

# In silico analysis of DNA variants

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## Abstract

In order to determine the consequence of *ILDR1* reported mutations on splicing and translation initiation, we used Human Splicing Finder (version 3.0) (<http://www.umd.be/HSF3/>) and ORF finder ([http://www.geneinfinity.org/sms/sms\\_orffinder.html](http://www.geneinfinity.org/sms/sms_orffinder.html)) respectively.

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## Protocol