# ZNF516 (m): 293T Lysate: sc-124804



The Power to Question

### **BACKGROUND**

Zinc-finger proteins contain DNA-binding domains and have a wide variety of functions, most of which encompass some form of transcriptional activation or repression. The majority of zinc-finger proteins contain a Krüppel-type DNA binding domain and a KRAB domain, which is thought to interact with KAP1, thereby recruiting histone modifying proteins. As a member of the krueppel  $C_2H_2$ -type zinc-finger protein family, ZNF516 (zinc-finger protein 516) is a 1,163 amino acid nuclear protein that contains ten  $C_2H_2$ -type zinc-fingers. The gene encoding ZNF516 maps to human chromosome 18, in a region that is frequently found to be affected in 18q deletion syndrome, a multiple-anomaly mental retardation syndrome that is associated with congenital aural atresia.

## **REFERENCES**

- Payre, F. and Vincent, A. 1988. Finger proteins and DNA-specific recognition: distinct patterns of conserved amino acids suggest different evolutionary modes. FEBS Lett. 234: 245-250.
- Rosenfeld, R. and Margalit, H. 1993. Zinc-fingers: conserved properties that can distinguish between spurious and actual DNA-binding motifs. J. Biomol. Struct. Dyn. 11: 557-570.
- Nagase, T., Seki, N., Ishikawa, K., Ohira, M., Kawarabayasi, Y., Ohara, O., Tanaka, A., Kotani, H., Miyajima, N. and Nomura, N. 1996. Prediction of the coding sequences of unidentified human genes. VI. The coding sequences of 80 new genes (KIAA0201-KIAA0280) deduced by analysis of cDNA clones from cell line KG-1 and brain. DNA Res. 3: 321-9, 341.
- Urrutia, R. 2003. KRAB-containing zinc-finger repressor proteins. Genome Biol. 4: 231.
- Nuijten, I., Admiraal, R., Van Buggenhout, G., Cremers, C., Frijns, J.P., Smeets, D. and van Ravenswaaij-Arts, C. 2003. Congenital aural atresia in 18q deletion or de Grouchy syndrome. Otol. Neurotol. 24: 900-906.
- Beausoleil, S.A., Jedrychowski, M., Schwartz, D., Elias, J.E., Villén, J., Li, J., Cohn, M.A., Cantley, L.C. and Gygi, S.P. 2004. Large-scale characterization of HeLa cell nuclear phosphoproteins. Proc. Natl. Acad. Sci. USA 101: 12130-12135.
- Dostal, A., Nemeckova, J., Gaillyova, R., Vranova, V., Zezulkova, D., Lejska, M., Slapak, I., Dostalova, Z. and Kuglik, P. 2006. Identification of 2.3-Mb gene locus for congenital aural atresia in 18q22.3 deletion: a case report analyzed by comparative genomic hybridization. Otol. Neurotol. 27: 427-432.
- 8. Riley, D.E., Jeon, J.S. and Krieger, J.N. 2007. Simple repeat evolution includes dramatic primary sequence changes that conserve folding potential. Biochem. Biophys. Res. Commun. 355: 619-625.

### **STORAGE**

Store at -20° C. Repeated freezing and thawing should be minimized. Sample vial should be boiled once prior to use. Non-hazardous. No MSDS required.

### **PROTOCOLS**

See our web site at www.scbt.com for detailed protocols and support products.

#### **CHROMOSOMAL LOCATION**

Genetic locus: Zfp516 (mouse) mapping to 18 E3.

#### **PRODUCT**

ZNF516 (m): 293T Lysate represents a lysate of mouse ZNF516 transfected 293T cells and is provided as 100 µg protein in 200 µl SDS-PAGE buffer.

#### **APPLICATIONS**

ZNF516 (m): 293T Lysate is suitable as a Western Blotting positive control for mouse reactive ZNF516 antibodies. Recommended use:  $10-20~\mu l$  per lane.

Control 293T Lysate: sc-117752 is available as a Western Blotting negative control lysate derived from non-transfected 293T cells.

## **RESEARCH USE**

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

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