

Figure 16. Sanger sequencing chromatogram for patient 13012708, showing a heterozygous intronic variant in the *ABCA4* gene (chr1:94,476,351; *ABCA4* c.5714+5G>A, NM_000350.2). c.5714+5G>A reduces the splicing efficiency of the exon40/intron40 donor splice site¹ and has previously been reported as pathogenic.^{2,3}

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