

Anti-Human ZNF701, monoclonal (clone R1015.1.1C5)

Recommended name: Zinc finger protein 701; Short name: ZNF701



Cat. No. m14-316
Lot. No. 20150924.IJVR

Quantity: 100 µg
Storage: -20 °C

FOR RESEARCH USE ONLY

NOT FOR USE IN HUMANS

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UniProt / NCBI Summary

UniProt

Primary Accession # [Q9NV72](#)
Secondary Accession # [Q66K42](#); [A2RRM8](#)

NCBI

GI # [289547638](#)
GenID [55762](#)
Accession # [NP_060730.2](#)
GenBank Nucleotide # [NM_018260.2](#)

Molecular Weight 60,903 Da (531 aa)

May be involved in transcriptional regulation. Belongs to the krueppel C2H2-type zinc-finger protein family. Contains 9 C2H2-type zinc fingers and 1 KRAB domain.

Subcellular location: nucleus

Physical Characteristics

Quantity: 100 µg

Concentration: 1.0 mg/ml

Host / Isotype: mouse IgG1

Clonality: monoclonal; ID JR1015.1.1C5

Immunogen: recombinant protein corresponding to aa residues 122-267 of human ZNF701

Purification: affinity-chromatography using Protein G

Formulation: 30% glycerol, 1x PBS, 0.02% sodium azide

Specificity: monospecific for human ZNF701; see microarray analysis below

Reactivity: human

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; if necessary, briefly centrifuge the vial on a tabletop centrifuge to dislodge any liquid in the container cap.

Tested Research Applications

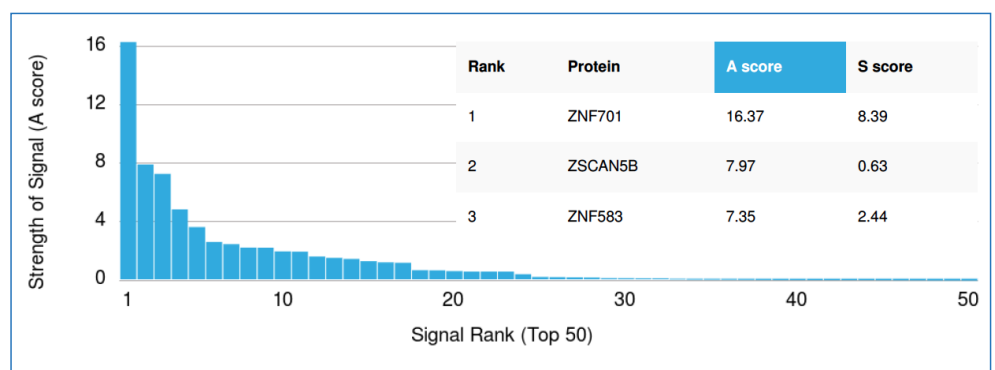
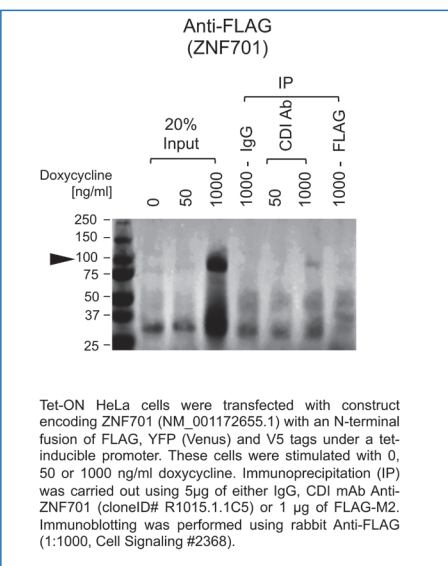
Immunoprecipitation: recommended; see below.

ChIP-Seq: recommended; see page 2

Western Blot: tested on cells transfected with a construct encoding ZNF701; utility on native cells under evaluation

Octet: Recommended.

Quality Assurance



Specificity Analysis with HuProt™ Human Proteome Microarray: Anti Human ZNF701 (clone R1015.1.1C5) was analyzed using the CDI HuProt™ Human Proteome Microarray.

For more information on A/S scores and how they relate to specificity, see page 2.

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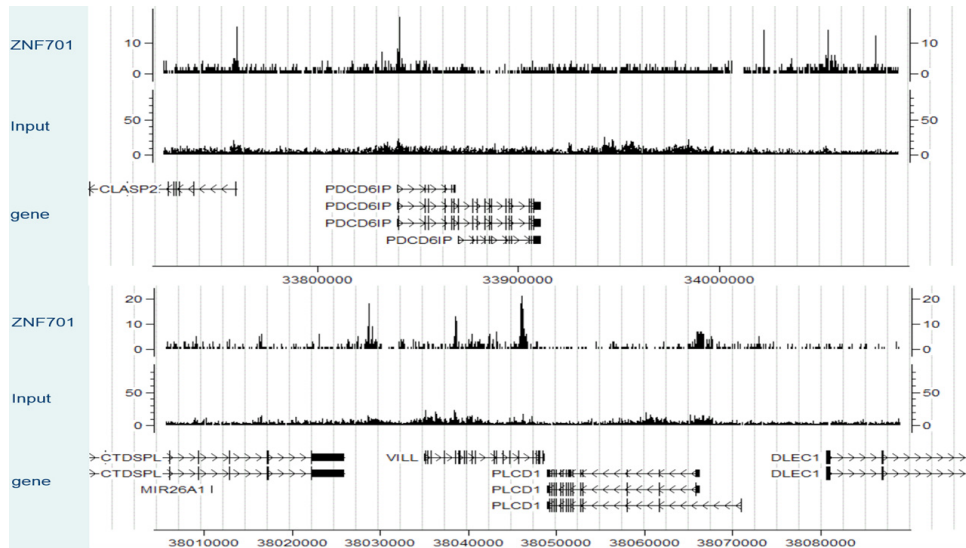
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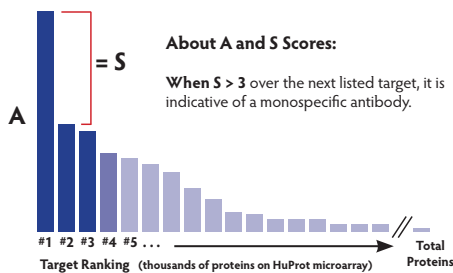
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Tested Research Applications

ChIP-Seq: Recommended



The ChIP was performed with chromatin from 10 million MCF7 cells and 3 µg of Anti-ZNF701 (cloneID # R1015.1.1C5) antibody. The ChIP DNA was sequenced on an Illumina HiSeq platform and read counts were calculated at consecutive 100 bp bins across the human genome hg19. Normalized read-count levels for ChIP-seq of ZNF701 (R1015.1.1C5) and control (Input) around the PDCD6IP and VILL loci are displayed in the CisGenome browser.



Statistical Analysis: Thousands of GenePix data points (from the microarray) are analyzed in terms of signal strength and ranked accordingly.

SUMMARY: The A-score indicates the number of standard deviations above background seen for the mean signal bound by the target antigen. The S-score represents the difference between the A-score of the target antigen and the next best hit on the array. S-scores **greater than 3 standard deviations over the next listed target** are deemed statistically significant and indicate **highly specific antibodies**. More info at cdi-lab.com/HighSpec.html

The development of this antibody was supported by the National Institutes of Health Protein Capture Reagent Program under award U54HG06434 to CDI Laboratories and Johns Hopkins University.

