

INTENDED USE

JAX SomaticSeq is a qualitative next generation sequencing based assay that uses targeted selection of DNA from 517 genes of interest, and RNA from 55 known drivers, for a total of 1.94Mb panel size. JAX SomaticSeq detects genomic signatures including microsatellite instability (MSI) and tumor mutational burden (TMB) using DNA and RNA isolated from formalin-fixed paraffin embedded (FFPE) tissue specimens in a single, streamlined workflow. Over 1,000 clinical trials are available for the range of variants targeted by the assay. JAX SomaticSeq was developed with the Illumina TruSight Oncology 500 Assay and its performance characteristics determined by The Jackson Laboratory. It has not been cleared or approved by the US Food and Drug Administration.

GENE LIST

Small Variants Only		Small Variants & CNVs		Small Variants & Fusions		Small Variants, CNVs, & Fusions	
ABL1	CD274	EPHB1	GATA6	KDM5A	NFE2L2	PRKN (PARK2)	SOX10
ABL2	CD276	ERBB2 (HER2)	GEN1	KDM5C	NFKBIA	PRSS8	SOX17
ABRAXAS1 (FAM175A)	CD 74	ERBB3	GID4	KDM6A	NKX2-1	PTCH1	SOX2
ACVR1	CD79A	ERBB4	GLI1	KDR	NKX3-1	PTEN	SOX9
ACVR1B	CD79B	ERCC1	GNA11	KEAP1	NOTCH1	PTPN11	SPEN
ADGRA2 (GPR124)	CDC73	ERCC2	GNA13	KEL	NOTCH2	PTPRD	SPOP
AKT1	CDH1	ERCC3	GNAQ	KIF5B	NOTCH3	PTPRS	SPTA1
AKT2	CDK12	ERCC4	GNAS	KIT	NOTCH4	PTPRT	SRC
AKT3	CDK4	ERCC5	GPS2	KLF4	NPM1	QKI	SRSF2
ALK	CDK6	ERG	GREM1	KLHL6	NRAS	RAB35	STAG1
ALOX12B	CDK8	ERRF1	GRIN2A	KMT2A (MLL)	NRG1	RAC1	STAG2
AMER1 (FAM123B)	CDKN1A	ESR1	GRM3	KRAS	NSD1	RAD21	STAT3
ANKRD11	CDKN1B	ETS1	GSK3B	LAMP1	NTRK1	RAD50	STAT4
ANKRD26	CDKN2A	ETV1	H1-2 (HIST1H1C)	LATS1	NTRK2	RAD51	STAT5A
APC	CDKN2B	ETV4	H2BC5 (HIST1H2BD)	LATS2	NTRK3	RAD51B	STAT5B
AR	CDKN2C	ETV5	H3-3A (H3F3A)	LMO1	NUP93	RAD51C	STK11
ARAF	CEBPA	ETV6	H3-3B (H3F3B)	LRP1B	NUTM1	RAD51D	STK40
ARFRP1	CENPA	EWSR1	H3-4 (HIST3H3)	LYN	PAK1	RAD52	SUFU
ARID1A	CHD2	EZH2	H3-5 (H3F3C)	LZTR1	PAK3	RAD54L	SUZ12
ARID1B	CHD4	FANCA	H3C1 (HIST1H3A)	MAGI2	PAK5 (PAK7)	RAF1	SYK
ARID2	CHEK1	FANCC	H3C10 (HIST1H3H)	MALT1	PALB2	RANBP2	TAF1
ARID5B	CHEK2	FANCD2	H3C11 (HIST1H3I)	MAP2K1	PARP1	RARA	TBX3
ASXL1	CIC	FANCE	H3C12 (HIST1H3J)	MAP2K2	PAX3	RASA1	TCF3
ASXL2	COP1 (RFWD2)	FANCF	H3C13 (HIST2H3D)	MAP2K4	PAX5	RB1	TCF7L2
ATM	CREBBP	FANCG	H3C14 (HIST2H3C)	MAP3K1	PAX7	RBM10	TENT5C (FAM46C)
ATR	CRKL	FANCI	H3C15 (HIST2H3A)	MAP3K13	PAX8	RECQL4	TERC
ATRX	CRLF2	FANCL	H3C2 (HIST1H3B)	MAP3K14	PBRM1	REL	TERT
AURKA	CSF1R	FAS	H3C3 (HIST1H3C)	MAP3K4	PDCD1	RET	TET1
AURKB	CSF3R	FAT1	H3C4 (HIST1H3D)	MAPK1	PDCD1LG2	RHEB	TET2
AXIN1	CSNK1A1	FBXW7	H3C6 (HIST1H3E)	MAPK3	PDGFRA	RHOA	TFE3
AXIN2	CTCF	FGF1	H3C7 (HIST1H3F)	MAX	PDGFRB	RICTOR	TFRC
AXL	CTLA4	FGF10	H3C8 (HIST1H3G)	MCL1	PDK1	RIT1	TGFBF1
B2M	CTNNA1	FGF14	HGF	MDC1	PDPK1	RNF43	TGFBF2
BAP1	CTNNB1	FGF19	HNF1A	MDM2	PGR	ROS1	TMEM127
BARD1	CUL3	FGF2	HNRNPK	MDM4	PHF6	RPS6KA4	TMPRSS2
BBC3	CUX1	FGF23	HOXB13	MED12	PHOX2B	RPS6KB1	TNFAIP3
BCL10	CXCR4	FGF3	HRAS	MEF2B	PIK3C2B	RPS6KB2	TNFRSF14
BCL2	CYLD	FGF4	HSD3B1	MEN1	PIK3C2G	RPTOR	TOP1
BCL2L1	DAXX	FGF5	HSP90AA1	*MET	PIK3C3	RUNX1	TOP2A
BCL2L11	DCUN1D1	FGF6	ICOSLG	MGA	PIK3CA	RUNX1T1	TP53
BCL2L2	DDR2	FGF7	ID3	MITF	PIK3CB	RYBP	TP63
BCL6	DDX41	FGF8	IDH1	MLH1	PIK3CD	SDHA	TRAF2
BCOR	DHX15	FGF9	IDH2	MLL3	PIK3CG	SDHAF2	TRAF7
BCORL1	DICER1	FGFR1	IFNGR1	MPL	PIK3R1	SDHB	TSC1
BCR	DIS3	FGFR2	IGF1	MRE11 (MRE11A)	PIK3R2	SDHC	TSC2
BIRC3	DNAJB1	FGFR3	IGF1R	MSH2	PIK3R3	SDHD	TSHR
BLM	DNMT1	FGFR4	IGF2	MSH3	PIM1	SETBP1	UZAF1
BMPR1A	DNMT3A	FH	IKBKE	MSH6	PLCG2	SETD2	VEGFA
BRAF	DNMT3B	FLCN	IKZF1	MST1	PLK2	SF3B1	VHL
BRCA1	DOT1L	FLI1	IL10	MST1R	PMAIP1	SH2B3	VTCN1
BRCA2	E2F3	FLT1	IL7R	MTOR	PMS1	SH2D1A	WT1
BRD4	EED	FLT3	INHA	MUTYH	PMS2	SHQ1	XIAP
BRIP1	EGFL7	FLT4	INHBA	MYB	PNRC1	SLIT2	XPO1
BTG1	*EGFR	FOXA1	INPP4A	MYC	POLD1	SLX4	XRCC2
BTK	EIF1AX	FOXO2	INPP4B	MYCL (MYCL1)	POLE	SMAD2	YAP1
CALR	EIF4A2	FOXO1	INSR	MYCN	PPARG	SMAD3	YES1
CARD11	EIF4E	FOXP1	IRF2	MYD88	PPM1D	SMAD4	ZBTB2
CASP8	ELOC (TCEB1)	FRS2	IRF4	MYOD1	PPP2R1A	SMARCA4	ZBTB7A
CBFβ	EML4	FUBP1	IRS1	NAB2	PPP2R2A	SMARCB1	ZFXH3
CBL	EMSY (C11orf30)	FYN	IRS2	NBN	PPP6C	SMARCD1	ZNF217
CCN6 (WISP3)	EP300	GABRA6	JAK1	NCOA3	PRDM1	SMC1A	ZNF703
CCND1	EPCAM	GATA1	JAK2	NCOR1	PREX2	SMC3	ZRSR2
CCND2	EPHA3	GATA2	JAK3	NEGR1	PRKAR1A	SMO	*EGFR exons 2-7 (vIII) & MET exon 14 skipping events also reported
CCND3	EPHA5	GATA3	JUN	NF1	PRKCI	SNCAIP	
CCNE1	EPHA7	GATA4	KAT6A	NF2	PRKDC	SOCS1	

