Table S1. The genetic association case-control studies and meta-analyses focused on polymorphisms in interleukin-1 (*IL-1*), *IL-6*, *IL-10*, *IL-17A*, and *IL-18* in patients with an aggressive form of periodontitis (AgP).

Gene	Polymorphism (rs number)	*N* (AgP/HC)	Form of AgP	Population	Association with AgP	Reference
IL-1A	–889 C/T (rs1800587)	91/210	Generalized	Czech	NS	Current study
		41+19/128	Generalized+localized	Algerian	T allele and TT genotype (risk)	[1]
		20/20	Generalized	Indian	TT genotype (risk)	[2]
		1266/2134 (meta-analysis including 19 studies)	Unknown	Mixed	NS	[3]
		80/80	Generalized	Iranian	NS	[4]
		55/41	Generalized	Brazilian	NS	[5]
IL-1B	+3953 (+3954) C/T (rs1143634)	91/210	Generalized	Czech	NS	Current study
		229/308 (meta-analysis including 3 studies)	Unknown	Mixed	NS	[6]
		43/50	Unknown	Brazilian	T allele in haplotype with *IL-1RN**S allele (risk)	[7]
		41+19/128	Generalized+localized	Algerian	T allele (risk)	[1]
		1594/2483 (meta-analysis including 25 studies)	Unknown	Mixed	NS	[8]
		26/26	Generalized	Iranian	TT genotype (risk)	[9]
		54/101	Unknown	Indian	NS	[10]
		28/33	Generalized	German	NS	[11]
IL-1RN	intron 2, 86 bp VNTR (rs2234663)	91/210	Generalized	Czech	NS	Current study
		43/50	Unknown	Brazilian	S allele, SS genotype, and haplotype with *IL-1B* +3953T allele (risk)	[7]
		577/904 (meta-analysis including 10 studies)	Unknown	Mixed (Caucasians 234/272)	NS (LS genotype protective for AgP in Caucasians)	[12]
		46/90	Generalized	Greek	NS	[13]
IL-6	–174 C/G (rs1800795)	91/210	Generalized	Czech	NS	Current study
		53/38	Generalized	Turkish	G allele and GG genotype (risk)	[14]
		53/50	Generalized	Turkish	G allele and GG genotype (risk)	[15]

		9+3/6	Generalized+localized	UK (white)	Haplotype was associated with inflammantory response in AgP patients	[16]
		122/246	Generalized	Italian	NS	[17]
		35/85	Generalized	Turkish	NS	[18]
		981/497 (meta-analysis including 6 studies)	Unknown (included also patients with CP)	Mixed (Caucasian 806/423)	G allele (risk)	[19]
		167 +57/231	Generalized+localized	UK (mixed)	NS	[20]
		91/210	Generalized	Czech	GG genotype (risk) GCC (rs1800896/ rs1800871/ rs1800872) haplotype (risk)	Current study
		1557/1447 (meta-analysis including 16 studies)	Unknown (included also patients with CP)	Mixed (non-Asian)	NS	[21]
		50/61	Generalized	Brazilian	NS	[22]
		85/86	Localized	Jordanian	NS	[23]
IL-10	–1087 (*–1082)* G/A (rs1800896)	197/502 (meta-analysis including 7 studies)	Unknown	Mixed (Caucasian 132/376)	NS	[24]
		122/246	Generalized	Italian	NS	[17]
		35/85	Generalized	Turkish	NS	[18]
		65/126	Generalized	Taiwan	Haplotype ATA/ATA (risk)	[25]
		55/43	Unknown	Brazilian	NS	[26]
		32/34	Generalized	German	Allele A (risk)	[27]
		52/61	Generalized	Iranian	NS	[28]
		51/100	Unknown	UK (white)	NS	[29]
		91/210	Generalized	Czech	GCC (rs1800896/ rs1800871/ rs1800872) haplotype (risk)	Current study
		1557/1447 (meta-analysis including 16 studies)	Unknown (included also patients with CP)	Mixed (non-Asian)	NS	[21]
	–829 (*–824)* C/T (rs1800871)	50/61	Generalized	Brazilian	NS	[22]
		97/269 (meta-analysis including 4 studies)	Unknown	Mixed (Caucasian 32/143)	NS	[24]
		35/85	Generalized	Turkish	NS	[18]
		65/126	Generalized	Taiwan	Haplotype ATA/ATA (risk)	[25]
		32/34	Generalized	German	Allele T (risk)	[27]

		6+12/21	Generalized+localized	German	NS	[30]
	–597 (*–592)* C/A (rs1800872)	91/210	Generalized	Czech	GCC (rs1800896/ rs1800871/ rs1800872) haplotype (risk)	Current study
		1557/1447 (meta-analysis including 16 studies)	Unknown (included also patients with CP)	Mixed	NS	[21]
		288/399 (meta-analysis including 6 studies)	Unknown	Mixed	Allele A and genotype AA (risk)	[31]
		53/38	Generalized	Turkish	Genotype AA	[14]
		53/50	Generalized	Turkish	Genotype AA risk	[15]
		85/86	Localized	Jordanian	NS	[23]
		97/442 (meta-analysis including 5 studies)	Unknown	Mixed (Caucasian 32/316)	NS	[24]
		122/246	Generalized	Italian	NS	[17]
		35/85	Generalized	Turkish	NS	[18]
		65/126	Generalized	Taiwan	Haplotype ATA/ATA (risk)	[25]
		32/34	Generalized	German	Allele A of haplotype ATA/ATA – present only in GAgP group	[27]
		6+12/21	Generalized+localized	German	NS	[30]
IL-17A	–197 A/G (rs2275913)	91/210	Generalized	Czech	NS	Current study
		188/655 (meta-analysis including 5 studies)	Unknown	Mixed	NS	[32]
		35/35	Localized	Indian	A allele and AA+AG genotypes risk	[33]
		45/72	Generalized+ localized	Brazilian	G allele and GG genotype associated with AgP in non-smokers	[34]
IL-18	–607 A/C (rs1946518)	91/210	Generalized	Czech	Haplotype AGC (rs1946518/rs187238/rs4988359) (risk in dominant model)	Current study
		576/458 (meta-analysis including 5 studies)	Unknown (included also patients with CP)	Mixed	Only 1 study with AgP patients was included.	[35]
		109/100	Generalized	Italian	Genotype AA with CG genotype of -137	[36]

				(rs187238) associated with AgP	
	122/246	Generalized	Italian	NS	[17]
	111/80	Generalized	German	NS	[37]
	123/121	Unknown	German	NS	[38]
	91/210	Generalized	Czech	Haplotype AGC (rs1946518/rs187238/rs 4988359) (risk in dominant model)	Current study
–137 C/G (rs187238)	576/458 (meta-analysis including 5 studies)	Unknown (included also patients with CP)	Mixed	Only 1 study with AgP patients was included.	[35]
	109/100	Generalized	Italian	Genotype CG with AA genotype of -607 (rs1946518) associated with AgP	[36]
	111/80	Generalized	German	NS	[37]
	123/121	Unknown	German	NS	[38]

AgP, aggressive periodontitis; CP, chronic periodontitis; HC, healthy control; *N*, number; NS, nonsignificant; for references, see the end of this file.

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