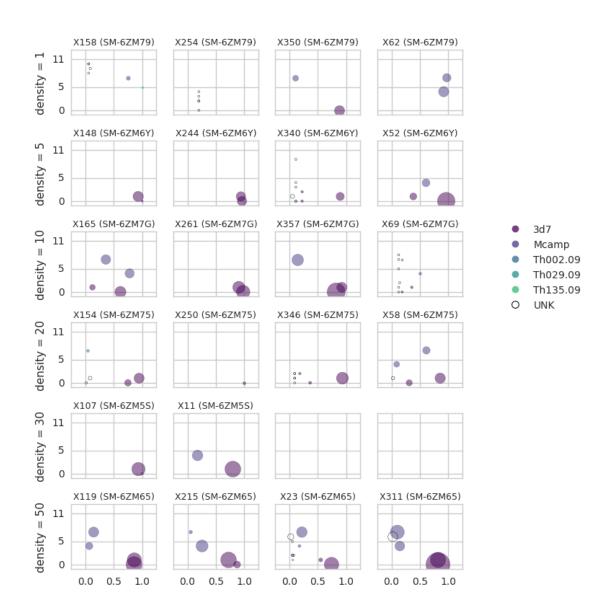
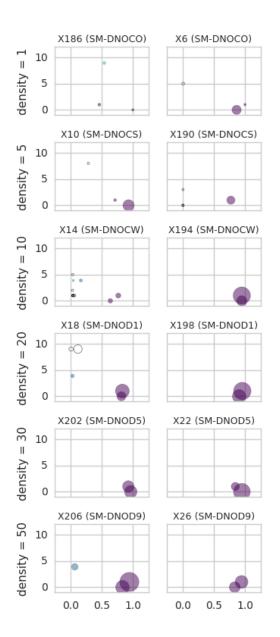
rtss amplicon seq

20171201

50:50 3d7;Mcamp





3d7

Dd2

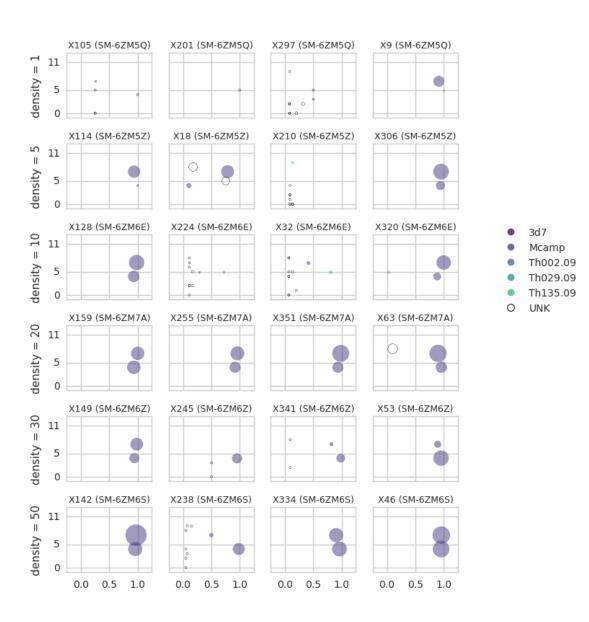
O UNK

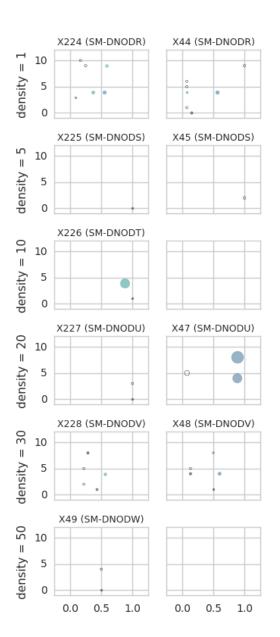
Mcamp

Th002.09

Th135.09

100 Mcamp





3d7

Dd2

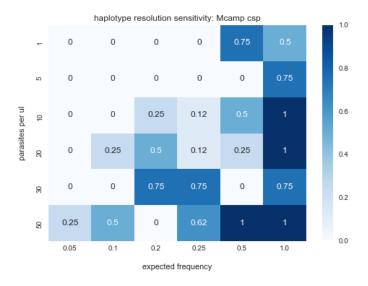
O UNK

Mcamp

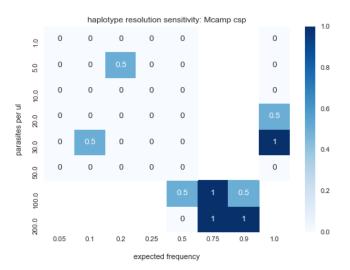
Th002.09

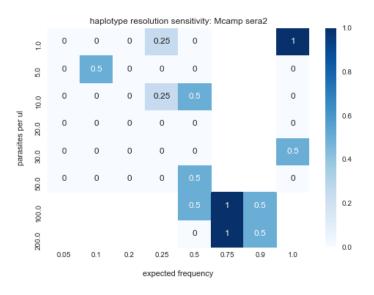
Th135.09

haplotype resolution

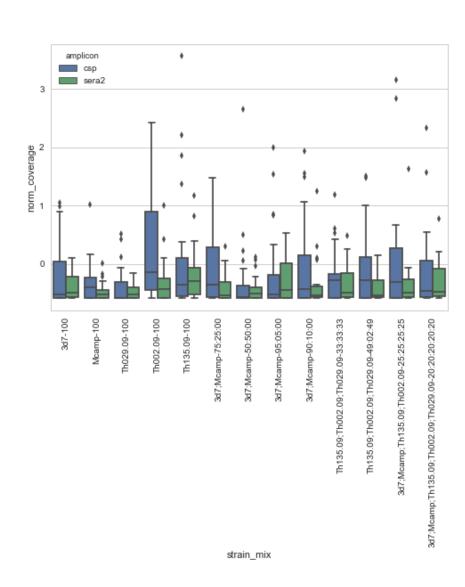


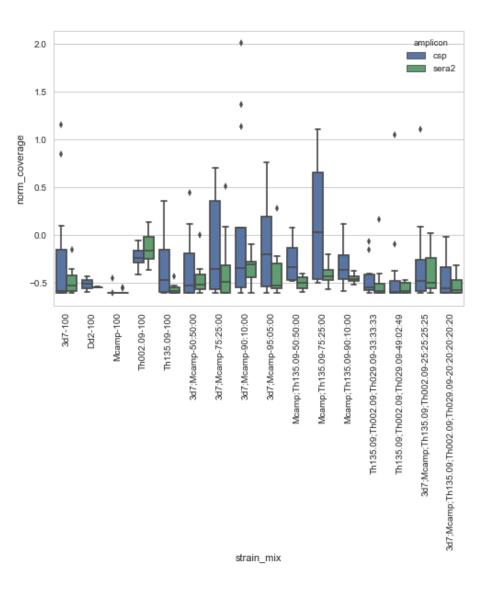






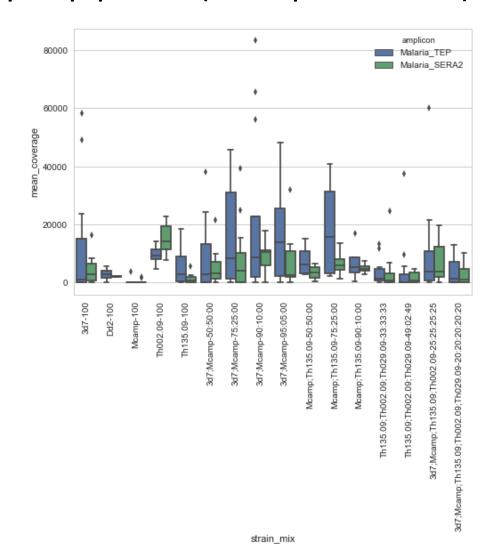
coverage by strain-mix





coverage by strain-mix

pre-pipeline (from pcr metrics)



post-pipeline

