		O4	End and the	Lava estla (lava)	# of top 40/ CNIDs within region
					# of top 1% SNPs within region
1	1	7792224	7896757	104533	15
2	1	29158117	29429574	271457	23
3	1	35056561	35336113	279552	25 47
4	1	35507170	35954991	447821	47
5	1	46855808	47044686	188878	13
6	1	48692244	48738338	46094	10
7	1	65202269	65234137	31868	13
8	1	66552479	66592850	40371	12
9	1	72439588	72466646	27058	20
10	1	75251851	75299066	47215	11
11	1	92152172	92376182	224010	17
12	1	144086558	144143214	56656	20
13	1	154307552	154353904	46352	12
14	1	165883926	166024771	140845	42
15	1	166346357	166519547	173190	19
16	1	166681780	166723544	41764	17
17	1	169968307	170102315	134008	16
18	1	170156251	170215393	59142	10
19	1	176923846	176967051	43205	14
20	1	195442649	195604958	162309	38
21	1	196340094	196397371	57277	10
22	1	200916214	200962221	46007	10
23	1	201923735	202026783	103048	23
24	1	202833886	202890572	56686	10
25 26	1	208774473	208800319	25846	13
26 27	1 1	229667496 231315554	229723531 231352079	56035 36525	29 18
28	1	231367326	231423057	55731	17
29	1	231866718	231423037	32740	10
30	1	242939710	243063807	124097	25
31	2	13851646	13918802	67156	24
32	2	16780080	16802058	21978	10
33	2	17114840	17206352	91512	17
34	2	17251570	17423034	171464	59
35	2	17446983	17667050	220067	25
36	2	17760353	17837901	77548	13
37	2	17957610	18024021	66411	28
38	2	19049791	19081464	31673	10
39	2	19684280	19756560	72280	26
40	2	25993269	26148038	154769	13
41	2	26406754	26653031	246277	26
42	2	38074516	38197596	123080	45
43	2	38416518	38458268	41750	16
44	2	40370011	40386252	16241	13
45	2	40735500	40810538	75038	26
46	2	40903943	40952139	48196	20
47	2	44179533	44234161	54628	19
48	2	55857206	55881997	24791	10
49	2	60615770	60695881	80111	14
50	2	69192866	69249852	56986	10
51	2	74612395	74799689	187294	13

52	2	74842869	74940127	97258	38
53	2	86570541	86615534	44993	10
54	2	104335362	104566945	231583	20
55	2	107789200	108010333	221133	46
56	2	108507295	108558105	50810	11
57	2	108686523	108877935	191412	19
58	2	109135001	109250374	115373	40
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60	2	123068296	123216160	147864	38
61	2	126123179	126413028	289849	72
62	2	135792281	135916327	124046	11
63	2	153350334	153425089	74755	16
64	2	153497060	153543393	46333	16
65	2	154991040	155035601	44561	10
66	2	164314128	164405200	91072	10
67	2	171563920	171643068	79148	23
68	2	173348329	173510862	162533	23
69	2	177817852	177847126	29274	21
70	2	179847398	179907516	60118	16
71	2	211791790	211840711	48921	10
72	2	212728567	212758365	29798	12
73	2	216479398	216532339	52941	19
74	2	216763322	216813395	50073	23
75	2	218061201	218104452	43251	14
76	2	223645065	223730976	85911	16
77	3	2622113	2636675	14562	10
78	3	3163016	3213726	50710	16
79	3	36074643	36111561	36918	16
80	3	49349262	49516063	166801	11
81	3	58052200	58075903	23703	10
82	3	64229886	64290858	60972	31
83	3	64309210	64375379	66169	17
84	3	71426913	71535395	108482	13
85	3	108953294	109010111	56817	14
86	3	109338409	109382830	44421	11
87	3	121253963	121369516	115553	13
88 89	3 3	121423609 124670392	121459597	35988	10
	3		124740474	70082	10
90	3	125262001	125277705	15704 28927	10 14
91	3	141599911	141628838		
92	3	161955802	161998153	42351	10
93		162026513	162069293	42780	10
94	3	163781804	163817888	36084	10
95 06	3	176938190	176989747	51557	14
96	3	179156942	179189986	33044	13
97	3	197925843	198044316	118473	12
98	4	6016995	6045112	28117	24
99	4	21397803	21425668	27865	12
100	4	22652230	22849845	197615	32
101	4	27166805	27232128	65323	14
102	4	29140421	29178507	38086	14
103	4	38128437	38177892	49455	13

104	4	38408315	38433315	25000	12
105	4	43253692	43328678	74986	21
106	4	57024995	57037233	12238	10
107	4	60998034	61064279	66245	16
108	4	63255385	63332752	77367	24
109	4	73375440	73517739	142299	26
110	4	73572135	73624879	52744	12
111	4	86950413	87070744	120331	33
112	4	87123335	87147766	24431	10
113	4	91071226	91160540	89314	20
114	4	111381500	111406702	25202	12
115	4	135431375	135868494	437119	76
116	4	153388383	153458454	70071	16
117	4	153545357	153677816	132459	43
118	4	160730848	160899142	168294	25
119	4	167774644	167835716	61072	34
120	4	171772452	171844514	72062	17
121	4	176751863	176813538	61675	20
122	4	180232731	180286178	53447	19
123	5	2910280	2936337	26057	10
124	5	4969220	4992993	23773	13
125	5	10314172	10341846	27674	10
126	5	13646073	13659894	13821	14
127	5	18126844	18168395	41551	11
128	5	28279274	28338829	59555	13
129	5	30074058	30085976	11918	14
130	5	30480325	30648051	167726	15
131	5	33992072	34011634	19562	11
132	5	39629840	39759458	129618	25
133	5	40266628	40301827	35199	10
134	5	66850378	66925736	75358	19
135	5	72863229	72914103	50874	26
136	5	104400223	104431728	31505	12
137	5	108720962	108883620	162658	24
138	5	128387641	128406390	18749	17
139	5	132085926	132178119	92193	22
140	5	136335816	136391843	56027	10
141	5	142273506	142436334	162828	40
142	5	142456316	142486468	30152	10
143	5	145865172	145924847	59675	13
144	5	147045949	147054555	8606	15
145	5	152830782	152856226	25444	12
146	5	153137695	153286146	148451	50
147	5	153524278	153582509	58231	25
148	5	167044409	167085287	40878	17
149	5	170102703	170180754	78051	16
150	6	6949860	7102039	152179	25
151	6	22151788	22203585	51797	10
152	6	38294473	38493205	198732	34
153	6	44388835	44464921	76086	12
154	6	54829488	54904124	74636	32
155	6	56163423	56182022	18599	11

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157	6	91357650	91374538	16888	17
158	6	93435249	93489970	54721	14
159	6	93635981	93690767	54786	22
160	6	97698965	97813189	114224	19
161	6	121227677	121295655	67978	36
162	6	122184180	122268449	84269	18
163	6	124122315	124129724	7409	11
164	6	126589385	127059355	469970	68
165	6	127766722	127935184	168462	30
166	6	127982571	128129811	147240	18
167	6	131077658	131113533	35875	12
168	6	144858907	144905759	46852	11
169	6	144931561	145084047	152486	18
170	6	145239076	145268835	29759	17
171	6	156759285	156772699	13414	13
172	6	159547865	159573610	25745	29
172	6	166430647	166459673	29026	22
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174	6	169751423	169920385	168962	
175	6	169951018	170034558	83540	20
176	6	170510772	170597840	87068	18
177	7	9840940	9868091	27151	12
178	7	14042164	14056377	14213	12
179	7	14405099	14456612	51513	13
180	7	25467843	25499533	31690	11
181	7	25531780	25547719	15939	14
182	7	27884458	27921718	37260	14
183	7	80306942	80347005	40063	12
184	7	83050708	83114306	63598	18
185	7	84825282	84954886	129604	17
186	7	85544786	85720004	175218	16
187	7	88273509	88339369	65860	20
188	7	101232338	101380336	147998	21
189	7	112615625	112700875	85250	11
190	7	112910324	113036180	125856	14
191	7	130226349	130283457	57108	14
192	8	11467894	11537376	69482	15
193	8	11539406	11636979	97573	14
194	8	12733069	12780372	47303	15
195	8	31840607	31868316	27709	20
196	8	37465204	37506646	41442	13
197	8	62194953	62241990	47037	32
198	8	66262374	66363868	101494	15
199	8	73867598	73940386	72788	33
200	8	111649715	111792276	142561	12
201	8	122558472	122574815	16343	10
202	8	122640877	122673774	32897	10
203	8	126840040	126938040	98000	27
204	9	9809049	9851006	41957	10
205	9	10363600	10423023	59423	11
206	9	12391209	12543171	151962	47
207	9	12568438	12765488	197050	79

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208	9	16651092	16735514	84422	20
209	9	16785286	16821748	36462	10
210	9	26089466	26157022	67556	18
211	9	26799022	27127654	328632	55
212	9	34965508	35079498	113990	13
213	9	83368043	83407378	39335	10
214	9	83463865	83813052	349187	48
215	9	86567954	86657492	89538	17
216	9	87802219	87830149	27930	41
217	9	98441204	98481793	40589	13
218	9	100471446	100518062	46616	11
219	9	109395321	109433839	38518	13
220	10	4254320	4286824	32504	14
221	10	5890155	5908924	18769	11
222	10	8259087	8316999	57912	14
223	10	8783468	8835711	52243	22
224	10	20378684	20394280	15596	10
225	10	23501565	23667419	165854	31
226	10	28361372	28446847	85475	22
227	10	37128805	37230563	101758	31
228	10	49042716	49140239	97523	32
229	10	54880153	54932740	52587	16
230	10	55214898	55242747	27849	13
231	10	55262962	55433826	170864	60
232	10	68929453	69024427	94974	20
233	10	73379900	73520018	140118	15
234	10	73549532	73962617	413085	61
235	10	77406138	77436812	30674	13
236	10	81834175	81877658	43483	14
237	10	94477764	94697070	219306	67
238	10	94773161	94842631	69470	17
239	10	109230427	109241819	11392	11
240	11	4849514	4889476	39962	13
241	11	5519812	5533605	13793	15
242	11	5557776	5572157	14381	15
243	11	23762166	23796832	34666	13
244	11	32341203	32375549	34346	12
245	11	38365710	38571109	205399	51
246	11	44700663	44749595	48932	14
247	11	61493075	61524162	31087	11
248	11	80985427	81010491	25064	12
249	11	99872041	99950416	78375	13
250	11	119638249	119708817	70568	19
251	12	4039860	4126366	86506	15
252	12	12024132	12053666	29534	11
253	12	22095425	22135645	40220	12
254	12	25230671	25275080	44409	14
255	12	32980008	33037082	57074	12
256	12	33137238	33309428	172190	40
257	12	33629037	33719472	90435	22
258	12	47070467	47172762	102295	19
259	12	52458569	52544208	85639	27

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260	12	53626666	53691996	65330	30
261	12	67137191	67173223	36032	10
262	12	73449504	73551061	101557	15
263	12	85692457	85785491	93034	13
264	12	92850691	92867117	16426	10
265	12	110154175	111455363	1301188	126
266	13	19072807	19247692	174885	47
267	13	23902547	23958467	55920	34
268	13	29743277	29760283	17006	11
269	13	32625602	32678316	52714	16
270	13	32720737	32737733	16996	15
271	13	41744257	41811922	67665	18
272	13	41946931	41980423	33492	11
273	13	57656354	57697478	41124	14
274	13	61097059	61192794	95735	20
275	13	61323097	61690524	367427	105
276	13	62001263	62015319	14056	23
277	13	70135944	70172850	36906	10
278	13	70185834	70229368	43534	22
279	13	71752804	71765917	13113	 10
280	13	71905253	71924535	19282	11
281	13	85996226	86067435	71209	10
282	13	87596501	87699922	103421	28
283	13	89475732	89508535	32803	17
284	13	91694613	91749890	55277	20
285	13	93244651	93271433	26782	14
286	13	96417299	96450726	33427	12
287	13	107161131	107194602	33471	13
288	13	109656667	109680616	23949	11
289	14	22745635	22846783	101148	20
290	14	30868687	30876603	7916	13
291	14	33124367	33201040	76673	10
292	14	41224268	41263424	39156	15
293	14	44786876	44916436	129560	37
294	14	47208922	47329393	120471	21
295	14	48003860	48102105	98245	20
296	14	63312060	63368417	56357	19
297	14	67351554	67398608	47054	22
298	14	67420411	67643874	223463	23
299	14	94067518	94073821	6303	20
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300	14	95232641	95346960		
301	14	99371605	99515197	143592	20
302	15	25852515	25976371	123856	14
303	15	43287724	43440039	152315	55
304	15	44586911	44596891	9980	16
305	15	45984817	46027466	42649	11
306	15	46092560	46149550	56990	10
307	15	48295500	48382496	86996	15
308	15	50201080	50269509	68429	19
309	15	61021149	61037344	16195	10
310	15	61415521	61877168	461647	88
311	15	61976761	62027981	51220	11

312	15	62820643	62914408	93765	17
313	15	69654175	69683364	29189	42
314	15	89228014	89298922	70908	10
315	15	89823285	89847368	24083	16
316	16	5737814	5773535	35721	15
317	16	10271364	10299314	27950	10
318	16	20587355	20646759	59404	13
319	16	59107572	59156992	49420	14
320	16	62221993	62235327	13334	10
321	16	65617179	65659731	42552	13
322	16	75338036	75562484	224448	40
323	16	75939634	75954187	14553	18
324	16	78169979	78210538	40559	16
325	16	85587338	85608958	21620	12
326	16	89539805	89609838	70033	19
327	17	14619706	14642944	23238	12
328	17	17862368	18255795	393427	39
329	17	20154543	20381537	226994	34
330	17	30771281	30975214	203933	21
331	17	48927414	49050363	122949	15
332	17	64522305	64746634	224329	42
333	17	69995635	70035019	39384	21
334	18	7585095	7617199	32104	10
335	18	31158770	31232197	73427	12
336	18	36382489	36415916	33427	10
337	18	39167190	39207577	40387	16
338	18	54929757	54972968	43211	15
339	18	67707371	67766235	58864	11
340	19	4900266	5012052	111786	10
341	19	39328872	39420460	91588	23
342	19	49588016	49605296	17280	13
343	20	1455272	1482850	27578	13
344	20	1904445	1969008	64563	30
345	20	2028591	2141407	112816	58
346	20	7843067	7900947	57880	11
347	20	18361623	18392366	30743	13
348	20	20169188	20224681	55493	21
		22371059			
349	20		22417151	46092	14
350	20	31747352	32140745	393393	52
351	20	52236817	52261680	24863	13
352	20	60202852	60209916	7064	11
353	20	60416190	60431631	15441	10
354	20	62957281	63049047	91766	14
355	21	23706934	23716305	9371	18
356	21	25832335	25882957	50622	10
357	21	28760070	28787610	27540	11
358	21	38837152	38857481	20329	17
359	21	41724902	41750242	25340	14
360	21	42875375	42914787	39412	26
361	22	21986644	22033028	46384	11
362	22	41104264	41270923	166659	14
363	22	41552747	41585275	32528	11

# of Perlegen SNPs within region	ratio of top 1% SNPs to total # of SNPs	average Fst	average Fst p-value
36	0.417	0.493	0.010
76	0.303	0.391	0.016
79	0.316	0.539	0.001
68	0.691	0.700	0.000
49	0.265	0.389	0.016
33	0.303	0.464	0.007
31	0.419	0.386	0.014
45	0.267	0.449	0.008
37	0.541	0.523	0.000
20	0.550	0.499	0.003
57	0.298	0.413	0.015
42	0.476	0.473	0.008
21	0.571	0.476	0.005
87	0.483	0.572	0.001
60	0.317	0.438	0.006
62	0.274	0.467	0.006
39	0.410	0.538	0.002
35	0.286	0.446	0.012
31	0.452	0.403	0.019
127	0.299	0.350	0.032
35	0.286	0.409	0.011
34	0.294	0.354	0.024
75	0.307	0.485	0.003
37	0.270	0.436	0.011
27	0.481	0.475	0.010
98	0.296	0.377	0.018
43	0.419	0.549	0.000
53	0.321	0.632	0.000
37	0.270	0.424	0.015
60	0.417	0.483	0.006
51	0.471	0.456	0.004
33	0.303	0.398	0.015
52	0.327	0.341	0.032
119	0.496	0.525	0.006
101	0.248	0.464	0.008
45	0.289	0.454	0.004
65	0.431	0.568	0.001
21	0.476	0.456	0.005
87	0.299	0.490	0.005
20	0.650	0.584	0.003
78	0.333	0.487	0.001
120	0.375	0.523	0.002
31	0.516	0.493	0.005
43	0.302	0.450	0.008
51	0.510	0.486	0.007
51	0.392	0.351	0.021
47	0.404	0.490	0.007
35	0.286	0.308	0.055
46	0.304	0.475	0.005
33	0.303	0.431	0.015
44	0.295	0.643	0.001

76	0.500	0.700	0.000
30	0.333	0.495	0.003
53	0.377	0.543	0.003
101	0.455	0.493	0.001
38	0.289	0.457	0.006
58	0.328	0.460	0.004
98	0.408	0.761	0.000
17	0.647	0.445	0.011
84	0.452	0.455	0.005
191	0.377	0.552	0.004
52	0.212	0.317	0.036
35	0.457	0.550	0.003
43	0.372	0.517	0.002
36	0.278	0.437	0.013
30	0.333	0.492	0.003
67	0.343	0.416	0.005
84	0.274	0.452	0.013
62	0.339	0.541	0.001
61	0.262	0.481	0.004
35	0.286	0.366	0.021
37	0.324	0.455	0.005
63	0.302	0.476	0.005
69	0.333	0.571	0.000
43	0.326	0.484	0.007
48	0.333	0.356	0.034
25	0.400	0.449	0.005
54	0.296	0.528	0.002
57	0.281	0.407	0.017
37	0.297	0.468	0.011
29	0.345	0.523	0.004
71	0.437	0.478	0.004
58	0.293	0.517	0.002
41	0.317	0.485	0.002
40	0.350	0.459	0.003
38	0.289	0.461	0.002
41	0.317	0.458	0.007
36	0.278	0.403	0.011
34	0.294	0.500	0.005
20	0.500	0.553	0.002
32	0.438	0.519	0.007
35	0.286	0.365	0.017
34	0.294	0.366	0.024
25	0.400	0.444	0.010
30	0.467	0.470	0.008
27	0.481	0.548	0.001
34	0.353	0.432	0.008
68	0.353	0.445	0.005
33	0.364	0.508	0.005
70	0.457	0.508	0.003
52	0.269	0.379	0.022
40	0.350	0.502	0.007
49	0.265	0.569	0.001

33	0.364	0.517	0.002
55	0.382	0.478	0.008
31	0.323	0.496	0.008
57	0.281	0.442	0.015
64	0.375	0.429	0.006
61	0.426	0.558	0.000
40	0.300	0.322	0.029
80	0.413	0.523	0.002
29	0.345	0.543	0.003
62	0.323	0.391	0.014
32	0.375	0.398	0.014
214	0.355	0.478	0.005
57	0.281	0.394	0.011
107	0.402	0.498	0.003
71	0.352	0.562	0.001
54	0.630	0.626	0.001
45	0.378	0.468	0.006
39	0.513	0.455	0.005
68	0.279	0.323	0.026
28	0.357	0.408	0.007
23	0.565	0.670	0.001
29	0.345	0.385	0.014
26	0.538	0.522	0.003
28	0.393	0.476	0.009
41	0.317	0.421	0.011
24	0.583	0.496	0.003
32	0.469	0.579	0.001
24	0.458	0.692	0.000
63	0.397	0.428	0.008
34	0.294	0.360	0.029
42	0.452	0.492	0.001
50	0.520	0.425	0.004
38	0.316	0.391	0.017
66	0.364	0.389	0.010
37	0.459	0.365	0.024
46	0.478	0.490	0.003
35	0.286	0.443	0.004
104	0.385	0.481	0.006
34	0.294	0.370	0.021
39	0.333	0.487	0.005
19	0.789	0.502	0.002
24	0.500	0.504	0.005
99	0.505	0.563	0.000
35 	0.714	0.554	0.000
37	0.459	0.516	0.003
52	0.308	0.438	0.008
54	0.463	0.483	0.003
37	0.270	0.352	0.028
93	0.366	0.435	0.015
32	0.375	0.465	0.011
76	0.421	0.494	0.003
20	0.550	0.515	0.002

37	0.270	0.392	0.007
18	0.944	0.620	0.000
40	0.350	0.505	0.004
65	0.338	0.463	0.002
64	0.297	0.411	0.006
82	0.439	0.496	0.004
43	0.419	0.457	0.009
16	0.688	0.480	0.008
109	0.624	0.604	0.000
91	0.330	0.390	0.001
57	0.316	0.469	0.017
31	0.387	0.420	0.012
31	0.355	0.385	0.012
72	0.250	0.409	0.009
37	0.459	0.475	0.000
15	0.867	0.566	0.000
55	0.527	0.515	0.002
54	0.407	0.400	0.015
51	0.275	0.395	0.015
51	0.392	0.435	0.007
41	0.439	0.451	0.004
21	0.571	0.508	0.004
21	0.571	0.484	0.007
48	0.271	0.410	0.009
35	0.314	0.464	0.006
41	0.341	0.492	0.003
39	0.359	0.586	0.000
35	0.343	0.435	0.002
43	0.419	0.465	0.004
43	0.395	0.403	0.015
60	0.267	0.371	0.023
48	0.417	0.374	0.023
67	0.313	0.499	0.009
36		0.499	
	0.306		0.016
39	0.359	0.419	0.010
24	0.583	0.506	0.001
44	0.341	0.424	0.006
47	0.298	0.392	0.013
42	0.357	0.436	0.007
40	0.500	0.508	0.001
21	0.619	0.549	0.003
81	0.395	0.554	0.000
48	0.313	0.448	0.002
94	0.351	0.563	0.001
41	0.293	0.457	0.005
21	0.476	0.518	0.005
36	0.278	0.409	0.013
67	0.403	0.486	0.005
34	0.294	0.462	0.002
26	0.423	0.517	0.004
133	0.353	0.462	0.008
187	0.422	0.473	0.003
101	U.722	0.770	0.000

61	0.328	0.480	0.006
32	0.313	0.510	0.003
56	0.321	0.413	0.010
162	0.340	0.486	0.000
45	0.289	0.383	0.012
34	0.294	0.393	0.014
134	0.358	0.424	0.006
24	0.708	0.732	0.000
43	0.953	0.732	0.000
23	0.565	0.500	0.007
38	0.289	0.364	0.019
39	0.333	0.463	0.004
33	0.424	0.466	0.008
11	1.000	0.534	0.001
51	0.275	0.467	0.003
59	0.373	0.454	0.003
32	0.313	0.320	0.031
79	0.392	0.566	0.002
54	0.407	0.627	0.000
65	0.477	0.478	0.004
101	0.317	0.380	0.015
57	0.281	0.468	0.003
38	0.342	0.497	0.001
160	0.375	0.522	0.002
37	0.541	0.481	0.000
47	0.319	0.484	0.004
114	0.535	0.519	0.003
39	0.333	0.529	0.003
36	0.389	0.455	0.008
140	0.479	0.573	0.002
65	0.262	0.478	0.002
27	0.407	0.479	0.007
39	0.333	0.463	0.002
28			0.005
	0.536	0.472	
38	0.395	0.403	0.011
23	0.565	0.524	0.000
41	0.293	0.481	0.003
122	0.418	0.439	0.001
27	0.519	0.617	0.001
29	0.379	0.500	0.004
40	0.300	0.326	0.037
42	0.310	0.368	0.016
34	0.559	0.554	0.001
42	0.357	0.519	0.002
20	0.550	0.527	0.002
35	0.343	0.442	0.005
18	0.778	0.526	0.001
41	0.293	0.440	0.004
98	0.408	0.543	0.001
76	0.289	0.438	0.010
59	0.322	0.351	0.014
73	0.370	0.429	0.008
. 5	0.070	3.120	5.555

56	0.536	0.649	0.000
35	0.286	0.296	0.038
42	0.357	0.419	0.007
40	0.325	0.419	0.007
18	0.556	0.473	0.014
281	0.448	0.473	0.000
109	0.431	0.565	0.001
71	0.479	0.521	0.000
25	0.440	0.465	0.002
34	0.471	0.501	0.002
37	0.405	0.563	0.003
52	0.346	0.480	0.003
39	0.282	0.353	0.015
48	0.292	0.430	0.008
56	0.357	0.515	0.005
252	0.417	0.506	0.001
40	0.575	0.566	0.002
36	0.278	0.447	0.005
55	0.400	0.432	0.007
36	0.278	0.384	0.013
18	0.611	0.461	0.007
33	0.303	0.413	0.013
48	0.583	0.618	0.001
51	0.333	0.615	0.000
56	0.357	0.466	0.001
22	0.636	0.501	0.003
40	0.300	0.429	0.002
24	0.542	0.531	0.002
33	0.333	0.392	0.009
63	0.317	0.433	0.007
17	0.765	0.517	0.007
26	0.765	0.502	0.000
23	0.652	0.596	0.002
79	0.468	0.444	0.006
66	0.318	0.428	0.012
42	0.476	0.483	0.002
49	0.388	0.388	0.012
45	0.489	0.516	0.001
74	0.311	0.471	0.005
24	0.833	0.502	0.004
67	0.269	0.467	0.003
47	0.426	0.427	0.006
58	0.241	0.492	0.005
108	0.509	0.525	0.005
21	0.762	0.651	0.000
41	0.268	0.400	0.010
36	0.278	0.577	0.000
50	0.300	0.303	0.031
47	0.404	0.514	0.002
23	0.435	0.443	0.002
225	0.391	0.661	0.001
37	0.297	0.497	0.002

34	0.500	0.490	0.005
52	0.808	0.568	0.001
37	0.270	0.396	0.011
25	0.640	0.510	0.001
45	0.333	0.403	0.010
36	0.278	0.546	0.002
39	0.333	0.377	0.013
40	0.350	0.306	0.042
14	0.714	0.557	0.002
34	0.382	0.539	0.004
104	0.385	0.506	0.001
23	0.783	0.563	0.001
41	0.390	0.472	0.002
39	0.308	0.453	0.005
45	0.422	0.465	0.003
32	0.375	0.519	0.002
101			
	0.386	0.434	0.005
95	0.358	0.405	0.002
40	0.525	0.545	0.000
60	0.250	0.449	0.005
126	0.333	0.498	0.003
60	0.350	0.367	0.010
24	0.417	0.512	0.000
41	0.293	0.407	0.008
21	0.476	0.591	0.000
25	0.640	0.482	0.004
52	0.288	0.379	0.008
30	0.367	0.431	0.004
36	0.278	0.415	0.005
58	0.397	0.514	0.001
21	0.619	0.590	0.000
27	0.481	0.563	0.000
87	0.345	0.448	0.006
89	0.652	0.610	0.000
40	0.275	0.460	0.002
26	0.500	0.406	0.004
39	0.538	0.537	0.000
52	0.269	0.442	0.002
154	0.338	0.474	0.001
28	0.464	0.526	0.000
15	0.733	0.622	0.000
32	0.733	0.468	0.000
33	0.424	0.453	0.006
33	0.545	0.423	0.003
37	0.270	0.375	0.007
35	0.314	0.438	0.004
35	0.486	0.451	0.002
36	0.389	0.378	0.012
62	0.419	0.517	0.001
31	0.355	0.453	0.001
46	0.304	0.486	0.002
36	0.306	0.470	0.000
- -	0.000	3	0.000

het-ratio (Chn/Fur)	het-ratio n-value	Eur_LRHtest_# of HapMap SNPs within region
0.662	0.131	19
1.574	0.972	42
7.628	1.000	36
3.993	0.998	63
0.804	0.274	48
0.906	0.435	16
1.180	0.819	11
0.418	0.032	20
1.143	0.808	9
0.128	0.008	0
0.131	0.004	55
0.351	0.024	0
1.046	0.669	12
0.207	0.004	39
1.409	0.949	45
0.647	0.109	19
0.512	0.060	30
0.391	0.033	20
1.064	0.703	13
1.594	0.982	55
0.540	0.068	15
0.949	0.515	13
2.126	0.992	30
0.836	0.323	0
1.395	0.913	7
1.763	0.986	18
0.403	0.027	15
0.459	0.040	10
0.159	0.007	11
1.028	0.686	25
1.211	0.864	18
0.729	0.189	6
1.324	0.919	33
0.835	0.304	56
0.394	0.016	63
0.388	0.028	22
0.223	0.008	29
1.003	0.606	11
0.588	0.061	25
0.662	0.163	33
1.135	0.838	64
0.722	0.144	37
0.927	0.479	12
1.078	0.742	6
4.561	0.999	28
1.560	0.960	20
0.168	0.007	13
0.992	0.597	12
0.728	0.185	20
3.825	0.997	17
2.515	0.992	33

1.184 0.393 1.009 1.545 0.538 0.048 0.254 1.480 1.703	0.872 0.038 0.636 0.972 0.071 0.001 0.007 0.915	19 0 73 0 0 14 9 3 42
0.371	0.007	97
0.735	0.182	32
0.609	0.099	26
0.781	0.237	12
0.911	0.456	13
2.700	0.991	22
1.531	0.964	20
0.675	0.114	49
0.201	0.007	12
1.068	0.748	14
0.957	0.535	22
0.507	0.056	10
0.222	0.007	22
1.670	0.978	19
0.249	0.012	9
1.036	0.681	30
0.426	0.048	5
1.377	0.932	17
1.130	0.816	18
0.227	0.012	35
0.304	0.020	11
1.146	0.837	24
1.176	0.849	22
0.110	0.004	44
0.379	0.024	20
1.764	0.967	12
1.199	0.848	42
0.465	0.045	18
2.058	0.984	19
3.474	0.994	8
0.890	0.417	8
0.667	0.136	14
0.683	0.153	17
0.278	0.021	0
1.135	0.782	18
1.047	0.681	7
0.639	0.120	34
0.270	0.008	8
0.698	0.162	8
0.911	0.455	43
0.887	0.404	17
1.050	0.694	11
0.822	0.298	18

1.692	0.962	7
2.095	0.987	22
1.722	0.962	3
2.037	0.986	15
0.764	0.204	30
3.790	0.999	32
1.015	0.638	12
0.215	0.005	30
1.433	0.917	8
1.539	0.962	28
2.429	0.990	6
0.262	0.003	112
0.615	0.088	19
0.437	0.022	26
0.176	0.006	35
2.258	0.992	0
0.274	0.013	17
1.843	0.977	14
0.908	0.438	20
0.650	0.140	7
0.075	0.004	4
0.510	0.069	14
0.905	0.444	3
0.910	0.442	14
1.784	0.974	24
0.524	0.077	3
1.074	0.721	35
56.264	1.000	0
1.513	0.962	33
0.457	0.044	12
0.525	0.060	19
1.059	0.719	19
0.971	0.563	7
0.520	0.051	45
1.666	0.968	9
1.025	0.661	23
1.023	0.734	11
0.831	0.734	55
1.031	0.660	11
0.789	0.252	22
3.216	0.232	4
0.495	0.991	9
0.493	0.004	45
0.239	0.005	17
1.282	0.891	8
	0.065	o 21
0.553		44
0.544	0.061	
1.279	0.890	11
1.081	0.773	53 10
2.070	0.983	19
0.359	0.016	28
0.944	0.508	6

0.702	0.161	27	
0.841	0.349	8	
0.944	0.510	15	
1.073	0.754	24	
0.809	0.259	36	
0.462	0.034	38	
2.554	0.993	28	
		20	
6.018	0.998		
0.113	0.002	103	
0.601	0.074	59 46	
0.294	0.013	46	
1.221	0.847	12	
1.561	0.947	13	
1.242	0.895	40	
2.802	0.996	9	
5.566	0.996	2	
1.014	0.645	9	
0.659	0.116	6	
0.810	0.268	0	
0.784	0.232	0	
0.887	0.406	0	
6.993	0.999	9	
0.504	0.071	7	
0.813	0.282	13	
0.865	0.377	12	
0.768	0.225	6	
0.790	0.255	15	
1.518	0.945	12	
1.421	0.937	13	
0.634	0.105	32	
0.996	0.624	47	
0.700	0.154	20	
0.050	0.001	11	
0.940	0.509	26	
0.546	0.071	28	
0.631	0.134	11	
0.463	0.043	23	
0.750	0.206	26	
1.625	0.960	16	
7.687	1.000	12	
3.747	0.995	14	
0.521	0.042	45	
0.975	0.570	50	
0.630	0.085	63	
1.373	0.924	44	
0.306	0.025	8	
0.788	0.259	22	
0.539	0.053	45	
0.746	0.208	23	
1.495	0.929	19	
0.627	0.068	84	
0.475	0.016	119	

1.513 0.853 0.492 0.561 1.013 0.973	0.961 0.356 0.045 0.033 0.646 0.558	53 20 30 113 46 28
1.170 3.888	0.884 0.997	123 0
0.016 0.524	0.001 0.082	0 14
0.903	0.441	25
0.905 0.640	0.442 0.116	19 11
0.206	0.024	12
0.671	0.124	26
0.618	0.097	16
0.494	0.056	3
0.529	0.042	44
1.970	0.986	36
1.797	0.980	16
0.511	0.034	26
0.601 0.208	0.080 0.009	16 9
0.278	0.004	66
1.175	0.833	30
0.246	0.010	30
0.196	0.003	89
1.127	0.789	9
0.226	0.012	18
0.537	0.037	66
0.659	0.111	30
1.452 0.556	0.925 0.075	8 11
0.494	0.060	11
0.433	0.039	8
2.142	0.978	11
1.327	0.912	6
0.695	0.119	54
0.878	0.394	15
1.051	0.684	12
0.372	0.023	8
0.636 0.691	0.115 0.160	23 24
0.571	0.079	14
0.515	0.081	10
1.325	0.900	14
0.311	0.028	10
0.799	0.266	19
0.752	0.170	57
0.968	0.565	17
1.203	0.866	23
0.479	0.040	27

0.929	0.483	29
1.136	0.791	15
1.069	0.732	32
0.656	0.119	33
0.122	0.007	6
1.444	0.988	199
0.493	0.988	59
0.493 0.167		
	0.006	23 7
0.949	0.520	
1.383	0.922	21
0.534	0.066	8
0.929	0.483	20
1.091	0.749	10
0.674	0.131	13
0.370	0.022	27
0.313	0.003	138
4.956	0.999	4
0.670	0.137	15
2.084	0.987	12
1.148	0.805	4
3.500	0.993	3
0.682	0.150	20
2.894	0.996	36
3.620	0.998	12
1.013	0.652	23
1.813	0.961	7
0.942	0.506	17
0.488	0.063	6
0.623	0.105	8
0.990	0.614	33
0.618	0.144	2
0.811	0.307	20
3.051	0.993	11
0.156	0.004	37
0.753	0.194	30
0.620	0.194	23
	0.965	23
1.602		
1.106	0.774	16
0.269	0.009	61
0.649	0.144	1
0.307	0.011	30
0.352	0.020	36
0.830	0.301	0
0.696	0.127	39
1.217	0.831	2
1.674	0.968	10
10.178	1.000	8
0.633	0.109	26
1.143	0.812	22
1.420	0.912	4
0.089	0.000	142
0.376	0.027	16

1.000 0.993 0.829 0.827 0.586 0.307 1.057 1.399	0.608 0.609 0.312 0.321 0.080 0.018 0.704 0.923	33 10 27 8 12 9 15
0.354	0.044	3
0.204	0.009	13
1.078	0.792	56
0.794	0.286	6
0.913	0.448	14
1.236	0.866	10
0.506	0.054	28
2.239	0.987	4
0.393	0.016	5
0.281	0.008	0
0.479	0.044	38
0.462	0.034	34
0.134	0.001	67
1.327	0.929	14
4.388	0.998	7
1.086	0.746	32
0.133	0.008	17
0.161	0.010	32
1.271	0.899	30
1.174	0.814	16
0.759	0.223	28
0.201	0.008	37
1.631	0.945	7
0.904	0.436	12
0.946	0.516	23
0.580	0.062	30
1.584	0.955	17
0.926	0.478	13
2.003	0.983	18
3.074	0.997	17
0.499	0.024	96
1.939	0.973	12
2.254	0.979	6
0.958	0.526	4
0.775	0.241	22
1.064	0.708	14
1.307	0.902	23
1.225	0.853	14
0.944	0.512	14
0.805	0.280	26
0.485	0.043	22
0.350	0.026	11
0.882	0.396	54
0.872	0.387	18

	Eur_LRHtest_proportion of extreme SNPs	Eur_LRHtest_p-value
0	0	
3	0.071428571	0.183
25	0.69444444	0.010
24	0.380952381	0.037
6	0.125	0.132
0	0	
0	0	
1	0.05	
0	0	
0	NA	
0	0	
0	NA	
0	0	
0	0	
0	0	
1	0.052631579	0.217
0	0	
0	0	
1	0.076923077	0.194
17	0.309090909	0.057
0	0	
0	0	
0	0	
0	NA	
0	0	
0	0	
1	0.06666667	0.200
0	0	
0	0	
12	0.48	0.018
1	0.05555556	0.222
0	0	
0	0	
0	0	
1	0.015873016	
0	0	
1	0.034482759	
5	0.454545455	0.018
0	0	
0	0	
0	0	
1	0.027027027	
0	0	
0	0	
1	0.035714286	
0	0	
0	0	
0	0	
0	0	
4	0.235294118	0.059
7	0.212121212	0.068

3 0	0.157894737	0.099
24	NA 0.328767123	0.042
0	NA	
0	NA	
0	0	
0	0	
0	0	
2	0.047619048	0.400
7	0.072164948	0.192
29 0	0.90625 0	0.001
4	0.333333333	0.046
0	0	0.040
2	0.090909091	0.198
0	0	01.00
13	0.265306122	0.055
0	0	
1	0.071428571	0.226
7	0.318181818	0.036
1	0.1	0.162
2	0.090909091	0.173
0	0	
0 0	0 0	
0	0	
1	0.058823529	0.228
0	0	0.220
1	0.028571429	
0	0	
0	0	
2	0.090909091	0.181
6	0.136363636	0.146
0	0	
0	0	
0	0	
0 0	0 0	
0	0	
3	0.375	0.024
3 5 5	0.357142857	0.027
5	0.294117647	0.033
0	NA	
0	0	
0	0	
0	0	
0	0	
0	0	
0	0	
0 0	0 0	
2	0.11111111	0.139
4	O. 111111111	0.108

	2	
0 10	0 0.454545455	0.016
0	0.43434343	0.016
8	0.533333333	0.014
5	0.166666667	0.117
0	0	0
0	0	
0	0	
0	0	
0	0	
2	0.333333333	0.032
4	0.035714286	
7	0.368421053	0.016
1	0.038461538	
0	0	
0	NA	
0	0	
0	0 0.05	
0	0	
1	0.25	0.049
0	0	0.049
3	1	0.000
0	0	0.000
0	0	
0	0	
12	0.342857143	0.027
0	NA	
0	0	
6	0.5	0.016
3	0.157894737	0.099
0	0	
0	0	
1	0.02222222	
1	0.111111111	0.166
4	0.173913043	0.076
0 4	0 0.072727273	0.217
4	0.363636364	0.217
4	0.181818182	0.014
0	0	0.007
0	0	
0	0	
1	0.058823529	0.263
0	0	
0	0	
9	0.204545455	0.073
0	0	
2	0.037735849	
8	0.421052632	0.019
0	0	
1	0.166666667	0.095

3 0	0.111111111	0.185
0 2 0	0 0.083333333	0.206
9	0 0.236842105	0.050
0 0	0 0	
0 5	0 0.084745763	0.177
0	0	
1 0	0.083333333	0.214
6	0.15	0.117
0	0 0	
0	0	
1	0.166666667	0.095
0	NA NA	
0	NA	
5	0.55555556	0.005
0	0	
0	0 0	
0	0	
0	0	
0	0	
0	0	
0	0 0	
0	0	
0	0	
0	0	
0	0	
0 2	0 0.086956522	0.186
0	0	0.100
0	0	
0	0	
2	0.142857143	0.132
0 6	0 0.12	0.163
5	0.079365079	0.194
1	0.022727273	
0	0	
0	0	
0	0 0.043478261	
2	0.105263158	0.170
24	0.285714286	0.042
61	0.512605042	0.007

11 1	0.20754717 0.05	0.067
2 3	0.066666667 0.026548673	0.216
0 4 12 0 0 0 1	0 0.142857143 0.097560976 NA NA 0 0.04	0.123 0.163
0 0 0 0 0	0 0 0 0 0 0	
0 0	0	
3 3 1	0.083333333 0.1875 0.038461538	0.184 0.077
1	0.0625	0.252
1 6	0.11111111 0.090909091	0.151 0.173
19 0	0.63333333 0	0.003
41	0.460674157	0.011
0 0	0 0	
3	0.0454545	
0	0	
1 0	0.125	0.123
2 0	0 0.181818182 0	0.100
0	0	
0	0	
0 0	0 0	
0	0	
1	0.125	0.120
1 13	0.043478261 0.541666667	0.006
0	0	0.000
0	0	
1 0	0.071428571 0	0.221
5	0.263157895	0.022
8	0.140350877	0.116
7	0.411764706	0.013
0 0	0 0	
U	U	

0	0	
0	0	
9	0.28125	0.046
3	0.090909091	0.179
0	0	
28	0.140703518	0.091
0	0	
0	0	
0	0	
0	0	
0	0	
0	0	
0 2	0 0.153846154	0.113
0	0.153646154	0.113
5	0.036231884	
4	1	0.000
0	0	0.000
2	0.166666667	0.087
0	0.10000007	0.067
0	0	
1	0.05	
0	0.05	
0	0	
1	0.043478261	
0	0.04047.0201	
0	0	
0	0	
0	0	
9	0.272727273	0.035
1	0.5	0.008
2	0.1	0.169
0	0	
1	0.027027027	
0	0	
0	0	
0	0	
2	0.125	0.130
11	0.180327869	0.070
0	0	
0	0	
0	0	
0	NA	
0	0	
0	0	
0	0	
0	0	
1_	0.038461538	
7	0.318181818	0.028
0	0	
0	0	2 22-
1	0.0625	0.225

0	0	
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0	0	
0	0	
5	0.33333333	0.019
0	0	
0	0	
6	0.461538462	0.003
0	0	
0	0	0.000
4	0.285714286	0.032
0	0	
0 0	0 0	
	0	
0 0	NA	
7	0.184210526	0.064
0	0.104210320	0.004
0	0	
2	0.142857143	0.094
2	0.285714286	0.025
0	0	0.025
0	0	
6	0.1875	0.053
0	0	0.000
0	0	
0	0	
0	0	
4	0.571428571	0.003
0	0	
1	0.043478261	
0	0	
5	0.294117647	0.024
0	0	
0	0	
0	0	
2	0.020833333	
2	0.16666667	0.070
0	0	
0	0	
0	0	
0	0	0.007
10	0.434782609	0.007
5	0.357142857	0.005
1 0	0.071428571	0.210
0	0 0	
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Chn_LRHtest_# of HapMap SNPs within region Chn_LRHtest_# of extreme SNPs 5 9 5

18 0		
53 0 0		
6 4 2		
43 76 31		
22 10 11		
24 27 52		
8 27 16		
5 19 15		
9 27 5		
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17 6 23 8 21 10 62	8 25 3 12 24 31 12 32 6 21 7 24 19 20 28 0 14 10 11 7 3 11 2 13 30 0 27 9 17		
	6 23 8 21 10		
26	7		

24 8		
15 32 39 34		
23 2 20		
37 27 11		
14 51 11 2		
12 8 0		
0 0 7		
2 7 10 5		
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34 41 18 6		
22 17 10		
19 24 17		
10 10 32 37		
60 29 5		
19 42 25		
20 69 69		

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42 14 22 96 49		
27 108 0 0 13		
23 14 12 14 22 18		
3 48 25 20 20 13		
8 60 30 31 93 16		
13 49 21 5 6		
11 9 9 5 38 15		
8 8 16 15 20 7		
10 12 14 47 17		
14 16		

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23 12 35 24 5 238 48 12 6 22 6 21 10 13 27 125 10 15 12 3 8 21		
38 14 21 7		
16 9 6 27 2 18 9		
27 31 19 16 14 37 1		
22 35 0 22 2 12 22 26		
22 26 23 3 97 12		

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35 2
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29 9 18 6		
12 9 12 15 3 8		
47 4 14 9 12 8		
4 0 26 31 58		
12 6 33 14 5 30		
19 24 33 5 8		
15 25 23 12 15 17		
78 6 4 5 16		
14 27 20 16 19 28		
6 36 19		

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0

Chn_LRHtest_proportion of extreme SNPs	Chn_LRHtest_p-value
0	
0.128205128	0.136
0.152542373	0.117
0	
0.040816327	
0	
0.5	0.015
0	
0.076923077	0.196
NA	
0.86666667	0.003
NA	
0	
0	
0.302325581	0.044
0.352941176	0.040
0.80555556	0.002
0.130434783	0.137
0	
0.053571429	0.186
0.538461538	0.009
0.0625	0.206
0	
0	
0.33333333	0.035
0.047619048	
0	
0.090909091	0.191
0.2	0.096
0.035714286	
0	
0.142857143	0.109
0.482758621	0.015
0.49122807	0.032
0.26666667	0.071
0.86666667	0.001
0.05	
0	
0.04	
0.147058824	0.100
0.078125	0.165
0.085714286	0.173
0	5 6
0	
0.130434783	0.128
0	5.120
0	
0	
0	
0.08	0.184
0	

0 NA		
0.037735849		
NA		
NA 0		
0		
0		
0.441860465	0.02	22
0.5	0.02	21
0		
0.272727273	0.05	
0.4	0.02	
0.272727273	0.05)4
0		
0.211538462	0.09	91
0	0.00	
0.037037037		
0.125	0.14	1 6
0		
0.315789474	0.03	38
0		
0		
0 0.2	0.09	7
0.2	0.08	11
0		
0.75	0.00)4
0		
0		
0.058823529	0.24	1 6
0		
0		
0 0.35483871	0.03	2.4
0.058823529	0.25	
0.030023329	0.20	וי
0		
0		
0.142857143	0.07	79
0.052631579	0.26	38
NA		
0		
0 0.357142857	0.03	22
0.337 142637	0.03 0.02	
0.575	0.02	
0		
0		
0.2	0.07	
0.0625	0.22	25

0 0.04	
0.04	
0	
0	
0	
0	
0	
0	
0	
0.142857143	0.127
0.142037143	0.127
0.157894737	0.093
0.2	0.071
0.25	0.034
NA	0.004
0.214285714	0.079
0	0.070
0	
0	
0.333333333	0.040
0.727272727	0.007
0	0.007
0	
0.238095238	0.056
0	0.000
0.033333333	
NA	
0	
0	
0	
0.058823529	0.246
0	
0	
0.625	0.003
0	
0	
0.048387097	
0.125	0.138
0	
0	
0	
0	
0	
0	
0.105263158	0.146
0.085714286	0.176
0.058823529	0.246
0.03125	
0	
0.038461538	
0.428571429	0.021

0.125 0	0.132
0.2 0.53125	0.069 0.013
0 0.176470588	0.093
0	
0.05 0	
0.037037037	
0	
0 0	
Ö	
0	
0	
0 NA	
NA	
NA	
0 0	
0	
0	
0	
0 0	
0	
0.323529412	0.024
0.341463415	0.023
0 0.333333333	0.024
0.090909091	0.193
0	
0	0.000
0.315789474 0.166666667	0.029 0.087
0	0.001
0	
0 0	
0.135135135	0.123
0.116666667	0.146
0	
0 0	
Ö	
0.28	0.029
0 0.028985507	
0.028985507	
•	

0 0	
0.136363636 0.104166667	0.107 0.137
0.020408163 0.037037037 0.074074074	0.210
NA NA 0	
0.043478261 0	
0 0.071428571	0.226
0.090909091	0.167
0.05555556	0.239
0.229166667 0	0.042
0.4	0.013
0.45	0.005
0.076923077	0.209
0.125	0.120
0.116666667 0	0.164
0.612903226	0.004
0.053763441	0.269
0 0.384615385	0.014
0.714285714	0.002
0.80952381	0.003
0	0.000
0	
0	
0	
0	
0	
0.263157895	0.037
0.2	0.072
0.125	0.142
0.375 0	0.019
0	
0	
0	
0	
0	
0.285714286	0.028
0.234042553	0.056
0.235294118	0.044
0	
0	

0.086956522	0.203
0 0	
0	
0	
0.025210084	
0.023210004	
0.25	0.043
0.333333333	0.022
0.590909091	0.003
0	0.000
0	
0	
0	
0	
0.224	0.054
0	
0.2	0.078
0	
0	
0	
0.047619048	
0.026315789	
0	
0 0	
0	
0	
0	
0.074074074	0.222
0	V
0	
0	
0.037037037	
0.032258065	
0.105263158	0.179
0	
0	
0.054054054	0.246
0	
0 0.257142857	0.022
0.257 142657 NA	0.023
0	
0	
0.25	0.036
0	0.000
0	
0.173913043	0.079
0	
0.360824742	0.016
0.166666667	0.091

0.310344828 0.222222222 0 0 0	0.025 0.061
0 0 0.0666666667 0 1 0.276595745 0.25 0	0.236 0.000 0.036 0.031
0 0 0 NA 0 0.290322581	0.028
0.120689655 0.083333333 0 0 0 0	0.130 0.188
0 0.052631579	0.267
0 0.272727273 0 0	0.023
0 0.32	0.011
0 0.166666667	0.081
0 0.058823529 0.076923077 0 0 0	0.282 0.212
0 0.074074074 0.5	0.242 0.004
0 0.052631579 0 0 0 0	0.321

Genbank mRNA identifier

BC025337

BC002781,BT007415,AF151821,X97198,U73727,X95712,U71075

X70944,BC004534,BC027708,AB007885,AF118068,AL834315,BC001858,BC007645,AK090878,AY163234,AK0;BC000268,BT007137,AK096303,AK096466,AF297866,BC038991,BC062215,AB046787,AK000040,AF093097,BC06921806,BC028102,AY358631,U21049

AK027348

AB007942

BC040516,BX640813,AL136561,AK125044,AK098192

AK093170,BC028743 BC001257,AK023212 BC053610 L16464,BC022868 BC033736 BC006983

M16967,AJ297255,M94010,AJ297254

BC009382 AB062450,Y00062

AY090784,AK000347,BC014268,BC048271,AB014563

AF130080,AK025133,AF212045

AB037810,AK093191 AK095031 AK095031

AY186742,BC017079,BC031010

BC038971

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AB084279,AB084281,AF035621,BC042486

BC009235,D16481,AY358172,AY140955,AY255611,AB051511,BC021229,AK057222,AL833892,BC009488,AK1: AY358269,AK095462,BC024243,AK057516,AF435956

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AY289212,M92439 BC053660,AK027437 AJ404611 AF279145

BC051848,BC051753,AF049227,AF290512,BC004558,BC017727,BC064425,BC010012,X87237,BC012163,AK0

BC025997,BC014602,AL136552,AF119878

AB008164,BT006951 AB002334,BC004487,AK095987,BC005341 AF130988

AK125008,BC016018,AF048731 AB067489 AB067489 AB078142,AB067505

AF369908,AF391554,AF391555,AF391556,AF391557,AF391558 X53586

X90568,X98114,X98115,BC058824,BC013396,AK096821,AK129847

L07868

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D84430,BC006502 AY090737 AL834397,BC017419,AF117230,AF130117

X13710,BC000742,L25080,BC000946,L41143,BC007546,D13811,BC050653,AF538150,L19711 AF042166,BX641085

AF146696,BC054815

AK000692,BC035971 AF488739,AK128198 BC000055,AF288537 AB037663,U48959 U94190 AJ278018,BC004871,BC007943 AK055115 AK055115

BC030705 AK128339 AK096400

AK126014

AF119815 AB002364,BC016451 AK091196,BC047918,AK130576 AK091196,BC047918 L08850 X04571

AB002311,AL117397

BC035499,BC002971,BC009454,AF275798,AJ278125

AF172849,BC064405

AB062126,X53280,BT007120,AF267862

AB007898,BC030826 BC041945 BC066278,M13982,AF043336,BC045542,BC039592,D86957 AF231124 NM_015071,AF141884 NM_015071,AF141884 AF017789,AY351676,AY255557 AB011127,AF273057

M81886

BC050375 AK127740,D49835

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AB033095,BC009381,BC013933,AF217795,AK131105

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Gene name

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MGC16664 NEK7,PTPRC

LAX,LAX,KIAA0663,KIAA0663

AF130080, NUCKS, AF212045

SIPA1L2,SIPA1L2 SLC35F3 SLC35F3

SMYD3,SMYD3,SMYD3

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VSNL1,SMC6L1,SMC6L1

ASXL2,AB084281,KIF3C,KIF3C

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LOC130951,LOC130951,SEMA4F,SEMA4F

SULT1C1,BT006951 GCC2,MGC10701,AK095987,LIMS1 EDAR

AK125008,ACMSD,CCNT2 FMNL2 FMNL2 GALNT13,GALNT13

MYO3B, MYO3B, MYO3B, MYO3B, MYO3B, MYO3B ITGA6

TTN,TTN,TTN,TTN,FLJ39502,FLJ39502

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FOXP1,BC054815

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GRIA1

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BTBD9,BTBD9 SPATS1,CDC5L,CDC5L,CDC5L,AF090903 C6orf143 COL21A1

C6orf152
C6orf167
3331107
TCBA1,TCBA1
AK125983,BC062712 BX640890,AK097903
UTRN UTRN
FNDC1,AJ586132
MGC43690,AK128656,AF451991,PHF10,PHF10,TCTE3,C6orf70,AK001369 AK056013,C6orf208 KIAA1838,KIAA1838,AY266458
DGKB DGKB
JAZF1
FLJ32110,FLJ32110 CUTL1,CUTL1,CUTL1,AK098270
C8orf14 GATA4,GATA4
KCNB2

C9orf82,PLAA,PLAA,AK002143,CCDC2,CCDC2,CCDC2,CCDC2,LRRC19,CCDC2,TEK,TEK,TEK,KIAA1045,DNAJB5,DNAJB5,AK090398,MGC41945,MGC41945,VCP,VCP,VCP,AF313486,AF313485,FANCG,FA

AK127390,AGTPBP1,AGTPBP1,AGTPBP1,AJ437018 LOC158046

INVS,INVS,TEX10

ANKRD16

PLXDC2,PLXDC2 C10orf67 MPP7,MPP7

ARHGAP22,ARHGAP22
AY388963,PCDH15
PCDH15,AL834134
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JDP1,SIRT1
DDIT4,AK123129,DNAJB12,CBARA1,CBARA1,CBARA1,CBARA1
CBARA1,CBARA1,CBARA1,CBARA1,C10orf42,C10orf42
C10orf11
MGC4248,MGC4248,MGC4248
SEC15L1,SEC15L1,SEC15L1,CYP26C1,CYP26C1,AK127234,CYP26A1
FER1L3,FER1L3,FER1L3

WT1

FADS2,VMD2,VMD2,FTH1,BC066341,AY033611,AK093779

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DACH, DACH DACH, DACH

SLITRK5

GPC6

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CFL2,CFL2

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RAI1,RAI1,SMCR5,RAI1,RAI1,RAI1,RAI1,RAI1,SREBF1,SREBF1,SREBF1,SREBF1,TOM1L2,AY358769,TOM1L: HCMOGT-1,HCMOGT-1,HCMOGT-1,HCMOGT-1,AF130065
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PTPRM

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GMEB2,STMN3,BC013933,C20orf41,TNFRSF6B

MRPL39, MRPL39

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SERHL,SERHL,CGI-96,CGI-96,CGI-96,BC063537,CGI-96,dJ222E13.1,SERHL,dJ222E13.2,POLDIP3,POLDIP3,FPACSIN2,

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¹AF2,AY203943,MGC3048,MGC3048,DRG2,DRG2,MYO15A

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