

Product datasheet for RC205715

RPA34 (RPA2) (NM_002946) Human Tagged ORF Clone

Product data:

OriGene Technologies, Inc.

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Expression Plasmids
RPA34 (RPA2) (NM_002946) Human Tagged ORF Clone
Myc-DDK
RPA34
REPA2; RP-A p32; RP-A p34; RPA32
Neomycin
pCMV6-Entry (PS100001)
Kanamycin (25 ug/mL)
<pre>>RC205715 ORF sequence Red=Cloning site Blue=ORF Green=Tags(s)</pre>
TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC GCC <mark>GCGATCGC</mark> C

AGCGGACCGACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC TGGATTACAAGGATGACGACGATAAGGTTTAA



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CRIGENE RPA34 (RPA2) (NM_002946) Human Tagged ORF Clone – RC205715

Reconstitution Method:	 Centrifuge at 5,000xg for 5min. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. Close the tube and incubate for 10 minutes at room temperature. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. 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.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	<u>NM 002946.5</u>
RefSeq Size:	1819 bp
RefSeq ORF:	813 bp
Locus ID:	6118
UniProt ID:	<u>P15927</u>
Cytogenetics:	1p35.3
Domains:	tRNA_anti
Protein Families:	Druggable Genome, Stem cell - Pluripotency
Protein Pathways:	DNA replication, Homologous recombination, Mismatch repair, Nucleotide excision repair
MW:	29.2 kDa
Gene Summary:	This gene encodes a subunit of the heterotrimeric Replication Protein A (RPA) complex, which binds to single-stranded DNA (ssDNA), forming a nucleoprotein complex that plays an important role in DNA metabolism, being involved in DNA replication, repair, recombination, telomere maintenance, and co-ordinating the cellular response to DNA damage through activation of the ataxia telangiectasia and Rad3-related protein (ATR) kinase. The RPA complex protects single-stranded DNA from nucleases, prevents formation of secondary structures that would interfere with repair, and co-ordinates the recruitment and departure

of different genome maintenance factors. The heterotrimeric complex has two different modes of ssDNA binding, a low-affinity and high-affinity mode, determined by which oligonucleotide/oligosaccharide-binding (OB) domains of the complex are utilized, and differing in the length of DNA bound. This subunit contains a single OB domain that participates in high-affinity DNA binding and also contains a winged helix domain at its

modifications of the RPA complex also plays a role in co-ordinating different damage

carboxy terminus, which interacts with many genome maintenance protein. Post-translational

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response pathways. [provided by RefSeq, Sep 2017]



Product images:



Circular map for RC205715



HEK293T cells were transfected with the pCMV6-ENTRY control (Cat# [PS100001], Left lane) or pCMV6-ENTRY RPA2 (Cat# RC205715, Right lane) cDNA for 48 hrs and lysed. Equivalent amounts of cell lysates (5 ug per lane) were separated by SDS-PAGE and immunoblotted with anti-RPA2(Cat# [TA500763]). Positive lysates [LY401031] (100ug) and [LC401031] (20ug) can be purchased separately from OriGene.

Western blot validation of overexpression lysate (Cat# [LY401031]) using anti-DDK antibody (Cat# [TA50011-100]). Left: Cell lysates from untransfected HEK293T cells; Right: Cell lysates from HEK293T cells transfected with RC205715 using transfection reagent MegaTran 2.0 (Cat# [TT210002]).

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Coomassie blue staining of purified RPA2 protein (Cat# [TP305715]). The protein was produced from HEK293T cells transfected with RPA2 cDNA clone (Cat# RC205715) using MegaTran 2.0 (Cat# [TT210002]).

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