Multiplex Families With Multiple System Atrophy

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Background: Multiple system atrophy (MSA) has been considered a sporadic disease, without patterns of inheritance.

Objective: To describe the clinical features of 4 multiplex families with MSA, including clinical genetic aspects.

Design: Clinical and genetic study.

Setting: Four departments of neurology in Japan.

Patients: Eight patients in 4 families with parkinsonism, cerebellar ataxia, and autonomic failure with age at onset ranging from 58 to 72 years. Two siblings in each family were affected with these conditions.

Main Outcome Measures: Clinical evaluation was performed according to criteria by Gilman et al. Trinucleotide repeat expansion in the responsible genes for the spinocerebellar ataxia (SCA) series and for dentatorubral-pallidoluysian atrophy (DRPLA) was evaluated by polymerase chain reaction. Direct sequence analysis of coding regions in the \( \alpha \)-synuclein gene was performed.

Results: Consanguineous marriage was observed in 1 of 4 families. Among 8 patients, 1 had definite MSA, 5 had probable MSA, and 2 had possible MSA. The most frequent phenotype was MSA with predominant parkinsonism, observed in 5 patients. Six patients showed pontine atrophy with cross sign or slitlike signal change at the posterolateral putaminal margin or both on brain magnetic resonance imaging. Possibilities of hereditary ataxias, including SCA1 (ataxin 1, \( \text{ATXN1} \)), SCA2 (\( \text{ATXN2} \)), Machado-Joseph disease/SCA3 (\( \text{ATXN3} \)), SCA6 (\( \text{ATXN6} \)), SCA7 (ATXN7), SCA12 (protein phosphatase 2, regulatory subunit B, \( \beta \) isofrom; \( \text{PP2R2B} \)), SCA17 (TATA box binding protein, \( \text{TBP} \)) and DRPLA (atrophin 1; \( \text{ATN1} \)), were excluded, and no mutations in the \( \alpha \)-synuclein gene were found.

Conclusions: Findings in these multiplex families suggest the presence of familial MSA with autosomal recessive inheritance and a genetic predisposition to MSA. Molecular genetic approaches focusing on familial MSA are expected to provide clues to the pathogenesis of MSA.

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Molecular genetic analysis

Genomic DNA was extracted from peripheral blood leukocytes after obtaining informed consent. Expansions of CAG repeats of the genes for dominant spinocerebellar ataxias (SCAs), including SCA1 (ataxin 1; ATXN1 [for gene nomenclature see http://www.gene.ucl.ac.uk/nomenclature]), SCA2 (ATXN2), Machado-Joseph disease/SCA3 (ATXN1), SCA6 (ATXN1), SCA7 (ATXN7), SCA12 (protein phosphatase 2, regulatory subunit B, β isoform, PP2R2B), SCA17 (TATA box binding protein, TBP), and dentatorubral-pallidoluysian atrophy (DRPLA) [atrophin 1, ATN1], were analyzed using an automated DNA sequencer (ABI 377; Applied Biosystems, Foster City, Calif) in the affected patients. Direct sequence analysis of 6 coding exons of the α-synuclein (SNCA) gene was performed using the ABI 3100 DNA sequencer in all patients.

Results

Two siblings were affected in each family. The parents of siblings in each family had no clinical signs of extrapyramidal or cerebellar disorders. Consanguineous marriage was present in family A, in which the parents were first-degree cousins. Among 8 patients, 1 (II-4 in family A) had definite MSA, 5 (II-8 in family A, II-2 and II-9 in family B, II-5 in family C, and II-7 in family D) had probable MSA, and 2 (II-4 in family C and II-3 in family D) had possible MSA. The most frequent phenotype was MSA with predominant parkinsonism (MSA-P), observed in 5 patients. One patient showed an MSA of the cerebellar type (MSA-C) phenotype, and 2 patients showed an MSA-P+C phenotype. The clinical phenotypes were concordant between the affected siblings in the 3 families (families A, B, and C). The mean age at onset was 65.9 years (age range, 58-72 years). Six patients showed pontine atrophy with cross sign or slitlike signal change at the posterolateral putaminal margin or both on brain MR imaging.

Testing for the trinucleotide repeat expansions in the responsible genes for SCA1 (ATXN1), SCA2 (ATXN2), Machado-Joseph disease/SCA3 (ATXN1), SCA6 (ATXN1), SCA7 (ATXN7), SCA12 (PP2R2B), SCA17 (TBP), and DRPLA (ATN1) gave normal results. No mutations in the SNCA gene were found in the family members.

Report of Cases

Family A

Patient II-4

Patient II-4 was diagnosed as having retinitis pigmentosa at age 33 years by an ophthalmologist. She noticed resting tremor in her left hand at age 68 years, followed by the development of bradykinesia and gait disturbance at age 69 years. Neurological examination at age 71 years revealed bradykinesia, a masklike face, urinary frequency, severe rigidity in the 4 limbs, and limb ataxia predominant on the left side. Parkinsonism was not improved by levodopa treatment. She did not show dementia, muscular atrophy, sensory disturbance, or limitation of ocular movement. Brain computed tomography showed distinct cerebellar atrophy. Brain MR imaging was not performed. She had recurrent pneumonia and died at age 73 years. On postmortem examination, the brain weighed 1030 g before fixation. On gross examination, the cerebellum and pontine base were moderately atrophic. The inferior olivary nuclei were slightly atrophic. Moderate depigmentation in the substantia nigra and locus coeruleus was observed. Microscopically, moderate to severe loss of neurons with gliosis was observed in the pontine nuclei. The transverse bundles in the pontine base...
showed severe loss of myelinated nerve fibers. The loss of Purkinje cells was mild to moderate, and the cerebellar white matter was severely degenerated. In the cerebellar dentate nucleus, neuronal shrinkage associated with gliosis was observed. Moderate neuronal loss associated with gliosis was observed in the substantia nigra and locus coeruleus. In the putamen, a small amount of GCIs was observed, but no obvious neuronal loss was observed. Mild neuronal loss was also detectable in the red nucleus and dorsal vagal and vestibular nuclei. In the thalamic medial nuclei, gliosis without distinct neuronal loss was observed. Numerous argyrophilic GCIs and an abnormal accumulation of α-synuclein–precursor of the non-Α4 component of Alzheimer disease amyloid (NACP) in glial cells were observed in the substantia nigra, pontine base, inferior olive, and cerebellar white matter (Figure 2). In the retina, severe loss of rods and cones and patchy disappearance of pigmentary epithelial cells were observed.

Patient II-8

Patient II-8 developed night blindness at age 48 years and was diagnosed as having retinitis pigmentosa at age 51 years. He had mild resting tremor in both hands, bradykinesia, and rigidity at age 62 years, which were symmetrical and refractory to a combined therapy of 200 mg of levodopa–dopadecarboxylase inhibitor. He had difficulty in urination at age 63 years, orthostatic hypotension at age 64 years, and truncal ataxia at age 65 years. He became bedridden at age 66 years because of the progression of his symptoms. On laboratory investigation, his protein level, cell counts, glucose concentration, IgG index, and 5-hydroxyindoleacetic acid level in cerebrospinal fluid were within normal ranges. His homovanillic acid level was decreased to 19.9 ng/mL (reference range, 28-77 ng/mL). Thyroid hormone levels, serum and urinary copper levels, serum ceruloplasmin level, lysosomal enzyme activities in leukocytes, and vitamin B12, B12, and E levels were normal. Nerve conduction studies revealed no abnormalities. Brain MR imaging showed mild atrophy of the brain. A summary of the clinical findings among 4 multiplex families with multiple system atrophy (MSA) is provided in Table 1. The criteria used for diagnosis were based on the guidelines by Gilman et al.1

Table. Clinical Findings Among 4 Multiplex Families With Multiple System Atrophy (MSA)

<table>
<thead>
<tr>
<th>Family and Affected Individuals</th>
<th>Finding</th>
<th>A</th>
<th>B</th>
<th>C</th>
<th>D</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Sex</strong></td>
<td>Female</td>
<td>Male</td>
<td>Male</td>
<td>Male</td>
<td>Female</td>
</tr>
<tr>
<td><strong>Age at onset, y</strong></td>
<td>68</td>
<td>62</td>
<td>72</td>
<td>63</td>
<td>68</td>
</tr>
<tr>
<td><strong>Age at examination, y</strong></td>
<td>71</td>
<td>66</td>
<td>66</td>
<td>66</td>
<td>72</td>
</tr>
<tr>
<td><strong>Initial symptoms</strong></td>
<td>Tremor</td>
<td>Ataxia</td>
<td>Tremor</td>
<td>Tremor</td>
<td>Tremor</td>
</tr>
<tr>
<td><strong>Parkinsonism</strong></td>
<td>*</td>
<td>*</td>
<td>*</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td><strong>Cerebellar sign</strong></td>
<td>††</td>
<td>††</td>
<td>††</td>
<td>††</td>
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</tr>
<tr>
<td><strong>Upper limb</strong></td>
<td>Enhanced</td>
<td>Normal</td>
<td>Normal</td>
<td>Normal</td>
<td>Normal</td>
</tr>
<tr>
<td><strong>Lower limb</strong></td>
<td>Enhanced</td>
<td>Normal</td>
<td>Normal</td>
<td>Normal</td>
<td>Normal</td>
</tr>
<tr>
<td><strong>Urinary dysfunction</strong></td>
<td>††</td>
<td>††</td>
<td>††</td>
<td>††</td>
<td>††</td>
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<tr>
<td><strong>Orthostatic hypotension</strong></td>
<td>††</td>
<td>††</td>
<td>††</td>
<td>††</td>
<td>††</td>
</tr>
<tr>
<td><strong>Response to levodopa</strong></td>
<td>Poor</td>
<td>Poor</td>
<td>Poor</td>
<td>Poor</td>
<td>Poor</td>
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<tr>
<td><strong>Brain magnetic resonance imaging</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>Cerebellar atrophy</strong></td>
<td>NE</td>
<td>NE</td>
<td>NE</td>
<td>NE</td>
<td>NE</td>
</tr>
<tr>
<td><strong>Pontine atrophy</strong></td>
<td>NE</td>
<td>NE</td>
<td>NE</td>
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</tr>
<tr>
<td><strong>Cross sign</strong></td>
<td>NE</td>
<td>NE</td>
<td>NE</td>
<td>NE</td>
<td>NE</td>
</tr>
<tr>
<td><strong>Slitlike signal change at the posterolateral putaminal margin</strong></td>
<td>NE</td>
<td>NE</td>
<td>NE</td>
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<td>NE</td>
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<tr>
<td><strong>Criteria by Gilman et al</strong></td>
<td>Definite</td>
<td>Probable</td>
<td>Probable</td>
<td>Probable</td>
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</tr>
<tr>
<td><strong>Phenotypes</strong></td>
<td>MSA-P</td>
<td>MSA-P</td>
<td>MSA-P</td>
<td>MSA-P</td>
<td>MSA-P</td>
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<tr>
<td><strong>Complication</strong></td>
<td>Rheumatoid arthritis</td>
<td>Rheumatoid arthritis</td>
<td>Rheumatoid arthritis</td>
<td>Rheumatoid arthritis</td>
<td>Rheumatoid arthritis</td>
</tr>
</tbody>
</table>

Abbreviations: MSA-C, MSA of the cerebellar type; MSA-P, MSA with predominant parkinsonism; NE, not examined; RP, retinitis pigmentosa; ellipsis, not applicable.

*Severe finding.
†Moderate finding.
‡Finding present.

Figure 2. Histopathologic features of the pons from patient II-4 in family A. A. Argyrophilic glial cytoplasmic inclusions (arrows) (Gallyas-Braak stain). B. Abnormal accumulation of α-synuclein–precursor of the non-Α4 component of Alzheimer disease amyloid (NACP) in glial cells (arrows).
pons and cerebral cortex on T1-weighted imaging. Atrophy of the pontine base accompanied by cross sign on T2-weighted and proton density–weighted imaging was observed. Slitlike signal changes with hyperintensity on T1-weighted imaging and hypointensity on T2-weighted imaging were observed in the posterolateral putaminal margin (Figure 3A). Single-photon emission computed tomography images showed mild hypoperfusion in the pons, basal ganglia, and right frontal and left parietal lobes.

Both affected siblings in family A had retinitis pigmentosa. The other siblings were unaffected.

FAMILY B

Patient II-2 began to have resting tremor in the right hand at age 72 years, followed by rigidity in the 4 extremities, parkinsonian gait, urinary incontinence, and orthostatic hypotension within a year. These symptoms responded poorly to a combined therapy of 900 mg of levodopa–dopadecarboxylase inhibitor. He required support in walking at age 76 years. Brain MR imaging revealed slitlike signal change at the posterolateral putaminal margin on T2-weighted imaging and pontine atrophy on T1-weighted imaging (Figure 3B).

Patient II-9

The brother of patient II-2, patient II-9, developed resting tremor in the right hand at age 63 years and in the right leg at age 65 years. Neurological examination at age 66 years revealed severe rigidity in the 4 extremities, bradykinesia, mild limb ataxia, urinary incontinence, and orthostatic hypotension. The symptoms responded poorly to a combined therapy of 900 mg of levodopa–dopadecarboxylase inhibitor. Brain MR imaging showed slitlike signal change at the posterolateral putaminal margin, atrophy of the cerebellar vermis, and cross sign in the pons on T2-weighted imaging (Figure 3C).

FAMILY C

Patient II-4 had cerebellar ataxia at age 68 years, followed by the development of resting tremor predominantly on the left side, neck rigidity, parkinsonian gait, and urinary frequency. These symptoms were progres-
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Introduction:

Corticobasal degeneration is a neurodegenerative disorder characterized by motor and sensory symptoms, with a high prevalence among first-degree or second-degree relatives of patients with MSA. In this study, we aimed to investigate the epidemiology and possible genetic factors of MSA in four families with autosomal dominant inheritance. We found that 6 of 8 patients in these families fulfilled the clinical criteria for progressive supranuclear palsy, while the other 2 patients developed rigidity predominantly on the left side and truncal ataxia. We also observed severe cerebellar ataxia, pyramidal signs such as Babinski sign, extrapyramidal signs such as dysarthria and urinary incontinence. We concluded that MSA susceptibility genes strongly contribute to the pathogenesis among these families.

Methods:

We examined the clinical, radiological, and genetic features of MSA in four families with autosomal dominant inheritance. We also performed a molecular diagnosis in all affected individuals.

Results:

We found that 6 of 8 patients in these families fulfilled the clinical criteria for progressive supranuclear palsy, while the other 2 patients developed rigidity predominantly on the left side and truncal ataxia. We also observed severe cerebellar ataxia, pyramidal signs such as Babinski sign, extrapyramidal signs such as dysarthria and urinary incontinence. We concluded that MSA susceptibility genes strongly contribute to the pathogenesis among these families.

Discussion:

We identified some characteristic features of MSA in these families, such as severe cerebellar ataxia, pyramidal signs such as Babinski sign, extrapyramidal signs such as dysarthria and urinary incontinence. We also found that MSA susceptibility genes strongly contribute to the pathogenesis among these families.

Conclusion:

We concluded that MSA susceptibility genes strongly contribute to the pathogenesis among these families. We also identified some characteristic features of MSA in these families, such as severe cerebellar ataxia, pyramidal signs such as Babinski sign, extrapyramidal signs such as dysarthria and urinary incontinence.

In conclusion, MSA is a complex disease that involves genetic factors, environmental factors, and epigenetic factors. Further studies are needed to elucidate the genetic and environmental factors that contribute to MSA.
This is inconsistent with the fact that the MSA-C phenotype is the most prevalent form of MSA in Japan. According to a recent nationwide survey in Japan, 82.5% of patients with MSA have the MSA-C phenotype, while 10.6% have the MSA-P phenotype. Fourth, the clinical phenotypes were concordant between the affected siblings in 3 of 4 families. Third, the mean age at onset (65.9 years) of these family members is later than that among patients with sporadic MSA (age range, 55.4-57.5 years). This observation is interesting. More pathologically proven cases of familial MSA are required to reveal the pathological features of familial MSA.

Because the SNCA gene is the major component of GCIs and of neuronal cytoplasmic inclusions in MSA,2,4 we considered this a candidate gene and conducted mutational analysis of the coding regions but failed to find any mutations in our patients. A previous study demonstrated no significant changes in the expression levels of the SNCA gene messenger RNA in the brains of patients with MSA compared with those of control subjects.5

Taken together, the SNCA gene is unlikely to be the causative gene for MSA. Other genetic or environmental factors associated with SNCA might be operable in the pathogenesis of MSA. To identify genetic factors related to MSA, investigations of additional multiplex families with MSA are necessary.

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