

## Product datasheet for **TL313172**

### ERG Human shRNA Plasmid Kit (Locus ID 2078)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	ERG Human shRNA Plasmid Kit (Locus ID 2078)
Locus ID:	2078
Synonyms:	erg-3; p55
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	ERG - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 2078). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">NM_001136154</a> , <a href="#">NM_001136155</a> , <a href="#">NM_001243428</a> , <a href="#">NM_001243429</a> , <a href="#">NM_001243432</a> , <a href="#">NM_001291391</a> , <a href="#">NM_004449</a> , <a href="#">NM_182918</a> , <a href="#">NR_111949</a> , <a href="#">NM_001331025</a> , <a href="#">NM_182918.1</a> , <a href="#">NM_182918.2</a> , <a href="#">NM_182918.3</a> , <a href="#">NM_004449.1</a> , <a href="#">NM_004449.2</a> , <a href="#">NM_004449.3</a> , <a href="#">NM_004449.4</a> , <a href="#">NM_001136154.1</a> , <a href="#">NM_001136155.1</a> , <a href="#">NM_001243432.1</a> , <a href="#">NM_001243432.2</a> , <a href="#">NM_001243429.1</a> , <a href="#">NM_001243428.1</a> , <a href="#">NM_001291391.1</a> , <a href="#">BC040168</a> , <a href="#">BC040168.1</a> , <a href="#">NM_001243433</a>
UniProt ID:	<a href="#">P11308</a>



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<b>Summary:</b>	<p>This gene encodes a member of the erythroblast transformation-specific (ETS) family of transcription factors. All members of this family are key regulators of embryonic development, cell proliferation, differentiation, angiogenesis, inflammation, and apoptosis. The protein encoded by this gene is mainly expressed in the nucleus. It contains an ETS DNA-binding domain and a PNT (pointed) domain which is implicated in the self-association of chimeric oncoproteins. This protein is required for platelet adhesion to the subendothelium, inducing vascular cell remodeling. It also regulates hematopoiesis, and the differentiation and maturation of megakaryocytic cells. This gene is involved in chromosomal translocations, resulting in different fusion gene products, such as TMPSSR2-ERG and NDRG1-ERG in prostate cancer, EWS-ERG in Ewing's sarcoma and FUS-ERG in acute myeloid leukemia. More than two dozens of transcript variants generated from combinatorial usage of three alternative promoters and multiple alternative splicing events have been reported, but the full-length nature of many of these variants has not been determined. [provided by RefSeq, Apr 2014]</p>
<b>shRNA Design:</b>	<p>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 <a href="mailto:techsupport@origene.com">techsupport@origene.com</a>. If you need a special design or shRNA sequence, please utilize our <a href="#">custom shRNA service</a>.</p>
<b>Performance Guaranteed:</b>	<p>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.</p> <p>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 <a href="mailto:techsupport@origene.com">techsupport@origene.com</a>. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).</p>