

Product datasheet for TP527370

OriGene Technologies, Inc.

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Twist1 (NM_011658) Mouse Recombinant Protein

Product data:

Product Type: Recombinant Proteins

Description: Purified recombinant protein of Mouse twist basic helix-loop-helix transcription factor 1 (Twist1),

with C-terminal MYC/DDK tag, expressed in HEK293T cells, 20ug

Species: Mouse

Expression Host: HEK293T

Expression cDNA

>MR227370 representing NM_011658

Red=Cloning site Green=Tags(s)

Clone or AA Sequence:

MMQDVSSSPVSPADDSLSNSEEEPDRQQPASGKRGARKRRSSRRSAGGSAGPGGATGGGIGGGDEPGSPA

QGKRGKKSAGGGGGGGGGGGGGGGSSSGGGSPQSYEELQTQRVMANVRERQRTQSLNEAFAALRKIIP

TLPSDKLSKIQTLKLAARYIDFLYQVLQSDELDSKMASCSYVAHERLSYAFSVWRMEGAWSMSASH

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Tag: C-MYC/DDK

Predicted MW: 21.6 kDa

Concentration: >0.05 μg/μL as determined by microplate BCA method

Purity: > 80% as determined by SDS-PAGE and Coomassie blue staining

Buffer: 25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol

Note: For testing in cell culture applications, please filter before use. Note that you may experience

some loss of protein during the filtration process.

Storage: Store at -80°C after receiving vials.

Stability: Stable for 12 months from the date of receipt of the product under proper storage and handling

conditions. Avoid repeated freeze-thaw cycles.

RefSeq: NP 035788

 Locus ID:
 22160

 UniProt ID:
 P26687

 RefSeq Size:
 1665

Cytogenetics: 12 14.81 cM





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RefSeq ORF: 618

Synonyms: bHLHa; bHLHa38; M-Twi; M-Twist; pd; Pde; pdt; Pluri; Ska; Ska10; Ska Twist

Summary: Basic helix-loop-helix (bHLH) transcription factors have been implicated in cell lineage

determination and differentiation. This gene encodes a bHLH transcription factor that is evolutionarily conserved from invertebrates to humans, and was originally identified in Drosophila as an essential gene involved in early mesoderm development and dorsal-ventral patterning in the embryo. This protein plays a role in cancer by regulating the epithelial-mesenchymal transition (EMT), a process that is critical for metastasis initiation, and promoting tumor progression. Mutations in the human gene are associated with Saethre-Chotzen

syndrome (SCS). Mice with heterozygous mutations in this gene exhibit cranofacial and structural

defects similar to those seen in human SCS patients. [provided by RefSeq, Sep 2015]