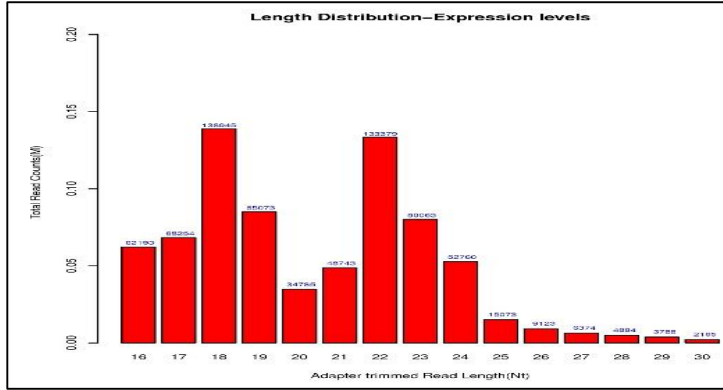
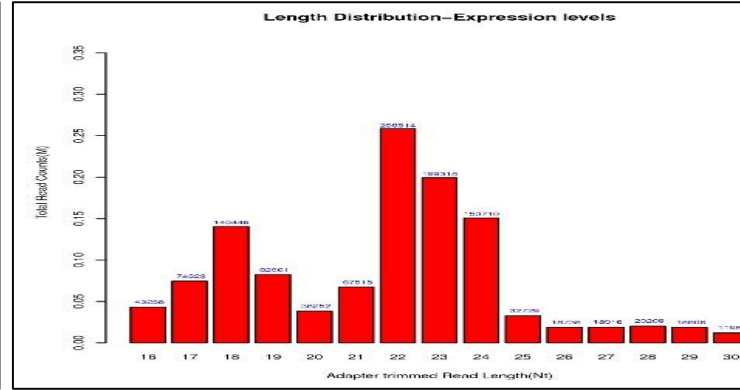


Control

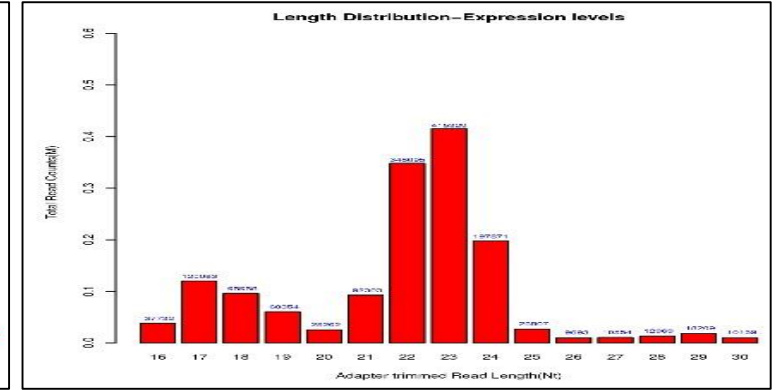
Donor 1



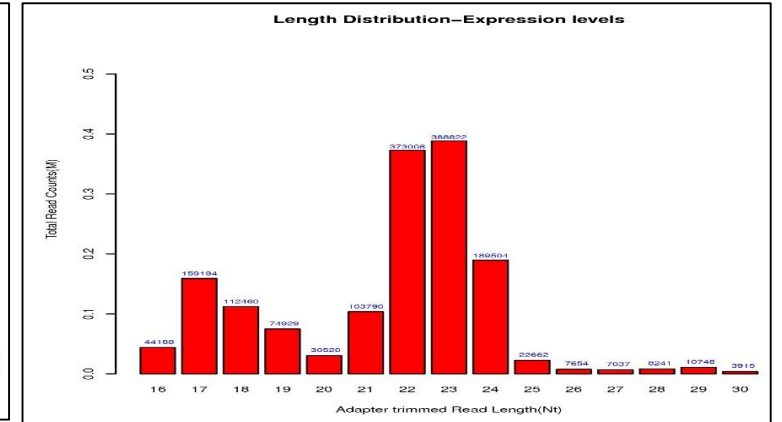
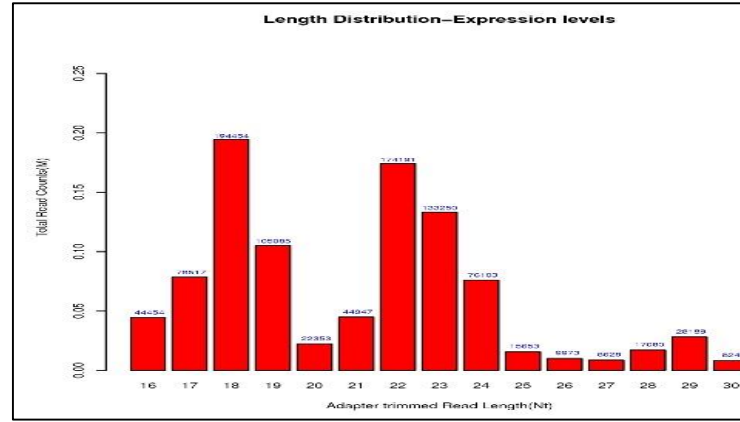
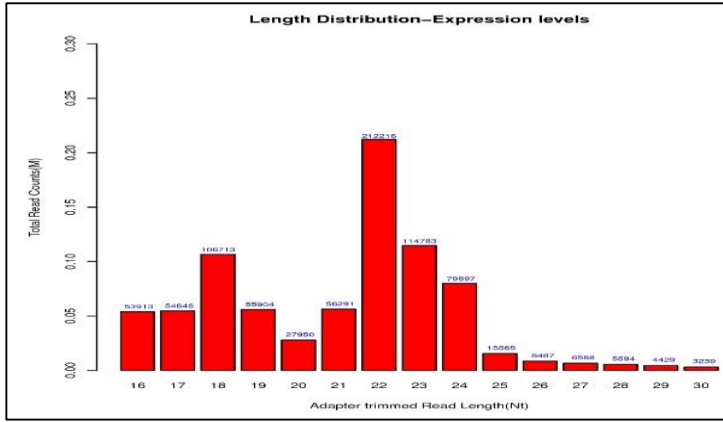
Donor 2



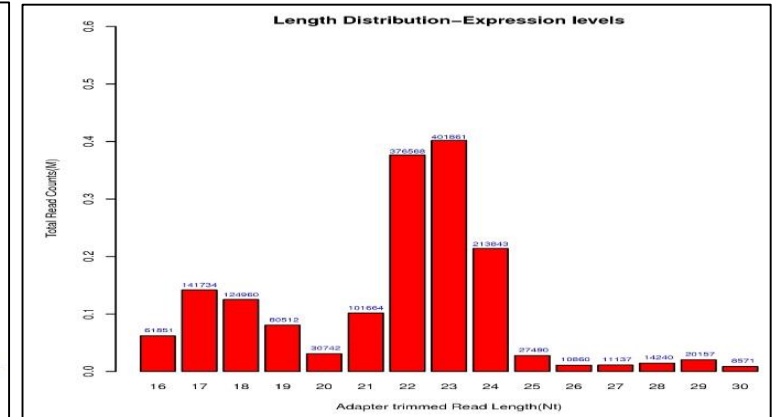
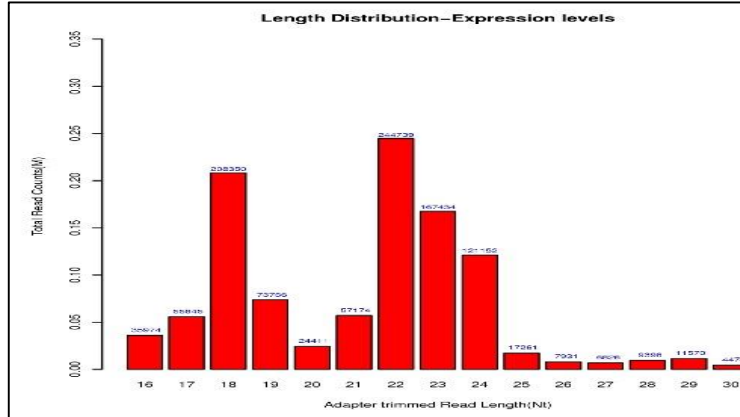
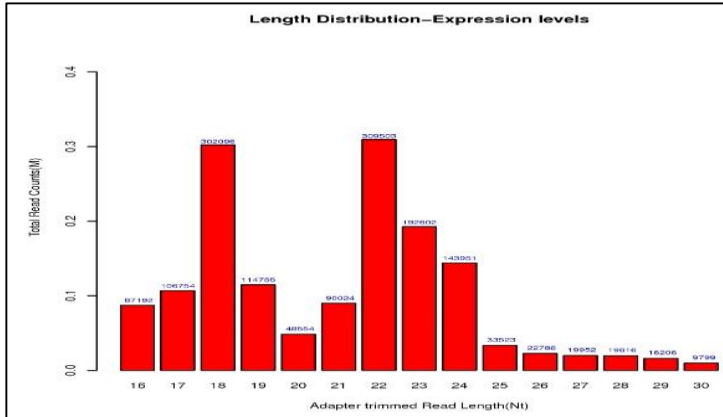
Donor 3



HIV-1

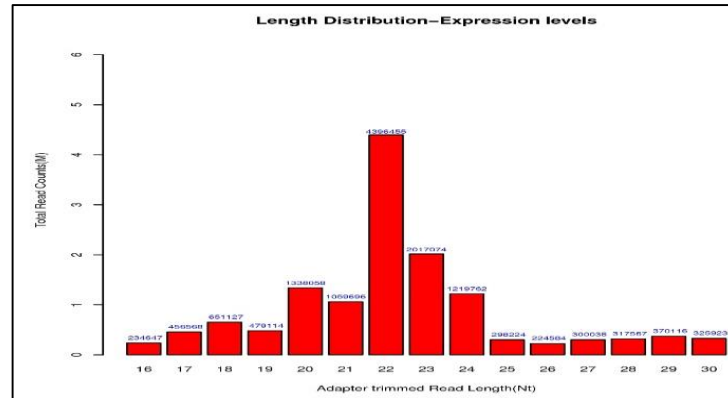


HIV-2

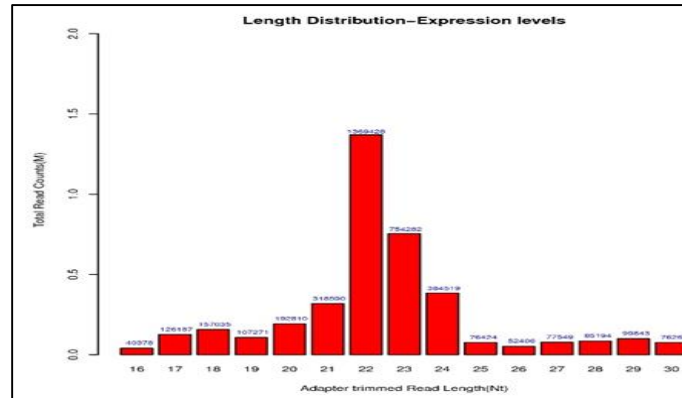


(A) MDM

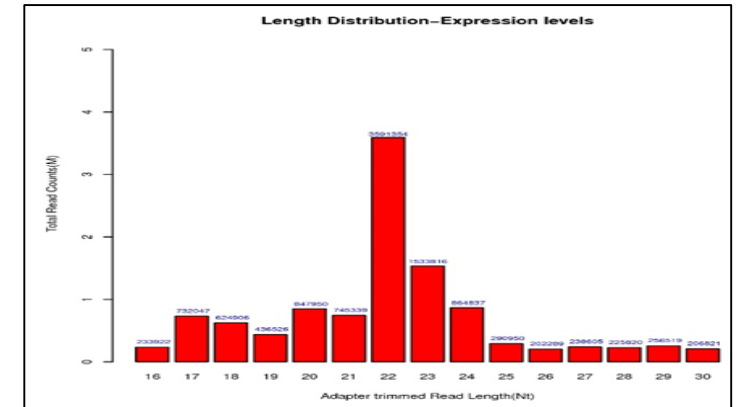
Control



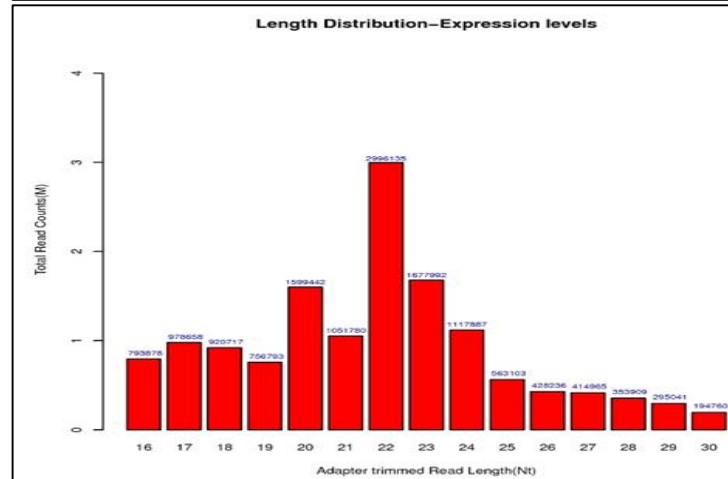
Donor 2



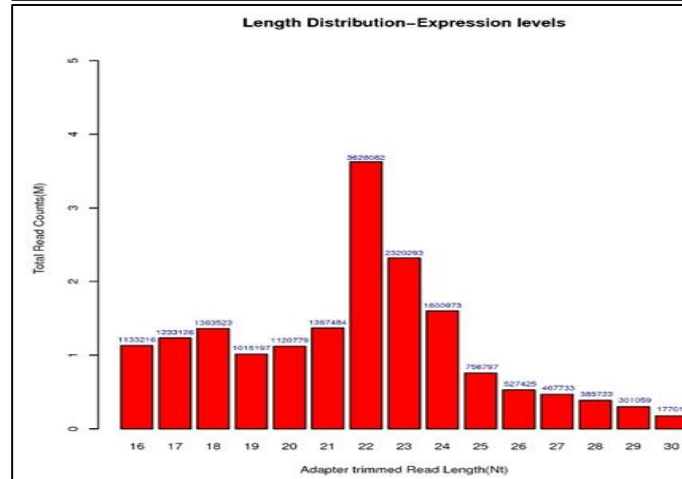
Donor 3



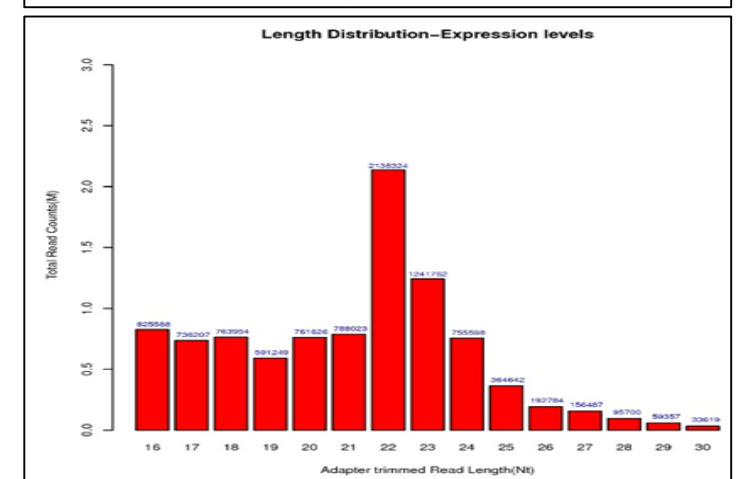
HIV-1



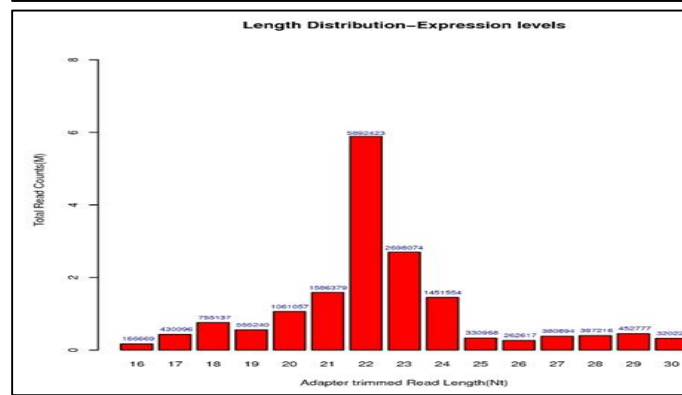
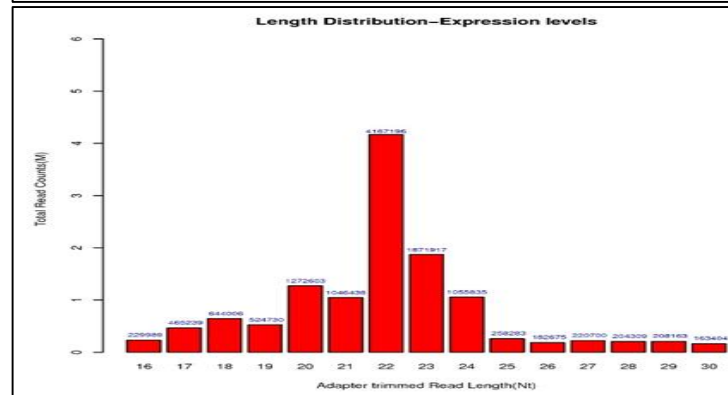
(B) PBMC



Length Distribution-Expression levels



HIV-2



Length Distribution-Expression levels

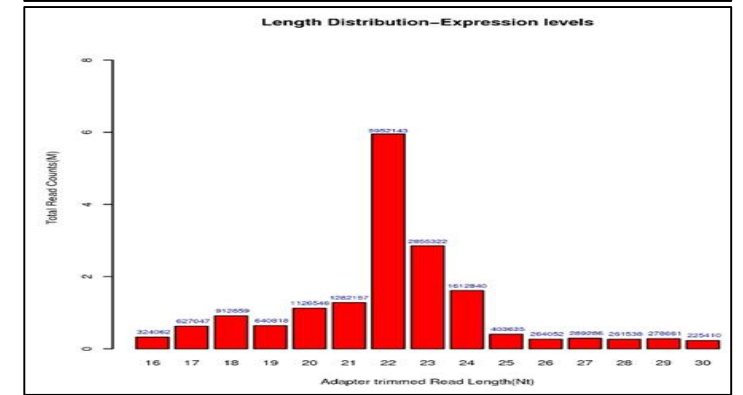


Figure S1: The length distribution of total small RNAs in all eighteen libraries.

The distribution of small RNAs in MDMs (A) and PBMCs (B) for each group. The vertical axis indicates the total read counts and the horizontal axis indicates the read lengths for the complete adapter-trimmed read sets.