
ACADVL Variant Allele Frequencies
2/19/2009

Notes: These summary statistics were compiled from clinical tests.
Only variants that are within the coding regions and about 50 base pairs of flanking sequence on each side are reported.
At this time, these summary statistics do NOT include insertion/deletion variants.

Exon	Left Flanking	Sequence Variation	Right Flanking	Coding Change	Documentation	Allele Frequencies
2	GGCCCTATGCCGGGG	c.128 G>A	TGCCGCTCAGGTAAG	p.Gly43Asp	rs2230178	G-96.2% A-3.8%
8	GGGGAGACTGTGGCC	c.637 G>A	CTTTCTGTCTAACCG	p.Ala213Thr		G-96.2% A-3.8%
13	GGGGTATGGGCTTCA	c.1328 T>G	GAAGGTACAGGACGG	p.Met443Arg		T-96.2% G-3.8%
16	TGGCGAGCTGGTAAG	c.1605+6 T>C	GGCCAGGGGTCCAGG	Intronic	rs17671352	C-69.2% T-30.8%