



Extended Fig. 67. SQANTI3 analysis of SIRV reads in manatee samples. a) SQANTI3 categories for reads mapping to SIRVs in cDNA–PacBio and cDNA–ONT replicates. b) Number of SIRV transcripts with at least one Reference Match (RM) read in cDNA–PacBio and cDNA–ONT replicates