

Product datasheet for TR302669

PARP9 Human shRNA Plasmid Kit (Locus ID 83666)

Product data:

OriGene Technologies, Inc.

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Product Type:	shRNA Plasmids
Product Name:	PARP9 Human shRNA Plasmid Kit (Locus ID 83666)
Locus ID:	83666
Synonyms:	ARTD9; BAL; BAL1; MGC:7868
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	PARP9 - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 83666). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	<u>NM 001146102</u> , <u>NM 001146103</u> , <u>NM 001146104</u> , <u>NM 001146105</u> , <u>NM 001146106</u> , <u>NM 031458</u> , <u>NM 031458.1</u> , <u>NM 031458.2</u> , <u>NM 001146106.1</u> , <u>NM 001146102.1</u> , <u>NM 001146103.1</u> , <u>NM 001146104.1</u> , <u>NM 001146105.1</u> , <u>BC039580</u> , <u>BC017463</u> , <u>NM 001146105.2</u> , <u>NM 001146103.2</u> , <u>NM 001146106.2</u> , <u>NM 001146104.2</u> , <u>NM 001146102.2</u>
UniProt ID:	<u>Q8IXQ6</u>



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PARP9 Human shRNA Plasmid Kit (Locus ID 83666) – TR302669

Summary:	ADP-ribosyltransferase which, in association with E3 ligase DTX3L, plays a role in DNA damage repair and in immune responses including interferon-mediated antiviral defenses (PubMed:16809771, PubMed:23230272, PubMed:26479788, PubMed:27796300). Within the complex, enhances DTX3L E3 ligase activity which is further enhanced by PARP9 binding to poly(ADP-ribose) (PubMed:28525742). In association with DTX3L and in presence of E1 and E2 enzymes, mediates NAD(+)-dependent mono-ADP-ribosylation of ubiquitin which prevents ubiquitin conjugation to substrates such as histones (PubMed:28525742). During DNA repair, PARP1 recruits PARP9/BAL1-DTX3L complex to DNA damage sites via PARP9 binding to ribosylated PARP1 (PubMed:23230272). Subsequent PARP1-dependent PARP9/BAL1-DTX3L-mediated ubiquitination promotes the rapid and specific recruitment of 53BP1/TP53BP1, UIMC1/RAP80, and BRCA1 to DNA damage sites (PubMed:23230272, PubMed:28525742). In response to DNA damage, PARP9-DTX3L complex is required for efficient non-homologous end joining (NHEJ); the complex function is negatively modulated by PARP9 activity (PubMed:28525742). Dispensable for B-cell receptor (BCR) assembly through V(D)] recombination and class switch recombination (CSR) (By similarity). In macrophages, positively regulates pro-inflammatory cytokines production in response to IFNG stimulation by suppressing PARP14-mediated STAT1 ADP-ribosylation and thus promoting STAT1 phosphorylation (PubMed:27796300). Also suppresses PARP14-mediated STAT6 ADP-
	ribosylation (PubMed:27796300).[UniProtKB/Swiss-Prot Function]
shRNA Design:	These shRNA constructs were designed against multiple splice variants at this gene locus. To be certain that your variant of interest is targeted, please contact <u>techsupport@origene.com</u> . If you need a special design or shRNA sequence, please utilize our <u>custom shRNA service</u> .
Performance Guaranteed:	OriGene guarantees that the sequences in the shRNA expression cassettes are verified to correspond to the target gene with 100% identity. One of the four constructs at minimum are guaranteed to produce 70% or more gene expression knock-down provided a minimum transfection efficiency of 80% is achieved. Western Blot data is recommended over qPCR to evaluate the silencing effect of the shRNA constructs 72 hrs post transfection. To properly assess knockdown, the gene expression level from the included scramble control vector must be used in comparison with the target-specific shRNA transfected samples.
	For non-conforming shRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the shRNA kit. To arrange for a free replacement with newly designed constructs, please contact Technical Services at techsupport@origene.com. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data

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preferred).