



PerkinElmer  
**GENOMICS**



# TANDEM REPEAT ANALYSIS BY WHOLE GENOME SEQUENCING

Historically, detection of repeat expansions has only been possible with polymerase chain reaction-based assays or Southern blots, which are costly and time-consuming methods. With new technology, we can now analyze STRs on NGS based assays, allowing WGS the potential to profile nearly all genetic variants simultaneously in a single assay.

Tandem repeat analysis by Whole Genome Sequencing can be used to rule out select repeat conditions by identifying and characterizing non-expanded (normal) alleles in the proband. The limit of detection of an expanded allele size on NGS based methods is currently not well established. Thus, if an allele is expanded, depending on the size, further confirmatory diagnostic testing may be required to fully characterize the repeat size to provide a definitive diagnosis.

## TANDEM REPEAT ANALYSIS BY WHOLE GENOME SEQUENCING

At PerkinElmer Genomics, we include the analysis of 31 genes for short tandem repeats in our Whole Genome Sequencing (WGS) test, listed below.

Gene	Condition
<i>AFF2</i>	Fragile X syndrome, FRAXE type
<i>AR</i>	Spinal bulbar muscular atrophy
<i>ATN1</i>	Dentatorubral-pallidoluysian atrophy
<i>ATXN1</i>	Spinocerebellar ataxia type 1
<i>ATXN10</i>	Spinocerebellar ataxia type 10
<i>ATXN2</i>	Spinocerebellar ataxia type 2, L-dopa responsive parkinsonism ALS type 13
<i>ATXN3</i>	Spinocerebellar ataxia type 3
<i>ATXN7</i>	Spinocerebellar ataxia type 7
<i>ATXN8OS</i>	Spinocerebellar ataxia type 8
<i>C9orf72</i>	Frontotemporal dementia, ALS type 1, hereditary ataxia
<i>CACNA1A</i>	Spinocerebellar ataxia type 6
<i>CNBP</i>	Myotonic dystrophy type 2
<i>COMP</i>	Pseudo-achondroplasia
<i>CSTB</i>	Unverricht-Lundborg disease
<i>DIP2B</i>	Mental retardation type 12A
<i>DMPK</i>	Myotonic dystrophy type 1

Gene	Condition
<i>FMR1</i>	Fragile X syndrome, fragile X-associated tremor/ataxia syndrome
<i>FOXL2</i>	Blepharophimosis, ptosis, epicanthus inversus
<i>FXN</i>	Friedreich ataxia
<i>HOXD13</i>	Syndactyly, type V
<i>NOP56</i>	Spinocerebellar ataxia type 36
<i>PABPN1</i>	Oculopharyngeal muscular dystrophy
<i>PHOX2B</i>	Congenital central hypoventilation syndrome
<i>PPP2R2B</i>	Spinocerebellar ataxia type 12
<i>RUNX2</i>	Cleidocranial dysplasia
<i>SOX3</i>	Panhypopituitarism and intellectual disability with growth hormone deficiency
<i>TBP</i>	Spinocerebellar ataxia type 17
<i>TBX1</i>	Tetralogy of Fallot
<i>TCF4</i>	Fuchs endothelial corneal dystrophy
<i>ZIC2</i>	Holoprosencephaly type 5
<i>ZIC3</i>	VACTERL

## HOW TO ORDER

Please reach out to your nearest PerkinElmer Genomics' representatives to order for any queries, feedback, or information you may require. You may also contact us on:

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