

Supplemental Material: A Tumor-Specific Neo-Antigen Caused by a Frameshift Mutation in *BAP1* is a Potential Personalized Biomarker in Malignant Peritoneal Mesothelioma

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Table S1. 725 targeted genes in GenCap™ oncocap_725 kit.

<i>ABI1</i>	<i>ABI3</i>	<i>ABL1</i>	<i>ABL2</i>	<i>ABLIM1</i>	<i>ACBD6</i>	<i>ACCS</i>	<i>ACSL3</i>	<i>ACSL6</i>	<i>ACVR1B</i>
<i>ADAM12</i>	<i>ADAM19</i>	<i>ADAM29</i>	<i>ADAMTS16</i>	<i>ADAMTS18</i>	<i>ADAMTS20</i>	<i>ADAMTSL3</i>	<i>AFF1</i>	<i>AFF3</i>	<i>AIM1</i>
<i>AKAP12</i>	<i>AKAP9</i>	<i>AKT1</i>	<i>AKT2</i>	<i>AKT3</i>	<i>ALK</i>	<i>APC</i>	<i>APC2</i>	<i>APOL1</i>	<i>ARF1</i>
<i>ARHGAP26</i>	<i>ARHGEF12</i>	<i>ARID1A</i>	<i>ARNT</i>	<i>ARRB2</i>	<i>ASF1A</i>	<i>ASPSCR1</i>	<i>ASXL1</i>	<i>ATF1</i>	<i>ATIC</i>
<i>ATM</i>	<i>ATP10B</i>	<i>ATP13A3</i>	<i>BAG2</i>	<i>BAP1</i>	<i>BCL10</i>	<i>BCL11A</i>	<i>BCL11B</i>	<i>BCL2</i>	<i>BCL2L1</i>
<i>BCL3</i>	<i>BCL6</i>	<i>BCL7A</i>	<i>BCL9</i>	<i>BCR</i>	<i>BHLHE40</i>	<i>BIN1</i>	<i>BIRC3</i>	<i>BLM</i>	<i>BMI1</i>
<i>BMPR1A</i>	<i>BNIP3L</i>	<i>BRAF</i>	<i>BRCA1</i>	<i>BRCA2</i>	<i>BRD3</i>	<i>BRD4</i>	<i>BRD7</i>	<i>BRIP1</i>	<i>BTG1</i>
<i>BUB1</i>	<i>BUB1B</i>	<i>C15orf55</i>	<i>C18orf32</i>	<i>C2orf40</i>	<i>CANT1</i>	<i>CAPN1</i>	<i>CARD10</i>	<i>CARD11</i>	<i>CARD8</i>
<i>CARS</i>	<i>CASC5</i>	<i>CASP1</i>	<i>CASP8</i>	<i>CBFA2T3</i>	<i>CBFB</i>	<i>CBL</i>	<i>CBLB</i>	<i>CBLC</i>	<i>CBX8</i>
<i>CCDC6</i>	<i>CCL2</i>	<i>CCL26</i>	<i>CCNA2</i>	<i>CCNB1</i>	<i>CCNB1IP1</i>	<i>CCND1</i>	<i>CCND2</i>	<i>CCND3</i>	<i>CCNDBP1</i>
<i>CD27</i>	<i>CD3EAP</i>	<i>CD74</i>	<i>CDC25C</i>	<i>CDC42EP1</i>	<i>CDC6</i>	<i>CDC73</i>	<i>CDH1</i>	<i>CDH10</i>	<i>CDH11</i>
<i>CDK1</i>	<i>CDK2</i>	<i>CDK4</i>	<i>CDK5</i>	<i>CDK6</i>	<i>CDKN1A</i>	<i>CDKN1B</i>	<i>CDKN1C</i>	<i>CDKN2A</i>	<i>CDKN2AIP</i>
<i>CDKN2B</i>	<i>CDKN2C</i>	<i>CDKN2D</i>	<i>CDX2</i>	<i>CEBPA</i>	<i>CEP110</i>	<i>CHCHD7</i>	<i>CHEK1</i>	<i>CHEK2</i>	<i>CHIC2</i>
<i>CHN1</i>	<i>CIC</i>	<i>CIITA</i>	<i>CIRBP</i>	<i>CITED2</i>	<i>CLDN1</i>	<i>CLP1</i>	<i>CLTC</i>	<i>CLTCL1</i>	<i>CNBP</i>
<i>COL1A1</i>	<i>COMMD2</i>	<i>COMMD3</i>	<i>COMMD4</i>	<i>COMMD5</i>	<i>COMMD6</i>	<i>COMMD8</i>	<i>COX6C</i>	<i>CRAF</i>	<i>CREB1</i>
<i>CREB3L2</i>	<i>CREBBP</i>	<i>CREG1</i>	<i>CRTC1</i>	<i>CRTC3</i>	<i>CSF1R</i>	<i>CTNNA1</i>	<i>CTNNA2</i>	<i>CTNNB1</i>	<i>CTSF</i>
<i>CTSW</i>	<i>CXCL1</i>	<i>CXCR7</i>	<i>CYB561</i>	<i>CYLD</i>	<i>DAP3</i>	<i>DARC</i>	<i>DCLRE1B</i>	<i>DDB2</i>	<i>DDIT3</i>
<i>DDX10</i>	<i>DDX5</i>	<i>DDX6</i>	<i>DEK</i>	<i>DICER1</i>	<i>DLEC1</i>	<i>DMTF1</i>	<i>DNMT3A</i>	<i>DUSP3</i>	<i>DUSP4</i>
<i>DUSP6</i>	<i>DYRK1A</i>	<i>E2F1</i>	<i>E2F3</i>	<i>EDF1</i>	<i>EFEMP1</i>	<i>EGFR</i>	<i>EIF4A2</i>	<i>ELF4</i>	<i>ELK4</i>
<i>ELL</i>	<i>ELN</i>	<i>EML4</i>	<i>ENDOG</i>	<i>EP300</i>	<i>EPHA3</i>	<i>EPHA4</i>	<i>EPHA7</i>	<i>EPHB6</i>	<i>EPS15</i>

ERBB2	ERC1	ERCC2	ERCC3	ERCC4	ERCC5	ERG	ETS1	ETS2	ETV4
ETV5	EWSR1	EXT1	EXT2	EZH2	EZR	FAM123B	FANCA	FANCC	FANCD2
FANCE	FANCF	FANCG	FAS	FBX031	FBXW7	FCGR2B	FCGR3A	FCHSD1	FCRL4
FEV	FGFR1OP	FGFR2	FGFR3	FH	FHL2	FIP1L1	FLCN	FLI1	FLT3
FLT4	FN1	FOXA1	FOXL2	FOXO1	FOXO3	FOXO4	FOXP1	FRA6F	FSTL3
FUS	GADD45A	GAS7	GATA1	GATA2	GLI1	GLI3	GMPS	GNAQ	GNAS
GNG11	GOLGA5	GOPC	GPC3	GPHN	GRIN2A	GUCY1A2	HEPACAM	HERPUD1	HIP1
HIRA	HIST1H4I	HLF	HMGA1	HMGA2	HMOX1	HMOX2	HNF1A	HNRNPA2B1	HOOK3
HOXA11	HOXA13	HOXA9	HOXC11	HOXC13	HOXD11	HOXD13	HPV6AI1	HRAS	HSP90AA1
HSP90AB1	HSPA9	ID1	ID2	IDH1	IDH2	IER3	IFI16	IGFBP3	IGFBP7
IKBKB	IKBKE	IKBKG	IL12A	IL1R1	IL2	IL20	IL21R	IL6ST	INF2
INK4A	INTS4	IRAK1	IRAK1	IRAK2	IRF1	IRF2	IRF3	IRF4	IRF5
IRF7	IRF8	IRS4	ISG15	ITCH	ITGAD	ITGAE	ITGB2	ITGB3BP	ITK
JAK2	JAK3	JAZF1	JUNB	JUND	KDM5C	KDM6A	KDR	KDSR	KIAA1549
KIT	KL	KLF6	KLK2	KLRG1	KRAS	KSR1	KSR2	KTN1	LASP1
LCK	LCP1	LDB3	LEO1	LHFP	LIFR	LIMS1	LIMS2	LIMS3	LIMS3-LOC440895
LIMS3L	LMNA	LMO1	LMO2	LOC651644	LOXL1	LOXL2	LOXL3	LOXL4	LPP
LYL1	LYL1	MAD2L1BP	MAF	MAFB	MALT1	MAML2	MAP2K3	MAP2K4	MAP2K7
MAP3K14	MAP3K3	MAP3K8	MAPK14	MAPKAPK5	MAPKBP1	MCM2	MDM4	MECOM	MEN1
MET	MIR146A	MIR17	MIR205	MIR217	MIR22	MIR29A	MIR30A	MIR34A	MIR34B
MIR34C	MIR93	MITF	MKL1	MLF1	MLH1	MLL	MLL2	MLL3	MLLT1
MLLT10	MLLT11	MLLT3	MLLT4	MLLT6	MMP2	MN1	MNX1	MORF4	MPL
MSH2	MSH6	MSI2	MSN	MT-CYB	MTCP1	MTOR	MUC1	MUTYH	MXD3
MXD4	MYC	MYCL1	MYCN	MYD88	MYH11	MYH9	MYST3	MYST4	NACA
NBN	NCOA1	NCOA2	NCOA4	NDFIP1	NF1	NF2	NFKB1	NFKB2	NFKBIA
NFKBIB	NFKBID	NFKBIE	NFKBIL1	NFKBIZ	NFRKB	NIN	NINJ1	NKAP	NKAPL
NKAPP1	NKIRAS1	NKIRAS2	NLRP2	NOD1	NOD2	NONO	NOTCH1	NOTCH2	NOX4
NPM1	NR4A3	NRAS	NSD1	NTRK1	NTRK3	NUAK1	NUMA1	NUP214	NUP98

NUTF2	OLIG2	OMD	PAFAH1B2	PALB2	PATZ1	PAX3	PAX5	PAX7	PAX8
PBRM1	PBX1	PCSK7	PDCD11	PDCD4	PDE4DIP	PDGFB	PDGFRA	PDGFRB	PEA15
PER1	PHOX2B	PIAS4	PICALM	PIK3CA	PIK3CG	PIK3R1	PIM1	PINK1	PLA2R1
PLAG1	PLEKHG5	PML	PMS1	PMS2	POU2AF1	POU5F1	PPARG	PPM1A	PPP1R13L
PPP1R3A	PPP2R1A	PPP2R2C	PPP4C	PRCC	PRDM16	PRF1	PRG4	PRKAR1A	PRKCZ
PRKD1	PRPF19	PSD	PTCH1	PTEN	PTK2	PTPN11	PTPN22	PTPRU	PTTG1
RABEP1	RAD51L1	RAF1	RANBP17	RANBP2	RAP1A	RAP1GAP	RAP1GDS1	RARA	RARB
RASEF	RASGRF2	RB1	RBBP8	RBL1	RBL2	RBM15	RBX1	REL	RELA
RELB	RET	RGN	RHOH	RIPK1	RNF25	RNF43	ROMO1	ROS1	RPL10L
RPL22	RPLP1	RPN1	RRAD	RSL1D1	RUNX1	RUNX1T1	S1PR2	SAP30	SBDS
SDHB	SDHC	SDHD	SEC31A	SEN2	SEN3	SEN6	SEN6A	SEN6B	SENP1
5-Sep	6-Sep	9-Sep	SET	SFPQ	SFRS3	SH3GL1	SHC1	SILV	SIRT1
SIRT6	SKIL	SLC22A9	SLC45A3	SLC50A1	SMAD1	SMAD2	SMAD3	SMAD4	SMAD9
SMARCA4	SMARCB1	SMO	SOCS1	SRC	SRGAP3	SS18	SS18L1	SSX1	SSX2
SSX4	STIL	STK11	SUFU	SUMO1	SUMO2	SUMO3	SUMO4	SUPT4H1	SYK
TAB2	TAB3	TAF15	TAGLN	TAL1	TAL2	TANK	TBK1	TBKBP1	TCEA1
TCF12	TCF3	TCF7L1	TCF7L2	TCL1A	TCL6	TEC	TEK	TERC	TERT
TET1	TFE3	TFEB	TFG	TFPT	TFRC	TGFB1I1	TGFBR2	TGM3	THRAP3
TICAM1	TICAM2	TIFAB	TIMM50	TLX1	TLX3	TMED4	TMEM158	TMPRSS2	TNFAIP3
TNFRSF10B	TNFRSF17	TNFSF13	TNFSF14	TNIP1	TNIP2	TOP1	TP53	TP53BP1	TPM3
TPM4	TPR	TRADD	TRAF1	TRAF3	TRAF3IP2	TRAF3IP2-AS2	TRIM24	TRIM27	TRIM33
TRIP11	TSC1	TSC2	TSC22D3	TSHR	TTL	TUBB4	TWIST1	UBN1	UNC5CL
UQCRC2	USP6	USP9X	USP9Y	VHL	WAS	WHSC1	WHSC1L1	WLS	WRN
WT1	XPA	XPC	YWHAB	YWHAE	YWHAG	YWHAH	YWHAQ	YWHAZ	ZBTB16
ZBTB7A	ZNF331	ZNF384	ZNF521	ZSCAN4	-	-	-	-	-

Table S2. Primer sequences for Sanger validation.

Gene	Chromosome	Position	Primer (5' to 3')	T _m (°C)	Product Size (bp)
<i>NOTCH2</i>	1	120497774	F: TCAATAAGAACTATCACATGGGGCAG	60.5	231
			R: CAGGGCAGAGATGTAACATTGACATT	60.4	
<i>PDE4DIP</i>	1	144930847	F: ACCGGAACAACTCCATCTGTCAT	60.3	224
			R: CTGAGAGAAGTGCAAAGGAACTTGG	62.0	
<i>MN1</i>	22	28193936	F: CTTTGGAGCCGCTGCTACTG	61.9	222
			R: CCACCGCTTGCCAGAACAT	59.7	
<i>BAP1</i>	3	52437592	F: GCAGGCCTGTGCTGATGAC	61.9	219
			R: GCAATGAGAGTACAGACACGGC	61.9	
<i>ATP10B</i>	5	160097517	F: AGGAGGCATCTACTACTTTCCCC	62.0	220
			R: CCTCCTTTTTCTCTGACCCC	61.9	
<i>NSD1</i>	5	176721814	F: TACTATCAGCTGTGGTCCAGAC	60.1	245
			R: CTCAACAGCTCTCGGCATATGC	61.9	
<i>RAF1</i>	3	12626480	F: ACACAGTCAGTACCAGCCTC	61.9	224
			R: GTGCGAAAGGACCCAACCTCG	61.9	
<i>NSD1</i>	5	176665298	F: TACAACAATTTTGGCCTGTGGACT	58.6	223
			R: GCATGTCCACCTCCAAGTTCTG	61.9	
<i>DICER1</i>	14	95599707	F: GCTCCAGTATTAGTGTTTCGATTAGTA	60.5	221
			R: GTAGTAAGCTGTGCTAGAACAAAAATGC	60.5	
<i>NF2</i>	22	30054224	F: CCCATCTCACTTAGCTCCAATGACA	61.9	225
			R: AAGAGTCTATCGCCTTGAATGAAGA	60.4	
<i>BIVM-ERCC5</i>	13	103524612	F: TCTCATTGCTGTGTAAGTAATTGTTTCC	60.5	240
			R: CTCTAATTTTGTGCGAGATCAGGTTTCCC	61.9	

Table S2. Cont.

Gene	Chromosome	Position	Primer (5' to 3')	Tm (°C)	Product Size (bp)
<i>RBBP8</i>	18	20572853	F: CTCCTCCTCAAGAAGAATTACCTACTCG	62.2	232
			R: GACCCAAGACTGTGATGTGTGAAAAG	62.4	
<i>PRG4</i>	1	186276044	F: GACACGTCTACCACCCAACA	59.6	2689
			R: TGCCTTGATTGGGTACTCTCG	59.8	