

## Product datasheet for PH313202

### Phospholipase C epsilon 1 (PLCE1) (NM\_016341) Human Mass Spec Standard

#### Product data:

Product Type:	Mass Spec Standards
Description:	PLCE1 MS Standard C13 and N15-labeled recombinant protein (NP_057425)
Species:	Human
Expression Host:	HEK293
Expression cDNA Clone or AA Sequence:	RC213202
Predicted MW:	259.2 kDa



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**Protein Sequence:** >RC213202 representing NM\_016341  
 Red=Cloning site Green=Tags(s)

MTSEEMTASVLIPTVQRKVVSAQSAADESSEKVSINISKAHTVRRSGETSHTISQLNKLKEEPSGSNLP  
 KILSIAREKIVSDENSNEKCEWIKIMPDSAKNLNINCNNILRNHQHGLPQRQFYEMYNSVAEEDLCLETGI  
 PSPLERKVFPGIQLELDPRSMGISPLGNQSVIIETGRAHPDSRRVAFHFHYEVDRRMSDTFCTLSENIL  
 DDCGNCVPLPGGEEKQKKNYVAYTCKLMELAKNCDNKNEQLQCDHCDTLNDKYFCFEGSCEKVDVMVYSGD  
 SFCRKDFDTSQAAKTFLSHFEDFPDNCDDVEEDAFKSKKERSTLLVRRFCCKNDREVKKSIVYTGTRAIVRT  
 LPSGHIGLTAWSYIDQKRNGLPLPCGRVMEPPSTVEIRQDGSQRLSEAQWYPIYNAVRREETENTVGSLL  
 HFLTKLPASETAHGRISVGPCLKQCVRDVTCEYRATLQRTSISQYITGSLLLEATTSLGARSGLLSTFGGS  
 TGRMMLKERQPGPSVANSNALPSSSAGISKEIDLQPLIQFPEEVASILMEQEQTIIYRRVLPVDYLCLFT  
 RDLGTPECQSSLPCLKASISASILTQNGEHNLEDLVMRFNEVSSWVTWILITAGSMEEKREVFSYLH  
 VAKCCWNMGYNVMEFLAGLRSRVLKMWQFMDQSDIETMRSLKDAMAQHESSCEYRKYVTRALHIPGC  
 KVVPFVCGVFLKELCEVLDGASGLMKLCPRYNSQEETLEFVADYSGQDNFLQRVGQNLKNSKESTVNSI  
 FQVIRSCNRSLETDDEEDSPSEGNSSRKSLLKDKSRWQFIIGDLLSDNDIFEQSKEYDSHGSEDSQKAFD  
 HGTELIPWYVLSIQADVHQFLQGATVIHYDQDTHLSARCFLLQPDNSTLTWVKPTTASPASSKAKLGV  
 LNNTAEPGKFPLLGNAGLSSLTEGVLDFAVKAVYMGHPGIDIHTVVCVQNKLGSMFLSETGVTLTYGLQT  
 TDNRLLHFVAPKHTAKMLFSGLLELTRAVERKMRKFPDQRQWLRKQYVSLYQEDGRYEGPTLAHVELFG  
 GRRWSARNPSPTSAKNAEKPNMQRNNTLGIISTTKKKKILMRGESGEVTDDEMATRKAKMHKECRSRG  
 SDPQDINEQEESEVNAIANPPNPLPSRRAHSLTTAGSNLAAGTSSPIRVSPPVSSSNKSPSSAWSS  
 SWHGRIKGGMKGFQSFVSDSNMSFVEFVELFKSFSVRSRKLKDLFDVYAVPCNRSGSESAPLYNLTI  
 DENTSDLQPDLLTRNVDLGLFIKSKQQLSDNQRQISDAIAAASIVTNGTGIESTSLGIFGVGILQLN  
 DFLVNCQGEHCTYDEILSIIQKFEPSISMCHQGLMSFEGFARFLMDKENFASKNDESQENIKELQLPLSY  
 YYIESSHNTYLTGHQLKGESSVELYSQVLLQGCRSVELDCWDGDDGMPIIYHGHLLTTKIPFKEVVEAID  
 RSAFINSDLPISIIENHCSLPQQRKMAEIFKTVFGEKLVTKFLFETDFSDPMLPSPDQLRKKVLLKKNK  
 KLKAHQTPVDILKQKAHQQLASMVQVQAYNGGNANPRPANNEEEDEEYDYDESLSDDNILEDPRPENKS  
 CNDKLFQFEYNEEIPKRIKKADNSACNKGKVVYDMELGEEFYLDQNKESRQIAPESDLVIYCAVQKFPGL  
 STLNASGSSRGKERSRKSIFGNPNRMSPGETASFNKTSGKSSCEGIRQTWEESSSPLNPTTSLSAIIR  
 TPKCYHISLNAAKRLCRRYSQKLQHTACQLLRTYPAATRIDSNNPMLFWLHGIQLVALNYQTD  
 LPLHLNAAMFEANGCGYVLPKPPVLDKNCMPYQKFSPLERDLSDMPAVVSLTIVSGQNVCPNSMGSP  
 CIEVDVLMPLDSCHFRKPIHRNLTNPMWNEQFLFHVHFDLVFLRFVAVVENNSSAVTAQRRIIPLKALK  
 RGYRHLQLRNLHNEVLEISSLFINSRRMEENSSGNTMSASSMFNTEERKCLQTHRVTVHGVPGPEPFTVF  
 TINGGTAKAKQLQQLTNEQDIKPVTTDYFLMEEKYFISKEKNECRKQPFQRAIGPEEIMQILSSWFPE  
 EGYMGRIVLKTQQENLEEKNIVQDDKEVILSSEEEFFVQVHDVSPEQPRTVIKAPRVSTAQDVIQQTLC  
 KAKYSYSILSNPNPSDYVLLLEEVKDTTNNKTTTPKSSQVLLDQECVFAQSKWKAGAGFILKLKEQVQ  
 ASREDKKKGISFASLKKLTKSTKQPRGLTSPSQLLTSESIQTKEEKPVGGLSSSDTMDYRQ

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

**Tag:** C-Myc/DDK

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

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

**Labeling Method:** Labeled with [U- <sup>13</sup>C<sub>6</sub>, <sup>15</sup>N<sub>4</sub>]-L-Arginine and [U- <sup>13</sup>C<sub>6</sub>, <sup>15</sup>N<sub>2</sub>]-L-Lysine

**Buffer:** 25 mM Tris-HCl, 100 mM glycine, pH 7.3

**Storage:** Store at -80°C. Avoid repeated freeze-thaw cycles.

**Stability:** Stable for 3 months from receipt of products under proper storage and handling conditions.

**RefSeq:** [NP\\_057425](#)

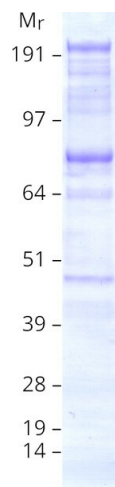
RefSeq Size:	7992
RefSeq ORF:	6906
Synonyms:	NPHS3; PLCE; PPLC
Locus ID:	51196
UniProt ID:	<a href="#">Q9P212</a>
Cytogenetics:	10q23.33

**Summary:** This gene encodes a phospholipase enzyme that catalyzes the hydrolysis of phosphatidylinositol-4,5-bisphosphate to generate two second messengers: inositol 1,4,5-triphosphate (IP3) and diacylglycerol (DAG). These second messengers subsequently regulate various processes affecting cell growth, differentiation, and gene expression. This enzyme is regulated by small monomeric GTPases of the Ras and Rho families and by heterotrimeric G proteins. In addition to its phospholipase C catalytic activity, this enzyme has an N-terminal domain with guanine nucleotide exchange (GEF) activity. Mutations in this gene cause early-onset nephrotic syndrome; characterized by proteinuria, edema, and diffuse mesangial sclerosis or focal and segmental glomerulosclerosis. Alternative splicing results in multiple transcript variants encoding distinct isoforms.[provided by RefSeq, Sep 2009]

**Protein Families:** Druggable Genome

**Protein Pathways:** Calcium signaling pathway, Inositol phosphate metabolism, Metabolic pathways, Phosphatidylinositol signaling system

### Product images:



Coomassie blue staining of purified PLCE1 protein (Cat# [TP313202]). The protein was produced from HEK293T cells transfected with PLCE1 cDNA clone (Cat# [RC213202]) using MegaTran 2.0 (Cat# [TT210002]).