

Table S16

Sample Name	HEK293T-CT	HEK293T-KD
Read count	90638165	83117053
Base count	13866969515	12314145819
Mean read length	152.99	148.15
Reads mapped by bowtie2	49198728	31046804
Reads remained unmapped	41439437	52070249
Base count for unmapped reads	5938568698	7395489494
Total A-U-B chimeric events in unmapped reads	4888	5472
Chimeric DNA events per billion bases sequenced	823.09	739.91