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Ⅲ. 研究成果の刊行物・別刷



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Research Report

Abnormal maturation and differentiation of neocortical neurons in epileptogenic cortical malformation: Unique distribution of layer-specific marker cells of focal cortical dysplasia and hemimegalencephaly

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ABSTRACT

Focal cortical dysplasia (FCD) and hemimegalencephaly (HME) are major causes of intractable epilepsy in children. The probable pathogenesis of FCD and HMG is the abnormal migration and differentiation of neurons. The aim of the present study was to clarify the abnormal cytoarchitecture, based on neuronal immaturation. Tissue samples were obtained from 16 FCD and seven HME patients, aged between 2 months and 12 years, who had been diagnosed as typical FCD and HME, following surgical treatment for intractable epilepsy. Paraffin-embedded sections were stained with the antibodies of three layer-markers that are usually present only during the fetal period, namely SATB2 (expressed in the upper layer of the normal fetal neocortex), FOXP1 (expressed in the 5th layer), and TBR1 (expressed in the 6th layer). In FCD, SATB2-positive (+) cells located in the middle and deep regions of FCD Ia and Ib, but only in the superficial region of FCD IIa and IIb. FOXP1+ cells diffusely located in the neocortex, especially the upper layer of FCD IIaand IIb. TBR1+ cells mainly located in the middle and deep regions, and also white matter. In FCD IIb, TBR1+ cells were in the superficial region. In HME, SATB2+ and FOXP1+ cells were found diffusely. TBR1+ cells were in the middle and deep regions. On the basis of continued expression of fetal cortical layer-specific markers in FCD and HME brains, the

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abnormal neocortical formation in both is likely to be the result of disrupted neuronal migration and dysmaturation. The expression pattern is different between FCD and HME.

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1. Introduction

Focal cortical dysplasia (FCD) and hemimegalencephaly (HME), rare epileptogenic brain malformations are usually accompanied by severe epilepsy and occasionally by mental retardation. The incidence of FCD epilepsy identified in surgical series varies between 12% and 40% (Blümcke et al., 1999; Nordborg et al., 1999; Prayson et al., 2002), while that of HME is lower. These diseases are of relatively high frequency in surgical epilepsy, but have a low morbidity rate (Prayson and Estes, 1995; Prayson, 2000). FCD and HME are usually diagnosed by neuropathological findings in specimens undergoing cortical resection for the treatment of refractory epilepsy. As common features, mental development problems sometimes remain, in spite of well-controlled epilepsy.

FCD was recognized as a pathologic substrate associated with epilepsy (Taylor et al., 1971). It is known that FCD has columnar and laminar disorganization with various cellular abnormalities, including dysmorphic neurons, giant neurons, and balloon cells (Prayson et al., 1996; Yamanouchi et al., 1996; Palmini et al., 2004; Alonso-Nanclares et al., 2005; Blümcke et al., 2011). On the other hand, HME mainly shows cortical laminar abnormality, such as polymicrogyria and neuronal heterotopia. Together with these pathological findings, it is thought that FCD and HME may result from erroneous migration, maturation, or cell death during ontogenesis (Crino and Eberwine, 1997; Cotter et al., 1999; Najm et al., 2007). However, a common pathogenesis remains unknown.

On the other hand, some molecules are useful to detect layer formation of human neocortex. We recently demonstrated that human malformed brains have unique layer patterns (Saito et al., 2010). In the present study, we seek to detect abnormal neuronal migration and differentiation in FCD and HME that will lead to greater understanding of the pathophysiology of the epileptogenic malformed brain.

2. Results

2.1. Histological distribution

Histopathological results were summarized in Table 1. We obtained 4 FCD Ia, 4 FCD Ib, 4 FCD IIa and 4 FCD IIb from the international classification (Blümcke et al., 2011), and seven HME. FCD Ic was relatively rare. Although we could not examine this subtype, it was enough to investigate FCD Ia and Ib for the aim of the present study because FCD Ic pathologically showed the combination of FCD Ia and Ib. All HME cases showed polymicrogyria and/or unlayered neocortex with neuronal heterotopia and mineralization.

2.2. Immunohistochemistry of FCD

Immunohistochemistry results were summarized in Table 2. SATB2+ cells were dominant in the middle and deep regions

of the neocortex in FCD Ia and Ib, although they were widely distributed (Figs. 1 and 2, Table 2). Interestingly, SATB2+ cell distribution of FCD IIa and IIb was limited to the superficial region of the neocortex (Figs. 3 and 4, Table 2). FOXP1 immunoreactivity was diffusely intense (Table 2). Only FCD Ib revealed FOXP1+ cells in the white matter. FCD IIa and IIb demonstrated no FOXP1+ cells in the white matter, and FOXP1+ cells tended to appear in the superficial region of the neocortex. TBR1+ cells were the most prominent in the middle and deep regions of the neocortex (Fig. 4). Notably, TBR1+ cells exhibited a unique localization of the superficial region of the neocortex in FCD IIb (Fig. 4, Table 2). The immunopositive cells for SATB2+, FOXP1 and TBR1 were confirmed as neurons with NeuN-immunopositivity (data not shown). There were no significant differences in the marker expression patterns in each lobe.

2.3. Immunohistochemistry of HME

SATB2+ cells in HME were diffused in the neocortex, but relatively dense in the superficial region of the neocortex (Fig. 5). FOXP1 immunoreactivity also diffusely distributed, but was occasionally negative (Fig. 5, Table 2). TBR1+ cells were limited to the middle and deep regions of the neocortex (Fig. 5, Table 2). The immunopositive cells for SATB2+, FOXP1, and TBR1 in HME were also confirmed as neurons with NeuNimmunopositivity (data not shown). There were also no significant differences in the marker expression patterns in each lobe.

3. Discussion

In the normal developing cortex, the localization of SATB2, FOXP1, and TBR1 is restricted to specific cortical layers, and the expression of all three markers disappears in the postnatal brain (Saito et al., 2011). In the present series, SATB2, FOXP1, and TBR1 were diffusely expressed throughout the cortex in samples from all cases. The result indicates that FCD and HME consist of immature cells. Moreover, we identified that these layer-marker immunopositive cells were neurons by a neuron marker, NeuN. There was an observable tendency for SATB2+cells to be distributed in the middle and deep regions of the neocortex of FCD Ia and Ib, and limited to the superficial region of FCD IIa and IIb. It is quite interesting that SATB2+ and FOXP1+ cells in HME were diffusely distributed and TBR1+ cells were localized in the middle and deep regions of the neocortex. To evaluate the expression patterns of those specific markers in FCD subtypes or HME, we divided them into three regions of the neocortex in terms of thickness.

Although there is little evidence regarding the mechanisms responsible for human FCD, it has been reported that FCD neurons originate from abnormal migration, maturation, and

| Case S | Sex | Age at | Age at seizure | Seizure | Intelligence | FCD location on | Pathological findings | | |
|-----------|-----|---------|----------------|---------|--------------|-----------------|-----------------------|-------------------|--|
| | | surgery | onset | | | imaging | Main pathology | Others | |
| FCD | | | | | | | | | |
| 1 | M | 2 Y | 2 m | CPS+GTC | 100 (IQ) | P | FCD Ia | Mild gliosis | |
| 2 | F | 3 Y | 3 m | CPS | 50 (DQ) | F | FCD Ia | HN, gliosis | |
| 3 | М | 6 Y | 4 m | CPS | 33 (DQ) | F | FCD Ia | Mild gliosis | |
| 4 | М | 7 Y | 11 m | CPS | 58 (IQ) | F | FCD Ia | Mild gliosis | |
| 5 | М | 6 M | 1 m | CPS | 40 (DQ) | P | FCD Ib | Mild gliosis | |
| 6 | M | 2 Y | 20 d | CPS | 18 (DQ) | F | FCD Ib | HN, gliosis | |
| 7 | F | 3 Y | 3 m | CPS | 15 (DQ) | P | FCD Ib | HN, gliosi | |
| 8 | F | 12 Y | 11 m | CPS | 43 (IQ) | F | FCD Ib | Mild gliosis | |
| 9 | F | 3 Y | 6 m | CPS+GTC | 40 (DQ) | F+P | FCD IIa | HN, gliosi | |
| 10 | F | 5 Y | 2 y 9 m | CPS | 81 (IQ) | F | FCD IIa | HN, gliosi | |
| 11 | M | 6 Y | 3 d | CPS | 16 (DQ) | T+P+O | FCD IIa | HN, gliosi | |
| 12 | M | 7 Y | 7 m | CPS | 50 (IQ) | P P | FCD IIa | HN, gliosi | |
| 13 | M | 10 Y | 1 y 11 m | CPS+GTC | 22 (IQ) | F+T+P | FCD IIb | HN, gliosi | |
| 14 | M | 3 Y | 2y 9 m | GTC | 15 (DQ) | T+P+O | FCD IIb | HN, gliosi | |
| 15 | M | 8 Y | 3 m | CPS+GTC | 30 (IQ) | F+P | FCD IIb | HN, gliosi | |
| 16 HME | F | 19 Y | 9 m | CPS+GTC | 25 (IQ) | P | FCD IIb | HN, gliosi | |
| 1 | F | 3 M | 2 d | CPS+GTC | 30 (DQ) | rt-hemisphere | Polymicrogyria | HN, M, gliosis | |
| 2 | F | 3 M | 7 d | CPS+GTC | 35 (DQ) | rt-hemisphere | DN, BC | HN, M, gliosis | |
| 3 | М | 3 M | 14 d | CPS+GTC | 30 (DQ) | lt-hemisphere | Polymicrogyria | HN, M, gliosis | |
| 4 | M | 6 M | 7 d | CPS+GTC | 50 (DQ) | lt-hemisphere | DN | HN, M, gliosis | |
| 5 | M | 3 M | 14 d | EIEE | 50 (DQ) | rt-hemisphere | DN, BC | HN, M, gliosis | |
| 6 | F | 4 M | 1 d | EIEE | 30 (DQ) | rt-hemisphere | Polymicrogyria | HN, M, gliosis | |
| 7 | М | 7 M | 3 d | EIEE | 20 (DQ) | lt-hemisphere | Polymicrogyria | HN, M, gliosis | |

M: male, F: female, M (m): month (s), Y (y): year (s), d: days, CPS: complex partial seizure, GTC: generalized tonic-clonic convulsion, EIEE: early infantile epileptic encephalopathy, IQ: intelligence quotient, DQ: development quotient, F: frontal lobe, P: parietal lobe, T: temporal lobe, O: occipital lobe, rt: right side, lt: left side, DN: dysmorphic neuron, BC: balloon cell, HN: heterotopic neuron, M: mineralization.

cell death during ontogenesis (Spreafico et al., 1998a; 1998b; Andres et al., 2005). Our results may support this theory, indicating the persistence of immature neurons in the white matter. A recent report shows that markers of neuronal immaturity were overexpressed to excess in FCD (Hanai et al., 2010). Moreover, the previous studies have also reported that FCD neurons exhibited various degrees of neuronal maturation, glial cells or a combination of neuronal and glial characteristics (Crino and Eberwine, 1997; Yamanouchi et al., 1998; Aronica et al., 2003; Fauser et al., 2004; Ying et al., 1999). FCD and HME may retain certain characteristics indicative of immaturity.

SATB2 is a DNA-binding protein that regulates chromatin organization and gene expression. In the developing brain, SATB2 is expressed in cortical projection neurons. In a previous study, SATB2 has expressed predominantly in the upper layer, and not in the deep layer, of the cortex (Britanova et al., 2008). This expression pattern suggests that SATB2 may

be involved in the control of early aspects of upper layer neuron specification. Interestingly, the SATB2 expression pattern can be clearly divided into three types in the present study. The first pattern is SATB2 expression in the middle and deep regions of the neocortex and some in the white matter. The second is SATB2 expression in the superficial region of the neocortex. The third pattern is diffused SATB2 expression. The first was identified in FCD Ia and Ib, the second in FCD IIa and IIb, and the third in HME.

FOXP1+ cells are supposed to be projection neurons. FOXP1 is a member of a conserved family of genes that shares a common DNA-binding domain, namely the T-box (Tamura et al., 2004). The T-box genes encode transcription factors involved in the regulation of developmental processes. A similar protein that is highly expressed in the 4th and 5th layers has been reportedly disrupted in mice and shown to be critical for early cortical developmental processes (Takahashi

| Case | Pathological classification | SATB2 | | | | FOXP1 | | | | TBR1 | | | |
|----------|-----------------------------|-------------|----------|----------|--|-----------------|--------|------|--------------|----------------|--------|------|------|
| | | Superficial | Middle | Deep | WM | Superficial | Middle | Deep | WM | Superficial | Middle | Deep | WM |
| FCD | | | | | | | | | | | | | |
| 1 | Ia | + | ++- | ++ | + | + | ++ | _ | _ | + | # | ++ | ++ |
| 2 | Ia | _ | ++- | ++ | _ | + | + | _ | | | ++ | # | ++- |
| 3 | Ia | _ | ++ | ++ | + | 4.2 | ++- | # | _ | _ | ++ | ++ | ++- |
| 4 | Ia | <u>-</u> | ++ | ++ | + | | ++ | ++ | - | _ | ++ | ++ | |
| 5 | Ib | - | ++- | ++ | + | + | ++- | + | + | _ | + | + | + |
| 6 | Ib | _ | ++- | # | + | + | ++- | ++ | + | _ | + | + | + |
| 7 | Ib | ++ | ++ | ++ | # | _ | ++ | ++ | ++ | _ | ++ | ++ | ++- |
| 8 | Ib | +1- | +1- | + | _ | + | +1- | _ | + | _ | # | ++- | _ |
| 9 | IIa | + | _ | <u> </u> | | ++ | + | | _ | | + | - | **** |
| 10 | IIa | ++ | _ | _ | - | + | + | + | _ | | + | _ | _ |
| 11 | IIa | + | + | _ | + | + | ++ | _ | _ | _ | + | _ | _ |
| 12 | IIa | ++ | | - | | ++ | + | + | _ | _ | + | _ | _ |
| 13 | IIb | ++ | ++ | _ | - | ++ | ++- | + | _ | + | + | ++ | + |
| 14 | IIb | + | - | - | 40 (40 (40 (40 (40 (40 (40 (40 (40 (40 (| ++ | + | + | - | + | + | ++ | + |
| 15 | IIb | + | <u> </u> | - | _ | ++ | + | _ | _ | + | + | ++ | + |
| 16 | IIb | + | - | | _ | ++ | _ | - | _ | + | + | ++ | + |
| HOME | | | | | | | | | | | | | |
| 1 | rt-ḥemisphere | ++ | + | + | + | + | + | + | + | + | ++ | + | _ |
| 2 | rt-hemisphere | ++ | + | + | + | + | + | + | + | + | ++ | + | + |
| 3 | lt-hemisphere | ++ | + | + | + | + | - | + | + | - | ++ | + | _ |
| 4 | lt-hemisphere | ++ | + | + | + | - | + | _ | _ | <u>-</u> | ++ | + | |
| 5 | rt-hemisphere | ++ | + | + | + | + | + | + | + | _ | ++ | + | - |
| 6 | rt-hemisphere | # | + | + | + | + | + | + | + | -11 | + | + | |
| 7 | lt-hemisphere | ++ | + | + | + | + | + | + | _ | + | ++ | + | _ |
| Control | | | | | | | | | | | | | |
| 23-29 GW | | + | -11- | _ | - | <u>-</u> 100000 | - " | ++ | - | ++ | ++ | + | _ |
| 1 M-8 Y | | <u>-</u> | _ | - 1 | - | | _ | | _ | _ | | _ | _ |

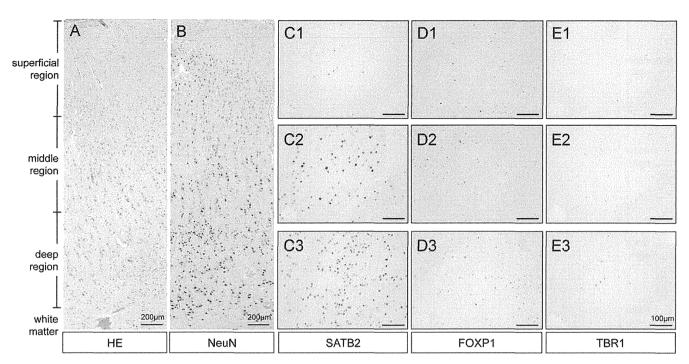


Fig. 1 – Histology and immunohistochemistry of FCD Ia. FCD Ia demonstrates abnormal radial lamination and abundant microcolumns of the neocortex (A, B). SATB2+ cells scattered in the superficial (C1), but at a high concentration in the middle (C2), and deep (C3) regions of the neocortex. FOXP1+ cells diffusely locate in the neocortex (D1-D3). TBR1+ cells diffusely locate in the neocortex (E1-E3) and those concentrations are low.

(A) Hematoxylin and eosin staining; (B) NeuN-immunostaining; (C) SATB2-immunostaining, superficial region (C1), middle region (C2) and deep region (C3); (D) FOXP1-immunostaining, superficial region (D1), middle region (D2), and deep region (D3); (E) TBR1-immunostaining, superficial region (E1), middle region (E2), and deep region (E3). Scales of A-B and C1-E3 indicate 200 µm and 100 µm, respectively.

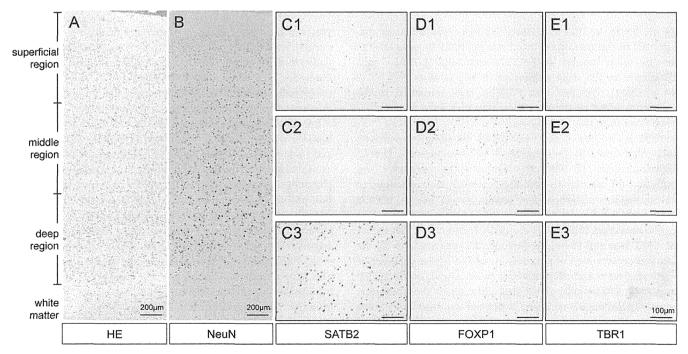


Fig. 2 – Histology and immunohistochemistry of FCD Ib. FCD Ib demonstrates abnormal tangential layer composition of the neocortex (A, B). SATB2+ cells scattered in the superficial (C1), but at a low concentration in the middle (C2) and a high concentration in the deep (C3) regions of the neocortex. FOXP1+ cells diffusely locate in the neocortex (D1-D3). TBR1+ cells diffusely locate in the neocortex, but those concentrations are low (E1-E3).

(A) Hematoxylin and eosin staining; (B) NeuN-immunostaining; (C) SATB2-immunostaining, superficial region (C1), middle region (C2), and deep region (C3); (D) FOXP1-immunostaining, superficial region (D1), middle region (D2), and deep region (D3); (E) TBR1-immunostaining, superficial region (E1), middle region (E2), and deep region (E3). Scales of A-B and C1-E3 indicate 200 μm and 100 μm, respectively.

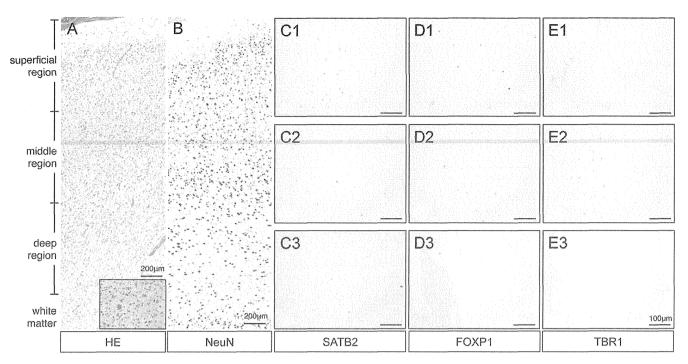


Fig. 3 – Histology and immunohistochemistry of FCD IIa. FCD IIa demonstrates unidentified layer-formation of the neocortex and a high neuronal concentration (A, B), and contains dysmorphic neurons (small window in A). SATB2+ cells scattered in the superficial (C1) and middle (C2) regions, but those concentrations are very low in the deep (C3) region of the neocortex. FOXP1+ cells diffusely locate in the neocortex (D1-D3), and evidence a relatively low concentration in the deep region (D3). TBR1+ cells diffusely locate in the neocortex, but those concentrations are low (E1-E3).

(A) Hematoxylin and eosin staining; (B) NeuN-immunostaining; (C) SATB2-immunostaining, superficial region (C1), middle

(A) Hematoxylin and eosin staining; (B) NeuN-immunostaining; (C) SATB2-immunostaining, superficial region (C1), middle region (D2), and deep region (D3); (E) TBR1-immunostaining, superficial region (E1), middle region (E2), and deep region (E3). Scales of A-B and C1-E3 indicate 200 μm and 100 μm, respectively.

et al., 2008). In FCD and HME, FOXP1+ cells diffusely distributed in the neocortex. Interestingly, FOXP1+ cells located in the white matter of FCD Ib and HME. FOXP1+ cells distribute in the deep neocortex in the fetal period (Saito et al., 2011). The fact that FOXP1+ cells remained in the postnatal white matter may indicate more delayed neuronal migration than the other types.

TBR1 is also one of the T-box genes encoding transcription factors involved in the regulation of development. The Cterminal region of TBR1 was found to be necessary and sufficient for association with the guanylate kinase domain of calcium/calmodulin-dependent serine protein kinase. TBR1 is highly expressed in early neurons of the preplate and deep layer of the neocortex (Bulfone et al., 1995). Furthermore, the cortex of TBR1 mutants shows developmental abnormalities in the laminar organization of neurons, as well as in the guidance of cortical afferent and efferent axons (Hevner et al., 2001). In FCD and HME, TBR1+ cells distributed diffusely in the neocortex, mainly in the middle and deep regions of the neocortex. These data may be supported by the previous study (Hadjivassiliou et al., 2010). Moreover, TBR1+ cells were observed in the superficial region of the neocortex of FCD IIb and in the white matter of FCD Ia and Ib.

SATB2, FOXP1, and TBR1 are normally expressed in immature neurons. Based on our results indicating the diffuse expression of all three markers throughout the neocortex of FCD and HME, it appears that immature cells are present in the cortex of both.

Furthermore, our results confirm that abnormal migration occurs, and this is supported by the well-known fact that epileptic malformed brains often have neuronal migration disruption. Our data may indicate that FCD and HME are developmental brain disorders characterized by abnormalities in neuronal migration and crucially differentiation.

It is of considerable interest that the immature transcription markers of the neocortex are useful to identify FCD subtypes and HME. SATB1-, FOXP1-, and TBR1-positive cells are expected to be projection neurons, which use excitatory neurotransmitter glutamate. Electrophysiological experiments have demonstrated that immature and potentiated excited-GABAergic neurons are strongly related to FCD epileptogenesis (Cepeda et al., 2007). Moreover, dysmorphic neurons and balloon cells are thought to originate from the neocortical subventricular zone (Lamparello et al., 2007). From our data, glutaminergic neurons of FCD may also have neuronal immaturity. The pathological causes of the intractable seizures in FCD and HME may thus include the immaturity and dysfunction of neurons.

4. Experimental procedures

4.1. Human tissue preparation

Human neocortical tissues were obtained from 16 FCD and seven HME patients (Table 1) who underwent surgical treatment