

Validation of consensus calling

Consensus Sequences

simulate2 – Thres: 90 %
simulate2 – Thres: 50 %
simulate2 – Thres: 25 %
simulate2 – Thres: 0 %
simulate1 – Thres: 90 %
simulate1 – Thres: 50 %
simulate1 – Thres: 25 %
simulate1 – Thres: 0 %
27a – Thres: 90 %
27a – Thres: 50 %
27a – Thres: 25 %
27a – Thres: 0 %
26a – Thres: 90 %
26a – Thres: 50 %
26a – Thres: 25 %
26a – Thres: 0 %

0	3370	3370
0	68	68
0	46	46
0	46	46
0	3213	3213
0	134	134
0	102	102
0	102	102
0	494	494
0	465	465
0	465	465
0	465	465
0	496	496
0	452	452
0	452	452
0	452	452

Geneious vs iVar

iVar vs Reference

Geneious vs Reference

Nucleotide Differences

