

S4 Table: Sample characteristics

Characteristic	PMN _{HITTIN} ^a , N = 17 [*]	PMN _{HIT} ^b , N = 11 [*]	p-value ^{**}
Sequencer			0.4
HiSeq	3/ 17 (18%)	4/ 11 (36%)	
NovaSeq	14/ 17 (82%)	7/ 11 (64%)	

^{*}n/ N (%); Mean (SD) ^{**}Fisher's exact test; Wilcoxon rank sum test

^aPMN_{HITTIN} (neutrophils from HIV-1-infected persistently TB, tuberculin and IGRA negative), ^bPMN_{HIT} (neutrophils from HIV-1-infected IGRA positive tuberculin positive)