

2005 (2)

1. Rojas RE, Chervenak KA, Thomas J, et al. Vdelta2+ gammadelta T cell function in *Mycobacterium* tuberculosis- and HIV-1-positive patients in the United States and Uganda: application of a whole-blood assay. *J Infect Dis* 2005;192 (10):1806-14.
2. Stein CM, Nshuti L, Chiunda AB, et al. Evidence for a major gene influence on tumor necrosis factor-alpha expression in tuberculosis: path and segregation analysis. *Hum Hered* 2005;60

1. Lancioni C, Nyendak M, Kiguli S, et al. CD8+ T cells provide an immunologic signature of tuberculosis in young children. *Am J Respir Crit Care Med* 2012;185 (2):206-12.
2. Mahan CS, Zalwango S, Thiel BA, et al. Innate and adaptive immune responses during acute M. tuberculosis infection in adult household contacts in Kampala, Uganda. *Am J Trop Med Hyg* 2012;86 (4):690-7.
3. Mupere E, Malone L, Zalwango S, et al. Lean tissue mass wasting is associated with increased risk of mortality among women with pulmonary tuberculosis in urban Uganda. *Ann Epidemiol* 2012;22 (7):466-73.

2013 (5)

3. Sobota RS, Stein CM, Kodaman N, et al. A Locus at 5q33.3 Confers Resistance to Tuberculosis in Highly Susceptible Individuals. *Am J Hum Gene* 2016;98 (3):514-24.

2017 (4)

1. Bark CM, Manceur AM, Malone LL, et al. Identification of Host Proteins Predictive of Early Stage Mycobacterium tuberculosis Infection. *EBioMedicin* 2017;21: 150-157.
2. Martinez L, Handel A, Shen Y, et al. A Prospective Validation of a Clinical Algorithm to Detect Tuberculosis in Child Contacts. *Am J Respir Crit Care Me* 2017.
3. Seshadri C, Sedaghat N, Campo M, et al. Transcriptional networks are associated with resistance to Mycobacterium tuberculosis infection. *PLoS On* 2017;12 (4):e0175844.
4. Sobota RS, Stein CM, Kodaman N, et al. A chromosome 5q31.1 locus associates with tuberculin skin test reactivity in HIV-positive individuals from tuberculosis hyper-endemic regions in east Africa. *PLoS Gene* 2017;13 (6):e1006710.

2018 (6)

1. Fluegge K, Malone LL, Nsereko M, et al. Impact of geographic distance on appraisal delay for active TB treatment seeking in Uganda: a network analysis of the Kawempe Community Health Cohort Study. *BMC Public Healt* 2018;18 (1):798.
2. Igo RP, Jr., Hall NB, Malone LL, et al. Fine-mapping analysis of a chromosome 2 region linked to resistance to Mycobacterium tuberculosis infection in Uganda reveals potential regulatory variants. *Genes Immur* 2018.
3. Martinez L, Handel A, Shen Y, et al. A Prospective Validation of a Clinical Algorithm to Detect Tuberculosis in Child Contacts. *Am J Respir Crit Care Me* 2018;197 (9):1214-1216.
4. Martinez L, Shen Y, Handel A, et al. Effectiveness of WHO's pragmatic screening algorithm for child contacts of tuberculosis cases in resource-constrained settings: a prospective cohort study in Uganda. *Lancet Respir Me* 2018;6(4):276-286.
5. Simmons JD, Stein CM, Seshadri C, et al. Immunological mechanisms of human resistance to persistent Mycobacterium tuberculosis infection. *Nat Rev Immunol* 2018.
6. Stein CM, Zalwango S, Malone LL, et al. Resistance and Susceptibility to Mycobacterium Tuberculosis Infection and Disease in Tuberculosis Households in Kampala, Uganda. *Am J Epidemiol* 2018.

2019 (7)

1. Handel A, Martinez L, Sekandi JN, et al. Evidence for supercoughers in an analysis of six tuberculosis cohorts from China, Peru, The Gambia and Uganda. *Int J Tuberc Lung Dis* 2019;23 (12):1286-1292.
2. Igo RP, Jr., Hall NB, Malone LL, et al. Fine-mapping analysis of a chromosome 2 region linked to resistance to Mycobacterium tuberculosis infection in Uganda reveals potential regulatory variants. *Genes Immur* 2019;20 (6):473-483.
3. Lu LL, Smith MT, Yu KKQ, et al. Publisher Correction: IFN-gamma-independent immune markers of Mycobacterium tuberculosis exposure. *Nat Mec* 2019;25 (7):1175.
4. Lu LL, Smith MT, Yu KKQ, et al. IFN-gamma-independent immune markers of Mycobacterium tuberculosis exposure. *Nat Mec* 2019;25 (6):977-987.
5. Muzanyi G, Mulumba Y, Mubiri P, Mayanja H, Johnson JL, Mupere E. Predictors of recurrent TB in sputum smear and culture positive adults: a prospective cohort study. *Afr Health Sci* 2019;19 (2):2091-2099.
6. Stein CM, Nsereko M, Malone LL, et al. Long-term Stability of Resistance to Latent Mycobacterium tuberculosis Infection in Highly Exposed Tuberculosis Household Contacts in Kampala, Uganda. *Clin Infect Dis* 2019;68 (10):1705-1712.
7. Wampande EM, Naniima P, Mupere E, et al. Genetic variability and consequence of Mycobacterium tuberculosis lineage 3 in Kampala-Uganda. *PLoS On* 2019;14 (9):e0221644.