

AibGenesis™ Mouse Anti-BACE2 Antibody (CBMOAB-36733FYA)

Cat. No.: CBMOAB-36733FYA

Specifications

Host species	Mouse (<i>Mus musculus</i>)
Clone	MO36733FYA
Specificity	This antibody binds to Rhesus BACE2.
Format	Liquid or Lyophilized
Storage	Store at 4°C: short-term (1-2weeks) Store at -20°C: long-term and future use
Purity	> 90% was determined by SDS-PAGE
Purification	Purified with Protein A or G affinity chromatography

Application-Information

Application	WB, ELISA
Application Notes	ELISA: 1:1000-1:3000 Other applications are to be developed. The optimal dilution should be determined by the end user.

Target

Introduction	This gene encodes an integral membrane glycoprotein that functions as an aspartic protease. The encoded protein cleaves amyloid precursor protein into amyloid beta peptide, which is a critical step in the etiology of Alzheimer's disease and Down syndrome. The protein precursor is further processed into an active mature peptide. Alternative splicing results in multiple transcript variants. (From NCBI)
Product Overview	Mouse Anti-Rhesus BACE2 Antibody is a mouse antibody against BACE2. It can be used for BACE2 detection in Western Blot, Enzyme-Linked Immunosorbent Assay.
Protein Refseq	The length of the protein is 518 amino acids long. The sequence is shown below: MGALARALLPLLAQWLLRAAPELAPAPFTLPLRVAAATNRVVAPTGGPGTPAERRADGLALALEPALESAAGAAANFLAMVDNLQGDSGRGYYLEMLIGTPPQKLQILVDTGSSNFAVAGAPH

SYIDTYFDTERSSTYRSKGFDTVKYTGQSWTGFVGEDLVTPKGFNSFLVNIATI FESENFFLPGIKWNGI
LGLAYATLAKPSSSLETFFDSLVTQANIPNVFSMQMCGAGLPVA GSGTNGGSLVLGGIEPSLYKGDIWYTP
KEEWYYQIEILKLEIGGQSLNLD CREYNADKA IVDSGTLLRLPQKVFDAVVEAVARASLIPEFSDGFWTGS
QLACWTNSETPWSYFPKISI YLRDENSSRSFRITILPQLYIQPMMGAGLNYECYRFGISPSTNALVIGATVME
GFYVIFD RARKRVGFAASPCAEIAGA AVSEVSGPFSTEDIASNCVPAQSLSEPILWIVSYALMSVCG AILLVLI
VLLLLPFRQCRRPRDPEVVNDESSLVRHRWK.

Sub Cat	Clonalit y	Species	Reactivity	Applicati on	Clone	Conjugate	Size
CBMOAB-67365 FYA	Monoclon al	Zebrafish (<i>Danio rerio</i>)	WB, ELISA		MO67365F YA	Unconjugated; also available for Biotin; HRP; FITC and ot her labeled form.	100 µ g

Protein Refseq: The length of the protein is 462 amino acids long.

The sequence is show below: LQKNEAAAKSGLSLASDPAGIVNFLDMINNLKGDSEGRGYMQMIIGTPGQTLNILVDTGSSNFAVAAAAHPYITHYF
NRALSSTYQSTEK AVAVKYTQGEWEGELGDTLITIEPGPSGAINIAAILTSEGFFLPGINWQGILGLAYPLLARPDPSPVEFFNSVVRQTGIPDVFS
LQMC GAGVPASTTGDPAGGSLIMGGVEPTLHRGPIWYTPVLEEWYYQVEVLKLEVGDNQNLNLDCKEYNSDKAIVDSGTLLRLPANVFTAVVDT I
MQTSLIDDFSAGFFDGTCLACWMRGESPWRRFPKVSILRGNTS QSFRITILPQLYIQPVTDIDGTLCDFRFGISPSANGLVIGATVM EGFYVIFD
RAQKRVGFAVSTCAENGGVPFAEIA GPFLAEGVASDCTSGMTLREPMWVIAYTLMGVCAVLIILLVLLILPCRRHRDDEI TDESSLVRHRIK.

MO-AB-00257Y	Monoclon al	Chicken (<i>Gallus gallus</i>)	WB, ELISA		MO00257Y	Unconjugated; also available for Biotin; HRP; FITC and ot her labeled form.	100 µ g
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Protein Refseq: The length of the protein is 392 amino acids long. The sequence is show below: HLLLNTLSSTYQSQGIEVTVKYSQGSWTG
VLGTDVVTIPKGIDGRYTINIATILESENFFLPGVKWHGILGLAYDTLAKPSSSVETFFD SLVKQAKIPNIFSLQMCGAGLPVSGSGTNGGSLVLGGIE
PSLYKGNIWYTPIKEEWYYQVEILKLEVGQNLELD CREYNADKAIVDSGT TLLRLPQKVF GAVVQAIARTSLIQEFSSGFWSGSQLACWDKTER
PWSLFPKLSIYMRDENSSRSFRISILPQLYIQPILGIGENLQCYRF GISSSTNALVIGATVM EGFYVIFDRAQRRVGF VSPCAEVDGSPVSEIEGPF
TTTDVASNCVSSITFHEPVLWIASYALMSLCGIILLIL IILLIPPRCQHRYTDNDVVNDESSLVRHRWK.

MO-AB-01795H	Monoclon al	Frog (<i>Xenopus laevis</i>)	WB, ELISA		MO01795C	Unconjugated; also available for Biotin; HRP; FITC and ot her labeled form.	100 µ g
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Protein Refseq: The length of the protein is 500 amino acids long.

The sequence is show below: MKVPLVRLLLLCAAACASNKFIVPLNVSPAIEKGTLPVAPATPKDKPGLLLASDPGGTINFFSMVDNLAGDSGRGYY
LELLIGSPPQKVN ILVDTGSSNFAVAGSPNPVDNTFFDSKLSTSYQSLNTEVTVRYTQGSWTGLLGKD VVSIPKGVNGTFLINIASIFQSESFFLPNI
NWQGI LGLAYSTLAKPSSSVPEFFDSLQVENIPDVFSMQMCGAGQSSPGNGINAGSLVLGGVEPSLYKGNIWYTPITEEWYYQVEVLKFEVGG
Q RLNLDCTVYNSDKAIVDSGTLLRLPDKVFNAMVDAIVQTS LIQNFNAEFWAGLQLACWDKTQQPWNYFPDISIYLRDNTSRSFRLTLK PQLYI

Sub Cat	Clonalit y	Species Reactivity	Applicati on	Clone	Conjugate	Size
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QSVLTFQESLNCFRFGISQSASTLVIGATVMEGFYVIFDRAEKRVGFAVSSCAEVSGITVSEIAGPFGTSDVSSNCIARNPLREP IMWIIISYLSMLC
GMILLVLVILLLLSNRQRHDDMETINDESSLVQHRWK.

MO-AB-17421W	Monoclonal	Chimpanzee (<i>Pan troglodytes</i>)	WB, ELISA	MO17421W	Unconjugated; also available for Biotin; HRP; FITC and other labeled form.	100 µg
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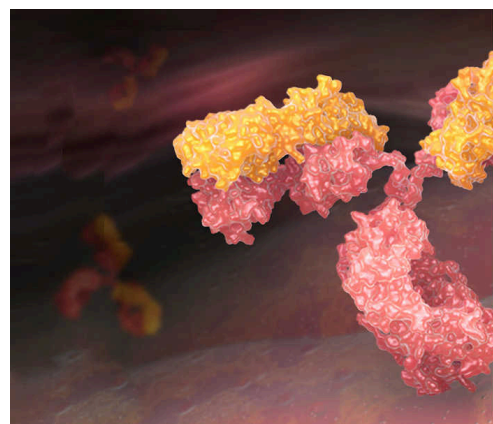
Protein Refseq: The length of the protein is 518 amino acids long.

The sequence is show below: MGALARALLPLLAQWLLRAAPELAPAPFTLPLRVAAATNRVVAPTGPGPSPAERHADGLALALEPALASPAGAAN
FLAMVDNLQGDSGR GYYLEMLIGTPPQKLQILVDTGSSNFAVAGTPHYSIDTYFDTERSSTYRSKGFDTVKYTGQSWTGFVGEDFVTIPKGFNT
SFLVNIATI FESENFPLPGIKWNGILGLAYATLAKPSSSLETFDLSVTQANIPNVFSMQMCGAGLPVAGSGTNGGSLVLGGIEPSLYKGDIWYTPIK
E EWYYQIEILKLEIGGQSLNLD CREYNADKAIVDSGTTLLRLPQKVFD AVEAVARASLIPEFSDGFWTGSQ LACWTNSETPWSYFPKISI YLRDE
NSSRSFRITILPQLYIQPMMGAGLNYECYRFGISPSTNALVIGATVMEGFYVIFDRAQKRVGFAASPCAIEIAGAAVSEISGPFST EDVASNCVPAQS
LSEPI LWIVSYALMSVCGAILLVLIVLLLPFRCQRRPRDPEVVNDESSLVRHRWK.

MO-AB-51712W	Monoclonal	Marmoset	WB, ELISA	MO51712W	Unconjugated; also available for Biotin; HRP; FITC and other labeled form.	100 µg
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Protein Refseq: The length of the protein is 518 amino acids long.

The sequence is show below: MGALARALLPLLAQWLLRAAPELAPAPFTLPLRVATATNRAAAPTSGPGTPAERRADGLALALEPALESAAGSAN
FLAMVDNLQGDSGR GYYLEMLIGTPPQKLQILVDTGSSNFAVAGAPHYSIDTYFDTERSSTYRSKGFDTVKYTGQSWTGFVGEDLVTIPKGFNS
SFLVNIATI FESENFPLPGIKWNGILGLAYATLAKPSSSLETFDSL VKANIPNVFSMQMCGAGLPVAGSGTNGGSLVLGGIEPSLYKGDIWYTPIK
E EWYYQIEILKLEIGGQSLNLD CREYNADKAIVDSGTTLLRLPQKVFD AVEAVARASLIPEFSDGFWTGSQ LACWANSETPWSYFPKISI YLRDE
NSSRSFRLTILPQLYIQPMMGAGLNYECYRFGISPSTNALVIGATVMEGFYVIFDRARKRVGFAASPCAIEIAGAAVSEISGPFST EDVASNCVPAQS
LSEPI LWIVSYALMSVCGAILLVLIVLLLPFRCQHRPRDPEIVNDESSLVRHRWK.



AibGenesis™

is an advanced AI-driven platform designed to create novel antibody sequences with unprecedented speed and precision. By integrating deep learning, structure prediction, and comprehensive immunological datasets, **AibGenesis™** intelligently designs antibodies optimized for affinity, stability, and developability. The platform generates antibody products that support basic scientific research, drug development, and diagnostic applications.