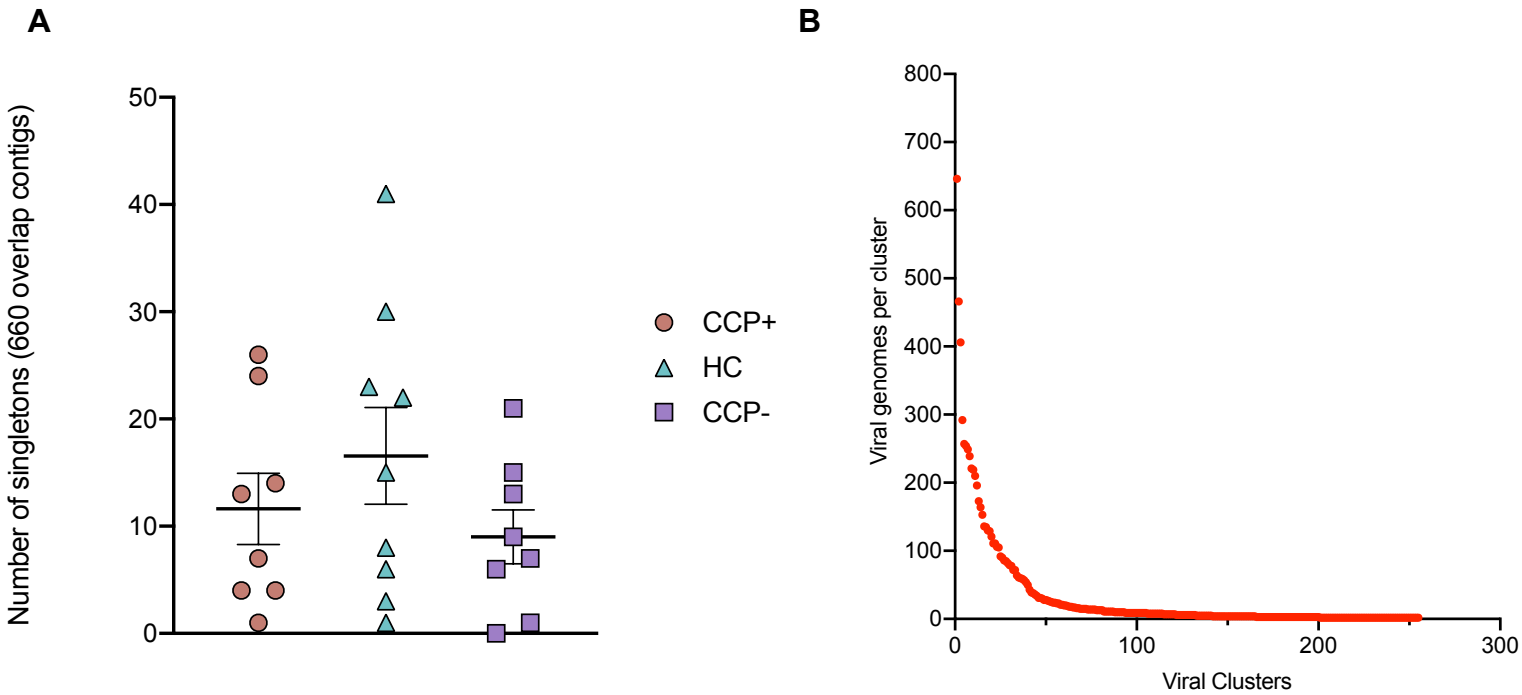
Figure S4


A
Number of singletons (660 overlap contigs)
50
40
30
20
10
0
CCP+
HC
CCP-
B
Viral genomes per cluster
800
700
600
500
400
300
200
100
0
0
100
200
300
Viral Clusters