

Human ACE-2 PE-conjugated Antibody

Monoclonal Rat IgG_{2B} Clone # 379131 Catalog Number: FAB9334P

100 Tests

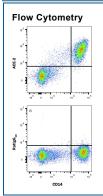
DESCRIPTION			
Species Reactivity	Human		
Specificity	Detects human ACE-2 in direct ELISAs.		
Source	Monoclonal Rat IgG _{2B} Clone # 379131		
Purification	Protein A or G purified from hybridoma culture supernatant		
Immunogen	Mouse myeloma cell line, NS0-derived human ACE-2 Gln18-Ser740 Accession # Q9BYF1		
Conjugate	Phycoerythrin Excitation Wavelength: 488 nm Emission Wavelength: 565-605 nm		
Formulation	Supplied in a saline solution containing BSA and Sodium Azide. See Certificate of Analysis for details.		
	*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

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.

	Recommended Concentration	Sample
Flow Cytometry	10 μL/10 ⁶ cells	Human PBMCs

DATA



Detection of ACE-2 in Human PBMC by Flow Cytometry. Human PBMC were stained with (A) Rat Anti-Human ACE-2 PE-conjugated Monoclonal Antibody (Catalog # FAB9334P) or (B) Rat IgG2b Isotype Control Antibody (Catalog # IC013P) and Mouse anti-Human CD14 APC-conjugated Monoclonal Antibody (Catalog # FAB3832A). Staining was performed using our Staining Membrane-associated Proteins protocol.

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.

Rev. 11/4/2022 Page 1 of 2





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BACKGROUND

Angiotensin I Converting Enzyme (ACE-2), also called ACEH (ACE homologue), is a dimeric, zinc-dependent metalloprotease of the ACE family that also includes somatic and germinal ACE (1, 2). ACE-2 mRNA is found at high levels in heart, testis, and kidney and at lower levels in a wide variety of tissues (1, 3). ACE-2 is the SARS-CoV and SARS-CoV2 Spike protein receptor *in vivo* (4-6), functions catalytically as a carboxypeptidase to cleave several substrates including angiotensins I and II, and acts as a partner for B0AT1-family amino acid transporters (1, 2). Through thesFAB9334Pe functions, ACE-2 has been shown to be involved in several diseases including SARS, COVID19, acute lung injury (4, 7), heart disease (8), liver and lung fibrosis (9), inflammatory lung disease (10), and cardiopulmonary disease (11). Full length ACE-2 protein includes an extracellular region composed of a single N-terminal peptidase domain and C-terminal collectrin-like domain (CLD), a transmembrane domain, and a short cytoplasmic tail (12). The N-terminal peptidase region is required for binding to SARS-CoV and SARSCoV2 spike proteins, while the CLD contains a region that promotes dimerization and association with amino acid transporters (2). The peptidase domain contains a long deep cleft that undergoes a large hinge-bending movement at substrate and inhibitor binding (12). Classical ACE inhibitors such as captopril and lisinopril do not inhibit ACE-2 activity and inhibitors of ACE-2 do not inhibit ACE activity (13).

References:

- 1. Kuba, K. et al. (2010) Pharmacol. Ther. 128:119.
- 2. Yan, et al. (2020) Science 367:1444.
- 3. Tipnis, S.R. et al. (2000) J. Biol. Chem. 275:33238.
- 4. Kuba, K. et al. (2005) Nature Med. 11:875.
- 5. Hoffman, M. et al. (2020) Cell. 181:1.
- 6. Wrapp, et al. (2020) Science 367:1260.
- 7. Imai, Y. et al. (2005) Nature 436:112.
- 8. Huang, L. et al. (2003) J. Biol. Chem. 278:15532.
- 9. Schrom, E. et al. (2017) Mol. Therapy Nuc. Acid 7:350.
- 10. Jia, H. et al. (2016) Shock. 46:239.
- 11. Cole-Jeffrey, C.T. et al. (2015) J. Cadiovasc. Pharmacol. 66:540.
- 12. Towler, P. et al. (2004) J. Biol. Chem. 279:17996.
- 13. Crackower, M.A. et al. (2002) Nature 417:822.