### **Supplemental Material**

# PCB 95 Modulates Calcium-Dependent Signaling Pathway Responsible for Activity-Dependent Dendritic Growth

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#### **Supplemental Methods**

#### **Materials**

PCB 95 (2,2',3,5',6-pentachlorobiphenyl, >99% purity) and PCB 66 (2,3',4,4'-tetrachlorobiphenyl, >99% purity) were purchased from AccuStandard, Inc. (New Haven, CT). FLA365, which blocks RyR channels (Mack et al. 1992) was a generous gift from Michele Chiesi (CIBA). Bicuculline was purchased from Sigma (St. Louis, MO); U0126, from Calbiochem (EMD4Biosciences, Rockland, MA); and STO-609, from Tocris Biosciences (Ellisville, Construction and characterization of the following expression vectors were Missouri). previously published (Wayman et al. 2006): Microtubule-associated protein 2B (MAP2B)enhanced green fluorescent protein (EGFP) fusion construct subcloned into the pCAGGS expression vector, pCAG-tomato fluorescent protein (TFP) and Wif expression vectors. Also previously described and characterized: plasmids encoding dominant negative (dn)CaMKI (K49E, T177A, IHQS286DDEE, F307A) (Wayman et al. 2004) and pCAGACREB (Arthur et al. 2004), and small hairpin (sh)RNA constructs targeting the  $\alpha$ ,  $\beta$ ,  $\gamma$  and  $\delta$  isoforms of Ca<sup>2+</sup>/calmodulin kinase (CaMK)I subcloned into the p702P vector (Wayman et al. 2006). A detailed description and characterization of RyR-specific and negative control siRNA primers (Invitrogen, Carlsbad, CA) is provided in the supplemental material for the companion paper (Wayman et al. 2012). Three different sets of siRNA primers with differing GC levels were designed for RyR1 and for RyR2 using the Invitrogen (Carlsbad, CA) stealth RNAi service. Each set (which was a mixture of 2 separate siRNAs) was tested for efficiency and specificity using HEK cells stably transfected with either Ryr1 or Ryr2. Data from these experiments was used to select one set against each RyR isoform to use in experiments in neuronal cell cultures (described in Table S1). The Invitrogen negative control kit were used as controls; sequences of these negative control siRNA are proprietary but are confirmed by Invitrogen to not be homologous to any expressed genes in the vertebrate transcriptome.

## Supplemental Material, Table 1. Sequences of RyR siRNA used in neuronal cells

| RyR1-specific siRNA           | RyR2-specific siRNA           |
|-------------------------------|-------------------------------|
| CGAGGAACUUCUACACGCUGCGAUU and | CAAGCGCAUCGAGAGGGUCUACUUU and |
| AAUCGCAGCGUGUAGAAGUUCCUCG     | AAAGUAGACCCUCUCGAUGCGCUUG     |

## WNT2 quantitative polymerase chain reaction (qPCR)

Total RNA was isolated from dissociated hippocampal neuron cultures (9 DIV) using Trizol (Invitrogen) according to manufacturer's instructions. RNA (50 ng to 3 mg) was reverse transcribed with Moloney murine leukemia virus reverse transcriptase (Invitrogen) and 50–250 ng random primers (Invitrogen). qPCR primers were designed by MIT's Primer3 software (http://frodo.wi.mit.edu/cgi-bin/primer3/primer3\_www.cgi) using default parameters with the following exceptions: rodent and simple repeat library was on, product size was 50–200bp, primer size was 18–27 bases, Tm was 66°C–72°C, maximum self-complementarity was 5, and maximum 30 complementarity was 3. Primer sequences for Wnt2 are: forward GGC GTT GTA TTT GCC ATC ACC AG; reverse GCT GTC CTT GCC ACT CCC TTT C. PCRs (10ml) contained 1 ml 103 PCR buffer (Invitrogen), 2.5 mM MgCl2, 200 mM dNTP (Roche), 0.125–0.25 mM primer (IDT), 13 SYBR green I (Invitrogen), and 1 U platinum Taq (Invitrogen). PCR was run on an Opticon OP346 (MJ Research) for one cycle at 95°C for 35 s, and 30–50 cycles at 94°C for 15 s; 68°C–70°C for 40 s. RT-PCR experiments were normalized to GAPDH RNA levels (other housekeeping genes showed similar results). All RT-PCR generated 100-fold higher levels of product relative to no reverse transcriptase controls.

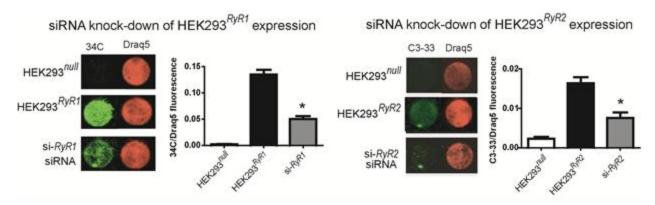
#### HEK cell culture and transfection

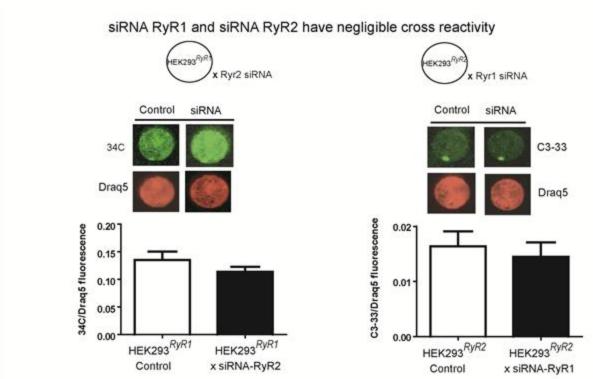
Human embryonic kidney (HEK 293) cells were maintained in DMEM medium supplemented with 2 mM glutamine,  $100 \mu g/ml$  streptomycin, 100 U/ml penicillin, 1 mM sodium pyruvate, and 10% FBS. Cells were transfected with full-length cDNA for rabbit Ryr1 or Ryr2 cloned into the

pCI-neo expression vector (Promega, Madison, WI) using Lipofectamine 2000 (Invitrogen) according to the manufacturer's instructions. Stably transfected cells were obtained by Geneticin sulfate (500  $\mu$ g/ml) selection for 2 wk. RyR1 and RyR2 expression were confirmed by immunocytochemistry using antibody 34C (University of Iowa Hybridoma Bank, Iowa City, IA) or C3-33 (generous gift of Dr. G. Meissner, University of North Carolina), respectively. Cultures were transfected with siRNA (25 pmol) using Lipofectamine RNAiMax (Invitrogen) per the manufacturer's instructions. RyR expression was quantified by in-cell western 48 h after transfection using anti-RyR1 antibody 34C (1:25) or anti-RyR2 antibody C3-33 (1:1000), and counterstaining with the nucleic acid binding dye DraQ5 (LICOR-Biosciences). Cultures were scanned using the LICOR Odyssey infrared imager. Total RyR signal (800 nm) was normalized to DraQ5 signal (700 nm) within the same well. Three wells were analyzed per treatment; significant differences were identified using one-way ANOVA with *post hoc* Newman-Keuls.

### **Supplemental References**

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Supplemental Material, Figure 1. Specificity of RyR siRNAs. Wild type HEK293 cell line (HEK293<sup>null</sup>), which lack detectable RyR expression, were stably transfected with *Ryr1* cDNA (HEK293<sup>RyR1</sup>, top left) or *Ryr2* cDNA (HEK293<sup>RyR2</sup>, top right).. RyR expression was quantified via densitometry of in-cell western blots 48 h after transfection with RyR isoform-specific siRNA constructs. Shown are representative images of in-cell western blots using antibodies 34C and C3-33 to probe RyR1 and RyR2, respectively, and Draq5 to stain nuclei; densitometric analyses of RyR1 and RyR2 expression normalized to Draq5 within each well are summarized in the bar graphs. As illustrated in bottom panels, siRNA 2-RyR1 did not knockdown RyR2 expression in HEK293<sup>RyR2</sup> cells; conversely, siRNA 1-RyR2 did not knockdown RyR1 expression in HEK293<sup>RyR1</sup> cells. Data expressed as the mean ± SD (n=3 wells).