

Product datasheet for TR513988

Fto Mouse shRNA Plasmid (Locus ID 26383)

Product data:

Product Type: shRNA Plasmids

Product Name: Fto Mouse shRNA Plasmid (Locus ID 26383)

Locus ID: 26383

Synonyms: AW743446; mKIAA1752

Vector: pRS (TR20003)

E. coli Selection: Ampicillin

Mammalian Cell Puromycin

Selection:

Format: Retroviral plasmids

Components: Fto - Mouse, 4 unique 29mer shRNA constructs in retroviral untagged vector (Gene ID =

26383). 5µg purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.

RefSeq: <u>BC022222</u>, <u>BC057008</u>, <u>NM 011936</u>, <u>NM 011936.1</u>, <u>NM 011936.2</u>

UniProt ID: Q8BGW1

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Summary:

RNA demethylase that mediates oxidative demethylation of different RNA species, such as mRNAs, tRNAs and snRNAs, and acts as a regulator of fat mass, adipogenesis and energy homeostasis (PubMed:17991826, PubMed:18775698, PubMed:28002401). Specifically demethylates N(6)-methyladenosine (m6A) RNA, the most prevalent internal modification of messenger RNA (mRNA) in higher eukaryotes (PubMed:28002401). M6A demethylation by FTO affects mRNA expression and stability (By similarity). Also able to demethylate m6A in U6 small nuclear RNA (snRNA) (By similarity). Mediates demethylation of N(6),2'-Odimethyladenosine cap (m6A(m)), by demethylating the N(6)-methyladenosine at the second transcribed position of mRNAs and U6 snRNA (PubMed:28002401). Demethylation of m6A(m) in the 5'-cap by FTO affects mRNA stability by promoting susceptibility to decapping (By similarity). Also acts as a tRNA demethylase by removing N(1)-methyladenine from various tRNAs (By similarity). Has no activity towards 1-methylguanine (By similarity). Has no detectable activity towards double-stranded DNA (By similarity). Also able to repair alkylated DNA and RNA by oxidative demethylation: demethylates single-stranded RNA containing 3methyluracil, single-stranded DNA containing 3-methylthymine and has low demethylase activity towards single-stranded DNA containing 1-methyladenine or 3-methylcytosine (PubMed:17991826, PubMed:18775698). Ability to repair alkylated DNA and RNA is however unsure in vivo (PubMed:17991826, PubMed:18775698). Involved in the regulation of fat mass, adipogenesis and body weight, thereby contributing to the regulation of body size and body fat accumulation (PubMed:19234441, PubMed:19680540, PubMed:21076408, PubMed:23817550, PubMed:23300482). Involved in the regulation of thermogenesis and the control of adipocyte differentiation into brown or white fat cells (PubMed:19234441, PubMed:19680540). Regulates activity of the dopaminergic midbrain circuitry via its ability to demethylate m6A in mRNAs (PubMed:23817550).[UniProtKB/Swiss-Prot Function]

shRNA Design:

Performance Guaranteed: 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 techsupport@origene.com. If you need a special design or shRNA sequence, please utilize our custom shRNA service.

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 preferred).