

Anti-Human MAX, monoclonal

Alternate Names: Protein max

Cat. No. 14-094

FOR RESEARCH USE ONLY

NOT FOR USE IN HUMANS



PRODUCT DESCRIPTION

UniProt Summary

UniProt

Primary Accession: [P61244](#)

Transcription regulator. Forms a sequence-specific DNA-binding protein complex with MYC or MAD which recognizes the core sequence 5'-CAC[GA]TG-3'. The MYC:MAX complex is a transcriptional activator, whereas the MAD:MAX complex is a repressor. May repress transcription via the recruitment of a chromatin remodeling complex containing H3 'Lys-9' histone methyltransferase activity. Efficient DNA binding requires dimerization with another bHLH protein. Binds DNA as a heterodimer with MYC or MAD. Part of the E2F6.com-1 complex in G0 phase composed of E2F6, MGA, MAX, TFDP1, CBX3, BAT8, EUHMTASE1, RING1, RNF2, MBLR, L3MBTL2 and YAF2. Component of some MLL1/MLL complex, at least composed of the core...

Physical Characteristics

Quantity: 1 ml (culture supernatant) or 100ug at 1mg/ml (purified)

Format: culture supernatant or purified material

Host/Isotype: mouse IgG2b

Clonality: monoclonal; ID R536.1.1E7

Formulation: culture supernatant contains 0.02% NaN₃. Purified material contains 30% glycerol, PBS and 0.02% NaN₃

Specificity: monospecific for human MAX ; see "Microarray Analysis" below

Reactivity: human; not tested for cross reactivity in other species

Stability/Storage: 12 months long term: -20°C; short term: 4°C; avoid freeze-thaw cycles; aliquot as required

Handling Notes: small volumes of antibody may occasionally become entrapped in the seal of the product vial during shipment and storage; briefly centrifuge the vial on a tabletop centrifuge to dislodge any liquid in the container cap

Tested Research Applications

- Western Immunoblotting - Not recommended
- Immunoprecipitation - PASS by SOP

Antibody tested as purified IgG. Optimal dilution to be determined by user.

Quality Assurance

Notes:

1. Please refer to the SOP manual ([click here](#)) to for detailed explanation of the experimental setup and thresholds used to evaluate the results shown below.
2. Predicted MW of the target protein expressed in the western immunoblotting (WB) and immunoprecipitation (IP) experiment is **11.44kDa** and the three fusion tags (venus, 3xFLAG, V5) adds another **38.7kDa** to this protein. Therefore the protein is expected to migrate **~50.14kDa** in a denatured SDS-PAGE gel.
3. See results below for any applicable cautionary notes.

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Lot-specific COA version tracker: v1.0.0



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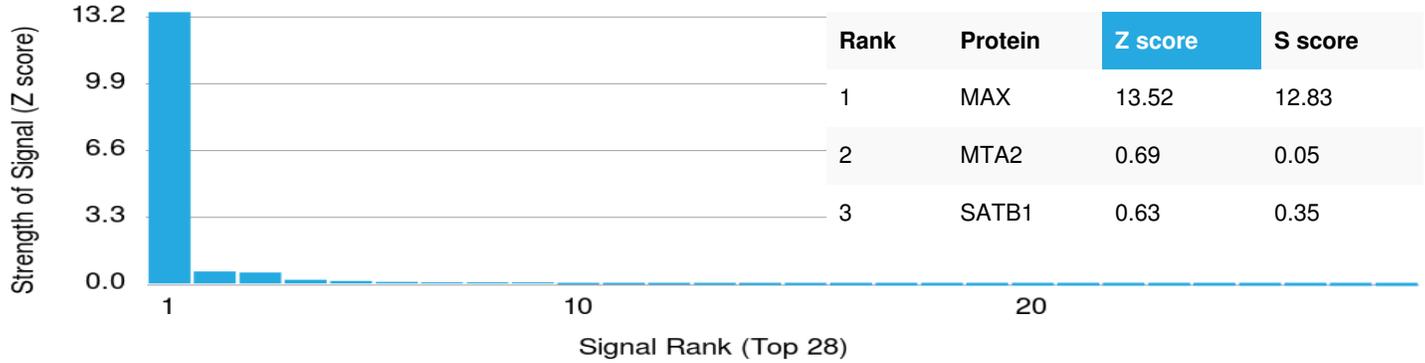
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NEXTGEN PROTEOMICS

PRODUCT DESCRIPTION

Quality Assurance (HuProt™ Array)



About Z and S Scores: The z-score represents the strength of a signal that a mAb (in combination with a fluorescently-tagged anti-IgG secondary antibody) produces when binding to a particular protein on the HuProt™ array. Z-scores are described in units of standard deviations (SDs) above the mean value of all signals generated on that array. To be considered a definite binding event between a mAb and a protein target, the z-score must have a value of at least 2.5 (SDs above the mean). The s-score represents the relative target specificity of a mAb, and is the difference (also in units of SDs) between the z-score a mAb generates upon binding to one protein and the next highest z-score that the mAb generates against another protein. For example, if a mAb binds to protein X with an z-score of 43 and to protein Y with an z-score of 14, then the s-score for the binding of that mAb to protein X is equal to 29.

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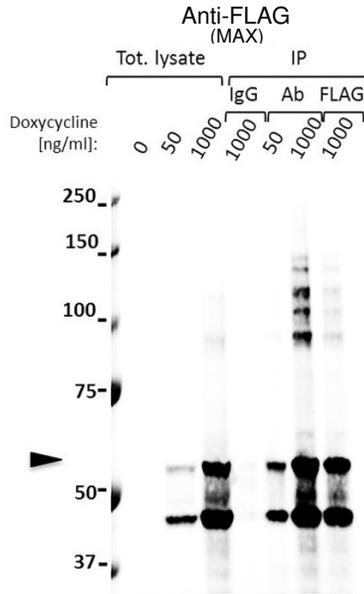


NEXTGEN PROTEOMICS

PRODUCT DESCRIPTION

Quality Assurance (continued)

IMMUNOPRECIPITATION - Predicted MW 50.14kDa



Tet-ON HeLa cells were transfected with construct encoding MAX (NM_145113.1) with an N-terminal fusion of FLAG, YFP (Venus) and V5 tags under a tet-inducible promoter. These cells were stimulated with 0, 50 or 1000 ng/ml doxycycline. Immunoprecipitation (IP) was carried out using 5 µg of either IgG, CDI mAb Anti-MAX (cloneID# R536.1.1E7) or 1 µg of FLAG-M2 (Sigma). Immunoblotting was performed using rabbit Anti-FLAG (1:1000, Cell Signaling #2368). We observe that such fusion proteins are generally expressed as a doublet, with the upper band corresponding to the expected size of the MAX with an N-terminal fusion of FLAG, YFP (Venus) and V5 tags.

Cautionary notes for IP and WB experiments:

1. Target protein runs substantially heavier or lighter than expected

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