

## **Product datasheet for SC333441**

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## PRAC2 (NM\_001282275) Human Untagged Clone

**Product data:** 

**Product Type:** Expression Plasmids

Product Name: PRAC2 (NM\_001282275) Human Untagged Clone

Tag: Tag Free Symbol: PRAC2

Synonyms: C17orf93; HOXB-AS5; HOXB13-AS1; HOXBAS5; NCRNA00253

Mammalian Cell

Selection:

Neomycin

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

Fully Sequenced ORF: >SC333441 representing NM\_001282275.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

**Restriction Sites:** Sgfl-Mlul

ACCN: NM\_001282275

**Insert Size:** 273 bp

**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 001282275.1</u>

 RefSeq Size:
 584 bp

 RefSeq ORF:
 273 bp

 Locus ID:
 360205

 UniProt ID:
 D3DTV9

 Cytogenetics:
 17q21.32

 MW:
 10.4 kDa

**Gene Summary:** This gene is highly expressed in prostate, rectum, colon, and testis. This gene may produce a

non-coding RNA or may encode a short protein that might localize to the nucleus. Alternative

splicing results in multiple transcript variants. [provided by RefSeq, Aug 2013]

Transcript Variant: This variant (2) differs in the 5' UTR, lacks a portion of the 5' coding region, and initiates translation at a downstream in-frame start codon, compared to variant 1. The

encoded isoform (b) has a shorter N-terminus, compared to isoform a.