

SUPPLEMENTARY DATA

Supplementary Table S1. Allele frequency

Locus	Allele	Count	Proportion	95% CI
<i>VEGFA / rs699947</i>	A	269	0.470	(0.429–0.512)
	C	303	0.530	(0.488–0.571)
<i>VEGFA / rs833061</i>	C	272	0.476	(0.434–0.517)
	T	300	0.524	(0.483–0.566)
<i>VEGFR1 / rs9554320</i>	A	228	0.399	(0.358–0.440)
	C	344	0.601	(0.560–0.642)
<i>VEGFR1 / rs9582036</i>	C	165	0.288	(0.252–0.327)
	A	407	0.712	(0.673–0.748)
<i>VEGFR2 / rs2071559</i>	T	312	0.545	(0.504–0.587)
	C	260	0.455	(0.413–0.496)
<i>VEGFR2 / rs1870377</i>	A	130	0.227	(0.194–0.264)
	T	442	0.773	(0.736–0.806)
<i>VEGFR3 / rs6877011</i>	C	520	0.909	(0.882–0.931)
	G	52	0.091	(0.069–0.118)
<i>LOXL2 / rs4872122</i>	A	232	0.406	(0.365–0.447)
	C	340	0.594	(0.553–0.635)
<i>eNOS / rs2070744</i>	T	365	0.638	(0.597–0.678)
	C	207	0.362	(0.322–0.403)
<i>SH3GL2 / rs10963287</i>	C	429	0.750	(0.712–0.785)

T	143	0.250	(0.215–0.288)	
<i>CCDC26</i> / <i>rs60315789</i>	–	267	0.467	(0.425–0.509)
TAT	305	0.533	(0.491–0.575)	

Abbreviations: CI, confidence interval; SNP, single nucleotide polymorphism; VEGFR, VEGF receptor.