

*Variant: NM\_004992.3(MECP2):c.806delG (p.Gly269Alafs)*

Version: 1.0

CA199475 [↗](#)

95202 (ClinVar) [↗](#)

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

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

**Inheritance Mode:** X-linked inheritance

**UUID:** 1e897fde-76a7-45dc-8f7f-6e1191668335

**Approved on:** 2021-03-09

**Published on:** 2021-05-07

### *HGVS expressions*

**NM\_004992.3:c.806delG**

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NM\_004992.3(MECP2):c.806delG (p.Gly269Alafs)

NC\_000023.11:g.154031025del

CM000685.2:g.154031025del

NC\_000023.10:g.153296476del

CM000685.1:g.153296476del

NC\_000023.9:g.152949670del

NG\_007107.2:g.111106del

NG\_007107.3:g.111082del

ENST00000303391.11:c.806del

ENST00000453960.7:c.842del

ENST00000637917.1:c.66-86del

ENST00000303391.10:c.806del

ENST00000407218.5:c.\*178del

ENST00000453960.6:c.842del

ENST00000619732.4:c.806del

ENST00000622433.4:c.794del

ENST00000628176.2:c.\*178del

NM\_001110792.1:c.842del

NM\_001316337.1:c.527del

NM\_001110792.2:c.842del

NM\_001316337.2:c.527del

NM\_001369391.2:c.527del

NM\_001369392.2:c.527del

NM\_001369393.2:c.527del

NM\_001369394.1:c.527del

NM\_001369394.2:c.527del

NM\_001386137.1:c.137del

NM\_001386138.1:c.137del

NM\_001386139.1:c.137del

NM\_004992.4:c.806del

**Pathogenic**

Met criteria codes **3**

PS2\_Very Strong

PM2\_Supporting

PVS1

Evidence Links **0**

Expert Panel

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

Criteria Specification Information **!**

[Criteria Specifications for this VCEP](#)

Evidence submitted by expert panel

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

The p.Gly269Alafs\*20 variant in MECP2 is predicted to cause a premature stop codon that leads to a truncated or absent protein in a gene where loss-of-function is an established mechanism. There is significant evidence that loss of this region of the gene is pathogenic (PVS1). This variant has been reported as a de novo occurrence (biological parentage confirmed) in at least 2 individuals with Rett syndrome (PMID 26984561, 10854091) (PS2\_VS). The p.Gly269Alafs\*20 variant in MECP2 is absent from gnomAD (PM2\_supporting). In summary, the p.Gly269Alafs\*20 variant in MECP2 is classified as Pathogenic for Rett syndrome based on the ACMG/AMP criteria (PVS1, PS2\_VS, PM2\_supporting).

#### Met criteria codes

**PS2\_Very Strong**



≥2 independent occurrences of De novo (both maternity and paternity confirmed) in patients with the Rett syndrome and no family history,( PMID 26984561,10854091)

**PM2\_Supporting**



Met- variant is absent in gnomAD

**PVS1**



Met- Null variant (frame-shift) affecting gene MECP2, which is a known mechanism of disease, PMID: 12481990

[Curation History](#)

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