

Recombinant Human PRD protein ab185411

Description	
Product name	Recombinant Human PRD protein
Purity	> 95 % SDS-PAGE. ab185411 is >95% pure, as determined by SEC-HPLC and reducing SDS-PAGE. Supplied as an 0.2 µM filtered solution.
Endotoxin level	< 1.000 Eu/µg
Expression system	Escherichia coli
Accession	<u>P12955</u>
Protein length	Full length protein
Animal free	No
Nature	Recombinant
Species	Human
Sequence	AAATGPSFWLGNETLKVPLALFALNRQRLCERLRKNPAV QAGSIVLQGG EETQRYCTDTGVLFRQESFFHWAFGVTEPGCYGVIDVDT GKSTLFPRLP ASHATWMGKIHSKEHFKEKYAVDDVQYVDEIASVLTSQKP SVLLTLRGVN TDSGSVCREASFDGISKFEVNNTILHPEMECRVFKTDMEL EVLRYTNKI SSEAHREVMKAVKVG MKEYELES LFEHYCYSRGGMRHS SYTCICGSGENS AVLHYGHAGAPNDRTIQNGDMCLFDMGGEYYCFASDITCS FPANGKFTAD QKAVYEAVLRSSRAVMGAMKPGVWW PDMHRLADRIHLEELAHMGILSGSVDAMVQAHLGAVFMP HGLGHFLGIDV HDVGGYPEGVERIDEPLRSLRTARHLQPGMVL TVEPGIY FIDHLLDEAL ADPARASFFNREVLQRFRGFGGVRIEEDVVVTDSGIELLT CVPRTVEEIE ACMAGCDKAFTPFSGPK
Predicted molecular weight	54 kDa
Amino acids	2 to 493
Additional sequence information	Mature protein

Specifications

Our **Abpromise guarantee** covers the use of **ab185411** in the following tested applications.

The application notes include recommended starting dilutions; optimal dilutions/concentrations should be determined by the end user.

Applications	SDS-PAGE HPLC
Form	Liquid
Additional notes	This product was previously labelled as PEPD

Preparation and Storage

Stability and Storage	Shipped on Dry Ice. Store at -20°C or -80°C. Avoid freeze / thaw cycle. pH: 7.40 Constituent: 100% PBS
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General Info

Function	Splits dipeptides with a prolyl or hydroxyprolyl residue in the C-terminal position. Plays an important role in collagen metabolism because the high level of iminoacids in collagen.
Involvement in disease	Defects in PEPD are a cause of prolydase deficiency (PD) [MIM:170100]. Prolidase deficiency is an autosomal recessive disorder associated with iminodipeptiduria. The clinical phenotype includes skin ulcers, mental retardation, recurrent infections, and a characteristic facies. These features, however are incompletely penetrant and highly variable in both age of onset and severity. There is a tight linkage between the polymorphisms of prolidase and the myotonic dystrophy trait.
Sequence similarities	Belongs to the peptidase M24B family. Eukaryotic-type prolidase subfamily.

Please note: All products are "FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC PROCEDURES"

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