

Table S5 Viral read counts in the mock-infected founders. Each strain's total comes from four mice. The counts are low, several orders of magnitude lower than infected samples. The fact we still get non-zero reads may be an artifact of mapping and select strains may have partial sequence similarity to viruses.

Strain	MA15	PR8
129S1	5084	11394
AJ	9549	2659
B6	521	1656
CAST	10490	3060
NOD	608	857
NZO	629	636
PWK	957	634
WSB	144	7491