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## **MEN1** Antibody

Hurification MethodAntigen Affinity purifiedIsotypeIgGAliasmultiple endocrine neoplasia I;MEN1;MEAI;SCG2 ;Product TypePurified Rabbit Anti human PolyClonal AntibodyImmunogen SpeciesHomo sapiens (Human)Target NamesMEN1Target DetailsThis gene encodes menin, a putative tumor suppressor associated with syndrome known as multiple endocrine neoplasia type 1. In vitro studie shown menin is localized to the nucleus, possesses two functional nucl localization signals, and inhibits transcriptional activation by JunD, how function of this protein is not known. Two messages have been detected northern blots but the larger message has not been characterized. Alternational participation		
Uniprot No.O00255ImmunogenHuman MEN1Raised InRabbitSpecies ReactivityHuman,Mouse,RatTested ApplicationsELISA,WBStorage BufferPBS with 0.02% Sodium Azide, 50% Glycerol, pH 7.320°C, Avoid free thaw cycles.Purification MethodAntigen Affinity purifiedIsotypeIgGAliasmultiple endocrine neoplasia I;MEN1;MEAI;SCG2 ;Product TypePurified Rabbit Anti human PolyClonal AntibodyImmunogen SpeciesMEN1Target DetailsThis gene encodes menin, a putative tumor suppressor associated with syndrome known as multiple endocrine neoplasia type 1. In vitro studie shown menin is localized to the nucleus, possesses two functional nucl localization signals, and inhibits transcriptional activation by JunD, how function of this protein is not known. Two messages have been detected northern blots but the larger message has not been characterized. Altern	Product Code	CSB-PA013694GA01HU
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splicing results in multiple transcript variants.	Target Details	This gene encodes menin, a putative tumor suppressor associated with a syndrome known as multiple endocrine neoplasia type 1. In vitro studies have shown menin is localized to the nucleus, possesses two functional nuclear localization signals, and inhibits transcriptional activation by JunD, however, the function of this protein is not known. Two messages have been detected on northern blots but the larger message has not been characterized. Alternative splicing results in multiple transcript variants.

1