

# **Product datasheet for PH302066**

### OriGene Technologies, Inc.

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## RPA70 (RPA1) (NM 002945) Human Mass Spec Standard

**Product data:** 

**Product Type:** Mass Spec Standards

RPA1 MS Standard C13 and N15-labeled recombinant protein (NP\_002936) **Description:** 

Species: Human **HEK293 Expression Host: Expression cDNA Clone** 

or AA Sequence:

RC202066

Predicted MW: 68.1 kDa

>RC202066 protein sequence **Protein Sequence:** 

Red=Cloning site Green=Tags(s)

MVGQLSEGAIAAIMQKGDTNIKPILQVINIRPITTGNSPPRYRLLMSDGLNTLSSFMLATQLNPLVEEEQ LSSNCVCQIHRFIVNTLKDGRRVVILMELEVLKSAEAVGVKIGNPVPYNEGLGQPQVAPPAPAASPAASS RPQPQNGSSGMGSTVSKAYGASKTFGKAAGPSLSHTSGGTQSKVVPIASLTPYQSKWTICARVTNKSQIR TWSNSRGEGKLFSLELVDESGEIRATAFNEQVDKFFPLIEVNKVYYFSKGTLKIANKQFTAVKNDYEMTF NNETSVMPCEDDHHLPTVQFDFTGIDDLENKSKDSLVDIIGICKSYEDATKITVRSNNREVAKRNIYLMD TSGKVVTATLWGEDADKFDGSRQPVLAIKGARVSDFGGRSLSVLSSSTIIANPDIPEAYKLRGWFDAEGQ ALDGVSISDLKSGGVGGSNTNWKTLYEVKSENLGQGDKPDYFSSVATVVYLRKENCMYQACPTQDCNKKV IDQONGLYRCEKCDTEFPNFKYRMILSVNIADFQENQWVTCFQESAEAILGQNAAYLGELKDKNEQAFEE

VFQNANFRSFIFRVRVKVETYNDESRIKATVMDVKPVDYREYGRRLVMSIRRSALM

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- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine

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

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

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

RefSeq: NP 002936

RefSeq Size: 4345 RefSeq ORF: 1848



#### RPA70 (RPA1) (NM\_002945) Human Mass Spec Standard - PH302066

Synonyms: HSSB; MST075; REPA1; RF-A; RP-A; RPA70

 Locus ID:
 6117

 UniProt ID:
 P27694

 Cytogenetics:
 17p13.3

**Summary:** This gene encodes the largest 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 nucleoprotein complex protects the 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. This subunit contains four oligonucleotide/oligosaccharide-binding (OB) domains, though the majority of ssDNA binding occurs in two of these domains. The heterotrimeric complex has two different modes of ssDNA binding, a low-affinity and high-affinity mode, determined by which ssDNA binding domains are utilized. The different binding modes differ in the length of DNA bound and in the proteins with which it interacts, thereby playing a role in regulating different genomic maintenance pathways. [provided by RefSeq, Sep 2017]

**Protein Families:** Druggable Genome, Stem cell - Pluripotency

**Protein Pathways:** DNA replication, Homologous recombination, Mismatch repair, Nucleotide excision repair

## **Product images:**



Coomassie blue staining of purified RPA1 protein (Cat# [TP302066]). The protein was produced from HEK293T cells transfected with RPA1 cDNA clone (Cat# [RC202066]) using MegaTran 2.0 (Cat# [TT210002]).