

TBL1XR1 (L-08): sc-100908

BACKGROUND

TBL1XR1 (transducin (β)-like 1 X-linked receptor 1), also known as C21, DC42, IRA1 or TBLR1 (TBL1-related protein 1), is a ubiquitously expressed protein that belongs to the WD repeat EBI family of proteins. Localizing to the cytoplasm and the nucleus, TBL1XR1 contains eight WD repeats, one LisH domain and one F-box-like domain. TBL1XR1 functions as a transcriptional regulator, acting as a component of the co-repressor machinery (NCoR/SMRT complex) that is required for the activation of many transcription factors. Specifically, TBL1XR1 is essential for the recruitment of proteasome machinery and, therefore, the subsequent degradation of co-repressors upon ligand binding. The knockdown of TBL1XR1 inhibits β-catenin-mediated transcription and greatly suppresses the growth of head and neck squamous cell carcinoma cells. This suggests that TBL1XR1 may be a useful target in anti-cancer therapy.

REFERENCES

1. Online Mendelian Inheritance in Man, OMIM™. 2002. Johns Hopkins University, Baltimore, MD. MIM Number: 608628. World Wide Web URL: <http://www.ncbi.nlm.nih.gov/omim/>
2. Yoon, H.G., et al. 2003. Purification and functional characterization of the human N-CoR complex: the roles of HDAC3, TBL1 and TBLR1. *EMBO J.* 22: 1336-1346.

CHROMOSOMAL LOCATION

Genetic locus: TBL1XR1 (human) mapping to 3q26.32.

SOURCE

TBL1XR1 (L-08) is a mouse monoclonal antibody raised against partial recombinant protein mapping to an internal region of TBL1XR1 of human origin.

PRODUCT

Each vial contains 50 µg IgG₁ kappa light chain in 0.5 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

APPLICATIONS

TBL1XR1 (L-08) is recommended for detection of TBL1XR1 of human origin by Western Blotting (starting dilution 1:200, dilution range 1:100-1:1000), immunoprecipitation [1-2 µg per 100-500 µg of total protein (1 ml of cell lysate)], immunofluorescence (starting dilution 1:50, dilution range 1:50-1:500), immunohistochemistry (including paraffin-embedded sections) (starting dilution 1:50, dilution range 1:50-1:500) and solid phase ELISA (starting dilution 1:30, dilution range 1:30-1:3000).

Suitable for use as control antibody for TBL1XR1 siRNA (h): sc-106601, TBL1XR1 shRNA Plasmid (h): sc-106601-SH and TBL1XR1 shRNA (h) Lentiviral Particles: sc-106601-V.

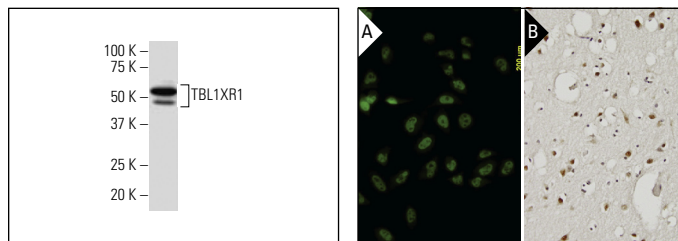
Molecular Weight of TBL1XR1: 55 kDa.

Positive Controls: HeLa whole cell lysate: sc-2200 or Y79 nuclear extract: sc-2126.

STORAGE

Store at 4° C, **DO NOT FREEZE**. Stable for one year from the date of shipment. Non-hazardous. No MSDS required.

DATA



TBL1XR1 (L-08): sc-100908. Western blot analysis of TBL1XR1 expression in HeLa whole cell lysate.

TBL1XR1 (L-08): sc-100908. Immunofluorescence staining of paraformaldehyde-fixed HeLa cells showing nuclear localization (A). Immunoperoxidase staining of formalin-fixed, paraffin-embedded human cerebral cortex tissue showing nuclear localization (B).

SELECT PRODUCT CITATIONS

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3. Jones, C.L., et al. 2014. Loss of TBL1XR1 disrupts glucocorticoid receptor recruitment to chromatin and results in glucocorticoid resistance in a B-lymphoblastic leukemia model. *J. Biol. Chem.* 289: 20502-20515.
4. Daniels, G., et al. 2016. Cytoplasmic, full length and novel cleaved variant, TBLR1 reduces apoptosis in prostate cancer under androgen deprivation. *Oncotarget* 7: 39556-39571.
5. van de Poel, S., et al. 2018. Identification and functional characterization of phosphorylation sites of the human papillomavirus 31 E8^AE2 protein. *J. Virol.* 92: e01743-17.
6. Zhang, T., et al. 2020. TBL1XR1 is involved in c-Met-mediated tumorigenesis of human non-small cell lung cancer. *Cancer Gene Ther.* 27: 136-146.
7. Tillotson, R., et al. 2021. Neuronal non-CG methylation is an essential target for MeCP2 function. *Mol. Cell* 81: 1260-1275.e12.
8. Zhou, Q., et al. 2021. Deficiency of TBL1XR1 causes asthenozoospermia. *Andrologia* 53: e13980.
9. Bray, D., et al. 2022. CASCADE: high-throughput characterization of regulatory complex binding altered by non-coding variants. *Cell Genom.* 2: 100098.
9. Carrasco Pro, S., et al. 2023. Widespread perturbation of ETS factor binding sites in cancer. *Nat. Commun.* 14: 913.

RESEARCH USE

For research use only, not for use in diagnostic procedures.