

SUPPLEMENTARY DATA

SUPPLEMENTARY TABLE S1. DOMINANT, RECESSIVE, AND ADDITIVE MODELS FOR THE FAMILY-BASED ASSOCIATION ANALYSIS OF ASTHMA AND RELATED TRAITS WITH *SPP1* POLYMORPHISMS IN PUERTO RICAN ASTHMA TRIOS

<i>Phenotype</i>	<i>Model</i>	<i>rs4660</i>	<i>rs6812524</i>	<i>rs7435825</i>	<i>rs1126772</i>	<i>rs1126616</i>
Asthma	Additive	0.42	0.52	0.47	0.21	0.003
	Dominant	0.46	0.5	0.51	0.04	0.000065
	Recessive	—	—	—	0.19	0.66
Moderate-severe asthma	Additive	0.43	0.56	0.41	0.75	0.01
	Dominant	0.48	0.54	0.54	0.34	0.00037
	Recessive	—	—	—	0.34	0.16
Asthma with elevated IgE (>100 IU/mL)	Additive	0.83	0.78	0.52	0.08	0.002
	Dominant	0.75	0.82	0.73	0.02	0.000084
	Recessive	—	—	—	0.498	0.89
Post-BD FEV ₁	Additive	0.98	0.44	0.33	0.37	0.10
	Dominant	0.46	0.29	0.32	0.07	0.02
	Recessive	—	—	—	0.21	0.55
Log ₁₀ IgE level	Additive	0.29	0.38	0.69	0.24	0.26
	Dominant	0.28	0.41	0.72	0.14	0.18
	Recessive	—	—	—	0.66	0.98

P values <0.05 are in bold. For 3 SNPs, the minor allele frequency was sufficiently rare that the recessive model had insufficient informative families to perform the analysis.

IgE, immunoglobulin E; Post-BD, post-bronchodilator lung function; SNP, single-nucleotide polymorphism.