

# WES HEREDITARY CANCER DG 3.7

<i>Gene</i>	<i>Twist X2 covered &gt;10x</i>	<i>Twist X2 covered &gt;20x</i>	<i>WGS covered &gt;10x</i>	<i>WGS covered &gt;20x</i>	<i>Associated Phenotype description and OMIM disease ID</i>
A2ML1	100.0%	100.0%	100.0%	99.6%	
ACD	100.0%	100.0%	100.0%	99.3%	?Dyskeratosis congenita, autosomal recessive 7, 616553 ?Dyskeratosis congenita, autosomal dominant 6, 616553
AIP	100.0%	100.0%	100.0%	100.0%	Pituitary adenoma 1, multiple types, 102200 Pituitary adenoma predisposition, 102200
AKT1	100.0%	100.0%	100.0%	100.0%	Breast cancer, somatic, 114480 Cowden syndrome 6, 615109 Colorectal cancer, somatic, 114500 Proteus syndrome, somatic, 176920 Ovarian cancer, somatic, 167000
ALK	100.0%	99.9%	99.9%	98.7%	
AMH	100.0%	100.0%	100.0%	100.0%	Persistent Mullerian duct syndrome, type I, 261550
AMHR2	100.0%	100.0%	100.0%	99.7%	Persistent Mullerian duct syndrome, type II, 261550
ANKRD26	97.2%	97.2%	100.0%	98.6%	Thrombocytopenia 2, 188000

APC	100.0%	100.0%	100.0%	99.2%	Colorectal cancer, somatic, 114500 Brain tumor-polyposis syndrome 2, 175100 Desmoid disease, hereditary, 135290 Adenoma, periampullary, somatic, 175100 Hepatoblastoma, somatic, 114550 Gastric cancer, somatic, 613659 Gastric adenocarcinoma and proximal polyposis of the stomach, 619182 Gardner syndrome, 175100 Adenomatous polyposis coli, 175100
ARMC5	100.0%	100.0%	100.0%	99.9%	ACTH-independent macronodular adrenal hyperplasia 2, 615954
ASXL1	100.0%	100.0%	100.0%	99.6%	Myelodysplastic syndrome, somatic, 614286 Bohring-Opitz syndrome, 605039
ATM	100.0%	100.0%	100.0%	99.0%	Ataxia-telangiectasia, 208900 Lymphoma, B-cell non-Hodgkin, somatic, T-cell prolymphocytic leukemia, somatic, Lymphoma, mantle cell, somatic,
ATR	100.0%	100.0%	100.0%	99.2%	Seckel syndrome 1, 210600 ?Cutaneous telangiectasia and cancer syndrome, familial, 614564
AXIN2	100.0%	100.0%	100.0%	99.7%	Colorectal cancer, somatic, 114500 Oligodontia-colorectal cancer syndrome, 608615
BAP1	100.0%	100.0%	100.0%	99.8%	Kury-Isidor syndrome, 619762 Tumor predisposition syndrome 1, 614327
BARD1	100.0%	100.0%	100.0%	99.4%	
BLM	100.0%	100.0%	100.0%	99.2%	Bloom syndrome, 210900

BMPR1A	100.0%	100.0%	100.0%	99.4%	Polyposis syndrome, hereditary mixed, 2, 610069 Polyposis, juvenile intestinal, 174900
BRAF	100.0%	100.0%	100.0%	99.5%	Melanoma, malignant, somatic, 155600 LEOPARD syndrome 3, 613707 Cardiofaciocutaneous syndrome, 115150 Adenocarcinoma of lung, somatic, 211980 Noonan syndrome 7, 613706 Colorectal cancer, somatic, 114500 Nonsmall cell lung cancer, somatic, 211980
BRCA1	100.0%	100.0%	100.0%	99.1%	Fanconi anemia, complementation group S, 617883
BRCA2	100.0%	100.0%	100.0%	98.6%	Fanconi anemia, complementation group D1, 605724 Wilms tumor, 194070
BRIP1	100.0%	100.0%	100.0%	99.1%	Fanconi anemia, complementation group J, 609054
BUB1	100.0%	100.0%	100.0%	99.4%	Colorectal cancer with chromosomal instability, somatic, 114500 Microcephaly 30, primary, autosomal recessive, 620183
BUB1B	100.0%	100.0%	100.0%	99.5%	Colorectal cancer, somatic, 114500 Mosaic variegated aneuploidy syndrome 1, 257300
BUB3	100.0%	100.0%	100.0%	99.5%	
CARD11	100.0%	100.0%	100.0%	99.7%	B-cell expansion with NFKB and T-cell anergy, 616452 Immunodeficiency 11B with atopic dermatitis, 617638 Immunodeficiency 11A, 615206

CBL	100.0%	100.0%	100.0%	99.7%	Noonan syndrome-like disorder with or without juvenile myelomonocytic leukemia, 613563 ?Juvenile myelomonocytic leukemia, 607785
CD27	100.0%	100.0%	100.0%	99.7%	Lymphoproliferative syndrome 2, 615122
CD70	100.0%	100.0%	100.0%	99.2%	Lymphoproliferative syndrome 3, 618261
CDC73	100.0%	100.0%	100.0%	99.7%	Hyperparathyroidism, familial primary, 145000 Parathyroid adenoma with cystic changes, 145001 Parathyroid carcinoma, 608266 Hyperparathyroidism-jaw tumor syndrome, 145001
CDH1	98.7%	98.7%	100.0%	99.5%	Ovarian cancer, somatic, 167000 Blepharocheilodontic syndrome 1, 119580 Diffuse gastric and lobular breast cancer syndrome with or without cleft lip and/or palate, 137215 Endometrial carcinoma, somatic, 608089 Breast cancer, lobular, somatic, 114480
CDH23	100.0%	100.0%	100.0%	99.8%	Usher syndrome, type 1D, 601067 Usher syndrome, type 1D/F digenic, 601067 Deafness, autosomal recessive 12, 601386
CDK4	100.0%	100.0%	100.0%	99.9%	
CDKN1A	100.0%	100.0%	100.0%	99.9%	
CDKN1B	100.0%	100.0%	100.0%	99.4%	Multiple endocrine neoplasia, type IV, 610755
CDKN1C	100.0%	100.0%	100.0%	99.8%	IMAGE syndrome, 614732 Beckwith-Wiedemann syndrome, 130650
CDKN2A	100.0%	100.0%	100.0%	99.3%	
CDKN2B	100.0%	100.0%	100.0%	99.9%	

CDKN2C	100.0%	100.0%	100.0%	99.2%	
CEBPA	100.0%	100.0%	99.9%	92.7%	Leukemia, acute myeloid, somatic, 601626 ?Leukemia, acute myeloid, 601626
CHEK2	100.0%	100.0%	100.0%	98.9%	Osteosarcoma, somatic, 259500 Li-Fraumeni syndrome 2, 609265
CREBBP	100.0%	100.0%	100.0%	99.2%	Menke-Hennekam syndrome 1, 618332 Rubinstein-Taybi syndrome 1, 180849
CTC1	100.0%	100.0%	100.0%	99.6%	Cerebroretinal microangiopathy with calcifications and cysts, 612199
CTLA4	100.0%	100.0%	100.0%	99.2%	Immune dysregulation with autoimmunity, immunodeficiency, and lymphoproliferation, 616100
CTNNA1	100.0%	100.0%	100.0%	99.4%	Macular dystrophy, patterned, 2, 608970
CTR9	100.0%	100.0%	100.0%	99.2%	
CYLD	100.0%	100.0%	100.0%	99.2%	Brooke-Spiegler syndrome, 605041 Cylindromatosis, familial, 132700 Trichoepithelioma, multiple familial, 1, 601606 ?Frontotemporal dementia and/or amyotrophic lateral sclerosis 8, 619132
DCLRE1B	100.0%	100.0%	100.0%	99.8%	Dyskeratosis congenita, autosomal recessive 8, 620133
DDB2	100.0%	100.0%	100.0%	99.1%	Xeroderma pigmentosum, group E, DDB-negative subtype, 278740
DDX11	100.0%	100.0%	100.0%	99.9%	Warsaw breakage syndrome, 613398
DDX41	100.0%	100.0%	100.0%	99.8%	
DGCR8	100.0%	100.0%	100.0%	99.6%	

DICER1	100.0%	100.0%	100.0%	99.2%	Pleuropulmonary blastoma, 601200 Goiter, multinodular 1, with or without Sertoli-Leydig cell tumors, 138800 GLOW syndrome, somatic mosaic, 618272 Rhabdomyosarcoma, embryonal, 2, 180295
DIS3L2	100.0%	100.0%	100.0%	99.4%	Perlman syndrome, 267000
DKC1	100.0%	100.0%	98.0%	73.8%	?Cataracts, hearing impairment, nephrotic syndrome, and enterocolitis 1, 301108 Dyskeratosis congenita, X-linked, 305000
DLST	100.0%	100.0%	100.0%	99.8%	Paragangliomas 7, 618475
DNAJC21	100.0%	100.0%	100.0%	98.5%	Bone marrow failure syndrome 3, 617052
EGFR	100.0%	100.0%	100.0%	99.6%	?Inflammatory skin and bowel disease, neonatal, 2, 616069 Nonsmall cell lung cancer, response to tyrosine kinase inhibitor in, 211980 Adenocarcinoma of lung, response to tyrosine kinase inhibitor in, 211980
EGLN1	100.0%	100.0%	100.0%	98.1%	Erythrocytosis, familial, 3, 609820
EGLN2	100.0%	100.0%	100.0%	100.0%	
ELANE	100.0%	100.0%	100.0%	99.9%	Neutropenia, cyclic, 162800 Neutropenia, severe congenital 1, autosomal dominant, 202700
ELP1	100.0%	100.0%	100.0%	99.6%	Dysautonomia, familial, 223900
EPAS1	100.0%	100.0%	99.9%	98.7%	Erythrocytosis, familial, 4, 611783
EPCAM	100.0%	100.0%	100.0%	99.4%	Diarrhea 5, with tufting enteropathy, congenital, 613217 Lynch syndrome 8, 613244
ERCC1	100.0%	100.0%	100.0%	99.5%	Cerebrooculofacioskeletal syndrome 4, 610758

ERCC2	100.0%	100.0%	100.0%	99.7%	Xeroderma pigmentosum, group D, 278730 Trichothiodystrophy 1, photosensitive, 601675 ?Cerebrooculofacioskeletal syndrome 2, 610756
ERCC3	100.0%	100.0%	100.0%	99.5%	Trichothiodystrophy 2, photosensitive, 616390 Xeroderma pigmentosum, group B, 610651
ERCC4	100.0%	100.0%	100.0%	98.8%	Xeroderma pigmentosum, type F/Cockayne syndrome, 278760 XFE progeroid syndrome, 610965 Xeroderma pigmentosum, group F, 278760 Fanconi anemia, complementation group Q, 615272
ERCC5	100.0%	100.0%	100.0%	99.3%	Xeroderma pigmentosum, group G, 278780 Cerebrooculofacioskeletal syndrome 3, 616570 Xeroderma pigmentosum, group G/Cockayne syndrome, 278780
ERCC6	100.0%	100.0%	100.0%	99.4%	UV-sensitive syndrome 1, 600630 Cerebrooculofacioskeletal syndrome 1, 214150 ?De Sanctis-Cacchione syndrome, 278800 Cockayne syndrome, type B, 133540 Premature ovarian failure 11, 616946
ESR2	100.0%	100.0%	100.0%	99.7%	?Ovarian dysgenesis 8, 618187
ETV6	100.0%	100.0%	100.0%	99.4%	Thrombocytopenia 5, 616216 Leukemia, acute myeloid, somatic, 601626
EXT1	100.0%	100.0%	100.0%	99.3%	Exostoses, multiple, type 1, 133700 Chondrosarcoma, 215300

EXT2	100.0%	100.0%	100.0%	99.6%	Seizures, scoliosis, and macrocephaly syndrome, 616682 Exostoses, multiple, type 2, 133701
EZH2	100.0%	100.0%	100.0%	99.6%	Weaver syndrome, 277590
FANCA	100.0%	100.0%	100.0%	99.5%	Fanconi anemia, complementation group A, 227650
FANCB	100.0%	100.0%	98.0%	71.0%	Fanconi anemia, complementation group B, 300514
FANCC	100.0%	100.0%	100.0%	99.6%	Fanconi anemia, complementation group C, 227645
FANCD2	100.0%	100.0%	100.0%	99.1%	Fanconi anemia, complementation group D2, 227646
FANCE	100.0%	100.0%	100.0%	99.6%	Fanconi anemia, complementation group E, 600901
FANCF	100.0%	100.0%	100.0%	99.3%	Fanconi anemia, complementation group F, 603467
FANCG	100.0%	100.0%	100.0%	98.4%	Fanconi anemia, complementation group G, 614082
FANCI	100.0%	100.0%	100.0%	99.2%	Fanconi anemia, complementation group I, 609053
FANCL	100.0%	100.0%	100.0%	99.3%	Fanconi anemia, complementation group L, 614083
FANCM	100.0%	100.0%	100.0%	98.7%	?Premature ovarian failure 15, 618096 Spermatogenic failure 28, 618086
FAS	100.0%	100.0%	100.0%	98.9%	Autoimmune lymphoproliferative syndrome, type IA, 601859 Squamous cell carcinoma, burn scar-related, somatic,

FASLG	100.0%	100.0%	100.0%	99.4%	Autoimmune lymphoproliferative syndrome, type IB, 601859
FBXW7	99.5%	98.2%	100.0%	99.4%	Developmental delay, hypotonia, and impaired language, 620012
FH	100.0%	100.0%	100.0%	99.2%	Leiomyomatosis and renal cell cancer, 150800 Fumarase deficiency, 606812
FLCN	100.0%	100.0%	100.0%	99.5%	Birt-Hogg-Dube syndrome, 135150 Colorectal cancer, somatic, 114500 Pneumothorax, primary spontaneous, 173600 Renal carcinoma, chromophobe, somatic, 144700
G6PC3	100.0%	100.0%	100.0%	99.8%	Dursun syndrome, 612541 Neutropenia, severe congenital 4, autosomal recessive, 612541
GALNT12	100.0%	100.0%	100.0%	99.4%	
GATA2	100.0%	100.0%	100.0%	99.8%	Emberger syndrome, 614038 Immunodeficiency 21, 614172
GDNF	100.0%	100.0%	100.0%	99.6%	
GFI1	100.0%	100.0%	100.0%	99.7%	?Neutropenia, nonimmune chronic idiopathic, of adults, 607847 Neutropenia, severe congenital 2, autosomal dominant, 613107
GPC3	99.6%	98.9%	98.1%	72.5%	Wilms tumor, somatic, 194070 Simpson-Golabi-Behmel syndrome, type 1, 312870
GPR161	100.0%	100.0%	100.0%	99.4%	
GREM1	100.0%	100.0%	100.0%	99.9%	

GRHL2	100.0%	100.0%	100.0%	99.7%	Deafness, autosomal dominant 28, 608641 Ectodermal dysplasia/short stature syndrome, 616029 Corneal dystrophy, posterior polymorphous, 4, 618031
HAVCR2	100.0%	100.0%	100.0%	99.7%	T-cell lymphoma, subcutaneous panniculitis-like, 618398
HAX1	100.0%	100.0%	100.0%	98.6%	Neutropenia, severe congenital 3, autosomal recessive, 610738
HOXB13	100.0%	100.0%	100.0%	99.7%	
IDH1	100.0%	100.0%	100.0%	99.6%	
IDH2	100.0%	100.0%	100.0%	99.6%	D-2-hydroxyglutaric aciduria 2, 613657
IKZF1	100.0%	100.0%	100.0%	99.7%	Immunodeficiency, common variable, 13, 616873
IPMK	100.0%	100.0%	100.0%	99.0%	
ITK	100.0%	100.0%	100.0%	99.4%	Lymphoproliferative syndrome 1, 613011
KIF1B	100.0%	100.0%	100.0%	99.4%	Charcot-Marie-Tooth disease, type 2A1, 118210
KIT	100.0%	100.0%	100.0%	99.4%	Gastrointestinal stromal tumor, familial, 606764 Mastocytosis, cutaneous, 154800 Piebaldism, 172800 Germ cell tumors, somatic, 273300 Mastocytosis, systemic, somatic, 154800 Leukemia, acute myeloid, somatic, 601626

KRAS	100.0%	100.0%	100.0%	99.8%	Gastric cancer, somatic, 613659 Oculoectodermal syndrome, somatic, 600268 Breast cancer, somatic, 114480 Noonan syndrome 3, 609942 RAS-associated autoimmune leukoproliferative disorder, 614470 Arteriovenous malformation of the brain, somatic, 108010 Lung cancer, somatic, 211980 Pancreatic carcinoma, somatic, 260350 Leukemia, acute myeloid, somatic, 601626 Schimmelpenning-Feuerstein-Mims syndrome, somatic mosaic, 163200 Cardiofaciocutaneous syndrome 2, 615278 Bladder cancer, somatic, 109800
LHCGR	100.0%	100.0%	100.0%	99.6%	Leydig cell adenoma, somatic, with precocious puberty, 176410 Leydig cell hypoplasia with pseudohermaphroditism, 238320 Leydig cell hypoplasia with hypergonadotropic hypogonadism, 238320 Luteinizing hormone resistance, female, 238320 Precocious puberty, male, 176410
LIG4	100.0%	100.0%	100.0%	99.4%	LIG4 syndrome, 606593
LZTR1	100.0%	100.0%	100.0%	99.8%	Noonan syndrome 2, 605275 Noonan syndrome 10, 616564
MAD2L2	100.0%	100.0%	100.0%	99.9%	?Fanconi anemia, complementation group V, 617243

MAP2K1	100.0%	100.0%	100.0%	99.2%	Cardiofaciocutaneous syndrome 3, 615279 Melorheostosis, isolated, somatic mosaic, 155950
MAP2K2	100.0%	100.0%	100.0%	99.8%	Cardiofaciocutaneous syndrome 4, 615280
MAX	100.0%	100.0%	100.0%	99.3%	
MBD4	100.0%	100.0%	100.0%	99.4%	Tumor predisposition syndrome 2, 619975
MCM8	94.4%	94.4%	100.0%	99.6%	?Premature ovarian failure 10, 612885
MCM9	100.0%	100.0%	100.0%	99.1%	Ovarian dysgenesis 4, 616185
MDH2	100.0%	100.0%	100.0%	99.3%	Developmental and epileptic encephalopathy 51, 617339
MEN1	100.0%	100.0%	100.0%	99.5%	Multiple endocrine neoplasia 1, 131100 Lipoma, somatic, Angiofibroma, somatic, Carcinoid tumor of lung, Adrenal adenoma, somatic, Parathyroid adenoma, somatic,
MET	100.0%	100.0%	100.0%	99.5%	Renal cell carcinoma, papillary, 1, familial and somatic, 605074 ?Arthrogryposis, distal, type 11, 620019 Hepatocellular carcinoma, childhood type, somatic, 114550 ?Deafness, autosomal recessive 97, 616705
MITF	99.9%	99.7%	100.0%	99.5%	Waardenburg syndrome, type 2A, 193510 Tietz albinism-deafness syndrome, 103500 COMMAD syndrome, 617306
MLH1	100.0%	100.0%	100.0%	99.1%	Lynch syndrome 2, 609310 Muir-Torre syndrome, 158320 Mismatch repair cancer syndrome 1, 276300

MPL	100.0%	100.0%	100.0%	99.5%	Myelofibrosis with myeloid metaplasia, somatic, 254450 Thrombocythemia 2, 601977 Thrombocytopenia, congenital amegakaryocytic, 604498
MRE11	100.0%	100.0%	100.0%	98.9%	Ataxia-telangiectasia-like disorder 1, 604391
MSH2	100.0%	100.0%	100.0%	98.9%	Lynch syndrome 1, 120435 Muir-Torre syndrome, 158320 Mismatch repair cancer syndrome 2, 619096
MSH3	100.0%	100.0%	100.0%	98.1%	Familial adenomatous polyposis 4, 617100 Endometrial carcinoma, somatic, 608089
MSH6	100.0%	100.0%	100.0%	99.4%	Lynch syndrome 5, 614350 Mismatch repair cancer syndrome 3, 619097
MTAP	100.0%	100.0%	100.0%	99.4%	Diaphyseal medullary stenosis with malignant fibrous histiocytoma, 112250
MUTYH	100.0%	100.0%	100.0%	99.7%	Adenomas, multiple colorectal, 608456 Gastric cancer, somatic, 613659
NBN	100.0%	100.0%	100.0%	98.8%	Leukemia, acute lymphoblastic, 613065 Aplastic anemia, 609135 Nijmegen breakage syndrome, 251260
NF1	100.0%	100.0%	100.0%	99.3%	Watson syndrome, 193520 Leukemia, juvenile myelomonocytic, 607785 Neurofibromatosis, familial spinal, 162210 Neurofibromatosis, type 1, 162200 Neurofibromatosis-Noonan syndrome, 601321

NF2	100.0%	100.0%	100.0%	99.3%	Meningioma, NF2-related, somatic, 607174 Schwannomatosis, vestibular, 101000 Schwannomatosis, somatic, 101000
NHP2	100.0%	100.0%	100.0%	99.3%	Dyskeratosis congenita, autosomal recessive 2, 613987
NOP10	100.0%	100.0%	100.0%	99.8%	?Pulmonary fibrosis and/or bone marrow failure syndrome, telomere-related, 9, 620400 ?Cataracts, hearing impairment, nephrotic syndrome, and enterocolitis 2, 620425 ?Dyskeratosis congenita, autosomal recessive 1, 224230
NPM1	100.0%	100.0%	100.0%	98.0%	Leukemia, acute myeloid, somatic, 601626
NRAS	100.0%	100.0%	100.0%	99.8%	Noonan syndrome 6, 613224 ?RAS-associated autoimmune lymphoproliferative syndrome type IV, somatic, 614470 Melanocytic nevus syndrome, congenital, somatic, 137550 Epidermal nevus, somatic, 162900 Schimmelpenning-Feuerstein-Mims syndrome, somatic mosaic, 163200 Thyroid carcinoma, follicular, somatic, 188470 Neurocutaneous melanosis, somatic, 249400 Colorectal cancer, somatic, 114500
NSD1	100.0%	100.0%	100.0%	99.3%	Sotos syndrome, 117550
NTHL1	100.0%	100.0%	100.0%	99.8%	Familial adenomatous polyposis 3, 616415
PALB2	100.0%	100.0%	100.0%	98.6%	Fanconi anemia, complementation group N, 610832

PARN	97.0%	95.9%	100.0%	99.5%	Dyskeratosis congenita, autosomal recessive 6, 616353 Pulmonary fibrosis and/or bone marrow failure syndrome, telomere-related, 4, 616371
PAX5	100.0%	100.0%	100.0%	99.7%	
PDGFB	100.0%	100.0%	100.0%	98.9%	Meningioma, SIS-related, 607174 Basal ganglia calcification, idiopathic, 5, 615483 Dermatofibrosarcoma protuberans, 607907
PDGFRA	100.0%	100.0%	100.0%	99.3%	Gastrointestinal stromal tumor/GIST-plus syndrome, somatic or familial, 175510 Hypereosinophilic syndrome, idiopathic, resistant to imatinib, 607685
PHOX2B	100.0%	100.0%	100.0%	99.2%	Neuroblastoma with Hirschsprung disease, 613013 Central hypoventilation syndrome, congenital, 1, with or without Hirschsprung disease, 209880

PIK3CA	100.0%	100.0%	100.0%	99.4%	CLOVE syndrome, somatic, 612918 Hepatocellular carcinoma, somatic, 114550 Breast cancer, somatic, 114480 Cerebral cavernous malformations 4, somatic, 619538 Ovarian cancer, somatic, 167000 Colorectal cancer, somatic, 114500 Macrodactyly, somatic, 155500 CLAPO syndrome, somatic, 613089 Keratosis, seborrheic, somatic, 182000 Nevus, epidermal, somatic, 162900 Gastric cancer, somatic, 613659 Nonsmall cell lung cancer, somatic, 211980 Megalencephaly-capillary malformation-polymicrogyria syndrome, somatic, 602501 Cowden syndrome 5, 615108
PMS2	100.0%	100.0%	99.7%	96.6%	Lynch syndrome 4, 614337 Mismatch repair cancer syndrome 4, 619101
PMS2CL	%	%	%	%	
POLD1	100.0%	100.0%	100.0%	99.7%	Mandibular hypoplasia, deafness, progeroid features, and lipodystrophy syndrome, 615381
POLE	100.0%	100.0%	100.0%	99.7%	FILS syndrome, 615139 IMAGE-I syndrome, 618336
POLH	100.0%	100.0%	100.0%	99.7%	Xeroderma pigmentosum, variant type, 278750

POT1	100.0%	100.0%	100.0%	99.5%	?Cerebroretinal microangiopathy with calcifications and cysts 3, 620368 ?Pulmonary fibrosis and/or bone marrow failure syndrome, telomere-related, 8, 620367
POU6F2	100.0%	100.0%	100.0%	99.3%	
PPM1D	100.0%	100.0%	100.0%	99.5%	Breast cancer, somatic, 114480 Jansen-de Vries syndrome, 617450
PRDM10	100.0%	100.0%	100.0%	99.8%	?Birt-Hogg-Dube syndrome 2, 620459
PRF1	100.0%	100.0%	100.0%	100.0%	Hemophagocytic lymphohistiocytosis, familial, 2, 603553 Aplastic anemia, 609135 Lymphoma, non-Hodgkin, 605027
PRKAR1A	100.0%	100.0%	100.0%	99.6%	Pigmented nodular adrenocortical disease, primary, 1, 610489 Acrodysostosis 1, with or without hormone resistance, 101800 Carney complex, type 1, 160980 Myxoma, intracardiac, 255960 Adrenocortical tumor, somatic,
PRKN	91.9%	91.1%	100.0%	99.2%	Adenocarcinoma of lung, somatic, 211980 Parkinson disease, juvenile, type 2, 600116 Ovarian cancer, somatic, 167000
PRSS1	100.0%	100.0%	100.0%	97.8%	Pancreatitis, hereditary, 167800
PTCH1	100.0%	100.0%	100.0%	99.4%	Basal cell nevus syndrome 1, 109400 Basal cell carcinoma, somatic, 605462 Holoprosencephaly 7, 610828

PTEN	100.0%	100.0%	100.0%	99.3%	Cowden syndrome 1, 158350 Lhermitte-Duclos disease, 158350 Prostate cancer, somatic, 176807 Macrocephaly/autism syndrome, 605309
PTPN11	100.0%	100.0%	100.0%	98.9%	Noonan syndrome 1, 163950 LEOPARD syndrome 1, 151100 Metachondromatosis, 156250 Leukemia, juvenile myelomonocytic, somatic, 607785
RAD50	100.0%	100.0%	100.0%	98.7%	Nijmegen breakage syndrome-like disorder, 613078
RAD51C	100.0%	100.0%	100.0%	98.9%	Fanconi anemia, complementation group O, 613390
RAD51D	100.0%	100.0%	100.0%	99.7%	
RAF1	100.0%	100.0%	100.0%	99.7%	Cardiomyopathy, dilated, 1NN, 615916 Noonan syndrome 5, 611553 LEOPARD syndrome 2, 611554
RB1	100.0%	99.7%	100.0%	98.8%	Small cell cancer of the lung, somatic, 182280 Bladder cancer, somatic, 109800 Retinoblastoma, trilateral, 180200 Osteosarcoma, somatic, 259500 Retinoblastoma, 180200
RECQL4	100.0%	100.0%	100.0%	100.0%	Baller-Gerold syndrome, 218600 Rothmund-Thomson syndrome, type 2, 268400 RAPADILINO syndrome, 266280

REST	98.2%	98.2%	100.0%	99.5%	Deafness, autosomal dominant 27, 612431 Fibromatosis, gingival, 5, 617626
RET	100.0%	100.0%	100.0%	99.8%	Multiple endocrine neoplasia IIA, 171400 Medullary thyroid carcinoma, 155240 Pheochromocytoma, 171300 Multiple endocrine neoplasia IIB, 162300
RHBDF2	100.0%	100.0%	100.0%	99.9%	Tylosis with esophageal cancer, 148500
RIT1	100.0%	100.0%	100.0%	99.8%	Noonan syndrome 8, 615355
RMRP	%	%	%	%	Anauxetic dysplasia 1, 607095 Metaphyseal dysplasia without hypotrichosis, 250460 Cartilage-hair hypoplasia, 250250
RNASEL	100.0%	100.0%	100.0%	98.9%	Prostate cancer 1, 601518
RNF43	100.0%	100.0%	100.0%	99.8%	Sessile serrated polyposis cancer syndrome, 617108
RPA1	100.0%	100.0%	100.0%	99.7%	Pulmonary fibrosis and/or bone marrow failure syndrome, telomere-related, 6, 619767
RPL11	100.0%	100.0%	100.0%	99.1%	Diamond-Blackfan anemia 7, 612562
RPL15	99.6%	96.8%	100.0%	99.7%	Diamond-Blackfan anemia 12, 615550
RPL18	100.0%	100.0%	100.0%	99.9%	?Diamond-Blackfan anemia 18, 618310
RPL27	100.0%	100.0%	100.0%	99.3%	?Diamond-Blackfan anemia 16, 617408
RPL35A	100.0%	100.0%	100.0%	99.5%	Diamond-Blackfan anemia 5, 612528
RPL5	100.0%	100.0%	100.0%	99.5%	Diamond-Blackfan anemia 6, 612561

RPS10	100.0%	100.0%	100.0%	99.6%	Diamond-Blackfan anemia 9, 613308
RPS15A	79.7%	79.7%	100.0%	97.6%	?Diamond-Blackfan anemia 20, 618313
RPS17	100.0%	100.0%	100.0%	99.0%	Diamond-Blackfan anemia 4, 612527
RPS19	100.0%	100.0%	100.0%	99.5%	Diamond-Blackfan anemia 1, 105650
RPS20	100.0%	100.0%	100.0%	98.2%	
RPS24	100.0%	100.0%	100.0%	99.3%	Diamond-blackfan anemia 3, 610629
RPS26	100.0%	98.8%	100.0%	99.0%	Diamond-Blackfan anemia 10, 613309
RPS27	100.0%	100.0%	100.0%	99.5%	?Diamond-Blackfan anemia 17, 617409
RPS28	100.0%	100.0%	100.0%	99.7%	Diamond Blackfan anemia 15 with mandibulofacial dysostosis, 606164
RPS29	100.0%	100.0%	100.0%	99.1%	Diamond-Blackfan anemia 13, 615909
RPS7	100.0%	100.0%	100.0%	99.0%	Diamond-Blackfan anemia 8, 612563
RTEL1	100.0%	100.0%	100.0%	99.9%	Dyskeratosis congenita, autosomal dominant 4, 615190 Dyskeratosis congenita, autosomal recessive 5, 615190 Pulmonary fibrosis and/or bone marrow failure syndrome, telomere-related, 3, 616373
RUNX1	100.0%	100.0%	100.0%	99.7%	Platelet disorder, familial, with associated myeloid malignancy, 601399 Leukemia, acute myeloid, 601626

SAMD9	100.0%	100.0%	100.0%	99.0%	Tumoral calcinosis, familial, normophosphatemic, 610455 Monosomy 7 myelodysplasia and leukemia syndrome 2, 619041 MIRAGE syndrome, 617053
SAMD9L	100.0%	100.0%	100.0%	98.9%	Ataxia-pancytopenia syndrome, 159550 Monosomy 7 myelodysplasia and leukemia syndrome 1, 252270 Spinocerebellar ataxia 49, 619806
SBDS	100.0%	100.0%	100.0%	99.1%	Shwachman-Diamond syndrome 1, 260400
SDHA	100.0%	100.0%	100.0%	99.9%	Cardiomyopathy, dilated, 1GG, 613642 Mitochondrial complex II deficiency, nuclear type 1, 252011 Neurodegeneration with ataxia and late-onset optic atrophy, 619259 Paragangliomas 5, 614165
SDHAF2	100.0%	98.3%	100.0%	99.4%	Paragangliomas 2, 601650
SDHB	100.0%	100.0%	100.0%	99.5%	Paragangliomas 4, 115310 Mitochondrial complex II deficiency, nuclear type 4, 619224 Gastrointestinal stromal tumor, 606764 Pheochromocytoma, 171300 Paraganglioma and gastric stromal sarcoma, 606864
SDHC	100.0%	100.0%	100.0%	99.7%	Paragangliomas 3, 605373 Paraganglioma and gastric stromal sarcoma, 606864 Gastrointestinal stromal tumor, 606764

SDHD	78.9%	78.9%	100.0%	99.6%	Paragangliomas 1, with or without deafness, 168000 Paraganglioma and gastric stromal sarcoma, 606864 Mitochondrial complex II deficiency, nuclear type 3, 619167 Pheochromocytoma, 171300
SEMA4A	100.0%	100.0%	100.0%	98.7%	Retinitis pigmentosa 35, 610282 Cone-rod dystrophy 10, 610283
SFTPA1	100.0%	100.0%	100.0%	99.9%	Interstitial lung disease 1, 619611
SFTPA2	100.0%	100.0%	100.0%	99.8%	Interstitial lung disease 2, 178500
SH2B3	100.0%	100.0%	100.0%	99.1%	Thrombocythemia, somatic, 187950 Myelofibrosis, somatic, 254450 Erythrocytosis, somatic, 133100
SH2D1A	100.0%	100.0%	99.8%	79.8%	Lymphoproliferative syndrome, X-linked, 1, 308240
SHOC2	100.0%	100.0%	100.0%	98.7%	Noonan syndrome-like with loose anagen hair 1, 607721
SLC25A11	100.0%	100.0%	100.0%	99.7%	Paragangliomas 6, 618464
SLX4	100.0%	100.0%	100.0%	99.6%	Fanconi anemia, complementation group P, 613951
SMAD4	100.0%	100.0%	100.0%	99.8%	Pancreatic cancer, somatic, 260350 Myhre syndrome, 139210 Polyposis, juvenile intestinal, 174900 Juvenile polyposis/hereditary hemorrhagic telangiectasia syndrome, 175050
SMAD9	100.0%	100.0%	100.0%	99.2%	Pulmonary hypertension, primary, 2, 615342

SMARCA4	100.0%	100.0%	100.0%	99.8%	Coffin-Siris syndrome 4, 614609
SMARCB1	100.0%	100.0%	100.0%	99.9%	Rhabdoid tumors, somatic, 609322 Coffin-Siris syndrome 3, 614608
SMARCE1	100.0%	100.0%	100.0%	99.5%	Coffin-Siris syndrome 5, 616938
SOS1	100.0%	100.0%	100.0%	99.0%	Noonan syndrome 4, 610733 ?Fibromatosis, gingival, 1, 135300
SPINK1	99.9%	99.2%	100.0%	99.4%	Tropical calcific pancreatitis, 608189 Pancreatitis, hereditary, 167800
SPRED1	100.0%	100.0%	100.0%	99.6%	Legius syndrome, 611431
SQSTM1	100.0%	100.0%	100.0%	99.7%	Neurodegeneration with ataxia, dystonia, and gaze palsy, childhood-onset, 617145 Frontotemporal dementia and/or amyotrophic lateral sclerosis 3, 616437 Myopathy, distal, with rimmed vacuoles, 617158 Paget disease of bone 3, 167250
STK11	100.0%	100.0%	100.0%	99.6%	Melanoma, malignant, somatic, 155600 Pancreatic cancer, somatic, 260350 Peutz-Jeghers syndrome, 175200 Testicular tumor, somatic, 273300
SUCLG2	100.0%	99.8%	100.0%	99.1%	
SUFU	100.0%	100.0%	100.0%	99.8%	Joubert syndrome 32, 617757 Basal cell nevus syndrome 2, 620343

TERC	%	%	%	%	Pulmonary fibrosis and/or bone marrow failure syndrome, telomere-related, 2, 614743 Dyskeratosis congenita, autosomal dominant 1, 127550
TERF2IP	99.7%	96.0%	100.0%	99.5%	
TERT	100.0%	100.0%	100.0%	100.0%	Dyskeratosis congenita, autosomal dominant 2, 613989 Dyskeratosis congenita, autosomal recessive 4, 613989 Pulmonary fibrosis and/or bone marrow failure syndrome, telomere-related, 1, 614742
TG	100.0%	100.0%	100.0%	99.6%	Thyroid dysmorphogenesis 3, 274700
THPO	100.0%	100.0%	100.0%	98.3%	Thrombocythemia 1, 187950 Thrombocytopenia 9, 620478 Amegakaryocytic thrombocytopenia, congenital, 2, 620481
TINF2	100.0%	100.0%	100.0%	99.4%	Dyskeratosis congenita, autosomal dominant 3, 613990 Revesz syndrome, 268130
TMEM127	100.0%	100.0%	100.0%	99.6%	
TNFRSF11A	100.0%	99.6%	100.0%	99.4%	Osteopetrosis, autosomal recessive 7, 612301 Osteolysis, familial expansile, 174810
TP53	94.7%	94.7%	100.0%	99.3%	Hepatocellular carcinoma, somatic, 114550 Breast cancer, somatic, 114480 Li-Fraumeni syndrome, 151623 Pancreatic cancer, somatic, 260350 Nasopharyngeal carcinoma, somatic, 607107 Bone marrow failure syndrome 5, 618165

TRIM28	100.0%	100.0%	100.0%	99.9%	
TRIM37	98.3%	98.3%	100.0%	99.4%	Mulibrey nanism, 253250
TRIP13	100.0%	100.0%	100.0%	99.6%	Oocyte/zygote/embryo maturation arrest 9, 619011 Mosaic variegated aneuploidy syndrome 3, 617598
TSC1	100.0%	100.0%	100.0%	99.4%	Focal cortical dysplasia, type II, somatic, 607341 Tuberous sclerosis-1, 191100 Lymphangi leiomyomatosis , 606690
TSC2	100.0%	100.0%	100.0%	99.8%	Lymphangi leiomyomatosis , somatic, 606690 ?Focal cortical dysplasia, type II, somatic, 607341 Tuberous sclerosis-2, 613254
USB1	100.0%	100.0%	100.0%	98.7%	Poikiloderma with neutropenia, 604173
VHL	100.0%	100.0%	100.0%	99.8%	Erythrocytosis, familial, 2, 263400 von Hippel-Lindau syndrome, 193300 Renal cell carcinoma, somatic, 144700 Pheochromocytoma, 171300 Hemangioblastoma, cerebellar, somatic,
WAS	100.0%	98.8%	98.8%	74.3%	Wiskott-Aldrich syndrome, 301000 Neutropenia, severe congenital, X-linked, 300299 Thrombocytopenia, X-linked, intermittent, 313900 Thrombocytopenia, X-linked, 313900
WRAP53	100.0%	100.0%	100.0%	99.6%	Dyskeratosis congenita, autosomal recessive 3, 613988
WRN	100.0%	100.0%	100.0%	98.9%	Werner syndrome, 277700

WT1	100.0%	100.0%	100.0%	99.3%	Mesothelioma, somatic, 156240 Meacham syndrome, 608978 Frasier syndrome, 136680 Nephrotic syndrome, type 4, 256370 Denys-Drash syndrome, 194080 Wilms tumor, type 1, 194070
XPA	100.0%	100.0%	100.0%	99.4%	Xeroderma pigmentosum, group A, 278700
XPC	100.0%	100.0%	100.0%	97.7%	Xeroderma pigmentosum, group C, 278720

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 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 3.7.0.

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