

IDH2 gene mutation

[Isocitrate dehydrogenase](#) [NADP], mitochondrial is an [enzyme](#) that in humans is encoded by the [IDH2](#) gene.

Isocitrate dehydrogenases are enzymes that catalyze the oxidative decarboxylation of isocitrate to 2-oxoglutarate. These enzymes belong to two distinct subclasses, one of which utilizes [NAD\(+\)](#) as the electron acceptor and the other as [NADP\(+\)](#). Five isocitrate dehydrogenases have been reported: three [NAD\(+\)](#)-dependent isocitrate dehydrogenases, which localize to the mitochondrial matrix, and two [NADP\(+\)](#)-dependent isocitrate dehydrogenases, one of which is mitochondrial and the other predominantly cytosolic. Each [NADP\(+\)](#)-dependent isozyme is a homodimer. The protein encoded by the [IDH2](#) gene is the [NADP\(+\)](#)-dependent isocitrate dehydrogenase found in the mitochondria. It plays a role in intermediary metabolism and energy production. This protein may tightly associate or interact with the pyruvate dehydrogenase complex.

Somatic mosaic mutations of this gene have also been found associated to Ollier disease and [Maffucci syndrome](#).

[IDH1](#) Arg132 mutations and [IDH2](#) Arg140 and Arg172 mutations accounting for >90% of aberrations ¹⁾ ²⁾. [IDH1](#) and [IDH2](#) mutations reduce the enzymatic capacity of these proteins to bind [isocitrate](#), their substrate, and convert it into [alpha-ketoglutaric acid](#) (α -KG), generating [carbon dioxide](#) and replenishing [NADH](#) and [NADPH](#) as side products ³⁾. This is one of the irreversible steps in the [tricarboxylic acid](#) cycle important for cellular respiration. Mutant [IDH1](#) (cytoplasmic) and [IDH2](#) (mitochondrial) enzymes also show a modified enzymatic capacity to convert α -KG into 2-hydroxyglutarate (2-HG), a small [oncometabolite](#). Equally important, [IDH1](#) and [IDH2](#) mutations stratify individuals into molecular subtypes with distinct clinical outcomes – the mutations are associated with lower-grade [astrocytomas](#), [oligodendrogliomas](#) (grade II/III) and secondary gliomas with better [overall survival](#), [progression-free survival](#) and [chemosensitivity](#) than [glioblastomas](#) that are [wild type](#) for both genes ⁴⁾ ⁵⁾ ⁶⁾.

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Yan H, Parsons DW, Jin G, McLendon R, Rasheed BA, Yuan W, Kos I, Batinic-Haberle I, Jones S, Riggins GJ, Friedman H, Friedman A, Reardon D, Herndon J, Kinzler KW, Velculescu VE, Vogelstein B, Bigner DD. [IDH1](#) and [IDH2](#) mutations in gliomas. *N Engl J Med*. 2009; 360:765–773

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