Medical Coverage Policy

Policy Number – MP23-033E Original review date – 02/09/2023 Effective date – 09/18/2024

Whole exome sequencing for autism and developmental delay

Background

Whole exome sequencing is sequencing of the all the protein coding regions of the genome. This represents approximately 1% of the entire genome, around 20,000 genes. High-throughput next generation sequencing techniques along with bioinformatics are used to identify, by comparison to a reference genome, variations associated with genetic disorders (such as SNPs, insertions, deletions, translocations and so on).

Policy statement

Disclaimer: This policy is applicable to TRICARE Prime and Select beneficiaries and may not apply to Active Duty Service Members (ADSM) under Supplemental Health Care Program (SHCP) or TRICARE Prime Remote (TPR) in accordance with TRICARE Operations Manual (TOM) Chapter 17, Section 3. Please review TOM Chapter 17, Section 3, Paragraph 2.0 onwards, regarding SHCP coverage and any TRICARE-specific exclusions included in this coverage policy to accurately determine the benefit for ADSMs.

Whole exome sequencing may be considered medically necessary for the <u>affected person</u> if the following criteria are met:

- I. Patient has unexplained neurodevelopmental disorder, manifesting as developmental delay or intellectual disability
- II. Clinical presentation does not fit a well-described syndrome, as indicated by one or more of the following:
 - a. Absence of clinically recognizable syndrome caused by single gene disorder or chromosomal disorder; OR
 - b. Phenotype suggestive of more than one genetic disorder
- III. Genetic etiology is considered the most likely explanation for the phenotype, as indicated by one or more of the following:
 - a. Multiple congenital abnormalities affecting unrelated organ systems; OR
 - b. Family history strongly suggestive of a genetic etiology (e.g., consanguinity);OR
 - c. Period of unexplained developmental regression; OR
 - d. Biochemical findings suggestive of an inborn error of metabolism
- IV. Results of the test will directly affect treatment and management of the patient
- V. Genetic counseling has been performed, as indicated by all of the following:
 - a. Counseling is provided by healthcare professional with education and training in genetic issues relevant to the genetic tests under consideration
 - b. Counselor is not an employee of the genetics testing laboratory
- VI. Family trio testing (whole exome sequencing of the biologic parents or sibling of the affected person) is considered medically necessary when above criteria are met for the affected person





Limitations of coverage

- I. Whole exome sequencing for isolated autism spectrum disorders, without signs and symptoms of another condition, is not considered medically necessary due to insufficient evidence of clinical benefit.
- II. Whole genome sequencing is not considered medically necessary for any indication due to insufficient evidence of clinical benefit.

Coding information

81415	Exome (e.g., unexplained constitutional or heritable disorder or syndrome); sequence analysis
81416	Exome (e.g., unexplained constitutional or heritable disorder or syndrome); sequence analysis, each comparator exome (e.g., parents, siblings) (List separately in addition to code for primary procedure)
81417	Exome (e.g., unexplained constitutional or heritable disorder or syndrome); reevaluation of previously obtained exome sequence (e.g., updated knowledge or unrelated condition/syndrome)

References

- MCG Health. Whole Genome/Exome Sequencing Developmental Delay and Intellectual Disability. Ambulatory Care 27th edition. ACG: A-0926 (AC). Last updated 9/21/2023
- MCG Health. Whole Genome/Exome Sequencing Autism Spectrum Disorders. Ambulatory Care 27th edition. ACG: A-0870 (AC). Last updated 9/21/2023
- 3. Uptodate Inc. Autism Spectrum Disorder: Evaluation and diagnosis. Last updated May 16, 2022
- 4. Uptodate Inc. Intellectual Disability in Children: Evaluation for a Cause. Last updated January 10, 2023

Review History

September 2024: Updated references





Approved by:

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