



# SZABO SCANDIC

Part of Europa Biosite

## Produktinformation



Forschungsprodukte & Biochemikalien



Zellkultur & Verbrauchsmaterial



Diagnostik & molekulare Diagnostik



Laborgeräte & Service

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See the following pages for more information!



### Lieferung & Zahlungsart

siehe unsere [Liefer- und Versandbedingungen](#)

### Zuschläge

- Mindermengenzuschlag
- Trockeneiszuschlag
- Gefahrgutzuschlag
- Expressversand

### SZABO-SCANDIC HandelsgmbH

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## Datasheet

### TSC22D3 (Human) Recombinant Protein (Q01)

**Catalog Number:** H00001831-Q01

**Regulation Status:** For research use only (RUO)

**Product Description:** Human TSC22D3 partial ORF (NP\_932174, 1 a.a. - 97 a.a.) recombinant protein with GST-tag at N-terminal.

**Sequence:**

MAQSKLDCRSPVGLDCCNCCLDLAHRSGLRGSSGE  
NNNPGSPTVSNFRQLQEKLVFENLNTDKLNSIMRQDS  
LEPVLRDPCYLINEGICNRNIDQT

**Host:** Wheat Germ (in vitro)

**Theoretical MW (kDa):** 36.41

**Applications:** AP, Array, ELISA, WB-Re  
(See our web site product page for detailed applications information)

**Protocols:** See our web site at  
<http://www.abnova.com/support/protocols.asp> or product page for detailed protocols

**Preparation Method:** [in vitro wheat germ expression system](#)

**Purification:** Glutathione Sepharose 4 Fast Flow

**Storage Buffer:** 50 mM Tris-HCl, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.

**Storage Instruction:** Store at -80°C. Aliquot to avoid repeated freezing and thawing.

**Entrez GeneID:** 1831

**Gene Symbol:** TSC22D3

**Gene Alias:** DIP, DKFZp313A1123, DSIPI, GILZ, TSC-22R, hDIP

**Gene Summary:** The protein encoded by this gene shares significant sequence identity with the murine TSC-22 and Drosophila shs, both of which are leucine zipper proteins, that function as transcriptional

regulators. The expression of this gene is stimulated by glucocorticoids and interleukin 10, and it appears to play a key role in the anti-inflammatory and immunosuppressive effects of this steroid and chemokine. Transcript variants encoding different isoforms have been identified for this gene. [provided by RefSeq]