Kinase	JQ1 10 µM	I-BET762 10 µM
ABL, ARG	8.0	-4.4
ACK	-20.1	10.8
AMPKa1, AMPKa2	-9.3	1.9
ATR	2.4	0.3
AurA	5.5	9.2
AurA, AurB, AurC	1.3	5.5
BRAF	-5.3	6.7
CaMK1d	-24.4	-24.5
CaMK2d	-12.0	-10.8
CaMK2g	-7.3	0.5
CaMK4	-16.1	-6.1
CDC2	-0.2	-15.0
CDK11, CDK8	13.6	14.6
CDK2	0.3	1.0
CDK5	0.6	3.1
CDK6	5.8	4.1
CDK9	10.0	2.9
CHK1	-0.8	-4.2
CHK2	-0.6	1.0
CK1a	7.4	6.9
CSK	-2.1	6.1
DNAPK	-2.2	-1.6
eEF2K	-10.4	-16.1
Erk1	-3.0	-2.9
Erk5	17.5	11.4
FER	8.0	13.3
FRAP	0.9	-1.1
GCK	3.9	14.3
GSK3B	3.7	-5.1
HPK1	-12.8	7.4
IKKa	-8.3	0.2
IKKe, TBK1	8.2	5.2
ILK	1.6	1.2
IRAK4	-2.3	8.3
IRE1	-6.6	20.6
JAK1 domain1	5.4	10.3
JAK1 domain2	-4.7	-4.0
JNK1, JNK2, JNK3	-8.1	3.9
KHS1	-0.7	-2.0
LATS1	0.2	4.7
LKB1	-5.6	2.0
LOK	5.3	4.7
MAP2K1, MAP2K2	0.0	-1.3
MAP2K3	0.0	5.3
MAP2K4	1.4	5.5
MAP2K6	-10.4	16.3
MAP3K2, MAP3K3	2.3	-0.1
MAP3K4	9.0	0.6
MARK2, MARK3	0.0	-11.9
MARK3, MARK4	4.7	0.2
MAST3	-1.8	0.3
MASTL	-7.7	-1.8
MLK3	14.6	8.9
MLKL	-1.6	-6.2
MSK1 domain1, MSK2 domain1	-1.9	8.5
MST1	2.8	2.9
MST2	17.6	15.1
MST3	-5.9	-20.8
MST4, YSK1	0.1	-4.4
NDR1	-0.3	2.3
NDR2	-11.9	-6.4
NEK1	0.6	-7.7
NEK6, NEK7	-6.5	2.3
NEK7	-1.0	3.2
NEK9	12.3	2.4
p38a	-25.4	-18.6
p38d, p38g	2.2	0.4
p70S6K	1.2	-9.5
p70S6K, p70S6Kb	-3.0	0.1
PCTAIRE2	9.7	-11.7
PEK	-1.3	-1.3
PHKg2	-12.9	3.2
PI4KA, PI4KAP2	9.7	6.3
PI4KB	2.9	4.2
PIK3C2B	3.6	16.2
PIK3C3	6.5	-1.7
PIK3CD	-17.1	-8.9
PIP4K2A	-6.5	-6.4
PIP4K2C	-5.0	1.7
PIP5K3	-9.0	0.4
PITSLRE	-3.5	0.0
PKCi	10.3	4.7
PKD2	0.3	-2.6
PKR	-10.8	2.2
PRPK	-3.2	11.1
ROCK1	-7.2	-9.1
RSK1 domain1, RSK2 domain1, RSK3 domain1	-1.5	7.5
RSK1 domain2	5.9	1.0
RSK2 domain1	-23.0	-1.4
RSK2 domain2	-28.6	-2.4
SGK3	0.7	5.0
SLK	-1.9	8.8
SMG1	-7.6	-7.1
SRPK1	-1.2	5.1
SRPK1, SRPK2	4.6	14.3
TAO1, TAO3	-13.5	-25.3
TAO2	-8.3	-0.3
TLK1	1.6	-12.8
TLK2	-3.5	-12.5
Wnk1, Wnk2, Wnk3	13.5	5.8
Wnk1, Wnk2, Wnk4	2.8	2.9
ZAK	-5.0	3.1
ZAP70	1.7	-1.0
ZC1/HGK, ZC2/TNIK, ZC3/MINK	1.4	-2.3