e0064  MICRO RNAS ARE INVOLVED IN THE PSYCHOLOGICAL STRESS-INDUCED CARDIAC DISORDERS IN RATS

doi:10.1136/hrt.2010.208967.64


**Objective** Psychological stress has become an important factor in the development of cardiac disorders. MicroRNAs (miRNA) have been implicated in regulation of cardiovascular diseases. Therefore, we investigated whether miRNAs are involved in the psychological stress-induced cardiac disorders.

**Methods** Stress rat models were established by complex stimulation at different times during the daytime for 2 weeks. Body weight, blood pressure and ECG were measured every 3 days during processing stress. Adrenocorticotropic hormone (ACTH) was measured by using ELISA. Cardiac changes were detected by HE staining and electronic microscopy. MicroRNA microarray was used for analysing the differential expression of miRNAs.

**Results** After psychological stress, rats displayed change in behaviour. Body weight increased slowly and systolic blood pressure increased significantly in stress group from the 6th day to the 15th day. The ECG of all rats was normal before experiment. 2 weeks after stimulation the ECG recorded of different individuals in stress group showed different arrhythmia, including sinus tachycardia, atrial premature contraction, ventricular arrhythmia, and ST-T changes. Hypothalamic-pituitary-adrenal activity increased in stress group compared to control group by detecting ACTH. And the ultrastructure and histology showed injury changes in stress group. Compared to the control group, there were 55 different miRNAs in stress model including upregulation of 20 and downregulation of left, among which miRNA-141, miRNA-382, miRNA-219-5p and miRNA-296 up-regulate, miRNA-153a and miRNA-466b are significant down-regulate.

**Conclusions** Complex stimulation can induce psychological stress, which can cause cardiac injury. miRNAs change in stress rat models, including upregulation of miR-141, miR-382, miR-219-5p and miR-296, and significant downregulation of miR-153a and miR-466b, which may play important roles in psychological stress-induced cardiac disorders. (Supported by research grant NSFC 30940041).

e0065  THE ASSOCIATION BETWEEN THE SINGLE NUCLEOTIDE POLYMORPHISMS OF MATRIX METALLOPROTEINASES AND THE CAROTID ATHEROSCLEROSIS IN PATIENTS WITH ESSENTIAL HYPERTENSION

doi:10.1136/hrt.2010.208967.65

1Yan-Ping Ma, 2Xin-Juan Xu, 2Fang Zhao, 2Guang-Mei Hu, 2Jun-Shi Zhang. 1Department of Hypertension, The First Affiliated Hospital of Medical University, Urumchi Xinjiang, China; 2Department of Hypertension, The Xinjiang Armed Police Hospital, Urumchi Xinjiang, China

**Objective** To investigate the relationship between the single nucleotide polymorphisms (SNP) of matrix metallo proteases (MMP-2 -735C/T; MMP-3 -1171 5A/6A) and the carotid atherosclerosis (CAS) in Chinese Han and Uygur populations with EH.

**Methods** The study comprised 276 Han nationality and 212 Uygur participants, who were divided into two groups: CAS (n=293) and NS (n=195). Genotypes were detected by PCR-RFLP and their frequencies were determined.

**Results** (1) The frequencies of MMP-2 TT genotype and T allele in CAS were higher than in NS (Han: X²=11.441, p=0.003; Uygur: X²=28.255, p=0.000). In NS, the frequencies of TT genotype and T allele in Han were higher than in Uygur (X²=12.809, p=0.001)). (2) The frequencies of MMP-3 6A/6A genotype and 6A allele in CAS were higher than NS (Han: X²=7.523, p=0.024; Uygur: X²=6.474, p=0.039). The frequencies of MMP-3 6A/6A genotype and 6A allele in Han were higher than Uygur (CAS: X²=26.230, p=0.000; NS: X²=18.809, p=0.000). (3) The single gene analysis showed Han individuals with CT or TT genotypes had 2.2-fold risk and Han individuals with 6A/6A genotypes had 1.85-fold risk suffering from CAS. Han individuals with both T allele and 6A/6A genotypes had 3.17-fold risk suffering from CAS. The single gene analysis showed that Uygur individuals with CT or TT genotypes had 5.04-fold risk suffering from CAS. Uygur individuals with 6A/6A genotypes had 2.20-fold risk suffering from CAS. Uygur individuals with both T allele and 6A/6A genotypes had 3.20-fold risk suffering from CAS. (4) According to MMP-2 genotypes, Han individuals with MMP-2 CT+TT genotypes had higher LDL and lower HDL levels than CAS genotype (LDL:2.9mmol/l vs 2.6 mmol/l; HDL:1.2 mmol/l vs 1.0 mmol/l). Uygur individuals with CT+TT genotypes had higher TG levels than CAS genotype (Uygur : 2.5 mmol/l vs 1.6 mmol/l; NS : 3.9 mmol/l vs 2.0 mmol/l). According to MMP-3, Han individuals with 6A/6A genotype had higher T-CHOL and LDL levels than 5A/5A+5A/6A genotypes in NS group (T-CHOL : 4.6 mmol/l vs 4.2 mmol/l; LDL : 2.3 mmol/l vs 2.2 mmol/l). (5) The binary logistic regression analysis showed MMP-2 CT+TT genotypes were the risk factors for CAS in individuals with EH (Uygur: OR=9.65; Han : OR=2.076). MMP-3 6A homogeneses were the risk factors for CAS in individuals with EH. The SNP of MMP-2 CT+TT and MMP-3 6A homogeneses had a combined influence on the incidence of CAS in Han individuals with EH.

**Conclusions** (1) Han and Uygur individuals had differential distribution of MMPs. (2) The SNP of MMP-2 -735C/T is associated with CAS in individuals with EH. The MMP-2 T allele may be a risk factors on CAS in individuals with EH. The SNP of MMP-3 -1171 5A/6A is associated with CAS in Han individuals with EH. The 6A allele may be a risk factors on CAS in Han individuals with EH.

e0066  ISCHAEMIC PRECONDITIONING COMBINED WITH GRADUAL REPERFUSION OFFERS NO ADDITIONAL BENEFIT ON MITOCHONDRIAL PERMEABILITY PORE OVER PRECONDITIONING OR GRADUAL REPERFUSION ALONE

doi:10.1136/hrt.2010.208967.66

Dong Chen, Lihuan Li. Fuwai Hospital

**Objective** Recent investigations demonstrate that ischaemic preconditioning and post-conditioning can reduce infarct size to the same degree. After reflow, opening of the mitochondrial permeability transition pore (mPTP) has been involved in lethal reperfusion injury. We hypothesised that the combination of ischaemic preconditioning and post-conditioning would result in greater...