

Total admissions, (2012-2019), **n=2001**



Consented & stool obtained, **n=1969,**
98.4%

Not consented, **n=32**

RVA positives, **n=415, 21.1%**

Not tested/RVA negative, **n=1554**

Successfully GP typed, **n=346, 83.4%**
G1P[8], n=172

Partially typed/Failed sequencing, **n=69**

Full genome sequencing, **n=393, 94.6%**

Preselected from total positives

Assembled and genotyped, **n=336, 85.5%**

Failed genome assembly, **n=57**

G1P[8] genomes, **n=171, 50.8%**

Non-G1P[8] genomes, **n=165**