

Recombinant Monoclonal Rabbit IgG Clone # 2959F Catalog Number: FAB55871S 100 µg

| Species Reactivity | Human |
|--------------------|---|
| Specificity | Detects human STEAP1 in direct ELISA. |
| Source | Recombinant Monoclonal Rabbit IgG Clone # 2959F |
| Purification | Protein A or G purified from hybridoma culture supernatant |
| lmmunogen | <i>E. coli-</i> derived recombinant human STEAP1 Met1-Trp71 Accession # Q9UHE8 |
| Conjugate | Alexa Fluor 750 Excitation Wavelength: 749 nm Emission Wavelength: 775 nm |
| Formulation | Supplied 0.2 mg/mL in a saline solution containing BSA and Sodium Azide. |
| | *Contains <0.1% Sodium Azide, which is not hazardous at this concentration according to GHS classifications. Refer to the Safety Data Sheet (SDS) for additional information and handling instructions. |

| APPLICATIONS | |
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| Please Note: Optimal dilutions should be | determined by each laboratory for each application. General Protocols are available in the Technical Information section on our website. |
| Flow Cytometry | Titration recommended for optimal concentration with starting range of 0.1-1 μg/1 million cells. Sample used for this experiment was HEK293 cells transfected with Human STEAP1 and eGFP vs irrelevant. |

| PREPARATION AND STORAGE | |
|-------------------------|---|
| Shipping | The product is shipped with polar packs. Upon receipt, store it immediately at the temperature recommended below. |
| Stability & Storage | Protect from light. Do not freeze. |
| | 12 months from date of receipt, 2 to 8 °C as supplied. |

BACKGROUND

STEAP1 (six-transmembrane epithelial antigen of the prostate-1) is a 40 kDa protein (predicted) of the STEAP family of metalloreductases. It is expressed mainly at cell-cell junctions between prostate secretory epithelium, and up-regulated in prostate and some bladder, colon and ovarian cancers and Ewing's sarcomas. Human STEAP1 is 339 amino acids (aa) in length. It contains a ferric oxidoreductase domain (aa 119-265) that includes transmembrane sequences (# 2, 3, 4, and part of 5, out of 6 sequences). Two splice forms diverge at aa 255, terminating after aa 58 or 259. Over aa 1-71, human and mouse STEAP1 share 68% aa identity.

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