

### What is fragile X syndrome?

Fragile X syndrome is an X-linked disease of intellectual disability with variable severity.<sup>1</sup> Expansions of CGG repeat sequences in the *FMR1* gene account for 99% of mutations causing fragile X syndrome. Interpretation of CGG repeat expansion results is based on the following ranges: Negative: <45 repeats; intermediate: 45-54 repeats; premutation: 55-200 repeats; full mutation: >200 repeats. Greater than 99% of males and approximately 50% of females with the full mutation are intellectually disabled.<sup>2</sup>

### What are the symptoms of fragile X syndrome and what treatment is available?

Fragile X syndrome is associated with a range of symptoms. Early signs include delayed speech and language skills.<sup>1</sup> Intellectual problems vary from mild learning disabilities to severe intellectual disability.<sup>3</sup> Behavioral characteristics include autism and hyperactivity.<sup>1</sup> Physical features, such as a long face and large or prominent ears, are usually more noticeable in adults than in children, and in males more than females.<sup>3</sup> There is no cure for fragile X syndrome. Treatment is supportive and focuses on educational and behavioral support and management of symptoms.<sup>2</sup>

Individuals with a premutation do not have fragile X syndrome, but may have an increased risk for fragile X-related disorders. Females may have fragile X-associated primary ovarian insufficiency (FXPOI), which can cause infertility or early menopause. Most males with a premutation and some females are at risk for fragile X-associated tremor and ataxia syndrome (FXTAS), which can affect balance and is associated with tremor and memory problems in older individuals.<sup>3</sup>

Intermediate numbers of CGG repeats are not associated with symptoms and there is no increased risk to have children with fragile X syndrome.<sup>2</sup>

### How is fragile X syndrome inherited?

Fragile X syndrome is an X-linked disease caused by mutations in the *FMR1* gene.<sup>1</sup> When a premutation is passed from a mother to her offspring, the number of CGG repeat sequences may expand in length from a premutation to a full mutation.<sup>2</sup>

If a female carries a full mutation, the risk for each son or daughter to inherit the full mutation is 50%.<sup>4</sup> If a female carries a premutation, the risk that the premutation will expand to a full mutation depends on the repeat size of the premutation.<sup>2</sup> Men who are carriers pass the premutation to all of their daughters but none of their sons.<sup>4</sup>

### Who is at risk for fragile X syndrome?

Fragile X syndrome can occur in individuals of all races and ethnicities. It affects approximately 1 in 4,000 males and 1 in 8,000 females.<sup>1</sup> One (1) in 259 females and 1 in 813 males are carriers of a premutation.<sup>5</sup>

Having a relative who is a carrier of fragile X syndrome or a relative who has been diagnosed with fragile X syndrome or a fragile X-associated disorder increases an individual's risk to be a carrier. Consultation with a genetics health professional may be helpful in determining carrier risk and appropriate testing.

### What does a positive test result mean?

If a fragile X premutation or full mutation is identified, an individual should speak to a physician or genetics health professional about the implications of the result and appropriate testing for at-risk family members.

### What does a negative test result mean?

A negative result means an individual does not have a premutation or full mutation and therefore is not at increased risk of having a child with fragile X syndrome due to an expansion mutation.

### Where can I get more information?

National Fragile X Foundation: <https://fragilex.org/fragile-x/fragile-x-syndrome/>

March of Dimes: <http://www.marchofdimes.org/complications/fragile-x-syndrome.aspx>

Genetics Home Reference: <http://ghr.nlm.nih.gov/condition/fragile-x-syndrome>

### References

1. Genetics Home Reference: Fragile X syndrome. Available at <http://ghr.nlm.nih.gov/condition/fragile-x-syndrome> . Accessed January 21, 2016.
2. Saul R and Tarleton J. FMR1-Related Disorders. *GeneReviews* Available at <http://www.ncbi.nlm.nih.gov/books/NBK1384/> Accessed January 21, 2016.
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5. Dombrowski C *et al*, Premutation and intermediate-size FMR1 alleles in 10572 males from the general population: loss of an AGG interruption is a late event in the generation of fragile X syndrome alleles. *Hum Mol Genet*. 2002 11(4):371-8