

Supplementary Information

A Siglec-like sialic acid binding motif found in an adenovirus capsid protein

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Supporting Tables S1 and S2

Supporting Figures S1 and S2

Table S1: List of all complexes analyzed:

1CT1; 1DBN; 1DFQ; 1DQF; 1E8U; 1F31; 1FV2; 1FV3; 1G1T; 1H7T; 1HGE; 1HGG; 1HGH; 1HGI; 1JSH; 1JSI; 1JSN; 1JSO; 1KQR; 1MQM; 1MQN; 1MWE; 1MX1; 1MX5; 1N1Y; 1NSC; 1QDD; 1QFO; 1QWJ; 1RVT; 1RVX; 1RVZ; 1S0I; 1S0J; 1SE3; 1URL; 1USR; 1USX; 1UXA; 1UXB; 1V0F; 1V3C; 1VPS; 1W0O; 1W0P; 1W1X; 1W20; 1W21; 1WGC; 1WW4; 1YA4; 1YA8; 1YAH; 1YAJ; 1YYN; 1Z4X; 2BAT; 2BER; 2BF6; 2C25; 2C4A; 2C4L; 2CHB; 2DF3; 2DQY; 2DQZ; 2DR0; 2DS0; 2EHI; 2H7C; 2HRL; 2HRQ; 2HRR; 2I2S; 2JH7; 2JHD; 2JHL; 2KMB; 2OSX; 2P3I; 2P3J; 2P3K; 2QWB; 2R61; 2RDG; 2RFT; 2RFU; 2V73; 2VU9; 2W68; 2W9L; 2WBV; 2WBW; 2WGC; 2WR1; 2WR2; 2WR3; 2WR4; 2WR7; 2WRB; 2WRE; 2WRF; 2WRG; 2WRH; 2Z8L; 2ZG1; 2ZG3; 2ZOE; 3B50; 3BWR; 3CHB; 3E81; 3F53; 3F5A; 3F5E; 3GVJ; 3GVK; 3GVL; 3H72; 3HMY; 3HN1; 3HTP; 3HTQ; 3HTT; 3JU4; 4HMG; 5HMG

Table S2: Oligosaccharide probes included in the glycan microarray

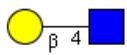
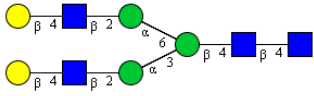
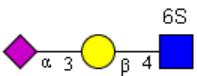
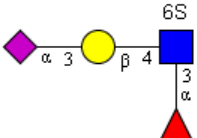
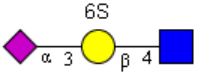
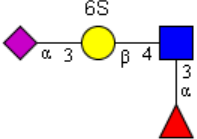
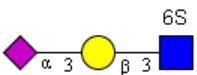

Chart #	Structure	IUPAC Nomenclature
1		Gal β (1-4)-GlcNAc β -ethyl-NH ₂
2		Gal β (1-4)-GlcNAc β (1-2)-Man α (1-3)-[Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂
3		NeuAc α (2-3)-Gal β (1-4)-6-sulfo-GlcNAc β -propyl-NH ₂
4		NeuAc α (2-3)-Gal β (1-4)-[Fuc α (1-3)]-6-O-sulfo-GlcNAc β -propyl-NH ₂
5		NeuAc α (2-3)-6-O-sulfo-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
6		NeuAc α (2-3)-6-O-sulfo-Gal β (1-4)-[Fuc α (1-3)]-GlcNAc β -propyl-NH ₂
7		NeuAc α (2-3)-Gal β (1-3)-6-O-sulfo-GlcNAc β -propyl-NH ₂
8		NeuAc α (2-3)-Gal β (1-4)-Glc β -ethyl-NH ₂


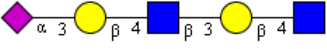
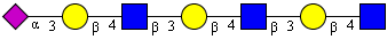
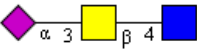

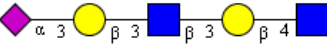
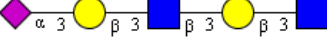
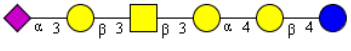
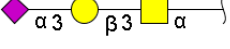
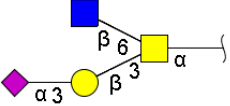
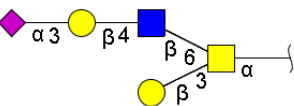
Chart #	Structure	IUPAC Nomenclature
9		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
10		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
11		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
12		NeuAc α (2-3)-GalNAc β (1-4)-GlcNAc β -ethyl-NH ₂
13		NeuAc α (2-3)-Gal β (1-3)-GlcNAc β -ethyl-NH ₂
14		NeuAc α (2-3)-Gal β (1-3)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
15		NeuAc α (2-3)-Gal β (1-3)-GlcNAc β (1-3)-Gal β (1-3)-GlcNAc β -ethyl-NH ₂
16		NeuAc α (2-3)-Gal β (1-3)-GalNAc β (1-3)-Gal α (1-4)-Gal β (1-4)-Glc β -ethyl-NH ₂
17		NeuAc α (2-3)-Gal β (1-3)-GalNAc α -Thr-NH ₂
18		NeuAc α (2-3)-Gal β (1-3)-[GlcNAc β (1-6)]-GalNAc α -Thr-NH ₂
19		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-6)-[Gal β (1-3)]-GalNAc α -Thr-NH ₂

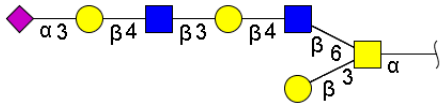
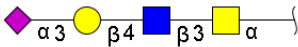
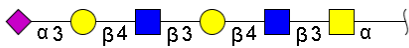
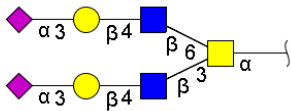
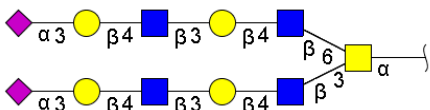
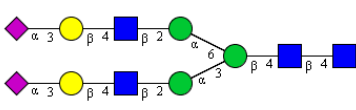
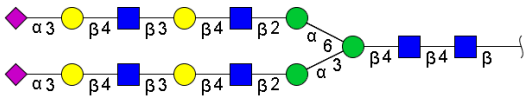
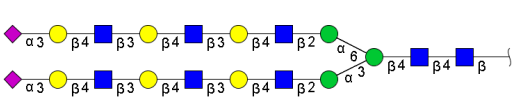
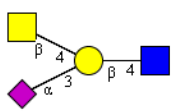
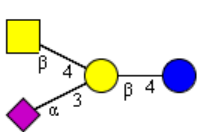
Chart #	Structure	IUPAC Nomenclature
20		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-6)-[Gal β (1-3)]-GalNAc α -Thr-NH ₂
21		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-GalNAc α -Thr-NH ₂
22		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-3)-GalNAc α -Thr-NH ₂
23		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-[NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-6)]-GalNAc α -Thr-NH ₂
24		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-3)-[NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-6)]-GalNAc α -Thr-NH ₂
25		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-3)-[NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂
26		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-3)-[NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂
27		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-3)-[NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂
28		NeuAc α (2-3)-[GalNAc β (1-4)]-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
29		NeuAc α (2-3)-[GalNAc β (1-4)]-Gal β (1-4)-Glc β -ethyl-NH ₂

Chart #	Structure	IUPAC Nomenclature
30		Gal β (1-3)-GalNAc β (1-4)-[NeuAc α (2-3)]-Gal β (1-4)-Glc β -ethyl-NH ₂
31		NeuAc α (2-3)-Gal β (1-4)-[Fuc α (1-3)]-GlcNAc β -propyl-NH ₂
32		NeuAc α (2-3)-Gal β (1-3)-[Fuc α (1-4)]-GlcNAc β (1-3)-Gal β (1-4)-[Fuc α (1-3)]-GlcNAc β -ethyl-NH ₂
33		NeuAc α (2-3)-Gal β (1-4)-[Fuc α (1-3)]-GlcNAc β (1-3)-Gal β (1-4)-[Fuc α (1-3)]-GlcNAc β -ethyl-NH ₂
34		NeuAc α (2-3)-Gal β (1-4)-[Fuc α (1-3)]-GlcNAc β (1-3)-Gal β (1-4)-[Fuc α (1-3)]-GlcNAc β (1-3)-Gal β (1-4)-[Fuc α (1-3)]-GlcNAc β -ethyl-NH ₂
35		NeuGc α (2-3)-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
36		NeuAc α (2-6)-Gal β (1-4)-6-sulfo-GlcNAc β -propyl-NH ₂
37		NeuAc α (2-6)-Gal β (1-4)-Glc β -ethyl-NH ₂


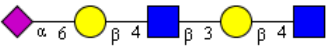
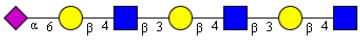
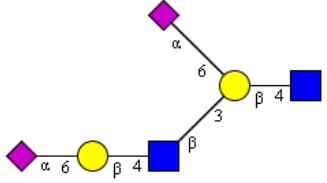
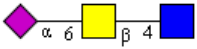
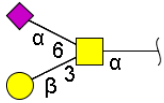
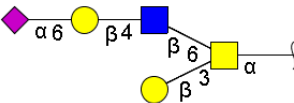
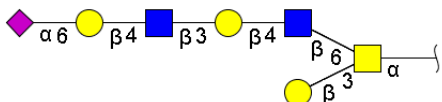
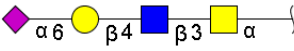
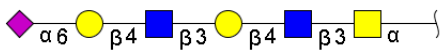
Chart #	Structure	IUPAC Nomenclature
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39		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
40		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
41		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-[NeuAc α (2-6)]-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
42		NeuAc α (2-6)-GalNAc β (1-4)-GlcNAc β -ethyl-NH ₂
43		NeuAc α (2-6)-[Gal β (1-3)]-GalNAc α -Thr-NH ₂
44		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-6)-[Gal β (1-3)]-GalNAc α -Thr-NH ₂
45		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-6)-[Gal β (1-3)]-GalNAc α -Thr-NH ₂
46		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-GalNAc α -Thr-NH ₂
47		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-3)-GalNAc α -Thr-NH ₂

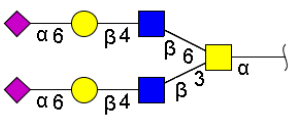
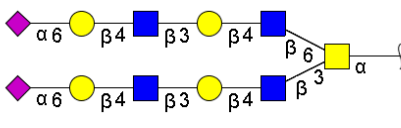
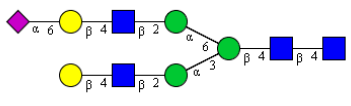
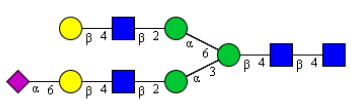
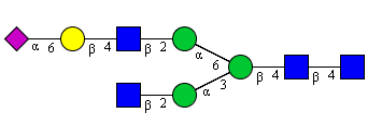
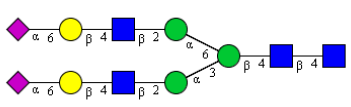
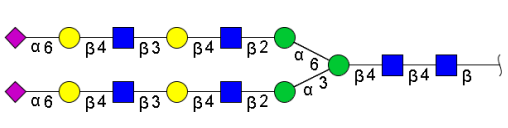
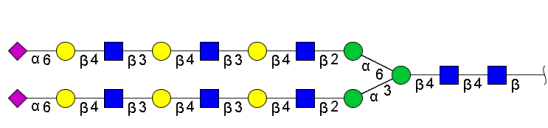
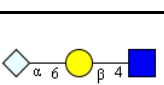
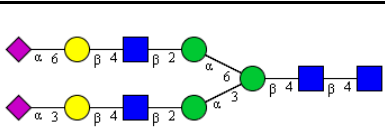
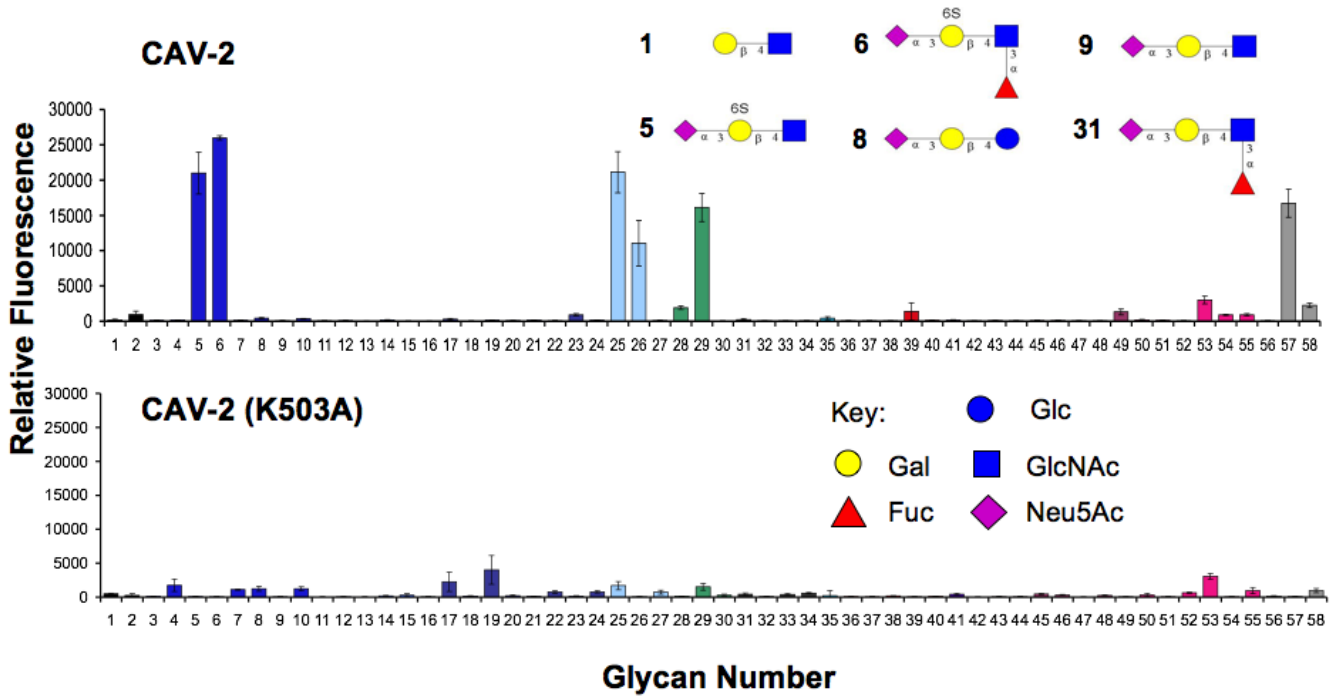
Chart #	Structure	IUPAC Nomenclature
48		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-[NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-6)]-GalNAc α -Thr-NH ₂
49		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-3)-[NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-6)]-GalNAc α -Thr-NH ₂
50		Gal β (1-4)-GlcNAc β (1-2)-Man α (1-3)-[NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂
51		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-3)-[Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂
52		GlcNAc β (1-2)-Man α (1-3)-[NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂
53		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-3)-[NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂
54		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-3)-[NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂
55		NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-3)-[NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂
56		NeuGc α (2-6)-Gal β (1-4)-GlcNAc β -ethyl-NH ₂
57		NeuAc α (2-3)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-3)-[NeuAc α (2-6)-Gal β (1-4)-GlcNAc β (1-2)-Man α (1-6)]-Man β (1-4)-GlcNAc β (1-4)-GlcNAc β -Asn-NH ₂

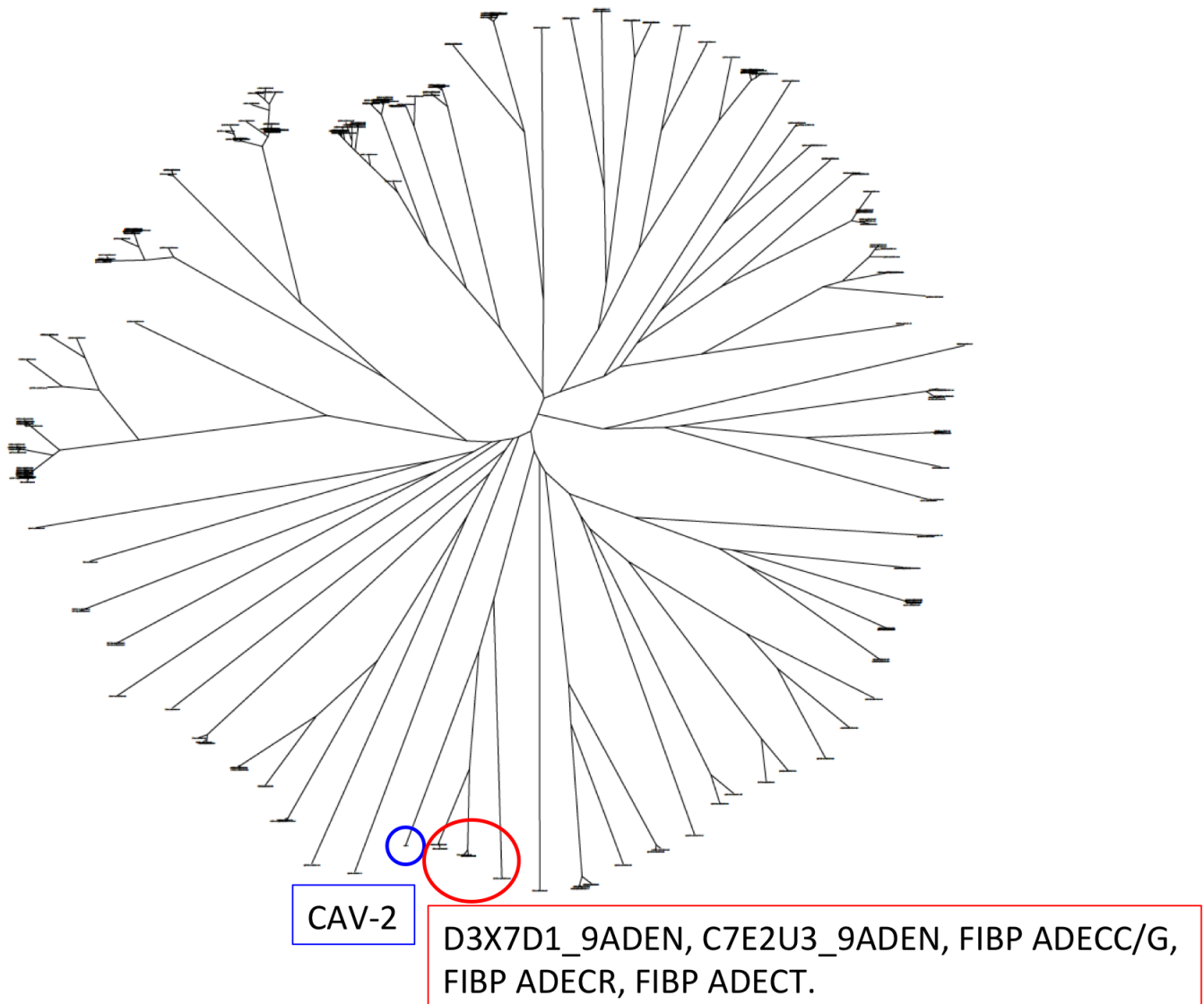
Chart #	Structure	IUPAC Nomenclature
58		<p>NeuAcα(2-6)-Galβ(1-4)-GlcNAcβ(1-2)-Manα(1-3)-[NeuAcα(2-3)-Galβ(1-4)-GlcNAcβ(1-2)-Manα(1-6)]-Manβ(1-4)-GlcNAcβ(1-4)-GlcNAcβ-Asn-NH₂</p>

Supporting Figure S1:



Glycan array analysis of the WT and K503A CAV-2 fiber knob protein at 500 ug/ml final concentration. (Top) In addition to sialosides **5** and **6** found at lower protein concentrations (50 ug/ml), structures bearing terminal α 2,3)-linked Neu5Ac **25**, **26**, **29** and **57** are detected by the fiber knob protein. (Bottom) The mutant CAV-2 (K503A) does not bind significantly to the printed sialoside array at high concentrations.

Supporting Figure S2:



Sequence alignment and family tree of adeno fiber knobs:

A multiple sequence alignment of the adenoviral fibre protein family (PF00541) provided by the PFAM database was extended by the CAV-2 sequence (Uniprot: Q65914) in ClustalX (version 2.0.9). The following sequences contain the essential arginine stretch FRFLN: D3X7D1_9ADEN, C7E2U3_9ADEN, FIBP ADECC, FIBP ADECG, FIBP ADECR, FIBP ADECT. The knobs also contain a lysine residue in a similar position. A family dendrogram resulting from the alignment file was visualized using drawtree as implemented on the mobyle server (Felsenstein, J. 1993. PHYLIP (Phylogeny Inference Package) version 3.5c. Distributed by the author. Department of Genetics, University of Washington, Seattle.).