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

Chair: Susan DORMAN (USA)
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)