

# **EXOSC7 Rabbit mAb**

货号: B29615

### 产品信息

| 反应    | Human  |
|-------|--|
| 宿主    | Rabbit   |
| 克隆性   | Monoclonal   |
| 预测反应  |  |
| 应用    | WB IHC IP  |
| 推荐浓度  | WB: 1:500 - 1:2000<br>IHC: 1:50 - 1:200<br>IP: 1:20 - 1:50                                 |
| 理论分子量 | 31kDa  |
| 实测分子量 | 32kDa  |
| 形式    | Liquid   |
| 保存条件  | Store at -20°C. Avoid freeze / thaw cycles. Buffer: PBS with 0.75% BSA,50% glycerol,pH7.3. |
| 偶联物   | Unconjugated   |
| 阳性对照  | U-87MG,SKOV3,K-562,Jurkat,MCF7,Mouse brain,Rat brain                                       |
| 细胞定位  | Cytoplasm,Nucleus,nucleolus  |
| 纯化    | Affinity purification  |

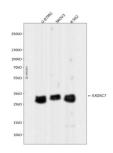
### 抗原信息

| 抗原信息 | Recombinant fusion protein corresponding to Human EXOSC7.  |
|------|--|
| 序列   | MASVTLSEAEKVYIVHGVQEDLRVDGRGCEDYRCVEVETDVVSNTSGSARVKLGHTDILVGVKAEMGTPKLEKPNEGYLE FFVDCSASATPEFEGRGGDDLGTEIANTLYRIFNNKSSVDLKTLCISPREHCWVLYVDVLLLECGGNLFDAISIAVKAALFNT RIPRVRVLEDEEGSKDIELSDDPYDCIRLSVENVPCIVTLCKIGYRHVVDATLQEEACSLASLLVSVTSKGVVTCMRKVGKGS LDPESIFEMMETGKRVGKVLHASLQSVVHKEESLGPKRQKVGFLG |

靶点信息

| 研究背景         | Non-catalytic component of the RNA exosome complex which has 3'->5' exoribonuclease activity and par ticipates in a multitude of cellular RNA processing and degradation events. In the nucleus, the RNA exoso me complex is involved in proper maturation of stable RNA species such as rRNA, snRNA and snoRNA, in the elimination of RNA processing by-products and non-coding 'pervasive' transcripts, such as antisense RNA species and promoter-upstream transcripts (PROMPTs, and of mRNAs with processing defects, there by limiting or excluding their export to the cytoplasm. The RNA exosome may be involved in Ig class swit ch recombination (CSR and/or Ig variable region somatic hypermutation (SHM by targeting AICDA deamin ation activity to transcribed dsDNA substrates. In the cytoplasm, the RNA exosome complex is involved in general mRNA turnover and specifically degrades inherently unstable mRNAs containing AU-rich element s (AREs within their 3' untranslated regions, and in RNA surveillance pathways, preventing translation of aberrant mRNAs. It seems to be involved in degradation of histone mRNA. The catalytic inactive RNA exo some core complex of 9 subunits (Exo-9 is proposed to play a pivotal role in the binding and presentation of RNA for ribonucleolysis, and to serve as a scaffold for the association with catalytic subunits and access sory proteins or complexes. |
|--------------|--|
| 基因 <b>ID</b> | 23016  |
| 基因名          | EXOSC7   |
| Swiss        | Q15024   |
| 别名           | EXOSC7;EAP1;RRP42;Rrp42p;hRrp42p;p8  |

### 产品验证



Western blot analysis of EXOSC7 expressed in U-87MG,SKOV3,K-562 using EXOSC7 Rabbit mAb at 1:1 000. Secondary antibody: HRP Goat Anti-Rabbit lgG (H+L) at 1:5000. Lysates/proteins: 30ug per lane. Blocking buffer: 5% non-fat dry milk in TBST. Detection: ECL Enhanced Kit. Exposure time: 120s.

## 实验步骤

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