

	thresholds	500	1000	2000	3000
5 snps	the 1st Qu.	5 (90248)	5 (90746)	5 (90658)	5 (90742)
	the median.	5 (60411)	5 (60506)	5 (60507)	5 (60507)
	the 3rd Qu.	4 (30247)	5 (30252)	4 (30254)	5 (30254)
25 snps	the 1st Qu.	25 (90690)	25 (90425)	25 (90721)	25 (90758)
	the median.	23 (60216)	20 (60506)	23 (60507)	22 (60501)
	the 3rd Qu.	14 (30222)	15 (30254)	15 (30254)	15 (30254)
100 snps	the 1st Qu.	94 (90273)	95 (90733)	94 (90739)	95 (90729)
	the median.	74 (57564)	77 (60487)	79 (60506)	74 (60427)
	the 3rd Qu.	49 (30225)	50 (30254)	51 (30254)	48 (30254)
300 snps	the 1st Qu.	273 (87892)	284 (90661)	283 (90681)	277 (90760)
	the median	219 (60479)	219 (60501)	219 (60506)	217 (60494)
	the 3rd Qu.	148 (30253)	145 (30250)	141 (30254)	144 (30254)

	threshold	500	1000	2000	3000
5 snps	the 1st Qu.	5 (90501)	5 (90755)	5 (90760)	5 (90759)
	the median	5 (60343)	5 (60506)	5 (60495)	5 (60505)
	the 3rd Qu.	3 (30239)	3 (30245)	3 (30254)	3 (30254)
25 snps	the 1st Qu.	25 (89728)	25 (90755)	25 (90743)	25 (90759)
	the median	22 (60046)	22 (60410)	22 (60501)	22 (60485)
	the 3rd Qu.	12 (30253)	13 (30254)	13 (30254)	12 (30254)
100 snps	the 1st Qu.	96 (90063)	96 (90757)	97 (90671)	97 (90752)
	the median.	86 (60489)	88 (60458)	94 (60507)	90 (60505)
	the 3rd Qu.	56 (30254)	55 (30240)	57 (30263)	57 (30254)
300 snps	the 1st Qu.	284 (90632)	291 (90760)	289 (90742)	289 (90760)
	the median	233 (60497)	254 (60497)	268 (60506)	264 (60507)
	the 3rd Qu.	165 (30250)	168 (30252)	164 (30252)	166 (60254)

	threshold	500	1000	2000	3000
50 snps (LD-prune)	the 1st Qu.	49 (90488)	50 (90595)	50 (90757)	50 (90737)
	the median	43 (60450)	45 (60438)	45 (60479)	41 (60491)
	the 3rd Qu.	32 (30254)	33 (30251)	33 (30253)	33 (30254)
50 snps (gene <i>pbpX</i>)	the 1st Qu.	49 (90589)	49 (90753)	49 (90759)	49 (90756)
	the median	43 (60054)	45 (60501)	46 (60507)	45 (60465)
	the 3rd Qu.	34 (30251)	32 (30243)	33 (30254)	32 (30253)
50 snps (50% gene <i>pbpX</i> , 50% gene <i>penA</i>)	the 1st Qu.	47 (90733)	48 (90753)	48 (90754)	48 (90750)
	the median.	40 (60495)	44 (60123)	43 (60506)	42 (60490)
	the 3rd Qu.	25 (30158)	27 (30253)	27 (30254)	26 (30254)
50 snps (16 from gene <i>pbpX</i> , 17 from gene <i>penA</i> , 17 from gene <i>penX</i>)	the 1st Qu.	49 (90720)	49 (90759)	49 (90752)	49 (90757)
	the median	46 (59328)	47 (60459)	46 (60471)	46 (60499)
	the 3rd Qu.	34 (30232)	34 (30254)	35 (30254)	35 (30254)

	threshold	$h^2 = 0.1$	$h^2 = 0.3$	$h^2 = 0.6$	$h^2 = 0.9$
50 snps (gene <i>pbpX</i>)	the 1st Qu.	44 (90760)	44 (90760)	45 (90760)	44 (90760)
	the median	44 (60507)	44 (60507)	44 (60507)	44 (60507)
	the 3rd Qu.	42 (29977)	41 (30234)	42 (30180)	43 (30254)
50 snps (50% gene <i>pbpX</i> , 50% gene <i>penA</i>)	the 1st Qu.	44 (90760)	45 (90760)	45 (90760)	45 (90760)
	the median.	43 (60507)	43 (60507)	43 (60494)	43 (60507)
	the 3rd Qu.	42 (30254)	41 (30254)	41 (30254)	41 (30254)
50 snps (16 from gene <i>pbpX</i> , 17 from gene <i>penA</i> , 17 from gene <i>penX</i>)	the 1st Qu.	49 (90760)	48 (90760)	48 (90760)	48 (60760)
	the median	48 (60507)	47 (60507)	47 (60507)	47 (60506)
	the 3rd Qu.	41 (30254)	42 (30253)	42 (30254)	43 (30254)