

Supplementary Materials: Nerve Growth Factor Signaling from Membrane Microdomains to the Nucleus: Differential Regulation by Caveolins

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Table S1. Summary of the experiments performed on different clones of PC12 cells stably expressing Cav-1 or Cav-2. Analyses of the different clones, performed as described in the Materials and Methods, yielded comparable results.

Clone	CAV-1						Clone	CAV-2				
	3	11	12	16	18	20		5	11	13	14	23
Proliferation			+	+			Proliferation	+				+
Differentiation	+	+	+	+	+	+	Differentiation	+	+	+	+	+
Western							Western					
Cav-1	++	+++	+	++	+++	+++	Cav-2	++	++	+++	++	+
TrkA PO4			+	+	+	+	TrkA PO4	+		+	+	+
pMAPK 42/44	+	+	+	+	+		pMAPK 42/44	+	+			+
pRSK	+	+		+	+	+	pRSK	+	+			
pCREB/CREB	+	+		+	+	+	pCREB/CREB	+	+			
p38				+			p38	+				
pAkt	+			+			pAkt	+	+			
pJNK	+			+			pJNK	+	+			
LRF		+		+		+	LRF	+	+			+
IF		+		+		+	IF	+	+			

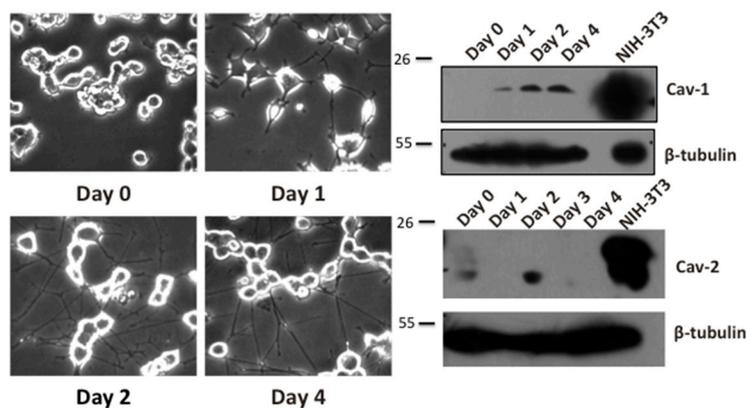


Figure S1. Expression of endogenous Cav-1 and Cav-2 in response to NGF. PC12 cells were exposed to NGF (50 ng/mL), harvested, and extracted for Western blot analysis for Cav-1 and Cav-2 as described in the Materials and Methods. Exponentially growing NIH-3T3 cells are used as a positive control. Beta-tubulin is used as loading control.

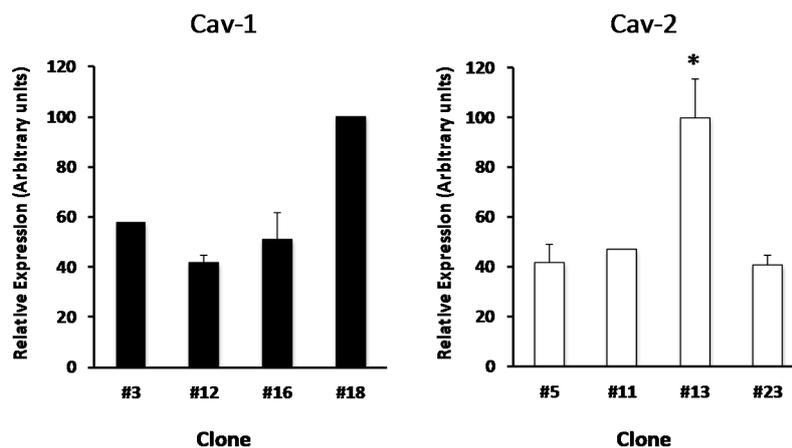


Figure S2. Relative expression levels of Cav-1 and Cav-2 in selected clones. PC12 clones expressing Cav-1 or Cav-2 were extracted for Western blot analysis and quantified using the gel analysis function of ImageJ as described in the Materials and Methods. Beta-tubulin was used as the loading control. The clone expressing the highest relative amount was considered 100%. (Mean \pm SEM derived from eight gels generated in four independent experiments. Statistical significance was ascertained using the unpaired, two-tail Student's *t*-Test comparing with clone Cav-1 #12 and Cav-2 #11 respectively. * $p < 0.05$.)