

Thrombocytopenia-absent radius syndrome

Description

Thrombocytopenia-absent radius (TAR) syndrome is characterized by the absence of a bone called the radius in each forearm and a shortage (deficiency) of blood cells involved in clotting (platelets). This platelet deficiency (thrombocytopenia) usually appears during infancy and becomes less severe over time; in some cases the platelet levels become normal.

Thrombocytopenia prevents normal blood clotting, resulting in easy bruising and frequent nosebleeds. Potentially life-threatening episodes of severe bleeding (hemorrhages) may occur in the brain and other organs, especially during the first year of life. Hemorrhages can damage the brain and lead to intellectual disability. Affected children who survive this period and do not have damaging hemorrhages in the brain usually have a normal life expectancy and normal intellectual development.

The severity of skeletal problems in TAR syndrome varies among affected individuals. The radius, which is the bone on the thumb side of the forearm, is almost always missing in both arms. The other bone in the forearm, which is called the ulna, is sometimes underdeveloped or absent in one or both arms. TAR syndrome is unusual among similar malformations in that affected individuals have thumbs, while people with other conditions involving an absent radius typically do not. However, there may be other abnormalities of the hands, such as webbed or fused fingers (syndactyly) or curved pinky fingers (fifth finger clinodactyly). Some people with TAR syndrome also have skeletal abnormalities affecting the upper arms, legs, or hip sockets.

Other features that can occur in TAR syndrome include malformations of the heart or kidneys. Some people with this disorder have unusual facial features including a small lower jaw (micrognathia), a prominent forehead, and low-set ears. About half of affected individuals have allergic reactions to cow's milk that may worsen the thrombocytopenia associated with this disorder.

Frequency

TAR syndrome is a rare disorder, affecting fewer than 1 in 100,000 newborns.

Causes

Mutations in the *RBM8A* gene cause TAR syndrome. The *RBM8A* gene provides instructions for making a protein called RNA-binding motif protein 8A. This protein is believed to be involved in several important cellular functions involving the production of other proteins.

Most people with TAR syndrome have a mutation in one copy of the *RBM8A* gene and a deletion of genetic material from chromosome 1 that includes the other copy of the *RBM8A* gene in each cell. A small number of affected individuals have mutations in both copies of the *RBM8A* gene in each cell and do not have a deletion on chromosome 1. *RBM8A* gene mutations that cause TAR syndrome reduce the amount of RNA-binding motif protein 8A in cells. The deletions involved in TAR syndrome eliminate at least 200,000 DNA building blocks (200 kilobases, or 200 kb) from the long (q) arm of chromosome 1 in a region called 1q21.1. The deletion eliminates one copy of the *RBM8A* gene in each cell and the RNA-binding motif protein 8A that would have been produced from it.

People with either an *RBM8A* gene mutation and a chromosome 1 deletion or with two gene mutations have a decreased amount of RNA-binding motif protein 8A. This reduction is thought to cause problems in the development of certain tissues, but it is unknown how it causes the specific signs and symptoms of TAR syndrome. No cases have been reported in which a deletion that includes the *RBM8A* gene occurs on both copies of chromosome 1; studies indicate that the complete loss of RNA-binding motif protein 8A is not compatible with life.

Researchers sometimes refer to the deletion in chromosome 1 associated with TAR syndrome as the 200-kb deletion to distinguish it from another chromosomal abnormality called a 1q21.1 microdeletion. People with a 1q21.1 microdeletion are missing a different, larger DNA segment in the chromosome 1q21.1 region near the area where the 200-kb deletion occurs. The chromosomal change related to 1q21.1 microdeletion is often called the recurrent distal 1.35-Mb deletion.

[Learn more about the gene and chromosome associated with Thrombocytopenia-absent radius syndrome](#)

- *RBM8A*
- chromosome 1

Inheritance

TAR syndrome is inherited in an autosomal recessive pattern, which means both copies of the gene in each cell are altered. In this disorder, either both copies of the *RBM8A* gene in each cell have mutations or, more commonly, one copy of the gene has a mutation and the other is lost as part of a deleted segment on chromosome 1. The affected individual usually inherits an *RBM8A* gene mutation from one parent. In about 75 percent of cases, the affected person inherits a copy of chromosome 1 with the 200-

kb deletion from the other parent. In the remaining cases, the deletion occurs during the formation of reproductive cells (eggs and sperm) or in early fetal development. Although parents of an individual with TAR syndrome can carry an *RBM8A* gene mutation or a 200-kb deletion, they typically do not show signs and symptoms of the condition.

Other Names for This Condition

- Chromosome 1q21.1 deletion syndrome, 200-KB
- Radial aplasia-amegakaryocytic thrombocytopenia
- Radial aplasia-thrombocytopenia syndrome
- TAR syndrome
- Thrombocytopenia absent radii

Additional Information & Resources

Genetic Testing Information

- Genetic Testing Registry: Radial aplasia-thrombocytopenia syndrome (<https://www.ncbi.nlm.nih.gov/gtr/conditions/C0175703/>)

Genetic and Rare Diseases Information Center

- Thrombocytopenia-absent radius syndrome (<https://rarediseases.info.nih.gov/diseases/5116/index>)

Patient Support and Advocacy Resources

- National Organization for Rare Disorders (NORD) (<https://rarediseases.org/>)

Catalog of Genes and Diseases from OMIM

- THROMBOCYTOPENIA-ABSENT RADIUS SYNDROME; TAR (<https://omim.org/entry/274000>)

Scientific Articles on PubMed

- PubMed (<https://pubmed.ncbi.nlm.nih.gov/?term=%28thrombocytopenia-absent+radius+syndrome%5BTIAB%5D%29+AND+english%5Bla%5D+AND+human%5Bmh%5D+AND+%22last+3600+days%22%5Bdp%5D>)

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