Case Report

Monoclonal Gammopathy of Undetermined Significance (MGUS) in a Man with Fragile X-associated Tremor/Ataxia Syndrome

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We report the clinical presentation and laboratory findings of a 69-year-old man with fragile X-associated tremor ataxia syndrome (FXTAS), a progressive neurodegenerative disorder, who was noted to have monoclonal gammopathy of undetermined significance (MGUS), a plasma cell proliferative disorder and a precursor disease of multiple myeloma. Both MGUS and FXTAS are associated with microRNA (miRNA) dysregulation. We speculate that individuals with FXTAS may be predisposed to MGUS and further studies are warranted regarding this association.

1. Introduction

Fragile X-associated tremor/ataxia syndrome (FXTAS) is an adult progressive neurodegenerative disorder that affects predominantly male carriers of the fragile X mental retardation 1 (FMR1) gene premutation (CGG expansion of 55–200 repeats). The clinical features of FXTAS are progressive intention tremor, cerebellar ataxia, autonomic dysfunction, peripheral neuropathy, cognitive decline, middle cerebellar peduncle signs (MCPs), and white matter disease combined with brain atrophy on MRI [1–4].

The penetrance of FXTAS in male carriers older than age 50 is approximately 30 to 45.5% and 8 to 16.5% in females [2, 5, 6]. Overproduction of the FMR1 mRNA, up to 8 fold times normal, is the basic mechanism of FXTAS [7]. The elevated FMR1 mRNA causes a gain-of-function toxicity to cells leading to the formation of intranuclear inclusions in neurons and astrocytes throughout the central nervous system (CNS) and subsequent sequestration of critical proteins necessary for normal cell function [8, 9]. Mitochondrial dysfunction also occurs in older premutation carriers both with and without FXTAS [10, 11]. This cellular dysfunction leads to neuronal cell death and white matter disease in the CNS [3, 12, 13] associated with the clinical features of FXTAS.

Monoclonal gammopathy of undetermined significance (MGUS) is a plasma cell proliferative disorder characterized by a plasma cell level of less than 10% in the bone marrow, a monoclonal paraprotein band less than 30 g/L, and no end organ damage (CRAB—hypercalcemia, renal insufficiency, anemia, and bone lesions) [14]. It is an age-related medical condition found predominantly in older persons [15]. The diagnosis of MGUS is usually an unexpected event during an evaluation, and because it is typically asymptomatic, it is underdiagnosed [16]. However, some abnormalities such as bone loss or osteoporosis, peripheral neuropathy, and thromboembolic events can be associated to this disorder [17]. Individuals with MGUS have a higher risk of developing multiple myeloma (MM) or related malignancy [18–20]. Race, ethnicity, advance age, male sex [15], family history [21], and exposure to certain pesticides [22] increase...
the risk of MGUS indicating that MGUS is a multifactorial condition.

The pathogenesis of MGUS is hypothesized to begin with an abnormal response to antigenic stimulation mediated by toll-like receptors (TLRs) and interleukin IL-6 receptors and IL-1β [23, 24]. This results in cytotenic abnormalities—either hyperdiploidy or immunoglobulin heavy chain (IgH) translocations [25]. When it is followed by the second hit including RAS or p53 mutation, p18 deletion, myc dysregulation, MGUS progression to MM then occurs [26].

Dysregulation of microRNAs (miRNAs), the small non-coding RNAs involved in posttranscriptional gene regulation, plays a role in MGUS progression to MM [27]. MiRNAs have been demonstrated to play a role in regulation of apoptosis, proliferation, differentiation, cell survival, and oncogenesis [28]. Abnormal miRNA regulation may lead to oncogenesis, particularly MM [29–31]. The miRNA profiles discrepantly expressed in MGUS also show similar aberrant expression in MM suggesting the role of miRNAs in early myelomagenesis [27].

MiRNAs are encoded throughout the genome and are transcribed into pri-miRNA. The pri-miRNA is cleaved in the nucleus by the Drosha RNase III endonuclease. PremiRNAs are transported to the cytoplasm. In cytoplasm, Dicer, a ribonuclease III, cleaves the pre-miRNA into the mature miRNA which forms a complex with RNA-induced silencing complex (RISC). This complex directs the miRNA to the target mRNA leading either to degradation of target mRNA or translational repression [32]. In FXTAS patients, there is Drosha/DGCR8 sequestration by the elevated FMR1 miRNAs that leads to global miRNA dysregulation [33]. We speculate that this global miRNA dysregulation may predispose individuals to MGUS. Here we report the clinical presentation and laboratory findings of a man with FXTAS and MGUS.

2. Clinical Report

A 69-year-old Caucasian man with FXTAS had a premutation of 98 to 103 CGG repeats and an FMR1 mRNA level of 2.33 ± 0.02 times normal. His intention tremor began at age 55 and has been progressive. Currently, it interferes with his activities of daily living (ADL) including eating and handwriting. His gait imbalance began at age 58 and he started using a walker at age 66. His short-term memory problems began at age 63. His stamina has decreased significantly over the last few years and he has some decreased sensation in his lower legs in addition to pain with walking. He has autonomic dysfunction including bladder incontinence and erectile dysfunction beginning at age 66. His past medical history includes hypertension, type 2 diabetes, hyperlipidemia, hypothyroidism, sleep apnea, and migraine headaches (Table 1).

<table>
<thead>
<tr>
<th>Type of gait</th>
<th>Ataxic and broad based</th>
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<tbody>
<tr>
<td>Type of gait</td>
<td>Ataxic and broad based</td>
</tr>
<tr>
<td>Fragile X DNA test (CGG repeat number)</td>
<td>98–103 (premutation range)</td>
</tr>
<tr>
<td>FMR1 mRNA level</td>
<td>2.33 ± 0.02 times normal</td>
</tr>
<tr>
<td>FXTAS diagnosis</td>
<td>Definite</td>
</tr>
<tr>
<td>FXTAS stage</td>
<td>4</td>
</tr>
</tbody>
</table>

On physical examination, his blood pressure was 167/62 mmHg, the heart rate was 71 bpm, the head circumference was 59.5 cm, his height was 172.7 cm, and his weight was 87.5 kg. The ears were slightly prominent. His neurological examination was abnormal with a severe intention tremor with finger to nose touching but absent at rest. There were no deep tendon reflexes but positive primitive reflexes including the snout reflex. His vibration sense was decreased in both feet. His gait was ataxic and broad based. He cannot tandem walk and he had a positive pull test. On MRI, he had significant brain atrophy and white matter disease including a positive MCP sign.

As a part of a workup for his mild normocytic anemia, serum protein electrophoresis with immunofixation revealed 1 g/dL IgG kappa monoclonal paraproteinemia. His serum creatinine was mildly elevated at 1.3 mg/dL with estimated GFR of 55 mL/min. Calcium and highly sensitive C-reactive protein were within the normal reference range.

3. Discussion

This is the first clinical report of MGUS in an individual with FXTAS. Although MGUS is an asymptomatic condition, it is associated with peripheral neuropathy, osteoporosis, and thromboembolic events. Symptoms shared by FXTAS and MGUS include peripheral neuropathy. His risk factors for
Table 2: MGUS laboratory workup of the patient.

<table>
<thead>
<tr>
<th></th>
<th>Levels</th>
<th>Reference range</th>
<th>Status</th>
</tr>
</thead>
<tbody>
<tr>
<td>Red cell count</td>
<td>3.82</td>
<td>4.5–5.9 M/MM3</td>
<td>Low</td>
</tr>
<tr>
<td>Hemoglobin</td>
<td>11.9</td>
<td>13.5–17.5 GM/DL</td>
<td>Low</td>
</tr>
<tr>
<td>Hematocrit</td>
<td>34.7</td>
<td>41–53%</td>
<td>Low</td>
</tr>
<tr>
<td>Glucose</td>
<td>358</td>
<td>70–110 mg/dL</td>
<td>High</td>
</tr>
<tr>
<td>Hemoglobin A1C</td>
<td>8.5</td>
<td>3.9–5.9%</td>
<td>High</td>
</tr>
<tr>
<td>Calcium</td>
<td>9.5</td>
<td>8.6–10.5 mg/dL</td>
<td>Normal</td>
</tr>
<tr>
<td>Serum protein electrophoresis</td>
<td>1 g/dL</td>
<td>IgG kappa monoclonal</td>
<td></td>
</tr>
<tr>
<td>with immunofixation</td>
<td></td>
<td>paraproteinemia</td>
<td></td>
</tr>
<tr>
<td>Serum creatinine</td>
<td>1.3 mg/dL</td>
<td>0.6–1.2 mg/dL</td>
<td>Mildly elevated</td>
</tr>
<tr>
<td>GFR</td>
<td>55</td>
<td>&gt;60</td>
<td>Low</td>
</tr>
<tr>
<td>IgM</td>
<td>17 mg/dL</td>
<td>37 mg/dl–50 mg/dl</td>
<td>Decreased</td>
</tr>
<tr>
<td>Free kappa light chain</td>
<td>7.81 mg/dL</td>
<td>0.33–1.92 mg/dl</td>
<td>Elevated</td>
</tr>
<tr>
<td>Lambda light chain</td>
<td>1.88 mg/dL</td>
<td>0.57–2.63 mg/dl</td>
<td>Normal</td>
</tr>
<tr>
<td>Kappa to lambda light chain</td>
<td>4.15</td>
<td>0.26–1.65</td>
<td>Elevated</td>
</tr>
<tr>
<td>Beta-2 microglobulin</td>
<td>3.38 mcg/mL</td>
<td>1.8 mcg/mL</td>
<td>Elevated</td>
</tr>
<tr>
<td>Bone survey</td>
<td></td>
<td>No lytic or sclerotic lesions</td>
<td></td>
</tr>
</tbody>
</table>

developing an MGUS included advancing age, male sex, and having a mother with non-Hodgkin’s lymphoma, a hematologic cancer. We also suggest that those with FXTAS have a greater risk for MGUS through the common finding of miRNA dysregulation in both disorders.

The DROSHA/DGCR8 sequestration mechanism in FXTAS leads to miRNA alteration, specifically downregulation of several miRNAs [33]. Thus, it may play a role in oncogenesis and tumor biology through posttranscriptional gene regulation impacting the mechanism of proliferation, differentiation, apoptosis, metastasis, and cell survival [28]. In MGUS, miRNAs play an important role in early changes associated with the development of the abnormal clonal plasma cell and have a potential role in altering the p53 pathway [29]. Downregulation or absence of miRNAs has been reported in many cancers [34], including other hematologic malignancies such as acute myeloid leukemia [35], chronic lymphocytic leukemia [36], and diffuse large B-cell lymphoma [37].

It is unknown whether there is an increased risk of having cancer in those with a gene premutation, although there is one case report of a man with lung tumor and full mutation where the tumor’s origin in a cell with premutation instead of full mutation, indicating that the neoplasia development may have been facilitated by the premutation [38]. There is a relationship between elevated mRNA and cAMP response-element binding protein (CREB). The elevated mRNA is due to increased transcription [7], while CREB is involved in FMRI transcriptional activity [39] and may play a role in oncogenesis by various mechanisms including gene amplification, chromosome translocation, interaction with viral oncoprotein, and inactivation of tumor suppressor genes [40].

The current report of MGUS and FXTAS combined with the commonality of miRNA mechanisms in these two conditions suggests that this association may be more common than previously realized and this deserves further study. One possibility is screening for the premutation when MGUS is identified. This would be very important if FXTAS symptoms are present or if there is a family history of fragile X-associated disorders including FXPOI or fragile X syndrome or autism.

Conflict of Interests

Dr. Hagerman has received funding from Novartis, Roche, Seaside Therapeutics, Forest and Curemark for therapeutic trials in fragile X syndrome and autism.

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