

- No Criteria Specification was provided in the publication event for the variant classifaction.
- The **Criteria Specification** shown below is a best-estimate based on the VCEP, gene, disease, and publication date.

Variant: NM_000277.2(PAH):c.500A>G (p.Asn167Ser)

CA229585

102703 (ClinVar)

Gene: PAH ([HGNC:5053](#))
Condition: phenylketonuria ([MONDO:0009861](#))
Inheritance Mode: Autosomal recessive inheritance
UUID: 33e1f5d5-20ab-45d2-8dcd-4eba431a8642
Approved on: 2018-08-13
Published on: 2018-12-10

HGVS expressions

NM_000277.2(PAH):c.500A>G (p.Asn167Ser)
NM_000277.2:c.500A>G
NC_000012.12:g.102866605T>C
CM000674.2:g.102866605T>C
NC_000012.11:g.103260383T>C
CM000674.1:g.103260383T>C
NC_000012.10:g.101784513T>C
NG_008690.1:g.55998A>G
NG_008690.2:g.96806A>G
NM_000277.1:c.500A>G
NM_001354304.1:c.500A>G
NM_000277.3:c.500A>G
ENST00000307000.7:c.485A>G
ENST00000549111.5:n.596A>G
ENST00000551988.5:n.530+10857A>G
ENST00000553106.5:c.500A>G
More

Uncertain Significance

Met criteria codes 2
PP4 BS1
Not Met criteria codes 1
PP3

Evidence Links 1

Expert Panel

[Phenylketonuria VCEP](#)

Criteria Specification Information

[Go to PubMed](#) Close