

CORRECTION

Correction: Investigation of Susceptibility Genes Triggering Lachrymal/Salivary Gland Lesion Complications in Japanese Patients with Type 1 Autoimmune Pancreatitis

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There are errors in the ninth, tenth and eleventh sentences of the Abstract. The correct sentences are: Individual genotyping of *KLF7* rs2284932 revealed that the frequency of the minor C allele was significantly increased ($P = 0.00062$, $P_c = 0.003$, OR = 2.98, 95%CI = 1.58–5.65) in group A. The minor T allele of rs4473559 in *FRMD4* demonstrated a significant association in the A group ($P = 0.00015$, OR = 3.38, 95%CI = 1.77–6.45). In the *LOC101928923* gene, the frequency of the minor T allele of rs4379306 was significantly decreased in group A in both TaqMan and GWAS analyses. Lastly, the minor C allele of *MPPED2* rs514644 carried a significantly increased risk of complications.

There is an error in the third sentence of the Genetic Analysis section of the Results. The correct sentence is: The results for strong signals ($P < 0.0001$) are shown in Table 1. There are also errors in the penultimate sentence of this section. The correct sentence is: In the *LOC101928923* gene, the frequency of the minor T allele of rs4379306 was significantly decreased ($P = 0.011$, OR = 0.43) in both TaqMan and GWAS analyses in the A group (Table 5).

There is an error in [Table 3](#). The P value for rs2284932 should be CC+TC/TT. Please see the corrected [Table 3](#) here.



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Table 3. Association analysis of single nucleotide polymorphisms in the KLF7 gene.

dbSNP ID	Chrom. location	Typing method	Alleles	Frequency (%)		P value	Pc value	OR (95% CI)		
				A group	B group					
rs2287505	207655331	GWAS	C>A			A	0.027	2.74(1.09–6.90)		
rs1263615	207667483	TaqMan	A>G	A	74.0	68.6	G	0.758	3.032	1.10(0.61–1.96)
				G	26.0	31.4	GG+GA/AA	0.567	2.268	1.26(0.57–2.81)
rs768090	207711824	TaqMan	A>T	A	74.0	83.3	T	0.091	0.364	1.80(0.91–3.57)
				T	26.0	16.7	TT+TA/AA	0.047	0.188	2.28(1.01–5.16)
rs10195536	207715065	GWAS	T>A			A	0.016	3.79(1.21–11.92)		
rs2284932	207720754	TaqMan	T>C	T	90.0	95.1	C	0.00062	0.003	2.98(1.58–5.65)
				C	10.0	4.9	CC+TC/TT	0.00037	0.002	4.37(1.90–10.02)
rs12466923	207721800	TaqMan	A>C	A	60.0	81.4	C	0.039	0.156	2.16 (1.03–4.54)
				C	40.0	18.6	CC+AC/AA	0.0093	0.037	3.10(1.30–7.38)

A group: with lachrymal/salivary gland lesions, B group: without lachrymal/salivary gland lesions, dbSNP ID: SNP database identification, Chrom: chromosome, Pc: corrected P

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Reference

1. Oguchi T, Ota M, Ito T, Hamano H, Arakura N, Katsuyama Y, et al. (2015) Investigation of Susceptibility Genes Triggering Lachrymal/Salivary Gland Lesion Complications in Japanese Patients with Type 1 Autoimmune Pancreatitis. PLoS ONE 10(5): e0127078. doi: [10.1371/journal.pone.0127078](https://doi.org/10.1371/journal.pone.0127078) PMID: [25985088](https://pubmed.ncbi.nlm.nih.gov/25985088/)