

	Sample 1	Sample 2	Sample 3	Sample 4
Reads Sequenced	6,227,749	6,202,253	6,233,646	5,067,711
Alignments at HpaII sites	4,536,673	4,602,001	4,614,571	4,465,175
Unique alignments at HpaII sites	4,160,629	4,221,245	4,225,504	4,066,062