

Region	This Study				CanMap Data			Other Selection Studies				Genes
	$\Delta\pi$	Fst	Δ TD	CMS _{1-FDR}	$\Delta\pi$	Fst	Joint	1	2	3	4	
chr18:15.49-15.78	4.03	0.63	1.74	0.97	0.76	0.7	99.25	94.8				LHFPL3
chr18:0.57-1.02	4.36	0.67	1.62	0.97							28	
chr19:3.55-3.72	3.16	0.51	2.09	0.93	0.58	0.83	99.22	93.1				CCRN4L
chr31:1.95-2.14	2.9	0.65	2.17	0.92	0.54	0.4	94.22					CADM2
chr12:59.02-59.17	3.93	0.64	1.32	0.92	-1.06	0.01	0.57					RPL3;ACA64
chr10:4.04-4.22	2.42	0.53	2.21	0.9	0.3	0.61	96.02				18	SNORA31
chr11:37.55-37.76	3.18	0.69	1.62	0.89	0.03	0.47	79.33					SH3GL2
chr24:4.37-4.59	2.17	0.59	1.91	0.89	0.12	0.12	48.36					DTD1
chr24:23.25-23.48	8.91	0.69	1.4	0.89					0.7	0.64		ASIP;AHCY;unknown
chr02:21.9-22.16	1.87	0.93	2.14	0.88	0.18	0.19	68.71					FAM107B
chr32:0.64-0.78	4.09	0.43	2.66	0.88								ART3;NUP54;unknown;SCARB2
chr13:46.85-47.02	3.54	0.42	2	0.87	0.19	0.54	92.4	92.5				
chr13:53.99-54.13	3.71	0.43	1.65	0.87	0.05	0.3	75.43					unknown
chr01:2.93-3.19	2.75	0.75	2.13	0.86	-0.04	0.13	40.36				1	MBP;ZNF236
chr01:28.36-28.52	2.83	0.6	1.52	0.85	0.1	0.51	87.23	93.3				PDE7B
chr02:46.35-46.53	3.19	0.43	2.4	0.85	-1.39	0.34	3.67					
chr06:28.39-28.49	2.1	0.45	1.79	0.83	0.24	0.52	93.8					unknown;PDXDC1;NTAN1;RRN3
chr04:57.36-57.48	2.49	0.45	1.87	0.82	0.16	0.62	91.71					GLRA1
chr37:16.14-16.27	1.9	0.46	2.5	0.82	0.47	0.67	98.18	91.5				PLEKHM3
chr04:60.02-60.27	3.71	0.6	1.01	0.81	-0.14	0.52	54.44					
chr15:9.83-9.93	3.48	0.47	1.12	0.81	-0.36	0.4	31.01					EPS15;TTC39A
chr08:71.21-71.31	3.11	0.4	1.79	0.8								MARK3;U2;CKB;TRMT61A;BAG5;APOPT1
chr04:61.87-62.1	4.16	0.43	1.89	0.8	0.04	0.21	64.58					unknown
chr16:39.84-39.99	3.44	0.39	2.15	0.8	-0.11	0.07	19.51					
chr04:4.17-4.34	2.54	0.42	2.04	0.8	-1.29	0.06	1.34					LYST;unknown;GNG4
chr17:5.62-5.74	1.79	0.55	1.78	0.78	0.74	0.37	93.72					
chr11:48.48-48.71	5.07	0.4	1.58	0.78								
chr10:3.72-3.98	5.07	0.39	1.44	0.78							18	
chr36:0.34-0.52	1.68	0.67	1.63	0.77								MINOS1
chr02:53.12-53.38	2.24	0.41	2.45	0.77	0.27	0.41	91.98					
chr26:8.68-8.79	2.91	0.38	1.86	0.75	-0.05	0.1	29.52					CUX2
chr01:46.69-46.83	4.02	0.4	1.17	0.75	-0.15	0.11	26.5				2	ARID1B
chr18:24.04-24.16	2.41	0.39	1.89	0.75				90.3				
chr28:8.2-8.3	1.55	0.42	1.81	0.74	0.31	0.56	95.59	97.4				PLCE1
chr14:50.91-51.34	2.49	0.43	2.66	0.74	-0.7	0.48	15.13					LRRN3
chr34:14.47-14.61	2.24	0.55	0.97	0.73								unknown
chr04:40.19-40.59	0.98	0.61	2.11	0.73	0.06	0.05	19.88					C5orf50
chr26:2.42-2.61	3.4	0.67	0.96	0.72	-0.01	0.65	73.88					TMEM132D
chr34:26.31-26.51	1.4	0.78	2.34	0.72	-0.09	0.24	51.37					IFT80;SMC4;cfa-mir-15b;cfa-mir-16-2;TRIM59
chr11:46.55-46.92	4.23	0.64	1.14	0.7								cfa-mir-876;unknown
chr16:4.83-4.93	2.24	0.56	0.86	0.69	0.81	0.76	99.56					
chr11:9.72-9.88	5.26	0.41	0.9	0.68	0.08	0.13	49.56					
chr04:40.74-41.09	1.2	0.87	1.26	0.68	-0.96	0.07	2.91				10	RANBP17
chr06:25.29-25.4	8.82	0.44	0.94	0.67								GPR139;GPRC5B
chr32:0.53-0.63	1.31	0.39	1.8	0.67								SDAD1;unknown;CXCL10
chr02:43.98-44.08	3.51	0.35	1.37	0.67	-0.08	0.02	7.12					MAP3K1;SETD9;MIER3
chr31:3.09-3.19	2.29	0.35	1.77	0.67				93.9				
chr24:23.83-23.93	3.38	0.45	0.73	0.66	0.38	0.27	84.69					NCOA6;GGT7;ACSS2
chr18:1.46-1.71	3.34	0.55	1.83	0.66								C7orf72;IKZF1
chr37:4.38-4.57	4.29	0.42	0.77	0.63								
chr25:6.73-6.86	4.14	0.37	0.97	0.63								unknown
chr18:24.26-24.38	3.64	0.34	1.21	0.62	0.29	0.55	95.28	90.3				SEMA3D
chr01:79.87-80.05	3.11	0.58	0.63	0.61							4	unknown
chr21:7.19-7.34	2.2	0.33	2.12	0.6	0.91	0.93	99.85					unknown;CCDC67
chr18:0.45-0.56	3.62	0.4	0.88	0.59							28	
chr04:41.7-41.83	1.47	0.58	0.73	0.59	-0.57	0.18	14.6					FOXI1;DOCK2
chr37:6.56-6.66	2.8	0.55	0.59	0.59	0.28	0.15	60.86					unknown
chr13:48.69-48.82	1.59	0.34	2.05	0.59	0.22	0.3	85.08					SRP72;ARL9;THEGL;HOPX
chr10:6.39-6.5	2.92	0.56	0.53	0.57	-0.11	0.14	35.56					
chr06:33.34-33.56	4.08	0.65	1.02	0.56	-0.11	0.29	52.43					ABAT;METTL22;unknown;TMEM114
chr11:37.8-37.94	3.48	0.5	0.59	0.54							19	
chr01:21.15-21.26	2.2	0.58	0.49	0.53	0.33	0.5	95.13					STARD6;POLI;MBD2
chr16:2.08-2.24	1.39	0.56	0.61	0.52	-0.01	0	0.5					unknown
chr12:58.03-58.13	3.91	0.34	0.74	0.5	0.45	0.19	72.03					MCHR2
chr24:20.91-21.02	1	0.32	1.96	0.5	0.15	0.15	57					DEFB122;CBD118;DEFB119;CBD121;CBD122;DEFB3L;CBD123;DEFB124;CBD124;REM1;
chr01:50.78-50.92	1.23	0.31	1.88	0.48	0.42	0.45	95	95.8				unknown
chr37:12.31-12.42	1.58	0.3	1.94	0.48	0.34	0.38	92.04					RAPH1;unknown;U2
chr16:18.74-18.88	1.64	0.31	1.9	0.48	0.37	0.64	97.09					