

The genetic basis of the disorder is [mutations](#) in the [TSC1](#) or [TSC2](#) gene, which leads to overactivation of the mammalian target of rapamycin ([mTOR](#)) protein complex and results in the development of benign tumors in different body systems such as the brain, skin, lungs, and kidney.

Autosomal dominant inheritance; however, spontaneous mutation accounts for the majority of cases.

[Focal cortical dysplasia](#) (FCD) is a localized [cortical malformation](#) and considerable morphological overlap exists between [Focal cortical dysplasia type II B](#) (FCD IIB) and neurological lesions associated with [Tuberous sclerosis complex](#) (TSC). Abnormal [mTOR pathway](#) secondary to somatic [mTOR mutation](#) and TSC gene mutation linked to PI3K/AKT/mTOR pathway have supported the hypothesis of common pathogenesis involved.

Two distinct tumor suppressor genes have been identified: the [TSC1](#) gene (located on chromosome 9q34) codes for TSC1 (AKA hamartin), and the [TSC2](#) gene (on chromosome 16p13.3) codes for TSC2 (tuberin). Only 1 gene needs to be affected to develop TSC. These proteins work together to inhibit the activation of rapamycin (mTOR). Genetic counseling for unaffected parents with one affected child: 1-2% chance of recurrence ^{1) 2)}.

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European Chromosome 16 Tuberous Sclerosis Consortium. Identification and characterization of the tuberous sclerosis gene on chromosome 16. *Cell*. 1993;75:1305-1315.

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van Slegtenhorst M, deHoogt R, Hermans C, et al. Identification of the tuberous sclerosis gene TSC1 on chromosome 9q34. *Science*. 1997;277:805-808.

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