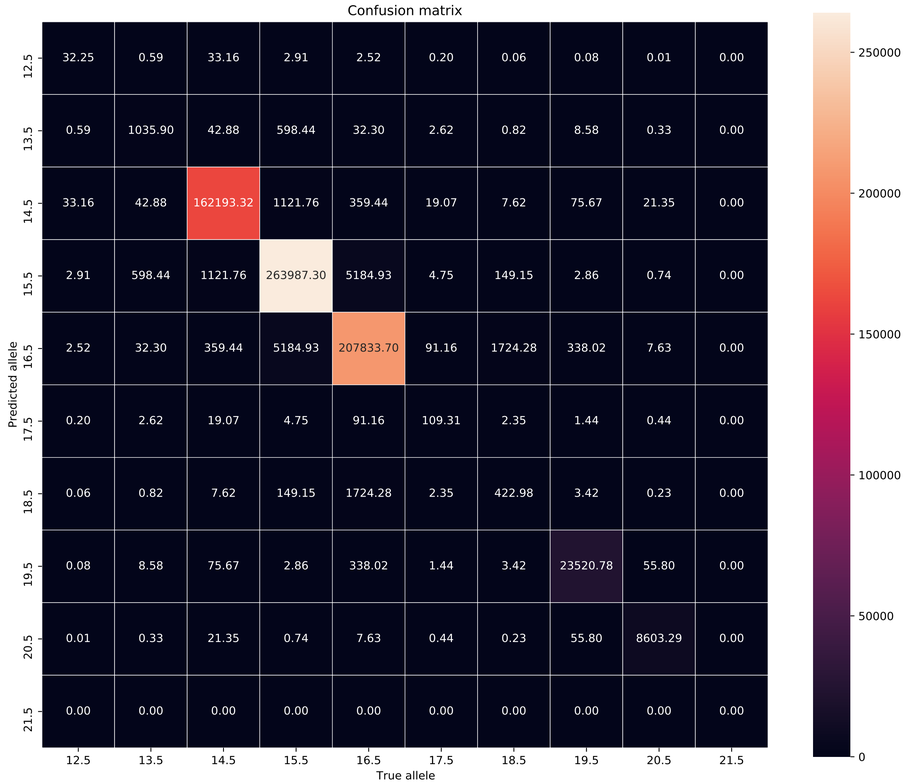
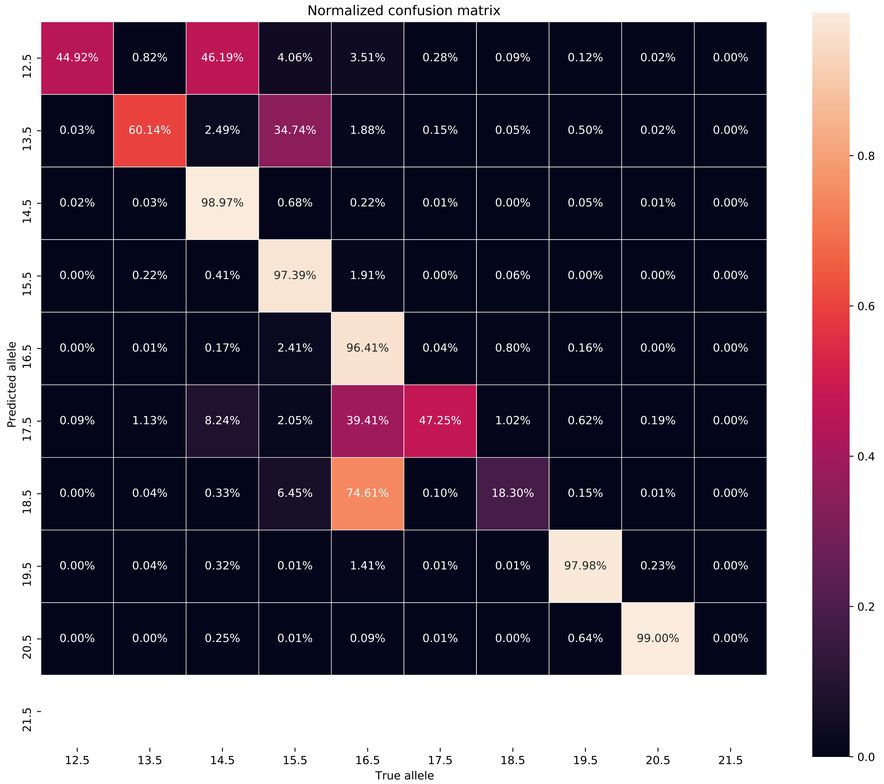

Confusion matrix
Predicted allele
True allele
12.5 13.5 14.5 15.5 16.5 17.5 18.5 19.5 20.5 21.5
32.25 0.59 33.16 2.91 2.52 0.20 0.06 0.08 0.01 0.00
0.59 1035.90 42.88 598.44 32.30 2.62 0.82 8.58 0.33 0.00
33.16 42.88 162193.32 1121.76 359.44 19.07 7.62 75.67 21.35 0.00
2.91 598.44 1121.76 263987.30 5184.93 4.75 149.15 2.86 0.74 0.00
2.52 32.30 359.44 5184.93 207833.70 91.16 1724.28 338.02 7.63 0.00
0.20 2.62 19.07 4.75 91.16 109.31 2.35 1.44 0.44 0.00
0.06 0.82 7.62 149.15 1724.28 2.35 422.98 3.42 0.23 0.00
0.08 8.58 75.67 2.86 338.02 1.44 3.42 23520.78 55.80 0.00
0.01 0.33 21.35 0.74 7.63 0.44 0.23 55.80 8603.29 0.00
0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.00
250000
200000
150000
100000
50000
0


Normalized confusion matrix
Predicted allele
True allele
12.5 13.5 14.5 15.5 16.5 17.5 18.5 19.5 20.5 21.5
44.92% 0.82% 46.19% 4.06% 3.51% 0.28% 0.09% 0.12% 0.02% 0.00%
0.03% 60.14% 2.49% 34.74% 1.88% 0.15% 0.05% 0.50% 0.02% 0.00%
0.02% 0.03% 98.97% 0.68% 0.22% 0.01% 0.00% 0.05% 0.01% 0.00%
0.00% 0.22% 0.41% 97.39% 1.91% 0.00% 0.06% 0.00% 0.00% 0.00%
0.00% 0.01% 0.17% 2.41% 96.41% 0.04% 0.80% 0.16% 0.00% 0.00%
0.09% 1.13% 8.24% 2.05% 39.41% 47.25% 1.02% 0.62% 0.19% 0.00%
0.00% 0.04% 0.33% 6.45% 74.61% 0.10% 18.30% 0.15% 0.01% 0.00%
0.00% 0.04% 0.32% 0.01% 1.41% 0.01% 0.01% 97.98% 0.23% 0.00%
0.00% 0.00% 0.25% 0.01% 0.09% 0.01% 0.00% 0.64% 99.00% 0.00%
0.8
0.6
0.4
0.2
0.0