

TEST CODE: PR21037

Overview

MyOme Proactive Health Plus Single-Gene Risk™ report, 151 Genes uses a PCR-free whole genome backbone that allows identification of a range of variant types. Whole genome sequencing (WGS) allows MyOme to re-query a patient's genome as healthcare needs change and new information about the genome is discovered.

Clinical Use

Test is intended for a wellness screening of germline heritable conditions in individuals from an asymptomatic population. MyOme annotates and interprets variants according to American College of Medical Genetics (ACMG) guidelines, and reports pathogenic or likely pathogenic variants. Genetic testing may provide information to assess individual risk, support a clinical diagnosis, and assist with the development of a personalized treatment and management strategy in conjunction with standard clinical assessment.

Method

PCR-free library prep followed by 2x150 bp pairedend whole genome sequencing is the backbone for this test. In-house pipeline allows identification of single-nucleotide variants (SNVs), small insertions and deletions (indels) and copy number variants (CNVs). Variant interpretation by qualified scientists based on guidelines by the ACMG.

Sample Types

- Blood (2 EDTA tubes)
- Saliva (2 tubes)
- Buccal (2 swabs)

Turn Around Time

 From initial sample received, approximately 6 to 8 weeks

Included

- Analysis of SNVs, indels and CNVs (deletions) and duplications)
- Confirmation of Pathogenic/Likely Pathogenic variants by orthogonal technology (e.g. Sanger sequencing)
- Cohesive report with actionable recommendations
- 151 Genes included: ABCD1, ACTA2, ACTC1, ACTN2, ACVRL1, APC, APOB, ATM, ATP7B, BAG3, BAP1, BARD1, BMPR1A, BMPR2, BRCA1, BRCA2, BRIP1, BTD, CACNA1C, CACNA1S, CALM1, CALM2, CALM3, CASQ2, CAV1, CDH1, CDK4, CDKN1B, CDKN2A, CHEK2, COL3A1, COL5A1, COL5A2, CRYAB, CSRP3, DES, DICER1, DMD, DSC2, DSG2, DSP, EGFR, EMD, ENG, EPCAM, F2, F5, F9, FBN1, FH, FHL1, FLCN, FLNC, G6PD, GAA, GCH1, GDF2, GLA, GREM1, HAMP, HFE, HJV, HMBS, HNF1A, HNF1B, HOXB13, JUP, KCNE1, KCNH2, KCNJ2, KCNQ1, KIT, LAMP2, LDLR, LDLRAP1, LMNA, LZTR1, MAX, MEFV, MEN1, MET, MITF, MLH1, MSH2, MSH3, MSH6, MUTYH, MYBPC3, MYH11, MYH7, MYL2, MYL3, MYLK, NF1, NF2, NTHL1, OTC, PALB2, PCSK9, PDGFRA, PKP2, PLN, PMS2, POLD1, POLE, POT1, PRKAG2, PRKG1, PROC, PROS1, PTEN, RAD51C, RAD51D, RB1, RBM20, RET, RPE65, RYR1, RYR2, SCN5A, SDHAF2, SDHB, SDHC, SDHD, SERPINA1, SERPINC1, SLC40A1, SMAD3, SMAD4, SMAD9, STK11, TFR2, TGFB2, TGFB3, TGFBR1, TGFBR2, TMEM127, TMEM43, TNNC1, TNNI3, TNNT2, TP53, TPM1, TRDN, TSC1, TSC2, TTN, TTR, VCL, VHL, WT1

Test Performance¹

- 30x average genome-wide coverage
- >99.5% of exonic regions at ≥10x depth
- >99.5% ClinVar P/LP variants covered by ≥10x depth
- >99% sensitivity for SNVs and indels
- 98% sensitivity for benchmark CNVs >1 kb in size

The test describe above was developed, and its performance characteristics were determined, by MyOme, Inc., a clinical laboratory certified under the Clinical Laboratory Improvement Amendments of 1988 (CLIA) to perform high complexity clinical laboratory testing. This test has not been cleared or approved by the U.S. Food and Drug Administration (FDA)

^{1.} MyOme Inc, Data on File.

CONDITION-GENE RELATIONSHIP

The genes listed below are analyzed in this report. MyOme selected them based on our Gene Inclusion Framework Guidelines. Genes are prioritized based on clinical validity, actionability, penetrance/prevalence, and feasibility.

Cardiovascular	
Condition	Gene(s)
Arrhythmogenic right ventricular cardiomyopathy	DES, DSC2, DSG2, DSP, JUP, PKP2, PLN, TMEM43
Brugada syndrome	SCN5A
CACNA1C-related disorders	CACNA1C
Catecholaminergic polymorphic ventricular tachycardia	CALM1, CALM2, CALM3, CASQ2, RYR2, TRDN
Danon disease	LAMP2
Dilated cardiomyopathy	ACTC1, BAG3, DES, FLNC, LMNA, MYH7, SCN5A, TNNC1, TNNI3, TNNT2, TPM1, TTN, VCL, RBM20
Ehlers-Danlos syndrome, vascular type	COL3A1
Emery-Dreifuss muscular dystrophy	EMD, FHL1, LMNA
Fabry disease	GLA
Familial hypercholesterolemia	APOB, LDLR, LDLRAP1, PCSK9
Familial thoracic aortic aneurysm and dissection	ACTA2, MYH11, MYLK, PRKG1, SMAD3, TGFB2, TGFB3
Hereditary transthyretin-related amyloidosis	TTR
Hypertrophic cardiomyopathy	ACTC1, CSRP3, MYBPC3, MYH7, MYL2, MYL3, PRKAG2, TNNC1, TNNI3, TNNT2, TPM1
Intrinsic Cardiomyopathy	ACTN2, PLN
Loeys-Dietz syndrome	TGFB2, TGFB3, TGFBR1, TGFBR2, SMAD3
Long QT syndrome	CALM1, CALM2, CALM3, KCNH2, KCNQ1, SCN5A, TRDN
Long QT syndrome, acquired	KCNE1
Marfan syndrome	FBN1
Myofibrillar myopathy	BAG3, CRYAB, DES, FLNC
Progressive muscular dystrophy	DMD
Short QT syndrome	KCNH2, KCNJ2, KCNQ1

Other	
Condition	Gene(s)
Acute intermittent porphyria	HMBS
Adrenoleukodystrophy	ABCD1
Alpha-1 antitrypsin defeciency	SERPINA1
Biotinidase deficiency	BTD
Ehlers-Danlos syndrome, classic type	COL5A1, COL5A2
Familial Mediterranean fever	MEFV
G6PD deficiency	G6PD
GTP cyclohydrolase I deficiency	GCH1
Hemophilia B	F9
Hereditary antithrombin deficiency	SERPINC1
Hereditary hemochromatosis	HAMP, HFE, HJV, SLC40A1, TFR2
Hereditary hemorrhagic telangiectasia	ACVRL1, ENG, GDF2, SMAD4
Hereditary thrombophilia due to congenital protein C deficiency	PROC
Hereditary thrombophilia due to congenital protein S deficiency	PROS1

Other (Cont.)	
Condition	Gene(s)
Malignant hyperthermia	CACNA1S, RYR1
Monogenic diabetes	HNF1A, HNF1B
Ornithine transcarbamylase deficiency	ОТС
Pulmonary arterial hypertension	BMPR2, CAV1, GDF2, SMAD9
RPE65-related retinopathy	RPE65
Thrombophilia	F2, F5
Wilson disease	ATP7B

Cancer	
Condition	Gene(s)
BAP1-related tumor predisposition syndrome	BAP1
Birt-Hogg-Dube syndrome	FLCN
CDH1-related diffuse gastric and lobular breast cancer syndrome	CDH1
DICER1-related tumor predisposition	DICER1
Familial adenomatous polyposis	APC, MSH3
Familial ovarian cancer	BRIP1, PALB2, RAD51C, RAD51D
Gastrointestinal stromal tumor	KIT, PDGFRA
Hereditary breast cancer	BARD1, ATM, CHEK2, PALB2
Hereditary breast and ovarian cancer	BRCA1, BRCA2
Hereditary leiomyomatosis and renal cell cancer	FH
Hereditary Mixed Polyposis Syndrome (HMPS)	GREM1
Hereditary nonpolyposis colon cancer	ATM
Hereditary paraganglioma-pheochromocytoma syndrome	MAX, SDHAF2, SDHB, SDHC, SDHD, TMEM127
Juvenile polyposis syndrome	BMPR1A
Juvenile polyposis with hereditary hemorrhagic telangiectasia	SMAD4
Li-Fraumeni syndrome	TP53
Lynch syndrome	EPCAM, MLH1, MSH2, MSH6, PMS2
Melanoma	CDK4 , MITF
Melanoma-pancreatic cancer syndrome	CDKN2A
Multiple endocrine neoplasia	CDKN1B, MEN1, RET
MUTYH-associated polyposis	MUTYH
Neurofibromatosis type 1	NF1
Neurofibromatosis type 2	NF2
Non-small cell lung carcinoma	EGFR
NTHL1-deficiency tumor predisposition syndrome	NTHL1
Papillary renal cell carcinoma	MET
Peutz-Jeghers syndrome	STK11
Polyposis and colorectal cancer	POLD1, POLE
POT1 Tumor predisposition	POT1
Prostate cancer	HOXB13
PTEN hamartoma tumor syndrome	PTEN
Retinoblastoma	RB1
Schwannomatosis	LZTR1
Tuberous sclerosis complex	TSC1, TSC2
Von Hippel-Lindau syndrome	VHL
WT1-related Wilms tumor	WT1