

## Supplementary data

**Supplementary Table S1.** List of genes analyzed with FoundationOne® CDx

A. gene list: the entire coding sequence was examined for the detection of base substitutions, insertion/deletions, and copy number alterations.

ABL1	ACVR1B	AKT1	AKT2	AKT3	ALK	ALOX12B	AMER1	APC	AR
ARAF	ARFRP1	ARID1A	ASXL1	ATM	ATR	ATRX	AURKA	AURKB	AXIN1
AXL	BAP1	BARD1	BCL2	BCL2L1	BCL2L2	BCL6	BCOR	BCORL1	BRAF
BRCA1	BRCA2	BRD4	BRIP1	BTG1	BTG2	BTK	C11orf30	CALR	CARD11
CASP8	CBFB	CBL	CCND1	CCND2	CCND3	CCNE1	CD22	CD274	CD70
CD79A	CD79B	CDC73	CDH1	CDK12	CDK4	CDK6	CDK8	CDKN1A	CDKN1B
CDKN2A	CDKN2B	CDKN2C	CEBPA	CHEK1	CHEK2	CIC	CREBBP	CRKL	CSF1R
CSF3R	CTCF	CTNNA1	CTNNB1	CUL3	CUL4A	CXCR4	CYP17A1	DAXX	DDR1
DDR2	DIS3	DNMT3A	DOT1L	EED	EGFR	EP300	EPHA3	EPHB1	EPHB4
ERBB2	ERBB3	ERBB4	ERCC4	ERG	ERRFI1	ESR1	EZH2	FAM46C	FANCA
FANCC	FANCG	FANCL	FAS	FBXW7	FGF10	FGF12	FGF14	FGF19	FGF23
FGF3	FGF4	FGF6	FGFR1	FGFR2	FGFR3	FGFR4	FH	FLCN	FLT1
FLT3	FOXL2	FUBP1	GABRA6	GATA3	GATA4	GATA6	GID4 (C17orf39)	GNA11	GNA13
GNAQ	GNAS	GRM3	GSK3B	H3F3A	HDAC1	HGF	HNF1A	HRAS	HSD3B1
ID3	IDH1	IDH2	IGF1R	IKBKE	IKZF1	INPP4B	IRF2	IRF4	IRS2
JAK1	JAK2	JAK3	JUN	KDM5A	KDM5C	KDM6A	KDR	KEAP1	KEL
KIT	KLHL6	KMT2A (MLL)	KMT2D (MLL2)	KRAS	LTK	LYN	MAF	MAP2K1	MAP2K2
MAP2K4	MAP3K1	MAP3K13	MAPK1	MCL1	MDM2	MDM4	MED12	MEF2B	MEN1
MERTK	MET	MITF	MKNK1	MLH1	MPL	MRE11A	MSH2	MSH3	MSH6
MST1R	MTAP	MTOR	MUTYH	MYC	MYCL	MYCN	MYD88	NBN	NF1
NF2	NFE2L2	NFKBIA	NKX2-1	NOTCH1	NOTCH2	NOTCH3	NPM1	NRAS	NT5C2
NTRK1	NTRK2	NTRK3	P2RY8	PALB2	PARK2	PARP1	PARP2	PARP3	PAX5
PBRM1	PDCD1	PDCD1L G2	PDGFRA	PDGFRB	PDK1	PIK3C2B	PIK3C2G	PIK3CA	PIK3CB
PIK3R1	PIM1	PMS2	POLD1	POLE	PPARG	PPP2R1A	PPP2R2A	PRDM1	PRKAR1A
PRKCI	PTCH1	PTEN	PTPN11	PTPRO	QKI	RAC1	RAD21	RAD51	RAD51B
RAD51C	RAD51D	RAD52	RAD54L	RAF1	RARA	RB1	RBM10	REL	RET
RICTOR	RNF43	ROS1	RPTOR	SDHA	SDHB	SDHC	SDHD	SETD2	SF3B1
SGK1	SMAD2	SMAD4	SMARCA4	SMARCB1	SMO	SNCAIP	SOC3	SOX2	SOX9
SPEN	SPOP	SRC	STAG2	STAT3	STK11	SUFU	SYK	TBX3	TEK
TET2	TGFBR2	TIPARP	TNFAIP3	TNFRSF14	TP53	TSC1	TSC2	TYRO3	U2AF1
VEGFA	VHL	WHSC1	WHSC1L1	WT1	XPO1	XRCC2	ZNF217	ZNF703	

B. Gene list: select rearrangements of the indicated genes were examined.

ALK intron 18, 19	BCL2 3' UTR	BCR intron 8, 13, 14	BRAF intron 7–10	BRCA1 intron 2, 7, 8, 12, 16, 19, 20
BRCA2 intron 2	CD74 intron 6–8	EGFR intron 7, 15, 24–27	ETV4 intron 5, 6	ETV5 intron 6, 7
ETV6 intron 5, 6	EWSR1 intron 7–13	EZR intron 9–11	FGFR1 intron 1, 5, 17	FGFR2 intron 1, 17
FGFR3 intron 17	KIT intron 16	KMT2A(MLL) intron 6–11	MSH2 intron 5	MYB intron 14
MYC intron 1	NOTCH2 intron 26	NTRK1 intron 8–10	NTRK2 intron 12	NUTM1 intron 1
PDGFRA intron 7, 9, 11	RAF1 intron 4–8	RARA intron 2	RET intron 7–11	ROS1 intron 31–35
RSPO2 intron 1	SDC4 intron 2	SLC34A2 intron 4	TERC non-coding RNA	TERT promoter
TMPRSS2 intron 1–3				

**Supplementary Table S2.** Details of MSI-high cases and MMR gene-mutant cases  
MSI-high case

Sample	Primary site	MMR gene	TMB (mut/Mb)	Other gene
SUR-758248	colorectal	Wild	24	ATM mutant (compound hetero)
SUR-758302	esophageal	Wild	19	BRCA2 truncation
SUR-758323	breast	MSH2	82	
SUR-758613	gastric	MSH3	26	

MMR: mismatch repair, TMB: tumor mutational burden

MMR gene-mutant case

Sample	Primary site	MMR gene	Alteration
SUR-758074	breast	<i>PMS2</i>	R315*
SUR-758323	breast	<i>MSH2</i>	Q409fs*7
SUR-758248	colorectal	<i>MSH3</i>	K383fs*32
SUR-758613	gastric	<i>MSH3</i>	K383fs*20
SUR-758613	gastric	<i>MSH3</i>	K383fs*32
SUR-758831	esophageal	<i>MSH2</i>	S612*
SUR-758072	other	<i>MSH3</i>	L290fs*10

**Supplementary Table S3.** The gene list of MMR genes and HRR genes  
MMR genes

<i>MLH1</i>	<i>MSH2</i>	<i>MSH6</i>	<i>PMS2</i>	<i>MSH3</i>
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HRR genes

<i>ARID1A</i>	<i>ATM</i>	<i>ATR</i>	<i>ATR</i>	<i>ATR</i>	<i>ATR</i>	<i>ATR</i>	<i>ATR</i>	<i>ATR</i>	<i>ATR</i>
<i>CDK12</i>	<i>CHEK1</i>	<i>CHEK2</i>	<i>FANCA</i>	<i>FANCC</i>	<i>FANCD2</i>	<i>FANCE</i>	<i>FANCF</i>	<i>FANCG</i>	<i>FANCL</i>
<i>MRE11A</i>	<i>NBN</i>	<i>PALB2</i>	<i>RAD50</i>	<i>RAD51</i>	<i>RAD51B</i>	<i>RAD51C</i>	<i>RAD51D</i>	<i>WRN</i>	