

Dataset	Replica set	ID	Replica	Original sample Type	Extraction method	Volume extracted ul	Elution volume ul	Diagnostic Ct value	Library Prep Protocol	Mean read depth following alignment
HSV1	1	C4	1	Cerebrospinal fluid	QIAMP DNA Mini	200	100	30.19	standard SureSelectXT 200 ng protocol	857
			2			135		32.75		273
			3			100		30.39		299
	2	C5	1			400	100	27.47		673
			2			60		31.35		384