MR-Revealed Myelination in the Cerebral Corticospinal Tract as a Marker for Pelizaeus-Merzbacher’s Disease with Proteolipid Protein Gene Duplication

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BACKGROUND AND PURPOSE: Pelizaeus-Merzbacher’s disease (PMD) is caused by mutations in the proteolipid protein (PLP) gene. Recent studies have shown that an increased PLP dosage, resulting from total duplication of the PLP gene, invariably causes the classic form of PMD. The purpose of this study was to compare the MR findings of PMD attributable to PLP duplication with those of PMD arising from a missense mutation.

METHODS: Seven patients with PMD, three with a PLP missense mutation in either exon 2 or 5 (patients 1–3), and four with PLP duplication (patient 4 having larger PLP duplication than patients 5–7) were clinically classified as having either the classic or connatal form of PMD. Cerebral MR images were obtained to analyze the presence of myelination and T1 and T2 shortening in the deep gray matter. Multiple MR studies were performed in six of the seven patients to analyze longitudinal changes.

RESULTS: Four patients (patients 1–4) were classified as having connatal PMD, whereas the other three (patients 5–7) were classified as having classic PMD. Myelination in the cerebral corticospinal tract, optic radiation, and corpus callosum was observed in three cases of classic PMD with PLP duplication. In patient 4, myelination extended to the internal capsule, corona radiata, and centrum semiovale over a 3-year period. No myelination was observed in three PMD cases with a PLP point mutation. T2 shortening in the deep gray matter was recognized in all patients with PMD.

CONCLUSION: The presence of myelination in the cerebral corticospinal tract with diffuse white matter hypomyelination on MR images could be a marker for PMD with PLP duplication. It is suggested that progression of myelination may be present in connatal PMD with large PLP duplication.

Pelizaeus-Merzbacher’s disease (PMD) is a rare X-linked inherited disorder affecting myelination of the central nervous system (1, 2). Pathologically, PMD, in contrast to other leukodystrophies like metachromatic leukodystrophy, adrenoleukodystrophy, and multiple sclerosis, is a dysmyelinating rather than a demyelinating disorder. In demyelinating disorders, myelin is formed, deposited around axons, and then destroyed later. In dysmyelinating disorders such as PMD, normal myelination never occurs.

Based on the time of onset and the clinical severity, PMD has traditionally been divided into four categories: the classic, connatal, transitional, and adult forms (1, 2). The classic and connatal forms are the most common. Classic PMD has its onset during late infancy. Early symptoms include nystagmoid, dancing or trembling eye movements, and delayed motor development followed by involuntary movements and spasticity. The course is usually protracted and it is often misdiagnosed as cerebral palsy. Connatal PMD is a rarer and more severe variant that begins at birth or in early infancy and has a more severe clinical course. Abnormal nystagmoid eye movements, extrapyramidal hyperkinesia, spasticity, optic atrophy, and seizures also occur during the early stage.

The basic defect in PMD patients is a defect in the proteolipid protein (PLP) gene. PLP is the most abundant protein in central nervous system myelin, accounting for approximately 50% of the total myelin.
elin protein (3). Traditionally, PLP and an isoform DM20, produced by alternative splicing of exon 3B, have been presumed to play a structural role in compact myelin, but more recent studies have suggested an additional role in glial cell development (4). A wide range of mutations in the PLP gene can result in PMD. The deletion of the 18q 22.3-ter region, which includes the myelin basic–protein gene, can also cause a variable myelin defect (5).

Hodes et al (6) reviewed 21 different missense mutations of the PLP gene, mostly located in exons 3 and 4. Missense mutations impair the cellular transport of PLP proteins to the membrane. The resulting accumulation of PLP in the endoplasmic reticulum finally leads to oligodendrocytic cell death. Mutations that inhibit the transport of both PLP and DM20 are found in patients with connatal PMD, whereas patients with classic PMD carry mutations that impair the transport of PLP only (7). Nonsense mutations, splice-site mutations, and small deletions have also been described (8). Sequential analysis of the entire PLP coding region, however, has failed to reveal mutations in approximately 80% of the patients with PMD.

Recent studies have shown that an increased PLP dosage, resulting from total duplication of the PLP gene, also causes PMD (8, 9). PLP over-expression caused by duplication of the gene also leads to glial cell degeneration. Duplication of the PLP gene has been shown to be a major cause of PMD (8, 9). PMD patients with PLP duplication invariably have the classic form with a relatively mild clinical course (8–10).

MR imaging is a useful method for assessing the dysmyelination of the cerebral white matter in PMD. MR imaging can show a hypomyelination pattern; ie, reversal of the white matter signal intensity on T1- and T2-weighted images (11). Nonetheless, there has been no report of a correlation between the genotype and MR imaging. The purpose of this study was to examine the MR findings of PMD attributable to PLP duplication compared with those arising from a missense PLP mutation.

**Methods**

Seven male patients with PMD, aged 2 to 17 years and from seven unrelated Japanese families, were involved in this study. The patients came from two hospitals in the city of Chiba. Genetic analysis of their PLP genes, obtained with previously reported methods (12), revealed missense point mutations in exon 2 (Leu45[CTA] → Arg[CGA]) in patient 1, exon 5 (Val209[GTT] → Asp[GAT]) in patient 2, and exon 5 (Pro210[CCA] → Leu[CTA]) in patient 3 (13). The proton MR spectroscopic findings in patients 1 and 3 were reported previously (14). These missense point mutations impaired the transport of both PLP and DM20. PLP gene duplications were identified in the other four patients (patients 4–7) by using an interphase fluorescence in situ hybridization (FISH) assay (9). To determine the size of the duplication and appropriate locations of the DNA rearrangement breakpoints, phage P1 artificial chromosomes (PAC) clones were used as FISH probes for breakpoint mapping (9). The size of the duplication in pa-

The patients were classified into the classic or connatal form based on the time of disease onset and the clinical severity (1). MR imaging was performed with a 0.5-T superconducting magnet in six patients (the exception being patient 6), or a 1.5-T superconducting magnet in five patients (patients 1, 3, 4, 5, and 6). At least two scans were obtained in 6 patients (patients 1–6). Axial or coronal T2-weighted images or both (2000/80/2 [TR/TE/excitations] for 0.5-T, 3000–4000/100/2 for 1.5-T) and T1-weighted images (500/30/2, TI = 700 for 0.5 T; 500/30/2 for 1.5-T) were obtained in all seven patients. The parameters were as follows: matrix size, 224 × 160 for 0.5-T, 256 × 256 or 256 × 192 for 1.5-T; field of view, 25 cm; slice thickness, 7 mm for 0.5-T, 6 mm for 1.5-T, and slice gap, 2 mm. The images were examined for signs of myelination (comparison of T1 and T2 shortening within the cortex) in parts of the brain; that is, the cerebral corticospinal tract (the internal capsule, corona radiata, centrum semiovale, pre- or postcentral cortex, and subcortical white matter). Optic radiation, corpus callosum, and cerebellar white matter. On the basis of the signal intensity, myelination was graded as “absent,” “present,” and “intermediate.” Absent myelination indicated a higher signal on T2-weighted and a lower signal on T1-weighted images in the cerebral cortex. Present myelination indicated a lower-to-equal signal on T2- and a higher signal on T1-weighted images in the cerebral cortex. Intermediate myelination showed a signal between that of absent and present myelination. When myelination was present, the “tigroid” appearance of preserved myelin was also evaluated.

T1 and T2 shortening within the cortex was evaluated in the deep gray matter regions; ie, the thalamus, striatum, and globus pallidus. Cerebral and cerebellar atrophy was also examined. Cerebral CT was performed at least once in all patients to identify low-density white matter areas and to rule out calcification, which could also shorten the T1 and T2 values.

**Results**

The clinical data and MR findings in each patient are presented in the following Table. Four patients (all three patients with a PLP point mutation and one with large PLP duplication [patients 1–4]) were classified as having connatal PMD, whereas the other three patients with PLP duplication (patients 5–7) were identified as having classic PMD. Nystagmus began at birth or soon after birth in cases of connatal PMD, and after 3 weeks in classic PMD. Patient 5 could crawl, and patients 6 and 7 could walk with assistance. On the other hand, the patients with connatal PMD never acquired head control.

Myelination in the cerebral corticospinal tract (the internal capsule, corona radiata, centrum semiovale, pre- or postcentral cortex, and subcortical white matter), optic radiation, and corpus callosum was observed in three cases of classic PMD with PLP duplication (patients 5–7, Fig 1). High signal intensity on T1-weighted images was more prominent than was low signal intensity on T2-weighted images. In a case of connatal PMD with large PLP duplication (patient 4), myelination was recognized only in the posterior limb of the internal capsule at age 1 year 9 months (Fig 2A–B). Myelination, however, extended into the internal capsule, corona radiata, and centrum semiovale 3 years later (Fig 2C–D). On the other hand, no myelination was ob-
Clinical and MR findings for all seven patients

<table>
<thead>
<tr>
<th>Patient</th>
<th>Age/sex</th>
<th>PLP gene mutation</th>
<th>Motor level</th>
<th>Clinical form</th>
<th>MRI (tesla)</th>
<th>Scan age</th>
<th>Cerebral atrophy</th>
<th>Cerebellar atrophy</th>
<th>High on T1WI</th>
<th>Low on T2WI</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>7Y/Male</td>
<td>Leu 45 Arg</td>
<td>no head control</td>
<td>connatal</td>
<td>0.5</td>
<td>9M</td>
<td>x</td>
<td>o</td>
<td>x</td>
<td>x</td>
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<tr>
<td>2</td>
<td>17Y/Male</td>
<td>Val 209 Asp</td>
<td>no head control</td>
<td>connatal</td>
<td>0.5</td>
<td>5Y</td>
<td>o</td>
<td>x</td>
<td>x</td>
<td>x</td>
</tr>
<tr>
<td>3</td>
<td>9Y/Male</td>
<td>Pro 210 Leu</td>
<td>no head control</td>
<td>connatal</td>
<td>1.5</td>
<td>10Y</td>
<td>ox</td>
<td>x</td>
<td>x</td>
<td>x</td>
</tr>
<tr>
<td>4</td>
<td>5Y/Male</td>
<td>large duplication</td>
<td>no head control</td>
<td>connatal</td>
<td>1.5</td>
<td>1Y9M</td>
<td>x</td>
<td>x</td>
<td>x</td>
<td>x</td>
</tr>
<tr>
<td>5</td>
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<td>crawl</td>
<td>classic</td>
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<td>5Y</td>
<td>x</td>
<td>x</td>
<td>x</td>
<td>x</td>
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<tr>
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<td>assist walk</td>
<td>classic</td>
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<td>10Y</td>
<td>o</td>
<td>x</td>
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<tr>
<td>7</td>
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<td>assist walk</td>
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<td>1Y2M</td>
<td>ox</td>
<td>x</td>
<td>x</td>
<td>x</td>
</tr>
</tbody>
</table>

Note.—o, indicates present; x, absent; ox, intermediate; Y, year; M, month; T1WI, T1-weighted image; IC, internal capsule; CS, centrum semiovale; WM, white matter; CC, corpus callosum; Th, thalamus; GP, globus pallidus; T2WI, T2-weighted image.
observed in the three PMD cases with PLP point mutations (patients 1–3, Fig 3). No progression of myelination was observed in six patients, the exception being patient 4. A tigroid pattern of myelination was not observed in any patient.

T1 and T2 shortening in the deep gray matter (ie, striatum, globus pallidus, and thalamus) was, to some degree, recognized in all patients in this study. Low signal intensity on T2-weighted images was more significant than high signal intensity on T1-weighted images. The longitudinal MR study showed progressive T2 shortening in the basal ganglionic regions.

Cerebral and cerebellar atrophy was present in connatal PMD. The atrophy was not present on initial scanning in patients 1 and 3 at the ages of 9 and 10 months. In the other four patients, no atrophy was observed.

Calcification was absent in all patients. Also, CT showed slightly low-density areas in the cerebral and cerebellar white matter in all patients; however, differences could not be seen between PMD with
PLP point mutations and PMD with PLP duplication.

Discussion

From the MR perspective, postnatal brain development consists primarily of changes in signal intensity secondary to the process of myelination (15). Brain myelination on MR images is shown to occur at different rates and at different times on T1-weighted images compared with those found on T2-weighted images. For example, at birth, high signal intensity on T1-weighted images is observed in the cerebellar peduncle and the posterior limb of the internal capsule, which shows low signal intensity on T2-weighted images during the interval between birth and 2 months after birth. Changes in white matter maturation are seen best on T1-weighted images during the first 6 to 8 months of life, and on T2-weighted images between 6 and 18 months. The exact reason for these differences has not been determined. It, however, is known that T1 shortening correlates temporally with the increased interaction of the hydrogen molecules of water binding to the member components of myelin as well as with the increases in cholesterol and glycolipids that accompany the formation of myelin from oligodendrocytes. T2 shortening correlates temporally with the tightening of the myelin spiral around the axon; that is, the maturation of the myelin sheath (15, 16). The seven PMD patients in this study were older than 9 months and therefore should have had an essentially adult pattern, at least on T1-weighted images.

Three patients with classic PMD with PLP duplication (patients 5–7) exhibited myelination on MR imaging in the cerebral corticospinal tract, optic radiation, and corpus callosum. In fact, pathologic analysis showed a relatively good state of myelination within the internal capsule (10) in cases of classic PMD. The T1 shortening in these regions was more prominent than was T2 shortening. This immature myelination pattern, also seen in the neonatal period (15), might reflect the pathologic findings in classic PMD; that is, the myelin sheath in the myelin islets is more or less damaged and shows varicose and bandlike distension, thinning, and pallor (10).

Patient 4 (conntal PMD with large PLP duplication) presented interesting neuroradiologic and clinical findings. Nezu et al (17) and van der Knaap and Valk (18) previously reported the absence of any progression of myelination in PMD. They suggested an arrest of myelination occurred before or soon after birth in PMD. MR imaging in patient 4, however, revealed longitudinal progression of myelination in the internal capsule, corona radiata, and centrum semiovale over a 3-year period. Myelination in the optic radiation and corpus callosum, which was present in the other three classic PMD cases with PLP duplication, was absent on follow-up MR imaging. The state of myelination in patient 4 was, therefore, between that of classic PMD with PLP duplication and conntal PMD with a PLP point mutation. From the clinical point of view, more than 40 PMD patients with PLP duplication had classic PMD with a relatively mild clinical course (19); however, patient 4 was classified as having conntal PMD with severe clinical manifestations, although he had PLP duplication.

We recently reported on 13 PMD patients with PLP duplication, who commonly had classic PMD (9). Among them, two exceptionally severe cases (clinically conntal PMD) carried large duplications, suggesting that either the size of the duplicated genomic region or the location of the breakpoint may affect the clinical severity. A duplicated large fragment may contain other genes as well as PLP. The duplication may affect these genes, resulting in the phenotypic variation and delayed myelination that was seen on MR images in patient 4.

On the other hand, myelination was totally absent on MR images in three cases of conntal PMD with a PLP point mutation. This finding coincides with the pathologic findings in conntal PMD; that is, dysmyelination spreads diffusely over all parts of the brain and is so intense that, in large areas, not a single myelinated fiber is to be found. Accordingly, the presence of myelination in the cor-
In the deep gray matter is maturational changes of the neuronal network (25). Maturational changes of neurons, particularly increasing synaptic density, may reduce the amount of free water in the brain, resulting in shortening of the T2 value (26). In PMD, axonal processes appear normal, and the neuronal architecture is unchanged pathologically. Proton MR spectroscopy revealed a normal N-acetylaspartate/creatine ratio (N-acetylaspartate is believed to be only present in neurons and is regarded as a neuronal marker) in two of our patients with PMD that was attributed to a PLP point mutation (14). Accordingly, normal neuronal maturation in the deep gray matter may cause signal changes without myelination.

Conclusion

Finally, the presence of myelination in the cerebral corticospinal tract, with diffuse white matter hypomyelination observed on MR images, could be a marker for PMD with PLP duplication and may help to establish a clinical prognosis. It is suggested that progression of myelination may occur in connatal PMD with large PLP duplication.

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References