

Supplementary Table 2. Missense *PNLIP* variants in Japanese subjects with chronic pancreatitis and healthy controls. Variants are heterozygous unless indicated as homozygous (hm).

Exon	Nucleotide change	Amino-acid change	Cases (%)	Controls (%)	<i>P</i> value
3	c.160C>T	p.R54C	2/223 (0.9)	1/1070 (0.1)	0.08
3	c.172T>C	p.Y58H	2/223 (0.9)	26/1070 (2.4)	0.21
3	c.175A>G	p.T59A	0/223 (0)	7/1070 (0.7)	0.61
6	c.557T>C	p.I186T	12/223 (5.4)	77 (2 hm)/1070 (7.2)	0.39
9	c.919G>A	p.V307I	0/223 (0)	1/1070 (0.1)	1.0
All variants			16/223 (7.2)	112/1070 (10.5)	0.14
All variants with p.I186T excluded			4/223 (1.8)	35/1070 (3.3)	0.29