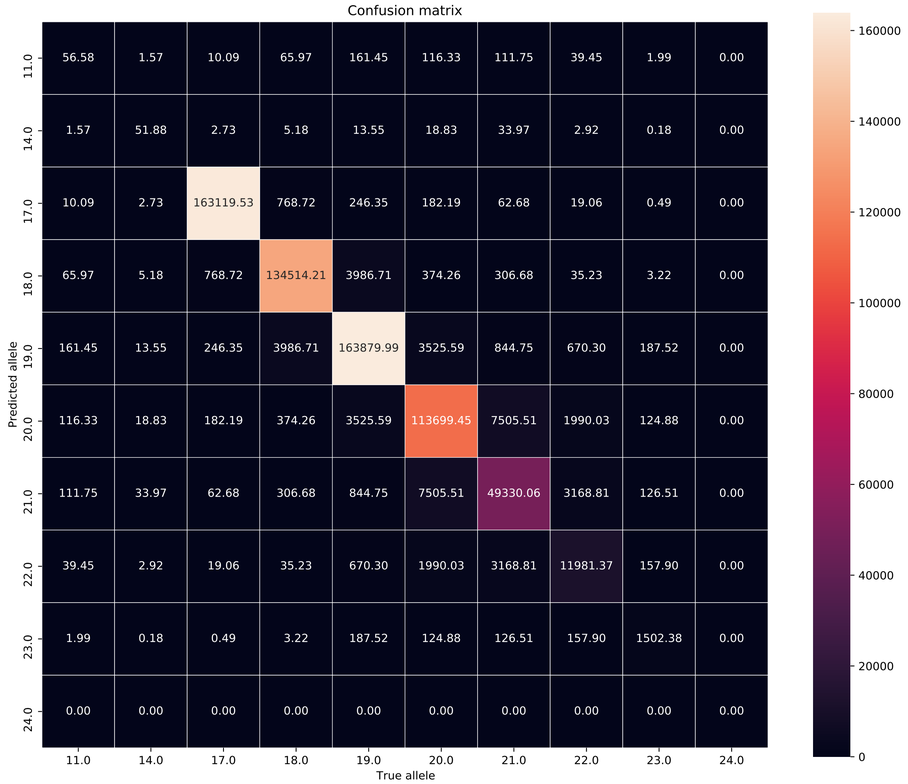
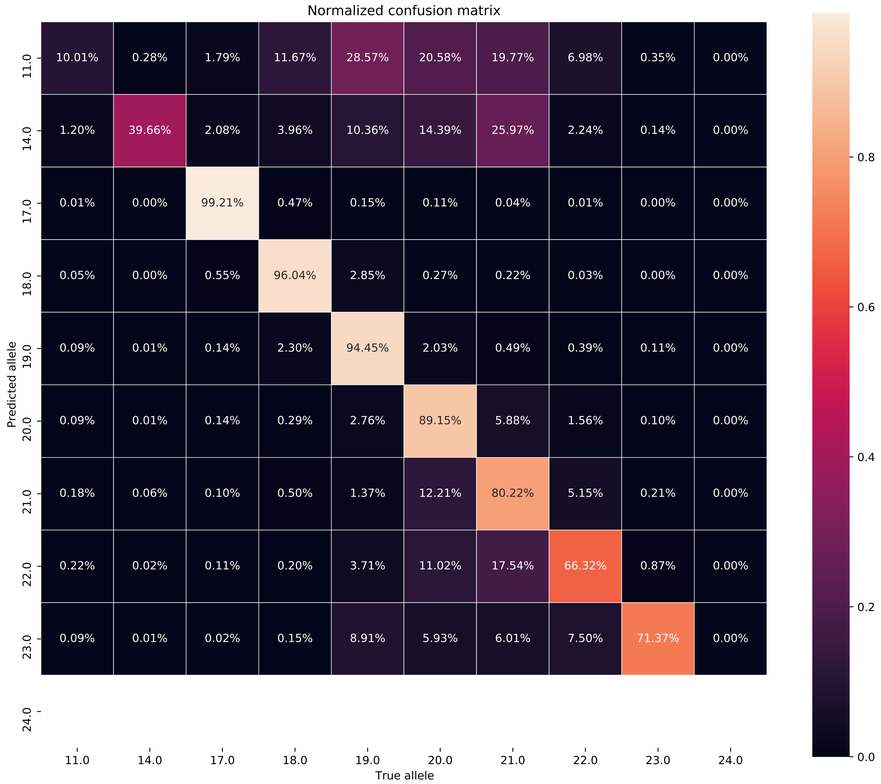

Confusion matrix
Predicted allele
True allele
11.0 14.0 17.0 18.0 19.0 20.0 21.0 22.0 23.0 24.0
11.0 56.58 1.57 10.09 65.97 161.45 116.33 111.75 39.45 1.99 0.00
14.0 1.57 51.88 2.73 5.18 13.55 18.83 33.97 2.92 0.18 0.00
17.0 10.09 2.73 163119.53 768.72 246.35 182.19 62.68 19.06 0.49 0.00
18.0 65.97 5.18 768.72 134514.21 3986.71 374.26 306.68 35.23 3.22 0.00
19.0 161.45 13.55 246.35 3986.71 163879.99 3525.59 844.75 670.30 187.52 0.00
20.0 116.33 18.83 182.19 374.26 3525.59 113699.45 7505.51 1990.03 124.88 0.00
21.0 111.75 33.97 62.68 306.68 844.75 7505.51 49330.06 3168.81 126.51 0.00
22.0 39.45 2.92 19.06 35.23 670.30 1990.03 3168.81 11981.37 157.90 0.00
23.0 1.99 0.18 0.49 3.22 187.52 124.88 126.51 157.90 1502.38 0.00
24.0 0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.00
160000 140000 120000 100000 80000 60000 40000 20000 0


Normalized confusion matrix
Predicted allele
True allele
11.0 14.0 17.0 18.0 19.0 20.0 21.0 22.0 23.0 24.0
11.0 10.01% 0.28% 1.79% 11.67% 28.57% 20.58% 19.77% 6.98% 0.35% 0.00%
14.0 1.20% 39.66% 2.08% 3.96% 10.36% 14.39% 25.97% 2.24% 0.14% 0.00%
17.0 0.01% 0.00% 99.21% 0.47% 0.15% 0.11% 0.04% 0.01% 0.00% 0.00%
18.0 0.05% 0.00% 0.55% 96.04% 2.85% 0.27% 0.22% 0.03% 0.00% 0.00%
19.0 0.09% 0.01% 0.14% 2.30% 94.45% 2.03% 0.49% 0.39% 0.11% 0.00%
20.0 0.09% 0.01% 0.14% 0.29% 2.76% 89.15% 5.88% 1.56% 0.10% 0.00%
21.0 0.18% 0.06% 0.10% 0.50% 1.37% 12.21% 80.22% 5.15% 0.21% 0.00%
22.0 0.22% 0.02% 0.11% 0.20% 3.71% 11.02% 17.54% 66.32% 0.87% 0.00%
23.0 0.09% 0.01% 0.02% 0.15% 8.91% 5.93% 6.01% 7.50% 71.37% 0.00%
24.0
0.8 0.6 0.4 0.2 0.0