

## ARG23254 anti-WC1 antibody [19.19] (FITC)

Package: 50 µg  
Store at: 4°C

### Summary

Product Description	FITC-conjugated Mouse Monoclonal antibody [19.19] recognizes WC1 Mouse anti Sheep WC1 antibody, clone 19. 19 recognizes the ovine WC1 cell surface antigen, a ~215 kDa glycoprotein, expressed by a population of gamma/delta T lymphocytes.
Tested Reactivity	Bov, Goat, Sheep
Tested Application	FACS
Host	Mouse
Clonality	Monoclonal
Clone	19.19
Isotype	IgG1
Target Name	WC1
Species	Sheep
Immunogen	Ovine thymocytes.
Conjugation	FITC
Alternate Names	M160; CD163B; CD163 antigen-like 1; Scavenger receptor cysteine-rich type 1 protein M160; CD antigen CD163b

### Application Instructions

Application table	Application	Dilution
	FACS	Neat

**Application Note** FACS: Use 10 µl of the suggested working dilution to label 10<sup>6</sup> cells in 100 µl.  
\* The dilutions indicate recommended starting dilutions and the optimal dilutions or concentrations should be determined by the scientist.

### Properties

Form	Liquid
Purification	Purification with Protein G.
Buffer	PBS, 0.09% Sodium azide and 1% BSA.
Preservative	0.09% Sodium azide
Stabilizer	1% BSA
Concentration	0.1 mg/ml
Storage instruction	Aliquot and store in the dark at 2-8°C. Keep protected from prolonged exposure to light. Avoid repeated freeze/thaw cycles. Suggest spin the vial prior to opening. The antibody solution should be gently mixed before use.

**Note**

For laboratory research only, not for drug, diagnostic or other use.

## Bioinformation

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Gene Symbol	CD163L1
Gene Full Name	CD163 molecule-like 1
Background	This gene encodes a member of the scavenger receptor cysteine-rich (SRCR) superfamily. Members of this family are secreted or membrane-anchored proteins mainly found in cells associated with the immune system. The SRCR family is defined by a 100-110 amino acid SRCR domain, which may mediate protein-protein interaction and ligand binding. The encoded protein contains twelve SRCR domains, a transmembrane region and a cytoplasmic domain. Alternative splicing results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Jul 2014]
Calculated Mw	159 kDa