

# Spinocerebellar Ataxia Type 36

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Spinocerebellar ataxia type 36 (SCA36) is a condition characterized by progressive problems with movement that typically begin in mid-adulthood. People with this condition initially experience problems with coordination and balance (ataxia). Affected individuals often have exaggerated reflexes (hyperreflexia) and problems with speech (dysarthria). They also usually develop muscle twitches (fasciculations) of the tongue and over time, the muscles in the tongue waste away (atrophy). These tongue problems can cause difficulties swallowing liquids. As the condition progresses, individuals with SCA36 develop muscle atrophy in the legs, forearms, and hands. Another common feature of SCA36 is the atrophy of specialized nerve cells that control muscle movement (motor neurons), which can contribute to the tongue and limb muscle atrophy in affected individuals.

Keywords: genetic conditions

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## 1. Introduction

Some people with SCA36 have abnormalities of the eye muscles, which can lead to involuntary eye movements (nystagmus), rapid eye movements (saccades), trouble moving the eyes side-to-side (oculomotor apraxia), and droopy eyelids (ptosis). Sensorineural hearing loss, which is hearing loss caused by changes in the inner ear, may also occur in people with SCA36.

Brain imaging of people with SCA36 shows progressive atrophy of various parts of the brain, particularly within the cerebellum, which is the area of the brain involved in coordinating movements. Over time, the loss of cells in the cerebellum causes the movement problems characteristic of SCA36. In older affected individuals, the frontal lobes of the brain may show atrophy resulting in loss of executive function, which is the ability to plan and implement actions and develop problem-solving strategies.

Signs and symptoms of SCA36 typically begin in a person's forties or fifties but can appear anytime during adulthood. People with SCA36 have a normal lifespan and are usually mobile for 15 to 20 years after they are diagnosed.

## 2. Frequency

Approximately 100 individuals with SCA36 have been reported in the scientific literature. Almost all of these individuals have been from two regions: western Japan and the Costa de Morte in Galicia, Spain.

## 3. Causes

SCA36 is caused by mutations in the *NOP56* gene. The *NOP56* gene provides instructions for making a protein called nucleolar protein 56, which is primarily found in the nucleus of nerve cells (neurons), particularly those in the cerebellum. This protein is one part (subunit) of the ribonucleoprotein complex, which is composed of proteins and molecules of RNA, DNA's chemical cousin. The ribonucleoprotein complex is needed to make cellular structures called ribosomes, which process the cell's genetic instructions to create proteins.

The *NOP56* gene mutations that cause SCA36 involve a string of six DNA building blocks (nucleotides) located in an area of the gene known as intron 1. This string of six nucleotides (known as a hexanucleotide) is represented by the letters GGCCTG and normally appears multiple times in a row. In healthy individuals, GGCCTG is repeated 3 to 14 times within the gene. In people with SCA36, GGCCTG is repeated at least 650 times. It is unclear if 15 to 649 repeats of this hexanucleotide cause any signs or symptoms.

To make proteins from the genetic instructions carried in genes, a molecule called messenger RNA (mRNA) is formed. This molecule acts as a genetic blueprint for protein production. However, a large increase in the number of GGCCTG repeats in the *NOP56* gene disrupts the normal structure of NOP56 mRNA. Abnormal NOP56 mRNA molecules form clumps called RNA foci within the nucleus of neurons. Other proteins become trapped in the RNA foci, where they cannot function. These proteins may be important for controlling gene activity or protein production.

Additionally, researchers believe that the large expansion of the hexanucleotide repeat in the *NOP56* gene may reduce the activity of a nearby gene called *MIR1292*. The *MIR1292* gene provides instructions for making a type of RNA that regulates the activity (expression) of genes that produce proteins called glutamate receptors. These proteins are found on the surface of neurons and allow these cells to communicate with one another. A decrease in the production of Mir1292 RNA can lead to an increase in the production of glutamate receptors. The increased receptor activity may overexcite neurons, which disrupts normal communication between cells and can contribute to ataxia.

The combination of RNA foci and overly excited neurons likely leads to the death of these cells over time. Because the *NOP56* gene is especially active in neurons in the cerebellum, these cells are particularly affected by expansion of the gene, leading to cerebellar atrophy. Deterioration in this part of the brain leads to ataxia and the other signs and symptoms of SCA36.

### 3.1 The gene associated with Spinocerebellar ataxia type 36

- [NOP56](#)

## 4. Inheritance

This condition is inherited in an autosomal dominant pattern, which means one copy of the altered gene in each cell is sufficient to cause the disorder.

In most cases, an affected person has one parent with the condition.

In conditions that are caused by repeated segments of DNA, the number of repeats often increases when the altered gene is passed down from one generation to the next. Additionally, a larger number of repeats is usually associated with an earlier onset of signs and symptoms. This phenomenon is called anticipation. Some families affected by SCA36 have demonstrated anticipation while others have not. When anticipation is observed in SCA36, the mutation is most often passed down from the affected father.

## 5. Other Names for This Condition

- Asidan ataxia
- Costa de Morte ataxia
- SCA36
- spinocerebellar ataxia 36

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## References

1. Abe K, Ikeda Y, Kurata T, Ohta Y, Manabe Y, Okamoto M, Takamatsu K, Ohta T, Takao Y, Shiro Y, Shoji M, Kamiya T, Kobayashi H, Koizumi A. Cognitive and affective impairments of a novel SCA/MND crossroad mutation Asidan. *Eur J Neurol*. 2012 Aug;19(8):1070-8. doi: 10.1111/j.1468-1331.2012.03669.x.
2. García-Murias M, Quintáns B, Arias M, Seixas AI, Cacheiro P, Tarrío R, Pardo J, Millán MJ, Arias-Rivas S, Blanco-Arias P, Dapena D, Moreira R, Rodríguez-Trelles F, Sequeiros J, Carracedo A, Silveira I, Sobrido MJ. 'Costa da Morte' ataxia is spinocerebellar ataxia 36: clinical and genetic characterization. *Brain*. 2012 May;135(Pt 5):1423-35. doi: 10.1093/brain/aws069.
3. Ikeda Y, Ohta Y, Kobayashi H, Okamoto M, Takamatsu K, Ota T, Manabe Y, Okamoto K, Koizumi A, Abe K. Clinical features of SCA36: a novel spinocerebellar ataxia with motor neuron involvement (Asidan). *Neurology*. 2012 Jul 24;79(4):333-41. doi: 10.1212/WNL.0b013e318260436f.
4. Kobayashi H, Abe K, Matsuura T, Ikeda Y, Hitomi T, Akechi Y, Habu T, Liu W, Okuda H, Koizumi A. Expansion of intronic GGCCTG hexanucleotide repeat in NOP56 causes SCA36, a type of spinocerebellar ataxia accompanied by motor neuron involvement. *Am J Hum Genet*. 2011 Jul 15;89(1):121-30. doi: 10.1016/j.ajhg.2011.05.015.

5. Liu W, Ikeda Y, Hishikawa N, Yamashita T, Deguchi K, Abe K. Characteristic RNAfoci of the abnormal hexanucleotide GGCCUG repeat expansion in spinocerebellar ataxia type 36 (Asidan). *Eur J Neurol*. 2014 Nov;21(11):1377-86. doi:10.1111/ene.12491.
  6. Sugihara K, Maruyama H, Morino H, Miyamoto R, Ueno H, Matsumoto M, Kaji R, Kitaguchi H, Yukitake M, Higashi Y, Nishinaka K, Oda M, Izumi Y, Kawakami H. The clinical characteristics of spinocerebellar ataxia 36: a study of 2121 Japanese ataxia patients. *Mov Disord*. 2012 Aug;27(9):1158-63. doi: 10.1002/mds.25092.
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