



# SZABO SCANDIC

Part of Europa Biosite

## Produktinformation



Forschungsprodukte & Biochemikalien



Zellkultur & Verbrauchsmaterial



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Laborgeräte & Service

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### Lieferung & Zahlungsart

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### Zuschläge

- Mindermengenzuschlag
- Trockeneiszuschlag
- Gefahrgutzuschlag
- Expressversand

### SZABO-SCANDIC HandelsgmbH

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## Datasheet

### SNX11 (Human) Recombinant Protein (P01)

**Catalog Number:** H00029916-P01

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

**Product Description:** Human SNX11 full-length ORF (AAH00768, 1 a.a. - 270 a.a.) recombinant protein with GST-tag at N-terminal.

**Sequence:**

MGFWCRMSENQEQEEVITVRVQDPRVQNEGSWNSY  
VDYKIFLHTNSKAFTAKTSCVRRRYREFVWLRKQLQR  
NAGLVPVPELPGKSTFFGTSDEFIEKRRQGLQHFLEKV  
LQSVVLLSDSQLHLFLQSQLSVPEIEACVQGRSTMTVS  
DAILRYAMSNCGWAQEERQSSSHLAKGDQPKSCCFL  
PRSGRRSSPPPPSEEKDHLEWAPVVDSEVPSLESP  
TLPPLSSPLCCDFGRPKEGTSTLQSVRRVAVGGDHAVP  
LDPGQLETVLEK

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

**Theoretical MW (kDa):** 55.44

**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:** 29916

**Gene Symbol:** SNX11

**Gene Alias:** MGC111019

**Gene Summary:** This gene encodes a member of the sorting nexin family. Members of this family contain a phox (PX) domain, which is a phosphoinositide binding domain, and are involved in intracellular trafficking. This protein does not contain a coiled coil region, like some family members. This gene encodes a protein of unknown function. This gene results in two transcript variants differing in the 5' UTR, but encoding the same protein. [provided by RefSeq]