

Online Appendix. The explicit values of the HR (95%CI) for type 1 diabetes in the HLA stratified analyses presented in Figure 1 in:

Complement gene variants in relation to autoantibodies to beta cell specific antigens and type 1 diabetes in the TEDDY Study

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HLA	SNP	ProbChisq	HazardRatio	HRLowerCL	HRUpperCL
DR3/3	rs1048971_A	0.7345	0.856	0.348	2.106
DR3/3	rs10818488_A	0.9489	1.023	0.516	2.025
DR3/3	rs1143678_A	0.9724	0.983	0.370	2.611
DR3/3	rs1143679_A	0.9110	0.936	0.292	2.999
DR3/3	rs1143683_A	0.9525	0.971	0.364	2.587
DR3/3	rs17615_A	0.8670	0.930	0.400	2.166
DR3/3	rs2230199_C	0.1924	0.477	0.156	1.452
DR3/3	rs2857009_C	0.7172	0.679	0.083	5.521
DR3/3	rs4308977_G	0.8792	0.937	0.403	2.176
DR3/3	rs4597342_A	0.6244	1.170	0.625	2.190
DR3/3	rs4844573_G	0.1146	1.866	0.860	4.050
DR3/3	rs665691_C	0.8336	0.918	0.413	2.040
DR3/3	rs7029078_G	0.6136	1.199	0.593	2.427
DR3/3	rs7951_A	0.4930	0.620	0.158	2.435
DR3/3	rs9332739_C	<.0001	0.000	0.000	0.000
DR3/4	rs1048971_A	0.1684	1.283	0.900	1.828
DR3/4	rs10818488_A	0.5967	1.089	0.794	1.494
DR3/4	rs1143678_A	0.3863	0.796	0.474	1.335
DR3/4	rs1143679_A	0.1495	0.640	0.349	1.174
DR3/4	rs1143683_A	0.3837	0.795	0.474	1.333
DR3/4	rs17615_A	0.0252	1.524	1.054	2.204
DR3/4	rs2230199_C	0.3396	1.228	0.806	1.871
DR3/4	rs2857009_C	0.5509	1.150	0.726	1.820
DR3/4	rs4308977_G	0.0588	1.425	0.987	2.058
DR3/4	rs4597342_A	0.7997	1.047	0.735	1.491
DR3/4	rs4844573_G	0.0169	0.628	0.429	0.920
DR3/4	rs665691_C	0.3754	1.173	0.824	1.668
DR3/4	rs7029078_G	0.6646	1.082	0.757	1.548
DR3/4	rs7951_A	0.9486	0.979	0.518	1.851
DR3/4	rs9332739_C	0.6070	1.203	0.595	2.433
DR4/4	rs1048971_A	0.9738	1.011	0.541	1.886
DR4/4	rs10818488_A	0.0733	1.675	0.952	2.945
DR4/4	rs1143678_A	0.8365	0.925	0.442	1.935
DR4/4	rs1143679_A	0.3145	0.571	0.192	1.702
DR4/4	rs1143683_A	0.8341	0.924	0.442	1.933
DR4/4	rs17615_A	0.5713	0.812	0.394	1.671
DR4/4	rs2230199_C	0.0002	3.200	1.750	5.851
DR4/4	rs2857009_C	0.9028	0.957	0.472	1.942
DR4/4	rs4308977_G	0.8859	0.952	0.489	1.856
DR4/4	rs4597342_A	0.9653	0.986	0.512	1.898
DR4/4	rs4844573_G	0.5362	1.216	0.654	2.262
DR4/4	rs665691_C	0.8878	1.041	0.596	1.820
DR4/4	rs7029078_G	0.3024	1.389	0.744	2.593
DR4/4	rs7951_A	0.8911	0.922	0.286	2.966
DR4/4	rs9332739_C	0.6760	1.190	0.526	2.691
DR4/8	rs1048971_A	0.9820	1.009	0.455	2.236
DR4/8	rs10818488_A	0.8802	0.939	0.417	2.118
DR4/8	rs1143678_A	0.7783	0.871	0.334	2.273
DR4/8	rs1143679_A	0.6221	0.756	0.249	2.297

DR4/8	rs1143683_A	0.7716	0.867	0.332	2.266
DR4/8	rs17615_A	0.9519	0.975	0.428	2.219
DR4/8	rs2230199_C	0.9470	0.972	0.419	2.256
DR4/8	rs2857009_C	0.7339	0.896	0.477	1.684
DR4/8	rs4308977_G	0.9823	0.991	0.435	2.255
DR4/8	rs4597342_A	0.9602	0.982	0.484	1.995
DR4/8	rs4844573_G	0.1782	0.670	0.373	1.201
DR4/8	rs665691_C	0.4927	1.151	0.770	1.721
DR4/8	rs7029078_G	0.4821	0.762	0.356	1.628
DR4/8	rs7951_A	0.8030	0.859	0.259	2.845
DR4/8	rs9332739_C	0.5119	0.519	0.073	3.680