

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**
Mk Vinokurova, N.e. Evdokimova, G.i. Alekseeva, A.f. Kravchenko, E.d. Savilov, O.b. Ogarkov, S Zhdanova (*Russian Federation*)
- 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*)