



AACI   MACI   MOSESTA   MACI   MOSESTA   MACI   MOSESTA   MACI	KINOMEscan Gene Symbol	Entrez Gene Symbol	Accession	scan MAX	scan EDGE	scan LIPID	scan TK	scanFLFCT	scan MODE	scan KINFTIC	Kd ELECT	Kd MAX
Mail   Principal	AAK1	AAK1	NP_055726.3	0				0	Jean WODL	0	0	0
Mail   Print	ABL1(E255K)-phosphorylated				0				_	0		
Mail: 1017-1017-2004-2004-200-200-200-200-200-200-200-2										$\cap$		
ABL   Propriet   ABL	ABL1(F317L)-phosphorylated ABL1(F317L)-nonphosphorylated							Ö				
ABL   1998   1	ABL1(F317L)-phosphorylated	ABL1	NP_005148.2	Ö			Ö	Ö	Ö		Ö	
Mail   Debt												
ABL   1928   1	ABL1(M351T)-phosphorylated ABL1(M351T)-phosphorylated								)	)		
ABILITY   Mark	ABL1(Q252H)-nonphosphorylated	ABL1	NP_005148.2	0			0	0			0	Ö
ABLITICATE   ABLITICATION   ABLITI	ABL1(Q252H)-phosphorylated											
ABLIVESTIT   ABL			_		0							
ABEL - INSERPONDERS   ABEL   NP 9051482   O O O O O O O O O O O O O O O O O O	ABL1(Y253F)-phosphorylated	ABL1	NP_005148.2	Ŏ			Ō	0	Ŭ	0	Ö	Ŏ
ABEZ  ABEZ  ABEZ  ABEZ  ABEZ  APEZ	ABL1-nonphosphorylated											
AGEVRII  ACVERII  ACV					0				O			
AGCYPEA ACYPEA A	ACVR1											
ACOPTES  ACO	ACVR1B				0							
ACCUPALL M. ACCUPALL M. P. 0500011 1			_							0		
ADDRIA MPC 991823 C	ACVRL1											
ANCT ANT BY COST BY CO	ADCK3	CABC1	NP_064632.2		0			0				
ART2 MAT2 MP 5001871	ADCK4											
AKT3   AKT3   AKT3   NP 0054561   O										O		
ALEXCE 1890')  ALEX NP 5049552 O O O O O O O O O O O O O O O O O O	AKT3	AKT3		0				0			Ö	0
ALSKL 198M) ALK MPFC aghal PRAAA1 ANSSE AN	ALK (21150V)				0							
AMERICA (1994)  AMERICA (1994)												
AMEN C 1997 AMEN C	AMPK-alpha1									)		
ARRIGINATION   NUMBER   NP 0598551   O	AMPK-alpha2	PRKAA2	NP_006243.2	Ō				0			Ö	Ö
ASKI	ANKK1											
MAPPIGE   NP 0048833   O										O		
AURING BURPAI BU	ASK2	MAP3K6	NP_004663.3	0				0			0	Ö
AUBRIC  AAXL  AXL  AXL  AXL  AXL  BARC  BMP2K  BMP2K  BMP2K  BMP1B  BMP1B  BMP1B  BMP1B  BMP1B  BMP1B  BMP1B  BMP1C  BMP1B  BMP1C  BMP1B  BMP1C  BMP1B  BMP1C  BMP1B  BMP1C  BMP1B  BMP1C  BMP1C  BMP1B  BMP1C  BMP1	AURKA											
AXL BRICE BMP2K BRICE BMP2K BROS00932  O O O O O O O O O O O O O O O O O O					O					O		
BIRE   SMP2K   NP-0600832   O   O   O   O   O   O   O   O   O	AXL				O		O			O		
BMPRIA   BMPRIA NP 004390	BIKE	BMP2K	NP_060063.2	0				0			0	Ö
BMPRIE   BMPRIE N. P. 001194.1   O	BLK BMDD1A						0			0		
BMPR 2   BMPR 2   NP 001195 2   O O O O O O O O O O O O O O O O O O	BMPR1B									0		
BRAFF   BRAF   NP 0043242   O O O O O O O O O O O O O O O O O O	BMPR2	BMPR2	NP_001195.2	0	0			0		0	0	0
BRAFY (1900E) BRAF NP 0005966; 1	BMX						0					
BRK   PTK6   NP 0095961   O O O O O O O O O O O O O O O O O O			_	_								
BRSK2 BRSK2 BRSK2 EAX024381 O O O O O O O D BUBI BITK BTK BTK BTK NP.0000521 O O O O O O O O BUBI BUBI NP.0004327.1 O O O O O O O O O O O CAMMY B NP.004327.1 O O O O O O O O O O O O CAMMY B PROCK EAX17235.1 O O O O O O O O O O CAMMY B PROCK EAX17235.1 O O O O O O O O O O O O O O O O O O O	BRK						0					
BIK BIK NP 000321 O O O O O O O O O O O O O O O O O O O	BRSK1	BRSK1		0	-			0	_	-	0	0
BUBI							$\cap$			$\circ$		
CAMKI	BUB1											
CAMKID	CAMK1	CAMK1	NP_003647.1	0				Ö			Ö	Ö
CAMKIG CAMKIG NP 0651721 O O O O O O O O O O O O O O O O O O O	CAMK1D										_	
CAMKZA  CAMKZB  CAMKZB  CAMKZB  CAMKZB  CAMKZD  CAMKZD  CAMKZD  CAMKZD  CAMKZD  CAMKZD  CAMKZC  CAMKZ  CAMKK1  CAMKK1  CAMKK1  CAMKK1  CAMKK1  CAMKK1  CAMKK1  CAMKK2  CAMKK2  PD0058403  CAMKK2  CAMKK2  CAMKK2  CAMKK2  CAMKK2  CAMKX2  CAMKK2  CAMKK1  CAMKX2  CAMKK2  CAMKK1  CAMKX2  CAMXX2  CAMXX2  CAMXX2  CAMXX2  CAMXX2  CAMXX2  CAMX				0				0		0	0	0
CAMK2D CAMK2D AAD20442.1 O O O O O O O O O O O O O O O O O O O	CAMK2A	CAMK2A	NP_741960.1									
CAMK2G CAMK4 CAMK4 CAMK4 CAMK4 CAMK1 CAMKK1 CAMKK1 CAMKK1 CAMKK1 CAMKK1 CAMKK2 CAMKK2 CAMKK2 CAMKK2 CAMKK2 CAMKK2 CAMKK2 CAMKX2 CAMKK2 CAMKC2 CAMKK2 CAMKC2 CAMKK2 CAMKC2 CAMCC2	CAMK2B									^		
CAMIKA  NP 115670.1  O  CAMIKA  CAMIKA  CAMIKA  CAMIKA  NP 006540.3  O  CAMIKA  CAMIKA  CAMIKA  CAMIKA  CAMIKA  NP 006540.3  O  CAMIKA  CAMIKA  CAMIKA  CAMIKA  NP 003679.2  O  COC2L.1  COC2L.1  COC2L.1  COC2L.1  COC2L.1  COC2L.2  COC2L.2  COC2L.2  COC2L.2  COC2L.2  COC2L.2  COC2L.2  COC2L.2  COC3L.3  COC										O		
CAMKK2	CAMK4		NP_001735.1	0				0		0	0	Ö
CASK	CAMKK1			_								
CDC2L1												
CDC212	CDC2L1											
CDK11	CDC2L2	CDC2L2	NP_076916.1	0				0		_	0	Ö
CDK2	CDC2L5											
CDK3	CDK11 CDK2				0					U		
CDK4-cyclinD1	CDK3	CDK3	NP_001249.1	Ö				0			0	Ö
CDK4-cyclinD3	CDK4											
CDK5         CDK5         NP 004926.1         O										0		
CDK7         CDK7         NP_001790.1         O         O         O         O         O         O         O         CDK8         O	CDK5											
CDK9         CDK9         NP_001252.1         O	CDK7	CDK7	NP_001790.1	0	0			0		0	0	0
CDKL1         CDKL1         NP_004187.2         O         O         O         O         O         C	CDK8									0		
CDKL2         CDKL2         NP_003939.1         O         O         O           CDKL3         CDKL3         NP_001107047.1         O         O         O         O           CDKL5         CDKL5         NP_003150.1         O         O         O         O           CHEK1         NP_001265.2         O         O         O         O         O           CHEK2         NP_001265.2         O         O         O         O         O           CHEK2         CHEK2         NP_001065.735.1         O         O         O         O           CIT         CIT         NP_009105.1         O         O         O         O         O           CIK1         CLK1         AAA61480.1         O         O         O         O         O         O           CLK2         GLK2         NP_003984.2         O	CDK9 CDKL1											
CDKL5         CDKL5         NP_003150.1         O	CDKL2	CDKL2	NP_003939.1	Ö				0			Ö	Ŏ
CHEK1         CHEK1         NP_001265.2         O	CDKL3											
CHEK2         CHEK2         NP_001005735.1         O         O         O         O           CIT         CIT         NP_009105.1         O         O         O         O           CLK1         CLK1         AAA61480.1         O         O         O         O           CLK2         CLK2         NP_003984.2         O         O         O         O           CLK3         NP_003983.2         O         O         O         O         O           CLK4         CLK4         NP_005983.2         O         O         O         O         O           CSF1R         CSF1R         NP_005202.2         O         O         O         O         O         O           CSF1R-autoinhibited         CSF1R         NP_005202.2         O         <					0					U		
CLK1         CLK1         AAA61480.1         O         O         O           CLK2         CLK2         NP 003984.2         O         O         O         O           CLK3         CLK3         NP 003983.2         O         O         O         O         O           CLK4         CLK4         NP 065717.1         O </td <td>CHEK2</td> <td></td> <td>NP_001005735.1</td> <td>0</td> <td></td> <td></td> <td></td> <td>0</td> <td></td> <td></td> <td>0</td> <td></td>	CHEK2		NP_001005735.1	0				0			0	
CLK2         CLK2         NP_003984.2         O	CIT	CIT	NP_009105.1	0	-			0	_		0	
CLK3         CLK3         NP_003983.2         O         O         O         O           CLK4         CLK4         NP_065717.1         O	CLK1									_		
CLK4         CLK4         NP_065717.1         O	CLK2 CLK3											
CSF1R-autoinhibited         CSF1R         NP_005202.2         O	CLK4	CLK4	NP_065717.1	Ö				0			0	Ö
CSK         CSK         NP_004374.1         O         O         O         O         O           CSNK1A1         CSNK1A1         NP_001883.4         O         O         O         O           CSNK1A1L         CSNK1A1L         NP_660204.1         O         O         O         O	CSF1R				0					0		
CSNK1A1         CSNK1A1         NP_001883.4         O         O         O           CSNK1A1L         CSNK1A1L         NP_660204.1         O         O         O									O	0		
CSNK1A1L   CSNK1A1L   NP_660204.1   O   O   O   O	CSNK1A1											
CSNKID	CSNK1A1L	CSNK1A1L	NP_660204.1	0				0			0	0
	CSNK1D	CSNK1D	NP_620693.1	0	0			0		0	0	0





KINOMEssan Cono Superbal	Entro: Care C	Accession	ccan NAAV	ccan EDCE	ccan LIDID	ccan TV	coapELECT	scan MODE	ccan KINETIC	VALLECT	Kd MAX
KINOMEscan Gene Symbol CSNK1E	Entrez Gene Symbol	NP 001885.1	scan MAX	scan EDGE	scan LIPID	scan TK	<i>scan</i> ELECT	scan MODE	scan KINETIC	Kd ELECT O	O
CSNK1G1	CSNK1G1	NP_071331.2	0				0			00	0
CSNK1G2	CSNK1G2	NP_001310.2	Ŏ	0			0			0	0
CSNK1G3	CSNK1G3	NP_004375.2	0				0			0	0
CSNK2A1	CSNK2A1	NP_001886.1	0				0		0	0	0
CSNK2A2 CTK	CSNK2A2 MATK	NP_001887.1	<u> </u>				0		0	00	0
DAPK1	DAPK1	NP_002369.2 NP_004929.2	0			0	0		0	00	0
DAPK2	DAPK2	NP_055141.2	0				Ö			Ö	Ö
DAPK3	DAPK3	NP_001339.1	ŏ				Ö			0	ŏ
DCAMKL1	DCLK1	NP_004725.1	0	0			0			0	0
DCAMKL2	DCLK2	NP_001035351.3	0				0			0	0
DCAMKL3	DCLK3	NP_208382.1	<u> </u>			_	Ŏ		Ŏ	Ŏ	Ŏ
DDR1 DDR2	DDR1 DDR2	NP_001945.3 CAA52777.1	0			0	0		0	00	0
DLK	MAP3K12	NP_006292.2	0			0	0			00	0
DMPK	DMPK	NP 004400.4	Ö				Ö		0	Ö	ŏ
DMPK2	CDC42BPG	NP_059995.2	Ö				Ö		Ö	Ö	Ŏ
DRAK1	STK17A	NP_004751.1	0				0			0	0
DRAK2	STK17B	NP_004217.1	0				0		0	0	0
DYRK1A	DYRK1A	NP_569121.1	<u> </u>				Ŏ		0	Ŏ	Ŏ
DYRK1B DYRK2	DYRK1B DYRK2	NP_004705.1 NP_003574.1	0	0			0		0	00	0
EGFR	EGFR	NP_005219.2	<del></del>	0		0	0		0	00	0
EGFR(E746-A750del)	EGFR	NP_005219.2	Ö			Ö	Ö			Ö	Ö
EGFR(G719C)	EGFR	NP_005219.2	Ö			Ŏ	Ŏ		0	Ö	Ŏ
EGFR(G719S)	EGFR	NP_005219.2	0			0	0		Ö	0	0
EGFR(L747-E749del, A750P)	EGFR	NP_005219.2	0			0	0			0	0
EGFR(L747-S752del, P753S)	EGFR	NP_005219.2	<u> </u>			0	0			0	0
EGFR(L747-T751del,Sins)	EGFR	NP_005219.2	<u> </u>	^		0	0		$\sim$	00	0
EGFR(L858R) EGFR(L858R,T790M)	EGFR EGFR	NP_005219.2 NP 005219.2	0	0		0	0		0	00	0
EGFR(L858R,1790M) EGFR(L861Q)	EGFR	NP_005219.2 NP_005219.2	0			0	0		0	0	0
EGFR(S752-I759del)	EGFR	NP_005219.2	0			0	0			0	0
EGFR(T790M)	EGFR	NP_005219.2	Ö			Ö	0		0	00	0
EIF2AK1	EIF2AK1	NP_055228.2	Ō				Ö		0	0	Ö
EPHA1	EPHA1	NP_005223.4	0			0	0		0	0	0
EPHA2	EPHA2	NP_004422.2	<u> </u>	0		0	0		_	0	0
EPHA3	EPHA3	NP_005224.2	0			0	0		0	0	0
EPHA4 EPHA5	EPHA4 EPHA5	NP_004429.1 NP_872272.2	0			0	0			00	0
EPHA6	EPHA6	NP 001073917.2	<del></del>			0	0		0	00	0
EPHA7	EPHA7	NP_004431.1	ŏ			ŏ	ŏ			ŏ	ŏ
EPHA8	EPHA8	NP_065387.1	Ö			Ö	Ö			0	Ö
EPHB1	EPHB1	NP_004432.1	0			0	0			0	0
EPHB2	EPHB2	NP_059145.2	0			0	0			0	0
EPHB3	EPHB3	NP_004434.2	0			0	0			0	0
EPHB4	EPHB4	NP_004435.3	<u> </u>			0	0		0	0	0
EPHB6 ERBB2	EPHB6 ERBB2	NP_004436.2 NP_001005862.1	0	0		0	0		0	00	0
ERBB3	ERBB3	NP_001003802.1	Ö	)		ŏ	ŏ			ŏ	ŏ
ERBB4	ERBB4	NP 001036064.1	Ö	0		Ö	Ö		0	0	ŏ
ERK1	MAPK3	NP_002737.2	Ö	Ö		Ĭ	Ö		Ĭ	Ö	Ö
ERK2	MAPK1	NP_620407.1	0				0			0	0
ERK3	MAPK6	NP_002739.1	0				0		0	0	0
ERK4	MAPK4	NP_002738.2	<u> </u>				Ŏ			Ŏ	Ŏ
ERK5	MAPK7 MAPK15	NP_002740.2	0				0			0	0
ERK8 ERN1	ERN1	NP_620590.2 NP 001424.3	0				00			00	0
FAK	PTK2	NP_722560.1	Ö	0		0	Ö		0	0	Ö
FER	FER	NP_005237.2	ŏ	)		Ö	Ö			0	ŏ
FES	FES	NP_001996.1	0			0	0			0	0
FGFR1	FGFR1	NP_075593.1	Ō			0	0		0	0	0
FGFR2	FGFR2	NP_075259.4	0	0		0	0		_	0	0
FGFR3	FGFR3	NP_000133.1	<u> </u>	0		0	0		0	00	0
FGFR3(G697C) FGFR4	FGFR3 FGFR4	NP_000133.1 NP_075252.2	0			0	0		0	0	0
FGR	FGR	NP_005239.1	<del></del>			0	0		0	00	0
FLT1	FLT1	NP_002010.2	Ö			00	00			00	Ö
FLT3	FLT3	NP_004110.2	0	0		0	0	0	0	0	0
FLT3(D835H)	FLT3	NP_004110.2	Q			0	0			0	0
FLT3(D835V)	FLT3	NP_004110.2	<u> </u>			0	0		0	0	0
FLT3(D835Y)	FLT3	NP_004110.2	0			0	0			00	0
FLT3(ITD) FLT3(ITD.D835V)	FLT3 FLT3	NP_004110.2 NP_004110.2	0			0	0		0	0	0
FLT3(ITD.D835V) FLT3(ITD.F691L)	FLT3	NP_004110.2	0			00	00		00	00	0
FLT3(K663Q)	FLT3	NP_004110.2	Ö			Ö	Ö			0	0
FLT3(N841I)	FLT3	NP_004110.2	Ö			0	0			0	0
FLT3(R834Q)	FLT3	NP_004110.2	0			0	0		0	0	0
FLT3-autoinhibited	FLT3	NP_004110.2	0			0	0	0	0	0	0
FLT4	FLT4	NP_002011.1	<u> </u>			0	0		0	0	0
FRK	FRK	NP_002022.1				0	0		0	0	0
FYN GAK	FYN GAK	NP_694592.1 NP_005246.2	0			0	0		0	0	0
GCN2(Kin.Dom.2,S808G)	EIF2AK4	NP 001013725.2	0				0		00	00	0
GRK1	GRK1	NP_002920.1	<del></del>				0		ŏ	0	Ö
GRK2	ADRBK1	NP_001610.2	0				)0			0	O
GRK3	ADRBK2	NP_005151.2	Ö				Ö			Ö	Ŏ
GRK4	GRK4	NP_892027.2	0				0			0	0
GRK7	GRK7	NP_631948.1	0				0		0	0	0
GSK3A	GSK3A	NP_063937.2	0				0		0	0	0
GSK3B HASPIN	GSK3B GSG2	NP_002084.2 NP_114171.2	0	0			0		0	00	0
HCK	HCK	NP_114171.2 NP_002101.2	0			0	0		0	00	0
HIPK1	HIPK1	NP_689909.2	0				0		0	0	Ö
コスエーバノナ体で会社											2014 04





SADDEGROED (September 1997)   September 1997   Septembe	DME <i>scan</i> Gene Symbol	Kd ELECT Kd N
BPSC		
PEPRE		
HUNK		
DEC		
GETER   GETER   MP 0008661   O O O O O O O O O O O O O O O O O O		
BIRCHESTS   DRING   NP 001871   O		
BRSC   BRSC   NP 064721.1   O		
RESPET   RISPISE N. P. G. 20530		
BRACI		
BRACS		
Title   Titl		
JAKE (JAPE (JAPE (JAPE )		
JAKE   JAMES   April		
JAKS_HI   Marker   JAKS_S   NP 0002012   O O O O O O O O O O O O O O O O O O		
JINKI		
JINE2		
JANES		
SIT		
KITCABEPD		
KITOBIEW)  KIT NP 000213.1		
KITOBINO   KIT		
KITUTSS9D   KIT	D816V)	0 0
KITUTSS9D, P6540)   KIT		O C
KITUTSSDQ.VSSAA		
KIT—autoninbited		
LATS		
LATS2 LOCK LOCK LOCK LOCK LOCK LOCK LOCK NP 00053473 O O O O O C LIMM(1 LIMM(1 LIMM(1 NP 002305.1 O O O O O O O O C LIMM(2 C LIRRIC2 C LIRRIC2 LIRRIC		
LCK		
LIMKI		
LIGH	(1 L	0 0
LORK		
LRRK2(Z019S)		
LRRK2(G2019S)		
LTK         LTK         NP 998844.1         O         O         O         O           LZK         MAPAIK13         NP 004712.1         O         O         O         O           MAK         MAK         MAK         NP 005912.1         O         O         O           MAP3K1         MAPAIK1         NP 005912.1         O         O         O           MAP3K1         NP 005912.1         O         O         O         O           MAP3K2         MAPAIK1         NP 006900.3         O         O         O         O           MAP3K2         MAPAIK2         NP 006903.3         O         O         O         O         O           MAP3K3         MAPAIK3         NP 003912.2         O		
LYN		
LZK         MAP3K13         NP 004712.1         O         O         O         O           MAK         MAK         NP 005912.1         O         O         O         O           MAP3K1         MAP3K15         NP 00101671.3         O         O         O         O           MAP3K2         MAP3K2         NP 00600.3         O         O         O         O           MAP3K3         MAP3K3         MAP3K3         NP 0059913.2         O         O         O         O           MAP4K4         MP0844         NP 0059913.2         O         O         O         O         O           MAP4K2         MAP4K4         NP 005901.2         O		
MAK         MAK         NP 005897.1         O         C         O           MAP3K15         MAP3K15         NP 001001671.3         O         O         O         O           MAP3K2         MAP08K15         NP 001001671.3         O         A         A         A         A         A         A         A         A         A         A         A         A		
MAP3K15	N	
MAP3K2		
MAP3K3		
MAP3K4		
MAP4K2		
MAP4K3		
MAPAK5		
MAPKAPK2	4K4 N	0 0
MAPKAPK5		
MARK1		
MARK2         MARK2         NP 059672.2         O         O         O           MARK3         MARK3         NP 002367.4         O         O         O         O           MARK4         MARK4         MARK4         NP 113055.2         O         O         O         O           MAST1         MAST1         NP 055790.1         O         O         O         O         O           MEK1         MAP2K1         NP 002746.1         O         O         O         O         O         O           MEK2         MAP2K3         NP 1095871.1         O	174	0 0
MARK3		0 0
MARK4		
MAST1         MAST1         NP.055790.1         O         O         O           MEK1         MAP2K1         NP.002746.1         O		
MEK2         MAP2K2         NP_109587.1         O		
MEK3         MAP2K3         NP.002747.2         O         O         O         O           MEK4         MAP2K4         NP.003001.1         O         O         O         O           MEK5         MAP2K5         NP.660143.1         O         O         O         O           MEK6         MAP2K6         NP.002749.2         O         O         O         O           MELK         MELK         NP.055606.1         O         O         O         O           METK         MERTK         AAB60430.1         O         O         O         O           MET         NP.000236.2         O         O         O         O         O           MET(Y1235D)         MET         NP.000236.2         O         O         O         O           MKNK1         MP.000236.2         O         O         O         O         O           MKK7         MAP2K7         NP.660186.1         O         O         O         O           MKNK1         MP.001129025.1         O         O         O         O         O           MKNK2         MAP3K10         NP.00129025.1         O         O         O         O	1 N	O C
MEK4         MAP2K4         NP_003001.1         O         O         O           MEK5         MAP2K5         NP_660143.1         O         O         O           MEK6         MAP2K6         NP_002749.2         O         O         O           MELK         MELK         NP_055606.1         O         O         O         O           METK         AAB60430.1         O         O         O         O         O         O           MET         MET         NP_000236.2         O		
MEK5         MAP2K5         NP_660143.1         O         O         O           MEK6         MAP2K6         NP_002749.2         O         O         O         O           MELK         NP_055606.1         O         O         O         O         O           MERTK         AAB60430.1         O         O         O         O         O           MET         NP_000236.2         O         O         O         O         O           MET(M1250T)         MET         NP_000236.2         O         O         O         O           MET(Y1235D)         MET         NP_000236.2         O         O         O         O           MET(Y1235D)         MET         NP_000236.2         O         O         O         O           MEXTY         NP_000236.2         O         O         O         O         O           MEXTY1235D)         MET         NP_000236.2         O         O         O         O           MEXTY1235D)         MET         NP_000236.2         O         O         O         O           MKK7         MAP2K7         NP_60120108.1         O         O         O         O		
MEK6         MAP2K6         NP_002749.2         O         O           MELK         MELK         NP_055606.1         O         O         O           MERTK         AAB60430.1         O         O         O         O         O           MET         NP_000236.2         O         O         O         O         O         O           MET(M1250T)         MET         NP_000236.2         O <td></td> <td></td>		
MELK         MELK         NP_055606.1         O         O           MERTK         MERTK         AAB60430.1         O         O         O           MET         NP_000236.2         O         O         O         O           MET         NP_000236.2         O         O         O         O           MET(Y1235D)         MET         NP_000236.2         O         O         O         O           MINK         MINKI         NP_0001020108.1         O         O         O         O           MKK7         MAP2K7         NP_660186.1         O         O         O         O           MKNK1         MRNK1         NP_060186.1         O         O         O         O           MKNK2         MP_660186.1         O         O         O         O         O           MKNK1         NP_060186.1         O         O         O         O         O           MKNK1         NP_060186.1         O         O         O         O         O           MKNK2         NP_060186.1         O         O         O         O         O           MLCK         MYLK3         NP_362299.2         O         O		
MERTK         MERTK         AAB60430.1         O         O         O         O         O           MET         MET         NP.000236.2         O         O         O         O         O           MET(M1250T)         MET         NP.000236.2         O         O         O         O         O           MET(Y1235D)         MET         NP.000236.2         O         O         O         O         O           MINK         MINKI         NP.001020108.1         O         O         O         O         O           MKK7         MAP2K7         NP.660186.1         O         O         O         O         O         O           MKNK1         NP.001129025.1         O		
MET         NP_000236.2         O         <	TK N	0 0
MET(Y1235D)         MET         NP_000236.2         O         O           MINK         MINK1         NP_001020108.1         O         O           MKK7         MAP2K7         NP_660186.1         O         O         O           MKNK1         MKNK1         NP_0601129025.1         O         O         O         O           MKNK2         MKNK2         NP_060042.2         O         O         O         O           MLCK         MYLK3         NP_872299.2         O         O         O         O           MLK1         MAP3K9         NP_149132.2         O         O         O         O           MLK2         MAP3K10         NP_002437.2         O         O         O         O           MLK3         MAP3K11         NP_002410.1         O         O         O         O           MRCKA         CDC42BPA         NP_003598.2         O         O         O         O           MST1         STK4         NP_006026.3         O         O         O         O           MST1R         MST1R         NP_002438.2         O         O         O         O		
MINK         MINK1         NP_001020108.1         O         O         O           MKK7         MAP2K7         NP_660186.1         O         O         O         O         O           MKNK1         MKNK1         NP_001129025.1         O		
MKK7         MAP2K7         NP_660186.1         O         O         O           MKNK1         MKNK1         NP_060129025.1         O         O         O         O           MKNK2         MKNK2         NP_060042.2         O         O         O         O           MLCK         MYLK3         NP_872299.2         O         O         O         O           MLK1         MAP3K9         NP_149132.2         O         O         O         O         O           MLK2         MAP3K10         NP_002437.2         O         O         O         O         O           MLK3         MAP3K11         NP_002437.2         O         O         O         O         O           MRCKA         CDC42BPA         NP_003598.2         O         O         O         O         O           MRCKB         CDC42BPB         NP_00626.3         O         O         O         O         O           MST1         STK4         NP_006273.1         O         O         O         O         O           MST1R         MST1R         NP_002438.2         O         O         O         O         O		
MKNK1         MKNK1         NP_001129025.1         O         O         O         O           MKNK2         MKNK2         NP_060042.2         O         O         O         O         O           MLCK         MYLK3         NP_872299.2         O         O         O         O         O           MLK1         MAP3K9         NP_149132.2         O         O         O         O         O         O           MLK2         MAP3K10         NP_002437.2         O		
MKNK2         MKNK2         NP_060042.2         O         O         O         O           MLCK         MYLK3         NP_872299.2         O         O         O         O           MLK1         MAP3K9         NP_149132.2         O         O         O         O           MLK2         MAP3K10         NP_002437.2         O         O         O         O           MLK3         MAP3K11         NP_002410.1         O         O         O         O           MRCKA         CDC42BPA         NP_003598.2         O         O         O         O           MRCKB         CDC42BPB         NP_006026.3         O         O         O         O           MST1         STK4         NP_006273.1         O         O         O         O           MST1R         MST1R         NP_002438.2         O         O         O         O		
MLCK         MYLK3         NP_872299.2         O         O         O           MLK1         MAP3K9         NP_149132.2         O         O         O         O           MLK2         MAP3K10         NP_002437.2         O         O         O         O           MLK3         MAP3K11         NP_002410.1         O         O         O         O           MRCKA         CDC42BPA         NP_003598.2         O         O         O         O           MRCKB         CDC42BPB         NP_006026.3         O         O         O         O           MST1         STK4         NP_006273.1         O         O         O         O           MST1R         MST1R         NP_002438.2         O         O         O         O		
MLK1         MAP3K9         NP_149132.2         O         O         O         O         O           MLK2         MAP3K10         NP_002437.2         O         O         O         O           MLK3         MAP3K11         NP_002410.1         O         O         O         O           MRCKA         CDC42BPA         NP_003598.2         O         O         O         O           MRCKB         CDC42BPB         NP_006026.3         O         O         O         O           MST1         STK4         NP_006273.1         O         O         O         O           MST1R         MST1R         NP_002438.2         O         O         O         O	K	0 0
MLK3         MAP3K11         NP_002410.1         O         O         O           MRCKA         CDC42BPA         NP_003598.2         O         O         O           MRCKB         CDC42BPB         NP_006026.3         O         O         O           MST1         STK4         NP_006273.1         O         O         O           MST1R         MST1R         NP_002438.2         O         O         O		O C
MRCKA         CDC42BPA         NP_003598.2         O         O         O           MRCKB         CDC42BPB         NP_006026.3         O         O         O           MST1         STK4         NP_006273.1         O         O         O           MST1R         MST1R         NP_002438.2         O         O         O		
MRCKB         CDC42BPB         NP_006026.3         O         O         O           MST1         STK4         NP_006273.1         O         O         O           MST1R         MST1R         NP_002438.2         O         O         O		
MST1         STK4         NP_006273.1         O         O         O           MST1R         MST1R         NP_002438.2         O         O         O         O		
MST1R		
MST3   STK24   NP_003567.2   O   O   O	3	0 0
MST4 NP 057626.2 O O O	4 N	0 0
MTOR MTOR NP_004949.1 O O O		
MUSK MVSK NP.005583.1 O O O O O		
MYLK         MYLK         NP_444254.3         O         O         O           MYLK2         MYLK2         NP 149109.1         O         O         O         O		
MYLK2         MYLK2         NP_149109.1         O         O         O           MYLK4         NP 001012418.1         O         O         O		
MYO3A MYO3A NP_059129.3 O O		
MY03B MY03B NP_001077084.2 O O		
NDR1 STK38 NP_009202.1 O O O		





STORY	KINOMEscan Gene Symbol	Entroz Gono Symbol	Accession	scan MAX	scan EDGE	scan LIPID	scan TK	SCAPELECT	scan MODE	scan KINETIC	Kd ELECT	Kd MAX
SECT   SECT   PROSESSED   O	·				SculledGE	SCUII LIPID	scan		SCUIT NIODE	SCUIT KINETIC		
SECOLO												
SEC												
SECOND   PROCESSED   PROCESSED   Company   C										0		
SESS										_		
MESS			_							0		
Select												
SECT												
SECTION   SECT										0		
MIST   MACK	NEK9	NEK9										
Section			_									
SSET   DOSE   PROSTOCE   DOSE   DOS												
Self-Sense   MAPPEL   MAPPEL   MP 200212   O O O O O O O O O O O O O O O O O O												
SEM-Brian   MAPK1					0					0		
SEP-DENE   MAPRICI   MAPRICI   MAPRICI   MAPRICI   MAPRICI   MAPRICI   MAPRICI   MARCINE   MAPRICI   MARCINE   MAR												
PARK					Ú							
MARCE   MARC	p38-gamma											
AMAG			_									
PAMES					0							
PARC												
ART					0							
ECTEX												
PECTICAL   CDENTAL NP 0025872   C			NP_006192.1		0			0		0		
## POOFFRA												
POGFRB												
PEPECPENT   PEPEC   PEPEC   PEPECPENT												
PRECENT   PREC							U			0		
PFFKQP   P												
PFTARIE2										0		
PHKG1	PFTAIRE2	CDK15	NP_631897.1	0				0			0	0
PHRGQ2												
PRISCISES PRISCI												
PRISCOA   PRISCOA NP 004591   2												
PIRSCA   P					0					0		
PRISCACLEGRED   PRISCA N. P. 006209.2   O O O O O O O O O O O O O O O O O O					0					0		
PRISCACEGNAN   PRISCA   NP 005092   O O O O O O O O O O O O O O O O O O												
PRISCACH (1947)	PIK3CA(E542K)	PIK3CA	NP_006209.2	0		0		0		0	0	0
PRISCACHHOPTY												
PINGSCA (HIGHYY)										0		
PRISCA MINO   PRISCA   N. P. 008209.2   O   O   O   O   O   O   O   O   O			_									
PIRSCAMMO48K												
PIRSCAGA   PIRSCAGA   NP 0082092   Q												
PIRSCO										0		
PIKAGG												
PIKKYE												
PINTY					O							
PIM1			_							0		
PIMS					0	)				0		
PIPSKIA	PIM2	PIM2		0	0			0			0	
PIPSKIC					0			0			0	0
PIPEK2B												
PIPSK2C												
PRAC-beta   PRKACB   NP 002721   O O O O O O O O O O O O O O O O O O												
PRAC-beta   PRKACB   NP 002722.1   O					0							
PKMYT1												
PKN1	PKMYT1	PKMYT1	NP_872629.1	0				0			0	Ö
PKNB(M.tuberculosis)												
PLK1												
PLK2         PLK2         NP 006613.2         O         O         O         O           PLK3         NP 104064.2         O <td< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td><math>\sim</math></td><td></td><td></td></td<>										$\sim$		
PLK3												
PLK4					0							
PRKCD	PLK4	PLK4	NP_055079.2	0				0			Ō	Ö
PRKCH         PPKCH         NP 006246.2         O         O         O         O         O         PRKCI         PRKCI         NP 002731.3         O												
PRKCI					0							
PRKCQ         PRKCQ         NP_006248.1         O										^		
PRKD1         PRKD1         NP.002733.2         O         O         O         O           PRKD2         PRKD2         NP.057541.2         O         O         O         O           PRKD3         PPKD3         NP.005804.1         O         O         O         O         O           PRKG1         PPKD3         NP.005804.1         O										0		
PRKD2         PRKD2         NP.057541.2         O										0		
PRKG1         PRKG1         NP.006249.1         O         O         O         O           PRKG2         PRKG2         NP.006250.1         O         O         O         O           PRKR         EIF2AK2         NP.002750.1         O         O         O         O           PRKX         PRKX         NP.005035.1         O         O         O         O           PRP4         PRPF4B         NP.003904.3         O         O         O         O           PYK2         PTK2B         NP.775267.1         O         O         O         O         O           QSK         KIAA0999         BAA76843.2         O         O         O         O         O           RET         NP.002871.1         O         O         O         O         O         O         O           RET         NP.065681.1         O		PRKD2	NP_057541.2					0			0	
PRKG2         PRKG2         NP.006250.1         O		PRKD3	NP_005804.1	0				0			0	0
PRKR         EIF2AK2         NP.002750.1         O												
PRKX         PRKX         NP.005035.1         O         O         O         O           PRP4         PRPF4B         NP.003904.3         O										0		
PRP4         PRPF4B         NP_003904.3         O												
PYK2         PTK2B         NP_775267.1         O												
QSK         KIAA0999         BAA76843.2         O							0					
RAF1	QSK											
RET	RAF1		NP_002871.1	0				0			0	0
RET (V804L)         RET         NP_065681.1         O	RET				0							
RET (V804M)         RET         NP_065681.1         O												
RIOK1   RIOK1   NP_113668.2   O												
RIOK2         RIOK2         NP_060813.2         O         O         O         O         O           RIOK3         NP_003822.2         O         O         O         O         O							U			0		
RIOK3 NP_003822.2 O O O					0					0		
RIPK1		RIOK3		0								
	RIPK1	RIPK1	NP_003795.2	0				0		0	0	0





SPECAL   S	KINOMEscan Gene Symbol	Entroz Gono Sumbol	Accession	scan MAX	scan EDGE	scan LIPID	scan TK	scanELECT	scan MODE	scan KINETIC	Kd ELECT	Kd MAX
BOOK	•				scan EDGE	scan LIPID	scan I K		scan MODE	scan KINETIC		
SPIRAL				0				0				
SPRING   S				0				0				
										0		
BOST	ROCK1	ROCK1	NP_005397.1	0				0			0	0
			NP_004841.2	0	0			0		0	0	
RESPECTATION   PROPERTY   PROPERTY   REPORT							0					
BEBBIASTOR   Den 1   M. termonia   DESPEAS   BE 7271881   O			_	_							•	
RESCHIED Dum   PETSHORAL   NP (200442   0												
REST (LEAD DATE 2**)										0		
BRACKER, Dent   February   BPSERCAL   NP 004571   O												
					0					0		
FERSIGNED Den 1-19t temmod   BPSSRAZ   DPSSRAZ   DPSSR										- Ŭ		
BESS/GER_DON-2**C-terminals   BPSS6A2   NP D0180911   O										0		
RSSAME   RPSGRASE   RPSGRASE   RP D05521   D				0				Ö			0	0
Sept	RSK4(Kin.Dom.1-N-terminal)	RPS6KA6	NP_055311.1	0				0		0	0	0
Service												
SQK												
Section												
SGNK2			_							0		
SOKA												
Six				_							•	
SR2												
SLK												
SHAPIK   SHIRK   SHI												
SHRK					C					0		
SRC												
SRMS					0		0					
SEPPKI										_		
SEPPK3				0				0			Ö	0
SIK16												
STR433	SRPK3	SRPK3	NP_055185.2	0	0			0			0	0
STR36										0		
STK36												
STK49												
SYK										0		
TAKIL							_					
TAOKI							0					
TAOK2												
TAOKS												
TBK1												
TEC. TEC. NP 0032061 O O O O O O O O O O O O O O O O O O O												
TESKI TESKI Nº 006278.2							0			- Ŭ		
TGFBR2			_				Ĭ					
TIEL   TIEL   NP 003415.1   O	TGFBR1	TGFBR1	NP_004603.1	0	0			Ö			Ö	Ö
TIEZ	TGFBR2	TGFBR2	NP_003233.4	0				0		0	0	0
TLK1	TIE1	TIE1	NP_005415.1	0			0	0		0	0	0
Tikk2					0		0			0		
TNIK												
TINK1												
TNK2										0		
TANISK				0			Ö	0			Ö	Ö
TRKA NTRK1 NP 001012331.1 O O O O O O O O O TRKB NTRK2 NP 006171.2 O O O O O O O O O O O O O O O O O O O				0			0	0		0	0	0
TRKB					$\cap$		$\cap$			0		
TRKC												
TRPM6												
TSSK1B												
TSSK3			NP_114417.1		0							
TTK         TTK         NP 003309.2         O			NP_443073.1	0								0
TYK2(JH1domain-catalytic)         TYK2         NP 003322.2         O			NP_003309.2							0		
TYK2(JH2domain-pseudokinase)         TYK2         NP_003322.3         O												
TYRO3					0							
ULK1										O		
ULK2         ULK2         NP_055498.3         O							O			_		
ULK3         ULK3         NP_001092906.1         O					_							
VEGFR2         KDR         NP 002244.1         O					U							
VPS34         PIK3C3         NP_002638.2         O					0		$\sim$					
VRK2         VPRV2         NP_006287.2         O         O         O           WEE1         WEE1         NP_001099028.1         O         O         O         O           WNK1         NP_001099028.1         O         O         O         O         O           WNK1         NP_061852.3         O         O         O         O         O           WNK2         WNK2         NP_006639.3         O         O         O         O           WNK3         WNK3         NP_065973.2         O         O         O         O           WNK4         WNK4         NP_115763.2         O         O         O         O           WNK4         WNK4         NP_115763.2         O         O         O         O           YANK4         STK32A         NP_001106195.1         O         O         O         O           YANK2         STK32B         NP_060871.1         O         O         O         O           YANK3         STK32C         NP_775846.2         O         O         O         O         O           YES         YES1         NP_005424.1         O         O         O         O         O						0						
WEE1         WE 2         WEE2         NP_001099028.1         O												
WEE2         WP_001099028.1         O         O         O           WNK1         WNK1         NP_061852.3         O         O         O           WNK2         NP_006639.3         O         O         O         O           WNK3         NP_065973.2         O         O         O         O           WNK4         WNK4         NP_115763.2         O         O         O         O           YANK1         STK32A         NP_001106195.1         O         O         O         O         O           YANK2         STK32B         NP_060871.1         O												
WNK1         WNK1         NP_061852.3         O         O         O         O           WNK2         WNK2         NP_06689.3         O         O         O         O         O           WNK3         NP_065973.2         O         O         O         O         O         O           WNK4         WNK4         NP_115763.2         O												
WNK2         WNK2         NP 006639.3         O												
WNK3         WNK3         NP_065973.2         O         O         O         O           WNK4         WNK4         NP_115763.2         O         O         O         O           YANK1         STK32A         NP_001106195.1         O         O         O         O           YANK2         STK32B         NP_060871.1         O         O         O         O           YANK3         STK32C         NP_775846.2         O         O         O         O           YES         YES1         NP_005424.1         O         O         O         O           YSK1         STK25         NP_006365.2         O         O         O         O           YSK4         YSK4         NP_079328.3         O         O         O         O           ZAK         ZAK         NP_598407.1         O         O         O         O           ZAP70         NP_997402.1         O         O         O         O         O												
YANK1         STK32A         NP_001106195.1         O         O         O         O           YANK2         STK32B         NP_060871.1         O	WNK3	WNK3	NP_065973.2							0	0	0
YANK2         STK32B         NP_060871.1         O         O         O         O           YANK3         STK32C         NP_775846.2         O												
YANK3         STK32C         NP_775846.2         O												
YES         YES1         NP 005424.1         O												
YSK1         STK25         NP_006365.2         O         O         O         O           YSK4         YSK4         NP_079328.3         O         O         O         O           ZAK         ZAK         NP_598407.1         O         O         O         O           ZAP70         NP_997402.1         O         O         O         O         O					0							
YSK4							O					
ZAK										U		
ZAP70												
					$\cap$		$\cap$					
	<u> </u>	12AF 10	[NT_00/402.1	468	97	22	135	468	18	233	469	469