

Genome-wide gene expression microarray identifies novel genes related to endothelial tight junctions and apoptosis in patients with obstructive sleep apnea

Yung-Che Chen^{1,2}, Meng-Chih Lin^{1,2*}, Mao-Chang Su^{1,2}, Chien-Hung Chin^{1,2}, Chia-Wei Liou⁴, Ya-Chun Chang^{1,2}, Kuo-Tung Huang^{1,2}, Kuang-Den Chen³, Chang-Chun Hsiao⁵, Chung-Jen Chen², Ting-Wen Chen⁵, Ting-Ya Wang¹, Division of Pulmonary and Critical Care Medicine¹, Sleep Center², Center of Translational Research in Biomedical Sciences³, Department of Neurology⁴, Kaohsiung Chang Gung Memorial Hospital and Chang Gung University College of Medicine, Kaohsiung, Taiwan; Graduate Institute of Clinical Medical Sciences⁵, Chang Gung University, Taiwan

Purpose: We aim to identify novel molecular mechanisms by which intermittent hypoxia with re-oxygenation (IHR) leads to adverse consequences, such as hypertension and excessive daytime sleepiness (EDS) in patients with obstructive sleep apnea (OSA).

Materials and Methods: We analyzed whole-genome gene expression profiles of peripheral blood mononuclear cells from 48 patients with sleep-disordered breathing stratified into four groups: primary snoring (PS), moderate to severe OSA (MSO), very severe OSA (VSO), and very severe OSA patients with long-term continuous positive airway pressure (CPAP) treatment (VSOC).

Results: Comparisons of the microarray gene expression data identified eight genes up-regulated with OSA and down-regulated with CPAP treatment, and five genes down-regulated with OSA and up-regulated with CPAP treatment (all p values < 0.05). Protein expression levels of 2 genes related to endothelial tight junctions (AMOT P130, and PLEKHH3; both p values < 0.001), and 3 genes related to anti-or pro-apoptosis (ADAR1 P150, BIRC3, and GALIG; all p values < 0.05) were all increased in the VSO group, while AMOT P130 was further increased, and PLEKHH3, BIRC3, and ADAR1 P150 were all decreased in the VSOC group. AMOT P130 protein expression was increased in OSA patients with EDS, BIRC3 protein expression was decreased in OSA patients with hypertension, and GALIG protein expression was increased in OSA patients with chronic kidney disease (CKD). In vitro IHR (7 cycles/day, 4 days) stimuli resulted in over-expression of ADAR1 P150.

Conclusions: We identified a novel association between AMOT/PLEKHH3/BIRC3/ADAR1/GALIG over-expressions and high Apnea-Hypnea Index in OSA patients. AMOT and GALIG may constitute an important determinant for the development of EDS and CKD, respectively, while BIRC3 may play a protective role in the development of hypertension in OSA.

中文題目 全基因體微陣基因表現全貌辨識出調控阻塞性睡眠呼吸中止症的新分子機轉：內皮細胞緊密接合和細胞凋零

作者 陳永哲^{1,2}，林孟志^{1,2}，蘇茂昌^{1,2}，陳廣典³，蕭長春⁵，陳忠仁⁵，秦建弘^{1,2}，劉嘉為⁴，張雅淳^{1,2}，黃國棟^{1,2}，張稔杰³，王亭雅¹，林詠詠¹，鄭宜欣¹，陳婷玟⁵

服務單位 高雄長庚呼吸胸腔內科，睡眠中心，轉譯醫學中心，風濕內科，神經內科；長庚大學臨床醫學研究所

發表方式 原著論文 病例報告論文