

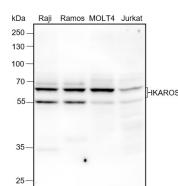
bsm-60278R**[Primary Antibody]****BioSS**
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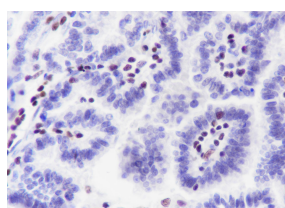
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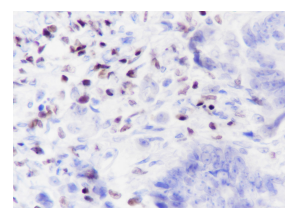
400-901-9800

IKZF1 Recombinant Rabbit mAb**— DATASHEET —****Host:** Rabbit**Isotype:** IgG**Clonality:** Recombinant**CloneNo.:** IK3C9**GeneID:** 10320**SWISS:** Q13422**Target:** IKZF1**Purification:** affinity purified by Protein A**Concentration:** 1mg/ml**Storage:** 0.01M TBS (pH7.4) with 1% BSA, 0.02% Proclin300 and 50% Glycerol.
Shipped at 4°C. Store at -20°C for one year. Avoid repeated freeze/thaw cycles.**Background:** This gene encodes a transcription factor that belongs to the family of zinc-finger DNA-binding proteins associated with chromatin remodeling. The expression of this protein is restricted to the fetal and adult hemo-lymphopoietic system, and it functions as a regulator of lymphocyte differentiation. Several alternatively spliced transcript variants encoding different isoforms have been described for this gene. Most isoforms share a common C-terminal domain, which contains two zinc finger motifs that are required for hetero- or homo-dimerization, and for interactions with other proteins. The isoforms, however, differ in the number of N-terminal zinc finger motifs that bind DNA and in nuclear localization signal presence, resulting in members with and without DNA-binding properties. Only a few isoforms contain the requisite three or more N-terminal zinc motifs that confer high affinity binding to a specific core DNA sequence element in the promoters of target genes. The non-DNA-binding isoforms are largely found in the cytoplasm, and are thought to function as dominant-negative factors. Overexpression of some dominant-negative isoforms have been associated with B-cell malignancies, such as acute lymphoblastic leukemia (ALL). [provided by RefSeq, May 2014]**Applications:** **WB** (1:500-2000)**IHC-P** (1:1000)**IHC-F** (1:50-100)**IF** (1:50-100)**ICC/IF** (1:50)**Reactivity:** Human**Predicted MW.:** 57 kDa**Subcellular Location:** Cytoplasm ,Nucleus**— VALIDATION IMAGES —**

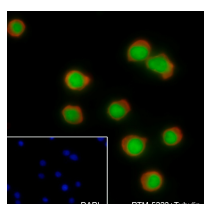
Blocking buffer: 5% NFDm/TBST Primary ab dilution: 1:2000 Primary ab incubation condition: 2 hours at room temperature Lysate: Raji, Ramos, MOLT4, Jurkat Protein loading quantity: 20 µg Exposure time: 60 s Predicted MW: 58 kDa Observed MW: 50-70 kDa



Tissue: Human lung adenocarcinoma Section type: Formalin fixed & Paraffin-embedded section Retrieval method: High temperature and high pressure Retrieval buffer: Tris/EDTA buffer, pH 9.0 Primary ab dilution: 1:1000 Primary ab incubation condition: 1 hour at room temperature Counter stain: Hematoxylin Comment: Color brown is the positive signal for bsm-60278R



Tissue: Human colon cancer Section type: Formalin fixed & Paraffin-embedded section Retrieval method: High temperature and high pressure Retrieval buffer: Tris/EDTA buffer, pH 9.0 Primary ab dilution: 1:1000 Primary ab incubation condition: 1 hour at room temperature Counter stain: Hematoxylin Comment: Color brown is the positive signal for bsm-60278R



Important Note: This product as supplied is intended for research use only, not for use in human, therapeutic or diagnostic applications.

Cell line: Jurkat Fixative: 4% Paraformaldehyde
Permeabilization: 0.1% TritonX-100 Primary ab
dilution: 1:50 Primary incubation condition: 4°C
overnight Secondary ab: Goat Anti-Rabbit IgG
Nuclear counter stain: DAPI (Blue) Comment:
Color green is the positive signal for bsm-60278R