Supplemental Figure 1: Bar plots of the total number of identifiable phage genes expressed in relative periodontal health or significant periodontal disease.
Supplemental Figure 2: Number of transcriptome reads with identifiable phage homologues in subjects with periodontal health (white bar) or disease (black bar).
Supplemental Figure 3: Percentage (±standard deviation) of transcriptome reads from subjects with periodontal health (white bar) or disease (black bar) that were homologous to virome sequences from all subjects.