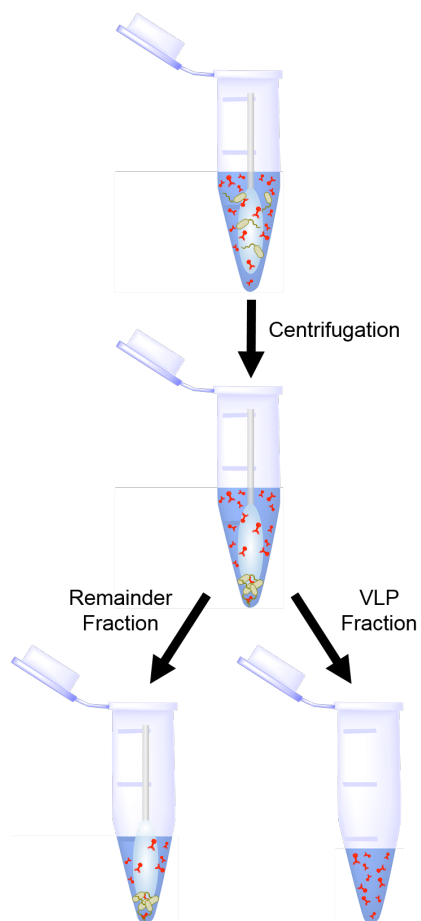
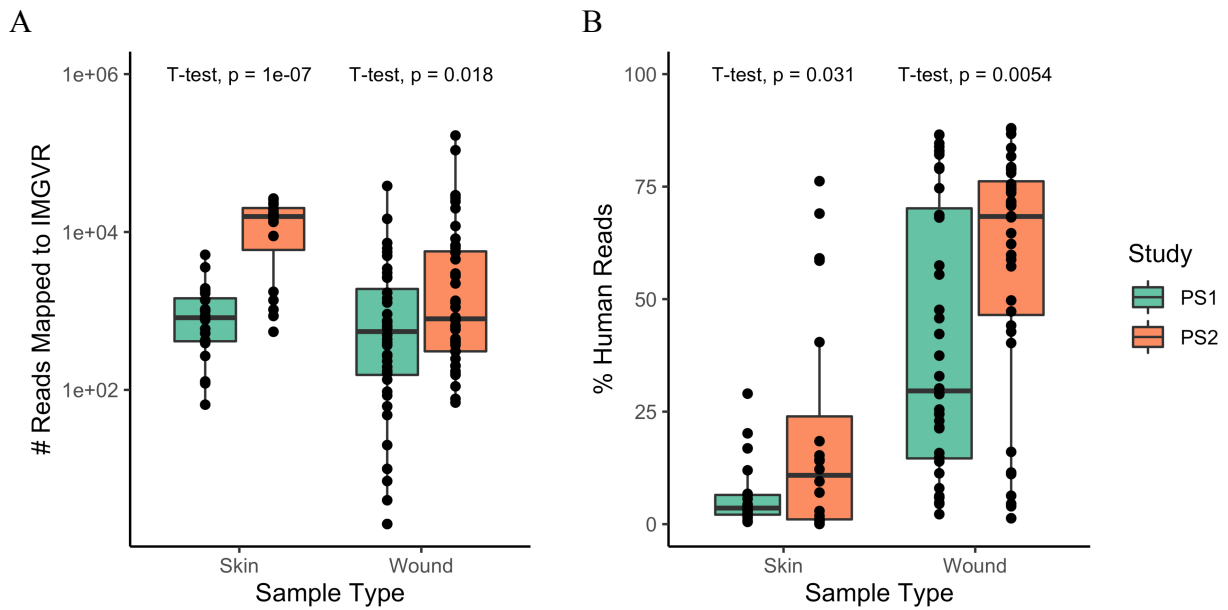


Supplementary Information

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Supplemental Figure S1. Graphical schematic of fractionation method.



Supplemental Figure S2. Additional comparisons of VLP-enriched DNA composition from clinical skin and wound swabs using kit-based extraction (PS1) and the method described here (PS2). VLP-enriched DNA was shotgun sequenced and mapped to the IMG/VR viral metagenome database (A) or the human genome (B). Means are compared by *t*-tests. Although PS2 samples contained more human DNA than PS1, the greater fraction of viral reads (Figure 6B) also translated to a greater absolute number of reads mapping to IMG/VR.