

## **Product datasheet for TP517263**

#### OriGene Technologies, Inc.

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### Arid3c (NM\_001017362) Mouse Recombinant Protein

**Product data:** 

**Product Type:** Recombinant Proteins

**Description:** Purified recombinant protein of Mouse AT rich interactive domain 3C (BRIGHT-like) (Arid3c),

with C-terminal MYC/DDK tag, expressed in HEK293T cells, 20ug

Species: Mouse Expression Host: HEK293T

expression nost: nex29.

**Expression cDNA Clone** >MR217263 protein sequence or AA Sequence: Red=Cloning site Green=Tags(s)

MEALQRQQAARLAQGVGPLAPPRLPLPQPPLLGARTLQAPEGAIGVVGAEEEEGAEDEEGETPLAEEETA EQSHPGARCPNSPSSQSPGIQPHEWTYEEQFKQLYELDADPKRKEFLDDLFSFMQKRGTPVNRVPIMAKQ VLDLYALFRLVTAKGGLVEVINRKVWREVTRGLSLPTTITSAAFTLRTQYMKYLYPYECETRALSSPGEL QAAIDSNRREGRRQAYTAVPLFNLAGPTPRGAPGPASSHGPAPTATPNCPGPTQGSASGLPAHACAQLSP SPVKKEESGIPPPRLALPMGLASEATREKLAPEEPPEKRAVLMGPVDSPRLGAPPSFLPRGKAPLREERL

 ${\tt DGPLNLAGSGISSINVALEINGVVYTGILFARRQPVPASLGPTNPPPLPSTGPPSSTLP}$ 

**TRTRPLEQKLISEEDLAANDILDYKDDDDKV** 

Tag: C-MYC/DDK
Predicted MW: 43.6 kDa

Concentration: >0.05 µg/µL as determined by microplate BCA method

**Purity:** > 80% as determined by SDS-PAGE and Coomassie blue staining

**Buffer:** 25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol

**Note:** For testing in cell culture applications, please filter before use. Note that you may experience

some loss of protein during the filtration process.

**Storage:** Store at -80°C after receiving vials.

Stability: Stable for 12 months from the date of receipt of the product under proper storage and

handling conditions. Avoid repeated freeze-thaw cycles.

**RefSeq:** NP 001017362

 Locus ID:
 550619

 UniProt ID:
 A6PWV5



# ORIGENE Arid

#### Arid3c (NM\_001017362) Mouse Recombinant Protein - TP517263

RefSeq Size: 1751

Cytogenetics: 4 A5 RefSeq ORF: 1230

Summary: This gene is a member of the ARID (AT-rich interaction domain) family of proteins. The ARID

domain is a helix-turn-helix motif-based DNA-binding domain. ARID family members have roles in embryonic patterning, cell lineage gene regulation, cell cycle control, transcriptional regulation and possibly in chromatin structure modification. [provided by RefSeq, Jul 2008]