Confusion matrix													Normalized confusion matrix													
34.0	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	- 160000		34.0											
35.0	0.00	170596.63	27.14	0.02	0.00	79.37	2.82	24.24	5.51	1.78	25.47	0.11			92.0 32.0 32.0	99.90%	0.02%	0.00%	0.00%	0.05%	0.00%	0.01%	0.00%	0.00%	0.01%	0.00%
36.0	0.00	27.14	223.37	0.00	0.00	0.00	0.00	0.01	0.00	0.00	0.00	0.00	- 140000		0.00%	10.83%	89.16%	0.00%	0.00%	0.00%	0.00%	0.00%	0.00%	0.00%	0.00%	0.00%
37.0	0.00	0.02	0.00	4722.69	0.00	2.44	0.00	402.82	6.26	1.25	7.53	1.23	- 120000		92.0	0.00%	0.00%	91.81%	0.00%	0.05%	0.00%	7.83%	0.12%	0.02%	0.15%	0.02%
38.0	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.24	0.04	0.00	0.00	- 100000	00	0.00%	0.00%	0.00%	0.00%	1.00%	0.00%	0.00%	2.41%	83.39%	12.86%	0.34%	0.00%
ed allele 39.0 '	0.00	79.37	0.00	2.44	0.00	166245.80	9.78	333.08	33.86	8.96	26.81	0.36	100000		39.0 39.0 	0.05%	0.00%	0.00%	0.00%	99.70%	0.01%	0.20%	0.02%	0.01%	0.02%	0.00%
Predicte 40.0		2.82	0.00	0.00	0.00	9.78	92.41	41.42	92.94	0.17	4.50	0.32	- 80000		Predicte 40.0	1.15%	0.00%	0.00%	0.00%	4.00%	37.82%	16.95%	38.03%	0.07%	1.84%	0.13%
41.0	0.00	24.24	0.01	402.82	0.01	333.08	41.42	95839.23	861.43	94.16	121.80	58.26	- 60000		0.00%	0.02%	0.00%	0.41%	0.00%	0.34%	0.04%	98.02%	0.88%	0.10%	0.12%	0.06%
42.0	0.00	5.51	0.00	6.26	0.24	33.86	92.94	861.43	117509.09	3241.77	507.65	12.74			75.0	0.00%	0.00%	0.01%	0.00%	0.03%	0.08%	0.70%	96.11%	2.65%	0.42%	0.01%
43.0	0.00	1.78	0.00	1.25	0.04	8.96	0.17	94.16	3241.77	58862.20	117.52	2.52	- 40000		6.000%	0.00%	0.00%	0.00%	0.00%	0.01%	0.00%	0.15%	5.20%	94.44%	0.19%	0.00%
44.0	0.00	25.47	0.00	7.53	0.00	26.81	4.50	121.80	507.65	117.52	52865.92	2141.55	- 20000		0.00%	0.05%	0.00%	0.01%	0.00%	0.05%	0.01%	0.22%	0.91%	0.21%	94.71%	3.84%
45.0	0.00	0.11	0.00	1.23	0.00	0.36	0.32	58.26	12.74	2.52	2141.55	4049.60			45.0 - 0.00%	0.00%	0.00%	0.02%	0.00%	0.01%	0.01%	0.93%	0.20%	0.04%	34.17%	64.62%
•	34.0	35.0	36.0	37.0	38.0	39.0	40.0	41.0	42.0	43.0	44.0	45.0	- 0		34.0	35.0	36.0	37.0	38.0	39.0	40.0	41.0	42.0	43.0	44.0	45.0

True allele

- 0.4

True allele