bs-41361R

[Primary Antibody]

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DATASHEET

Host: Rabbit Isotype: IgG

Clonality: Polyclonal

TMPRSS2 Rabbit pAb

GenelD: 7113 **SWISS:** 015393

Target: TMPRSS2

Immunogen: Recombinant human TMPRSS2 protein: 108-492/492.

Purification: affinity purified by Protein A

Concentration: 1mg/ml

Storage: 0.01M TBS (pH7.4) with 1% BSA, 0.02% Proclin300 and 50%

Shipped at 4°C. Store at -20°C for one year. Avoid repeated

freeze/thaw cycles.

Background: Extracellular proteases mediate the digestion of neighboring extracellular matrix components in initial tumor growth, allow desquamation of tumor cells into the surrounding environment, provide the basis for invasion of basement membranes in targeted metastatic organs and are required for release and activation of many growth and angiogenic factors (1,2). The TMPRSS2 gene encodes a 492 amino acid multimeric serine protease, which is mainly expressed in the mouse prostate and kidney, and is also expressed in the human small intestine, prostate, colon, stomach and salivary gland (3,4). TMPRSS2 contains several domains, including a serine protease domain of the S1 family, a scavenger receptor cysteine-rich domain of group A, an LDL receptor class A domain and a transmembrane domain (3). TMPRSS2 is expressed as a full length form and a cleaved protease domain (5) and its expression is increased by androgenic hormones. TMPRSS2 is also expressed in prostate carcinoma, suggesting that it may play a role in prostate carcinogenesis (6).

Applications: WB (1:500-2000)

ELISA (1:5000-10000)

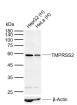
Reactivity: Human (predicted: Mouse,

Predicted 54 kDa

MW.:

Subcellular Location: Secreted ,Cell membrane

- VALIDATION IMAGES -



Sample: Lane 1: Human HepG2 cell lysates Lane 2: Human HeLa cell lysates Primary: Anti-TMPRSS2 (bs-41361R) at 1/1000 dilution Secondary: IRDve800CW Goat Anti-Rabbit IgG at 1/20000 dilution Predicted band size: 54 kDa Observed band size: 58 kDa

- SELECTED CITATIONS -

• [IF=3.8] Mohamed Elbadawy. et al. Establishment of a bat lung organoid culture model for studying bat-derived infectious diseases. SCIENTIFIC REPORTS. 2025 Feb 3;15(1):4035. IHC, IF; 39900611