

58,776 virus-like contigs
from the 4 Dutch cohort

41,284 genomes
from the databases

Dereplication

30,461 vOTUs

Read mapping

15,196 detected vOTUs

Caudoviricetes genomes with terminal repeats

1,899 selected vOTUs

Detected in > 5% samples of a Dutch cohort

54 selected vOTUs

