

		children whites										
		COHRA (N=608)					IHS (N=41)			IFS (N=136)		
gene	SNP	chromosome	position	beta	SE	p-value	beta	SE	p-value	beta	SE	p-value
AMBN	rs17149026	4	71440940	-0.2575	0.3198	0.4211	0.7944	3.98	0.8429	-0.5944	0.5199	0.255
	rs17733915	4	71447093	0.3792	0.1831	0.03877	1.531	1.089	0.1681	-0.01916	0.2758	0.9447
	rs7439186	4	71469604	0.1557	0.3516	0.6581	NA	NA	NA	0.9038	0.5743	0.1179
ENAM	rs1967376	4	71501744	-0.2772	0.481	0.5646	-0.2508	2.983	0.9334	-0.3243	0.6962	0.6421
	rs12640848	4	71506412	0.03201	0.1989	0.8722	-1.536	1.318	0.2513	0.1473	0.2924	0.6152
TFIP11	rs17402286	22	26894879	-0.1145	0.4388	0.7942	-2.703	4.138	0.5176	0.8311	0.5863	0.1587
	rs6005060	22	26895736	-0.7235	0.3152	0.02204	1.745	2.493	0.4884	-0.3488	0.4324	0.4213
	rs713900	22	26898242	0.2828	0.3235	0.3823	-2.505	2.591	0.3399	-0.4605	0.6775	0.4979
	rs134134	22	26898891	0.7205	0.2728	0.008467	-1.143	1.74	0.5155	0.01542	0.4826	0.9746
	rs134135	22	26898962	0.7549	0.2572	0.003	-0.3296	1.85	0.8596	-0.05748	0.4569	0.9001
	rs2097470	22	26904965	-0.01612	0.2643	0.9514	0.9028	1.799	0.6187	-0.2738	0.4658	0.5577
	rs134145	22	26909750	-0.4084	0.1856	0.02817	0.4312	1.256	0.7333	-0.3864	0.2816	0.1723
TUFT1	rs2337359	1	151495796	-0.2623	0.2496	0.2936	0.5034	1.709	0.77	0.2465	0.3613	0.4963
	rs1045298	1	151510825	-0.07064	0.2633	0.7885	-1.283	1.774	0.4741	0.5529	0.4247	0.1952
	rs10158855	1	151515654	0.0002306	0.1805	0.999	0.9176	1.334	0.4957	-0.2914	0.292	0.3202
	rs17640579	1	151521933	0.09479	0.205	0.644	0.02316	1.625	0.9887	0.195	0.3094	0.5296
	rs16833391	1	151547253	-0.003429	0.3197	0.9914	1.476	2.322	0.529	-0.1521	0.493	0.7581
	rs12749	1	151555741	-0.2763	0.2391	0.2484	-2.215	1.887	0.248	-0.1394	0.422	0.7417

gene	SNP	blacks			IHS (N=23)			all children		adults whites		
		Meta (N=785)	COHRA (N=81)					Meta (N=889)		COHRA (N=994)		
		p-value	beta	SE	p-value	beta	SE	p-value	p-value	beta	SE	p-value
AMBN	rs17149026	0.2557	-1.105	0.5134	0.03458	2.37	2.142	0.284	0.1251	-0.2639	0.5207	0.6124
	rs17733915	0.0354	0.7376	0.4929	0.1388	-1.836	2.731	0.5104	0.02044	0.1407	0.2915	0.6294
	rs7439186	0.2849	0.5933	0.5125	0.2507	-0.7093	2.334	0.7649	0.191	0.02106	0.579	0.971
ENAM	rs1967376	0.4717	-0.1703	0.4321	0.6945	0.5522	1.482	0.7141	0.4618	0.6368	0.7114	0.371
	rs12640848	0.9295	-0.4748	0.5699	0.4074	-12.77	5.309	0.02783	0.6018	-0.3537	0.3106	0.2551
TFIP11	rs17402286	0.8338	3.052	1.213	0.01405	-2.163	5.33	0.69	0.3815	-0.03027	0.7207	0.9665
	rs6005060	0.02842	-0.6081	0.4349	0.1661	-1.544	2.482	0.5421	0.01001	-0.1716	0.5094	0.7363
	rs713900	0.7896	0.714	1.127	0.5283	NA	NA	NA	0.6547	0.2158	0.5441	0.6917
	rs134134	0.02913	-0.06444	0.4155	0.8772	0.6345	1.726	0.7176	0.03924	0.1334	0.4514	0.7676
	rs134135	0.01308	0.03342	0.3895	0.9319	-0.1206	1.651	0.9426	0.019	0.2136	0.4049	0.5979
	rs2097470	0.8539	1.14	0.975	0.2461	-6.263	5.212	0.246	0.9887	-0.4447	0.4372	0.3093
	rs134145	0.01545	-0.0331	0.3804	0.9309	-0.8127	1.659	0.6306	0.01737	-0.07848	0.3009	0.7943
	rs2337359	0.1392	0.04209	0.4189	0.9202	-1.082	3.081	0.7299	0.2026	-0.1572	0.4027	0.6964
TUFT1	rs1045298	0.2279	1.353	1.516	0.3749	-9.557	5.809	0.1183	0.3478	0.06459	0.3935	0.8697
	rs10158855	0.7971	0.307	0.4081	0.4542	-1.44	1.931	0.4662	0.8943	0.1188	0.2946	0.6869
	rs17640579	0.5021	0.6475	0.6885	0.35	-1.495	3.487	0.6739	0.3969	0.01757	0.3378	0.9585
	rs16833391	0.995	0.1503	0.5984	0.8023	2.78	2.939	0.3575	0.8185	-0.0101	0.4591	0.9825
	rs12749	0.1564	0.02552	0.4101	0.9505	-2.884	1.622	0.09329	0.1134	0.2667	0.3688	0.4697

		Dental SCORE (N=277)			DRDR (N=702)			CEDAR (N=173)			Meta (N=2,146)
gene	SNP	beta	SE	p-value	beta	SE	p-value	beta	SE	p-value	p-value
AMBN	rs17149026	1.346	0.89	0.1315	-0.2631	0.7752	0.7344	-0.3235	1.119	0.7729	0.9371
	rs17733915	-0.1228	0.4626	0.7909	-0.67	0.3929	0.08857	-0.1934	0.4814	0.6884	0.3931
	rs7439186	-0.7887	0.8542	0.3567	-0.17	0.7427	0.8191	1.182	1.192	0.3229	0.8756
ENAM	rs1967376	0.5442	1.042	0.6019	-0.5996	0.8686	0.4902	3.588	1.436	0.01344	0.2698
	rs12640848	-0.8397	0.4627	0.07061	-0.8115	0.4149	0.05086	0.66	0.5131	0.2001	0.02948
TFIP11	rs17402286	0.168	1.229	0.8914	-0.01529	1.006	0.9879	-0.8841	1.196	0.4609	0.8434
	rs6005060	0.1846	0.763	0.809	0.1369	0.687	0.842	-0.2524	0.8649	0.7708	0.9116
	rs713900	-1.038	0.8875	0.2434	-1.223	0.8702	0.1602	0.1945	1.153	0.8663	0.3654
	rs134134	-0.6246	0.726	0.3903	0.4272	0.7168	0.5514	0.1036	0.8036	0.8976	0.7875
	rs134135	-0.7697	0.6634	0.2469	-0.4313	0.6486	0.5063	-0.3524	0.8015	0.6607	0.5743
	rs2097470	0.2652	0.6578	0.6871	0.4208	0.578	0.4668	1.722	0.6799	0.01225	0.5642
	rs134145	0.1328	0.4507	0.7685	-0.3867	0.418	0.3552	0.4329	0.5078	0.3951	0.7207
TUFT1	rs2337359	-0.4823	0.5524	0.3833	0.7187	0.5223	0.1693	2.126	0.6789	0.00205	0.0251
	rs1045298	0.4418	0.6005	0.4626	-0.2548	0.5401	0.6373	-0.5268	0.7174	0.4638	0.3935
	rs10158855	0.04054	0.451	0.9284	0.8299	0.3963	0.0366	-0.3023	0.4997	0.546	0.1833
	rs17640579	0.59	0.5346	0.2708	-0.3696	0.4498	0.4116	0.01667	0.5237	0.9746	0.9772
	rs16833391	0.2909	0.8206	0.7233	-0.2636	0.7843	0.737	0.6255	0.9546	0.5131	0.9157
	rs12749	0.4469	0.6292	0.4782	0.1425	0.5192	0.7837	-0.2265	0.6595	0.7317	0.42

		blacks										
		COHRA (N=86)			Dental SCORE (N=225)			DRDR (N=173)			CEDAR (N=68)	
gene	SNP	beta	SE	p-value	beta	SE	p-value	beta	SE	p-value	beta	SE
AMBN	rs17149026	-0.1213	1.287	0.9252	-0.3748	0.6728	0.5781	0.5683	1.003	0.5718	-0.1276	0.9733
	rs17733915	-0.1218	1.246	0.9223	-0.03969	0.6025	0.9475	-0.3394	0.8545	0.6918	0.3678	0.9164
	rs7439186	-0.4931	1.23	0.6897	-0.2626	0.7068	0.7106	0.6878	1.149	0.5502	1.051	1.091
ENAM	rs1967376	0.01998	1.073	0.9852	-0.4254	0.5101	0.4052	0.4444	0.7263	0.5414	1.112	0.7634
	rs12640848	-0.4908	1.297	0.7062	0.4459	0.6959	0.5223	-0.4705	1.099	0.669	1.374	1.259
TFIP11	rs17402286	-2.793	3.755	0.4592	-4.572	2.754	0.09826	0.2702	3.119	0.9311	1.04	4.321
	rs6005060	-0.9236	1.524	0.5462	-1.12	0.7476	0.1357	-0.1171	0.9661	0.9037	-0.7022	0.9542
	rs713900	4.478	3.697	0.2294	0.984	1.626	0.5458	-2.909	3.228	0.3688	0.4523	4.471
	rs134134	-0.01515	1.135	0.9894	0.2372	0.5367	0.659	-0.431	0.8619	0.6177	-1.217	0.9216
	rs134135	-0.5615	1.034	0.5885	0.5078	0.5328	0.3416	-0.5073	0.8085	0.5312	-0.05391	0.8066
	rs2097470	-1.208	2.295	0.6003	-0.04539	1.277	0.9717	-2.267	2.125	0.2877	-0.6754	2.095
	rs134145	-0.138	1.209	0.9094	-1.338	0.5848	0.02305	-0.2611	0.7517	0.7288	-0.2288	0.8767
	rs2337359	-0.147	0.958	0.8784	0.8818	0.5104	0.08546	0.855	0.7238	0.2392	-0.7691	0.8053
TUFT1	rs1045298	-2.301	3.24	0.4798	0.1038	1.406	0.9412	0.4091	2.009	0.8389	-3.374	2.243
	rs10158855	0.4046	1.021	0.6931	-0.4502	0.518	0.3857	0.791	0.7524	0.2947	0.6085	0.8418
	rs17640579	-5.091	2.302	0.02982	0.2599	0.8352	0.756	0.1394	1.245	0.911	0.3638	1.483
	rs16833391	-0.7912	1.271	0.5354	-0.5567	0.755	0.4617	0.4528	1.09	0.6783	-0.04215	1.072
	rs12749	-0.8506	1.056	0.4232	-0.08519	0.5808	0.8835	-0.5297	0.7737	0.4945	0.9539	0.8365

		all adults		
		Meta (N=552)	Meta (N=2,698)	
gene	SNP	p-value	p-value	p-value
AMBN	rs17149026	0.8961	0.9033	0.9003
	rs17733915	0.6896	0.8714	0.4037
	rs7439186	0.3389	0.7828	0.9881
ENAM	rs1967376	0.1503	0.7469	0.2584
	rs12640848	0.2792	0.6874	0.07837
TFIP11	rs17402286	0.8106	0.2243	0.4679
	rs6005060	0.4646	0.1297	0.4328
	rs713900	0.9198	0.6952	0.5287
	rs134134	0.1914	0.6445	0.9748
	rs134135	0.9469	0.9854	0.6222
	rs2097470	0.7483	0.3482	0.9294
	rs134145	0.7949	0.07471	0.2605
	rs2337359	0.3433	0.0317	0.0029
TUFT1	rs1045298	0.1376	0.3365	0.2322
	rs10158855	0.4725	0.6594	0.1657
	rs17640579	0.807	0.6095	0.7975
	rs16833391	0.9688	0.6198	0.8966
	rs12749	0.2585	0.6924	0.589