



POLH gene

DNA polymerase eta

Normal Function

The *POLH* gene provides instructions for making a protein called DNA polymerase eta. DNA polymerases are a group of enzymes that "read" sequences of DNA and use them as templates to produce new DNA. These enzymes are important for copying (replicating) cells' genetic material in preparation for cell division. DNA polymerases also play critical roles in DNA repair.

The major function of DNA polymerase eta is to replicate DNA that has been damaged, particularly by ultraviolet (UV) rays from sunlight. Most other DNA polymerases are unable to replicate DNA with this type of damage. When they reach a segment of damaged DNA, they get stuck and the replication process stalls. However, when DNA polymerase eta encounters damaged DNA, it skips over the abnormal segment and continues copying. This activity, which is known as translesion synthesis, allows cells to tolerate some abnormalities created by UV exposure. Without this tolerance, unrepaired DNA damage would block DNA replication and cause the cell to die. Therefore, DNA polymerase eta plays an essential role in protecting cells from some of the effects of DNA damage.

DNA polymerase eta is a relatively "error-prone" polymerase. When it bypasses damaged DNA, it often inserts an incorrect DNA building block (nucleotide). This type of error results in a mutation in the replicated DNA.

Health Conditions Related to Genetic Changes

Xeroderma pigmentosum

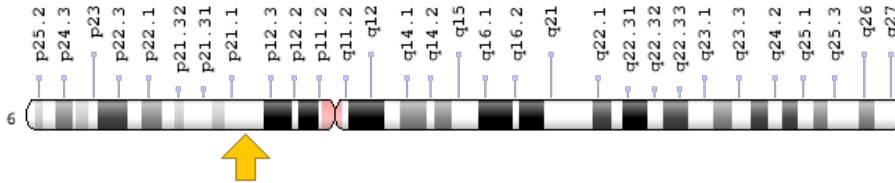
More than 30 mutations in the *POLH* gene have been found to cause the variant type of xeroderma pigmentosum (XP-V). Like the other forms of xeroderma pigmentosum, XP-V is characterized by an increased sensitivity to UV rays from sunlight. However, this form of the disorder is typically not associated with neurological abnormalities such as delayed development and hearing loss.

Most *POLH* gene mutations prevent the production of any detectable DNA polymerase eta. A loss of this enzyme prevents cells from replicating damaged DNA effectively. Without this mechanism for tolerating DNA damage, errors resulting from exposure to UV rays accumulate in genes that control cell growth and division. These errors cause cells to grow too fast and in an uncontrolled way. As a result, people with XP-V have an increased risk of developing skin cancer in areas where the skin is exposed to sunlight.

Chromosomal Location

Cytogenetic Location: 6p21.1, which is the short (p) arm of chromosome 6 at position 21.1

Molecular Location: base pairs 43,576,147 to 43,620,523 on chromosome 6 (Homo sapiens Updated Annotation Release 109.20190905, GRCh38.p13) (NCBI)



Credit: Genome Decoration Page/NCBI

Other Names for This Gene

- FLJ16395
- FLJ21978
- POLH_HUMAN
- polymerase (DNA directed), eta
- polymerase (DNA) eta
- RAD30
- RAD30 homolog A
- RAD30A
- xeroderma pigmentosum variant type protein
- XP-V
- XPV

Additional Information & Resources

Educational Resources

- Genomes (second edition, 2002): The DNA polymerases of bacteria and eukaryotes
<https://www.ncbi.nlm.nih.gov/books/NBK21113/#A8138>

Clinical Information from GeneReviews

- Xeroderma Pigmentosum
<https://www.ncbi.nlm.nih.gov/books/NBK1397>

Scientific Articles on PubMed

- PubMed
<https://www.ncbi.nlm.nih.gov/pubmed?term=%28%28POLH%5BTIAB%5D%29+OR+%28DNA+polymerase+eta%5BTIAB%5D%29%29+OR+%28%28XPV%5BTIAB%5D%29+OR+%28XP-V%5BTIAB%5D%29%29+AND+%28%28Genes%5BMH%5D%29+OR+%28Genetic+Phenomena%5BMH%5D%29%29+AND+english%5Bla%5D+AND+human%5Bmh%5D+AND+%22last+1800+days%22%5Bdp%5D>

Catalog of Genes and Diseases from OMIM

- POLYMERASE, DNA, ETA
<http://omim.org/entry/603968>

Research Resources

- Atlas of Genetics and Cytogenetics in Oncology and Haematology
<http://atlasgeneticsoncology.org/Genes/XPVID303.html>
- ClinVar
<https://www.ncbi.nlm.nih.gov/clinvar?term=POLH%5Bgene%5D>
- HGNC Gene Symbol Report
https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:9181
- Monarch Initiative
<https://monarchinitiative.org/gene/NCBIGene:5429>
- NCBI Gene
<https://www.ncbi.nlm.nih.gov/gene/5429>
- UniProt
<https://www.uniprot.org/uniprot/Q9Y253>

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Reprinted from Genetics Home Reference:
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Reviewed: May 2010
Published: October 15, 2019

Lister Hill National Center for Biomedical Communications
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