

PARKINSON DISEASE PANEL DG-4.0.0 (36 GENES)

<i>Gene</i>	<i>Twist X2 covered >10x</i>	<i>Twist X2 covered >20x</i>	<i>WGS covered >10x</i>	<i>WGS covered >20x</i>	<i>Associated Phenotype description and OMIM disease ID</i>
ATP13A2	100.0%	100.0%	100.0%	99.3%	Spastic paraplegia 78, autosomal recessive, 617225;Kufor-Rakeb syndrome, 606693
ATP1A3	100.0%	100.0%	100.0%	98.9%	Alternating hemiplegia of childhood 2, 614820;Dystonia-12, 128235;CAPOS syndrome, 601338;Developmental and epileptic encephalopathy 99, 619606
C19orf12	100.0%	99.8%	100.0%	98.4%	Neurodegeneration with brain iron accumulation 4, 614298;?Spastic paraplegia 43, autosomal recessive, 615043
CHCHD2	100.0%	100.0%	100.0%	99.6%	Parkinson disease 22, autosomal dominant, 616710
CHMP2B	89.9%	88.0%	100.0%	96.3%	Frontotemporal dementia and/or amyotrophic lateral sclerosis 7, 600795

CSF1R	100.0%	100.0%	100.0%	99.3%	Brain abnormalities, neurodegeneration, and dysosteosclerosis, 618476;Leukoencephalopathy, diffuse hereditary, with spheroids 1, 221820
DCTN1	100.0%	100.0%	100.0%	99.5%	Perry syndrome, 168605;{Amyotrophic lateral sclerosis, susceptibility to}, 105400;Neuronopathy, distal hereditary motor, autosomal dominant 14, 607641
DNAJC6	100.0%	100.0%	100.0%	98.6%	Parkinson disease 19a, juvenile-onset, 615528;Parkinson disease 19b, early-onset, 615528
FBXO7	100.0%	100.0%	100.0%	98.6%	Parkinson disease 15, autosomal recessive, 260300
FTL	100.0%	100.0%	100.0%	96.5%	Hyperferritinemia-cataract syndrome, 600886;L-ferritin deficiency, dominant and recessive, 615604;Neurodegeneration with brain iron accumulation 3, 606159

GBA1	100.0%	100.0%	100.0%	99.5%	{Lewy body dementia, susceptibility to}, 127750;Gaucher disease, type II, 230900;Gaucher disease, type IIIC, 231005;Gaucher disease, type III, 231000;Gaucher disease, type I, 230800;Gaucher disease, perinatal lethal, 608013;{Parkinson disease, late-onset, susceptibility to}, 168600
GCH1	100.0%	100.0%	99.9%	98.2%	Dystonia, DOPA-responsive, 128230;Hyperphenylalaninemia, BH4-deficient, B, 233910
GRN	100.0%	100.0%	100.0%	99.7%	Aphasia, primary progressive, 607485;Frontotemporal lobar degeneration with ubiquitin-positive inclusions, 607485;Ceroid lipofuscinosis, neuronal, 11, 614706
LRRK2	100.0%	100.0%	100.0%	97.8%	{Parkinson disease 8}, 607060

MAPT	95.4%	95.4%	100.0%	98.0%	Supranuclear palsy, progressive, 601104;Supranuclear palsy, progressive atypical, 260540;Dementia, frontotemporal, with or without parkinsonism, 600274;{Parkinson disease, susceptibility to}, 168600;Pick disease, 172700
MYORG	100.0%	100.0%	100.0%	100.0%	Basal ganglia calcification, idiopathic, 7, autosomal recessive, 618317
PARK7	100.0%	100.0%	100.0%	98.9%	Parkinson disease 7, autosomal recessive early-onset, 606324
PDGFB	100.0%	100.0%	99.7%	96.7%	Meningioma, SIS-related, 607174;Basal ganglia calcification, idiopathic, 5, 615483;Dermatofibrosarcoma protuberans, 607907
PDGFRB	100.0%	100.0%	100.0%	99.2%	Premature aging syndrome, Penttinen type, 601812;Kosaki overgrowth syndrome, 616592;Myofibromatosis, infantile, 1, 228550;Basal ganglia calcification, idiopathic, 4, 615007;Myeloproliferative disorder with eosinophilia, 131440

PINK1	100.0%	100.0%	100.0%	98.1%	Parkinson disease 6, early onset, 605909
PLA2G6	100.0%	99.9%	100.0%	99.2%	Parkinson disease 14, autosomal recessive, 612953;Neurodegeneration with brain iron accumulation 2B, 610217;Infantile neuroaxonal dystrophy 1, 256600
POLG	100.0%	100.0%	100.0%	99.4%	Mitochondrial recessive ataxia syndrome (includes SANDO and SCAE), 607459;Mitochondrial DNA depletion syndrome 4B (MNGIE type), 613662;Mitochondrial DNA depletion syndrome 4A (Alpers type), 203700;Progressive external ophthalmoplegia, autosomal dominant 1, 157640;Progressive external ophthalmoplegia, autosomal recessive 1, 258450
PRKN	100.0%	100.0%	100.0%	99.1%	Adenocarcinoma of lung, somatic, 211980;Parkinson disease, juvenile, type 2, 600116;Ovarian cancer, somatic, 167000
PRKRA	100.0%	100.0%	99.9%	97.3%	Dystonia 16, 612067

PSEN1	100.0%	100.0%	100.0%	99.2%	Pick disease, 172700;Alzheimer disease, type 3, with spastic paraparesis and apraxia, 607822;Dementia, frontotemporal, 600274;?Acne inversa, familial, 3, 613737;Cardiomyopathy, dilated, 1U, 613694;Alzheimer disease, type 3, with spastic paraparesis and unusual plaques, 607822;Alzheimer disease, type 3, 607822
SLC20A2	100.0%	100.0%	100.0%	99.0%	Basal ganglia calcification, idiopathic, 1, 213600
SLC30A10	100.0%	100.0%	100.0%	98.5%	Hypermanganesemia with dystonia 1, 613280
SLC39A14	93.6%	93.6%	100.0%	99.3%	?Hyperostosis cranialis interna, 144755;Hypermanganesemia with dystonia 2, 617013
SLC6A3	100.0%	100.0%	100.0%	99.5%	Parkinsonism-dystonia, infantile, 1, 613135;{Nicotine dependence, protection against}, 188890
SNCA	100.0%	100.0%	100.0%	98.6%	Dementia, Lewy body, 127750;Parkinson disease 1, 168601;Parkinson disease 4, 605543

TAF1	98.7%	98.6%	97.5%	69.2%	Intellectual developmental disorder, X-linked syndromic 33, 300966;Dystonia-Parkinsonism, X-linked, 314250
TH	100.0%	100.0%	100.0%	98.4%	Segawa syndrome, recessive, 605407
VPS13C	100.0%	100.0%	100.0%	98.3%	Parkinson disease 23, autosomal recessive, early onset, 616840
VPS35	100.0%	100.0%	100.0%	98.1%	{Parkinson disease 17}, 614203
WDR45	100.0%	100.0%	98.9%	76.4%	Neurodegeneration with brain iron accumulation 5, 300894
XPR1	100.0%	100.0%	100.0%	98.4%	Basal ganglia calcification, idiopathic, 6, 616413

Gene symbols used follow HGNC guidelines: Gray KA, Yates B, Seal RL, Wright MW, Bruford EA. *Nucleic Acids Res.* 2015 Jan 43(Database issue):D1079-85.

TWIST X2 Covered 10x describes the percentage of a gene's coding sequence that is covered at least 10x when analyzed by WES using TWIST X2 chemistry.

TWIST X2 Covered 20x describes the percentage of a gene's coding sequence that is covered at least 20x when analyzed by WES using TWIST X2 chemistry.

srWGS GRCh38 Covered 10x describes the percentage of a gene's coding sequence that is covered at least 10x when analyzed by WGS mapped against GRCh38.

srWGS GRCh38 Covered 20x describes the percentage of a gene's coding sequence that is covered at least 20x when analyzed by WGS mapped against GRCh38.

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 : March 17th, 2023.

This list is accurate for panel version DG 4.0.0