Journal publications: (from Pubmed: http://paed.hku.hk/genome)
(*: Corresponding or co-corresponding author. #: Equal contribution/co-first author.)


12. Jing Zhang, Lu Zhang, Yan Zhang, Jing Yang, Mengbiao Guo, Liangdan Sun, Hai-Feng Pan, Nattiya Hirankarn, Dingge Ying, Shuai Zeng, Tsu Leung Lee, Chak Sing Lau, Tak Mao Chan, Alexander Moon Ho Leung, Chi Chiu Mok, Sik Nin Wong, Ka Wing Lee, Marco Hok Kung Ho, Pamela Pui Wah Lee, Brian Hon-Yin Chung, Chun Yin Chong, Raymond Woon Sing Wong, Mo Yin Mok, Wilfred Hing Sang Wong, Kwok Lung Tong, Niko Kei Chiu Tse, Xiang-Pei Li, Yingyos Avihingsanon, Pornpimol Riantavorn, Thavatchai Deekajorndej, Kanya Suphapeetiporn, Vorasuk Shotelersuk, Shirley King Yee Ying, Samuel Ka Chun Fung, Wai Ming Lai, Maria-Mercè Garcia-Barceló, Stacey S. Cherny, Pak Chung Sham, Yong Cui, Sen Yang, Dong Qing Ye, Xue-Jun Zhang, *Yu Lung Lau, *Yang W. Gene-based meta-analysis of GWAS data identifies independent SNPs in ANXA6 as associated with SLE in Asian populations, Arthritis Rheumatol. 2015 Nov;67(11):2966-77. doi: 10.1002/art.39275.


62. **Yang W**, Wang Z, Wang L, Sham PC, Huang P, Lau YL. Predicting the number and sizes of IBD regions among family members and evaluating the family size.


