



FigS7. Fragment recruitment plots of contigs representing proviruss of a Bacteroidetes, Alphaproteobacteria, Deltaproteobacteria and a Gammaproteobacteria from selected metaviromic and metagenomic datasets (name shown at bottom right). Percentage identity (nucleotides) is shown on the Y-axis and the reads are shown in blue (metagenomes) and red (viromes)