

Variant Analysis Report

Generated Version: v1.0.0 (January 2025)

List of All Variants

- * NM_000051.4:c.5075_5076del (Pathogenic)
- * NM_007294.4:c.3835G>A (Benign)
- * NM_000059.4:c.341A>G (Likely Benign)
- * NM_000059.4:c.7877G>C (VUS)
- * NM_007194.4:c.1375+2T>G (Pathogenic)
- * NM_024675.4:c.721A>G (Benign)
- * NM_000051.4:c.902G>A (VUS)
- * NM_007294.4:c.4357+6T>C (Pathogenic)
- * NM_000059.4:c.7879A>T (Pathogenic)
- * NM_000059.4:c.9234C>T (Benign)
- * NM_024675.4:c.3556del (VUS)
- * NM_007294.4:c.5213G>C (VUS)
- * NM_007294.4:c.661G>T (Benign)
- * NM_000059.4:c.9271G>A (Likely Benign)
- * NM_000059.4:c.9648+1G>C (Likely Pathogenic)
- * NM_000059.4:c.9883C>T (Pathogenic)
- * NM_024675.4:c.172_175del (Pathogenic)
- * NM_000051.4:c.7638_7646del (Pathogenic)
- * NM_007294.4:c.3891_3893del (Benign)
- * NM_007294.4:c.4185G>A (Pathogenic)
- * NM_007294.4:c.628C>T (VUS)
- * NM_002878.4:c.565_568del (Likely Pathogenic)
- * NC_000013.10(NM_000059.4):c.8953+2T>C (VUS)