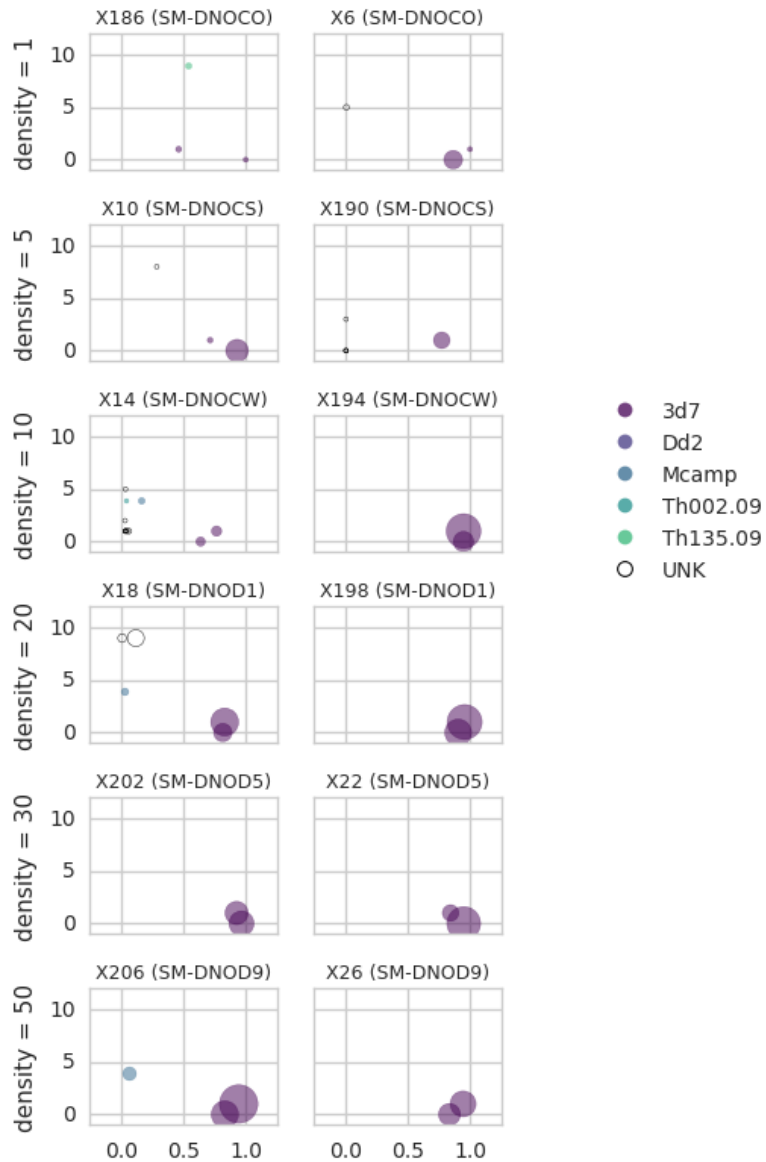
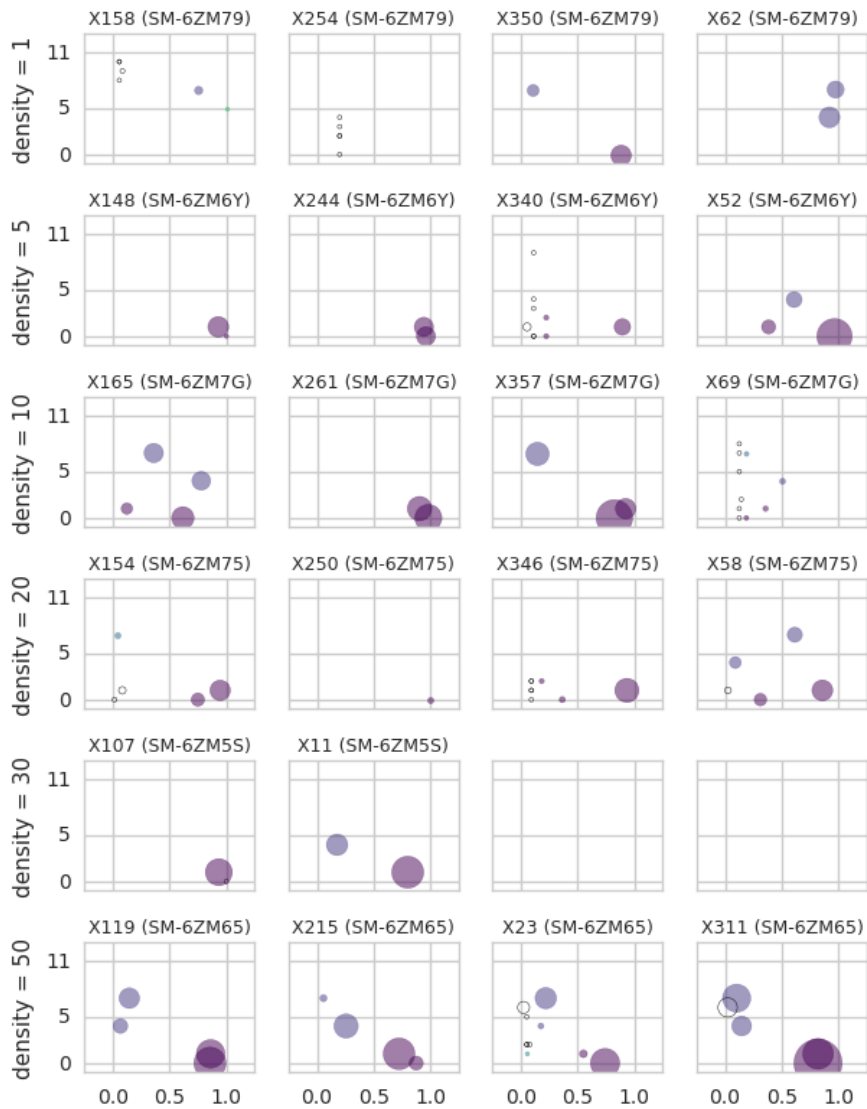


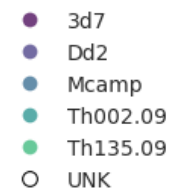
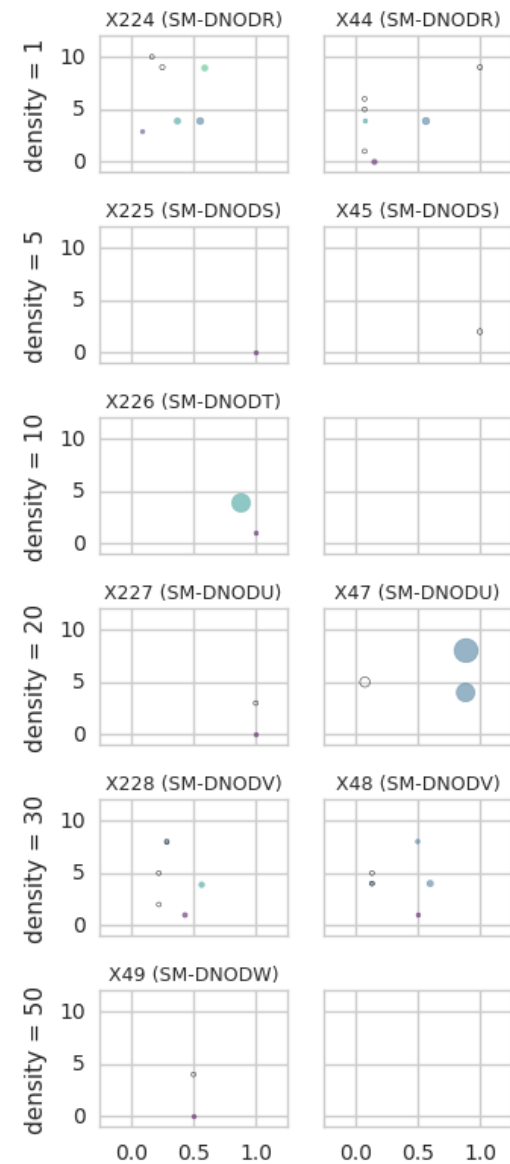
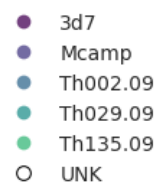
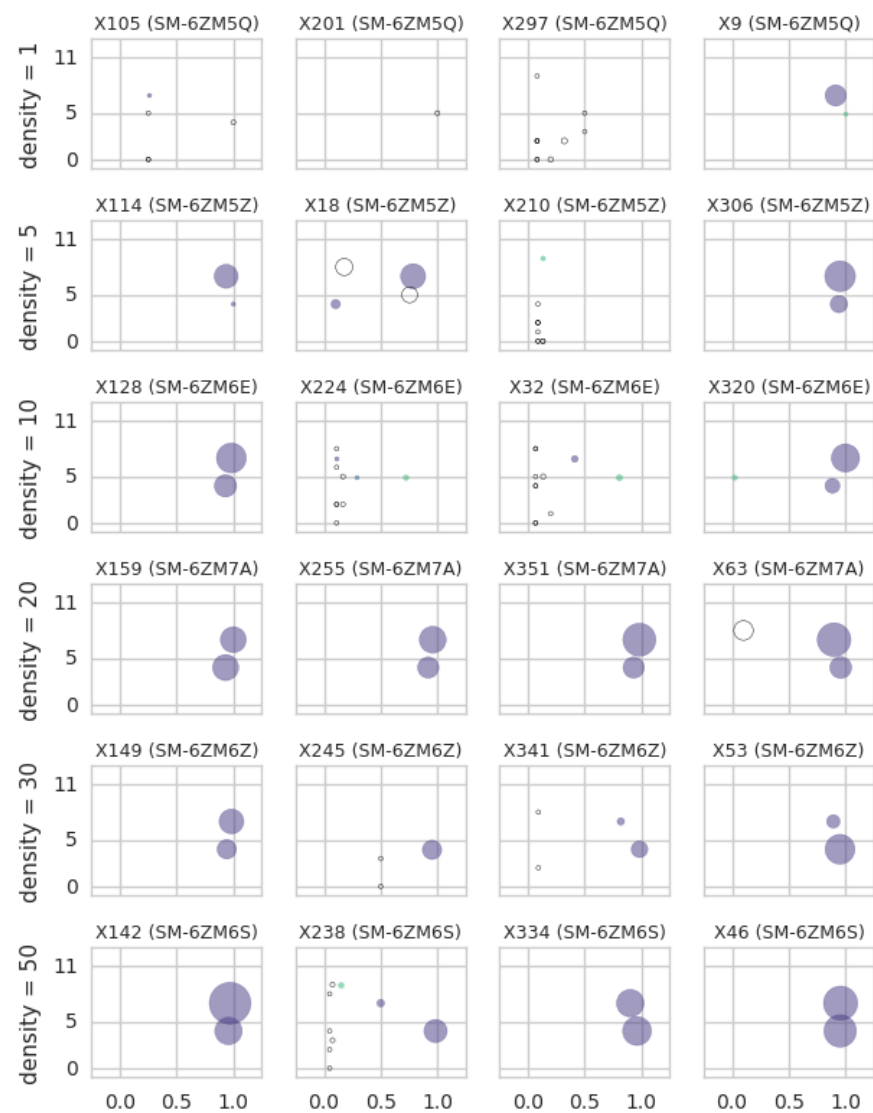
rtss amplicon seq

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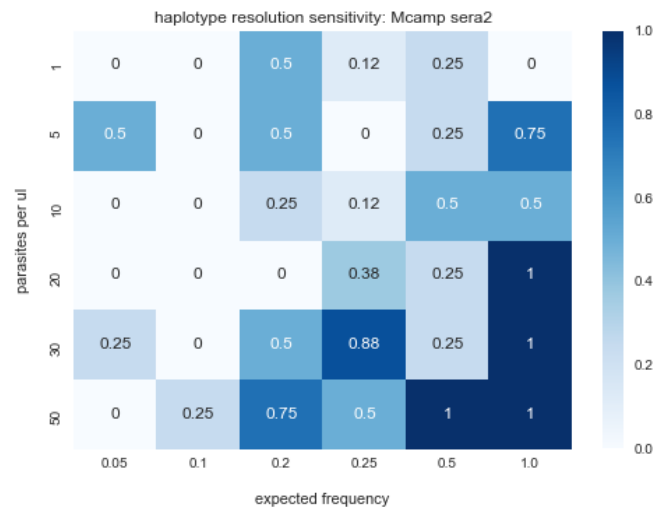
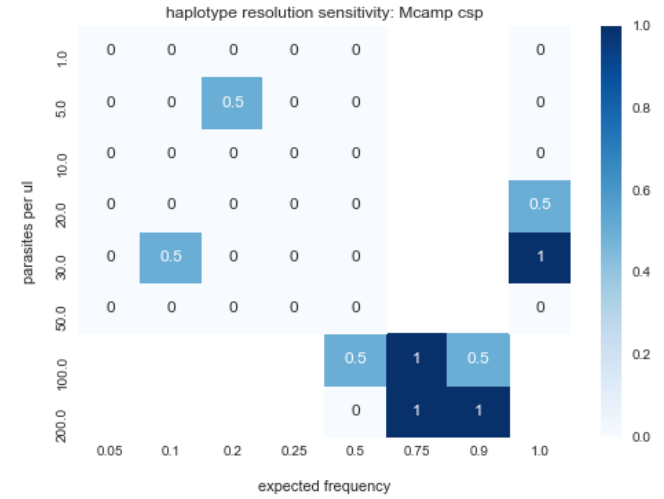
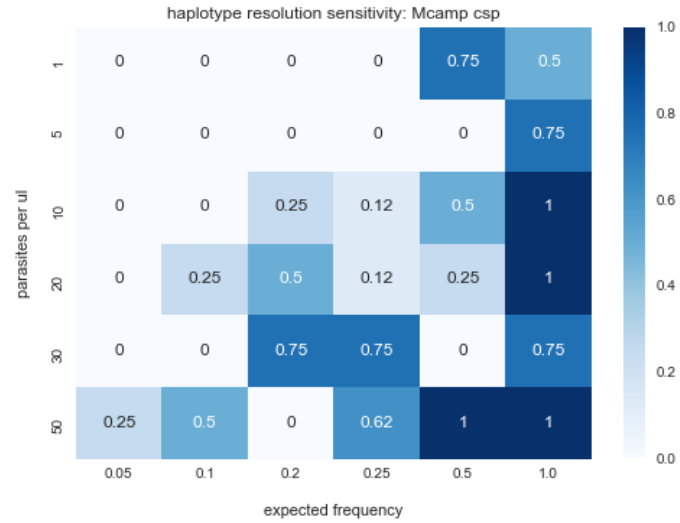
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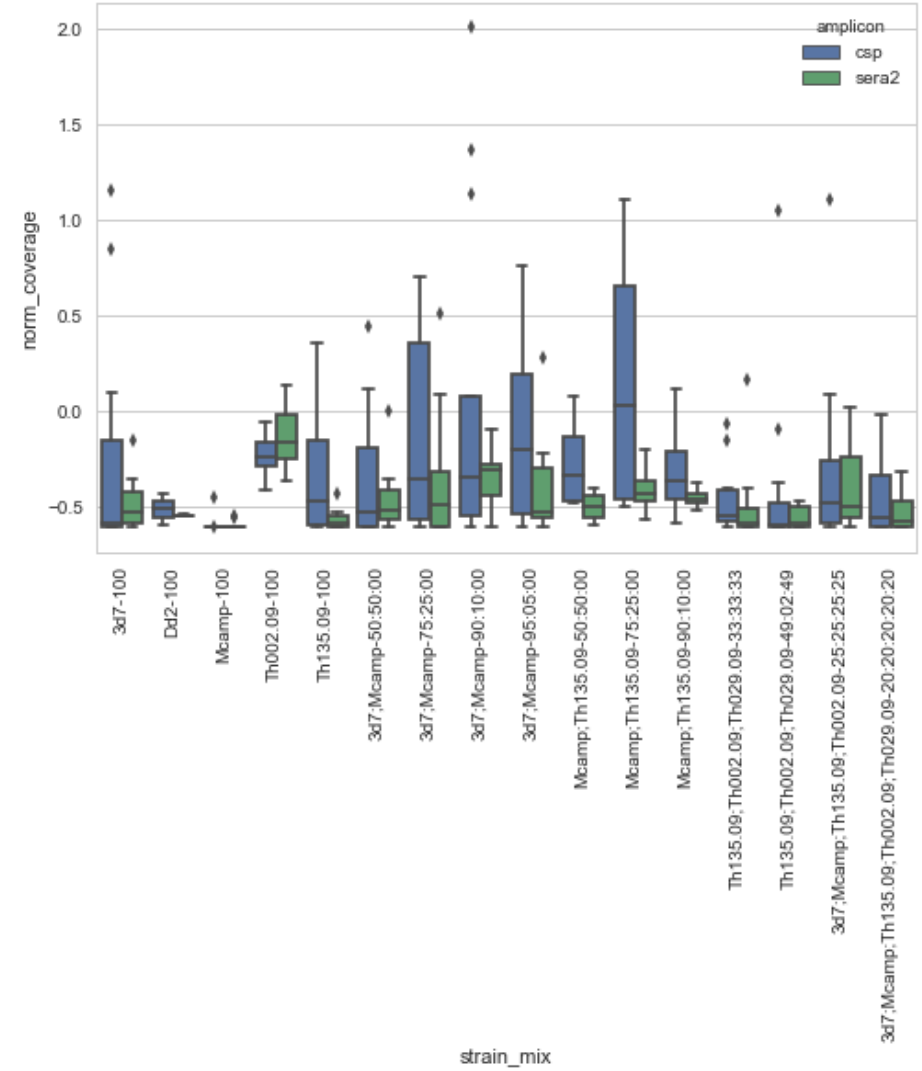
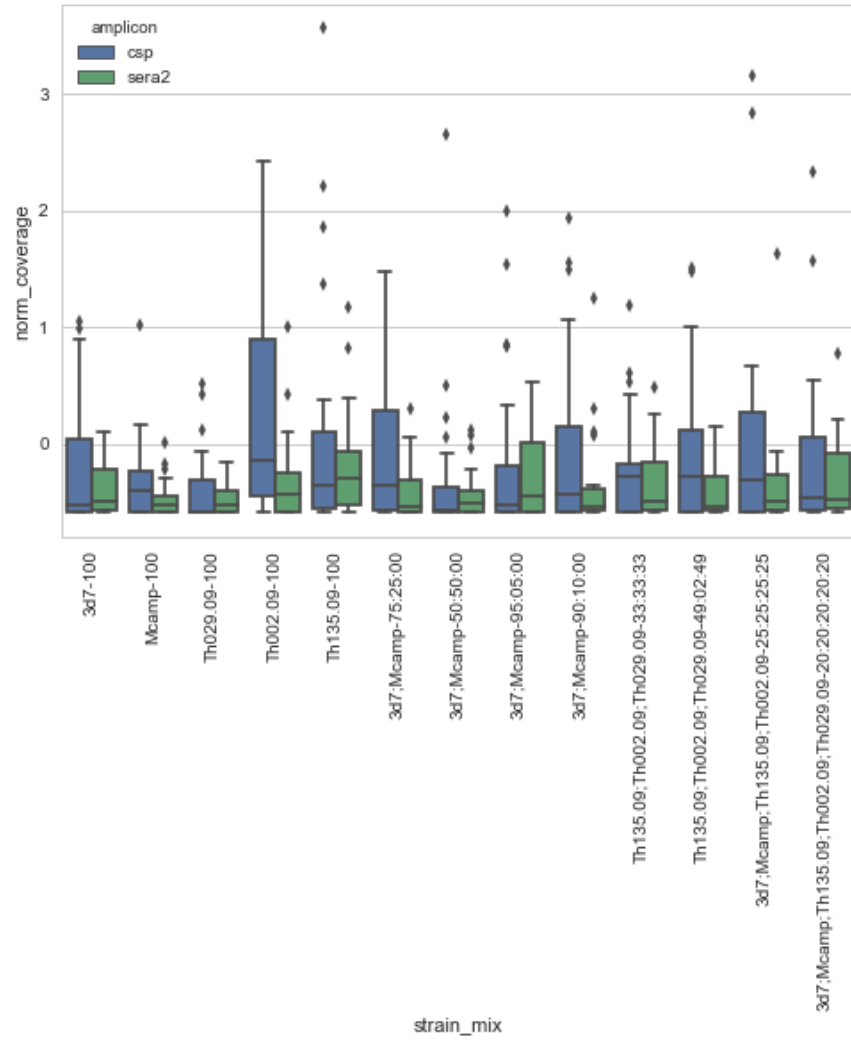
100 Mcamp



haplotype resolution

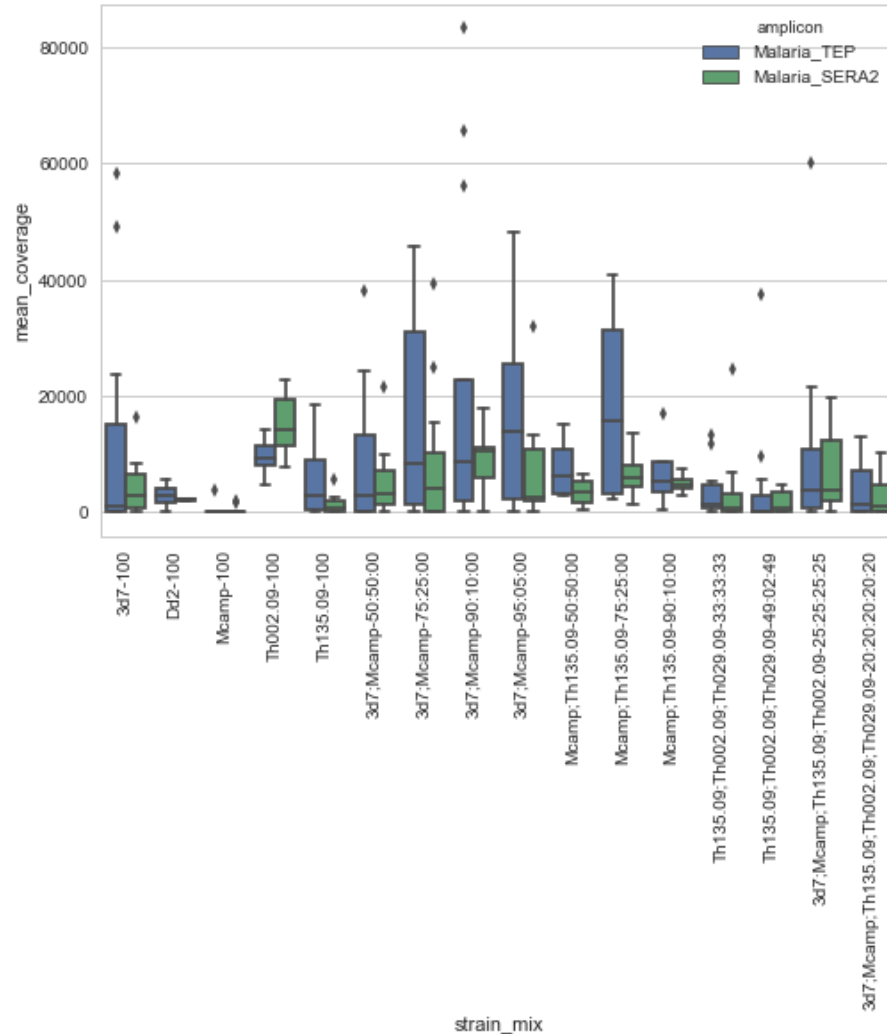


coverage by strain-mix



coverage by strain-mix

pre-pipeline (from pcr metrics)



post-pipeline

