# Mlx (h): 293 Lysate: sc-111152



The Power to Question

### **BACKGROUND**

Max is a nuclear localized bHLH-Zip protein that forms homodimers or heterodimers with Myc family members, including Myc, Mad 1, Mad 3, Mad 4, Mxi1 and Mnt (or Rox). These dimers bind to the E-box sequence CACGTG in order to regulate cell growth, proliferation and apoptosis. Mlx (Max-like protein X) is a bHLH-Zip protein that is structurally and functionally related to Max. Like Max, Mlx is broadly expressed in many tissues and has a long half-life. Mlx also forms homodimers or heterodimers with members of the Myc family, specifically Mad 1, Mad 4 and Rox, and members of the Mondo family, to repress or activate transcription from CACGTG E-boxes. MondoA forms weak homodimers and preferentially forms heterodimers with Mlx. The MondoA/Mlx complex is primarily localized to the cytoplasm, but will translocate to the nucleus in response to leptomycin B. Mlx can also dimerize with WBSCR14, a protein involved in Williams-Beuren syndrome (WBS), to repress E-box transcription, which provides further evidence that Mlx is a critical element in a transcription factor network.

#### **REFERENCES**

- Blackwood, E.M., et al. 1991. Max: a helix-loop-helix zipper protein that forms a sequence-specific DNA-binding complex with Myc. Science 251: 1211-1217.
- 2. Amati, B., et al. 1992. Oncogenic activity of the c-Myc protein requires dimerization with Max. Cell 72: 233-245.
- Billin, A.N., et al. 1999. Mlx, a novel Max-like bHLH-Zip protein that interacts with the Max network of transcription factors. J. Biol. Chem. 274: 36344-36350.
- Meroni, G., et al. 2000. Mlx, a new Max-like bHLH-Zip family member: the center stage of a novel transcription factors regulatory pathway? Oncogene 19: 3266-3277.
- 5. Billin, A.N., et al. 2000. MondoA, a novel basic helix-loop-helix-leucine zipper transcriptional activator that constitutes a positive branch of a Max-like network. Mol. Cell. Biol. 20: 8845-8854.
- 6. Cairo, S., et al. 2001. WBSCR14, a gene mapping to the Williams-Beuren syndrome deleted region, is a new member of the Mlx transcription factor network. Hum. Mol. Genet.10: 617-627.
- Stoeckman, A.K., et al. 2004. Mlx is the functional heteromeric partner of the carbohydrate response element-binding protein in glucose regulation of lipogenic enzyme genes. J. Biol. Chem. 279: 15662-15669.
- 8. Ma, L., et al. 2005. Direct role of ChREBP.Mlx in regulating hepatic glucose-responsive genes. J. Biol. Chem. 280: 12019-12027.
- Ma, L., et al. 2007. A critical role for the loop region of the basic helix-loophelix/leucine zipper protein Mlx in DNA binding and glucose-regulated transcription. Nucleic Acids Res. 35: 35-44.

# **CHROMOSOMAL LOCATION**

Genetic locus: MLX (human) mapping to 17q21.2.

# **RESEARCH USE**

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

# **PRODUCT**

MIx (h): 293 Lysate represents a lysate of human MIx transfected 293 cells and is provided as 100 µg protein in 200 µl SDS-PAGE buffer.

### **APPLICATIONS**

MIx (h): 293 Lysate is suitable as a Western Blotting positive control for human reactive MIx antibodies. Recommended use:  $10-20 \mu l$  per lane.

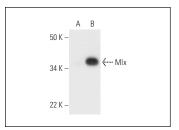
Control 293 Lysate: sc-110760 is available as a Western Blotting negative control lysate derived from non-transfected 293 cells.

Mlx (F-12): sc-393086 is recommended as a positive control antibody for Western Blot analysis of enhanced human Mlx expression in Mlx transfected 293 cells (starting dilution 1:100, dilution range 1:100-1:1,000).

#### **RECOMMENDED SUPPORT REAGENTS**

To ensure optimal results, the following support reagents are recommended: 1) Western Blotting: use m-lgG $\kappa$  BP-HRP: sc-516102 or m-lgG $\kappa$  BP-HRP (Cruz Marker): sc-516102-CM (dilution range: 1:1000-1:10000), Cruz Marker<sup>TM</sup> Molecular Weight Standards: sc-2035, UltraCruz® Blocking Reagent: sc-516214 and Western Blotting Luminol Reagent: sc-2048.

#### DATA



Mlx (F-12): sc-393086. Western blot analysis of Mlx expression in non-transfected: sc-110760 (**A**) and human Mlx transfected: sc-111152 (**B**) 293 whole cell bestes

# 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.