

Table S1. Logistic regression modelling identifies demographic and clinical variables independently associating with HIV-SN in Africans and Indonesians.

Variable	Odds Ratio	P Value	95% CI
African Optimal Model: $n = 71^a$, $p = 0.0007$, Pseudo $R^2 = 0.18$			
Weight (kg)	1.04	0.03	1.00–1.08
History of Tuberculosis	4.26	0.07	0.90–20.03
Nadir CD4 T-cells/ μ L	1.00	0.03	0.99–1.00
Indonesian Optimal Model: $n = 195^a$, $p = 0.0006$, Pseudo $R^2 = 0.08$			
Current CD4 T-cells/ μ L	1.00	0.01	0.99–1.00
>500 copies HIV RNA/mL	3.80	0.04	1.10–13.15

^a excluding samples with missing demographic, clinical and/or genotype data.