

*Variant: NM\_001110792.2(MECP2):c.1199C>T  
(p.Pro400Leu)*

Version: 1.0

CA270246 [↗](#)

143398 (ClinVar) [↗](#)

**Gene:** MECP2 ([HGNC:4204](#))

**Condition:** Rett syndrome ([MONDO:0010726](#))

**Inheritance Mode:** X-linked inheritance

**UUID:** c9f8a43d-6228-4bee-8416-b26cbcfdb0e4

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### *HGVS expressions*

**NM\_001110792.2:c.1199C>T**

NM\_001110792.2(MECP2):c.1199C>T (p.Pro400Leu)

NC\_000023.11:g.154030665G>A

CM000685.2:g.154030665G>A

NC\_000023.10:g.153296116G>A

CM000685.1:g.153296116G>A

NC\_000023.9:g.152949310G>A

NG\_007107.2:g.111463C>T

NG\_007107.3:g.111439C>T

ENST00000303391.11:c.1163C>T

ENST00000453960.7:c.1199C>T

ENST00000303391.10:c.1163C>T

ENST00000407218.5:c.\*535C>T

ENST00000453960.6:c.1199C>T

ENST00000619732.4:c.1163C>T

ENST00000628176.2:c.\*535C>T

NM\_001110792.1:c.1199C>T

NM\_001316337.1:c.884C>T

NM\_004992.3:c.1163C>T

NM\_001316337.2:c.884C>T

NM\_001369391.2:c.884C>T

NM\_001369392.2:c.884C>T

NM\_001369393.2:c.884C>T

NM\_001369394.1:c.884C>T

NM\_001369394.2:c.884C>T

NM\_001386137.1:c.494C>T

NM\_001386138.1:c.494C>T

NM\_001386139.1:c.494C>T

NM\_004992.4:c.1163C>T

**Benign**

Met criteria codes **2**

BS1 BS2

Evidence Links **0**

Expert Panel



Rett and Angelman-like Disorders VCEP [↗](#)

## Evidence submitted by expert panel

***Rett and Angelman-like Disorders VCEP***

The allele frequency of the p.Pro388Leu (NM\_004992.3) variant in MECP2 is 0.014% in Latino sub population in gnomAD, which is high enough to be classified as likely benign based on thresholds defined by the ClinGen Rett/Angelman-like Expert Panel for Rett/AS-like conditions (BS1). The p.Pro388Leu variant is observed in at least 2 unaffected individuals (GeneDx internal database) (BS2). In summary, the p.Pro388Leu variant in MECP2 is classified as benign based on the ACMG/AMP criteria (BS1, BS2).

**Met criteria codes**

<b>BS1</b>		The allele frequency of the p.Pro388Leu variant in MECP2 is 0.014% in Latino sub population in gnomAD, which is high enough to be classified as likely benign based on thresholds defined by the ClinGen Rett/Angelman-like Expert Panel for Rett/AS-like conditions.
<b>BS2</b>		The p.Pro388Leu variant is observed in at least 2 unaffected individuals (GeneDx internal database)

Curation History 




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