

Type of study	Gene/Locus	Phenotype	Population(s)	References	SNPs	Result	# SNPs with P<0.05
Candidate gene	<i>ULK1</i>	TST positivity	Seattle	(Horne, Graustein et al. 2016, Shah, Musvosvi et al. 2017)	rs12297124 and rs7300908	Untested	
	<i>SLC6A3</i>	RSTR	Uganda	(Hall, Igo et al. 2015)	rs409588, rs456082, rs464061	p=0.9, Untested, Untested	
	<i>IL10</i>	TST positivity	Ghana	(Thye, Browne et al. 2009)	rs6703630, rs1800896, rs1800871	Untested, 0.2, 0.31	
Genome-wide	2q21-q24 (<i>GTDC1</i> and <i>ZEB2</i>)	RSTR (longitudinal)	Uganda	(Stein, Zalwango et al. 2008) (Cobat, Gallant et al. 2009)	rs74533435	1.09E-04	404
	5p13-q22			(Igo, Hall et al. 2018)	rs62375908	1.12E-04	885
	11p14 (<i>TST1</i>)	TST positivity (cross-sectional)	South Africa	(Cobat, Gallant et al. 2009)	rs293983	1.20E-03	67
	5p15 (<i>TST2</i>)	Reactivity (quantitative)		(Cobat, Hoal et al. 2013)	rs56345976	1.66E-04	346
	5q31 (including <i>IL9</i>)	TST positivity and reactivity	Tanzania and Uganda	(Sobota, Stein et al. 2017)	rs466157	6.99E-04	155
	10q26.2	TST and IGRA positivity	South Africa, France, Vietnam	(Quistretbet et al. pre-print)	rs17155120	6.20E-01	