

Datasheet for ABIN6655575

anti-RUNX1T1 antibody (Internal Region, N-Term)



Images



Go to Product page

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| Quantity: | 100 μL |
|----------------------|---|
| Target: | RUNX1T1 |
| Binding Specificity: | Internal Region, N-Term |
| Reactivity: | Human |
| Host: | Rabbit |
| Clonality: | Polyclonal |
| Conjugate: | This RUNX1T1 antibody is un-conjugated |
| Application: | ELISA, Chromatin Immunoprecipitation (ChIP) |
| Product Details | |
| lmmunogen: | Immunogen: Anti-ETO Antibody was produced in rabbits by repeated immunizations with |
| | human ETO using two synthetic peptides containing sequences from the N-terminal and |
| | internal region of the protein respectively. |
| | Immunogen Type: Peptide |
| Purification: | Anti-ETO Antibody is monospecific antiserum processed by delipidation and defibrination |
| | followed by sterile filtration. This product reacts with human ETO. Cross reactivity with ETO |
| | from other sources is not known. |
| Target Details | |
| Target: | RUNX1T1 |
| Alternative Name: | ETO (RUNX1T1 Products) |
| | |

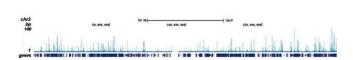
Target Details

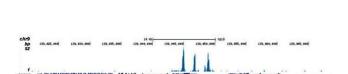
| Background: | Synonyms: Protein CBFA2T1, Cyclin-D-related protein, Eight twenty one protein, Protein ETO, Protein MTG8, Zinc finger MYND domain-containing protein 2, AML1T1, CBFA2T1, CDR, ETO, MTG8, ZMTND2 |
|---------------------|---|
| | Background: ETO is a transcriptional regulator which belongs to the myeloid translocation gene |
| | family. ETO exerts its function by interaction with transcription factors bound to promoters and |
| | binding to histone deacetylases. It recruits a range of corepressors to facilitate transcriptional |
| | repression. The t(8,21)(q22,q22) translocation is one of the most frequent karyotypic |
| | abnormalities in acute myeloid leukaemia. This translocation produces a chimeric gene made |
| | up of the 5'-region of AML1 and the 3'-region of the ETO gene. The chimeric protein is thought |
| | to associate with the nuclear corepressor/histone deacetylase complex to block hematopoietic |
| | differentiation. Anti-ETO Antibody is ideal for research in Gene Expression, Transcription and |
| | Cancer. |
| | Gene Name: RUNX1T1 |
| Gene ID: | 862 |
| NCBI Accession: | NP_001185554 |
| UniProt: | Q06455 |
| Application Details | |
| Application Notes: | Application Note: Anti-ETO Antibody is suitable for Chromatin Immunoprecipitation and ELISA. |
| | Specific conditions for reactivity should be optimized by the end user. |
| | ChIP Dilution: 4 μL/ChIP |
| | ELISA Dilution: 1:100 |
| Restrictions: | For Research Use only |
| Handling | |
| Format: | Liquid |
| Buffer: | 0.01 % (w/v) Sodium Azide |
| | Stabilizer: None |
| Preservative: | Sodium azide |
| Precaution of Use: | This product contains Sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE which |
| | should be handled by trained staff only. |
| Storage: | RT,4 °C,-20 °C |
| | |

Storage Comment:

Store vial at -20° C prior to opening. Aliquot contents and freeze at -20° C or below for extended storage. Avoid cycles of freezing and thawing. Centrifuge product if not completely clear after standing at room temperature. This product is stable for several weeks at 4° C as an undiluted liquid. Dilute only prior to immediate use.

Images



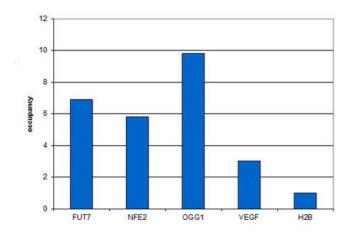


ChIP DNA-Sequencing

Image 1. ChIP-seq results of anti-ETO ChIP-seq results of anti-ETO antibody. ChIP was performed as described in figure 1. The IP'd DNA of 6 ChIP's were pooled and analyzed with an Illumina Genome Analyzer. Library preparation, cluster generation, and sequencing were performed according to the manufacturer's instructions. The 32 bp tags were aligned to the human reference genome (hg18) using the ELAND algorithm. Figure 2 shows the results of the complete chromosome 3. Figures 3-5 shows three genomic regions surrounding the OGG1, FUT7 and NFE2 genes, respectively. The position of the PCR amplicon is indicated with an arrow.

ChIP DNA-Sequencing

Image 2. ChIP-seq results of anti-ETO ab ChIP-seq results of anti-ETO antibody. ChIP was performed as described in figure 1. The IP'd DNA of 6 ChIP's were pooled and analysed with an Illumina Genome Analyzer. Library preparation, cluster generation, and sequencing were performed according to the manufacturer's instructions. The 32 bp tags were aligned to the human reference genome (hg18) using the ELAND algorithm. Figure 2 shows the results of the complete chromosome 3. Figures 3-5 shows three genomic regions surrounding the OGG1, FUT7 and NFE2 genes, respectively. The position of the PCR amplicon is indicated with an arrow.



Chromatin Immunoprecipitation

Image 3. Chromatin Immunoprecipitation of Anti-human ETO Antibody Chromatin Immunoprecipitation results of Rabbit Anti-human ETO Antibody. Chromatin from 1.25 million formaldehyde cross-linked SKNO-1 cells was used with 4ul of Anti-human ETO Antibody and 20ul of magnetic IgG beads per immunoprecipitation. QPCR was performed using primers specific for the FUT7, NFE2, OGG1 and VEGF genes. ChIP results shows the occupancy, calculated as the ratio + control/background for which the H2B gene was used.

Please check the product details page for more images. Overall 6 images are available for ABIN6655575.