

## Product datasheet for TP507296

## OriGene Technologies, Inc.

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## Pax8 (NM 011040) Mouse Recombinant Protein

**Product data:** 

**Expression Host:** 

**Product Type: Recombinant Proteins** 

Description: Purified recombinant protein of Mouse paired box 8 (Pax8), with C-terminal MYC/DDK tag,

expressed in HEK293T cells, 20ug

Species: Mouse

**Expression cDNA Clone** 

>MR207296 protein sequence or AA Sequence: Red=Cloning site Green=Tags(s)

HEK293T

MPHNSIRSGHGGLNQLGGAFVNGRPLPEVVRQRIVDLAHQGVRPCDISRQLRVSHGCVSKILGRYYETGS IRPGVIGGSKPKVATPKVVEKIGDYKRQNPTMFAWEIRDRLLAEGVCDNDTVPSVSSINRIIRTKVQQPF NLPMDSCVATKSLSPGHTLIPSSAVTPPESPQSDSLGSTYSINGLLGIAQPGNDNKRKMDDSDQDSCRLS IDSQSSSSGPRKHLRTDTFSQHHLEALECPFERQHYPEAYASPSHTKGEQGLYPLPLLNSALDDGKATLT SSNTPLGRNLSTHQTYPVVADPHSPFAIKQETPELSSSSSTPSSLSSSAFLDLQQVGSGGPAGASVPPFN AFPHAASVYGQFTGQALLSGREMVGPTLPGYPPHIPTSGQGSYASSAIAGMVAGSEYSGNAYSHTPYSSY

SEAWRFPNSSLLSSPYYYSSTSRPSAPPTSATAFDHL

**TRTRPLEQKLISEEDLAANDILDYKDDDDKV** 

C-MYC/DDK Tag: Predicted MW: 48.8 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.

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

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 035170

Locus ID: 18510





## Pax8 (NM\_011040) Mouse Recombinant Protein - TP507296

UniProt ID: Q00288

RefSeq Size: 2569

Cytogenetics: 2 16.43 cM

RefSeq ORF: 1374

Synonyms: Pax; Pax-8

**Summary:** This gene encodes a member of a family of transcription factors that contain a characteristic

N-terminal paired DNA-binding domain. The encoded protein is important for proper differentiation of the thyroid and the kidney. Alternatively spliced transcript variants of this gene have been described, but their full-length nature is not known. [provided by RefSeq, Mar

2013]