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Supplemental Data

Large-Scale Gene-Centric Meta-analysis

across 32 Studies Identifies Multiple Lipid Loci

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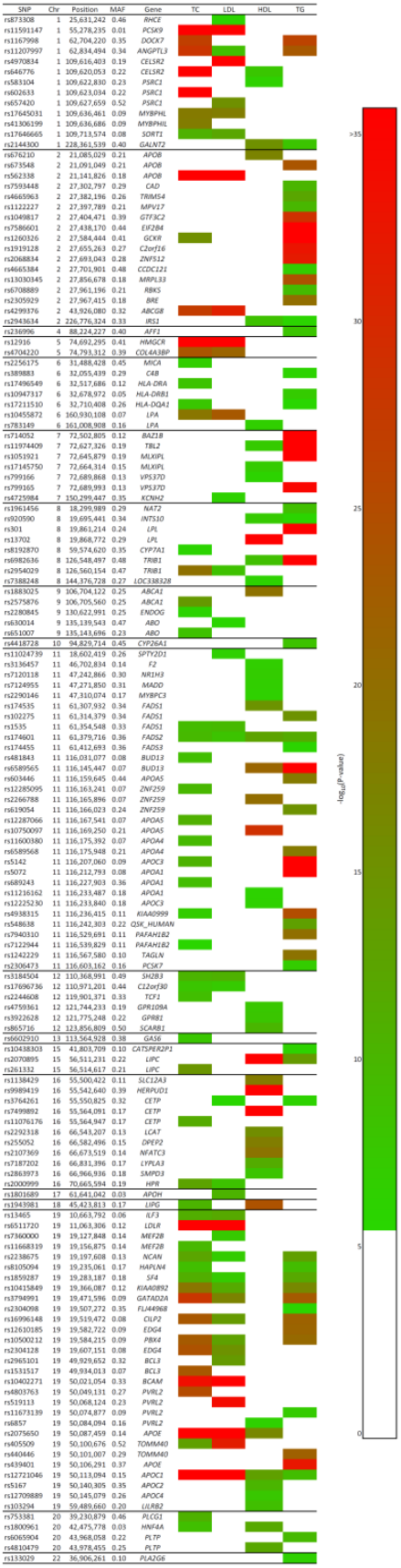

Figure S1. **Heat Map Summarizing the Signals Identified from the International IBC Lipid Meta-analysis and Their Relationship with All Four, HDL-C, LDL-C, TC, and TG Phenotypes Tested**

Only SNPs significant at the array wide p value threshold of 2.4×10^{-6} and filtered for heterogeneity <35% are shown.