PAINLESS PERIPHERAL NEUROPATHIES GENE PANEL DG 3.5.0 (2 genes)

Releasedate: 05-12-2022

| Gene | TWIST X2 | TWIST X2 | Associated Phenotype description and OMIM disease ID |
|--------|--------------|--------------|--|
| | covered >10x | covered >20x | |
| SCN11A | 100% | 100% | Episodic pain syndrome, familial, 3, 615552 |
| | | | Neuropathy, hereditary sensory and autonomic, type VII, 615548 |
| SCN9A | 100% | 100% | Erythermalgia, primary, 133020 |
| | | | Insensitivity to pain, congenital, 243000 |
| | | | Small fiber neuropathy, 133020 |
| | | | Paroxysmal extreme pain disorder, 167400 |
| | | | Neuropathy, hereditary sensory and autonomic, type IID, 243000 |

Gene symbols used follow HGCN guidelines: Gray KA, Yates B, Seal RL, Wright MW, Bruford EA. Nucleic Acids Res. 2015 Jan 43(Database issue):D1079-85. Gene symbols used follow HGCN guidelines: Gray KA, Yates B, Seal RL, Wright MW, Bruford EA. Nucleic Acids Res. 2015 Jan 43(Database issue):D1079-85. TWIST X2 is the chemistry used for WES analysis.

Covered 10x describes the percentage of a gene's coding sequence that is covered at least 10x.

Covered 20x describes the percentage of a gene's coding sequence that is covered at least 20x.

Genes with coverage denoting NC are non-protein coding genes.

non-protein coding genes are covered, but as coverage statistics are based on protein coding regions, statistics could not be generated.

OMIM release used for OMIM disease identifiers and descriptions: November 28th, 2022.

This list is accurate for panel version DG 3.5.0

Ad 1. "No OMIM Disease ID" signifies a gene without a current OMIM association Ad 2. OMIM phenotype descriptions between {} signify risk factors