

A

### Impact Prediction ✕

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Type	Interpretation
Broken WT Donor Site	Alteration of the WT Donor site, most probably affecting splicing

  

Algorithm/Matrix	position	sequences	variation
HSF Donor site (matrix GT)	chr3:123327620	- REF : CGG <b>G</b> TCTGT - ALT : CGG <b>T</b> TCTGT	80.48 > 53.34 => -33.72%
MaxEnt Donor site	chr3:123327620	- REF : CGG <b>G</b> TCTGT - ALT : CGG <b>T</b> TCTGT	3.55 > -4.95 => -239.44%

B

