Supplementary Table II **Association of the *IL1A* variants in the European imputed data, conditioned on rs6542095.**

Chr	SNP	Pos (bp)	RA	OA	QIMRHCS + OX ('All Endo')		QIMRHCS + OX ('Grade_B')	
					OR (95% CI)	*P*		*P*
2	rs11677416	113 529 240	T	C	1.05 (0.97–1.13)	2.05E−01	1.08 (0.97–1.2)	1.55E−01
2	rs2856836	113 532 083	A	G	1.05 (0.97–1.13)	2.04E−01	1.08 (0.97–1.2)	1.55E−01
2	rs1304037	113 532 236	T	C	1.05 (0.97–1.13)	2.04E−01	1.08 (0.97–1.2)	1.48E−01
2	rs17561	113 537 223	C	A	1.05 (0.97–1.13)	2.02E−01	1.08 (0.97–1.2)	1.55E−01

Chr, chromosome; Position, chromosomal position (bp) based on Human Build 37 (GRCh37/hg19); RA, risk allele; OA, other allele; OR, odds ratio; CI, confidence interval.