

Product Description

Pioneering GTPase and Oncogene Product Development since 2010

PSMB4 RABBIT PAB

货号: S218610 产品全名: PSMB4 兔多抗 基因符号 HN3; HsN3; PRAAS3; PROS26; PROS-26 UNIPROT ID: P28070 (Gene Accession - BC017307)

背景: The proteasome is a multicatalytic proteinase complex with a highly ordered ring-shaped 20S core structure. The core structure is composed of 4 rings of 28 non-identical subunits; 2 rings are composed of 7 alpha subunits and 2 rings are composed of 7 beta subunits. Proteasomes are distributed throughout eukaryotic cells at a high concentration and cleave peptides in an ATP/ubiquitin-dependent process in a non-lysosomal pathway. An essential function of a modified proteasome, the immunoproteasome, is the processing of class I MHC peptides. This gene encodes a member of the proteasome B-type family, also known as the TIB family, that is a 20S core beta subunit. 抗原: Fusion protein of human PSMB4

统原: Fusion protein of nume 经过测试的应用: ELISA, IHC

推荐稀释比: IHC: 50-200; ELISA: 5000-10000

种属反应性: Rabbit

克隆性: Rabbit Polyclonal

· 亚型: Immunogen-specific rabbit IgG

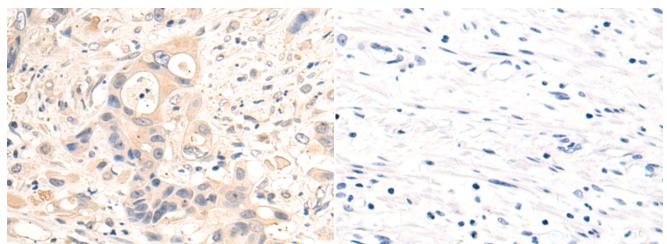
纯化: Antigen affinity purification

种属反应性: Human, Mouse, Rat

成分: PBS (without Mg2+ and Ca2+), pH 7.4, 150 mM NaCl, 0.05% Sodium Azide and 40% glycerol

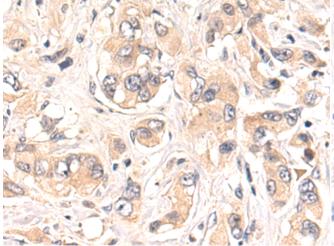
研究领域: Cell Biology

储存和运输: Store at -20°C. Avoid repeated freezing and thawing



Immunohistochemistry analysis of paraffin embedded Human colorectal cancer tissue using 218610(PSMB4 Antibody) at a dilution of 1/80(Cytoplasm).

In comparision with the IHC on the left, the same paraffin-embedded Human colorectal cancer tissue is first treated with the fusion protein and then with 218610(Anti-PSMB4 Antibody) at dilution 1/80.



The image on the left is immunohistochemistry of paraffinembedded Human breast cancer tissue using 218610(Anti-PSMB4 Antibody) at a dilution of 1/80.

In comparision with the IHC on the left, the same paraffin-embedded Human breast cancer tissue is first treated with fusion protein and then with D224804(Anti-PSMB4 Antibody) at dilution 1/80.



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