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anti-MTA2 antibody (Middle Region)

2 Images



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Target:

| Quantity: | 100 μL | |
|-----------------------|--|--|
| Target: | MTA2 | |
| Binding Specificity: | Middle Region | |
| Reactivity: | Human, Mouse, Rat, Dog, Cow, Rabbit, Horse, Goat, Guinea Pig, Pig, Zebrafish (Danio rerio) | |
| Host: | Rabbit | |
| Clonality: | Polyclonal | |
| Conjugate: | This MTA2 antibody is un-conjugated | |
| Application: | Western Blotting (WB) | |
| Product Details | | |
| Immunogen: | The immunogen is a synthetic peptide directed towards the middle region of human MTA2 | |
| Sequence: | WKKYGGLKTP TQLEGATRGT TEPHSRGHLS RPEAQSLSPY TTSANRAKLL | |
| Predicted Reactivity: | Cow: 93%, Dog: 93%, Goat: 93%, Guinea Pig: 86%, Horse: 93%, Human: 100%, Mouse: 93%, Pig: 93%, Rabbit: 93%, Rat: 93%, Zebrafish: 79% | |
| Characteristics: | This is a rabbit polyclonal antibody against MTA2. It was validated on Western Blot using a cell lysate as a positive control. | |
| Purification: | Affinity Purified | |
| Target Details | | |

MTA2

Alternative Name:

MTA2 (MTA2 Products)

Background:

MTA2 has been identified as a component of NuRD, a nucleosome remodeling deacetylase complex identified in the nucleus of human cells. It shows a very broad expression pattern and is strongly expressed in many tissues. It may represent one member of a small gene family that encode different but related proteins involved either directly or indirectly in transcriptional regulation. Their indirect effects on transcriptional regulation may include chromatin remodeling. This gene encodes a protein that has been identified as a component of NuRD, a nucleosome remodeling deacetylase complex identified in the nucleus of human cells. It shows a very broad expression pattern and is strongly expressed in many tissues. It may represent one member of a small gene family that encode different but related proteins involved either directly or indirectly in transcriptional regulation. Their indirect effects on transcriptional regulation may include chromatin remodeling. It is closely related to another member of this family, a protein that has been correlated with the metastatic potential of certain carcinomas. These two proteins are so closely related that they share the same types of domains. These domains include two DNA binding domains, a dimerization domain, and a domain commonly found in proteins that methylate DNA. One of the proteins known to be a target protein for this gene product is p53. Deacteylation of p53 is correlated with a loss of growth inhibition in transformed cells supporting a connection between these gene family members and metastasis. This gene encodes a protein that has been identified as a component of NuRD, a nucleosome remodeling deacetylase complex identified in the nucleus of human cells. It shows a very broad expression pattern and is strongly expressed in many tissues. It may represent one member of a small gene family that encode different but related proteins involved either directly or indirectly in transcriptional regulation. Their indirect effects on transcriptional regulation may include chromatin remodeling. It is closely related to another member of this family, a protein that has been correlated with the metastatic potential of certain carcinomas. These two proteins are so closely related that they share the same types of domains. These domains include two DNA binding domains, a dimerization domain, and a domain commonly found in proteins that methylate DNA. One of the proteins known to be a target protein for this gene product is p53. Deacteylation of p53 is correlated with a loss of growth inhibition in transformed cells supporting a connection between these gene family members and metastasis. Publication Note: This RefSeq record includes a subset of the publications that are available for this gene. Please see the Entrez Gene record to access additional publications. Alias Symbols: DKFZp686F2281, MTA1L1, PID Protein Interaction Partner: vpr, SUMO2, SUMO3, DNMT3B, DNMT1, NACC2, ZBTB16, UBC, RARA, ZBTB1, RNF2, GPHN, HNRNPR, TRIM28, PPP4R1, ROCK2, SPAG9, POLR2C, POLR2A,

Target Details

| | MAP4, RPP25, LSM8, GET4, LARP7, CD2AP, NR3C1, MTOR, EHBP1L1, RABEP2, BRCC3, SMEK1, ZGPAT, HDAC1, HDAC2, CSNK1E, MTA3, UBE3 Protein Size: 668 |
|-------------------|--|
| Molecular Weight: | 75 kDa |
| Gene ID: | 9219 |
| NCBI Accession: | NM_004739, NP_004730 |
| UniProt: | 094776 |
| Pathways: | Chromatin Binding |

Application Details

| Application Notes: | Optimal working dilutions should be determined experimentally by the investigator. | |
|--------------------|--|--|
| Comment: | Antigen size: 668 AA | |
| Restrictions: | For Research Use only | |

Handling

| Format: | Liquid |
|--------------------|---|
| Concentration: | Lot specific |
| Buffer: | Liquid. Purified antibody supplied in 1x PBS buffer with 0.09 % (w/v) sodium azide and 2 % sucrose. |
| Preservative: | Sodium azide |
| Precaution of Use: | This product contains Sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE which should be handled by trained staff only. |
| Handling Advice: | Avoid repeated freeze-thaw cycles. |
| Storage: | -20 °C |
| Storage Comment: | For short term use, store at 2-8°C up to 1 week. For long term storage, store at -20°C in small aliquots to prevent freeze-thaw cycles. |



Western Blotting

Image 1. WB Suggested Anti-MTA2 Antibody Titration: 0.2-1 ug/ml ELISA Titer: 1:62500 Positive Control: 721_B cell lysate MTA2 is supported by BioGPS gene expression data to be expressed in 721_B

Western Blotting

Image 2. WB Suggested Anti-MTA2

Antibody Titration: 0.2-1 µg/mL ELISA Titer: 1:.2500

Positive Control:.21_B cell lysate

MTA2 is supported by BioGPS gene expression data to be expressed in 721_B