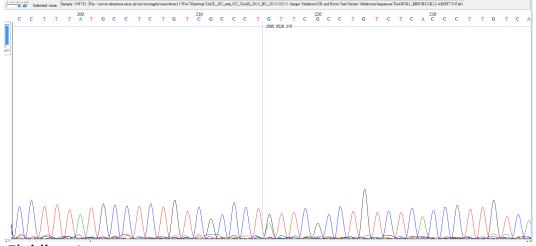
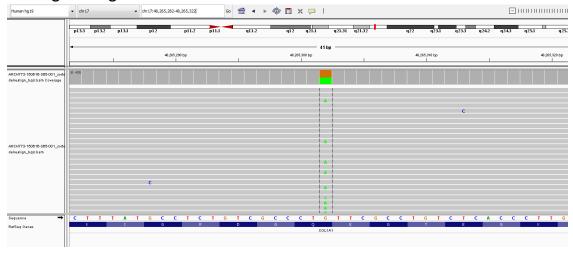
Variant Confirmation Report

ARCH773
COL1A1
45/51
ENST00000225964.5:c.3304C>T
ENSP00000225964.5:p.Gln1102Ter
17:48265302
0.42
144,106
Υ
This mutation introduces a prematurestop codon and is likely to be pathogenic

Validation:



Fluidigm Image:



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