

WASHC5 gene

WASH complex subunit 5

Normal Function

The *WASHC5* gene provides instructions for making a protein called strumpellin. Strumpellin is active (expressed) throughout the body, although its exact function is unknown. The protein's structure suggests that strumpellin may interact with the structural framework inside cells (the cytoskeleton) and may attach (bind) to other proteins.

Health Conditions Related to Genetic Changes

Spastic paraplegia type 8

At least three mutations in the *WASHC5* gene have been found to cause spastic paraplegia type 8. These mutations change single building blocks (amino acids) in the strumpellin protein. One mutation that has been seen in multiple families replaces the amino acid valine with the amino acid phenylalanine at position 626 in strumpellin (written Val626Phe or V626F). *WASHC5* gene mutations are thought to change the structure of the strumpellin protein. It is unknown how the altered strumpellin protein causes muscle weakness, muscle stiffness, and other features of spastic paraplegia type 8.

Cancers

Research has shown that the *WASHC5* gene is abnormally active (overexpressed) in certain types of prostate cancer. Scientists do not know what causes this abnormal expression and have not determined whether the *WASHC5* gene plays a role in the development of prostate cancer.

Other Names for This Gene

- KIAA0196
- MGC111053
- SPG8
- STRUM_HUMAN
- strumpellin

Additional Information & Resources

Tests Listed in the Genetic Testing Registry

- Tests of WASHC5 ([https://www.ncbi.nlm.nih.gov/gtr/all/tests/?term=9897\[geneid\]](https://www.ncbi.nlm.nih.gov/gtr/all/tests/?term=9897[geneid]))

Scientific Articles on PubMed

- PubMed (<https://pubmed.ncbi.nlm.nih.gov/?term=%28KIAA0196%5BTIAB%5D%29+AND+%28%28Genes%5BMH%5D%29+OR+%28Genetic+Phenomena%5BMH%5D%29%29+AND+english%5Bla%5D+AND+human%5Bmh%5D+AND+%22last+3600+days%22%5Bdp%5D%29>)

Catalog of Genes and Diseases from OMIM

- WASH COMPLEX, SUBUNIT 5; WASHC5 (<https://omim.org/entry/610657>)

Gene and Variant Databases

- NCBI Gene (<https://www.ncbi.nlm.nih.gov/gene/9897>)
- ClinVar ([https://www.ncbi.nlm.nih.gov/clinvar?term=WASHC5\[gene\]](https://www.ncbi.nlm.nih.gov/clinvar?term=WASHC5[gene]))

References

- Hedera P, Rainier S, Alvarado D, Zhao X, Williamson J, Otterud B, Leppert M, Fink JK. Novel locus for autosomal dominant hereditary spastic paraplegia, on chromosome 8q. *Am J Hum Genet.* 1999 Feb;64(2):563-9. doi: 10.1086/302258. Citation on PubMed (<https://pubmed.ncbi.nlm.nih.gov/9973294>) or Free article on PubMed Central (<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC1377766/>)
- Meijer IA, Valdmanis PN, Rouleau GA. Spastic Paraplegia 8. 2008 Aug 13[updated 2020 May 21]. In: Adam MP, Feldman J, Mirzaa GM, Pagon RA, Wallace SE, Bean LJH, Gripp KW, Amemiya A, editors. *GeneReviews(R)* [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2024. Available from <http://www.ncbi.nlm.nih.gov/books/NBK1827/> Citation on PubMed (<https://pubmed.ncbi.nlm.nih.gov/20301727>)
- Porkka KP, Tammela TL, Vessella RL, Visakorpi T. RAD21 and KIAA0196 at 8q24 are amplified and overexpressed in prostate cancer. *Genes Chromosomes Cancer.* 2004 Jan;39(1):1-10. doi: 10.1002/gcc.10289. Citation on PubMed (<https://pubmed.ncbi.nlm.nih.gov/14603436>)
- Rocco P, Vainzof M, Froehner SC, Peters MF, Marie SK, Passos-Bueno MR, Zatz M. Brazilian family with pure autosomal dominant spastic paraplegia maps to 8q: analysis of muscle beta 1 syntrophin. *Am J Med Genet.* 2000 May 15;92(2):122-7. doi: 10.1002/(sici)1096-8628(20000515)92:23.0.co;2-b. Citation on PubMed (<https://pubmed.ncbi.nlm.nih.gov/10797436>)

- Valdmanis PN, Meijer IA, Reynolds A, Lei A, MacLeod P, Schlesinger D, Zatz M, Reid E, Dion PA, Drapeau P, Rouleau GA. Mutations in the KIAA0196 gene at the SPG8 locus cause hereditary spastic paraplegia. *Am J Hum Genet.* 2007 Jan;80(1):152-61. doi: 10.1086/510782. Epub 2006 Dec 1. Citation on PubMed (<https://pubmed.ncbi.nlm.nih.gov/17160902>) or Free article on PubMed Central (<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC1785307/>)
- van Duin M, van Marion R, Vissers K, Watson JE, van Weerden WM, Schroder FH, Hop WC, van der Kwast TH, Collins C, van Dekken H. High-resolution array comparative genomic hybridization of chromosome arm 8q: evaluation of genetic progression markers for prostate cancer. *Genes Chromosomes Cancer.* 2005 Dec;44(4):438-49. doi: 10.1002/gcc.20259. Citation on PubMed (<https://pubmed.ncbi.nlm.nih.gov/16130124>)

Genomic Location

The *WASHC5* gene is found on chromosome 8 (<https://medlineplus.gov/genetics/chromosome/8/>).

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