

Full name:	Cancer (Breast, ovary, colon,...) (26 genes) - ULG
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Related Diseases

- [Ataxia-telangiectasia](#)
- [Ataxia-telangiectasia-like disorder](#)
- [Bloom syndrome](#)
- [Constitutional mismatch repair deficiency syndrome](#)
- [Familial pancreatic carcinoma](#)
- [Familial prostate cancer](#)
- [Fanconi anemia](#)
- [Hereditary breast and/or ovarian cancer syndrome](#)
- [Hereditary breast cancer](#)
- [Hereditary diffuse gastric cancer](#)
- [Hereditary site-specific ovarian cancer syndrome](#)
- [Inherited cancer-predisposing syndrome due to biallelic BRCA2 mutations](#)
- [Juvenile polyposis of infancy](#)
- [Li-Fraumeni syndrome](#)
- [Lynch syndrome](#)
- [MUTYH-related attenuated familial adenomatous polyposis](#)
- [Muir-Torre syndrome](#)
- [Multiple endocrine neoplasia type 1](#)
- [Nijmegen breakage syndrome](#)
- [Nijmegen breakage syndrome-like disorder](#)
- [Peutz-Jeghers syndrome](#)

Related Analytes

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
ABRAXAS1			

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>ATM</u>			
<u>BARD1</u>			
<u>BLM</u>			
<u>BRCA1</u>			
<u>BRCA2</u>			
<u>BRIP1</u>			
<u>CDH1</u>			
<u>CHEK2</u>			
<u>EPCAM</u>			
<u>MEN1</u>			
<u>MLH1</u>			
<u>MRE11</u>			
<u>MSH2</u>			
<u>MSH6</u>			
<u>MUTYH</u>			
<u>NBN</u>			
<u>PALB2</u>			
<u>PMS2</u>			
<u>PTEN</u>			
<u>RAD50</u>			

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>RAD51C</u>			
<u>RAD51D</u>			
<u>STK11</u>			
<u>TP53</u>			
<u>XRCC2</u>			