33. Molecules, polymorphisms and resistance to TB
Saturday, 05 December 2015, 12:30-13:30

Chair: To be confirmed

Track: TB epidemiology

**PC-973-05**  Standardized mycobacterial interspersed repetitive units-variable number of tandem repeat genotyping of Mycobacterium tuberculosis in Taiwan
P-C. Chuang, R Jou  (*Taiwan*)

**PC-974-05**  Polymorphisms in the STAT4 gene and tuberculosis susceptibility in Chinese Han people
Q Wu, Yi Ji, Can Wu, Yu Wang, Qing He  (*China*)

**PC-975-05**  Genetic polymorphisms of glutathione S-transferase P1 and the incidence of anti-tuberculosis drug-induced hepatitis in a Chinese cohort
J He, Quan Wu, Yu Wang, Can Wu, Yi Ji  (*China*)

**PC-976-05**  Key population for tuberculosis intensified case finding identified through national TB prevalence survey
Db Lolong, Os Simamarta, B Dwihardiani, M Farid, Lam Pangaribuan, A V Siagian  (*Indonesia*)

**PC-977-05**  Trends of fluoroquinolone resistance in Mycobacterium tuberculosis mirror resistance trends among community pathogens
S Shakoor, J Farooqi, K Jabeen, F Malik, R Hasan  (*Pakistan*)

**PC-978-05**  Key genotypes of M. tuberculosis responsible for the spread of multi- and extensively drug-resistant disease in the Sakha Republic, Yakutia

**PC-979-05**  WGS reveals genetic heterogeneity and suggests the role of selective bottleneck in defining the population structure of Mycobacterium tuberculosis
M De Vos, P Black, E Louw, R Van Der Merwe, S Sampson, P Van Helden, R M Warren, P Arnab  (*South Africa*)

**PC-980-05**  Treatment-based risk factors for tuberculosis drug resistance in Bolivia
A Flynn, J Carlos Mollinedo, C L Daley, A Faino, R Flick, E Casso, M Camacho De Colque  (*USA, Malawi, Bolivia*)