

Product datasheet for TP520912

Neil2 (NM_201610) Mouse Recombinant Protein

Product data:

Product Type:	Recombinant Proteins
Description:	Purified recombinant protein of Mouse nei like 2 (E. coli) (Neil2), with C-terminal MYC/DDK tag, expressed in HEK293T cells, 20ug
Species:	Mouse
Expression Host:	HEK293T
Expression cDNA Clone or AA Sequence:	>MR220912 representing NM_201610 Red =Cloning site Green =Tags(s)

MPEGPSVRKFHHHLVSPFVGQKVVKTGGSSKKLHPPAFQSLWLQDAQVHGKFLRFDPDEEMEPLNSSPQ
PIQGMWQKEAVDRELALGPSAQEPSAGPSGSGEPVPSRSAETYNLGKIPSADAQRWLEVRFLGFGSIWVN
DFSRAKKANKKGDWRDPVPRVLVHFGGGFLVFNQCMSWSPPPVIEPTCDILSEKFHRGQALEALSQAQ
PVCYTLLDQRYFSGLGNIKNEALYRARIHPLSLGSCSSSSREALVDHVVFEFSKDWLRDKFQGKERHTQ
IYQKEQCPSGHQVMKETFGPPDGLQRLTWWCPQCQPQLSSKGPQNLPS

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Tag:	C-MYC/DDK
Predicted MW:	36.8 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.
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_963904
Locus ID:	382913
UniProt ID:	Q6R2P8 , Q1LZM6



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RefSeq Size: 1914

Cytogenetics: 14 D1

RefSeq ORF: 987

Synonyms: Gm1212; NEH2

Summary: Involved in base excision repair of DNA damaged by oxidation or by mutagenic agents. Has DNA glycosylase activity towards 5-hydroxyuracil and other oxidized derivatives of cytosine with a preference for mismatched double-stranded DNA (DNA bubbles). Has low or no DNA glycosylase activity towards thymine glycol, 2-hydroxyadenine, hypoxanthine and 8-oxoguanine. Has AP (apurinic/aprimidinic) lyase activity and introduces nicks in the DNA strand. Cleaves the DNA backbone by beta-delta elimination to generate a single-strand break at the site of the removed base with both 3'- and 5'-phosphates (By similarity).
[UniProtKB/Swiss-Prot Function]