

European Respiratory Society Annual Congress 2012

Abstract Number: 1687

Publication Number: P1496

Abstract Group: 4.3. Pulmonary Circulation and Pulmonary Vascular Disease

Keyword 1: Pulmonary hypertension **Keyword 2:** Biomarkers **Keyword 3:** Molecular pathology

Title: Circulating microRNA signature and its novel involvement in pathogenesis of chronic thromboembolic pulmonary hypertension

Dr. Lijuan 11422 Guo lg83330102@163.com^{1,2,4}, Prof. Dr Jun 11423 Cai caijun7879@yahoo.com.cn MD⁵, Dr. Lei 11424 Wang wl860806wb@163.com^{1,2,4}, Dr. Jifeng 11425 Li lijifengcyh@163.com MD^{1,2,4}, Mr. Jie 11426 Liu huxishl@ccmu.edu.cn², Prof. Dr Yuanhua 11433 Yang yyh1031@sina.com MD^{1,4}, Prof. Baosen 11435 Pang pangbaosen@163.com^{1,4}, Prof. Dr Zhenguo 11441 Zhai zhaizhenguo@gmail.com MD^{1,4}, Dr. Yan 11443 Liu liuyanok2@hotmail.com MD^{4,5}, Dr. Song 11444 Gu gusong@163.com MD^{4,5}, Prof. Dr Chen 11445 Wang cyh-birm@263.net MD^{1,3,4} and Prof. Jun 11541 Wang wang_jun@ccmu.edu.cn^{1,2,4}. ¹ Beijing Key Laboratory of Respiratory and Pulmonary Circulation Disorders, Beijing Chao-Yang Hospital, Beijing, China, 100020 ; ² Department of Physiology, Capital Medical University, Beijing, China, 100069 ; ³ Department of Beijing Hospital, Ministry of Health, Beijing, China, 100730 ; ⁴ Beijing Institute of Respiratory Medicine, Beijing Chao-Yang Hospital, Beijing, China, 100020 and ⁵ Cardiology Center, Beijing Chao-Yang Hospital, Beijing, China, 100020 .

Body: Chronic thromboembolic pulmonary hypertension (CTEPH) is a progressive disease characterized by multiple etiology and mechanism. Circulating miRNA is partially derived from cells affected by disease and therefore can serve as potential biomarker and reflect the pathogenesis of this disease. In present study, we compared miRNA expression in plasma from 10 CTEPH patients and 10 healthy control subjects by microarray, and fourteen miRNAs were identified to be differentially expressed.

Selectively, five of the differentially expressed miRNAs were further validated in an independent 40 pairs of subjects by stem-loop qRT-PCR, among which let-7b and miR-22 were downregulated to about 25% in CTEPH patients. Endothelin-1 (ET-1) and transforming growth factor beta receptor 1 (TGFB1) was the direct targets of let-7b by reporter assay, and plasma ET-1 level was reversely correlated to let-7b. TGFB1 was further required for induction of ET-1 in endothelial cells.