

# Supplementary Materials: Multigene Panel Testing Increases the Number of Loci Associated with Gastric Cancer Predisposition

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**Table S1.** List of the 31 missense variants classified as probably damaging by both SIFT and PolyPhen2 identified in our case series and patients' characteristics.

Gene	Exon	cDNA	Protein	dbSNP	ClinVar	SIFT	Polyphen2 HVAR	Patient ID	Sex	Selection criteria	Cancer(s)	Age at Diagnosis
<i>FH</i>	2	c.234C>A	p.Asn78Lys	-	-	D	D	BM32	F	I	DGC	49
<i>EPCAM</i>	6	c.605A>C	p.Lys202Thr	-	-	D	D	BM67	F	I	DGC	41
<i>PMS1</i>	3	c.224C>T	p.Thr75Ile	rs61756360	-	D	D	BM31	F	I	DGC	47
<i>FANCD2</i>	3	c.195G>C	p.Gln65His	rs36084488	VUS/Benign	D	D	BM71	M	II	DGC	35
<i>GATA2</i>	5	c.1010G>A	p.Arg337Gln	-	-	D	D	BM67	F	I	DGC	41
<i>KIT</i>	21	c.2867G>A	p.Arg956Gln	rs139694927	VUS	D	D	BM85	M	I	IGC	66
<i>PMS2</i>	2	c.52A>G	p.Ile18Val	rs63750123	VUS/Likely benign/Benign	D	D	BM56	F	I	DGC	55
<i>MET</i>	14	c.2975C>T	p.Thr992Ile	rs56391007	VUS/Likely benign/Benign	D	D	BM93	M	I	DGC	65
<i>MET</i>	14	c.2975C>T	p.Thr992Ile	rs56391007	VUS/Likely benign/Benign	D	D	BM50	F	IV	LBC, LBC	53, 58
<i>EZH2</i>	19	c.2151G>C	p.Gln717His	-	-	D	D	BM62	M	I	DGC	45
<i>WRN</i>	14	c.1717A>G	p.Thr573Ala	rs150148567	VUS/Likely benign	D	D	BM58	M	I	DGC	54
<i>WRN</i>	14	c.1717A>G	p.Thr573Ala	rs150148567	VUS/Likely benign	D	D	BM75	F	I	IGC, OC, LBC	57, 60, 72
<i>WRN</i>	17	c.1909C>T	p.Arg637Trp	rs148286708	VUS	D	D	BM58	M	I	DGC	54
<i>WRN</i>	30	c.3523C>A	p.Pro1175Thr	-	-	D	D	BM84	F	I	LBC, DGC	61, 66
<i>NBN</i>	3	c.283G>A	p.Asp95Asn	rs61753720	VUS/Likely benign/Benign	D	D	BM41	F	III	LBC	45
<i>NBN</i>	3	c.283G>A	p.Asp95Asn	rs61753720	VUS/Likely benign/Benign	D	D	BM96	F	II	DGC	33
<i>RET</i>	11	c.1946C>T	p.Ser649Leu	rs148935214	VUS/Likely benign/Benign	D	D	BM69	F	I	GC	77

<i>RET</i>	11	c.1997A>T	p.Lys666Met	rs377767439	VUS	D	D	BM48	F	I	GC	47
<i>BMPRI1A</i>	11	c.1243G>A	p.Glu415Lys	rs140592056	VUS/Likely benign/Benign	D	D	BM79	F	III	LBC	52
<i>EXT2</i>	5	c.889C>T	p.Arg297Cys	rs146098187	-	D	D	BM93	M	I	DGC	65
<i>CEP57</i>	3	c.333G>C	p.Gln111His	rs117321017	Benign	D	D	BM75	F	I	IGC, OC, LBC	57, 60, 72
<i>BRCA2</i>	14	c.7225C>T	p.Pro2409Ser	-	-	D	D	BM72	M	II	DGC	25
<i>ERCC5</i>	14	c.2890C>T	p.Arg964Trp	rs574826021	VUS	D	D	BM61	F	I	DGC	49
<i>FANCM</i>	20	c.4931G>A	p.Arg1644Gln	rs138151018	Likely benign/Benign	D	D	BM61	F	I	DGC	49
<i>FANCI</i>	18	c.1813C>T	p.Leu605Phe	rs117125761	Likely benign/Benign	D	D	BM122	M	I	GC	59
<i>BLM</i>	12	c.2474C>T	p.Pro825Leu	rs749632465	VUS	D	D	BM118	F	I	DGC	54
<i>TSC2</i>	17	c.1747G>A	p.Ala583Thr	rs1800729	Benign/Likely benign	D	D	BM21	F	I	DGC	47
<i>TSC2</i>	17	c.1747G>A	p.Ala583Thr	rs1800729	Benign/Likely benign	D	D	BM63	F	I	DGC, DBC	41, 50
<i>SLX4</i>	6	c.1192C>T	p.Arg398Trp	rs138799572	VUS	D	D	BM114	F	IV	LBC, GC, LBC	36, 38, 52
<i>ERCC4</i>	7	c.1135C>T	p.Pro379Ser	rs1799802	VUS/Likely benign	D	D	BM121	F	I	DGC	40
<i>ERCC4</i>	11	c.2117T>C	p.Ile706Thr	rs1800069	VUS	D	D	BM117	F	II	DGC	36
<i>FLCN</i>	6	c.503G>A	p.Arg168His	-	-	D	D	BM118	F	I	DGC	54
<i>NF1</i>	34	c.4526G>A	p.Arg1509His	rs546073780	VUS	D	D	BM124	M	I	DGC	51
<i>RAD51D</i>	10	c.932T>A	p.Ile311Asn	rs145309168	VUS/Likely benign	D	D	BM122	M	I	GC	59
<i>RHBDF2</i>	5	c.478C>T	p.Arg160Cys	rs751482282	VUS	D	D	BM120	F	II	DGC	36

VUS: variant of uncertain significance; D: damaging; GC: gastric cancer; DGC: diffuse-type gastric cancer; IGC: intestinal-type gastric cancer; LBC: lobular breast cancer; DBC: ductal breast cancer; OC: ovarian cancer. Selection criteria I–IV correspond to those established by the IGCLC for HDGC (see Materials and Methods).



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