

**Table S1.** Datasets included in the analysis<sup>a</sup>.

Year	Study	Accession	Countries
2012	Blouin <i>et al</i> , PLoS One (1)	ERP001885	Republic of Djibouti, Somalia, Sudan, Kenya
2013	Bryant <i>et al</i> , BMC (2)	ERP000111	Netherlands
2013	Clark <i>et al</i> , PLoS One (3)	ERP000520	Uganda
2013	Comas <i>et al</i> , Nat. Genet (4)	ERP001731	UK, Germany, USA, Africa(multiple countries), S. Asia, S.E. Asia, India, Uzbekistan, Netherlands Switzerland, Russia, Nepal, China, S. America
2013	Roetzer <i>et al</i> , PLoS One (5)	ERP006619	Germany
2013	Zhang <i>et al</i> , Nat. Genet (6)	SRP018402	China
2014	Casali <i>et al</i> , Nat. Genet (7)	ERP000192	Russia
2014	Perdigao <i>et al</i> , BMC Genomics (8)	ERP002611	Portugal
2015	Ali <i>et al</i> , PLoS One (9)	PRJEB7798	Pakistan
2015	Assuncao <i>et al</i> , e-Life (10)	ERP000436, ERP001072,	Malawi
2015	Merker <i>et al</i> , Nat. Genet (11)	PRJEB7281	Eurasia
2015	Walker <i>et al</i> , Lancet ID (12)	PRJNA282721	UK, Sierra Leone, S Africa, Germany, Uzbekistan
2015	Lee <i>et al</i> , PNAS (13)	SRP039605	Inuit
2016	Guimaraes <i>et al</i> , Microb Genom (14)	PRJEB8432	Ethiopia
2016	Stucki <i>et al</i> , Nat. Genet (15)	PRJEB11460	Africa, Nepal, Papua New Guinea, Switzerland
2017	Chatterjee <i>et al</i> , Tuberculosis (16)	SRP101835	India
2018	Chernyaeva <i>et al</i> , EID (17)	PRJNA352769	Russia

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