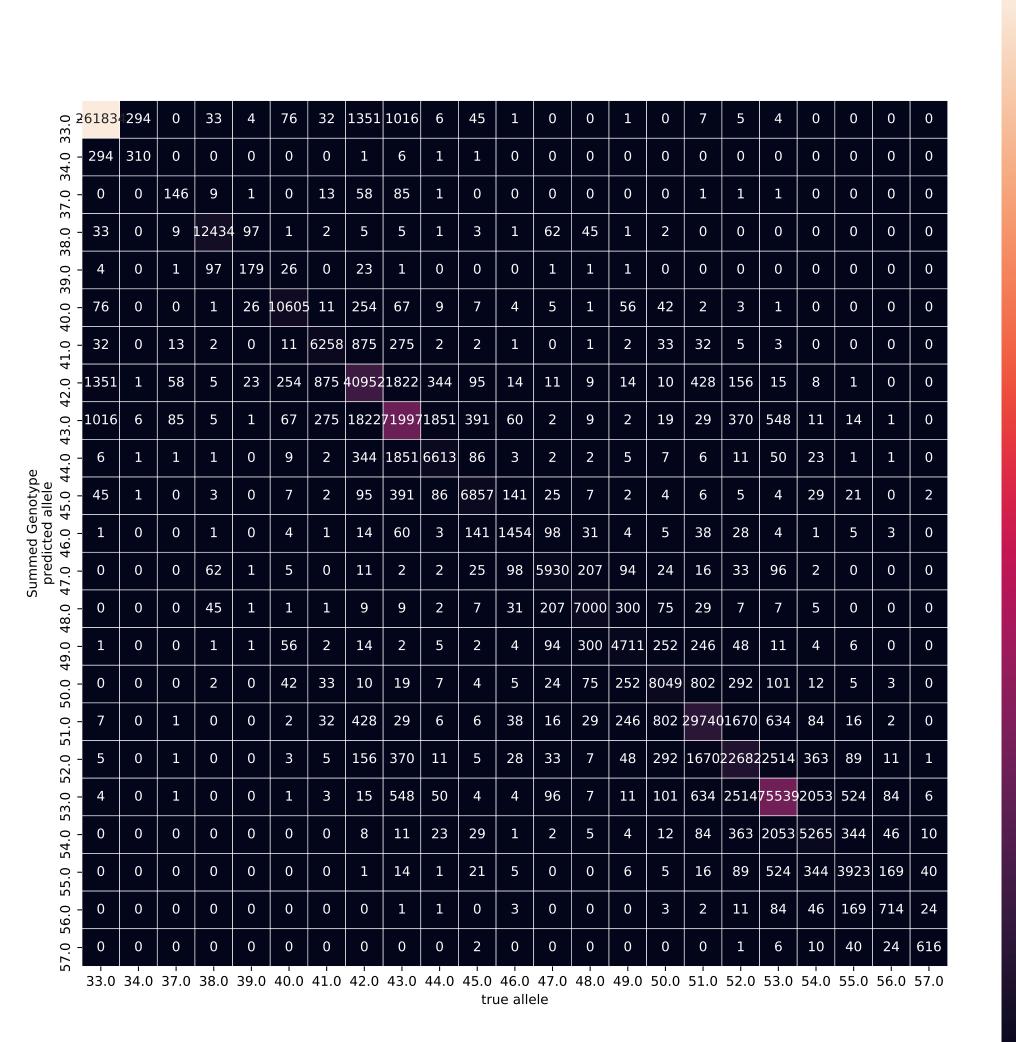
					N	ormalize	ed confus	sion mat	ix				
16.5	98.93%	0.11%	0.00%	0.00%	0.01%	0.00%	0.03%	0.01%	0.51%	0.38%	0.00%	0.02%	0.00%
17.5	30.41%	68.43%	0.00%	0.00%	0.08%	0.00%	0.00%	0.00%	0.06%	0.78%	0.00%	0.22%	0.03%
18.5	3.88%	0.00%	95.42%	0.00%	0.65%	0.01%	0.00%	0.00%	0.02%	0.00%	0.00%	0.01%	0.00%
20.5	0.04%	0.00%	0.00%	45.48%	3.11%	0.31%	0.00%	3.99%	18.87%	28.03%	0.15%	0.00%	0.00%
21.5	0.13%	0.00%	0.00%	0.08%	99.02%	0.69%	0.00%	0.02%	0.04%	0.01%	0.01%	0.00%	0.00%
ele 22.5	0.53%	0.00%	0.00%	0.32%	28.98%	53.18%	9.11%	0.03%	7.36%	0.22%	0.17%	0.09%	0.00%
predicted allele 23.5 22	0.32%	0.00%	0.00%	0.00%	0.00%	0.25%	96.28%	0.00%	2.37%	0.60%	0.10%	0.05%	0.03%
pre 24.5	0.21%	0.00%	0.00%	0.17%	0.03%	0.00%	0.01%	83.90%	11.83%	3.80%	0.04%	0.02%	0.00%
25.5	1.49%	0.00%	0.00%	0.13%	0.01%	0.05%	0.58%	1.94%	90.87%	3.91%	0.78%	0.22%	0.03%
26.5	0.67%	0.01%	0.00%	0.12%	0.00%	0.00%	0.09%	0.38%	2.38%	93.36%	2.38%	0.53%	0.08%
27.5	0.03%	0.00%	0.00%	0.01%	0.01%	0.01%	0.12%	0.03%	4.07%	20.46%	74.45%	0.80%	0.02%
28.5	0.34%	0.02%	0.00%	0.00%	0.01%	0.00%	0.08%	0.02%	1.52%	6.02%	1.05%	89.02%	1.92%
29.5	0.03%	0.01%	0.00%	0.00%	0.00%	0.00%	0.33%	0.00%	1.24%	5.89%	0.15%	12.31%	80.04%
	16.5	17.5	18.5	20.5	21.5	22.5	23.5 true allele	24.5	25.5	26.5	27.5	28.5	29.5

- 0.8

- 0.8

- 0.2



- 250000

- 200000

- 150000

- 100000

- 50000

