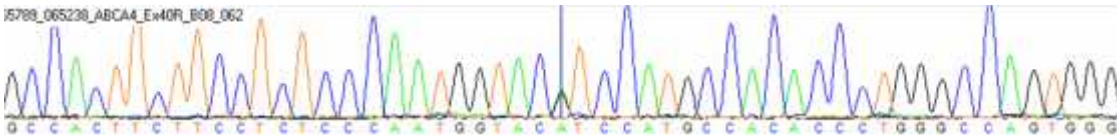

Figure 15. Sanger sequencing chromatogram for patient 065238, showing a heterozygous intronic variant in the *ABCA4* gene (chr1:94,476,351; *ABCA4* c.5714+5G>A, NM_000350.2). c.5714+5G>A reduces the splicing efficiency of the exon40/intron40 donor splice site[1] and has previously been reported as pathogenic.[2,3]

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