

Dataset	Replica set	ID	Replica	Original sample Type	Diagnostic CT	Volume extracted uL	Elution Volume uL	Library Prep Protocol	Estimated target genomes input	No. of pre-hyb PCR cycles	No. of post-hyb PCR cycles	Mean read depth following alignment
Noro	1	NOR_2048	1 2	Faeces	25	200	90	cDNA synthesis followed by standard SureSelectXT 200 ng protocol	1800	12 12	18 22	296 1429