

Supplementary Table 1: Probe sequences for SNPs at the 21q22 CHD risk locus

SNP - Allele	Sequence
rs9982601 – C	CACAGGGCTGCT <u>C</u> CATGGCCTTGGA
rs9982601 – C	TCCAAGGCCATGGAGCAGCCCTGTG
rs9982601 – T	CACAGGGCTGCTT <u>C</u> CATGGCCTTGGA
rs9982601 – T	TCCAAGGCCATGAAGCAGCCCTGTG
rs28451064 – G	CCAGGCCAAAGTGGACACCAAATAC
rs28451064 – G	GTATTTGGTGTCCACTTTGGCCTGG
rs28451064 – A	CCAGGCCAAAGTAGACACCAAATAC
rs28451064 – A	GTATTTGGTGTCTACTTTGGCCTGG
rs9980618 – C	AGGGTGTCTGCTCAGCACACCATG
rs9980618 – C	CATGGTGTGCTGGAGCAGACACCCT
rs9980618 – T	AGGGTGTCTGCTCAGCACACCATG
rs9980618 – T	CATGGTGTGCTGAAGCAGACACCCT
rs60687299 – T	CACTGTATTGAAT <u>A</u> CTGGAGGCAAC
rs60687299 – T	GTTGCCTCCAGT <u>A</u> TTCAATACAGTG
rs60687299 – C	CACTGTATTGAAC <u>A</u> CTGGAGGCAAC
rs60687299 – C	GTTGCCTCCAGT <u>G</u> TTCAATACAGTG
rs9977419 – T	TGTGATAGTGAGT <u>G</u> AGTTCTTACGA
rs9977419 – T	TCGTAAGAACTC <u>A</u> CTCACTATCACA
rs9977419 – A	TGTGATAGTGAG <u>A</u> GAGTTCTTACGA
rs9977419 – A	TCGTAAGAACTC <u>T</u> CTCACTATCACA
rs9977093 – G	CCATGCAGAACTGTGAATCAATTAA
rs9977093 – G	TTAATTGATTCA <u>C</u> AGTTCTGCATGG
rs9977093 – A	CCATGCAGAACTATGAATCAATTAA
rs9977093 – A	TTAATTGATTCA <u>T</u> AGTTCTGCATGG

The SNP position is underlined. EMSA=electrophoretic mobility shift assay.