

## SUPPLEMENTARY DATA

### STRUCTURE AND MUTAGENESIS OF NEURAL CELL ADHESION MOLECULE DOMAINS: EVIDENCE FOR FLEXIBILITY IN THE PLACEMENT OF POLYSIALIC ACID ATTACHMENT SITES.

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#### SUPPLEMENTAL EXPERIMENTAL PROCEDURES

*Analytical Ultracentrifugation of purified Ig5-FN1-Ig5-FN1* purified from *Escherichia coli* was dialyzed in 50 mM Tris, pH 8.0, 250 mM NaCl. Sedimentation velocity experiments were performed at the Center for Structural Biology at University of Illinois at Chicago. Briefly, 400  $\mu$ l of protein sample, ranging from 0.15 mg/ml to 1.5 mg/ml, was loaded into Epon double-sector centerpieces and centrifuged at 50,000 rpm in a Beckman ProteomeLab XL-1 centrifuge using an An-60 Ti rotor. Sedimentation was monitored by measuring absorbance at 280 nm. Buffer density and viscosity were determined using the program SEDNTERP (1) and data analyzed with the program SEDFIT (2).

*Immunofluorescence analysis of NCAM localization*-Cos-1 cells grown on 12-mm coverslips were transfected using 3  $\mu$ l Lipofectin in 300  $\mu$ l Opti-MEM 1, and 0.5  $\mu$ g NCAM cDNA. After a 6 h incubation, 1 ml DMEM, 10% FBS, was added to each well. Sixteen hours later, cell media was removed and cells were washed twice with 1 ml PBS. One milliliter of -20°C methanol was used to permeabilize cells to visualize both internal and cell surface structures. The coverslips were then incubated with 1 ml immunofluorescence blocking buffer (5% normal goat serum in PBS). To evaluate localization, cells were incubated with anti-V5 tag antibody diluted 1:250 in blocking buffer, washed with PBS four times for 5 min, followed by incubation with fluorescence isothiocyanate-conjugated goat anti-mouse IgG (Jackson ImmunoResearch) diluted 1:100 in blocking buffer. After further washing, coverslips were mounted on glass slides using mounting medium (15% (w/v) Vinol 205 polyvinyl alcohol, 33% (w/v) glycerol, 0.1% azide in PBS, pH 8.5). A Nikon Axiophot microscope equipped with epifluorescence illumination and a 60X oil immersion Plan Apochromat objective was used to visualize the cells and pictures taken with a SPOT RT color digital camera and processed with SPOT RT software version 3.5.1 (Diagnostic Instruments Inc, Sterling Heights, MI).

#### SUPPLEMENTAL FIGURE LEGENDS

**FIGURE 1. Analytical ultracentrifugation analysis of bacterially-expressed Ig5-FN1 indicates concentration-dependent dimer formation.** Sedimentation velocity analysis of purified Ig5-FN1 at a concentration of 0.15 mg/ml (blue) and 0.8 mg/ml (pink) using a 50 mM Tris, pH 8.0, 250 mM NaCl buffer.

**FIGURE 2. NCAM proteins containing engineered N-linked glycosylation sites localize to the Golgi and cell surface.** Cos-1 cells transiently expressing NCAM or various mutants were fixed with methanol and subcellular localization determined by indirect immunofluorescence microscopy using the anti-V5 epitope tag antibody. *Bar*, 10  $\mu$ m.

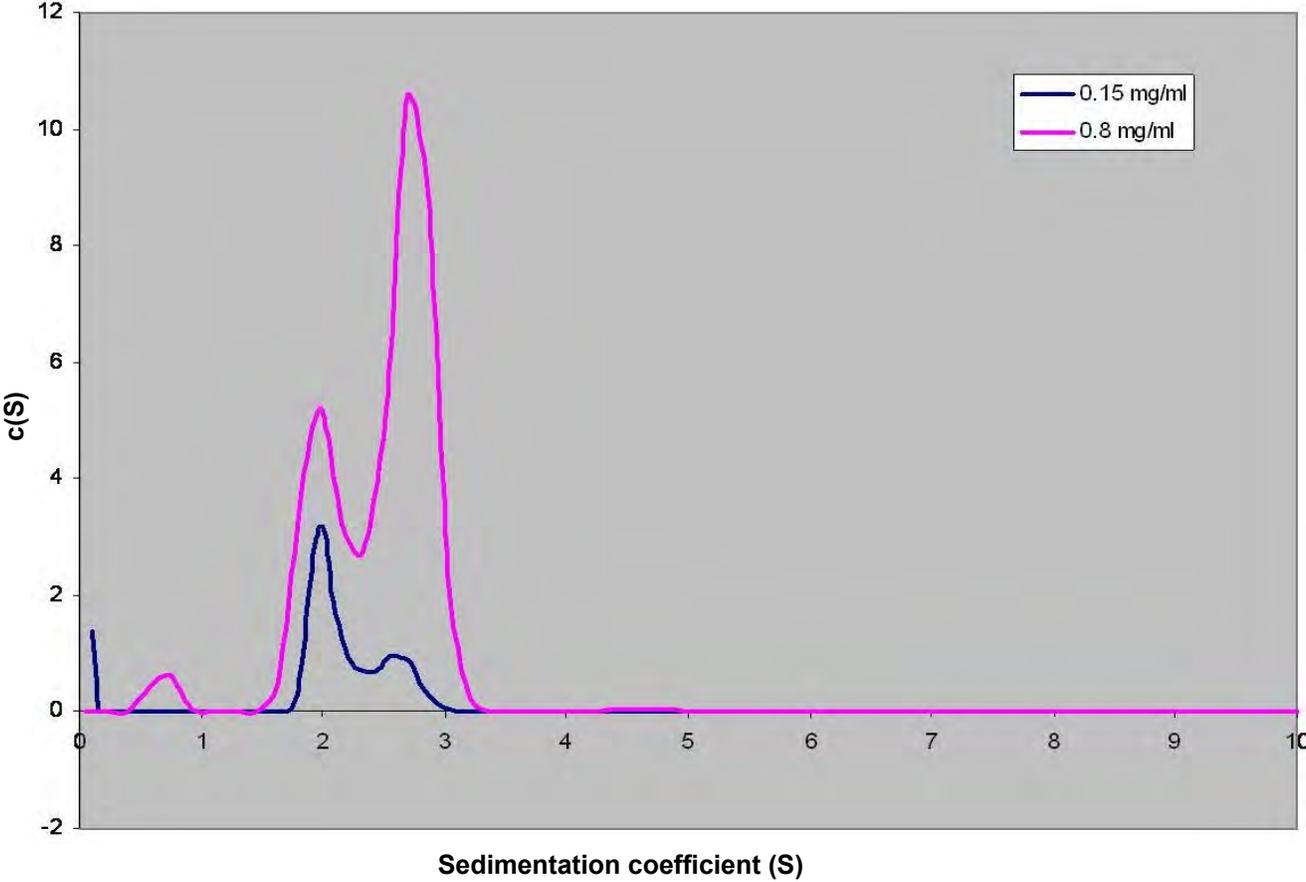
#### SUPPLEMENTAL REFERENCES

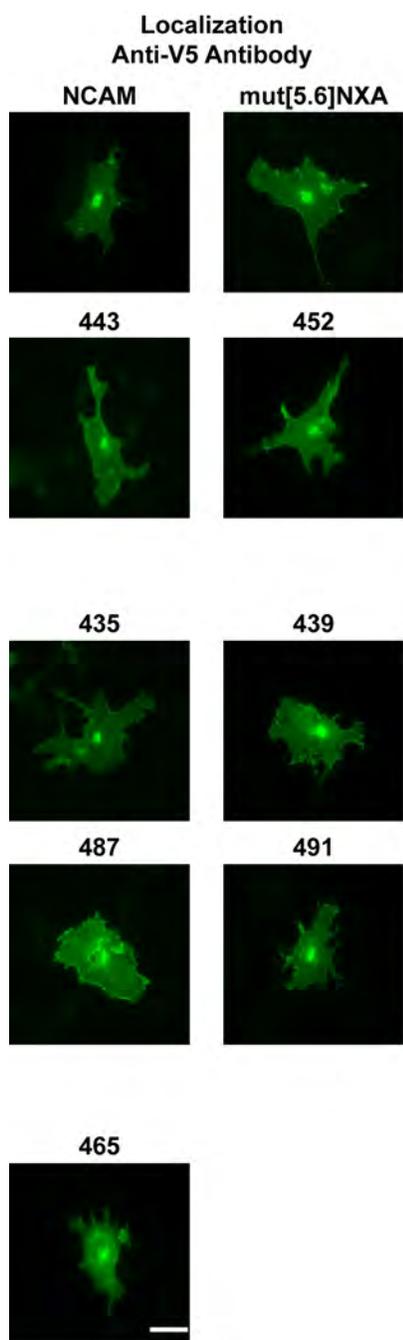
1. Hayes, D., Laue, T., and Philo J. <http://www.rasmb.bbri.org>
2. Schuck, P. *Biophys. J.* (2000) **78**, 1606-1619

**Supplemental Table 1:** List of primers used for site-directed mutagenesis of NCAM and Ig5-FN1

MUTANT	TEMPLATE	PRIMER	PRIMER SEQUENCE
Ig5-FN1 Asn[5.6]Gln	Ig5-FN1	N449Q	5'-GCCAAGCTCCCAGTACAGCAATATCAAG-3' 5'-CTTGATATTGCTGTACTGGGAGCTTGGC-3'
	Ig5-FN1 Asn[5]Gln	N478Q	5'-GGGAACTACCAATGTACTGCAGTGAAC-3' 5'-GTTCACTGCAGTACATTGGTAGTTCCC-3'
Ig5-FN1 Asn[4.5.6]Gln	Ig5-FN1 Asn[5.6]Gln	N423Q	5'-GGAACCAGGTGCAAATCACCTGCGAG-3' 5'-CTCGCAGGTGATTGTCACCTGGTTCC-3'
mut[5.6]NXA	NCAM	S451A	5'-CAAGCTCCAATTACGCCAATATCAAGATC-3' 5'-GATCTTGATATTGGCGTAATTGGAGCTTG-3'
	NCAM Asn[5]NXA	T480A	5'-GGGAACTACAACCTGTGCTGCAGTGAACCGC-3' 5'-GCGGTTCACTGCAGCACAGTTGTAGTTCCC-3'
sNCAM mut4 sIg5-FN1 mut4	sNCAM sIg5-FN1	T425A	5'-GAACCAGGTGAACATCGCCTGCGAGGTATTTG-3' 5'-CAAATACCTCGCAGGCGATGTTACCTGGTTC-3'
mut[5.6]NXA E465N	mut[5.6]NXA	E465N	5'-CTGCCAGCTATCTGAACGTGACCCCAGACTC-3' 5'-GAGTCTGGGGTCACGTTTCAGATAGCTGGCAG'-3'
mut[5.6]NXA Q443LL-NLS	mut[5.6]NXA	Q443LL-NLS	5'-CATGGTTTCGGGATGGCAACCTGTGCGCAAGCTCCAATTACG-3' 5'-CGTAATTGGAGCTTGGCGACAGGTTGCCATCCCGAAACCATG-3'
mut[5.6]NXA N452IK-NIS	mut[5.6]NXA	N452IK-NIS	5'-CAATTACGCCAATATCTCGATCTACAACACCCCC-3' 5'-GGGGGTGTTGTAGATCGAGATATTGGCGTAATTG-3'
mut[5.6]NXA F439RD-NRS	mut[5.6]NXA	F439RD-NRS	5'-GTGCCACGATCTCATGGAATCGGTCTGGCCAGCTGCTGCCAAG-3' 5'-CTTGGCAGCAGCTGGCCAGACCGATTCCATGAGATCGTGGCAC-3'
mut[5.6]NXA Q487N	mut[5.6]NXA	Q487N	5'-GTGAACCGCATTGGGAACGAGTCCTTGGAATTC-3' 5'-GAATCCAAGGACTCGTTCCCAATGCGGTTACAC-3'
mut[5.6]NXA E491FI-NFT	mut[5.6]NXA	E491FI-NFT	5'-GGGCAGGAGTCCTTGAACCTCACCTTGTTC AAGCAGAC-3' 5'-GTCTGCTTGAACAAGGGTGAAGTTCAAGGACTCCTGCCC-3'

### Interference Fits





Supplemental Figure 2