

Product datasheet for RN202443

Pds5a (NM_001083624) Rat Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Pds5a (NM_001083624) Rat Untagged Clone
Tag:	Tag Free
Symbol:	Pds5a
Synonyms:	RGD1307094; Scc-112
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>RN202443 representing NM_001083624 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGGACTTCACGCAGCCGAAGCCTGCCACGGCCCTCTGTGGCGTCGTGAGTGCGGACGGGAAGATCGCTT
ACCCTCCGGGAGTAAAAGAGATCACGGACAAGATCACCACCGACGAAATGATCAAACGTTTGAAGATGGT
AGTGAAAACCTTTATGGATATGGATCAGGATTCAGAAGATGAAAAACAACAGTATCTCCACTAGCCTTG
CATCTTGCATCTGAATCTTTCTCAGGAACCTAATAAAGATGTGCGCCTTCTTGAGCTTGTGTTTGG
CTGATATATTTTGAATCTATGCCCCAGAAGCTCCATACACTTCCCACGATAAACTTAAGGACATATTTCT
CTTTATTACCAGACAATTAAGGTTTGGAGGATACAAAGAGTCCACAGTTTAATAGATATTTTTATTTA
TTAGAGAAATTTAGCTTGGGTTAAATCATATAACATCTGCTTTGAATTGGAAGATTGCAATGAAATTTTTA
TTCAGCTTTTTAGGACTCTCTTCTCAGTAATCAACAATAGCCACAATAAGAAGGTACAAATGCACATGTT
AGACTTGATGAGTTCTATCATCATGGAAGGTGATGGAGTTACTCAAGAATTATTGGATTCCATTCTTATC
AATCTCATCCCTGCACACAAGAACTTAAATAAACAGTCCCTTGACCTTGCAGAAAGTTCTGTTGAAAAGGA
CAGTCCAGACTATTGAAGCATGTATTGCCAATTTTTCAATCAAGTGTGGTGTGGGCAGATCATCAGT
CAGTGATCTCTCCGAACAGTATTGATCTGATTCAGGAACTTTTGCTATAGATCTCATTATTTGTTA
TCTGTCTATGCCACAGCTTGAATTCAAACTGAAGAGCAATGACGGTGAGGAGCGGCTAGCTGTGGTGGCAG
TACTAGCAAAATTTGTTGGCTCTAAAGATTCAGATTTAGCAACACAGAATCGGCCTCTTTGGCAGTGCTT
TCTTGGGCGATTTAATGACATTCATGTTCTGTGAGGTTAGAAAGTGTGAAGTTTGCCAGTCACTGTTTA
ATGAATCACCTGATTTAGCAAAAGATCTCACAGAATATTTAAAAGTTAGGTACATGATCCGGAAGAAG
CCATTCGTATGATGTCATTGTTACTATAATAACAGCTGCCAAAAGAGACCTTGCCTTAGTAAATGATCA
GTTGCTTGGCTTTGTGAGGAAAGGACACTGGATAAACGGTGGCGAGTAAGAAAAGAAGCCATGATGGGT
CTGGCTCAACTCTATAAGAAATACTGTCTTCATGGTGAAGCAGGAAAGGAAAGCTGCAGAGAAAGTCAGTT
GGATAAAGGACAACTTCTACACATCTATTATCAGAATAGCATTGATGACAAACTGTTGGTAGAGAAAAT
CTTTGCTCAGTATCTTGTCCCATAACTGGAACAGAAAGAGAATGAAATGCTTATATTATTTATAT



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GCTAGCTTGGATCCAAATGCTGTAAAAGCTCTCAATGAAATGTGGAAGTGCAGAATATGCTTCGGAGTC
 ACGTGCGGGAAGTGTGGACTTGCAACAAGCAGCCTACATCGGAGGCTAACTGTTCTGCCATGTTGGGAA
 ACTGATGACCATAGCAAAGAATTTGCTGACCCCTGGGAAAGCACAAGATTTTGTAAAGAAATTTAACCA
 GTTCTTGGTGATGATGAAAACTGAGGTCTCAGTTGGAATTTAATCAGCCCAACCTGTTTCATGCAAGC
 AAGCTGACATTTGTGTGAGGGAATAGCTCGGAACTTGCAAATCCTAAGCAGCCAACCAACCCCTTTCT
 AGAGATGGTCAAGTTTCTGTGGAAGAATCGCTCCTGTGCACATTGATTGAGAAGCCATAAGTGCACTA
 GTGAAACTGATGAACAAGTCGATAGAAGGGACAGCCGATGATGAAGAGGAGGGTGTGTCAGCCAGACTCAG
 CCATTCGCTCAGGGCTTGAAGCTTCTTAAGGTTCTGTCTTTCACACATCCTACCTCGTTCCTACTCTGCAGA
 GACGTATGAGTCCTTGTACAGTGCCTAAGAATGGAGGACGACAAGGTAGCGGAAGCAGCAATACAGATT
 TTTAGAAACACAGGCCACAAAATAGAGACTGACCTTCCCCAGATACGGTCAACCTTGATCCCCATTTTAC
 ATCAGAAAGCCAAGCGGGGCACTCCTCACCAAGCAAAGCAGGCTGTGCACTGCATCCATGCCATCTTCTC
 AAACAAGGAAGTCCAGTTGGCACAGATTTTGGCCACTCAGTCGGAGTCTGAATGCTGATGTACCAGAG
 CAACTTAACTCCATTAGTTTCACTGGCCACATTTCCATGTTAGCACCAGATCAGTTTGCCTCCCCAA
 TGAATCTGTAGTGGCAAATTCATTGTTAAGGATCTTCTAATGAATGACAGGTCAACAGGTGAGAAGAA
 TGGAAAATATGGTCTCCGGATGAGGAAGTATCTCCTGAAGTGTAGCAAAGGTACAGGCAATTAACCTT
 CTGGTAAGGTGGCTGTTGGGTATGAAGAACAATCAGTCCAAATCTGCCAACTCAACTCTGCGGCTGTAT
 CAGCCATGCTGGTCAGTGAAGGGGACCTGACAGAGCAGAAGAGGATCAGTAAATCTGATATGTCTCGCTT
 GCGATTAGCTGCTGGTAGTGCCATAATGAAGCTTGTCTCAGGAACCTTGCTACCATGAAATATTACCCCA
 GAACAGTTTCAGCTCTGTGCACTGGTTATTAACGATGAGTGCTACCAAGTAAGGCAGATATTTGCCAGA
 AGCTTCATAAAGCCCTTGTGAAGTACTTCTCCATTGGAGTATATGGCGATCTTTGCTTTGTGTGCCAA
 AGATCCTGTGAAGGAAAGAGACATGCTCGCCAGTGTATTAAGAAGCATCAGCATACGCAGGGAG
 TACATTAACAGAACCCCATGGCTACTGAGAAGTATTATCATTGCTGCCTGAATATGTGGTTCATATA
 TGATTCACCTGTAGCCCATGATCCTGATTTTACAAGATCACAAGATGTTGATCAACTTCGTGATATAAA
 AGAGTGCCTGTGGTTTATGCTTGAAGTTTTAATGACAAAGAATGAGAACAACAGCCATGCATTCATGAAG
 AAGATGGCAGAGAATATCAAGCTGACCAGAGACGCCAGTCTCCAGATGAAGCCAAGACAATGAAAAAC
 TTTATACAGTTTGTGATGTGGCTCTGTGTGTTATAAATAGTAAGAGTGCTTTGTGCAATGCAGACTCACC
 AAAGGACCCAGTCTCCCAATGAAATCTTTACACAACCTGAAAAGGACTTCTCTAATGACAAAAGCTAT
 ATTTCCGGAAGAGACAAGAGTTCTTCTGTTGACAGGAAAGCCAAAGCCTACTGGAGTACTAGGTACAGTAA
 ACAAGCCCTTATCAGCAACGGGAAGGAAGCCTTATGTTAGGAGTGCCGGCACAGAGACTGGAAGCAATAT
 TAACGCCAGTTCAGAGCTGAGTCTTCCGCCGAAATCGTTCAAGGGAACAGAGTTCAGAGGCATCAGAA
 ACTGGAGTTAGTAAAAAGGAGAATCCTGTGAGAATAATTTCTGTACACCTGTAAGAGTATTGATA
 CGGTAAAGAATAAGGAAATTAATTTCTGATCAGTCTACCAAGGCAACATCAGCAGTGACCGAGGAAAGAA
 AAGAATTGTAACAGCAGCTGGTGTGAGAATATCCAACAAAAACCAGATGAGAAAGCAGATGAGTACGGA
 CCCCTGCCCTTCCAAACCCAGGAGAGGACGTCGCCCAAAATCTGAATCTCAGGGCAATGCAACAAAAA
 ACGATGATCTAAGTAAACCTGTTAGCAAGGGAAGGAGAGCTGCAGGCAGCCAGGAGAGTCTGGAGGC
 AGGCAATGCCAAAGCGCCCAAGCTACAAGATGGAGCCAAAAGGCAGTTCGCCCGAGAGACAAATTTGAT
 TTACAAAGGTA

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites:

SgfI-MluI

ACCN:

NM_001083624

Insert Size:

4002 bp

OTI Disclaimer:

Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).

Components:	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).
Reconstitution Method:	<ol style="list-style-type: none">1. Centrifuge at 5,000xg for 5min.2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.3. Close the tube and incubate for 10 minutes at room temperature.4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.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.
RefSeq:	<u>NM_001083624.1, NP_001077093.1</u>
RefSeq Size:	4030 bp
RefSeq ORF:	4002 bp
Locus ID:	305343
UniProt ID:	<u>A4L9P7</u>
Cytogenetics:	14p11
Gene Summary:	Probable regulator of sister chromatid cohesion in mitosis which may stabilize cohesin complex association with chromatin. May couple sister chromatid cohesion during mitosis to DNA replication. Cohesion ensures that chromosome partitioning is accurate in both meiotic and mitotic cells and plays an important role in DNA repair (By similarity).[UniProtKB/Swiss-Prot Function]