

Table S1 Primers used for PCR amplification and PCR conditions

Exon		Primer (5'-3')	Annealing T (°C)
2	Forward	GGACAGCCCCAGTAGTTAGTA	60°C
	Reverse	AAAGGAAAAAGCCTCAGGTGG	
3	Forward	AAGGTCTAACGCCCTCAGCT	60°C
	Reverse	CAGTCCACGAAGGATCTG	

Table S2. Summary of human PAX9 variants associated with non-syndromic oligodontia

Exon	Nucleotide change	Protein change	Type of variation	Right quadrants							Left quadrants							Reference	
				Max	8	7	6	5	4	3	2	1	1	2	3	4	5	6	
				Mand	8	7	6	5	4	3	2	1	1	2	3	4	5	6	
2 (PD)	c.218_219insG	p.Ser74fs	frameshift mutation	*	*	*	*	*	*						*	*	*	*	Stockton (2000) Nat Genet 24, 18
				*	*	*	*	*	*				*		*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	
				*	*	*	*								*	*	*	*	

					*	*	*	*	*	*	*
					*	*				*	*
					*		*	*		*	*
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					*	*	*	*		*	*
					*	*	*	*		*	*
					*		*			*	*
					*	*	*			*	*
					*	*				*	*
					*	*				*	*
2 (PD)	c.175_183delG ATACAAins28 8bp	p.Arg59fs	frameshift mutation		*	*	*	*		*	*
2 (PD)	c.271A > G	p.Lys91Glu	missense mutation		*	*	*			*	*
2 (PD)	c.76C > T	p.Arg26Trp	missense mutation		*	*	*	*		*	*
2	c.151G > A	p.Gly51Ser			*	*	*	*	*	*	*

Lammi (2003) Eur J Hum Genet 11, 866

(PD)			missense mutation		*	*			*	*	*	*	*	Mostowska (2003) Eur J Oral Sci 111, 272
2 (PD)	c.83G > C	p.Arg28Pro	missense mutation		*	*	*	*	*	*	*	*	*	Jumlongras (2004) Hum Genet 114, 242
1	c.1A > G	p.Met1Val	missense mutation		*	*	*	*	*	*	*	*	*	Klein (2005) J Dent Res 84, 43
					*	*	*	*			*	*	*	
					*	*	*	*			*	*	*	
					*	*	*	*		*	*	*	*	
2 (PD)	c.108_109insG	p.Ile37fs	frameshift mutation		*	*		*	*	*	*	*	*	Zhao (2005) Zhonghua Kou Qiang Yi Xue Za Zhi 40, 266
2 (PD)	c.139C > T	p.Arg47Trp	missense mutation		*	*	*		*	*	*	*	*	
2 (PD)	c.259A > T	p.Ile87Phe	missense mutation		*	*	*	*			*	*	*	Kapadia (2006) Eur J Hum Genet 14, 403
					*	*	*	*			*	*	*	
					*	*	*	*			*	*	*	
					*	*	*	*			*	*	*	
2	c.619_621delAT Cins24bp	p.Ile207fs	frameshift mutation		*	*	*	*			*	*	*	Mostowska (2006) Eur J Hum Genet 14, 173
					*	*	*	*		*	*	*	*	
					*	*	*	*			*	*	*	
					*	*	*	*		*	*	*	*	
					*	*	*	*			*	*	*	
					*	*	*	*		*	*	*	*	
					*	*	*	*			*	*	*	
					*	*	*	*		*	*	*	*	
					*	*	*	*		*	*	*	*	
2	c.433C > T	p.Gln145*	nonsense mutation		*	*	*			*	*	*	*	Hansen (2007) Eur J Oral Sci 115, 330
					*	*	*		*	*	*	*	*	
					*									
					*	*								
					*									
					*									
					*	*	*				*	*	*	

					*	*	*	*	*	*	*	*	*	*	*	*	*	
					*	*				*					*	*	*	
3	c.718G > C	p.Ala240Pro	missense mutation				*	*		*	*							Wang (2009) Hua Xi Kou Qiang Yi Xue Za Zhi 27, 606
2 (PD)	c.16G > A	p.Gly6Arg	missense mutation		*							*			*	*		Wang (2009) Cells Tissues Organs 189, 80
	c.128G > A and c.129C > A	p.Ser43Lys			*			*				*		*	*	*		
					*			*				*			*			
					*										*			
					*										*			
5'- UTR	g.-1258G > A				*	*	*							*	*	*		Mendoza-Fandino (2011) Clin Genet 80, 265
					*	*	*							*		*	*	
					*										*			
					*										*			
					*										*			
					*			*	*					*		*		
					*			*	*					*		*		
					*			*	*					*		*		
					*			*	*					*		*		
					*			*	*					*		*		
					*			*	*					*		*		
					*			*	*					*		*		
					*			*	*					*		*		
					*			*	*					*		*		
					*			*	*					*		*		
2 (PD)	c.321_322insG	p.Ala108fs	frameshift mutation		*	*	*	*						*		*	*	Suda (2011) J Dent Res 90, 382
					*	*	*	*						*		*	*	
					*	*	*								*	*		
					*	*	*	*							*	*		
					*	*	*	*							*	*		
					*	*	*	*							*	*		

					*	*	*			*			*	*	*	
2 (PD)	c.80T > C	p.Leu27Pro	missense mutation		*	*	*		*			*	*	*	*	
					*	*	*						*	*	*	
					*	*	*		*			*	*	*	*	
					*	*	*		*							
2 (PD)	c.86T > C	p.Ile29Thr	missense mutation		*	*	*	*	*			*		*	*	
					*	*			*			*			*	
					*	*	*		*			*	*	*	*	
					*	*								*	*	
2 (PD)	c.43T > A	p.Phe15Ile	missense mutation		*		*					*			*	
					*				*	*					*	
					?	*	*					*	*	*	*	?
					?	*	*		*					*	?	
2 (PD)	c.218_219insG	p.Ser74fs	frameshift mutation		?	*	*	*	*	*		*	*	*	*	?
					?	*	*		*					*	*	?
2	c.480C > G	p.Tyr160*	nonsense mutation		?	*	*	*	*			*	*	*	*	?
2	c.480C > G	p.Tyr160*	nonsense mutation		?	*	*		*			*	*	*	*	?
2 (PD)	c.140G > C	p.Arg47Pro	missense mutation		*	*						*		*	*	
					*		*					*		*		
2 (PD)	c.167T > C	p.Ile56Thr	missense mutation		*	*	*							*	*	
					*									*	*	
2 (PD)	c.340A > T	p.Lys114*	nonsense mutation		*	*	*							*	*	
					*									*	*	
2	c.406C > T	p.Gln136*	nonsense mutation		*	*	*	*						*	*	
					*									*	*	
2	c.503C > G	p.Ala168Gly	missense mutation		*									*		
					*									*		
					*									*		
					*		*							*		
					*		*							*		
					*		*							*		
					*									*		
					*									*		
					*									*		

Liang (2012) Arch Oral Biol 57, 784

Wang (2012) PLoS One 7, e51533

Zhu (2012) Mutagenesis 27, 313

Arte (2013) PLoS One 8, e73705

Boeira Junior (2013) J Oral Pathol Med 42, 99

					*	*		*		
					*	*		*		
					*					
2 (PD)	c.353_354insTG CC	p.Ser119fs	frameshift mutation		?	*	*		*	*
2 (PD)	c.73-75delATC	p.Ile25del	frameshift mutation		*	*	*		*	*
2 (PD)	c.146C > T	p.Ser49Leu	missense mutation		*	*	*	*	*	*
IVS2	g.9527G > T		splicing		*	*	*	*	*	*
					*	*			*	*
					*	*	*	*	*	*
					*	*			*	*
					*	*			*	*
					*	*			*	*
2 (PD)	c.59C > T	p.Pro20Leu	missense mutation		*	*	*	*	*	*
					*	*	*	*	*	*
					*		*			*
					*					*
					*	*	*	*	*	*
					*			*		*
					*					*
2	c.592_596dup	p.Asp200fs	frameshift mutation		*	*	*	*	*	*
					*	*	*	*	*	*
					*				*	*
					*		*		*	*
					*	*	*	*	*	*
					*					*
					*				*	*
1	c.2T > G	p.Met1Arg	missense mutation		*	*	*	*	*	*
					*	*	*		*	*
					*	*	*		*	*

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2 (PD)	g.10672A > T	p.Asn116Ile	missense mutation			*																	
2 (PD)	c.289_296del	p.Ile97fs	frameshift mutation																				
1	c.3G > A	p.Met1Ile	missense mutation			* * * * * * * *																	
2 (PD)	c.76C > T	p.Arg26Trp	missense mutation			* *																	
2 (PD)	c.140G > C	c.140G>C	missense mutation			* * * *																	
2 (PD)	c.146delC	p.Ser49fs	frameshift mutation			* * * * *																	
2 (PD)	c.167T > C	p.Ile56Asn	missense mutation			* * * *																	
2 (PD)	c.185_189dup	p.Gly64fs	frameshift mutation			* *																	
2	c.194C > A	p.Ser65*			* * * * *																		

Shahid (2016) Eur J Med Genet 59, 377

Yu (2016) Am J Hum Genet 99, 195

Sarkar (2017) Gene 635, 69

Wong (2018) J Dent Res 97, 155

(PD)			nonsense mutation		*	*	*	*	*	*	*	*	*	*	*	*	*	*	
2 (PD)	c.218dupG	p.Ser74fs	frameshift mutation		*	*	*	*	*	*	*	*	*	*	*	*	*	*	
2 (PD)	c.256_262dup	p.Arg88fs	frameshift mutation		*	*	*						*	*	*		*		
2 (PD)	c.271A > T	p.Lys91*	c		*	*	*	*	*	*	*	*	*	*	*	*	*	*	
2 (PD)	c.322G > C	p.Ala108Pro	missense mutation		*				*	*					*	*	*		
2	c.592delG	p.Val198fs	frameshift mutation		Cannot find more details														
1	c.2T > A	p.Met1*	nonsense mutation		*	*	*	*	*				*	*	*	*	*	*	
2(PD) 3	c.331G > A and c.718G > C	p.Val111Met p.Ala240Pro	missense mutation		*	*	*	*	*				*						Zhang(2019) Oral Dis 25, 234
2 (PD)	c.211_212insA	p.Ile71fs	frameshift mutation		*	*	*	*	*	*	*	*	*	*	*	*	*	*	
2 (PD)	c.229C > G	p.Arg77Gly	missense mutation		*		*	*					*	*	*			*	Sun(2021) Oral Dis 27,1468
2 (PD)	c.236_237insAC	p.Ter80fs	frameshift mutation		*	*	*	*					*		*	*	*		
2 (PD)	c.59C > T	p.Pro20Leu	missense mutation		*	*	*		*	*	*				*		*	*	Intarak (2022) Eur J Oral Sci 130,2
2 (PD)	c.230G>A	p.Arg77Gln	missense mutation		*	*	*								*	*	*		
2	c.491- 510delGCCCT ATCACGGCG GCAGGCC	p.Pro165fs	frameshift mutation		*	*	*	*	*	*	*	*	*	*	*	*	*		Sun(2022) Oral Dis May 20

					*	*	*	*	*	*	*	*	*	*	*
					*	*	*		*	*	*	*	*	*	*
					*	*	*		*	*	*	*	*	*	*
					*	*	*	*	*	*	*	*	*	*	*
					*	*	*	*	*	*	*	*	*	*	*
2 (PD)	c.350T>G	p.Val117Gly	missense mutation		*										*
					*	*									*
					*										*
3	c.648_649insC	p.Tyr217Leuf s*100	frameshift mutation		*	*	*	*	*		*	*	*	*	*
2 (PD)	c.352delC	p.Ser119Prof s*2	frameshift mutation		*	*	*	*	*	*	*	*	*	*	*
2 (PD)	c.191G>T	p. Gly64Val	missense mutation		*	*	*		*	*	*	*	*	*	*
					*	*	*	*	*	*	*	*	*	*	*

this study