

Table S2. Crude and adjusted analysis comparing HIV-infected and HIV-negative tuberculosis patients born in Europe (n=233) across the four most frequent *Mycobacterium tuberculosis* lineages.

Main lineage	Unadjusted		Adjusted	
	OR (95% CI)	P value	OR (95% CI)	P value
Lineage 1	7.0 (0.42-114.6)	0.17	6.15 (0.30-125.53)	0.24
Lineage 2	4.6 (1.23-17.51)	0.023	3.22 (0.80-13.04)	0.1
Lineage 3	13.93 (2.43-79.71)	0.003	12.54 (2.04-77.21)	0.006
Lineage 4	1 (ref)	-	1 (ref)	-

95% CI, 95% confidence interval; OR, Odds Ratio

Model was adjusted for age, sex, and being born in Switzerland

Lineage 1: Indo-Oceanic lineage; Lineage 2: East-Asian lineage (includes Beijing strains); Lineage 3: Delhi/CAS; Lineage 4: Euro-American lineage