

# Genetic Variation Underpinning ADHD Risk in a Caribbean Community

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## Supplementary Material:

**Table S1.** Single nucleotide polymorphisms (SNPs) genotyped in 386 individuals belonging to 113 nuclear families from Barranquilla, Colombia.

Marker	Alleles	Chr	Position	Closest gene(s)	QC result	Observations
rs143020947	A/G	2	143,643,356	<i>KYNU</i>	NO PASS	Monomorphic
rs207462497	G/T	2	143,689,722	<i>KYNU</i>	NO PASS	Monomorphic
rs1565902	C/T	4	62,091,215	<i>ADGRL3</i>	PASS	-
rs10001410	A/C	4	62,156,824	<i>ADGRL3</i>	PASS	-
rs2122642	G/A	4	62,380,859	<i>ADGRL3</i>	PASS	-
rs6551660	C/G	4	62,390,745	<i>ADGRL3</i>	NO PASS	HW criterion
rs207464763	A/G	4	62,416,514	<i>ADGRL3</i>	NO PASS	Monomorphic
rs1510921	C/T	4	62,578,187	<i>ADGRL3</i>	NO PASS	HW criterion
rs6551678	A/G	4	62,705,646	<i>ADGRL3</i>	NO PASS	HW criterion
rs207464765	A/T	4	62,720,720	<i>ADGRL3</i>	NO PASS	Monomorphic
rs748519607	A/T	4	63,888,775	<i>ADGRL3</i>	NO PASS	HW criterion
rs28363183	G/A	5	1,444,398	<i>SCL6A3</i>	NO PASS	Monomorphic
rs2282794	A/G	5	141,961,893	<i>FGF1</i>	PASS	-
rs761010945	A/C	9	136,499,907	<i>DBH</i>	NO PASS	Monomorphic
rs569034387	C/G	9	136,500,791	<i>DBH</i>	NO PASS	Monomorphic
rs199926239	A/T	9	136,512,962	<i>DBH</i>	NO PASS	Monomorphic
rs916457	C/T	11	627,014	<i>DRD4</i>	PASS	-
rs1800443	G/T	11	629,830	<i>DRD4</i>	NO PASS	HW criterion
rs2236225	G/A	14	63,978,598	<i>MTHFD1</i>	NO PASS	Monomorphic
rs11568324	C/T	16	54,283,559	<i>SLC6A2</i>	NO PASS	MAF criterion
rs371002154	A/C	16	55,726,127	<i>SLC6A2</i>	NO PASS	Monomorphic
rs79777590	C/T	17	4,298,478	<i>UBE2G1/SPNS3</i>	NO PASS	Monomorphic
rs565399294	A/G	17	4,395,222	<i>SPNS3/SPNS2</i>	NO PASS	Monomorphic
rs2228130	C/T	17	7,345,715	<i>POLR2A</i>	NO PASS	MAF criterion
rs6108461	A/G	20	10,215,270	<i>SNAP25</i>	NO PASS	HW criterion
rs362990	A/T	20	10,224,221	<i>SNAP25</i>	PASS	-

<sup>a</sup> UCSC GRCh37/hg19 coordinates. Chr: Chromosome; QC: Quality control; HW: Hardy-Weinberg. Markers highlighted in blue passed quality control filters and were further included for analyses.