

HOXC4

This [gene](#) belongs to the [homeobox](#) family of genes. The homeobox genes encode a highly conserved family of transcription factors that play an important role in [morphogenesis](#) in all multicellular organisms. Mammals possess four similar homeobox gene clusters, [HOXA](#), [HOXB](#), [HOXC](#), and [HOXD](#), which are located on different chromosomes and consist of 9 to 11 genes arranged in tandem. This gene, [HOXC4](#), is one of several homeobox [HOXC](#) genes located in a cluster on [chromosome 12](#). Three genes, [HOXC5](#), [HOXC4](#), and [HOXC6](#), share a 5' non-coding exon. Transcripts may include the shared exon spliced to the gene-specific exons, or they may include only the gene-specific exons. Two alternatively spliced variants that encode the same protein have been described for [HOXC4](#). Transcript variant one includes the shared exon, and transcript variant two includes only gene-specific [exons](#).

Analysis of unique and powerful [Utah](#) genetic resources allowed the identification of 38 strong candidate Chiari malformation predisposition [gene](#) variants. These variants should be pursued in independent [populations](#). One of the candidates, a rare [HOXC4](#) variant, was identified in 2 high-risk CM pedigrees, with this variant possibly predisposing patients to a Chiari phenotype with [craniocervical kyphosis](#) ¹⁾.

¹⁾

Brockmeyer DL, Cheshier SH, Stevens J, Facelli JC, Rowe K, Heiss JD, Musolf A, Viskochil DH, Allen-Brady KL, Cannon-Albright LA. A likely [HOXC4](#) predisposition variant for Chiari malformations. J Neurosurg. 2022 Nov 25:1-9. doi: 10.3171/2022.10.JNS22956. Epub ahead of print. PMID: 36433874.

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