

'Genetics Home Reference': Helping Patients Understand the Role of Genetics in Health and Disease

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Abstract

The surge of information generated by the Human Genome Project has left many health professionals and their patients struggling to understand the role of genetics in health and disease. To aid the lay public and health professionals, the US National Library of Medicine developed an online resource called 'Genetics Home Reference' (GHR), located at <http://ghr.nlm.nih.gov/>. Launched in April 2003, GHR's goal is to help the public interpret the health implications of the Human Genome Project. It bridges the clinical questions of consumers and the rich technical data emerging from the sequenced human genome. The GHR web site is designed for easy navigation among summaries for genetic conditions and the related gene(s) and chromosome(s). This design strategy enhances the user's appreciation of how genes, chromosomes, and conditions are interrelated.

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Introduction

The Human Genome Project amplified interest in genetics and propelled medicine into a new era in which genetic knowledge will help determine optimal health care [1, 2]. The surge of genetic information generated by the Human Genome Project is overwhelming, however, and often leaves patients and health professionals struggling to understand the implications of genetic findings in medical practice [1, 3]. Several organizations, such as the World Health Organization and the International Society of Bioethics, recognize that the goals of medical genetics can be fulfilled only in the context of an educated and informed public [4–6]. The healthcare community is challenged to communicate the complex developments in human genetics in a way that the public can freely access, easily understand, and appropriately apply [7].

Increasingly, the public is seeking health information online [8, 9], including information about inherited disorders [10, 11]. Consumers, however, report that genetics web sites are often confusing, difficult to navigate, and hard to understand [7, 12]. To address this confusion, the US National Library of Medicine developed an online resource called the Genetics Home Reference (GHR), located at <http://ghr.nlm.nih.gov/>.

Goal and General Overview

Launched in April 2003, GHR's goal is to help the public interpret the health implications of the Human Genome Project. It bridges the clinical questions of consumers and the rich technical data emerging from the sequenced human genome. Using a question and answer format, GHR describes genetic conditions and the gene or chromosome alterations associated with those conditions. Information for more than 160 genetic conditions and 250 genes is currently available. New topics are added continuously, and the information is monitored regularly to ensure that it is accurate and up-to-date. Experts in clinical genetics review each topic before it is posted to the GHR site and annually thereafter (for details see <http://ghr.nlm.nih.gov/ghr/ExpertReviewers>).

GHR's target audience is members of the general public who are motivated to learn about genetic conditions. GHR, however, also serves as an educational site for students and journalists. For health professionals, the site provides information suitable for patients as well as links to genetic resources with primary data and detailed clinical and technical discussions. By integrating a spectrum of resources, from patient-friendly fact sheets to technical databases, GHR serves the public, academic, clinical, and research communities.

General Design

The GHR web site is designed for easy navigation among summaries for genetic conditions and the related gene(s) and chromosome(s) [13]. This design strategy enhances the user's appreciation of how genes, chromosomes, and conditions are interrelated. Using muscular dystrophy (Duchenne and Becker types) as an example, users can navigate from the condition summary for muscular dystrophy (fig. 1a) to the DMD gene (fig. 1b) and then to a summary for the corresponding X chromosome (fig. 1c). From the summary about the X chromosome, there is a link to related chromosomal disorders such as Turner syndrome (fig. 1c). Navigation is not unidirectional. Users can begin with a gene or chromosome summary and follow the links to related conditions.

The GHR design also accommodates four possible relationships between genetic alterations (on a gene or chromosome level) and particular conditions. These relationships are: (1) causal, as HBB mutations cause sickle cell anemia or trisomy 21 causes Down syndrome; (2) predisposing, as a particular COL1A1 polymorphism appears

to increase the risk of developing osteoporosis; (3) associated, as TP53 mutations or chromosome translocations are associated with certain cancers, and (4) modifying, as HFE mutations modify the course of X-linked sideroblastic anemia. A particular gene may have more than one relationship, such as FGFR3 mutations in the germ line can cause achondroplasia, but FGFR3 mutations in somatic cells may increase the risk for bladder cancer.

Content

The content for each topic summary is derived from the primary literature and other sources such as US federal agencies, academic sites, and professional organizations. Each summary includes a bibliography from the link to references.

Medical and genetics guidelines [14, 15] are used to name conditions, and alternate names are listed as synonyms. Each condition summary provides a concise description of the disorder and an explanation of incidence or prevalence, inheritance pattern, and related genetic factors such as gene mutations or chromosome alterations. Condition summaries link to a range of educational and patient support resources such as MedlinePlus [16], ClinicalTrials.gov [17] and GeneTests [18].

Gene summaries use the official gene name and symbol, as determined by the HUGO Gene Nomenclature Committee [19]. Each summary explains the gene's normal function and how mutations in the gene cause, modify, or are otherwise associated with particular genetic conditions. Gene summaries link to resources such as Entrez Gene [20] and GeneTests [18].

Chromosome summaries provide an estimate of the number of genes on each chromosome, and some include a discussion of related chromosomal disorders. These summaries link to chromosome-specific resources such as MapViewer [21], Ensembl [22], Chromosome Launchpad [23], and other chromosome databases.

Learning Aids

Recognizing that the target audience may have a limited science background, GHR offers tools to help the motivated learner. Each web page links to a glossary of genetic and medical terms. In addition, each summary provides a list of glossary terms used on the page, with a direct link to their definitions. The GHR glossary is built from the following established sources: the Unified Med-

Fig. 1. GHR is designed to ease navigation among summaries for genetic conditions and the related gene(s) and chromosome(s). **a** The summary for muscular dystrophy, Duchenne and Becker types, links to the related gene, DMD. **b** The DMD gene summary provides links to related conditions and also to the X chromosome, where the DMD gene is located. **c** The X chromosome summary links to chromosomal conditions such as Turner syndrome.

a

Muscular dystrophy, Duchenne and Becker types

On this page:

- What is muscular dystrophy, Duchenne and Becker types?
- How common is muscular dystrophy, Duchenne and Becker types?
- What genes are related to muscular dystrophy, Duchenne and Becker types?
- How do people inherit muscular dystrophy, Duchenne and Becker types?
- Where can I find additional information about muscular dystrophy, Duchenne and Becker types?
- What other names do people use for muscular dystrophy, Duchenne and Becker types?
- What if I still have specific questions about muscular dystrophy, Duchenne and Becker types?
- What glossary definitions help with understanding muscular dystrophy, Duchenne and Becker types?

What is muscular dystrophy, Duchenne and Becker type?

Muscular dystrophies are a group of genetic conditions that cause muscle weakness and wasting. The Duchenne and Becker types are the most common. They primarily affect the skeletal muscles, which are the muscles of the body. Duchenne muscular dystrophy is a severe form of the disease that usually begins in early childhood and progresses rapidly. Becker muscular dystrophy is a milder form that typically begins in the teenage years. The signs and symptoms of Duchenne muscular dystrophy are usually milder and exhibit a large range of variation. In general, they become apparent later in childhood or adolescence and progress at a much slower rate.

How common is muscular dystrophy, Duchenne and Becker types?

Duchenne and Becker muscular dystrophies together affect 1 in 3,500 to 5,000 male children.

Related Gene(s)

- **DMD:** dystrophin (muscular dystrophy, Duchenne and Becker types)

b

DMD

On this page:

- What is the official name of the DMD gene?
- What is the normal function of the DMD gene?
- What conditions are related to the DMD gene?
- Where is the DMD gene located?
- Where can I find additional information about DMD?
- What other names do people use for the DMD gene or condition?
- What glossary definitions help with understanding DMD?

What is the official name of the DMD gene?

The official name of this gene is "dystrophin (muscular dystrophy protein)." DMD is the gene's official symbol. The DMD gene is also listed below.

What is the normal function of the DMD gene?

The DMD gene provides instructions for making a protein called dystrophin. This protein is found in skeletal muscle cells, where it helps anchor the cells to the surrounding tissue. This anchoring is important for the normal function of the muscle cells. Mutations in the DMD gene can lead to a condition called Duchenne muscular dystrophy (DMD), which is a severe form of muscular dystrophy.

Where is the DMD gene located?

Xp21.2

The DMD gene is located on the short (p) arm of the X chromosome at position 21.2.

Related Condition(s)

- muscular dystrophy, Duchenne and Becker types
- other disorders

c

X chromosome

On this page:

- What is the X chromosome?
- What chromosomal conditions are related to the X chromosome?
- Is there a standard way to diagram the X chromosome?
- Where can I find additional information about the X chromosome?
- What glossary definitions help with understanding the X chromosome?

What is the X chromosome?

The X chromosome is one of the two sex chromosomes in humans (the other is the Y chromosome). The sex chromosomes are one of the 23 pairs of human chromosomes. The X chromosome is larger than the Y chromosome and contains more genes. Each person normally has two X chromosomes. In females, both X chromosomes are active. In males, one X chromosome is active and the other is permanently inactivated. This phenomenon is called X-inactivation. Identifying genes on the X chromosome is important for researchers because many genetic disorders are caused by changes in the structure or number of copies of the X chromosome.

Related Chromosomal Condition(s)

The following conditions are caused by changes in the structure or number of copies of the X chromosome.

- Klinefelter syndrome
- triple X syndrome
- Turner syndrome
- other chromosomal conditions

ical Language System (UMLS) [24], GeneTests [18], the US Department of Energy Human Genome Project [25], and three components of the US National Institutes of Health [26–28].

An online tutorial is available through a feature called the ‘Help Me Understand Genetics Handbook’, an illustrated primer that explains the basics of genetics. The handbook is organized by chapters that provide information about how genes work, types of gene mutations, patterns of inheritance, the role of a genetics professional, genetic testing, gene therapy, pharmacogenomics, and the Human Genome Project.

Additional Resources

The GHR site also serves as a conduit to many additional online resources from the US and other countries. Links to condition-, gene-, or chromosome-specific information on other web sites are available from each summary page. Additionally, from the ‘Resources’ link in the tool bar on each web page, GHR users will find links to genetics news, policies and ethics, online exhibits, the Human Genome Project, genetics and health, and educational tools. GHR makes available its criteria in selecting web sites as additional resources. Briefly, these sites should be educational and free of advertisement for commercial products. There should be no charge or registration required to use the site. The source of the site’s content should be clearly visible, respected, and dependable, and the information should be current.

Evaluation and Usage

Preliminary surveys, conducted in early 2003, helped to assess GHR’s layout, navigational design, and level of content. Feedback from these surveys influenced the question-answer format of summaries, glossary content, Handbook chapters, and the types and placement of links to additional resources. In early 2004, a formal survey was conducted among members of the Genetic Alliance [29], an international coalition that represents individuals with genetic conditions. The survey provided feedback about GHR’s design and content. The results [30], based on 374 respondents, indicated overall user satisfaction – 88% of users were satisfied or very satisfied with the site. Respondents found GHR to be authoritative, accurate, unbiased, up-to-date, and informative.

Since its launch two years ago, GHR site traffic has increased nearly 10-fold. The median daily use based on unique Internet protocol (IP) addresses rose from 466 in May 2003 to 4,213 in May 2005. Tracking data show that the most popular features of the site are summaries of genetic conditions, the genetics tutorial handbook, and the glossary.

Summary and Future Plans

GHR fills a unique niche by using plain language to interpret the health implications of the Human Genome Project. The site provides consumer-friendly information about genetic conditions and the genes and/or chromosomes that play a role in those conditions, along with a range of resources – from basic tutorials to web sites with clinical and technical information. Long-range plans include addressing multifactorial chronic conditions, epigenetics, and the role of single nucleotide polymorphisms. Additional evaluations among motivated health-seeking Internet consumers will help assess GHR’s usability and its content readability and impact. GHR will grow and evolve to help consumers navigate the complex world of genetics.

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