

## **Product datasheet for SC334216**

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## TRA2A (NM\_001282759) Human Untagged Clone

**Product data:** 

**Product Type:** Expression Plasmids

Product Name: TRA2A (NM\_001282759) Human Untagged Clone

Tag: Tag Free Symbol: TRA2A

Synonyms: AWMS1; HSU53209

Mammalian Cell Neomycin

Selection:

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >NCBI ORF sequence for NM\_001282759, the custom clone sequence may differ by one or

more nucleotides

Restriction Sites: Sgfl-Mlul

**ACCN:** NM 001282759

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

**Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).





**Reconstitution Method:** 

- 1. Centrifuge at 5,000xg for 5min.
- 2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
- 3. Close the tube and incubate for 10 minutes at room temperature.
- 4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.

5. 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.

RefSeq: <u>NM 001282759.1</u>, <u>NP 001269688.1</u>

RefSeq Size: 2254 bp
RefSeq ORF: 543 bp
Locus ID: 29896
UniProt ID: Q13595
Cytogenetics: 7p15.3

**Protein Pathways:** Spliceosome

**Gene Summary:** This gene is a member of the transformer 2 homolog family and encodes a protein with

several RRM (RNA recognition motif) domains. This phosphorylated nuclear protein binds to specific RNA sequences and plays a role in the regulation of pre-mRNA splicing. Alternatively spliced transcript variants have been found for this gene. [provided by RefSeq, Sep 2013] Transcript Variant: This variant (4) contains two additional exons in the 5' region and uses an alternate in-frame acceptor splice site in the 3' coding region compared to variant 1. The former difference results in translation initiation from an in-frame downstream start codon,

and an isoform (3) with a shorter N-terminus compared to isoform 1.