Confusion matrix

Predicted allele	10.0	11.0	12.0	13.0	14.0	15.0	16.0	17.0	18.0
10.0	233.94	67.32	300.04	19.52	7.15	2.51	0.22	0.06	0.00
11.0	67.32	280489.75	911.94	321.01	66.36	628.72	3.03	35.27	28.86
12.0	300.04	911.94	123766.92	3835.45	1771.47	176.05	20.44	4.64	0.43
13.0	19.52	321.01	3835.45	30985.26	5954.35	1721.15	93.37	0.32	0.02
14.0	7.15	66.36	1771.47	5954.35	106929.38	3329.65	258.50	44.98	5.44
15.0	2.51	628.72	176.05	1721.15	3329.65	87363.87	4700.30	625.61	45.98
16.0	0.22	3.03	20.44	93.37	258.50	4700.30	6844.99	80.27	9.78
17.0	0.06	35.27	4.64	0.32	44.98	625.61	80.27	799.94	40.11
18.0	0.00	28.86	0.43	0.02	5.44	45.98	9.78	40.11	23.66

True allele

Normalized confusion matrix

Predicted allele	10.0	11.0	12.0	13.0	14.0	15.0	16.0	17.0	18.0
10.0	37.09%	10.67%	47.57%	3.09%	1.13%	0.40%	0.04%	0.01%	0.00%
11.0	0.02%	99.27%	0.32%	0.11%	0.02%	0.22%	0.00%	0.01%	0.01%
12.0	0.23%	0.70%	94.63%	2.93%	1.35%	0.13%	0.02%	0.00%	0.00%
13.0	0.05%	0.75%	8.93%	72.18%	13.87%	4.01%	0.22%	0.00%	0.00%
14.0	0.01%	0.06%	1.50%	5.03%	90.34%	2.81%	0.22%	0.04%	0.00%
15.0	0.00%	0.64%	0.18%	1.75%	3.38%	88.61%	4.77%	0.63%	0.05%
16.0	0.00%	0.03%	0.17%	0.78%	2.15%	39.13%	56.99%	0.67%	0.08%
17.0	0.00%	2.16%	0.28%	0.02%	2.76%	38.35%	4.92%	49.04%	2.46%
18.0	0.00%	18.71%	0.28%	0.02%	3.53%	29.80%	6.34%	26.00%	15.33%

True allele