

From Clinical Manifestations of Primary Mitochondrial Myopathies (PMM) to Diagnosis: Results from a Patient Journey Analysis Shows Limited Utilization of Genetic Testing

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BACKGROUND

- Primary mitochondrial myopathies (PMM) are a group of disabling and underdiagnosed rare genetic disorders characterized by a range of clinical presentations and multisystemic impact¹
- Diagnosis and management of PMM can be challenging due to the heterogeneity of clinical manifestations, including age of onset¹⁻³
- Patients may engage a variety of healthcare professionals to manage their accumulating symptoms and can receive their first clinical diagnosis from a wide spectrum of specialists³⁻⁵
- Diagnosis with genetic testing is recommended by the Mitochondrial Medicine Society,¹ but utilization remains low
- With no approved treatments for PMM, current treatment involves symptom management^{2,6}
- This approach is not optimal for treating the underlying cause or enabling patients to improve their physical and social functioning⁷
- To better understand the path to PMM diagnosis and management, a patient journey analysis was conducted using Komodo closed-claims data—one of the richest longitudinal data sets available for patient-level analyses

METHODS

- The Komodo database was analyzed for US patients with suspected PMM between 2016-2021
- Due to the absence of a PMM-specific ICD-10 diagnosis code, patients with suspected PMM were identified in payer-complete closed claims via a stepwise approach as follows: mitochondrial disorder identified → myopathy presentation confirmed → secondary mitochondrial disorders excluded
- In patients with suspected PMM, healthcare resource utilization (HCRU) claims were categorized by physician visits before and after the first mitochondrial diagnosis
- Patients included in the analysis were continuously enrolled in medical and pharmacy coverage through the duration of study, ensuring complete longitudinal visibility across all healthcare encounters

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The diagnostic journey for patients with PMM is often long and convoluted; low engagement with geneticists and genetic testing represents a significant gap that creates a suboptimal and burdensome diagnostic approach

