

Title: *PROP1*-Related Combined Pituitary Hormone Deficiency *GeneReview* – *HESX1* Variants

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Note: The following information is provided by the authors listed above and has not been reviewed by *GeneReviews* staff.

Specific variants in *HESX1*

- Dattani et al [1998] found a homozygous missense *HESX1* variant (p.Arg160Cys) in a brother and sister with septooptic dysplasia, agenesis of the corpus callosum, and CPHD (OMIM [182230](#)).
- Individuals homozygous for the recessive variant p.Arg160Cys had septo-optic dysplasia (SOD/Morsier syndrome) with agenesis of the corpus callosum and CPHD [Brickman et al 2001].
- Individuals with a monoallelic variant had milder phenotypes suggesting that they result from haploinsufficiency of the *HESX1* protein [Tajima et al 2003].
- In 2011, Vivenza et al describe a novel mutation (c.357+3G>A), identified at the heterozygous state in an IGHD patient. This mutation prevents the generation of one of the alternative isoforms normally produced by the wild-type allele, predicting a truncated *HESX1* protein. The mutation is likely to cause IGHD in the heterozygous patient by interfering with the downregulation of *HESX1* expression mediated by alternative splicing and nonsense-mediated decay.
- In 2011, Durmaz et al found a novel homozygous mutation (R160H) within the homeodomain of *HESX1*, which, is the first to be described in humans. Neuroimaging studies revealed anterior pituitary aplasia, a normal posterior pituitary gland, and a thin pituitary stalk but no midline abnormalities. Optic nerve studies showed no pathology. *HESX1* mutation causing an R160H substitution can result in panhypopituitarism without midline defects.
- Also in 2011, Reynaud et al studied 83 patients with pituitary stalk interruption syndrome and found one novel *HESX1* homozygous nonsense mutation generating a truncated protein, resulting in total loss of homeodomain and co-repressor binding. This *HESX1* p.Arg109X mutation leads to a severely truncated protein including the homeodomain that is involved in DNA binding and required to recruit components of N-CoR-associated co-repressors
- In 2013, a novel heterozygous sequence variant (c.142A>T, p.T48S) was found in *HESX1* in one PSIS patient by Yan Yang et al.

References

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