

高血压相关的线粒体DNA突变

薛凌¹, 陈红¹, 孟燕子¹, 王燕¹, 卢中秋¹, 吕建新², 管敏鑫^{1,2,3}

1. 温州医学院Attardi线粒体生物医学研究院, 温州 325035 2. 温州医学院浙江省医学遗传学重点实验室, 温州 325035 3. Cincinnati Children's Hospital Medical Center, Division of Human Genetics, Cincinnati, OH 45229, USA

XUE Ling¹, CHEN Hong¹, MENG Yan-Zi¹, WANG Yan¹, LU Zhong-Qiu¹, LU Jian-Xin², GUAN Min-Xin^{1,2,3}

1. Attardi Institute of Mitochondrial Biomedicine, Wenzhou Medical College, Wenzhou 325035, China 2. Zhejiang Provincial Key Laboratory of Medical Genetics, Wenzhou Medical College, Wenzhou 325035, China 3. Division of Human Genetics, Cincinnati Children's Hospital Medical Center, Cincinnati, OH 45229, USA

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摘要 线粒体DNA(mtDNA)突变是高血压发病的分子机制之一。已经报道的与原发性高血压相关的mtDNA突变包括: tRNA^{Met}A4435G, tRNA^{Met}/tRNA^{Gln}A4401G, tRNA^{Ile}A4263G, T4291C 和 A4295G突变。这些高血压相关的mtDNA突变改变了相应的线粒体tRNA的结构, 导致线粒体tRNA的代谢障碍。而线粒体tRNAs的代谢缺陷则影响蛋白质合成, 造成氧化磷酸化缺陷, 降低ATP的合成, 增加活性氧的产生。因此, 线粒体的功能缺陷可能在高血压的发生发展中起一定的作用。mtDNA突变发病的组织特异性则可能与线粒体tRNAs的代谢以及核修饰基因相关。目前发现的这些高血压相关的mtDNA突变则应该作为今后高血压诊断的遗传风险因子。高血压相关的线粒体功能缺陷的深入研究也将进一步诠释母系遗传高血压的分子致病机制, 为高血压的预防、控制和治疗提供依据。文章对高血压相关的mtDNA突变进行了综述。

关键词: 高血压 母系遗传 mtDNA突变 线粒体tRNA

Abstract: Mutations in mitochondrial DNA (mtDNA) are one of the molecular bases of hypertension. Among these, the tRNA^{Met}A4435G, tRNA^{Met}/tRNA^{Gln}A4401G, tRNA^{Ile}A4263G, T4291C and A4295G mutations have been reported to be associated with essential hypertension. These mutations alter the structure of the corresponding mitochondrial tRNAs and cause failures in tRNA metabolism. These shortages of these tRNAs lead to an impairment of mitochondrial protein synthesis and a failure in the oxidative phosphorylation function. These result in a deficit in ATP synthesis and an increase of generation of reactive oxygen species. As a result, these mitochondrial dysfunctions may contribute to the development of hypertension. Furthermore, the tissue specificity of these pathogenic mtDNA mutations might be associated with tRNA metabolism and nuclear modifier genes. These mtDNA mutations should be considered as inherited risk factors for future molecular diagnosis. Thus, these findings provide new insights into the molecular mechanism, management and treatment of maternally inherited hypertension. This review summarized the association between mtDNA mutations and hypertension.

Keywords: hypertension, maternal inheritance, mtDNA mutation, mitochondrial tRNA

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通讯作者 管敏鑫 Email: gminxin88@gmail.com


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[1] Gu DF, Reynolds K, Wu XG, Chen J, Duan XF, Muntner P, Huang GY, Reynolds RF, Su SY, Whelton PK, He J. Prevalence, awareness, treatment, and control of hyper-tension in China. *Hypertension*, 2002, 40(6): 920-927. 

[2] 中国高血压防治指南修订委员会. 中国高血压防治指南(2005年修订版). 高血压杂志, 2005, 134(增刊): 2-41.

[3] Whitworth JA, World Health Organization, International Society of Hypertension Writing Group. 2003 World Health Organization (WHO)/International Society of Hypertension (ISH). *J Hypertension*, 2003, 21(11): 432.

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- [4] Brandão AP, Brandão AA, Araújo EM, Oliveira RC. Familial aggregation of arterial blood pressure and possible genetic influence. *Hypertension*, 1992, 19(Suppl.2): 11214-11217.
- [5] Watson B Jr, Khan MA, Desmond RA, Bergman S. Mitochondrial DNA mutations in black Americans with hypertension-associated end-stage renal disease. *Am J Kidney Dis*, 2001, 38(3): 529-536. 
- [6] Hirano M, Davidson M, DiMauro S. Mitochondria and the heart. *Curr Opin Cardiol*, 2001, 16(3): 201-210. 
- [7] Schwartz F, Duka A, Sun FZ, Cui J, Manolis A, Gavras H. Mitochondrial genome mutations in hypertensive individuals. *Am J Hypertens*, 2004, 17(7): 629-635. 
- [8] Wallace DC. Mitochondrial diseases in man and mouse. *Science*, 1999, 283(5407): 1482-1488.
- [9] Schon EA, Bonilla E, DiMauro S. Mitochondrial DNA mutations and pathogenesis. *J Bioenerg Biomembr*, 1997, 29(2): 131-149. 
- [10] Anderson S, Bankier AT, Barrell BG, de Bruijn MHL, Coulson AR, Drouin J, Eperon IC, Nierlich DP, Roe BA, Sanger F, Schreier PH, Smith AJH, Staden R, Young IG. Sequence and organization of the human mitochondrial genome. *Nature*, 1981, 290(5806): 457-465.
- [11] Attardi G, Schatz G. Biogenesis of mitochondria. *Annu Rev Cell Biol*, 1988, 4: 289-333. 
- [12] Wallace DC. Diseases of the mitochondrial DNA. *Annu Rev Biochem*, 1992, 61(1): 1175-1212. 
- [13] Giles RE, Blanc H, Cann HM, Wallace DC. Maternal inheritance of human mitochondrial DNA. *Proc Natl Acad Sci USA*, 1980, 77(11): 6715-6719. 
- [14] Jenuth JP, Peterson AC, Fu K, Shoubridge EA. Random genetic drift in the female germline explains the rapid segregation of mammalian mitochondrial DNA. *Nat Genet*, 1996, 14(2): 146-151. 
- [15] 严庆丰, 管敏鑫. 线粒体疾病与核基因—线粒体基因的表达调控. *生命科学*, 2008, 20(4): 496-505.
- [16] Wallace DC. Mitochondrial defects in cardiomyopathy and neuromuscular disease. *Am Heart J*, 2000, 139(2-3): S70-S85.
- [17] Hutchin T, Cortopassi G. A mitochondrial DNA clone is associated with increased risk for Alzheimer disease. *Proc Natl Acad Sci USA*, 1995, 92(15): 6892-6895. 
- [18] Mayr-Wohlfarth U, Rödel G, Henneberg A. Mitochondrial tRNA(Gln) and tRNA(Thr) gene variants in Parkinson's disease. *Eur J Med Res*, 1997, 2(3): 111-113.
- [19] Perucca-Lostanlen D, Narbonne H, Hernandez JB, Staccini P, Saunieres A, Paquis-Flucklinger V, Vialettes B, Desnuelle C. Mitochondrial DNA variations in patients with maternally inherited diabetes and deafness syndrome. *Biochem Biophys Res Commun*, 2000, 277(3): 771-775. 
- [20] Li XM, Li RH, Lin XH, Guan MX. Isolation and characterization of the putative nuclear modifier gene MTO1 involved in the pathogenesis of deafness-associated mitochondrial 12 S rRNA A1555G mutation. *J Biol Chem*, 2002, 277(30): 27256-27264. 
- [21] Brown MD, Starikovskaya E, Derbeneva O, Hosseini S, Allen JC, Mikhailovskaya IE, Sukernik RI, Wallace DC. The role of mtDNA background in disease expression: a new primary LHON mutation associated with Western Eurasian haplogroup J. *Hum Genet*, 2002, 110(2): 130-138. 
- [22] Prezant TR, Agopian JV, Bohlman MC, Bu XD, Öztas S, Qiu WQ, Arnos KS, Cortopassi GA, Jaber L, Rotter JI, Shohat M, Fischel-Ghodsian N. Mitochondrial ribosomal RNA mutation associated with both antibiotic-induced and non-syndromic deafness. *Nat Genet*, 1993, 4(3): 289-294. 
- [23] Santorelli FM, Tanji K, Manta P, Casali C, Krishna S, Hays AP, Mancini DM, DiMauro S, Hirano M. Maternally inherited cardiomyopathy: an atypical presentation of the mtDNA 12S rRNA gene A1555G mutation. *Am J Hum Genet*, 1999, 64(1): 295-300. 
- [24] Zeviani M, Gellera C, Antozzi C, Rimoldi M, Morandi L, Villani F, Tiranti V, DiDonato S. Maternally inherited myopathy and cardiomyopathy: association with mutation in mitochondrial DNA tRNA^{Leu(UUR)}. *Lancet*, 1991, 338(8760): 143-147.
- [25] Silvestri G, Santorelli FM, Shanske S, Whitley CB, Schimmenti LA, Smith SA, DiMauro S. A new mtDNA mutation in the tRNA^{Leu(UUR)} gene associated with maternally inherited cardiomyopathy. *Hum Mutat*, 1994, 3(1): 37-43. 
- [26] Terasaki F, Tanaka M, Kawamura K, Kanzaki Y, Okabe M, Hayashi T, Shimomura H, Ito T, Suwa M, Gong JS, Zhang J, Kitaura Y. A case of cardiomyopathy showing progression from the hypertrophic to the dilated form: association of Mt8348A→G mutation in the mitochondrial tRNA^{Lys} gene with severe ultrastructural alterations of mitochondria in cardiomyocytes. *Jpn Circ J*, 2001, 65(7): 691-694. 
- [27] Wilson FH, Hariri A, Farhi A, Zhao HY, Petersen KF, Toka HR, Nelson-Williams C, Raja KM, Kashgarian M, Shulman GI, Scheinman SJ, Lifton RP. A cluster of metabolic defects caused by mutation in a mitochondrial tRNA. *Science*, 2004, 306(5699): 1190-1194.
- [28] Merante F, Myint T, Tein I, Benson L, Robinson BH. An additional mitochondrial tRNA^{Ile} point mutation (A-to-G at nucleotide 4295) causing hypertrophic cardiomyopathy. *Hum Mutat*, 1996, 8(3): 216-222. 3.0.CO;2-7 target="_blank"> 
- [29] Casali C, Santorelli FM, D'Amati G, Bernucci P, DeBiase L, DiMauro S. A novel mtDNA point mutation in maternally inherited cardiomyopathy. *Biochem Biophys Res Commun*, 1995, 213(2): 588-593. 
- [30] Tanaka M, Ino H, Ohno K, Hattori K, Sato W, Ozawa T, Tanaka T, Itoyama S. Mitochondrial mutation in fatal infantile cardiomyopathy. *Lancet*, 1990, 336(8728): 1452.
- [31] Zhu HY, Wang SW, Liu L, Chen R, Wang L, Gong XL, Zhang ML. Genetic variants in mitochondrial tRNA genes are associated with essential

- hypertension in a Chinese Han population. *Clin Chim Acta*, 2009, 410(1-2): 64-69. [crossref](#)
- [32] Taniike M, Fukushima H, Yanagihara I, Tsukamoto H, Tanaka J, Fujimura H, Nagai T, Sano T, Yamaoka K, Inui K, Okada S. Mitochondrial tRNA^{Leu} mutation in fatal car-diomyopathy. *Biochem Biophys Res Commun*, 1992, 186(1): 47-53. [crossref](#)
- [33] Merante F, Tein I, Benson L, Robinson BH. Maternally inherited hypertrophic cardiomyopathy due to a novel T-to-C transition at nucleotide 9997 in the mitochondrial tRNA(glycine) gene. *Am J Hum Genet*, 1994, 55(3): 437-446.
- [34] Shin WS, Tanaka M, Suzuki J, Hemmi C, ToyoOka T. A novel homoplasmic mutation in mtDNA with a single evolutionary origin as a risk factor for cardiomyopathy. *Am J Hum Genet*, 2000, 67(6): 1617-1620. [crossref](#)
- [35] Liu YQ, Li RH, Li ZB, Wang XJ, Yang L, Wang SW, Guan MX. Mitochondrial transfer RNAMet 4435A>G mutation is associated with maternally inherited hypertension in a Chinese pedigree. *Hypertension*, 2009, 53(6): 1083-1090. [crossref](#)
- [36] Li RH, Liu YQ, Li ZB, Yang L, Wang SW, Guan MX. Failures in mitochondrial tRNAMet and tRNAGln metabolism caused by the novel 4401A>G mutation are involved in essential hypertension in a Han Chinese Family. *Hypertension*, 2009, 54(2): 329-337. [crossref](#)
- [37] Opdal SH, Vege ÅS, Egeland T, Musse MA, Rognum TO. Possible role of mtDNA mutations in sudden infant death. *Pediatr Neurol*, 2002, 27(1): 23-29. [crossref](#)
- [38] Hattori Y, Nakajima K, Eizawa T, Ehara T, Koyama M, Hirai T, Fukuda Y, Kinoshita M. Heteroplasmic mito-chondrial DNA 3310 mutation in NADH dehydrogenase subunit 1 associated with type 2 diabetes, hypertrophic cardiomyopathy, and mental retardation in a single patient. *Diabetes Care*, 2003, 26(3): 952-953. [crossref](#)
- [39] Jaksch M, Hofmann S, Kaufhold P, Obermaier-Kusser B, Zierz S, Gerbitz KD. A novel combination of mitochon-drial tRNA and ND1 gene mutations in a syndrome with MELAS, cardiomyopathy, and diabetes mellitus. *Hum Mutat*, 1996, 7(4): 358-360. 3.0.CO;2-1 target="_blank">
[crossref](#)
- [40] Takagi K, Yamada Y, Gong JS, Sone T, Yokota M, Tanaka M. Association of a 5178C→A(Leu237Met) polymor-phism in the mitochondrial DNA with a low prevalence of myocardial infarction in Japanese individuals. *Athero-sclerosis*, 2004, 175(2): 281-286.
- [41] Ware SM, El-Hassan N, Kahler SG, Zhang Q, Ma YW, Miller E, Wong B, Spicer RL, Craigen WJ, Kozel BA, Grange DK, Wong LJ. Infantile cardiomyopathy caused by a mutation in the overlapping region of mitochondrial ATPase 6 and 8 genes. *J Med Genet*, 2009, 46(5): 308-314. [crossref](#)
- [42] Jonckheere AI, Hogeveen M, Nijtmans LG, van den Brand MA, Janssen AJ, Diepstra JH, van den Brandt FC, van den Heuvel LP, Hol FA, Hofste TG, Kapusta L, Dillmann U, Shamdeen MG, Smeitink JA, Rodenburg RJ. A novel mi-tochondrial ATP8 gene mutation in a patient with apical hypertrophic cardiomyopathy and neuropathy. *J Med Genet*, 2008, 45(3): 129-133.
- [43] Valnot I, Kassis J, Chretien D, de Lonlay P, Parfait B, Munnich A, Kachaner J, Rustin P, Rötig A. A mitochondrial cytochrome *b* mutation but no mutations of nuclearly encoded subunits in ubiquinol cytochrome *c* reductase (complex III) deficiency. *Hum Genet*, 1999, 104(6): 460-466. [crossref](#)
- [44] Khogali SS, Mayosi BM, Beattie JM, McKenna WJ, Watkins H, Poulton J. A common mitochondrial DNA variant associated with susceptibility to dilated cardiomyopathy in two different populations. *Lancet*, 2001, 357(9264): 1265-1267.
- [45] Levinger L, Mörl M, Florentz C. Mitochondrial tRNA 3' end metabolism and human disease. *Nucl Acids Res*, 2004, 32(18): 5430-5441. [crossref](#)
- [46] Lopez-Campistrous A, Hao L, Xiang W, Ton D, Semchuk P, Sander J, Ellison MJ, Fernandez-Patron C. Mitochondrial dysfunction in the hypertensive rat brain: respiratory complexes exhibit assembly defects in hypertension. *Hypertension*, 2008, 51(2): 412-4129. [crossref](#)
- [47] Postnov YV, Orlov SN, Budnikov YY, Doroschuk AD, Postnov AY. Mitochondrial energy conversion disturbance with decrease in ATP production as a source of systemic arterial hypertension. *Pathophysiology*, 2007, 14(3-4): 195-204. [crossref](#)
- [48] Sprinzl M, Horn C, Brown M, Ioudovitch A, Steinberg S. Compilation of tRNA sequences and sequences of tRNA genes. *Nucl Acids Res*, 1998, 26(1): 148-153. [crossref](#)
- [49] Björk GR. Biosynthesis and function of modified nucleotides. In: Söll D, RajBhandary UL, eds. tRNA: Structure, Biosynthesis and Function. Washington, DC: ASM Press, 1995: 165-206.
- [50] Sundaram M, Durant PC, Davis DR. Hypermodified nucleosides in the anticodon of tRNA(Lys) stabilize a ca-nonical U-turn structure. *Biochemistry*, 2000, 39(41): 12575-12584. [crossref](#)
- [51] Li JN, Esberg B, Curran JF, Björk GR. Three modified nucleosides present in the anticodon stem and loop influ-ence the *in vivo* aa-tRNA selection in a tRNA-dependent manner. *J Mol Biol*, 1997, 271(2): 209-221. [crossref](#)
- [52] Yarus M, Cline SW, Wier P, Breeden L, Thompson RC. Actions of the anticodon arm in translation on the phenotypes of RNA mutants. *J Mol Biol*, 1986, 192(2): 235-255. [crossref](#)
- [53] Florentz C, Sohm B, Tryoen-Tóth P, Pütz J, Sissler M. Human mitochondrial tRNAs in health and disease. *Cell Mol Life Sci*, 2003, 60(7): 1356-1375. [crossref](#)
- [54] Ojala D, Montoya J, Attardi G. tRNA punctuation model of RNA processing in human mitochondria. *Nature*, 1981, 290(5806): 470-474.
- [55] Guan MX, Enriquez JA, Fischel-Ghodsian N, Puranam RS, Lin CP, Maw MA, Attardi G. The deafness-associated mitochondrial DNA mutation at position 7445, which affects tRNASer(UCN) precursor processing, has long-range effects on NADH dehydrogenase ND6 subunit gene expression. *Mol Cell Biol*, 1998, 18(10): 5868-5879.

- [56] Wang S, Li R, Fettermann A, Li Z, Qian Y, Liu Y, Wang X, Zhou A, Mo JQ, Yang L, Jiang P, Taschner A, Rossmanith W, Guan MX. Maternally inherited essential hypertension is associated with the novel 4263A>G mutation in the mitochondrial tRNA^{Ile} gene in a large Han Chinese family. *Circ Res*, 2011, 108(7): 862-870. 
- [57] Holzmann J, Frank P, Löffler E, Bennett KL, Gerner C, Rossmanith W. RNase P without RNA: identification and functional reconstitution of the human mitochondrial tRNA processing enzyme. *Cell*, 2008, 135(3): 462-474.
- [58] Levinger L, Jacobs O, James M. *In vitro* 3' -end endonucleolytic processing defect in a human mitochondrial tRNA^{Ser(UCN)} precursor with the U7445C substitution, which causes non-syndromic deafness. *Nucl Acids Res*, 2001, 29(21): 4334-4340. 
- [59] Hofhaus G, Shakeley RM, Attardi G. Use of polarography to detect respiration defects in cell cultures. *Methods Enzymol*, 1996, 264: 476-483. 
- [60] Quigley GJ, Rich A. Structural domains of transfer RNA molecules. *Science*, 1976, 194(4267): 796-806.
- [61] Kim SH, Quigley GJ, Suddath FL, McPherson A, Sneden D, Kim JJ, Weinzierl J, Rich A. Three-dimensional structure of yeast phenylalanine transfer RNA: folding of the polynucleotide chain. *Science*, 1973, 179(70): 285-288.
- [62] Tomari Y, Hino N, Nagaïke T, Suzuki T, Ueda T. Decreased CCA-addition in human mitochondrial tRNAs bearing a pathogenic A4317G or A10044G mutation. *J Biol Chem*, 2003, 278(19): 16828-16833. 
- [63] Li XM, Fischel-Ghodsian N, Schwartz F, Yan QF, Friedman RA, Guan MX. Biochemical characterization of the mitochondrial tRNA^{Ser(UCN)} T7511C mutation associated with nonsyndromic deafness. *Nucl Acids Res*, 2004, 32(3): 867-877. 
- [1] 许睿玮, 严卫丽.原发性高血压全基因组关联研究进展[J]. 遗传, 2012,34(7): 793-809
- [2] 吴志俊, 金玮, 张凤如, 刘艳.利钠肽家族基因与心血管疾病研究新进展[J]. 遗传, 2012,34(2): 127-133
- [3] 李宗斌, 刘昱圻, 李彦华, 陈瑞, 王琳, 朱庆磊, 李泱, 王士雯.中国汉族原发性高血压患者线粒体tRNA基因突变[J]. 遗传, 2011,33(6): 601-606
- [4] 张子波, 于丽军, 杨康鹏, 徐良慰, 盛天听, 郝萍, 王玉萍, 孟繁平.延边朝鲜族和汉族脂联素启动子SNPs与原发性高血压的相关性[J]. 遗传, 2011,33(1): 54-59
- [5] 马厚勋, 谢正祥, 牛永红, 李章勇, 周平.汉族人群NOS3 A-922G、NOS3 T-786C 与NOS3 G894T SNP的等位基因及其组合分布与高血压的相关性[J]. 遗传, 2006,28(1): 3-10
- [6] 余运贤, Scott A. Venners, 牛天华, 陈常中, 黄爱群, 张岩, 冯雁, 李栋, 邢厚询, 吴迪, 彭少杰, 徐希平, .原发性高血压患者 α -Adducin基因单核苷酸多态性 (*Gly460Trp*) 与血清胆红素含量相关性[J]. 遗传, 2004,31(9): 941-949
- [7] 刘玲玲, 谭端军, 王士雯.重建邻接关系树评估原发性高血压患者的遗传性[J]. 遗传, 2004,26(4): 441-445
- [8] 王云英, 张七一, 王培林.ACE基因多态性与高血压肾脏损害及PAI-1的关系[J]. 遗传, 2003,25(6): 641-644
- [9] 汪振诚, 王学敏, 金由辛, 缪明永, 韩伟国, 焦炳华.人线粒体tRNA^{Leu(UUR)}基因A3243G点突变对其亮氨酰化活性的影响[J]. 遗传, 2003,25(4): 383-387
- [10] 王爱玲, 余元勋, 徐岩, 刘萍, 程自平, 张立, 吴继雄, 孙国梅, 陈森.高血压患者血管紧张素原基因调控及与细胞因子关系的研究[J]. 遗传, 2003,30(10): 978-982
- [11] 戴书萍, 时景璞, 丁茜, 王海龙, 董凌月, 孙迪, 房凯, 赵彦艳.中国东北高血压病高发地区人群中G蛋白 β -3|亚单位825C/T多态分析[J]. 遗传, 2002,29(4): 294-298
- [12] 牟奕, 单祥年, 严明, 刘宁生, 鲁晓萱, 邢光前, 卜行宽, 杨焕明.一个母系遗传非综合征耳聋大家系mtDNA序列分析[J]. 遗传, 2001,23(5): 401-724
- [13] 杨文杰, 顾东风.人类原发性高血压候选基因的研究进展[J]. 遗传, 2001,23(5): 487-277
- [14] 李少英, 李月秋, 吴雅冬, 罗佳滨, 王秀岩, 马继红, 吕学洗.血管紧张素原基因多态位点AGT174与原发性高血压相关性的研究[J]. 遗传, 1998,20(4): 18-20