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**ACTA1 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.

<b>Exon</b>	<b>Left Flanking</b>	<b>Sequence Variation</b>	<b>Right Flanking</b>	<b>Coding Change</b>	<b>Documentation</b>	<b>Allele Frequencies</b>
3	GGTGCGTGGTGTCTC	c.130-10 G>C	GCTCYGCAGGGCGTC	Intronic	rs41271481	G-81.2% C-18.8%
3	GTGGTGTCTCSGCTC	c.130-5 T>C	GCAGGGCGTCATGGT	Intronic	rs11803533	T-81.2% C-18.8%
6	CTGCTCACGCTCCCC	c.809-15dupC	SCCGCGGTCCCCAGG	Intronic		dup C-30%
6	TGCTCACGCTCCCC	c.809-14 G>C	CCGCGGTCCCCAGGT	Intronic	rs6673359	G-85% C-15%