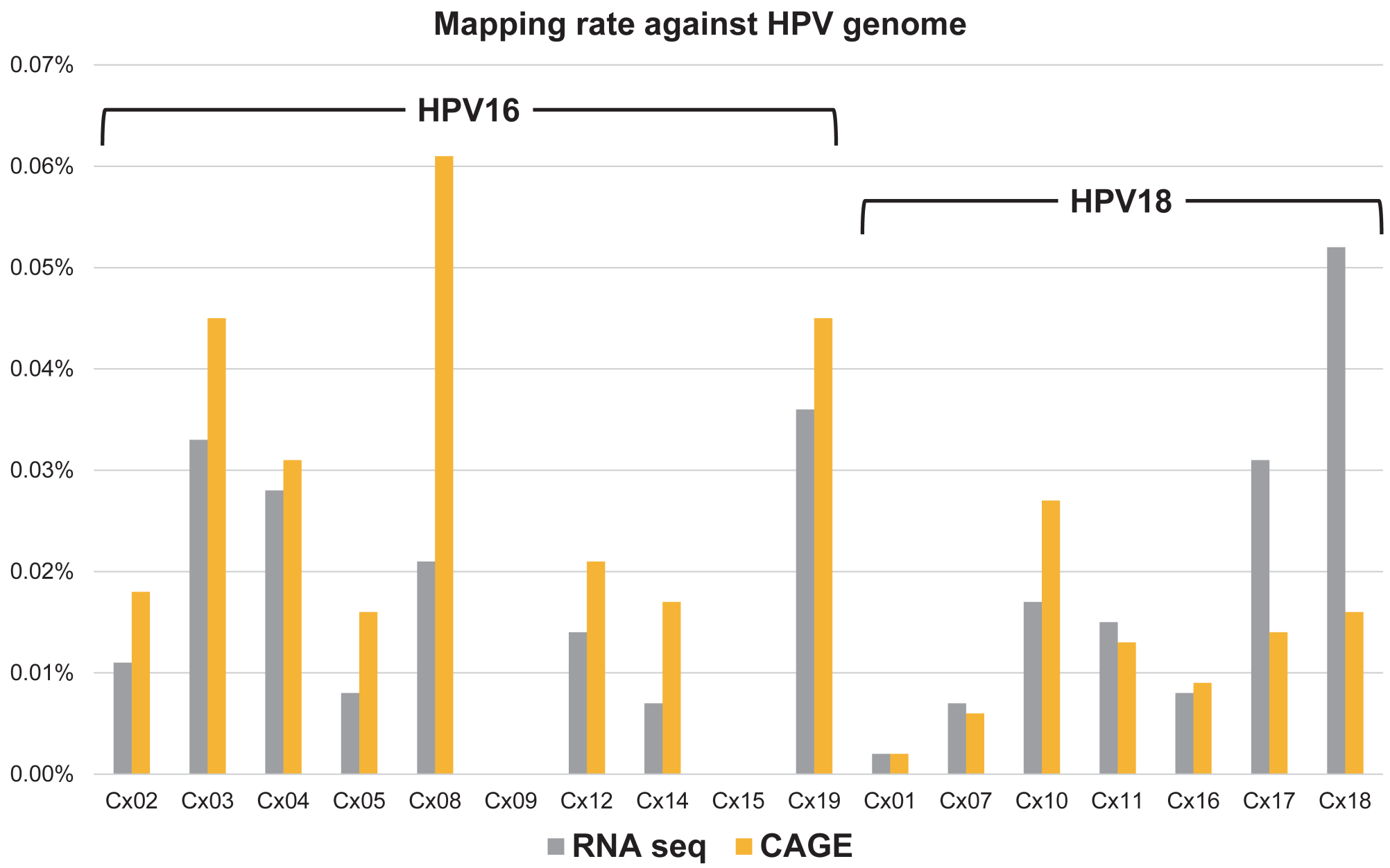
Fig. S1


Mapping rate against HPV genome
0.07%
0.06%
0.05%
0.04%
0.03%
0.02%
0.01%
0.00%
HPV16
HPV18
Cx02
Cx03
Cx04
Cx05
Cx08
Cx09
Cx12
Cx14
Cx15
Cx19
Cx01
Cx07
Cx10
Cx11
Cx16
Cx17
Cx18
RNA seq
CAGE