

## Product datasheet for **RG235348**

### PTPRB (NM\_001206972) Human Tagged ORF Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** PTPRB (NM\_001206972) Human Tagged ORF Clone  
**Tag:** TurboGFP  
**Symbol:** PTPRB  
**Synonyms:** HPTP-BETA; HPTPB; PTPB; R-PTP-BETA; VEPTP  
**Mammalian Cell Selection:** Neomycin  
**Vector:** pCMV6-AC-GFP (PS100010)  
**E. coli Selection:** Ampicillin (100 ug/mL)  
**ORF Nucleotide Sequence:** >RG235348 representing NM\_001206972  
Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGATCGCC**

ATGCTGAGCCATGGAGCCGGTTGGCCTTGTTGGATCACACTGAGCCTGCTGCAGACTGGACTGGCGGAGC  
CAGAGAGATGTAACCTCACCTGGCGGAGTCCAAGGCCTCCAGCCATTCTGTGTCTATCCAGTGGAGAAT  
TTTGGGCTCACCTGTAACCTTAGCCTCATCTATAGCAGTGACACCCTGGGGCCCGTTGTGCCCTACC  
TTTCGGATAGACAACACCACATACGGATGAACCTTCAAGATTTACAAGCAGGAACCATCTATAACTTCA  
GGATTATTTCTCTGGATGAAGAGAGAACAGTGGTCTTGCAAACAGATCCTTTACCTCCTGCTAGGTTTGG  
AGTCAGTAAAGAGAAGACGACTTCAACCAGCTTGCATGTTTGGTGGACTCCTTCTCCGAAAAGTCAAC  
TCATATGAGGTGCAATTATTTGATGAAAATAACAAAAGATACAGGGGTTCAAATTCAGAAAAGTACTT  
CATGGAATGAATACACTTTTTTCAATCTCACTGCTGGTAGTAAATACAATATTGCCATCACAGCTGTTTC  
TGGAGGAAAACGTTCTTTTTAGTTTATACCAATGGATCAACAGTGCCATCTCCAGTGAAGATATTGGT  
ATTTCCACAAAAGCCAATTCTCTCCTGATTTCTGGTCCCATGGTTCTGGGAATGTGGAACGATACCGGC  
TGATGCTAATGGATAAAGGGATCCTAGTTCATGGCGGTGTTGTGGACAAACATGCTACTTCTATGCTTT  
TCACGGGCTGACCCCTGGCTACCTCTACAACCTCACTGTTATGACTGAGGCTGCAGGGCTGCAAAACTAC  
AGGTGAAAACAGTACAGGACAGCCCCATGGAAGTCTCAAATCTGAAGGTGACAAATGATGCGAGTTTGA  
CCTCTCTAAAAGTCAAATGGCAAAGACCTCCTGAAAATGTGGATTCTTACAATATCACCTGTCTCACAA  
AGGGACCATCAAGGAATCCAGAGTATTAGCACCTTGGATTACTGAACTCACTTTAAAGAGTTAGTCCCC  
GGTCGACTTTATCAAGTTACTGTCAGCTGTGTCTCTGGTGAAGTCTGCTCAGAAGATGGCAGTGGGCA  
GAACATTTCCAGACAAAGTTGCAAACCTGGAGGCAACAATAATGGCAGGATGAGGTCTCTTGTAGTGAG  
CTGGTCGCCCCCTGCTGGAGACTGGGAGCAGTATCGGATCCTACTCTTCAATGATTCTGTGGTGTCTC  
AACACTACTGTGGAAAGGAAGAAACACAGTATGTCATGGATGACACGGGCTCGTACCGGAAGACAGT  
ATGAGGTGGAAGTCATTGTTGAGAGTGGAAATTTGAAGAATTCTGAGCGTTGCCAAGGCAGGACAGTCCC  
CCTGGCTGCTCCAGCTTCGTGTCAAACATGCCAATGAAACCTCACTGAGTATCATGTGCAGACCCCT



[View online »](#)

GTAGCAGAATGGGAGAAATACATCATTTCCCTAGCTGACAGAGACCTCTTACTGATCCACAAGTCACTCT  
 CCAAAGATGCCAAAGAATTCACITTTTACTGACCTGGTGCCTGGACGAAAAACATGGCTACAGTCAACAG  
 TATTAGTGGAGACTTAAAAAATTCCTCTTCAGTAAAAGGAAGAACAGTGCCTGCCAAAGTACTGACTTG  
 CATGTGGCCAACCAAGGAATGACCAGTAGTCTGTTTACTAACTGGACCCAGGCACAAGGAGACGTAGAAT  
 TTTACCAAGTCTTACTGATCCATGAAAAATGTGGTCATTAATAAATGAAAGCATCTCCAGTGAGACCAGCAG  
 ATACAGCTTCCACTCTCAAGTCCGGCAGCCTGTACTCCGTGGTGAACAACAGTGAAGTGGAGGGGATC  
 TCTTCCCGACAAGTGGTGTGGAGGGAAGAACAGTCCCTCCAGTGTGAGTGGAGTAACGGTGAACAAAT  
 CCGGTGTAATGACTACCTCAGCGTTTCCCTGGCTGCTGGCGCCGGAGATGTGGATAACTATGAGGTAAC  
 ATTGTCTCATGACGGCAAGGTGGTTCAGTCCCTTGTGATTGCCAAGTCTGTCAGAGAATGTTCCCTCAGC  
 TCCCTCACCCAGGCCGCTCTACACCGTGACCATAACTACAAGGAGTGGCAAGTATGAAAACTACTCCT  
 TCAGCCAAGAGCGGACAGTGCCTGACAAAGTCCAGGGAGTCAAGTGTAGCAACTCAGCCAGGAGTGACTA  
 TTTAAGGGTATCCTGGGTGCATGCCACTGGAGACTTTGATCACTATGAAGTCAACATTAATAAATAA  
 AACTTCATTCAAATAAAAGCATTCCCAAGTCAGAAAACGAATGTGATTTGTTTCACTAGTCCCTGGAC  
 GGTGTACAGTGTCACTGTTACTACAAAAGTGGACAATATGAAGCCAATGAACAAGGGAAATGGGAGAAC  
 AATTCCAGAGCCTGTTAAGGATCTAACATTGCGCAACAGGAGCACTGAGGACTTGCATGTGACTTGGTCA  
 GGAGCTAATGGGGATGTGACCAATATGAGATCCAGCTGCTTCAATGACATGAAAGTATTTCCCTCCTT  
 TTCACCTTGTAATAACCGCAACCGAGTATCGATTTACTTCCCTAACACCAGGCCGCAATACAAAATTTCT  
 TGTCTTGACGATTAGCGGGGATGTACAGCAGTCAAGCTTCACTGAGGGCTTACAGTTCCTAGTGTGTC  
 AAAAAATTTACATTTCTCCCAATGGAGCAACAGATAGCCTGACGGTGAAGTGGACTCCTGGTGGGGGAG  
 ACGTTGATTTCTACACGGTGTGGCATTTCAGGCACAGTCAAAGGTTGACTCTCAGACTATTTCCAAAGCA  
 CGTCTTTGAGCACAGTTCACAGACTGGAGGCCGGGGAGCAGTACCAGATCATGATTGCCTCAGTCAGC  
 GGGTCCCTGAAGAATCAGATAAATGTGGTTGGGCGGACAGTCCCAGCAGCTGTACCAGACTGAGGATCA  
 CAGAGAACTCCACCAGGCACCTGTCTTCCGCTGGACCGCTCAGAGGGGGAGCTCAGCTGTACAACAT  
 CTTTTTGTACAACCCAGATGGGAATCTCCAGGAGAGAGCTCAAGTTGACCCACTAGTCCAGAGCTTCTCT  
 TTCAGAACTTGCTACAAGGCAGAAATGTACAAGATGGTATTGTAACCTCAGAGTGGGAGCTGTCTAATG  
 AGTCTTTTCAATTTGGTAGAACAGTCCCAGCCTCTGTGAGTCACTCAGGGGGTCCAATCGGAACACGAC  
 AGACAGCCTTTGGTTCAACTGGAGTCCAGCCTCTGGGGACTTTGACTTTTATGAGCTGATTCTCTATAAT  
 CCCAATGGCACAAGAAGGAAAATGGAAGACAAGGACCTGACGGAGTGGCGGTTTCAAGGCCCTTGTTT  
 CTGGAAGGAAGTACGTGCTGTGGTGGTAACTCACAGTGGAGATCTCAGCAATAAAGTACAGCGGAGAG  
 CAGAACAGCTCCAAGTCCCTCCAGTCTTATGTCATTTGCTGACATTGCAAACACATCCTTGCCATCAG  
 TGGAAAGGGCCCCAGACTGGACAGACTACAACGACTTTGAGCTGCAGTGGTTGCCAGAGATGCACTTA  
 CTGCTTCAACCCCTACAACAACAGAAAATCAGAAGGACGCAATTGTGTATGGTCTTCGTCCAGGGAGATC  
 CTATCAATTCAACGTCAAGACTGTCAAGTGGTATTCTGGAAAATTTACAGCAAACCAATTTTTGGATCT  
 GTGAGGACAAAGCCTGACAAGATACAAAACCTGCATTGCCGGCCTCAGAACTCCACGGCCATTGCCTGTT  
 CTTGGATCCCTCCTGATTCTGACTTTGATGGTTATAGTATTGAATGCCGGAAAAATGGACACCCAAGAAGT  
 TGAGTTTTCCAGAAAGCTGGAGAAAGAAAAATCTCTGCTCAACATCATGATGCTAGTGCCCCATAAGAGG  
 TACCTGGTGTCCATCAAAGTGCAGTCCGGCCGCATGACCAGCGAGGTGGTTGAAGACAGCACTATCAAA  
 TGATAGACCGCCCCCTCCTCCACCCCAACACATTGCGTGAATGAAAAGGATGTGCTAATTAGCAAAGTC  
 TTCCATCAACTTTACTGTCAACTGCAGTGGTTCAGCGACACCAATGGAGCTGTGAAATACTTTCACAGT  
 GTGGTGAGAGAGGCTGATGGCAGTGTGAGCTGAAGCCAGAACAGCAGCACCCCTCCTCCTTCTACCTGG  
 AGTACAGGCACAATGCCTCCATTCCGGTGTATCAGACTAATTATTTTGGCAGCAAATGTGCCGAAAAATCC  
 TAACAGCAACTCCAAGAGTTTTAACATTAAGCTTGGAGCAGAGATGGAGAGCCTAGGTGGAAAAATGCGAT  
 CCCACTCAGCAAAAAATCTGTGATGGACCACTGAAGCCACACACTGCCTACAGAATCAGCATTGAGCTT  
 TTACACAGCTCTTTGATGAGGACCTGAAGGAATTCACAAAGCCACTCTATTACAGACATTTTTTTCTTT  
 ACCCATCACTACTGAATCAGAGCCCTTGTGGAGCTATTGAAGGTGTGAGTGTGGTCTGTTTTTAATT  
 GGCATGCTAGTGGCTGTGTGGCTTATTGATCTGCAGACAGAAAGTGAAGCATGGTTCGAGAAAGACCCCT  
 CTGCCCTGCTGAGCATTCTGAGGATCGACCATTATCTGTCCACTTAAACCTGGGCCAGAAAGGTAACCG  
 GAAAACTTCTGTCCAATAAAAAATAAATCAGTTTGAAGGGCATTTCATGAAGCTACAGGCTGACTCCAAC  
 TACCTTCTATCCAAGGAATACGAGGAGTTAAAAGACGTGGGCCGAAACAGTCAATGTGACATTGCACTCT  
 TGCCGGAGAATAGAGGGAAAAATCGATACAACAATATATTGCCCTATGATGCCACGCGAGTGAAGCTCTC  
 CAATGTAGATGATGATCCTTGTCTGACTACATCAATGCCAGCTACATCCCTGGCAACAACCTTCAGAAGA  
 GAATACATTGTCACTCAGGGACCGCTTCTGGCACCAAGGATGACTTCTGGAAAAATGGTGTGGGAACAAA

ACGTTACAACATCGTCATGGTGACCCAGTGTGTTGAGAAGGGCCGAGTAAAGTGTGACCATTACTGGCC  
 AGCGGACCAGGATTCCCTCTACTATGGGGACCTCATCCTGCAGATGCTCTCAGAGTCCGCTGCCTGAG  
 TGGACCATCCGGGAGTTAAGATATGCGGTGAGGAACAGCTTGTGCACACAGACTCATCCGCCACTTTC  
 ACTATACGGTGTGGCCAGACCATGGAGTCCCAGAAACCACCCAGTCTCTGATCCAGTTTGTGAGAACTGT  
 CAGGGACTACATCAACAGAAGCCCGGTGCTGGGCCACTGTGGTGCAGTGCAGTGTGGTGGTGGTGGT  
 ACTGGAACCTTTATTGCATTGGACCGAATCCTCCAGCAGTTAGACTCAAAGACTCTGTGGACATTTATG  
 GAGCAGTGCACGACCTAAGACTTCACAGGGTTACATGGTCCAGACTGAGTGTGAGTGTGATGTCTACCTACA  
 TCAGTGTGTAAGAGATGCTCTCAGAGCAAGAAAGCTACGGAGTGAACAAGAAAACCCCTGTTTCCAATC  
 TATGAAAATGTGAATCCAGAGTATCACAGAGATCCAGTCTATTCAAGGCAT

ACGCGTACGCGGCCGCTCGAG - GFP Tag - GTTTAA

**Protein Sequence:**

>RG235348 representing NM\_001206972  
 Red=Cloning site Green=Tags(s)

MLSHGAGLALWITLSLLQTGLAEPERCNFTLAESKASSHSVSIQWRILGSPCNFLIYSSDTLGAALCPT  
 FRIDNTTYGCNLQDLQAGTIYNFRIISLDEERTVVLQTDPLPPARFGVSKEKTTSTSLHVVWTPSSGKVT  
 SYEVQLFDENNQIQGVQIQESTSWNEYTFNLTAGSKYNIATAVSGGKRSFSVYTNSTVPSPVKDIG  
 ISTKANSLLISWSHSGSNVERYRLMLMDKGILVHGGVVDKHAISYAFHGLTPGYLYNLVMTAAGLQNY  
 RWKLVRTAPMEVSNLKVNDGSLTSLKVKWQRPVGNVDSYNITLSHKGTIKESRVLAPWITETHFKELVP  
 GRLYQVTVSCVSGELSAQKMAVGRTFPDKVANLEANNGRMRSLLVSWSPAPGDWEQYRILLFNDSVLL  
 NITVIGKEETQYVMDTGLVPGRQYEVVIVE SGNLKNSERCQGRVPLAVLQLRVKHANETSLSIMWQTP  
 VAEWEKYIISLADRDLILLHKSLSKDAKEFTFDLVPGRKYMATVTSISGDLKNSSSVKGRVPAQVTDL  
 HVANQGMTSSLFTNWTQAQGDVEFYQVLLIHENVVIKNEISSSETSRYSFHSLKSGSLYSVVVTTVSGGI  
 SSRQVVVEGRTVPSSVSGVTVNNSGRNDYLSVSWLLAPGDVDNYEVTLSHDGKVVQSLVIAKSVRECSFS  
 SLTPGRLYTVTITTRSGKYENHSFSQERTVPDKVQGVSVNSARSVDYLRVSWVHATGDFDHYEVTIKNKN  
 NFIQTKSIPKSENECVFVQLVPGRLYSVTVTTKSGQYEAENEQNGRTIPEPVKDLTLRNRSTEDLHVTWS  
 GANGVDVQYEQIQLLFNDMKVFPFHLVNTATEYRFTSLTPGRQYKILVLTISGDVQQSAFIEGFTVPSAV  
 KNIHISPNGATDSLTVNWTPGGGDVDSYTVSAFRHSQKVDSTIPKHVFEHTFHRLEAGEQYQIMIASVS  
 GSLKNQINVVGRTPAAVTDLRITENSTRHL SFRWTA SEGELSWYNI FL YNPDGNLQERAQVDPLVQSFS  
 FQNLQGRMYKMVIVTHSGELSNESFIFGRTVPASVSHLRGSRNRTTDSLWFWNSPASGDFDFYELILYN  
 PNGTKKENWKDKDLTEWRFQGLVPGRKYVLLVWVTHSGDL SNKVTAESRTAPSPPLMSFADIANTSLAIT  
 WKGPPDWDYNDFELQWLPRDALTVFNPNRNRKSEGRIVYGLRPGRSYQFNVKTVSGDSWKTYSKPIFGS  
 VRTKPKDIQNLHCRPQNSTAIACSWIPDSDFDGYSIECRKMDTQEVFESRKLEKEKSLNIMMLVPHKR  
 YLVSIVKVSAGMTSEVVEDSTITMIDRPPPPPHIRVNEKDVLISKSSINFTVNCVSWFSDTNGAVKYFTV  
 VVREADGSDELKPEQQHPLPSYLEYRHNASIRVYQTNFYASKCAENPNSNSKSFNIKLAGEMESLGGKCD  
 PTQQKFCDGPLKPHYAYRISIRAF TQLFDEDLKEFTKPLYSDTFFSLPITTESEPLFGAIEGVSAGLFLI  
 GMLVAVVALLICRQKVSHGRERPSARLSIRRRDRPLSVHLNLGQKGNRKTSCPIKINQFEGHFMKLQADSN  
 YLLSKEYEELKDVGRNQSCDIALLPENRGNRYNNILPYDATRVKLSNVDDDPDSYINASYIPGNFRFR  
 EYIVTQGPLPGTKDDFWKMWVQNVHNI VMTQCVEKGRVKCDHYWPAQDQSLYGGDLILQMLSESVLPE  
 WTIREFKICGEEQLDAHRLIRHFHYTVWPDHGVPETTQSLIQFVRTVRDYINRSPGAGPTVVHCSAGVGR  
 TGTFIALDRILQQLDSKDSVDIYGAVHDLRLHRVHMVQTECQYVYLHQCVRDVLRARKLRSEQENPLFPI  
 YENVNPEYHRDPVYSRH

TRTRPLE - GFP Tag - V

**Restriction Sites:**

Sgfl-Mlul



<b>OTI Disclaimer:</b>	The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. <a href="#">More info</a>
<b>OTI Annotation:</b>	This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.
<b>Components:</b>	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).
<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.</li></ol>
<b>RefSeq:</b>	<a href="#">NM_001206972.1</a> , <a href="#">NP_001193901.1</a>
<b>RefSeq Size:</b>	10393 bp
<b>RefSeq ORF:</b>	5724 bp
<b>Locus ID:</b>	5787
<b>UniProt ID:</b>	<a href="#">P23467</a>
<b>Cytogenetics:</b>	12q15
<b>Protein Families:</b>	Druggable Genome, Phosphatase
<b>Protein Pathways:</b>	Adherens junction
<b>Gene Summary:</b>	<p>The protein encoded by this gene is a member of the protein tyrosine phosphatase (PTP) family. PTPs are known to be signaling molecules that regulate a variety of cellular processes including cell growth, differentiation, mitotic cycle, and oncogenic transformation. This PTP contains an extracellular domain, a single transmembrane segment and one intracytoplasmic catalytic domain, thus belongs to receptor type PTP. The extracellular region of this PTP is composed of multiple fibronectin type_III repeats, which was shown to interact with neuronal receptor and cell adhesion molecules, such as contactin and tenascin C. This protein was also found to interact with sodium channels, and thus may regulate sodium channels by altering tyrosine phosphorylation status. The functions of the interaction partners of this protein implicate the roles of this PTP in cell adhesion, neurite growth, and neuronal differentiation. Alternate transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, May 2011]</p>