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**MT** OLWRFKRQGULD

**MutationTaster** in silico prediction tools used to evaluate the significance of identified amino acid changes

**Nomenclature** +\*96 QRP HQFODWRUETDULDQWKEVFOHRWDCENKEUHGLFWHBFWDYDULDQWKEURWHLDYHO  
OMIM 2QOLQHGHOLDKCHULWDQBP

**PolyPhen** in silico prediction tool used to evaluate the significance of amino acid changes

**POS** JHORPLSRVLWIRQKMDULDQWKRUPDMFKURPRVRPHSRVLWLRQ

**SIFT** in silico prediction tool used to evaluate the significance of amino acid changes

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