# Anti-CD49E (Integrin Alpha 5) [SAM1]

Catalogue number: 154737 Sub-type: Primary antibody

Images:

#### Contributor

**Inventor:** Arnoud Sonnenberg

Institute: Netherlands Cancer Institute

Images:

### **Tool details**

#### \*FOR RESEARCH USE ONLY

Zancer Tools.org Name: Anti-CD49E (Integrin Alpha 5) [SAM1]

Alternate name: ITGA5

Class: Monoclonal

Conjugate: Unconjugated

Description: The ITGA5 gene encodes the integrin alpha 5 chain. Alpha chain 5 undergoes posttranslational cleavage in the extracellular domain to yield disulfide-linked light and heavy chains that join with beta 1 to form a fibronectin receptor. In addition to adhesion, integrins are known to participate

in cell-surface mediated signalling

Purpose: Parental cell: Organism: Tissue: Model: Gender:

Isotype: IgG2b Reactivity: Human

Selectivity: Host: Mouse

Immunogen: U 937 cells **Immunogen UNIPROT ID:** 

Sequence:

**Growth properties: Production details:** 

Formulation:

Recommended controls: **Bacterial resistance:** 

#### Selectable markers: Additional notes:

# **Target details**

Target: CD49E

#### **Target alternate names:**

Target background: The ITGA5 gene encodes the integrin alpha 5 chain. Alpha chain 5 undergoes post-translational cleavage in the extracellular domain to yield disulfide-linked light and heavy chains that join with beta 1 to form a fibronectin receptor. In addition to adhesion, integrins are known to participate in cell-surface mediated signalling

Molecular weight: 135 kDa

Ic50:

# **Applications**

Cancer Tools.org Application: FACS; IHC; IP

**Application notes:** 

## Handling

Format: Liquid

Concentration: 0.9-1.1 mg/ml

Passage number: **Growth medium:** Temperature: **Atmosphere:** Volume:

Storage medium:

Storage buffer: PBS with 0.02% azide Storage conditions: -15° C to -25° C Shipping conditions: Shipping at 4° C

### Related tools

Related tools:

### References

**References:** van der Velden et al. 1993. Head Neck. 15(2):133-46. PMID: 7680025. ; Ivanyi et al. 1989. J Pathol. 159(1):7-12. PMID: 2478685.

