Autosomal Dominant Mental Retardation 1: \textit{MBD5} Gene Sequencing

\textbf{Test Code: SMBD5}
\textbf{Turnaround time: 4 weeks}
\textbf{CPT Codes: 81479 x1}

\section*{Condition Description}

Talkowski \textit{et al.} (2011) mapped the \textit{MBD5} gene (2q23.1) to the critical region of the 2q23.1 deletion syndrome. Haploinsufficiency of the \textit{MBD5} gene causes Autosomal Dominant Mental Retardation syndrome type 1. Overall, of the features evaluated in individuals with 2q23.1 deletion syndrome and \textit{MBD5}-specific deletions, approximately 84\% were observed in both groups. Features associated with the haploinsufficiency of the \textit{MBD5} gene include intellectual disability, developmental delay, motor delay, significant speech impairment, craniofacial manifestations, seizures, constipation, and behavioral problems.

For patients with suspected Autosomal Dominant Mental Retardation syndrome type 1, deletion/duplication analysis is recommended as the first step in mutation identification. For patients in whom mutations are not identified by deletion/duplication analysis, full gene sequencing is appropriate.

\textbf{References:}

\begin{itemize}
  \item OMIM \#611472: \textit{MBD5} gene
  \item OMIM \#156200: Autosomal Dominant Mental Retardation Syndrome Type 1
\end{itemize}

\section*{Genes}

\textbf{MBD5}

\section*{Indications}

This test is indicated for:

\begin{itemize}
  \item Confirmation of a clinical diagnosis of Autosomal Dominant Mental Retardation syndrome type 1 in whom deletion/duplication analysis was negative.
  \item Carrier testing in adults with a family history of Autosomal Dominant Mental Retardation syndrome type 1 in whom deletion/duplication analysis was negative.
\end{itemize}

\section*{Methodology}

PCR amplification of 10 exons contained in the \textit{MBD5} gene is performed on the patient's genomic DNA. Direct sequencing of amplification products is performed in both forward and reverse directions, using automated fluorescence dideoxy sequencing methods. The patient's gene sequences are then compared to a normal reference sequence. Sequence variations are classified as mutations, benign variants unrelated to disease, or variations of unknown clinical significance. Variants of unknown clinical significance may require further studies of the patient and/or family members. This assay does not interrogate the promoter region, deep intronic regions, or other regulatory elements, and does not detect large deletions.

\section*{Detection}

Clinical Sensitivity: Unknown. Mutations in the promoter region, some mutations in the introns and other regulatory element mutations cannot be detected by this analysis. Large deletions will not be detected by this analysis. Results of molecular analysis should be interpreted in the context of the patient's clinical and/or biochemical phenotype.

Analytical Sensitivity: \textasciitilde99\%

\section*{Specimen Requirements}

Submit only 1 of the following specimen types

* Preferred specimen type: Whole Blood

\textbf{Type: Whole Blood}

Specimen Requirements:

In EDTA (purple top) tube:
Infants (2 years): 3-5 ml
Older Children & Adults: 5-10 ml

Specimen Collection and Shipping: Refrigerate until time of shipment. Ship sample within 5 days of collection at room temperature with overnight delivery.

\textbf{Type: Saliva}
Specimen Requirements:

Oragene™ Saliva Collection kit (available through EGL) used according to manufacturer instructions.

Specimen Collection and Shipping: Store sample at room temperature. Ship sample within 5 days of collection at room temperature with overnight delivery.

**Special Instructions**

Deletion/duplication analysis is required before sequence analysis. If deletion/duplication analysis is performed outside of EGL Genetics, please submit a copy of the deletion/duplication report with the test requisition.

**Related Tests**

- Deletion/duplication analysis of the MBD5 gene by CGH array is available and is required before sequence analysis.
- Custom diagnostic mutation analysis (KM) is available to family members if mutations are identified by targeted mutation testing or sequencing analysis.
- Prenatal testing is available only for known familial mutations to individuals who are confirmed carriers of mutations. Please contact the laboratory genetic counselor to discuss appropriate testing prior to collecting a prenatal specimen.