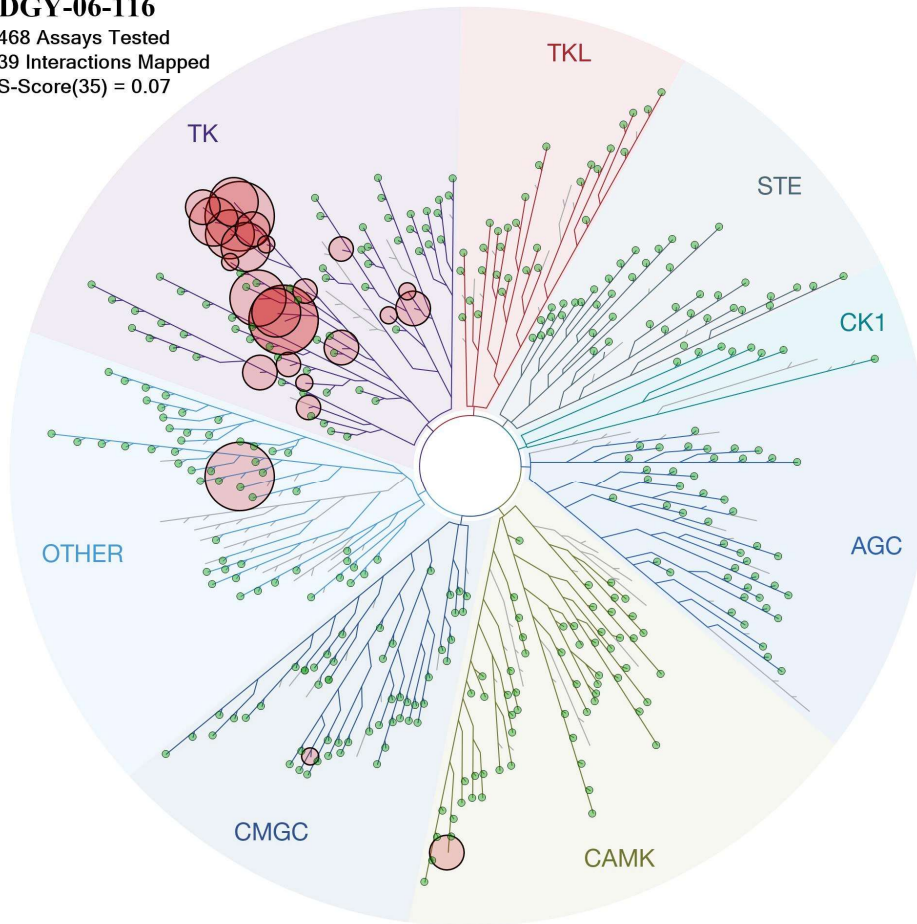
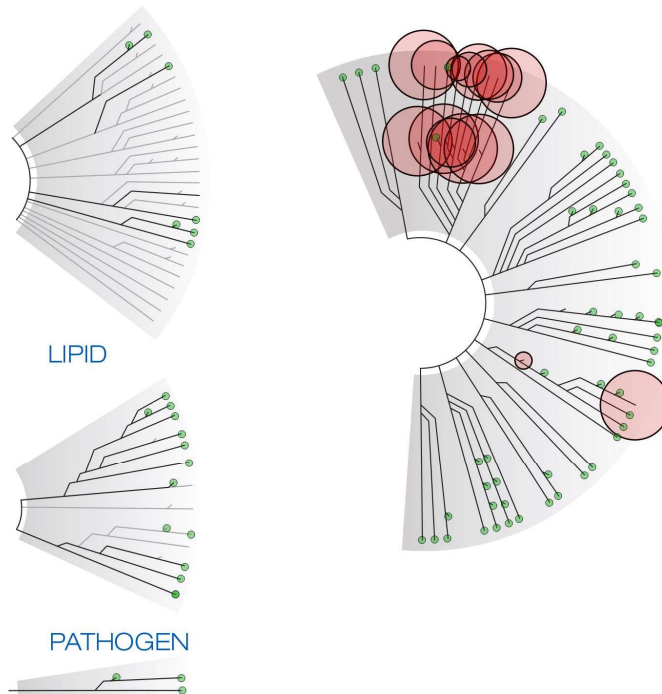


DGY-06-116
468 Assays Tested
39 Interactions Mapped
S-Score(35) = 0.07



ATYPICAL

MUTANT



Compound Name	DiscoverX Gene Symbol	Entrez Gene Symbol	Percent Control	Compound Concentration (nM)
DGY-06-116	AAK1	AAK1	100	1000
DGY-06-116	ABL1(E255K)-phosphorylated	ABL1	0.15	1000
DGY-06-116	ABL1(F317I)-nonphosphorylated	ABL1	0	1000
DGY-06-116	ABL1(F317I)-phosphorylated	ABL1	6.9	1000
DGY-06-116	ABL1(F317L)-nonphosphorylated	ABL1	0.9	1000
DGY-06-116	ABL1(F317L)-phosphorylated	ABL1	1.3	1000
DGY-06-116	ABL1(H396P)-nonphosphorylated	ABL1	0	1000
DGY-06-116	ABL1(H396P)-phosphorylated	ABL1	0	1000
DGY-06-116	ABL1(M351T)-phosphorylated	ABL1	0.8	1000
DGY-06-116	ABL1(Q252H)-nonphosphorylated	ABL1	0	1000
DGY-06-116	ABL1(Q252H)-phosphorylated	ABL1	0.7	1000
DGY-06-116	ABL1(T315I)-nonphosphorylated	ABL1	100	1000
DGY-06-116	ABL1(T315I)-phosphorylated	ABL1	100	1000
DGY-06-116	ABL1(Y253F)-phosphorylated	ABL1	0.1	1000
DGY-06-116	ABL1-nonphosphorylated	ABL1	0	1000
DGY-06-116	ABL1-phosphorylated	ABL1	0	1000
DGY-06-116	ABL2	ABL2	0.35	1000
DGY-06-116	ACVR1	ACVR1	100	1000
DGY-06-116	ACVR1B	ACVR1B	93	1000
DGY-06-116	ACVR2A	ACVR2A	100	1000
DGY-06-116	ACVR2B	ACVR2B	100	1000
DGY-06-116	ACVRL1	ACVRL1	100	1000
DGY-06-116	ADCK3	CABC1	100	1000
DGY-06-116	ADCK4	ADCK4	100	1000
DGY-06-116	AKT1	AKT1	89	1000
DGY-06-116	AKT2	AKT2	100	1000
DGY-06-116	AKT3	AKT3	100	1000
DGY-06-116	ALK	ALK	100	1000
DGY-06-116	ALK(C1156Y)	ALK	99	1000
DGY-06-116	ALK(L1196M)	ALK	90	1000
DGY-06-116	AMPK-alpha1	PRKAA1	100	1000
DGY-06-116	AMPK-alpha2	PRKAA2	100	1000
DGY-06-116	ANKK1	ANKK1	98	1000
DGY-06-116	ARK5	NUAK1	100	1000
DGY-06-116	ASK1	MAP3K5	100	1000
DGY-06-116	ASK2	MAP3K6	64	1000
DGY-06-116	AURKA	AURKA	100	1000
DGY-06-116	AURKB	AURKB	79	1000
DGY-06-116	AURKC	AURKC	98	1000
DGY-06-116	AXL	AXL	100	1000
DGY-06-116	BIKE	BMP2K	94	1000
DGY-06-116	BLK	BLK	0.15	1000
DGY-06-116	BMPR1A	BMPR1A	100	1000
DGY-06-116	BMPR1B	BMPR1B	100	1000
DGY-06-116	BMPR2	BMPR2	100	1000

DGY-06-116	BMX	BMX	48	1000
DGY-06-116	BRAF	BRAF	100	1000
DGY-06-116	BRAF(V600E)	BRAF	100	1000
DGY-06-116	BRK	PTK6	100	1000
DGY-06-116	BRSK1	BRSK1	100	1000
DGY-06-116	BRSK2	BRSK2	100	1000
DGY-06-116	BTK	BTK	0.1	1000
DGY-06-116	BUB1	BUB1	88	1000
DGY-06-116	CAMK1	CAMK1	100	1000
DGY-06-116	CAMK1B	PNCK	98	1000
DGY-06-116	CAMK1D	CAMK1D	100	1000
DGY-06-116	CAMK1G	CAMK1G	100	1000
DGY-06-116	CAMK2A	CAMK2A	100	1000
DGY-06-116	CAMK2B	CAMK2B	100	1000
DGY-06-116	CAMK2D	CAMK2D	100	1000
DGY-06-116	CAMK2G	CAMK2G	100	1000
DGY-06-116	CAMK4	CAMK4	100	1000
DGY-06-116	CAMKK1	CAMKK1	100	1000
DGY-06-116	CAMKK2	CAMKK2	100	1000
DGY-06-116	CASK	CASK	73	1000
DGY-06-116	CDC2L1	CDK11B	100	1000
DGY-06-116	CDC2L2	CDC2L2	100	1000
DGY-06-116	CDC2L5	CDK13	92	1000
DGY-06-116	CDK11	CDK19	96	1000
DGY-06-116	CDK2	CDK2	100	1000
DGY-06-116	CDK3	CDK3	100	1000
DGY-06-116	CDK4	CDK4	97	1000
DGY-06-116	CDK4-cyclinD1	CDK4	100	1000
DGY-06-116	CDK4-cyclinD3	CDK4	100	1000
DGY-06-116	CDK5	CDK5	100	1000
DGY-06-116	CDK7	CDK7	100	1000
DGY-06-116	CDK8	CDK8	100	1000
DGY-06-116	CDK9	CDK9	100	1000
DGY-06-116	CDKL1	CDKL1	72	1000
DGY-06-116	CDKL2	CDKL2	100	1000
DGY-06-116	CDKL3	CDKL3	100	1000
DGY-06-116	CDKL5	CDKL5	100	1000
DGY-06-116	CHEK1	CHEK1	100	1000
DGY-06-116	CHEK2	CHEK2	90	1000
DGY-06-116	CIT	CIT	100	1000
DGY-06-116	CLK1	CLK1	100	1000
DGY-06-116	CLK2	CLK2	100	1000
DGY-06-116	CLK3	CLK3	100	1000
DGY-06-116	CLK4	CLK4	100	1000
DGY-06-116	CSF1R	CSF1R	7.3	1000
DGY-06-116	CSF1R-autoinhibited	CSF1R	79	1000
DGY-06-116	CSK	CSK	1.3	1000

DGY-06-116	CSNK1A1	CSNK1A1	86	1000
DGY-06-116	CSNK1A1L	CSNK1A1L	100	1000
DGY-06-116	CSNK1D	CSNK1D	100	1000
DGY-06-116	CSNK1E	CSNK1E	100	1000
DGY-06-116	CSNK1G1	CSNK1G1	100	1000
DGY-06-116	CSNK1G2	CSNK1G2	100	1000
DGY-06-116	CSNK1G3	CSNK1G3	100	1000
DGY-06-116	CSNK2A1	CSNK2A1	68	1000
DGY-06-116	CSNK2A2	CSNK2A2	100	1000
DGY-06-116	CTK	MATK	100	1000
DGY-06-116	DAPK1	DAPK1	100	1000
DGY-06-116	DAPK2	DAPK2	100	1000
DGY-06-116	DAPK3	DAPK3	100	1000
DGY-06-116	DCAMKL1	DCLK1	100	1000
DGY-06-116	DCAMKL2	DCLK2	100	1000
DGY-06-116	DCAMKL3	DCLK3	100	1000
DGY-06-116	DDR1	DDR1	1.1	1000
DGY-06-116	DDR2	DDR2	26	1000
DGY-06-116	DLK	MAP3K12	97	1000
DGY-06-116	DMPK	DMPK	100	1000
DGY-06-116	DMPK2	CDC42BPG	100	1000
DGY-06-116	DRAK1	STK17A	100	1000
DGY-06-116	DRAK2	STK17B	100	1000
DGY-06-116	DYRK1A	DYRK1A	100	1000
DGY-06-116	DYRK1B	DYRK1B	74	1000
DGY-06-116	DYRK2	DYRK2	89	1000
DGY-06-116	EGFR	EGFR	80	1000
DGY-06-116	EGFR(E746-A750del)	EGFR	67	1000
DGY-06-116	EGFR(G719C)	EGFR	100	1000
DGY-06-116	EGFR(G719S)	EGFR	95	1000
DGY-06-116	EGFR(L747-E749del, A750P)	EGFR	66	1000
DGY-06-116	EGFR(L747-S752del, P753S)	EGFR	74	1000
DGY-06-116	EGFR(L747-T751del,Sins)	EGFR	73	1000
DGY-06-116	EGFR(L858R)	EGFR	62	1000
DGY-06-116	EGFR(L858R,T790M)	EGFR	100	1000
DGY-06-116	EGFR(L861Q)	EGFR	55	1000
DGY-06-116	EGFR(S752-I759del)	EGFR	92	1000
DGY-06-116	EGFR(T790M)	EGFR	100	1000
DGY-06-116	EIF2AK1	EIF2AK1	100	1000
DGY-06-116	EPHA1	EPHA1	100	1000
DGY-06-116	EPHA2	EPHA2	100	1000
DGY-06-116	EPHA3	EPHA3	68	1000
DGY-06-116	EPHA4	EPHA4	75	1000
DGY-06-116	EPHA5	EPHA5	78	1000
DGY-06-116	EPHA6	EPHA6	100	1000
DGY-06-116	EPHA7	EPHA7	100	1000
DGY-06-116	EPHA8	EPHA8	1	1000

DGY-06-116	EPHB1	EPHB1	66	1000
DGY-06-116	EPHB2	EPHB2	59	1000
DGY-06-116	EPHB3	EPHB3	97	1000
DGY-06-116	EPHB4	EPHB4	100	1000
DGY-06-116	EPHB6	EPHB6	6	1000
DGY-06-116	ERBB2	ERBB2	63	1000
DGY-06-116	ERBB3	ERBB3	89	1000
DGY-06-116	ERBB4	ERBB4	6.6	1000
DGY-06-116	ERK1	MAPK3	100	1000
DGY-06-116	ERK2	MAPK1	100	1000
DGY-06-116	ERK3	MAPK6	100	1000
DGY-06-116	ERK4	MAPK4	100	1000
DGY-06-116	ERK5	MAPK7	92	1000
DGY-06-116	ERK8	MAPK15	100	1000
DGY-06-116	ERN1	ERN1	98	1000
DGY-06-116	FAK	PTK2	100	1000
DGY-06-116	FER	FER	100	1000
DGY-06-116	FES	FES	100	1000
DGY-06-116	FGFR1	FGFR1	100	1000
DGY-06-116	FGFR2	FGFR2	95	1000
DGY-06-116	FGFR3	FGFR3	100	1000
DGY-06-116	FGFR3(G697C)	FGFR3	100	1000
DGY-06-116	FGFR4	FGFR4	100	1000
DGY-06-116	FGR	FGR	30	1000
DGY-06-116	FLT1	FLT1	100	1000
DGY-06-116	FLT3	FLT3	99	1000
DGY-06-116	FLT3(D835H)	FLT3	95	1000
DGY-06-116	FLT3(D835V)	FLT3	91	1000
DGY-06-116	FLT3(D835Y)	FLT3	100	1000
DGY-06-116	FLT3(ITD)	FLT3	100	1000
DGY-06-116	FLT3(ITD,D835V)	FLT3	89	1000
DGY-06-116	FLT3(ITD,F691L)	FLT3	85	1000
DGY-06-116	FLT3(K663Q)	FLT3	100	1000
DGY-06-116	FLT3(N841I)	FLT3	100	1000
DGY-06-116	FLT3(R834Q)	FLT3	100	1000
DGY-06-116	FLT3-autoinhibited	FLT3	100	1000
DGY-06-116	FLT4	FLT4	100	1000
DGY-06-116	FRK	FRK	5.1	1000
DGY-06-116	FYN	FYN	2	1000
DGY-06-116	GAK	GAK	0	1000
DGY-06-116	GCN2(Kin.Dom.2,S808G)	EIF2AK4	87	1000
DGY-06-116	GRK1	GRK1	69	1000
DGY-06-116	GRK2	ADRBK1	85	1000
DGY-06-116	GRK3	ADRBK2	100	1000
DGY-06-116	GRK4	GRK4	100	1000
DGY-06-116	GRK7	GRK7	100	1000
DGY-06-116	GSK3A	GSK3A	100	1000

DGY-06-116	GSK3B	GSK3B	98	1000
DGY-06-116	HASPIN	GSG2	100	1000
DGY-06-116	HCK	HCK	0.8	1000
DGY-06-116	HIPK1	HIPK1	100	1000
DGY-06-116	HIPK2	HIPK2	100	1000
DGY-06-116	HIPK3	HIPK3	99	1000
DGY-06-116	HIPK4	HIPK4	100	1000
DGY-06-116	HPK1	MAP4K1	100	1000
DGY-06-116	HUNK	HUNK	100	1000
DGY-06-116	ICK	ICK	96	1000
DGY-06-116	IGF1R	IGF1R	100	1000
DGY-06-116	IKK-alpha	CHUK	100	1000
DGY-06-116	IKK-beta	IKBKB	100	1000
DGY-06-116	IKK-epsilon	IKBKE	100	1000
DGY-06-116	INSR	INSR	100	1000
DGY-06-116	INSRR	INSRR	100	1000
DGY-06-116	IRAK1	IRAK1	100	1000
DGY-06-116	IRAK3	IRAK3	100	1000
DGY-06-116	IRAK4	IRAK4	100	1000
DGY-06-116	ITK	ITK	100	1000
DGY-06-116	JAK1(JH1domain-catalytic)	JAK1	100	1000
DGY-06-116	JAK1(JH2domain-pseudokinase)	JAK1	100	1000
DGY-06-116	JAK2(JH1domain-catalytic)	JAK2	90	1000
DGY-06-116	JAK3(JH1domain-catalytic)	JAK3	88	1000
DGY-06-116	JNK1	MAPK8	88	1000
DGY-06-116	JNK2	MAPK9	94	1000
DGY-06-116	JNK3	MAPK10	100	1000
DGY-06-116	KIT	KIT	40	1000
DGY-06-116	KIT(A829P)	KIT	0	1000
DGY-06-116	KIT(D816H)	KIT	41	1000
DGY-06-116	KIT(D816V)	KIT	100	1000
DGY-06-116	KIT(L576P)	KIT	65	1000
DGY-06-116	KIT(V559D)	KIT	21	1000
DGY-06-116	KIT(V559D,T670I)	KIT	100	1000
DGY-06-116	KIT(V559D,V654A)	KIT	100	1000
DGY-06-116	KIT-autoinhibited	KIT	92	1000
DGY-06-116	LATS1	LATS1	100	1000
DGY-06-116	LATS2	LATS2	100	1000
DGY-06-116	LCK	LCK	0.25	1000
DGY-06-116	LIMK1	LIMK1	100	1000
DGY-06-116	LIMK2	LIMK2	100	1000
DGY-06-116	LKB1	STK11	100	1000
DGY-06-116	LOK	STK10	96	1000
DGY-06-116	LRRK2	LRRK2	100	1000
DGY-06-116	LRRK2(G2019S)	LRRK2	100	1000
DGY-06-116	LTK	LTK	100	1000
DGY-06-116	LYN	LYN	2.8	1000

DGY-06-116	LZK	MAP3K13	91	1000
DGY-06-116	MAK	MAK	70	1000
DGY-06-116	MAP3K1	MAP3K1	99	1000
DGY-06-116	MAP3K15	MAP3K15	80	1000
DGY-06-116	MAP3K2	MAP3K2	94	1000
DGY-06-116	MAP3K3	MAP3K3	100	1000
DGY-06-116	MAP3K4	MAP3K4	65	1000
DGY-06-116	MAP4K2	MAP4K2	100	1000
DGY-06-116	MAP4K3	MAP4K3	100	1000
DGY-06-116	MAP4K4	MAP4K4	100	1000
DGY-06-116	MAP4K5	MAP4K5	100	1000
DGY-06-116	MAPKAPK2	MAPKAPK2	100	1000
DGY-06-116	MAPKAPK5	MAPKAPK5	100	1000
DGY-06-116	MARK1	MARK1	99	1000
DGY-06-116	MARK2	MARK2	100	1000
DGY-06-116	MARK3	MARK3	90	1000
DGY-06-116	MARK4	MARK4	95	1000
DGY-06-116	MAST1	MAST1	82	1000
DGY-06-116	MEK1	MAP2K1	93	1000
DGY-06-116	MEK2	MAP2K2	93	1000
DGY-06-116	MEK3	MAP2K3	69	1000
DGY-06-116	MEK4	MAP2K4	91	1000
DGY-06-116	MEK5	MAP2K5	81	1000
DGY-06-116	MEK6	MAP2K6	99	1000
DGY-06-116	MELK	MELK	100	1000
DGY-06-116	MERTK	MERTK	95	1000
DGY-06-116	MET	MET	100	1000
DGY-06-116	MET(M1250T)	MET	100	1000
DGY-06-116	MET(Y1235D)	MET	100	1000
DGY-06-116	MINK	MINK1	81	1000
DGY-06-116	MKK7	MAP2K7	96	1000
DGY-06-116	MKNK1	MKNK1	92	1000
DGY-06-116	MKNK2	MKNK2	97	1000
DGY-06-116	MLCK	MYLK3	100	1000
DGY-06-116	MLK1	MAP3K9	100	1000
DGY-06-116	MLK2	MAP3K10	100	1000
DGY-06-116	MLK3	MAP3K11	100	1000
DGY-06-116	MRCKA	CDC42BPA	100	1000
DGY-06-116	MRCKB	CDC42BPB	100	1000
DGY-06-116	MST1	STK4	93	1000
DGY-06-116	MST1R	MST1R	100	1000
DGY-06-116	MST2	STK3	100	1000
DGY-06-116	MST3	STK24	100	1000
DGY-06-116	MST4	MST4	85	1000
DGY-06-116	MTOR	MTOR	98	1000
DGY-06-116	MUSK	MUSK	100	1000
DGY-06-116	MYLK	MYLK	100	1000

DGY-06-116	MYLK2	MYLK2	100	1000
DGY-06-116	MYLK4	MYLK4	100	1000
DGY-06-116	MYO3A	MYO3A	100	1000
DGY-06-116	MYO3B	MYO3B	91	1000
DGY-06-116	NDR1	STK38	90	1000
DGY-06-116	NDR2	STK38L	93	1000
DGY-06-116	NEK1	NEK1	100	1000
DGY-06-116	NEK10	NEK10	100	1000
DGY-06-116	NEK11	NEK11	72	1000
DGY-06-116	NEK2	NEK2	100	1000
DGY-06-116	NEK3	NEK3	88	1000
DGY-06-116	NEK4	NEK4	94	1000
DGY-06-116	NEK5	NEK5	100	1000
DGY-06-116	NEK6	NEK6	100	1000
DGY-06-116	NEK7	NEK7	100	1000
DGY-06-116	NEK9	NEK9	100	1000
DGY-06-116	NIK	MAP3K14	100	1000
DGY-06-116	NIM1	MGC42105	100	1000
DGY-06-116	NLK	NLK	76	1000
DGY-06-116	OSR1	OXS1	100	1000
DGY-06-116	p38-alpha	MAPK14	28	1000
DGY-06-116	p38-beta	MAPK11	98	1000
DGY-06-116	p38-delta	MAPK13	100	1000
DGY-06-116	p38-gamma	MAPK12	97	1000
DGY-06-116	PAK1	PAK1	100	1000
DGY-06-116	PAK2	PAK2	97	1000
DGY-06-116	PAK3	PAK3	76	1000
DGY-06-116	PAK4	PAK4	100	1000
DGY-06-116	PAK6	PAK6	81	1000
DGY-06-116	PAK7	PAK7	100	1000
DGY-06-116	PCTK1	CDK16	100	1000
DGY-06-116	PCTK2	CDK17	100	1000
DGY-06-116	PCTK3	CDK18	98	1000
DGY-06-116	PDGFRA	PDGFRA	90	1000
DGY-06-116	PDGFRB	PDGFRB	29	1000
DGY-06-116	PDPK1	PDPK1	93	1000
DGY-06-116	PFCDPK1(P.falciparum)	CDPK1	72	1000
DGY-06-116	PFPK5(P.falciparum)	MAL13P1.279	100	1000
DGY-06-116	PFTAIRE2	CDK15	100	1000
DGY-06-116	PFTK1	CDK14	100	1000
DGY-06-116	PHKG1	PHKG1	100	1000
DGY-06-116	PHKG2	PHKG2	100	1000
DGY-06-116	PIK3C2B	PIK3C2B	100	1000
DGY-06-116	PIK3C2G	PIK3C2G	100	1000
DGY-06-116	PIK3CA	PIK3CA	100	1000
DGY-06-116	PIK3CA(C420R)	PIK3CA	100	1000
DGY-06-116	PIK3CA(E542K)	PIK3CA	100	1000

DGY-06-116	PIK3CA(E545A)	PIK3CA	94	1000
DGY-06-116	PIK3CA(E545K)	PIK3CA	79	1000
DGY-06-116	PIK3CA(H1047L)	PIK3CA	71	1000
DGY-06-116	PIK3CA(H1047Y)	PIK3CA	100	1000
DGY-06-116	PIK3CA(I800L)	PIK3CA	86	1000
DGY-06-116	PIK3CA(M1043I)	PIK3CA	100	1000
DGY-06-116	PIK3CA(Q546K)	PIK3CA	100	1000
DGY-06-116	PIK3CB	PIK3CB	100	1000
DGY-06-116	PIK3CD	PIK3CD	100	1000
DGY-06-116	PIK3CG	PIK3CG	100	1000
DGY-06-116	PIK4CB	PI4KB	91	1000
DGY-06-116	PIKFYVE	PIKFYVE	100	1000
DGY-06-116	PIM1	PIM1	100	1000
DGY-06-116	PIM2	PIM2	95	1000
DGY-06-116	PIM3	PIM3	100	1000
DGY-06-116	PIP5K1A	PIP5K1A	100	1000
DGY-06-116	PIP5K1C	PIP5K1C	100	1000
DGY-06-116	PIP5K2B	PIP4K2B	99	1000
DGY-06-116	PIP5K2C	PIP4K2C	100	1000
DGY-06-116	PKAC-alpha	PRKACA	100	1000
DGY-06-116	PKAC-beta	PRKACB	100	1000
DGY-06-116	PKMYT1	PKMYT1	100	1000
DGY-06-116	PKN1	PKN1	83	1000
DGY-06-116	PKN2	PKN2	100	1000
DGY-06-116	PKNB(M.tuberculosis)	pknB	100	1000
DGY-06-116	PLK1	PLK1	100	1000
DGY-06-116	PLK2	PLK2	100	1000
DGY-06-116	PLK3	PLK3	100	1000
DGY-06-116	PLK4	PLK4	92	1000
DGY-06-116	PRKCD	PRKCD	100	1000
DGY-06-116	PRKCE	PRKCE	100	1000
DGY-06-116	PRKCH	PRKCH	100	1000
DGY-06-116	PRKCI	PRKCI	94	1000
DGY-06-116	PRKCQ	PRKCQ	86	1000
DGY-06-116	PRKD1	PRKD1	100	1000
DGY-06-116	PRKD2	PRKD2	99	1000
DGY-06-116	PRKD3	PRKD3	99	1000
DGY-06-116	PRKG1	PRKG1	100	1000
DGY-06-116	PRKG2	PRKG2	100	1000
DGY-06-116	PRKR	EIF2AK2	100	1000
DGY-06-116	PRKX	PRKX	100	1000
DGY-06-116	PRP4	PRPF4B	81	1000
DGY-06-116	PYK2	PTK2B	100	1000
DGY-06-116	QSK	KIAA0999	69	1000
DGY-06-116	RAF1	RAF1	100	1000
DGY-06-116	RET	RET	100	1000
DGY-06-116	RET(M918T)	RET	87	1000

DGY-06-116	RET(V804L)	RET	100	1000
DGY-06-116	RET(V804M)	RET	100	1000
DGY-06-116	RIOK1	RIOK1	100	1000
DGY-06-116	RIOK2	RIOK2	100	1000
DGY-06-116	RIOK3	RIOK3	92	1000
DGY-06-116	RIPK1	RIPK1	100	1000
DGY-06-116	RIPK2	RIPK2	100	1000
DGY-06-116	RIPK4	RIPK4	100	1000
DGY-06-116	RIPK5	DSTYK	100	1000
DGY-06-116	ROCK1	ROCK1	100	1000
DGY-06-116	ROCK2	ROCK2	100	1000
DGY-06-116	ROS1	ROS1	100	1000
DGY-06-116	RPS6KA4(Kin.Dom.1-N-terminal)	RPS6KA4	100	1000
DGY-06-116	RPS6KA4(Kin.Dom.2-C-terminal)	RPS6KA4	100	1000
DGY-06-116	RPS6KA5(Kin.Dom.1-N-terminal)	RPS6KA5	100	1000
DGY-06-116	RPS6KA5(Kin.Dom.2-C-terminal)	RPS6KA5	100	1000
DGY-06-116	RSK1(Kin.Dom.1-N-terminal)	RPS6KA1	100	1000
DGY-06-116	RSK1(Kin.Dom.2-C-terminal)	RPS6KA1	100	1000
DGY-06-116	RSK2(Kin.Dom.1-N-terminal)	RPS6KA3	75	1000
DGY-06-116	RSK2(Kin.Dom.2-C-terminal)	RPS6KA3	100	1000
DGY-06-116	RSK3(Kin.Dom.1-N-terminal)	RPS6KA2	100	1000
DGY-06-116	RSK3(Kin.Dom.2-C-terminal)	RPS6KA2	100	1000
DGY-06-116	RSK4(Kin.Dom.1-N-terminal)	RPS6KA6	77	1000
DGY-06-116	RSK4(Kin.Dom.2-C-terminal)	RPS6KA6	96	1000
DGY-06-116	S6K1	RPS6KB1	95	1000
DGY-06-116	SBK1	SBK1	100	1000
DGY-06-116	SGK	SGK1	95	1000
DGY-06-116	SgK110	SgK110	60	1000
DGY-06-116	SGK2	SGK2	100	1000
DGY-06-116	SGK3	SGK3	86	1000
DGY-06-116	SIK	SIK1	1	1000
DGY-06-116	SIK2	SIK2	67	1000
DGY-06-116	SLK	SLK	100	1000
DGY-06-116	SNARK	NUAK2	100	1000
DGY-06-116	SNRK	SNRK	100	1000
DGY-06-116	SRC	SRC	0	1000
DGY-06-116	SRMS	SRMS	87	1000
DGY-06-116	SRPK1	SRPK1	100	1000
DGY-06-116	SRPK2	SRPK2	100	1000
DGY-06-116	SRPK3	SRPK3	100	1000
DGY-06-116	STK16	STK16	100	1000
DGY-06-116	STK33	STK33	100	1000
DGY-06-116	STK35	STK35	100	1000
DGY-06-116	STK36	STK36	47	1000
DGY-06-116	STK39	STK39	65	1000
DGY-06-116	SYK	SYK	100	1000
DGY-06-116	TAK1	MAP3K7	99	1000

DGY-06-116	TAOK1	TAOK1	100	1000
DGY-06-116	TAOK2	TAOK2	100	1000
DGY-06-116	TAOK3	TAOK3	100	1000
DGY-06-116	TBK1	TBK1	70	1000
DGY-06-116	TEC	TEC	95	1000
DGY-06-116	TESK1	TESK1	100	1000
DGY-06-116	TGFBR1	TGFBR1	100	1000
DGY-06-116	TGFBR2	TGFBR2	100	1000
DGY-06-116	TIE1	TIE1	83	1000
DGY-06-116	TIE2	TEK	100	1000
DGY-06-116	TLK1	TLK1	100	1000
DGY-06-116	TLK2	TLK2	100	1000
DGY-06-116	TNIK	TNIK	100	1000
DGY-06-116	TNK1	TNK1	29	1000
DGY-06-116	TNK2	TNK2	81	1000
DGY-06-116	TNNI3K	TNNI3K	83	1000
DGY-06-116	TRKA	NTRK1	100	1000
DGY-06-116	TRKB	NTRK2	100	1000
DGY-06-116	TRKC	NTRK3	100	1000
DGY-06-116	TRPM6	TRPM6	81	1000
DGY-06-116	TSSK1B	TSSK1B	86	1000
DGY-06-116	TSSK3	TSSK3	90	1000
DGY-06-116	TTK	TTK	94	1000
DGY-06-116	TXK	TXK	17	1000
DGY-06-116	TYK2(JH1domain-catalytic)	TYK2	100	1000
DGY-06-116	TYK2(JH2domain-pseudokinase)	TYK2	100	1000
DGY-06-116	TYRO3	TYRO3	89	1000
DGY-06-116	ULK1	ULK1	100	1000
DGY-06-116	ULK2	ULK2	100	1000
DGY-06-116	ULK3	ULK3	100	1000
DGY-06-116	VEGFR2	KDR	94	1000
DGY-06-116	VPS34	PIK3C3	98	1000
DGY-06-116	VRK2	VRK2	71	1000
DGY-06-116	WEE1	WEE1	100	1000
DGY-06-116	WEE2	WEE2	100	1000
DGY-06-116	WNK1	WNK1	100	1000
DGY-06-116	WNK2	WNK2	64	1000
DGY-06-116	WNK3	WNK3	100	1000
DGY-06-116	WNK4	WNK4	97	1000
DGY-06-116	YANK1	STK32A	100	1000
DGY-06-116	YANK2	STK32B	100	1000
DGY-06-116	YANK3	STK32C	100	1000
DGY-06-116	YES	YES1	0.55	1000
DGY-06-116	YSK1	STK25	100	1000
DGY-06-116	YSK4	MAP3K19	83	1000
DGY-06-116	ZAK	ZAK	98	1000
DGY-06-116	ZAP70	ZAP70	100	1000