto F,et al.Angiogenesis inhibitors for the treatment of hepatocellular carcinoma [J].Frontiers in Pharmacology,2016(7):428.
- [6] Mitsuhashi N,Kobayashi S,Doki T,et al.Clinical significance of alpha-fetoprotein:Involvement in proliferation,angiogenesis, and apoptosis of hepatocellular carcinoma [J].Journal of Gastroenterology and Hepatology,2008,23(7 Pt 2):e189-197.
- [7] Zhang C,Chen X,Liu H,et al.Alpha fetoprotein mediates HBx induced carcinogenesis in the hepatocyte cytoplasm [J].International Journal of Cancer,2015,137(8):1818-1829.
- [8] Wang S,Jiang W,Chen X,et al.Alpha-fetoprotein acts as a novel signal molecule and mediates transcription of Fn14 in human hepatocellular carcinoma [J].Journal of Hepatology,2012,57(2):322-329.
- [9] Li M,Li H,Li C,et al.Alpha-fetoprotein:A new member of intracellular signal molecules in regulation of the PI3K/AKT signaling in human hepatoma cell lines [J].International Journal of Cancer,2011,128(3):524-532.
- [10] Zhu M,Xia H,Li W,et al.Alpha-fetoprotein activates AKT/mTOR signaling to promote CXCR4 expression and migration of hepatoma cells [J].Oncoscience,2015,2(1):11.
- [11] Yang X,Zhang Y,Zhang L,et al.Silencing alpha-fetoprotein expression induces growth arrest and apoptosis in human hepatocellular cancer cell [J].Cancer letters,2008,271(2):281-293.
- [12] Sauzay C,Petit A,Bourgeois AM,et al.Alpha-foetoprotein (AFP):A multi-purpose marker in hepatocellular carcinoma [J].Clin Chim Acta,2016(463):39-44.
- [13] Wang S,Zhu M,Wang Q,et al.Alpha-fetoprotein inhibits autophagy to promote malignant behaviour in hepatocellular carcinoma cells by activating PI3K/AKT/mTOR signalling [J].Cell Death & Disease,2018,9(10):1027.
- [14] Viallard C,Larivée B.Tumor angiogenesis and vascular normalization:Alternative therapeutic targets [J].Angiogenesis,2017,20(4):409-426.
- [15] Sajib S,Zahra FT,Lionakis MS,et al.Mechanisms of angiogenesis in microbe-regulated inflammatory and neoplastic conditions [J].Angiogenesis,2018,21(1):1-14.
- [16] Mentzer SJ,Konerding MA.Intussusceptive angiogenesis:Expansion and remodeling of microvascular networks [J].Angiogenesis,2014,17(3):499-509.
- [17] Mallo M,Alonso CR.The regulation of Hox gene expression during animal development [J].Development,2013,140(19):3951-3963.
- [18] Bahrami SB,Veiseh M,Dunn AA,et al.Temporal changes in Hox gene expression accompany endothelial cell differentiation of embryonic stem cells [J].Cell Adhesion & Migration,2014,5(2):133-141.
- [19] Jia Y,Bleicher F,Merabet S.A systematic survey of HOX and TALE expression profiling in human cancers [J].The International Journal of Developmental Biology,2018,62(11-12):865-876.
- [20] Cantile M,Schiavo G,Terracciano L,et al.Homeobox genes in normal and abnormal vasculogenesis [J].Nutrition,Metabolism, and Cardiovascular Diseases:NMCD,2008,18(10):651-658.
- [21] Fontijn D,Bosch LJ,Duyndam MC,et al.Basic fibroblast growth factor-mediated overexpression of vascular endothelial growth factor in 1F6 human melanoma cells is regulated by activation of PI3K and p38 MAPK [J].Cellular Oncology,2009,31(3):1179-1190.

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