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# $\alpha$ II Spectrin Forms a Periodic Cytoskeleton at the Axon Initial Segment and Is Required for Nervous System Function

Claire Yu-Mei Huang,<sup>1</sup> Chuansheng Zhang,<sup>1</sup> Tammy Szu-Yu Ho,<sup>1</sup> Juan Oses-Prieto,<sup>3</sup> Alma L. Burlingame,<sup>3</sup> Joshua Lalonde,<sup>2</sup> Jeffrey L. Noebels,<sup>1,2</sup>  Christophe Leterrier,<sup>4</sup> and  Matthew N. Rasband<sup>1</sup>

<sup>1</sup>Department of Neuroscience and <sup>2</sup>Department of Neurology, Baylor College of Medicine, Houston, Texas 77030, <sup>3</sup>Department of Pharmaceutical Chemistry, University of California, San Francisco, California 94158, and <sup>4</sup>NeuroCyto, NICN UMR7259, Aix Marseille Université, CNRS, 13344 cedex 15, Marseille, France

Spectrins form a submembranous cytoskeleton proposed to confer strength and flexibility to neurons and to participate in ion channel clustering at axon initial segments (AIS) and nodes of Ranvier. Neuronal spectrin cytoskeletons consist of diverse  $\beta$  subunits and  $\alpha$ II spectrin. Although  $\alpha$ II spectrin is found in neurons in both axonal and somatodendritic domains, using proteomics, biochemistry, and superresolution microscopy, we show that  $\alpha$ II and  $\beta$ IV spectrin interact and form a periodic AIS cytoskeleton. To determine the role of spectrins in the nervous system, we generated *Sptan1*<sup>fl/fl</sup> mice for deletion of CNS  $\alpha$ II spectrin. We analyzed  $\alpha$ II spectrin-deficient mice of both sexes and found that loss of  $\alpha$ II spectrin causes profound reductions in all  $\beta$  spectrins.  $\alpha$ II spectrin-deficient mice die before 1 month of age and have disrupted AIS and many other neurological impairments including seizures, disrupted cortical lamination, and widespread neurodegeneration. These results demonstrate the importance of the spectrin cytoskeleton both at the AIS and throughout the nervous system.

**Key words:** ankyrin; axon; axon initial segment; cytoskeleton; node of Ranvier; spectrin

## Significance Statement

Spectrin cytoskeletons play diverse roles in neurons, including assembly of excitable domains such as the axon initial segment (AIS) and nodes of Ranvier. However, the molecular composition and structure of these cytoskeletons remain poorly understood. Here, we show that  $\alpha$ II spectrin partners with  $\beta$ IV spectrin to form a periodic cytoskeleton at the AIS. Using a new  $\alpha$ II spectrin conditional knock-out mouse, we show that  $\alpha$ II spectrin is required for AIS assembly, neuronal excitability, cortical lamination, and to protect against neurodegeneration. These results demonstrate the broad importance of spectrin cytoskeletons for nervous system function and development and have important implications for nervous system injuries and diseases because disruption of the spectrin cytoskeleton is a common molecular pathology.

## Introduction

Neurons have complex cytoskeletons regulating neuronal morphology, trafficking of cargoes, and assembly of specialized membrane domains and compartments. Spectrins comprise

a submembranous cytoskeleton proposed to confer strength and flexibility to the membrane (Bennett and Baines, 2001; Xu et al., 2013). Spectrins are widely expressed and exist as tetramers consisting of two  $\alpha$  and two  $\beta$  subunits. There are five vertebrate  $\beta$  spectrins:  $\beta$ I– $\beta$ IV are found in neurons and  $\beta$ V is found in hair cells (Legendre et al., 2008; Zhang et al., 2014).  $\beta$  spectrins are not uniformly distributed throughout cells, but are often located in specialized domains. For example,  $\beta$ IV spectrin forms a complex with the scaffolding protein ankyrinG (ankG) at axon initial segments (AIS) and nodes of Ranvier to cluster the voltage-gated ion channels necessary for action potential initiation and propagation (Berghs et al., 2000; Yang et al., 2007). Due to their high degree of homology, some  $\beta$  spectrins may compensate for one another (Ho et al., 2014). However,  $\alpha$ II spectrin is the only  $\alpha$  spectrin detected in the nervous system (Zhang et al., 2014), suggesting that, depending on its  $\beta$  spectrin binding partner, it must play important roles in a variety of cellular contexts.

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Correspondence should be addressed to Matthew N. Rasband, Department of Neuroscience, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030. E-mail: rasband@bcm.edu.

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Human mutations in  $\alpha$ II spectrin cause West syndrome, a severe infantile epilepsy that includes seizures, hypomyelination, and brain atrophy (Saito et al., 2010; Writzl et al., 2012; Tohyama et al., 2015). Mice and zebrafish lacking  $\alpha$ II spectrin are embryonic and larval lethal due to cardiac and nervous system malformations (Voas et al., 2007; Stankewich et al., 2011). Many nervous system injuries and diseases including stroke and traumatic brain injury converge on spectrin degradation because spectrins are potent substrates for the  $\text{Ca}^{2+}$ -dependent protease calpain (Siman et al., 1984; Schafer et al., 2009). Therefore, breakdown of spectrins may be a common molecular pathology in neurodegenerative diseases and injuries.

$\beta$ IV spectrin is enriched at AIS and nodes and forms a periodic lattice with actin (Xu et al., 2013; Leterrier et al., 2015; D'Este et al., 2017).  $\alpha$ II spectrin is found at paranodal junctions of myelinated axons and is transiently located at developing zebrafish nodes of Ranvier (Ogawa et al., 2006; Voas et al., 2007). Although  $\alpha$ II spectrin can coimmunoprecipitate  $\beta$ IV spectrin from the brain (Uemoto et al., 2007), it has not been found at AIS. Although loss of  $\alpha$ II spectrin disrupts node formation in zebrafish (Voas et al., 2007), whether loss of spectrins is sufficient to cause axon degeneration in vertebrates remains unknown. Furthermore, because  $\alpha$ II spectrin-deficient zebrafish and mice are larval and embryonic lethal, respectively, the postnatal functions of  $\alpha$ II spectrin remain poorly understood. Using proteomics, biochemistry, and superresolution microscopy, we found that  $\alpha$ II spectrin forms a periodic cytoskeleton and protein complex with  $\beta$ IV spectrin at AIS. To determine the function of neuronal spectrins, we generated mice lacking CNS  $\alpha$ II spectrin. We found the  $\alpha$ II spectrin-dependent submembranous cytoskeleton is required for many aspects of brain development and function, including assembly and maintenance of the AIS, cortical lamination, axon integrity, and proper neuronal excitability. Our results begin to reveal the many functions of the spectrin cytoskeleton in nervous system development and function.

## Materials and Methods

**Animals.** To generate a conditional allele for *Sptan1*, a targeting vector was designed to replace an 8.5 kb genomic fragment with loxP sites flanking exon 8 of *Sptan1*. This targeting vector construct was electroporated into embryonic stem cells (ESCs) derived from 129/Sv mice. Colonies were selected and screened by Southern blot. ESCs with correct homologous recombinations were injected into blastocysts and then transferred to foster female mice. Chimeric mice were crossed with C57BL/6 mice to confirm germline transmission. Upon crossing with cre recombinase-expressing mice, exon 8 will be excised and results in the premature termination of the *Sptan1* transcript. *Sptan1*<sup>fl/fl</sup> mice were crossed with *Nestin-cre* mice to spatially eliminate  $\alpha$ II-spectrin in the CNS. The *Sptan1*<sup>fl/fl</sup> mice were maintained on a 129/Sv and C57BL/6 mixed background. All experiments involving animals complied with National Institutes of Health (NIH) guidelines and were approved by the Animal Care and Use Committee at Baylor College of Medicine. PCR genotyping of mice was performed using the following primers: Floxed *Sptan1*: forward, 5'-AACAGTCACACCCTCTGAGTGCCA-3'; reverse, 5'-ATTCAGTGAAAGCTGAGAAG CCAG; *Nestin-cre*: forward, 5'-TGCTGTTTCACTGGTTATGCGG-3'; reverse, 5'-TTGCCCTGTTC ACTATCCAG-3'.

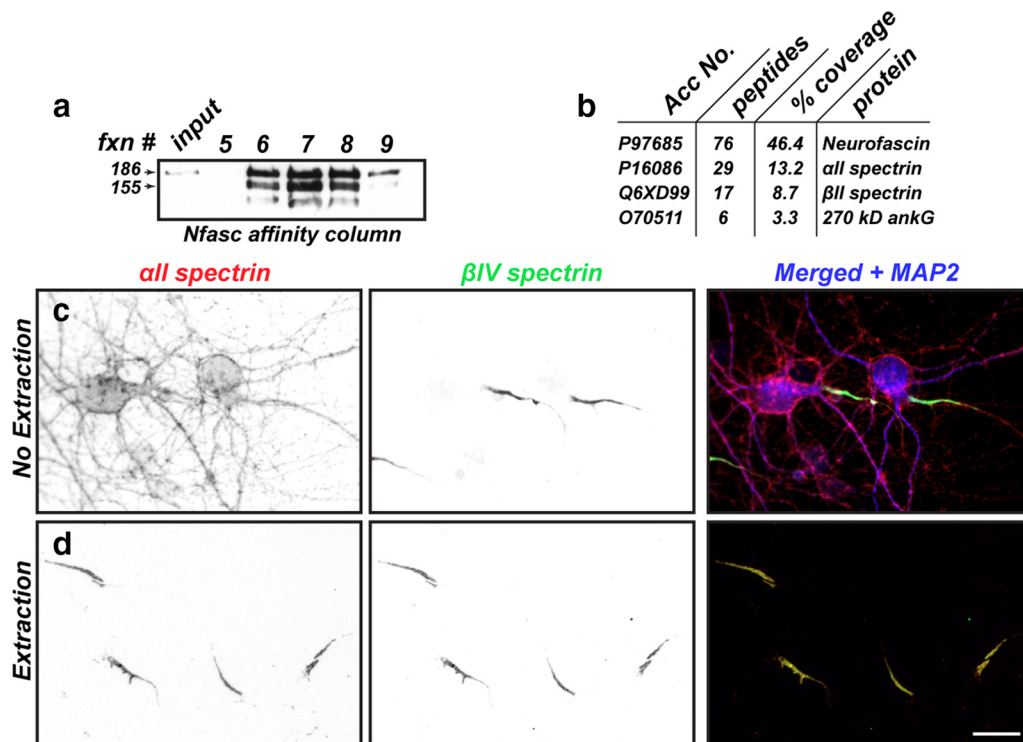
**DNA constructs.**  $\alpha$ II-spectrin-EGFP was a gift from Dr. Michael Stankewich (Yale University). The full-length  $\beta$ IV $\Sigma$ 6 spectrin with N-terminal Myc tag was provided by M. Komada (Tokyo Institute of Technology). The truncated myc-tagged  $\beta$ IV $\Sigma$ 6 spectrin was subcloned as described previously (Yang et al., 2007). The full-length  $\beta$ IV $\Sigma$ 1 spectrin with N-terminal Myc tag was subcloned and inserted into pCS3 + MT. The full-length rat 270 kDa ankG with C-terminal GFP tag (AnkG-GFP) was a gift from Dr. V. Bennett (Duke University).

**Antibodies.** The following primary antibodies were used: mouse monoclonal antibodies to  $\alpha$ II-spectrin (clone D8B7, Biologend; RRID:AB\_2564660),  $\beta$ II-spectrin (BD Bioscience; RRID:AB\_399853),  $\alpha$ I-spectrin (clone 17C7, Abcam; RRID:AB\_2194328), NeuN (EMD Millipore; RRID:AB\_2298772), and Reelin (EMD Millipore; RRID:AB\_2179313). The pan-NF monoclonal antibody (clone L11A/41; RRID:AB\_10672370) was made by immunizing mice with a glutathione S-transferase fusion protein containing the intracellular domain of rat neurofascin. The Kv1.2 (K14/16; RRID:AB\_2296313) antibodies were purchased from the University of California–Davis/NIH NeuroMab Facility. Rabbit polyclonal antibodies were as follows:  $\beta$ I-spectrin (Novus Biologicals; RRID:AB\_11036737),  $\beta$ III-spectrin (Novus Biologicals; RRID:AB\_10005990), neurofilament M (EMD Millipore; RRID:AB\_912011), Cux1 (Santa Cruz Biotechnology; RRID:AB\_2087003), Foxp2 (Abcam; RRID:AB\_2107107), Calbindin (D-28K, Swant; RRID:AB\_2314111),  $\beta$ APP (Thermo Fisher Scientific; RRID:AB\_2533902), GFP (A-11122, Thermo Fisher Scientific; RRID:AB\_221569), and  $\beta$ IV spectrin (generated against the peptide sequence DRAEELPRRRR ERQE found in the C-terminal “specific domain”; RRID:AB\_2315634). Rat monoclonal antibody to Ctip2 (25B6; ab18465; RRID:AB\_2064130) was from Abcam. Goat polyclonal antibodies against connective tissue growth factor (ctgf, sc-14939; RRID:AB\_638805) were from Santa Cruz Biotechnology. Secondary antibodies Alexa Fluor 350, 488, and 594 were from Thermo Fisher Scientific.

**Immunofluorescence microscopy.** Cultured neurons were fixed in 4% paraformaldehyde (PFA), pH 7.2–7.4, and immunostained as described previously (Hedstrom et al., 2008). For immunostaining of nervous system tissues, brains were collected at the indicated time points and fixed in 4% PFA on ice for 1 h. Brains were then immersed in 20% sucrose in 0.1 M PB overnight at 4°C and then embedded in optimal cutting temperature compound (Tissue-Tek; #4583) and sectioned using a cryostat (Thermo Fisher Scientific; Cryostar NX70) on glass coverslips for immunostaining. Procedures for immunofluorescence labeling were performed as described previously (Ogawa et al., 2006). Images of immunofluorescence were captured using an Axio-imager Z1 microscope (Carl Zeiss) or Axio-observer Z1 microscope (Carl Zeiss) fitted with an AxioCam digital camera (Carl Zeiss). Images were taken using 20 $\times$ /0.8 numerical aperture (NA), 40 $\times$ /1.0 NA, 40 $\times$ /0.75 NA, or 63 $\times$ /1.4 NA objectives. Images were then collected by Zen (Carl Zeiss) acquisition software. Measurement of fluorescence intensity and Sholl analysis was performed using FIJI and Zen software. In some cases, Z-stacks and 3D reconstructions were performed using Zen software.

**Superresolution Stochastic Optical Reconstruction Microscopy (STORM) and DNA-PAINT.** Twelve days *in vitro* (DIV) neurons were fixed using 4% PFA for 10 min. After blocking, they were incubated with primary antibodies overnight at 4°C, then with secondary antibodies for 1 h at room temperature. STORM imaging was performed on an N-STORM microscope (Nikon Instruments). Coverslips were imaged in STORM buffer containing the following: 50 mM Tris, pH 8, 10 mM NaCl, 10% glucose, 100 mM MEA, 3.5 U/ml pyranose oxidase, and 40 mg/ml catalase. The sample was illuminated continuously at 647 nm (full power) and 30,000–60,000 images were acquired at 67 Hz, with progressive reactivation by simultaneous 405 nm illumination (Leterrier et al., 2015). For DNA-PAINT labeling, fixed coverslips were incubated with primary antibodies and then with DNA-conjugated secondary antibodies (Ultivue) for 1 h at room temperature. Samples were imaged in imaging buffer with 0.32 nM Imager-650 and 0.16 nM Imager-560 (Ultivue). The sample was alternatively illuminated at 647 and 561 nm (full laser power) and 20,000–30,000 images of each channel were acquired at 25–33 Hz.

**Immunoprecipitation and immunoblotting.** COS-7 cells were transfected with  $\alpha$ II-spectrin-EGFP along with different Myc-tagged  $\beta$ IV-spectrin  $\Sigma$ 1,  $\Sigma$ 6, and truncated plasmids (Yang et al., 2007). For a negative control, cells were transfected with GFP along with  $\beta$ IV-spectrin  $\Sigma$ 1 or  $\beta$ IV-spectrin  $\Sigma$ 6. Two days after transfection, cells were collected and lysed with 250  $\mu$ l of lysis buffer (20 mM Tris-HCl, pH 8.0, 10 mM EDTA, pH 8.0, 150 mM NaCl) containing 1% Triton X-100 and protease inhibitors at 4°C for 1 h with agitation. Cell lysates were then centrifuged at 13,000  $\times$  g at 4°C for 10 min. The soluble supernatants were incubated overnight with GFP antibody-coupled protein A or G agarose beads (GE Healthcare). The beads were washed 7 times with 1 ml of lysis buffer and



**Figure 1.**  $\alpha$ II spectrin is found at the AIS. **a**, Immunoblot of neurofascin (Nfasc) purified from rat brain using an anti-Nfasc affinity column. Fractions 5–9 are shown. **b**, Proteins purified from the Nfasc affinity column identified by mass spectrometry. **c**, Immunostaining of 12 DIV rat hippocampal neurons using antibodies against  $\alpha$ II spectrin (red),  $\beta$ IV spectrin (green), and MAP2 (blue). **d**, Detergent-extracted 12 DIV rat hippocampal neurons immunostained using antibodies against  $\alpha$ II spectrin (red),  $\beta$ IV spectrin (green), and MAP2 (blue). Scale bar, 20  $\mu$ m.

then eluted with 20  $\mu$ l of 2 $\times$  reducing sample buffer at 100°C for 5 min. The immunoprecipitates and the total supernatant as input were separated by SDS-PAGE, transferred to nitrocellulose membrane (GE Healthcare), and detected by immunoblotting using Myc or GFP antibody.

**Neurofascin affinity purification.** The neurofascin affinity column was made by coupling mouse monoclonal antibodies to neurofascin (clone A12/18.1, RRID:AB\_2282826) using the Affi-Gel Hz Immunoaffinity kit (Bio-Rad) according to the manufacturer's instructions. Rat brain membrane homogenates were solubilized using RIPA buffer (1% Triton X-100, 1% sodium deoxycholate, 0.1% SDS, 0.15 M NaCl, 20 mM HEPES, 10 mM EDTA, pH 8.0, and protease inhibitors) at 4°C. The detergent-soluble fraction was then run over the affinity column three times. The column was washed with 50 ml of RIPA buffer, followed by elution of bound proteins using 10 ml of 0.1 M glycine, pH 2.5. Eluted proteins were collected in 1 ml fractions containing 100  $\mu$ l of 1 M Tris, pH 8.7. The proteins in each fraction were then precipitated by ethanol precipitation.

**Reverse-phase LC-MS/MS analysis.** Purified proteins were size fractionated by SDS PAGE and protein bands were excised. Purified proteins were then identified using mass spectrometry as described previously (Ogawa et al., 2010).

**In utero electroporation.** *In utero* electroporation was performed as described previously (Hedstrom et al., 2007). GFP and  $\alpha$ II spectrin shRNA plasmids (Galiano et al., 2012) were introduced into rat embryos at embryonic day 16 (E16). After recovery of the mother, newborn pups were killed at postnatal day 7 (P7) for immunostaining.

**EEG.** Mice were implanted with bilateral silver wire cortical EEG electrodes under Avertin anesthesia as described previously (Olivetti et al., 2014). Recordings were obtained after recovery with 8 h monitoring periods for 1 week after recovery using a Stellate Systems Harmonie 10 video EEG system. EEGs were analyzed by two observers blinded to genotype.

**Hippocampal neurons.** Primary cultures of hippocampal neurons were obtained from E18.5 Sprague Dawley rat embryos or P1–P2 mouse neonates. Hippocampi were dissected and dissociated. Neurons were then plated onto poly-D-lysine- and laminin-coated glass coverslips. Hippocampal neurons were cultured in Neurobasal medium (Life Technologies) containing 1% Glutamax (Life Technologies), 1% penicillin/

streptomycin (Life Technologies), and 2% B27 supplement (Life Technologies) in a 5% CO<sub>2</sub> incubator. For detergent extraction studies, hippocampal neurons and teased nerves were incubated with ice-cold PBS containing 0.5% Triton X-100 at 4°C for 30 min or 1 h, respectively. Neurons or nerves were then washed with 1 $\times$  PBS 3 times and then processed using the regular immunofluorescence procedures. In some cases, we infected hippocampal neurons using AAV-cre-GFP virus (AAV-EF1 $\alpha$ -EGFP-T2A-iCre (serotype: AAV-DJ8)) or adenovirus to express control shRNA or  $\alpha$ II spectrin shRNA. The AAV virus was obtained from the Intellectual and Developmental Disabilities Research Center Neuroconnectivity Core at Baylor College of Medicine and produced by Dr. B. Arenkiel (Baylor College of Medicine); the adenovirus was as described previously (Galiano et al., 2012).

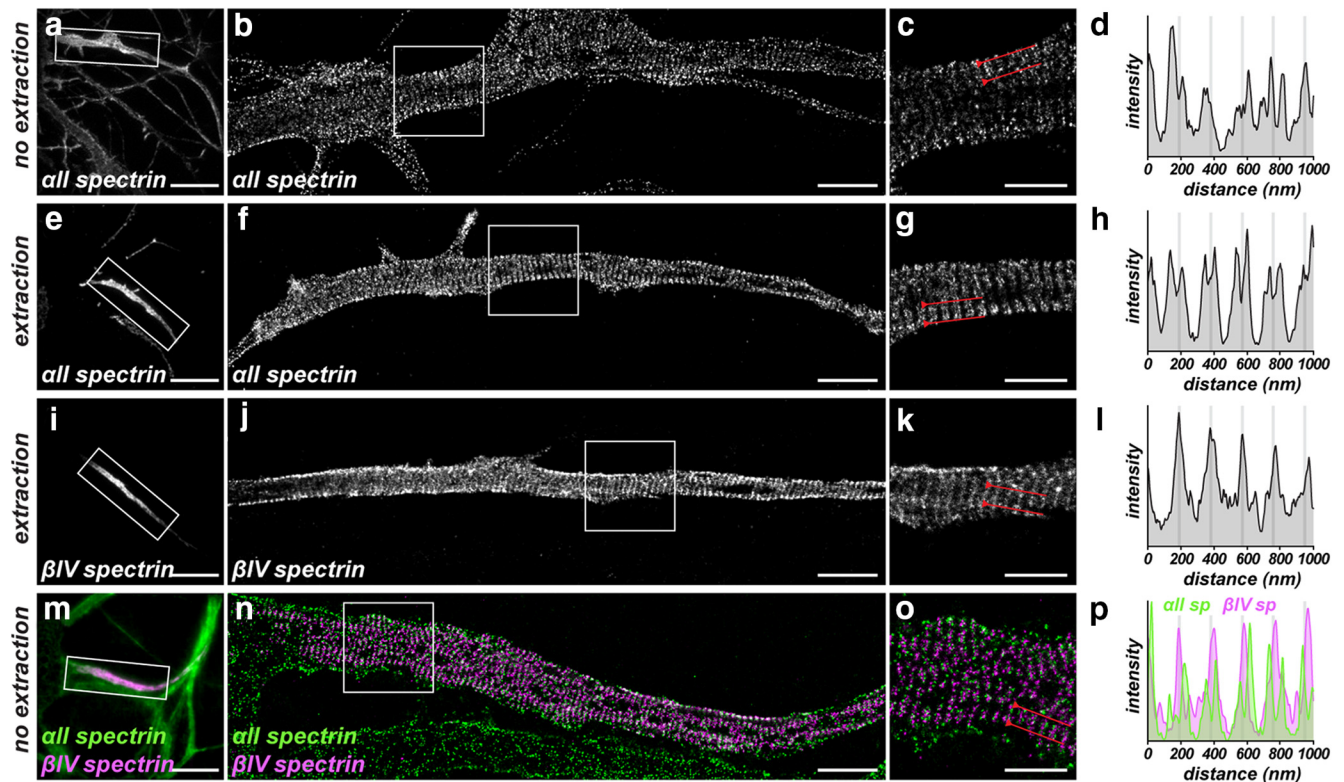
**Statistics.** No statistical methods were used to predetermine sample sizes, but our sample sizes are similar to those reported previously (Suzuki et al., 2013). All statistical analyses were performed using GraphPad Prism software or Microsoft Excel. Error bars in figures are  $\pm$  SEM. Unpaired, two-tailed Student's *t* test was used for statistical analyses unless otherwise specified. Except for EEG, data collection and analyses were not performed blinded to the conditions of the experiments. Data distribution was assumed to be normal.

## Results

### $\alpha$ II spectrin forms a periodic cytoskeleton with $\beta$ IV spectrin at the AIS

AIS have two main functions: to initiate action potentials and to regulate neuronal polarity. These properties require the ankG-dependent AIS cytoskeleton (Hedstrom et al., 2008; Kole et al., 2008). However, apart from ankG and  $\beta$ IV spectrin, the composition of the AIS cytoskeleton remains poorly understood. To identify new AIS proteins that may contribute to AIS structure and function, we made an anti-neurofascin antibody-conjugated affinity column to purify NF186 and its associated proteins from rat brain homogenate; NF186 is an ankyrin-binding cell adhesion molecule clustered at the AIS. Immunoblots of fractions eluted





**Figure 2.**  $\alpha$ II spectrin forms a periodic cytoskeleton with  $\beta$ IV spectrin at AIS. **a–d**, Conventional fluorescence (**a**) and STORM (**b, c**) imaging of 12 DIV cultured hippocampal neurons labeled with antibodies against  $\alpha$ II spectrin. The box in **a** corresponds to the STORM image shown in **b**. The box in **b** corresponds to the STORM image shown in **c**. The region between the lines in **c** was used to generate an  $\alpha$ II spectrin intensity profile **d**. Scale bars: **a**, 10  $\mu$ m; **b**, 2  $\mu$ m; **c**, 1  $\mu$ m. **e–h**, Conventional fluorescence (**e**) and STORM (**f, g**) imaging of detergent-extracted 12 DIV cultured hippocampal neurons labeled with antibodies against  $\alpha$ II spectrin. The box in **e** corresponds to the STORM image shown in **f**. The box in **f** corresponds to the STORM image shown in **g**. The region between the lines in **g** was used to generate an  $\alpha$ II spectrin intensity profile (**h**). Scale bars: **e**, 10  $\mu$ m; **f**, 2  $\mu$ m; **g**, 1  $\mu$ m. **i–l**, Conventional fluorescence (**i**) and STORM (**j, k**) imaging of detergent-extracted 12 DIV cultured hippocampal neurons labeled with antibodies against  $\beta$ IV spectrin. The box in **i** corresponds to the STORM image shown in **j**. The box in **j** corresponds to the STORM image shown in **k**. The region between the lines in **k** was used to generate a  $\beta$ IV spectrin intensity profile (**l**). Scale bars: **i**, 10  $\mu$ m; **j**, 2  $\mu$ m; **k**, 1  $\mu$ m. **m–p**, Conventional fluorescence (**m**) obtained by projecting frames from the PAINT acquisition sequence and DNA-PAINT (**n, o**) imaging of 12 DIV cultured hippocampal neurons labeled with antibodies against  $\alpha$ II spectrin and  $\beta$ IV spectrin. The box in **m** corresponds to the PAINT image shown in **n**. The box in **n** corresponds to the PAINT image shown in **o**. The region between the lines in **o** was used to generate  $\alpha$ II spectrin (green) and  $\beta$ IV spectrin (magenta) intensity profiles (**p**). Scale bars: **m**, 10  $\mu$ m; **n**, 2  $\mu$ m; **o**, 1  $\mu$ m.

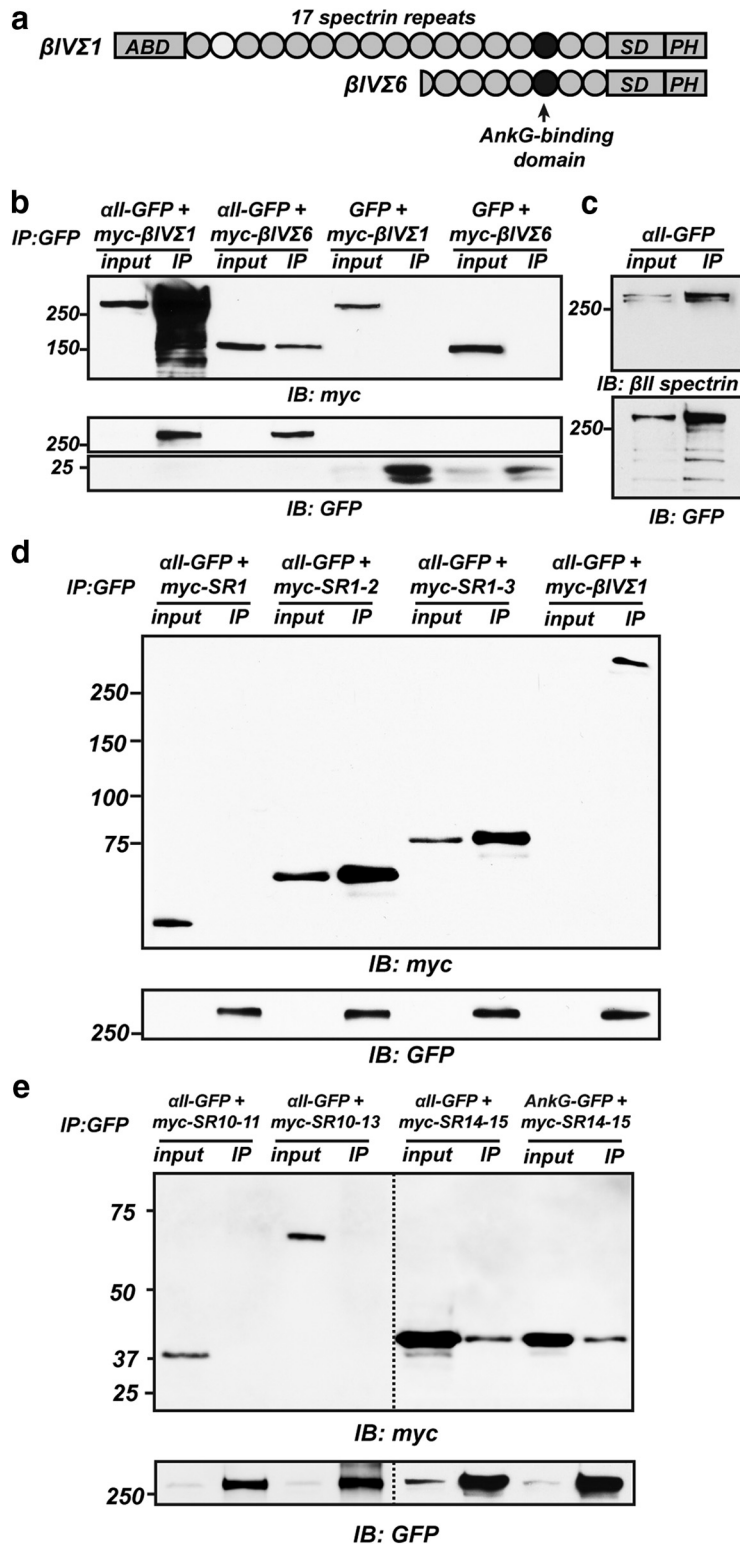
from this affinity column showed a strong chromatographic peak for both neuronal NF186 and oligodendroglial NF155 (Fig. 1*a*). We pooled fractions 6–8, precipitated the proteins, and then size fractionated them by SDS-PAGE. We excised proteins from the gels and then performed mass spectrometry and found, as expected, that the most abundant protein purified was neurofascin (Fig. 1*b*). The next most abundant protein was  $\alpha$ II spectrin.  $\alpha$ II spectrin attracted our attention because  $\beta$ IV spectrin is highly enriched at the AIS (Fig. 1*c*) and can form a complex with  $\alpha$ II spectrin (Uemoto et al., 2007). However,  $\alpha$ II spectrin has not been reported at the AIS. Consistent with our previous study (Galiano et al., 2012), immunostaining of  $\alpha$ II spectrin in 12 DIV rat hippocampal neurons revealed widespread and diffuse labeling rather than specific enrichment at the AIS (Fig. 1*c*). Remarkably, detergent extraction of 12 DIV neurons before fixation revealed a pool of detergent-resistant  $\alpha$ II spectrin only at the AIS that also colocalized with  $\beta$ IV spectrin (Fig. 1*d*).

To define the precise distribution of  $\alpha$ II spectrin in the AIS, we used STORM (Rust et al., 2006). STORM imaging of nondetergent extracted and  $\alpha$ II spectrin-labeled AIS revealed a periodic organization like that previously reported for ankG,  $\beta$ IV spectrin, Na<sup>+</sup> channels, and NF186 (Fig. 2*a–d*; Xu et al., 2013; Zhong et al., 2014; Leterrier et al., 2015). However, detergent extraction showed an even clearer periodicity (Fig. 2*e–h*) with two resolvable peaks of immunofluorescence very close to one another (Fig.

2*g, h*). In contrast, STORM imaging of detergent-extracted AIS using antibodies against the C terminus of  $\beta$ IV spectrin showed only one major peak with a periodicity of  $\sim$ 190 nm (Fig. 2*i–l*). To visualize  $\alpha$ II spectrin and  $\beta$ IV spectrin simultaneously, we combined DNA-PAINT labeling (Jungmann et al., 2014) with STORM imaging (Fig. 2*m–p*). Intensity profiles revealed that the two peaks of  $\alpha$ II spectrin immunoreactivity flank  $\beta$ IV spectrin labeling (Fig. 2*p*). The labeling patterns are consistent with a model in which spectrin tetramers space actin rings because the  $\alpha$ II spectrin antibodies were generated against the SH3 domain located after spectrin repeat (SR) 9, resulting in two resolvable peaks of fluorescence intensity surrounding the single peak resulting from labeling of the C terminus of  $\beta$ IV spectrin. Furthermore, these two-color DNA-PAINT images demonstrate directly *in situ* that  $\alpha$  spectrins are arranged head-to-head in the spectrin tetramer. Therefore,  $\alpha$ II spectrin is found at the AIS, where it forms a periodic cytoskeleton with  $\beta$ IV spectrin.

#### $\alpha$ II spectrin interacts with $\beta$ IV spectrin splice variants

How do  $\alpha$ II and  $\beta$ IV spectrin interact? Alternative splicing generates up to six  $\beta$ IV spectrin splice variants (Berghs et al., 2000). Among these,  $\beta$ IV $\Sigma$ 1 and  $\beta$ IV $\Sigma$ 6 are at AIS and nodes of Ranvier (Komada and Soriano, 2002; Lacas-Gervais et al., 2004; Yoshimura et al., 2016).  $\beta$ IV $\Sigma$ 1 is the longest splice variant and includes an N-terminal actin-binding domain, 17 SRs, a “spe-



**Figure 3.**  $\alpha$ II spectrin and  $\beta$ IV spectrin splice variants can interact through SR1-2 and SR14-15. **a**, Diagram of  $\beta$ IV spectrin splice variants and their domains. ABD, actin-binding domain; SD, specific domain; PH, pleckstrin homology domain. **b**, Cotransfection of COS-7 cells with GFP-tagged  $\alpha$ II spectrin and myc-tagged  $\beta$ IV $\Sigma$ 1 or  $\beta$ IV $\Sigma$ 6. Immunoprecipitation of  $\alpha$ II spectrin-GFP coimmunoprecipitates both myc- $\beta$ IV $\Sigma$ 1 and myc- $\beta$ IV $\Sigma$ 6. Immunoblots were performed using antibodies against myc and GFP. Molecular weights are shown in kDa. **c**, Transfection of GFP-tagged  $\alpha$ II spectrin in COS-7 cells. Immunoprecipitation of  $\alpha$ II spectrin-GFP coimmunoprecipitates endogenous  $\beta$ II spectrin. **d**, Cotransfection of COS-7 cells with GFP-tagged  $\alpha$ II spectrin and myc-tagged SR1, SR1-2, or SR1-3. Immunoprecipitation using anti-GFP antibodies coimmunoprecipitates myc-SR1-2 and myc-SR1-3. Molecular weights are shown in kilodaltons (kDa). Coimmunoprecipitation between  $\alpha$ II spectrin-GFP and myc- $\beta$ IV $\Sigma$ 1 is included as a positive control. **e**, Cotransfection of COS-7 cells with  $\alpha$ II spectrin-GFP and myc-tagged SR10-11, SR10-13 or SR14-15.  $\alpha$ II spectrin-GFP coimmunoprecipitates myc-SR14-15. Molecular weights are shown in kilodaltons (kDa). Coimmunoprecipitation between Ankg-GFP and myc-SR14-15 is included as a positive control.

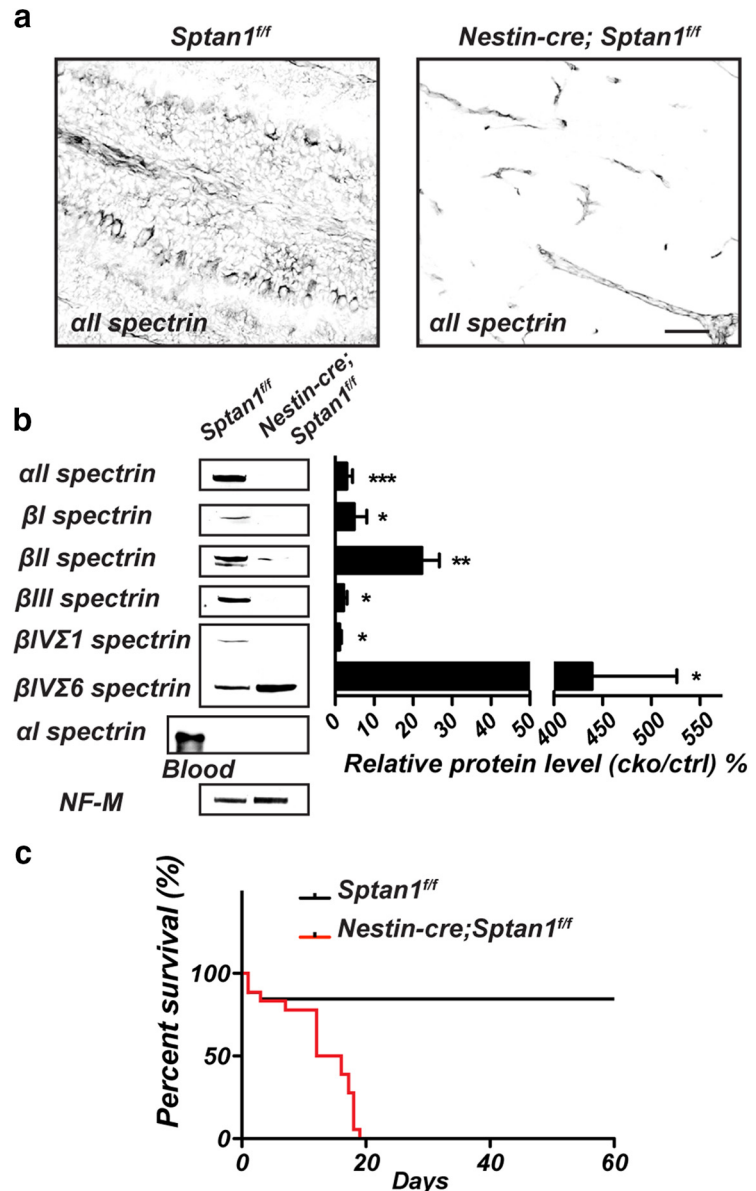
cific” domain (SD), and a C-terminal pleckstrin homology domain. In contrast,  $\beta$ IV $\Sigma$ 6 is an N-terminal truncation lacking the actin-binding domain and the first 9 SRs (Fig. 3a). Although  $\beta$ IV $\Sigma$ 1 and  $\beta$ IV $\Sigma$ 6 are both expressed throughout brain development,  $\beta$ IV $\Sigma$ 6 is expressed at the highest levels and is the major splice variant at mature AIS and nodes of Ranvier (Yoshimura et al., 2016).  $\alpha$ II spectrin was shown previously to coimmunoprecipitate both  $\beta$ IV $\Sigma$ 1 and  $\beta$ IV $\Sigma$ 6 from the brain (Uemoto et al., 2007). To confirm that  $\alpha$ II spectrin can interact with  $\beta$ IV spectrin splice variants, we cotransfected  $\alpha$ II spectrin-GFP ( $\alpha$ II-GFP) and myc- $\beta$ IV spectrins (myc- $\beta$ IV $\Sigma$ 1 or myc- $\beta$ IV $\Sigma$ 6) into COS-7 cells. Immunoprecipitation using antibodies against GFP confirmed that  $\alpha$ II spectrin binds to both  $\beta$ IV $\Sigma$ 1 and  $\beta$ IV $\Sigma$ 6 (Fig. 3b), but  $\beta$ IV $\Sigma$ 1 bound to  $\alpha$ II spectrin much more strongly than  $\beta$ IV $\Sigma$ 6 and neither interacted with GFP alone. Endogenous  $\beta$ II spectrin was also detected in COS-7 cells and could be coimmunoprecipitated with  $\alpha$ II-GFP (Fig. 3c).  $\alpha$ II spectrin’s ability to bind both  $\beta$ IV $\Sigma$ 1 and  $\beta$ IV $\Sigma$ 6 is surprising because  $\alpha$  and  $\beta$  spectrins are thought to interact through  $\beta$  spectrin’s first two SRs (Speicher et al., 1992). To determine how  $\alpha$ II spectrin and  $\beta$ IV spectrin splice variants interact, we constructed different truncated myc-tagged  $\beta$ IV spectrin proteins consisting of SR1, SR1-2, SR1-3, SR10-11, SR10-13, and SR14-15 and cotransfected these with  $\alpha$ II-GFP (Fig. 3d,e). Consistent with previous models for  $\alpha$  and  $\beta$  spectrin interactions (Speicher et al., 1992), we found that SR1-2 strongly bound  $\alpha$ II spectrin (Fig. 3d). However, these experiments also revealed a second, lower-affinity binding site (SR14-15) found in both  $\beta$ IV $\Sigma$ 1 and  $\beta$ IV $\Sigma$ 6 splice variants (Fig. 3e). Intriguingly, SR14-15 is also the binding site for ankG and is required for  $\beta$ IV spectrin recruitment to AIS and nodes (Yang et al., 2007). Therefore,  $\beta$ IV $\Sigma$ 1 binds strongly to  $\alpha$ II spectrin through two distinct SRs and  $\beta$ IV $\Sigma$ 6, despite lacking the N-terminal SR1-2, can also interact with  $\alpha$ II spectrin through SR14-15.

**Loss of  $\alpha$ II spectrin causes widespread CNS abnormalities**

Constitutive knock-out of  $\alpha$ II spectrin causes embryonic lethality in mice (Stanke-wich et al., 2011) and larval lethality in zebrafish (Voas et al., 2007). To determine the function of  $\alpha$ II spectrin in the postnatal brain, we generated  $\alpha$ II spectrin floxed mice (*Sptan1*<sup>flf</sup>) and crossed them with *Nestin-cre* mice to eliminate  $\alpha$ II spectrin

selectively in the CNS. We introduced *LoxP* sites flanking exon 8 of *Sptan1*. Upon recombination by Cre recombinase, exon 8 is excised, resulting in a frameshift mutation and premature stop codon near the end of SR3. We confirmed loss of  $\alpha$ II spectrin protein by immunostaining of cerebellar sections (Fig. 4*a*);  $\alpha$ II spectrin was detected only in endothelial cells in *Nestin-cre;Sptan1<sup>fl/fl</sup>* brains. To further confirm loss of  $\alpha$ II spectrin protein and to examine the consequence for  $\beta$  spectrins, we performed immunoblots on brain homogenates (Fig. 4*b*). As expected,  $\alpha$ II spectrin protein was nearly undetectable. Furthermore, we did not detect any  $\alpha$ I spectrin in either control or *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice, consistent with the conclusion that there is no compensation by  $\alpha$ I spectrin and  $\alpha$ II spectrin is the main  $\alpha$  spectrin detected in the brain (Zhang et al., 2014). Importantly, we measured a profound reduction in the amount of all  $\beta$  spectrins (with the exception of  $\beta$ IV $\Sigma$ 6, which was increased 4.5-fold; Fig. 4*b*). *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice were smaller than their littermate controls and loss of  $\alpha$ II spectrin resulted in juvenile lethality (Fig. 4*c*). *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice also had smaller brains with enlarged ventricles (data not shown). Therefore, loss of  $\alpha$ II spectrin disrupts spectrin cytoskeletons throughout the CNS, making the *Sptan1<sup>fl/fl</sup>* mice a powerful tool with which to study the importance of submembranous spectrin cytoskeletons in nervous system function.

Because  $\alpha$ II spectrin forms a complex with  $\beta$ IV spectrin at the AIS, we examined AIS in the cortex of *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice and found that, compared with controls, there were far fewer AIS and those present were fragmented and disrupted (Fig. 5*a*). We measured a reduction in the number of  $\beta$ IV spectrin-labeled AIS/1000  $\mu$ m<sup>2</sup> from  $30.2 \pm 2.8 \mu$ m in *Sptan1<sup>fl/fl</sup>* cortex to  $2.85 \pm 0.28$  in *Nestin-cre;Sptan1<sup>fl/fl</sup>* cortex ( $n = 3$  for each genotype,  $p = 0.0006$ ,  $t = 9.916$ ,  $df = 4$ ). The reduction in  $\beta$ IV spectrin-labeled AIS occurred despite the increased expression of the  $\beta$ IV $\Sigma$ 6 splice variant (Fig. 4*b*). This is consistent with the fact that  $\beta$ IV $\Sigma$ 6 alone is not sufficient to rescue AIS properties (Lacas-Gervais et al., 2004). To examine the consequence of loss of  $\alpha$ II spectrin on the organization of the AIS and axonal cytoskeletons, we performed STORM imaging on cultured *Sptan1<sup>fl/fl</sup>* hippocampal neurons transduced with AAV-Cre-GFP. STORM imaging showed that, whereas control (GFP<sup>-</sup>) neurons had intact  $\beta$ IV spectrin staining and periodicity at the AIS (Fig. 5*b–e*), neurons lacking  $\alpha$ II spectrin (GFP<sup>+</sup>) had significantly less clustered  $\beta$ IV spectrin and the periodicity was disrupted dramatically (Fig. 5*f–i*).  $\alpha$ II spectrin interacts with  $\beta$ II spectrin in the distal axon (Galiano et al., 2012) and  $\beta$ II spectrin forms a periodic cytoskeleton there (Fig. 5*j–m*) (Xu et al., 2013).

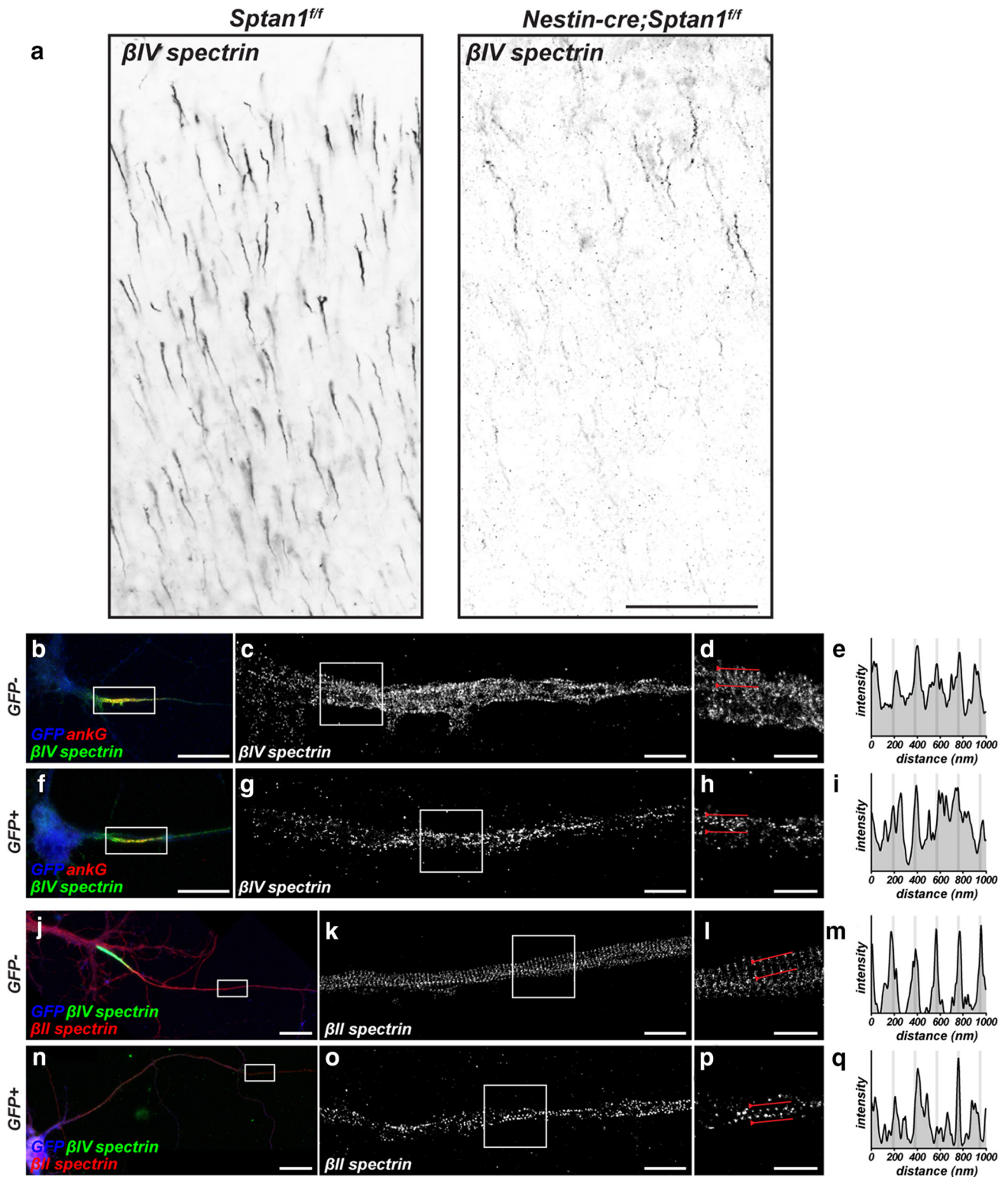


**Figure 4.** Mice lacking nervous system  $\alpha$ II spectrin. *a*,  $\alpha$ II spectrin immunostaining in cerebellum of P12 *Sptan1<sup>fl/fl</sup>* and *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice. *b*, Immunoblots of brain membrane homogenates from P7 *Sptan1<sup>fl/fl</sup>* and *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice using antibodies against the indicated spectrins. Blood was used as the positive control for  $\alpha$ I spectrin. Neurofilament-M (NF-M) served as the loading control. Protein levels were calculated as the percentage of *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice compared with *Sptan1<sup>fl/fl</sup>* mice.  $n = 3$  mice per genotype. Mean  $\pm$  SEM  $\alpha$ II:  $p = 0.0009$ ;  $\beta$ I:  $p = 0.0234$ ;  $\beta$ II:  $p = 0.0042$ ;  $\beta$ III:  $p = 0.032$ ;  $\beta$ IV- $\Sigma$ 1:  $p = 0.0112$ ;  $\beta$ IV- $\Sigma$ 6:  $p = 0.0174$ .  $\alpha$ II:  $t_{(4)} = 21.47$ ;  $\beta$ I:  $t_{(4)} = 3.57$ ;  $\beta$ II:  $t_{(4)} = 5.89$ ;  $\beta$ III:  $t_{(4)} = 3.228$ ;  $\beta$ IV- $\Sigma$ 1:  $t_{(4)} = 4.46$ ;  $\beta$ IV- $\Sigma$ 6:  $t_{(4)} = 3.912$ . *c*, Survival curve for *Sptan1<sup>fl/fl</sup>* ( $n = 26$ ) and *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice ( $n = 18$ ). \* $p < 0.05$ ; \*\* $p < 0.01$ ; \*\*\* $p < 0.001$ .

Similar to the AIS, loss of  $\alpha$ II spectrin disrupted the periodicity of the  $\beta$ II spectrin-containing distal axon cytoskeleton (Fig. 5*n–q*). Therefore, loss of  $\alpha$ II spectrin affects multiple  $\beta$  spectrins and has widespread effects on the organization of the axonal cytoskeleton.

Human mutations in *SPTAN1* cause severe epilepsy (Saitou et al., 2010). The *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice recapitulated this phenotype with generalized seizures lasting from 3 to 20 s and involving nearly continuous fine limb movements. EEG recordings (Fig. 6*a*) revealed spike frequencies of  $0.25 \pm 0.1$  ( $n = 4$ ,  $\pm$ SEM) and  $40.3 \pm 17.0$  ( $n = 5$ ,  $\pm$ SEM) spikes/h in control and *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice, respectively ( $p = 0.0159$ , Fig. 6*b*); control mice displayed no abnormal EEG discharges.





**Figure 5.** Loss of  $\alpha$ II spectrin disrupts the AIS cytoskeleton. **a**, AIS immunostained for  $\beta$ IV spectrin in P18 *Sptan1<sup>f/f</sup>* and *Nestin-cre;Sptan1<sup>f/f</sup>* cerebral cortex. Scale bar, 20  $\mu$ m. **b–e**, Conventional fluorescence (**b**) and STORM (**c, d**) imaging of GFP<sup>-</sup> 12 DIV cultured hippocampal neurons labeled with antibodies against GFP (blue), ankG (red), and  $\beta$ IV spectrin (green). The box in **b** corresponds to the STORM image shown in **c**. The box in **c** corresponds to the STORM image shown in **d**. The region between the lines in **d** was used to generate a  $\beta$ IV spectrin intensity profile (**e**). Scale bars: **b**, 10  $\mu$ m; **c**, 2  $\mu$ m; **d**, 1  $\mu$ m. **f–i**, Conventional fluorescence (**f**) and STORM (**g, h**) imaging of GFP<sup>+</sup> 12 DIV cultured hippocampal neurons labeled with antibodies against GFP (blue), ankG (red), and  $\beta$ IV spectrin (green). The box in **f** corresponds to the STORM image shown in **g**. The box in **g** corresponds to the STORM image shown in **h**. The region between the lines in **h** was used to generate a  $\beta$ IV spectrin intensity profile (**i**). Scale bars: **f**, 10  $\mu$ m; **g**, 2  $\mu$ m; **h**, 1  $\mu$ m. **j–m**, Conventional fluorescence (**j**) and STORM (**k, l**) imaging of GFP<sup>-</sup> 12 DIV cultured hippocampal neurons labeled with antibodies against GFP (blue),  $\beta$ II spectrin (red), and  $\beta$ IV spectrin (green). The box in **j** corresponds to the STORM image shown in **k**. The box in **k** corresponds to the STORM image shown in **l**. The region between the lines in **l** was used to generate a  $\beta$ II spectrin intensity profile (**m**). Scale bars: **j**, 10  $\mu$ m; **k**, 2  $\mu$ m; **l**, 1  $\mu$ m. **n–q**, Conventional fluorescence (**n**) and STORM (**o, p**) imaging of GFP<sup>+</sup> 12 DIV cultured hippocampal neurons labeled with antibodies against GFP (blue),  $\beta$ II spectrin (red), and  $\beta$ IV spectrin (green). The box in **n** corresponds to the STORM image shown in **o**. The box in **o** corresponds to the STORM image shown in **p**. The region between the lines in **p** was used to generate a  $\beta$ II spectrin intensity profile (**q**). Scale bars: **n**, 10  $\mu$ m; **o**, 2  $\mu$ m; **p**, 1  $\mu$ m.

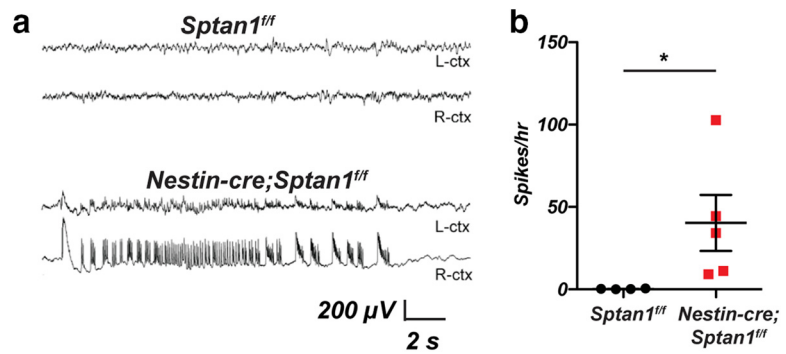


In addition to intrinsic axonal excitability changes arising from AIS disruption, epilepsy may also result from cortical malformations and aberrant neuronal layering (Guerrini and Dobyns, 2014). Immunostaining of *Nestin-cre;Sptan1<sup>fl/fl</sup>* cortex with antibodies against neurons normally restricted to distinct cortical layers revealed disrupted cortical lamination. We found that, at both P7 and P12, Reelin<sup>+</sup>, Ctip2<sup>+</sup>, Foxp2<sup>+</sup>, and ctgf<sup>+</sup> neurons corresponding to distinct cortical layers were frequently dispersed among other layers (Fig. 7*a,b*). For example, at P12, many Ctip2-labeled neurons (layer V) were found in layers II–IV (Cux1<sup>+</sup> region), whereas Reelin<sup>+</sup> neurons were distributed more broadly throughout the cortex at both P7 and P12 (Fig. 7*c,d*).

To gain insight into the causes of disrupted cortical lamination, we performed *in utero* electroporation to label migrating neurons and to silence expression of  $\alpha$ II spectrin using a highly efficient short hairpin RNA (shRNA) (Galiano et al., 2012). This allowed us to examine the cell-autonomous consequence of loss of  $\alpha$ II spectrin during development in single neurons. We found that, in contrast to electroporation with GFP alone (Fig. 7*e*), at P7, electroporated neurons lacking  $\alpha$ II spectrin were frequently arrested in their migration through the cortical plate.  $\alpha$ II spectrin-deficient neurons also lacked dendrites and remained in a migrating bipolar morphology (Fig. 7*f*). To examine more directly the effect of loss of  $\alpha$ II spectrin on dendrite morphology, we transduced cultured rat hippocampal neurons at 1 DIV using adenovirus to express a control shRNA or  $\alpha$ II spectrin shRNA (Galiano et al., 2012). At 17 DIV, we measured the extent of dendritic arborization and found a reduced complexity ( $n = 3$  independent experiments for each condition; Fig. 7*g*). Therefore, loss of  $\alpha$ II spectrin causes severe epilepsy that may reflect disrupted AIS, reduced dendritic complexity, and abnormal development of neural circuits due to impaired neuronal migration.

Consistent with widespread abnormalities in brain organization, when we examined the cerebellum, we found significantly fewer Purkinje neurons ( $53.3 \pm 1.3$  cells/mm and  $29.3 \pm 0.8$  cells/mm in WT and *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice, respectively;  $p = 0.0001$ ,  $t_{(4)} = 2.5$ ) and the *Nestin-cre;Sptan1<sup>fl/fl</sup>* Purkinje neuron AIS were fragmented and much shorter than those in control *Sptan1<sup>fl/fl</sup>* mice (Fig. 7*h,i*). Cerebellar pinneaux, labeled for Kv1.2 and corresponding to basket cell terminals, were also disrupted in  $\alpha$ II spectrin-deficient cerebellum (Fig. 7*h*). Furthermore, control AIS were usually oriented  $\pm 30^\circ$  perpendicular to the plane of the Purkinje neuron layer, but *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice had AIS that deviated much further from the perpendicular axis (Fig. 7*h,i*), indicating that altered axonal projections may also contribute to circuit dysfunction in  $\alpha$ II spectrin-deficient mice. Finally, the molecular layer at P18 was also significantly thinner in *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice ( $97.3 \pm 4.8 \mu\text{m}$  in *Sptan1<sup>fl/fl</sup>* mice and  $70.0 \pm 3.3 \mu\text{m}$  in *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice;  $n = 3$  for each genotype,  $p = 0.006$ ,  $t = 5.15$ ,  $df = 4$ ).

What accounts for the reduced numbers of Purkinje neurons, thinner molecular layer, and disrupted brain organization? Because neuronal cytoskeletons are thought to play important roles in neurons, one possibility is that  $\alpha$ II spectrin-deficient neurons may degenerate. To test this, we immunostained brain sections using antibodies against beta-amyloid precursor protein ( $\beta$ -APP),



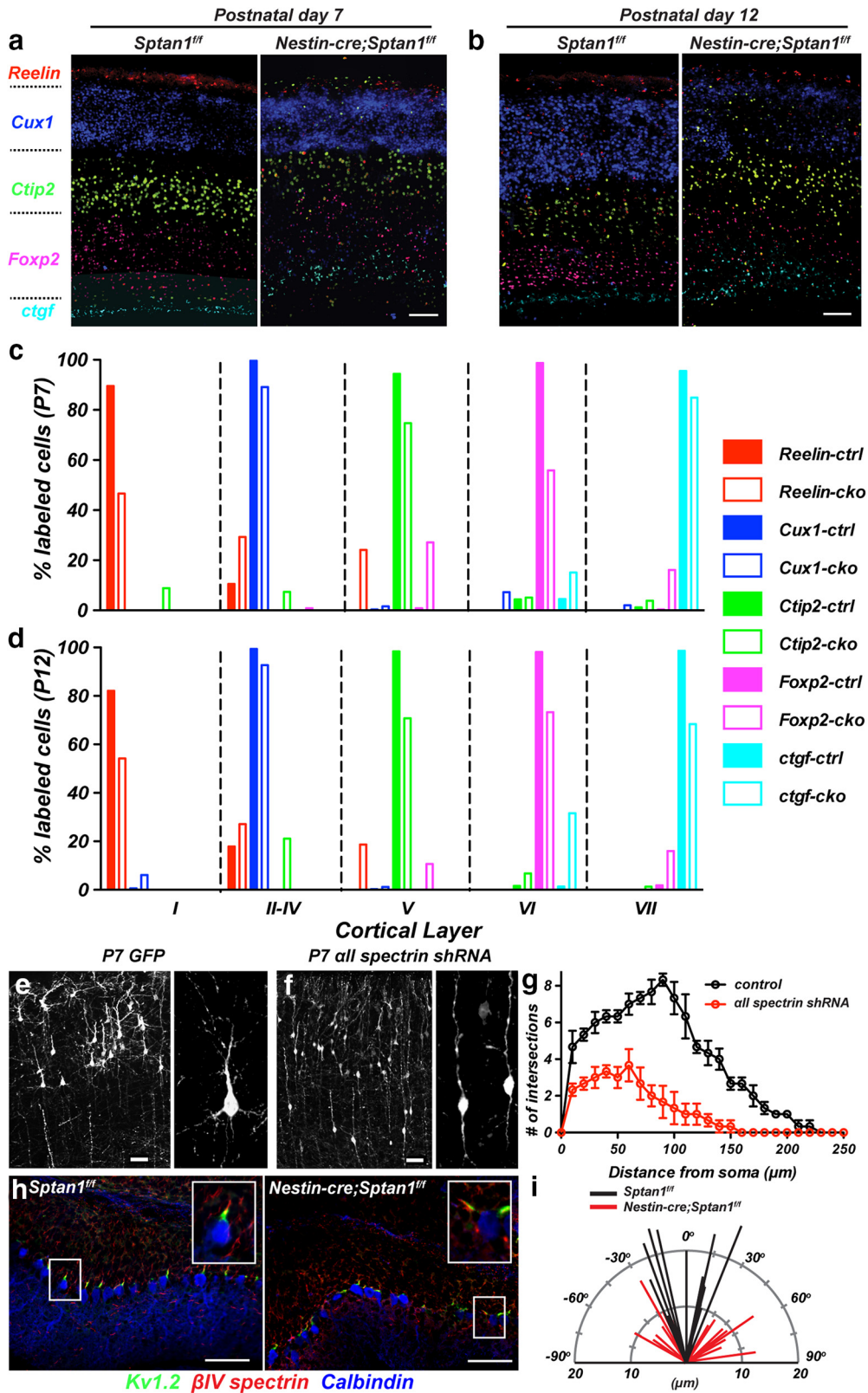
**Figure 6.**  $\alpha$ II spectrin-deficient mice have seizures. *a*, Video EEG monitoring of awake and behaving P12–P14 mice revealed generalized seizure discharges in *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice that were not detected in *Sptan1<sup>fl/fl</sup>* littermates. *b*, Quantification of spikes/h recorded in *Sptan1<sup>fl/fl</sup>* and *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice. \* $p = 0.0159$ .

a robust marker of axonal injury and degeneration (Gentleman et al., 1993). We found widespread immunostaining for  $\beta$ -APP in *Nestin-cre;Sptan1<sup>fl/fl</sup>* Purkinje neurons in the cerebellum and throughout the thalamus and cortex (Fig. 8*a–c*). These results show that loss of  $\alpha$ II spectrin leads to widespread CNS neurodegeneration.

## Discussion

Submembranous spectrin cytoskeletons are found in all eukaryotic cells and are thought to confer both flexibility and strength to cell membranes and to participate in organizing and maintaining the distribution of membrane proteins. These conclusions are based on detailed studies of erythrocyte spectrins and human  $\alpha$ I spectrin mutations that cause spherocytic anemia due to a fragile cell membrane (Perrotta et al., 2008). However, much less is known about spectrins in the nervous system. Although human mutations in  $\alpha$ II spectrin cause severe epilepsy (Saitsu et al., 2010; Tohyama et al., 2015), previous studies of spectrin function have mostly focused on the role of  $\beta$  spectrins. For example, human mutations in  $\beta$ III spectrin cause spinocerebellar ataxia type 5 (Ikeda et al., 2006) and mouse and human mutations in  $\beta$ IV spectrin cause deafness, neuropathy, and ataxia (Parkinson et al., 2001; Komada and Soriano, 2002; Yang et al., 2004; Knierim et al., 2017). Determining the function of the spectrin cytoskeleton in vertebrates has been complicated by the diversity of  $\beta$  spectrins and by the fact  $\beta$  spectrins may compensate for one another (Ho et al., 2014). The function of  $\alpha$ II spectrin has been studied in mice and zebrafish, but animals lacking  $\alpha$ II spectrin die at the embryonic and larval stages, respectively (Voas et al., 2007; Stankewich et al., 2011), precluding any studies of physiology or behavior or determination of the role of  $\alpha$ II spectrin in maintenance of neuronal integrity and function.

Here, we confirmed that  $\alpha$ II spectrin is the major  $\alpha$  spectrin found in the brain and is required for proper expression of all other  $\beta$  spectrins. Therefore, our  $\alpha$ II spectrin conditional knock-out mouse represents an attractive model to study, not only West syndrome, a severe form of infantile spasms for which only one other nonlethal genetic mouse model has been reported (Olivetti et al., 2014), but more broadly the function of the neuronal spectrin-based submembranous cytoskeleton. We demonstrate that  $\alpha$ II spectrin plays diverse roles in the nervous system including, but not limited to, AIS assembly, neuronal migration, excitability, and health. These pleiotropic functions of  $\alpha$ II spectrin likely reflect different roles depending on the cell type and which of the neuronal  $\beta$  spectrins it is partnered with. Future experiments will be required to further define the contributions of



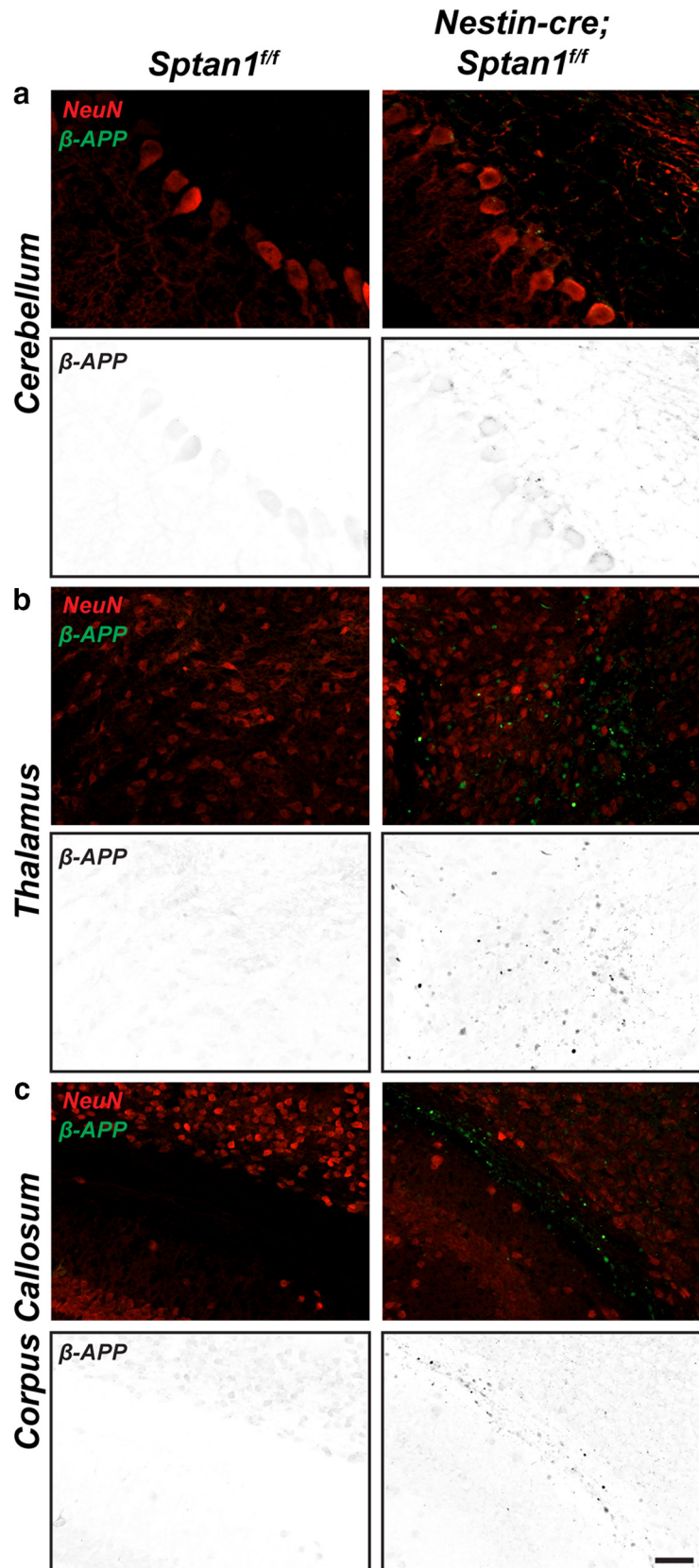
**Figure 7.**  $\alpha$ II spectrin-deficient mice have disrupted cortical lamination and cerebellar organization. **a, b**, Immunostaining of cerebral cortex using antibodies against Reelin (layer I), Cux1 (layers II–IV), Ctip2 (layer V), FoxP2 (layer VI), and ctgf (subplate) at P7 (**a**) and P12 (**b**). Scale bar, 50  $\mu$ m. **c, d**, Quantification of percentage of labeled cells (Reelin, red; Cux1, blue; Ctip2, green; Foxp2, magenta; ctgf, cyan) in their respective cortical layers (I, II–IV, V, VI, VII) in P7 (**c**) or P12 (**d**) control (close column) or  $\alpha$ II spectrin-deficient (open column) cerebral cortex. **e, f**, GFP immunoreactivity in P7 rats after *in utero* electroporation using GFP alone (**e**) or GFP and  $\alpha$ II spectrin shRNA (**f**) plasmids. Scale bar, 50  $\mu$ m. **g**, Sholl analysis of dendritic complexity in control or  $\alpha$ II spectrin-deficient hippocampal neurons *in vitro* ( $n = 3$ ). Error bars indicate  $\pm$  SEM. **h**, Immunostaining of P18 *Sptan1<sup>fl/fl</sup>* and *Nestin-cre;Sptan1<sup>fl/fl</sup>* cerebella using antibodies against Kv1.2 (green),  $\beta$ IV spectrin (red), and calbindin (blue). Scale bar, 50  $\mu$ m. **i**, Orientation and length of Purkinje neuron AIS in *Sptan1<sup>fl/fl</sup>* and *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice.

specific  $\alpha$ II/ $\beta$  spectrin combinations to nervous system function in distinct cell types (e.g., neurons and glia) and may be complicated by the ability of  $\beta$  spectrins to substitute for one another (Ho et al., 2014).

#### $\alpha$ II spectrin forms a periodic cytoskeleton with $\beta$ IV spectrin at the AIS and nodes of Ranvier

Among the  $\beta$  spectrins, one of the most extensively studied in the vertebrate nervous system is  $\beta$ IV spectrin because it is highly enriched at AIS and nodes of Ranvier (Berghs et al., 2000; Komada and Soriano, 2002). Previous studies in mice demonstrated the ability of  $\alpha$ II spectrin to coimmunoprecipitate both  $\beta$ IV $\Sigma$ 1 and  $\beta$ IV $\Sigma$ 6 (Uemoto et al., 2007) and experiments in zebrafish larvae revealed a transient localization of  $\alpha$ II spectrin to nodes of Ranvier (Voas et al., 2007). However,  $\beta$ IV spectrin's function at AIS and nodes remained enigmatic for the following reasons: (1)  $\alpha$ II spectrin has not been shown at AIS or nodes of mature axons, (2) the major  $\beta$ IV spectrin splice variant found in mature AIS and nodes ( $\beta$ IV $\Sigma$ 6; Yoshimura et al., 2016) lacks the canonical actin and  $\alpha$ II spectrin-binding domains (SR1-2), and (3) loss of  $\beta$ IV spectrin can be partially compensated for by  $\beta$ I spectrin (Ho et al., 2014). Our results help to resolve these problems. First, we found that detergent extraction of unfixed, cultured hippocampal neurons reveals a detergent insoluble pool of  $\alpha$ II spectrin restricted to the AIS. Superresolution microscopy shows that AIS  $\alpha$ II spectrin exists in a periodic distribution with a spacing of  $\sim$ 190 nm, corresponding to the expected length of a spectrin tetramer (Shotton et al., 1979). Second, we showed that  $\alpha$ II spectrin can bind to both  $\beta$ IV $\Sigma$ 1 and  $\beta$ IV $\Sigma$ 6, albeit with different affinities. Remarkably, in addition to SR1-2, we found that SR14-15, the same domain in  $\beta$ IV spectrin that binds to ankG (Yang et al., 2007), can also bind to  $\alpha$ II spectrin. It will be interesting to determine whether the addition of ankG results in cooperative or competitive binding. Finally, loss of  $\alpha$ II spectrin disrupts AIS and nodes (Huang et al., 2017), demonstrating the requirement for the spectrin cytoskeleton in their assembly and function.

Why does loss of  $\alpha$ II spectrin cause seizures in the *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice? It is possible that disrupted AIS in *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice causes seizures because the AIS is the main determinant of the output of a neuron and is responsible for the initiation of axonal action potentials (Kole et



**Figure 8.** *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice have widespread axon degeneration. **a–c**, Immunostaining of cerebellum (**a**), thalamus (**b**), and corpus callosum (**c**) in *Sptan1<sup>fl/fl</sup>* and *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice using antibodies against  $\beta$ -APP (green) and NeuN (red). Scale bar, 50  $\mu$ m.



al., 2008). However, because both excitatory and inhibitory neurons have AIS, our experiments cannot distinguish between the contributions to seizures from excitatory or inhibitory neurons. In addition, the severely disrupted cortical lamination observed in the *Nestin-cre;Sptan1<sup>fl/fl</sup>* mice may also contribute to seizures. We found that loss of  $\alpha$ II spectrin impaired the migration of cortical neurons to their proper locations. We speculate that, because  $\alpha$ II spectrin plays important roles in regulating the surface expression of membrane proteins and cell adhesion molecules (Susuki et al., 2011), migration may be arrested due to an impaired response to guidance cues. This view is consistent with the observation that the axons of *Nestin-cre;Sptan1<sup>fl/fl</sup>* Purkinje neurons have significantly altered trajectories. This may reflect an impaired ability of axons to respond to guidance cues because of a loss or reduction in their receptors.

In addition to their roles in maintaining ion channel clusters at AIS, neuronal spectrins have been proposed to help axons withstand the mechanical forces to which they are exposed (Xu et al., 2013). Although the experiments described here do not test this idea directly, the observation that loss of the spectrin cytoskeleton results in widespread axon degeneration is consistent with this model. More direct evidence will require experiments that uncouple the contribution of AIS and dendritic  $\alpha$ II spectrin from their role in long axons.

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