

## **Product datasheet for PH301653**

### OriGene Technologies, Inc.

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### EBP50 (SLC9A3R1) (NM\_004252) Human Mass Spec Standard

**Product data:** 

**Product Type:** Mass Spec Standards

**Description:** SLC9A3R1 MS Standard C13 and N15-labeled recombinant protein (NP\_004243)

Species:HumanExpression Host:HEK293

Expression cDNA Clone

RC201653

or AA Sequence: Predicted MW:

38.9 kDa

Protein Sequence: >RC201653 protein sequence

Red=Cloning site Green=Tags(s)

MSADAAAGAPLPRLCCLEKGPNGYGFHLHGEKGKLGQYIRLVEPGSPAEKAGLLAGDRLVEVNGENVEKE THQQVVSRIRAALNAVRLLVVDPETDEQLQKLGVQVREELLRAQEAPGQAEPPAAAEVQGAGNENEPREA DKSHPEQRELRPRLCTMKKGPSGYGFNLHSDKSKPGQFIRSVDPDSPAEASGLRAQDRIVEVNGVCMEGK QHGDVVSAIRAGGDETKLLVVDRETDEFFKKCRVIPSQEHLNGPLPVPFTNGEIQKENSREALAEAALES PRPALVRSASSDTSEELNSQDSPPKQDSTAPSSTSSSDPILDFNISLAMAKERAHQKRSSKRAPQMDWSK

KNELFSNL

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Tag: C-Myc/DDK

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

Concentration:  $>0.05 \mu g/\mu L$  as determined by microplate BCA method

Labeling Method: Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine

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

Storage: Store at -80°C. Avoid repeated freeze-thaw cycles.

**Stability:** Stable for 3 months from receipt of products under proper storage and handling conditions.

**RefSeq:** NP 004243

RefSeq Size: 2032 RefSeq ORF: 1074

Synonyms: EBP50; NHERF; NHERF-1; NHERF1; NPHLOP2

**Locus ID:** 9368



#### EBP50 (SLC9A3R1) (NM\_004252) Human Mass Spec Standard - PH301653

 UniProt ID:
 014745

 Cytogenetics:
 17q25.1

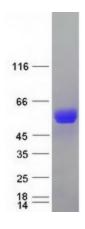
**Summary:** This gene encodes a sodium/hydrogen exchanger regulatory cofactor. The protein interacts

with and regulates various proteins including the cystic fibrosis transmembrane conductance regulator and G-protein coupled receptors such as the beta2-adrenergic receptor and the parathyroid hormone 1 receptor. The protein also interacts with proteins that function as linkers between integral membrane and cytoskeletal proteins. The protein localizes to actinrich structures including membrane ruffles, microvilli, and filopodia. Mutations in this gene result in hypophosphatemic nephrolithiasis/osteoporosis type 2, and loss of heterozygosity

of this gene is implicated in breast cancer.[provided by RefSeq, Sep 2009]

**Protein Families:** Druggable Genome

# **Product images:**



Coomassie blue staining of purified SLC9A3R1 protein (Cat# [TP301653]). The protein was produced from HEK293T cells transfected with SLC9A3R1 cDNA clone (Cat# [RC201653]) using MegaTran 2.0 (Cat# [TT210002]).