

## PMP22 Gene Deletion/Duplication Analysis in Hereditary Neuropathy with Liability to Pressure Palsy (HNPP) and Charcot-Marie-Tooth 1A (CMT1A)

### Clinical Features:

**Hereditary neuropathy with liability to pressure palsy (HNPP):** is characterized by recurrent episodes of sensory and motor neuropathy in a single nerve. While any nerve in the peripheral nervous system can be affected, the ulnar, peroneal, median, brachial plexus and radial nerves are most commonly affected<sup>1</sup>. Almost all affected individuals show prolongation of distal nerve conduction latencies. Other features include: reduced or absent tendon reflexes, pes cavus, episodic foot drop, atrophy and weakness of the hands, carpal tunnel syndrome, and pain, while less common features include: motor brachial paralysis, proximal muscle atrophy, respiratory insufficiency, white matter lesions on brain MRI, hypoglossal nerve paralysis of the tongue, and scapulo-peroneal syndrome. An episode can last from minutes to months. Individuals typically present in the 2<sup>nd</sup> or 3<sup>rd</sup> decade, although the age of onset can range from neonatal period into the 7<sup>th</sup> decade.<sup>1</sup>

**Charcot-Marie-Tooth type 1A (CMT1A):** is a progressive disorder characterized by slow nerve conduction velocity (less than 38m/s), distal muscle weakness and atrophy, depressed deep tendon reflexes, sensory loss, pes cavus, hammertoes, and bilateral foot drop. Hearing loss and hip dysplasia may also be present<sup>2,5</sup>. Approximately 85% of individuals with CMT1A present with initial symptoms before age 20.<sup>3</sup>

### Genetics:

Autosomal dominant with variable expressivity and reduced penetrance; approximately 80% of individuals with *PMP22* deletions inherited the deletion from a parent, while approximately 66% of individuals with *PMP22* duplications inherited the duplication from a parent<sup>1,2</sup>. HNPP is most commonly caused by a 1.5 Mb deletion on the short arm of chromosome 17, which includes the *PMP22* gene. While approximately 80% of individuals with HNPP have this recurrent deletion, the remaining 20% have point variants in the *PMP22* gene<sup>1</sup>. The reciprocal duplication of *PMP22* is the most common cause of Charcot-Marie-Tooth disease. Approximately 70% of CMT1 is caused by the recurrent *PMP22* duplication<sup>2</sup>. Other causes of demyelinating CMT include: point variants in *PMP22*, and pathogenic variants in *MPZ*, *LITAF*, *EGR2*, and *NEFL*. Sequence and deletion/duplication analysis of the 53 genes associated with inherited neuropathy, including *PMP22*, is available at GeneDx (see the Hereditary Neuropathy Panel).

## Test Methods:

Targeted array CGH analysis with exon-level resolution is performed to evaluate for a deletion or duplication of one or more exons of the gene. The presence of any potentially disease-associated copy number alteration(s) is confirmed by quantitative PCR or another appropriate method. Sequencing and deletion/duplication analysis of the remaining genes on the Hereditary Neuropathy Panel is available as a separate test if this test negative.

## Test Sensitivity:

Exon-level array CGH will detect partial and whole gene deletions and duplications of the *PMP22* gene. Approximately 80% of individuals with a HNPP will have a deletion of *PMP22*, while approximately 70% of individuals with CMT1 will have a duplication of the *PMP22* gene<sup>1,4</sup>.

## References:

1. Bird (Updated September 2014). Hereditary Neuropathy with Liability to Pressure Palsies. In: GeneReviews at GeneTests: Medical Genetics Information Resource (database online). Copyright, University of Washington, Seattle. 1997-2010. Available at <http://www.genetests.org>.
2. Bird (Updated May 2015). Charcot-Marie-Tooth Hereditary Neuropathy Overview. In: GeneReviews at GeneTests: Medical Genetics Information Resource (database online). Copyright, University of Washington, Seattle. 1997-2010. Available at <http://www.genetests.org>.
3. Li, J (2012) Semin Neurol. 32(3): 204-214.
4. Nelis et al. (2006) Eur J Hum Genet 4(1): 25-33.
5. Siskind et al. (2013) J Genet Counsel 22:422-436.