	Confusion matrix											
34.5	27.31	0.02	7.51	115.35	0.50	13.79	11.77	2.19	5.32	28.85	1.47	0.48
35.0	0.02	5.82	0.15	33.45	0.05	0.48	1.28	0.15	29.44	4.38	0.01	0.01
35.5	7.51	0.15	7753.81	382.86	5.60	16.26	52.67	4.77	30.35	70.33	0.62	1.46
36.5	115.35	33.45	382.86	336713.12	286.36	1263.03	5804.56	660.14	2252.30	7026.10	26.10	78.90
37.5	0.50	0.05	5.60	286.36	936.71	78.61	377.29	10.26	11.25	166.19	0.31	0.31
38.5	13.79	0.48	16.26	1263.03	78.61	17935.39	852.29	60.05	134.11	664.25	11.57	5.90
39.5 38.5	11.77	1.28	52.67	5804.56	377.29	852.29	101734.94	129.90	544.93	3129.61	10.86	10.71
40.0	2.19	0.15	4.77	660.14	10.26	60.05	129.90	9500.61	131.49	1585.96	0.86	2.39
40.5	5.32	29.44	30.35	2252.30	11.25	134.11	544.93	131.49	32780.33	2437.50	4.33	16.71
41.5	28.85	4.38	70.33	7026.10	166.19	664.25	3129.61	1585.96	2437.50	122844.34	71.58	110.94
42.5	1.47	0.01	0.62	26.10	0.31	11.57	10.86	0.86	4.33	71.58	78.22	0.46
43.5	0.48	0.01	1.46	78.90	0.31	5.90	10.71	2.39	16.71	110.94	0.46	91.21
	34.5	35.0	35.5	36.5	37.5	38.5	39.5	40.0	40.5	41.5	42.5	43.5

True allele

- 300000

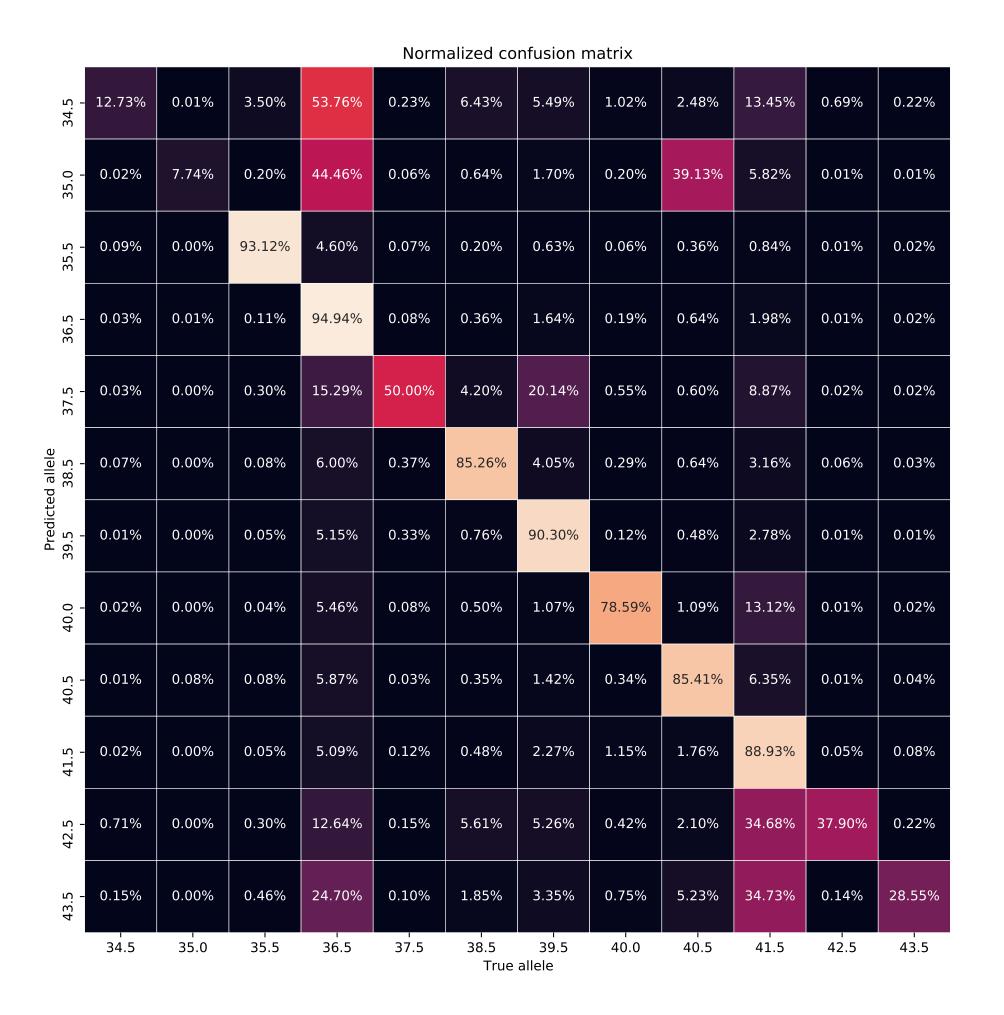
- 250000

- 200000

- 150000

- 100000

- 50000



- 0.8

- 0.6

- 0.4

- 0.2