

Sequencing techniques	Host cell reads	IVV reads	Average length of viral RNAs in bp	Average coverage of viral reads
MinION (ONT)_VSV_cap_mix	6,302,618	559,408	1,004.27	50,335.6
MinION (ONT) Glia_dcDNA_mix	29,888	53,305	775.42	3,703.39
MinION (ONT) Fibroblast_dcDNA_mix	80,344	32,360	763.40	22,13.39
MinION (ONT) Glia_amplified_cDNA	3,114,992	54,5547	781.81	4,251.85
MinION (ONT) Fibroblast_ amplified_cDNA	5,872,662	494,827	709.90	3,067.90

Additional file 2: Table S1. Reads and read counts obtained by using different sequencing techniques