

Product datasheet

anti-AAV2 (intact particle) mouse monoclonal, A20, lyophilized, purified

Short overview

 Cat. No.
 61055

 Quantity
 50 μg

Concentration 50 µg/ml after reconstitution with 1 ml dist. water

Product description

HostMouseAntibody TypeMonoclonalIsotypeIgG3CloneA20

Immunogen AAV2 capsids

Formulation Lyophilized; reconstitute in 1 ml dist. water (final solution contains 0.09% sodium azide, 0.5% BSA

in PBS buffer, pH 7.4)

Binding affinity KD value (AAV2) = 2.6E-11 M

KD value (AAV3) = <1.0E-12 M

Synomym Adeno-associated virus 2; AAV-2

Conjugate Unconjugated

Purification Affinity chromatography

Storage before 2-8°C until indicated expiry date

reconstitution

Storage after 2-8°C

reconstitution

Intended use Research use only

Application Affinity chromatography, Dot blot, ELISA, ICC/IF, IP, Neutralization assay

Reactivity AAV2, AAV2 7m8, AAV3

No reactivity AAV1, AAV11, AAV12, AAV4, AAV5, AAV6, AAV7, AAV8, AAV9, AAVDJ, AAVrh10, AAVrh74

Applications

Affinity Chromatography Assay dependent

Dot Blot 1:500 (0.1 μg/ml; non-denaturing conditions)

ELISA Assay dependent

Immunocytochemistry (ICC) 1:20
Immunoprecipitation (IP) 1:5

Neutralization Assay EC50 ~5 ng/ml (AAV2) and ~3 ng/ml (AAV3) - assay dependent

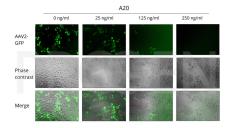
Background

For characterization of different stages of infection and very useful for the analysis of the AAV2 assembly process. A20 specifically reacts with AAV2 and AAV3, empty and full capsids. Recognizes a conformational epitope of assembled capsids, not present in denatured capsid proteins and native but unassembled capsid proteins. The antibody cannot be used for immunoblotting. Epitope mapping experiments (Wobus et al. 2000) identified four immunoreactive (discontinous) regions. The major reaction was attributed to sequence aa 369 to aa378 of AAV2 capsids. The antibody is also useful for neutralizing experiments.

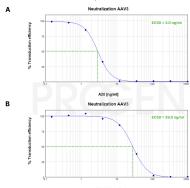
Wobus, C. E. et al. Monoclonal antibodies against the adeno-associated virus type 2 (AAV-2) capsid: epitope mapping and identification of capsid domains involved in AAV-2-cell interaction and neutralization of AAV-2 infection. J. Virol. 74, 9281-93 (2000). Moskalenko, M. et al. Epitope Mapping of Human Anti-Adeno-Associated Virus Type 2 Neutralizing Antibodies: Implications for Gene Therapy and Virus Structure. Journal Virol. 74, 1761-1766 (2000).

Limited Use Label License: Research Use OnlyProduct is exclusively licensed to PROGEN Biotechnik GmbH. The use of these products for the development, manufacturing and sale of secondary products/derivatives which are based on the purchased products and/or which include the purchased product require a royalty based sub-license agreement.

Product images



Neutralization of AAV2-GFP vectors with the A20 antibody (Cat. No. 61055). AAV infection was shown in HeLa cells and photos (GFP, CPE, merge) were taken ~48 h post infection. Neutralization was enhanced with increasing A20 concentration.



Neutralization of AAV3 with mouse monoclonal AAV2 antibody clone A20 (A) and human chimeric AAV2 antibody clone A20-h1 (B) by using AAV3-NanoLuc® viral particles from Promega. (A) anti-AAV2 (intact particle) mouse monoclonal, A20 (Cat. No. 61055) or (B) anti-AAV2, human chimeric, A20-h1 (Cat. No. 692379) were preincubated with AAV3-NanoLuc® viral particles for 30 min at RT at 300 rpm (antibody concentrations 0.2-3,000 ng/ml). HEK293 cells (100 μl) were plated at 200,000 cells/ml in DMEM + 1% FCS. Virus-antibody-mix (20 μl) was added to the cells and incubated for 16-24 h at 37°C. Extracellular NanoLuc Inhibitor and Nano-Glo® Live Cell Assay System (Promega) was added to the wells and incubated for 5 min at RT at 300 rpm. Luminescence was measured using an ID5-Reader and plotted with Softmax Pro 7.1 software to determine the EC50 values.

Serotype	Clone	Method	Residues						Reference
			200 - 299	300 - 399	400 - 499	500 - 599	600 - 699	700 - 731	Reference
AAV2	A20	Cryo-EM	253, 254, 258, 261, 262, 264	384, 385		548, 556	658 - 660	708, 717	1
		Peptide scanning	272 - 281	369 - 378		560 - 573			2
		Peptide insertion	261	381		534, 573			2, 3
		Site-directed mutagenesis	263, 264	384, 385		548		708	4

Several publications cited below describe the analysis of binding sites for the A20 antibody using different techniques. Multiple amino acids have been identified, that are very likely to be part of the binding site, especially those that were identified with more than one method (green boxes).

The amino acids of each binding site are located in different parts of the protein chains and are recognized as the epitope of the antibody only in the assembled capsid where they are in close proximity to each other and in the correct conformation. Note that they do not react with denaturated AAV2-VP.

1 McCraw et al. Structureof adeno-associated virus-2 in complex with neutralizing monoclonal antibody A20. Virology (2012) 431:40-9.

2 Wobus, C. E. et al. Monoclonal antibodies against the adeno-associated virus type 2 (AAV-2) capsid: epitope mapping and identification of capsid domains involved in AAV-2-cell interaction and neutralization of AAV-2 infection. J. Virol. 74, 928193 (2000).

3 Huttner et al. Genetic modifications of the adeno-associated virus type 2 capsid reduce the affinity and the neutralizing effects of human serum antibodies. Gene Ther (2003) 10:2139-47.

4 Lochrie et al. Mutations on the external surfaces of adeno-associated virus type 2 capsids that affect transduction and neutralization. J Virol. (2006) 80:821-34.

¹ McCraw et al. Structure of adeno-associated virus-2 in complex with neutralizing monoclonal antibody A20. Virology (2012) 431:40-9.

² Wobus et al. Monoclonal antibodies against the adeno-associated virus type 2 (AAV-2) capsid: epitope mapping and identification of capsid domains involved in AAV-2-cell interaction and neutralization of AAV-2 infection. J Virol (2000) 74:2951-93.

² Huttner et al. Genetic modifications of the adeno-associated virus type 2 capsid reduce the affinity and the neutralizing effects of human serum antibodies. Gene Ther (2003) 10:113:123-17.

^{*}Residues hoxed in green have been identified with at least two independent methods

References

Publication	Species	Application
Wistuba, A., Weger, S., Kern, A., Rgen, J. & Kleinschmidt, A. Intermediates of Adeno-Associated Virus Type 2 Assembly: Identification of Soluble Complexes Containing Rep and Cap Proteins. J. Virol. 69, 5311–5319 (1995).	AAV2	IP
Moskalenko, M. et al. Epitope mapping of Human Anti-Adeno-Associated Virus Type 2 Neutralizing Antibodies: Implications for Gene Therapy and Virus Structure. Jounal Virol. 74, 1761–1766 (2000).	AAV2	neutralization, epitope mapping
McCraw, D. M., O'Donnell, J. K., Taylor, K. A., Stagg, S. M. & Chapman, M. S. Structure of adeno-associated virus-2 in complex with neutralizing monoclonal antibody A20. Virology 431, 40–49 (2012).	AAV2	cryoEM
McCraw, D. M., O'Donnell, J. K., Taylor, K. A., Stagg, S. M. & Chapman, M. S. Structure of adeno-associated virus-2 in complex with neutralizing monoclonal antibody A20. Virology 431, 40–49 (2012).	AAV2	сгуоЕМ
Huttner, N. A. et al. Genetic modifications of the adeno-associated virus type 2 capsid reduce the affinity and the neutralizing effects of human serum antibodies. Gene Ther. 10, 2139–2147 (2003).	AAV2	epitope mapping