



Figure 2-figure supplement 1. Regression analyses to identify variables associated with cell-associated HIV unspliced (US) RNA and total HIV DNA levels in the AIMS cohort, taking into account either (A) duration of cumulative virological suppression on ART, or (B) duration of current ART regimen, prior to the measurements. Effect sizes and 95% confidence intervals for US RNA are plotted as log₁₀ copies per microgram of total cellular RNA and for total DNA as log₁₀ copies per million PBMC. Effect sizes were obtained by fitting generalized linear models. Variables associated with HIV RNA or DNA with p values <0.1 in the univariable analyses were included in the multivariable analyses.