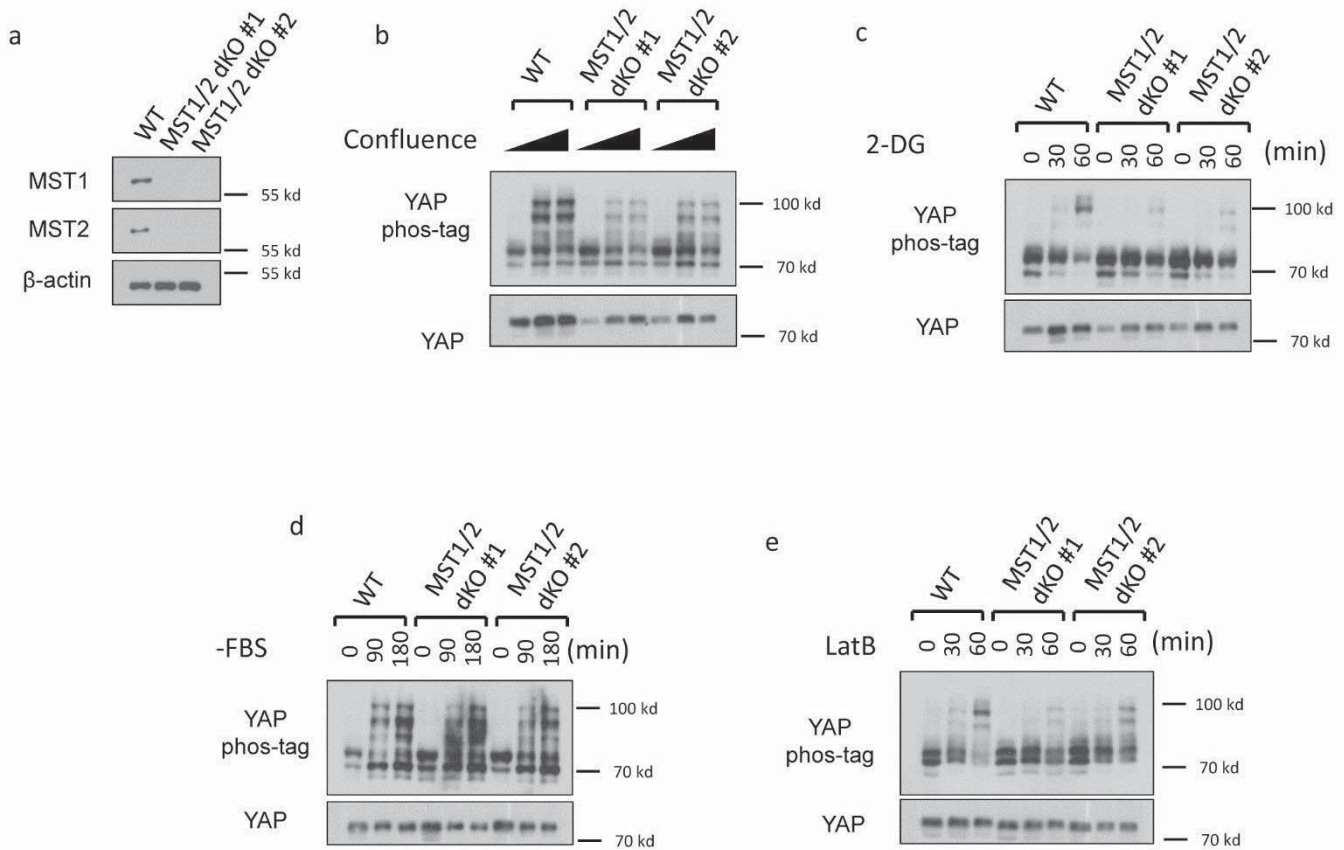


Supplementary fig. 1.

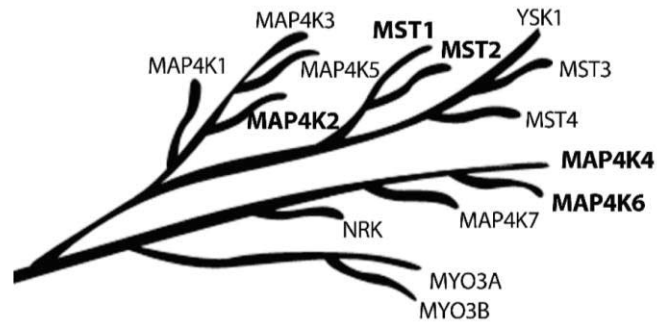


Supplementary fig. 1. Related to Figure 1. MST1/2 are not required for YAP regulation by upstream signals in U2OS cells.

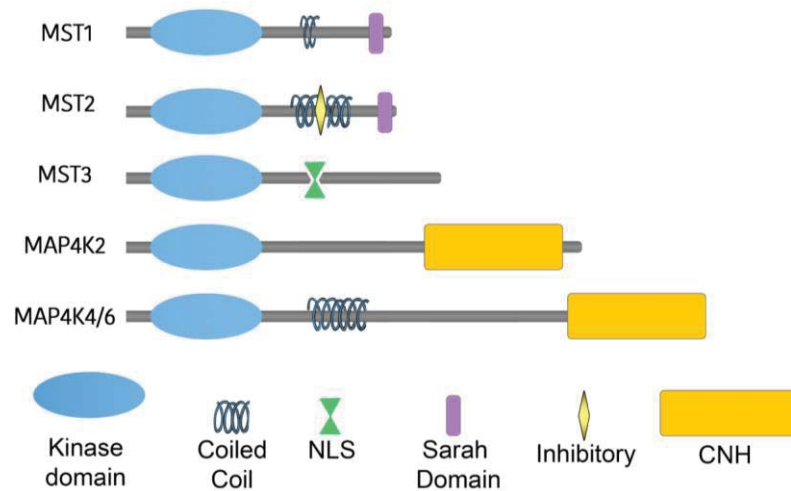
- Deletion of MST1/2 in U2OS cells.
- Contact inhibition-induced YAP phosphorylation. 1.5×10^5 , 6×10^5 , and 8×10^5 U2OS Cells per well were seeded onto 6-well plates. The cells were harvested for immunoblot 24 hours later.
- 2-DG induced YAP phosphorylation. Cells at low confluence were treated with 25 mM 2-DG for 30 or 60 minutes.
- Serum depletion-induced YAP phosphorylation. U2OS cells at low confluence were incubated with serum-free culture medium for 1.5 or 3.0 hours.
- Latrunculin B (LatB)-mediated YAP phosphorylation. Cells at low confluence were treated with 0.2 μ g/ml LatB for 30 or 60 minutes.

Supplementary fig. 2.

a



b

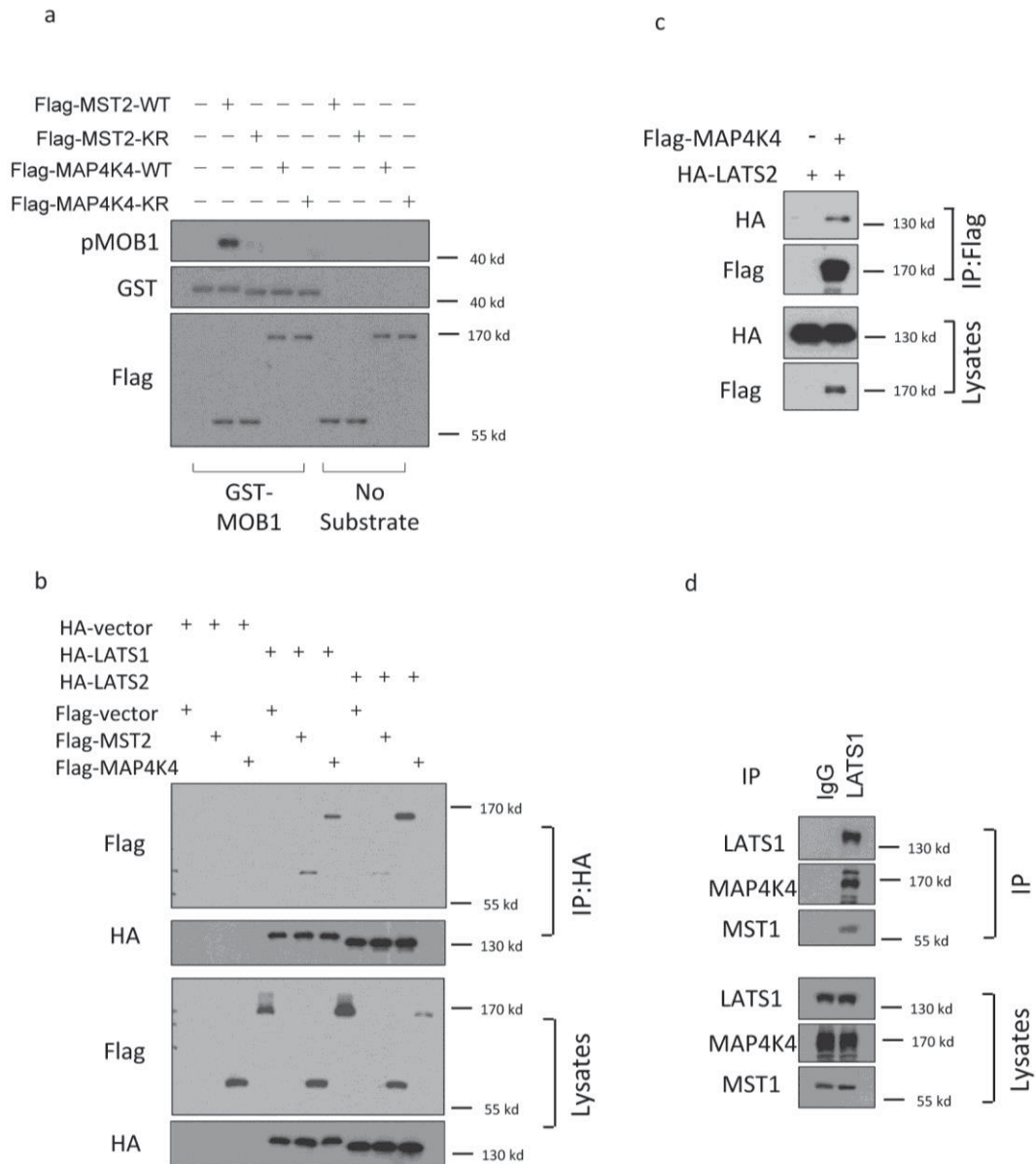


Supplementary fig. 2. Related to Figure 2. MAP4Ks are structurally related to MST1/2 and interact with LAT1/2.

(a) The branch of the human kinome tree where MST1/2 and MAP4K2/4/6 are located.

(b) Protein structures of MST1/2/3, MAP4K2, and MAP4K4/6. NLS, nuclear localization signal. CNH, citron homology domain.

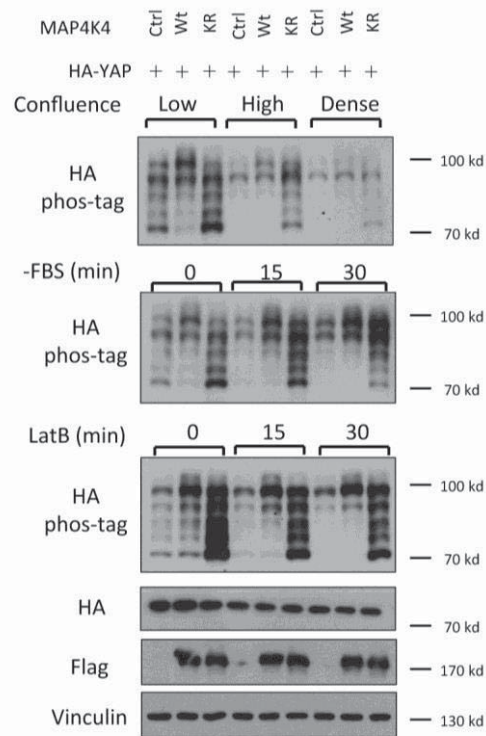
Supplementary fig. 3.



Supplementary fig. 3. Related to Figure 2. MAP4K4 is a specific kinase for LATS1/2.

- (a) MAP4K4 does not significantly phosphorylate MOB1. Flag-tagged MST2 and MAP4K4 were expressed and purified from HEK293A cells. GST-tagged MOB1 was purified from E coli.
- (b) Interaction of MAP4K4 and LATS1/2 shown by immunoprecipitation of LATS1/2. The plasmids for HA-tagged LATS1 or LATS2, and Flag-tagged MST2 or MAP4K4 were transfected into HEK293A cells. HA antibodies were used to immunoprecipitate HA-LATS1/2.
- (c) Interaction of MAP4K4 and LATS2 shown by immunoprecipitation of MAP4K4. The plasmids Flag-tagged MAP4K4 and HA-tagged LATS2 were co-transfected into HEK293A cells. Flag antibodies were used for immunoprecipitation.
- (d) Interaction of endogenous LATS1 and MAP4K4. LATS1 antibody and rabbit IgG were used for immunoprecipitation of HEK293A cell lysates.

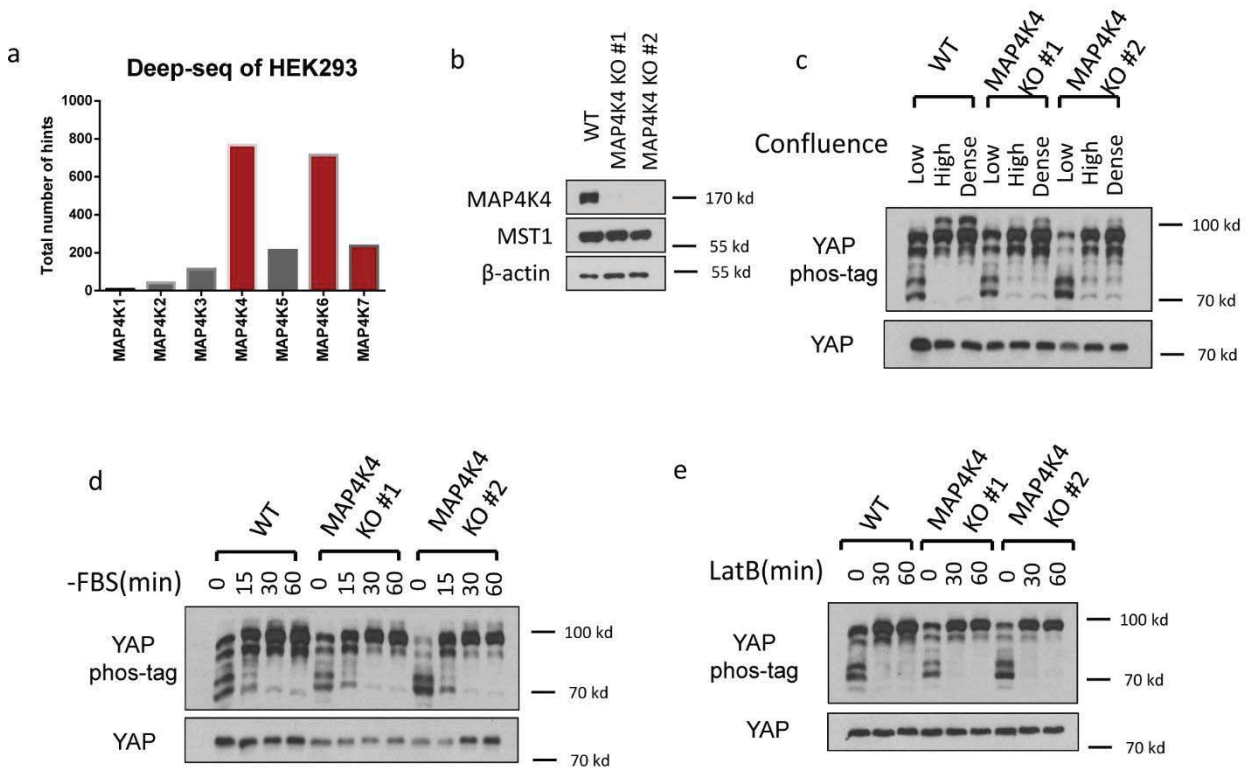
Supplementary fig. 4.



Supplementary fig. 4. Related to Figure 3. MAP4K4 induces YAP phosphorylation while its dominant-negative form promotes YAP dephosphorylation.

The Kinase dead mutant (KR) and wild-type (WT) MAP4K4 plasmids were transfected to HEK293A cells. 6 hours after transfection, the cells were split and seeded for testing effects of different conditions on YAP phosphorylation, including high cell density, serum deprivation, and LatB treatment.

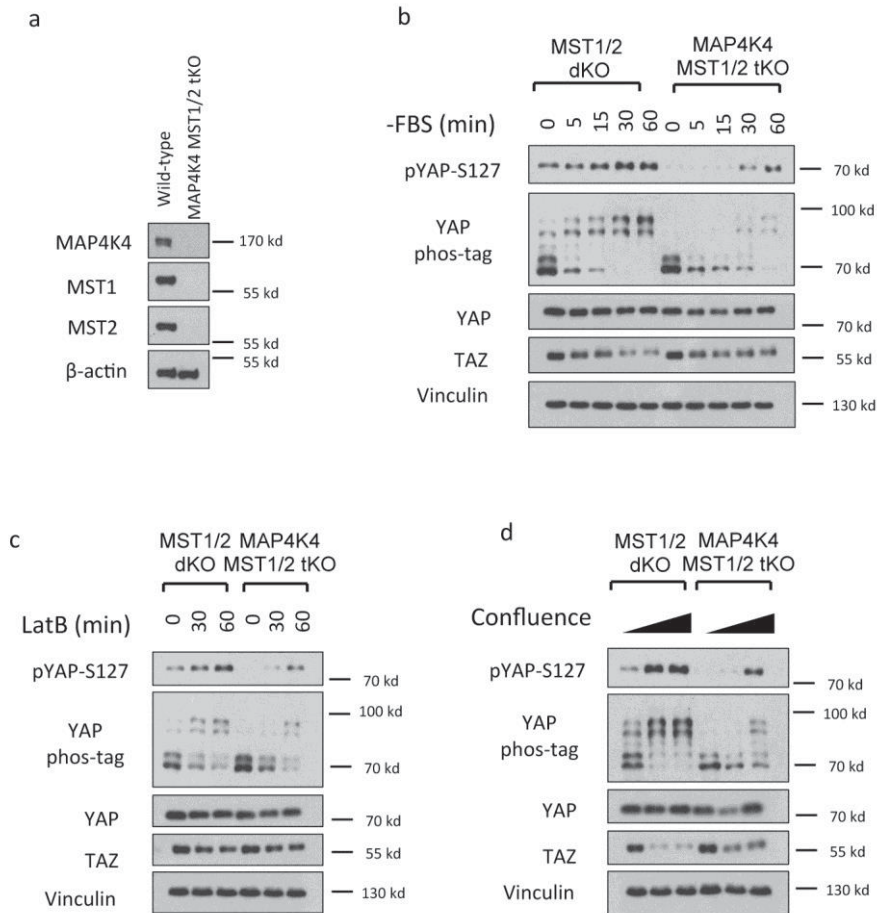
Supplementary fig. 5.



Supplementary fig. 5. Related to Figure 4. Deleting MAP4K4 alone is not sufficient to compromise any YAP phosphorylation signals.

- The RNA abundance of MAP4Ks based on a deep sequencing of HEK293A cells (Ref 35).
- Deletion of MAP4K4 in the HEK293A cells. Two independent clones (#1 and #2) are shown.
- Contact inhibition-induced YAP phosphorylation in two different clones of MAP4K4 KO cells.
- Serum deprivation-induced YAP phosphorylation in MAP4K4 KO cells.
- Actin depolymerization-induced YAP phosphorylation in MAP4K4 KO cells.

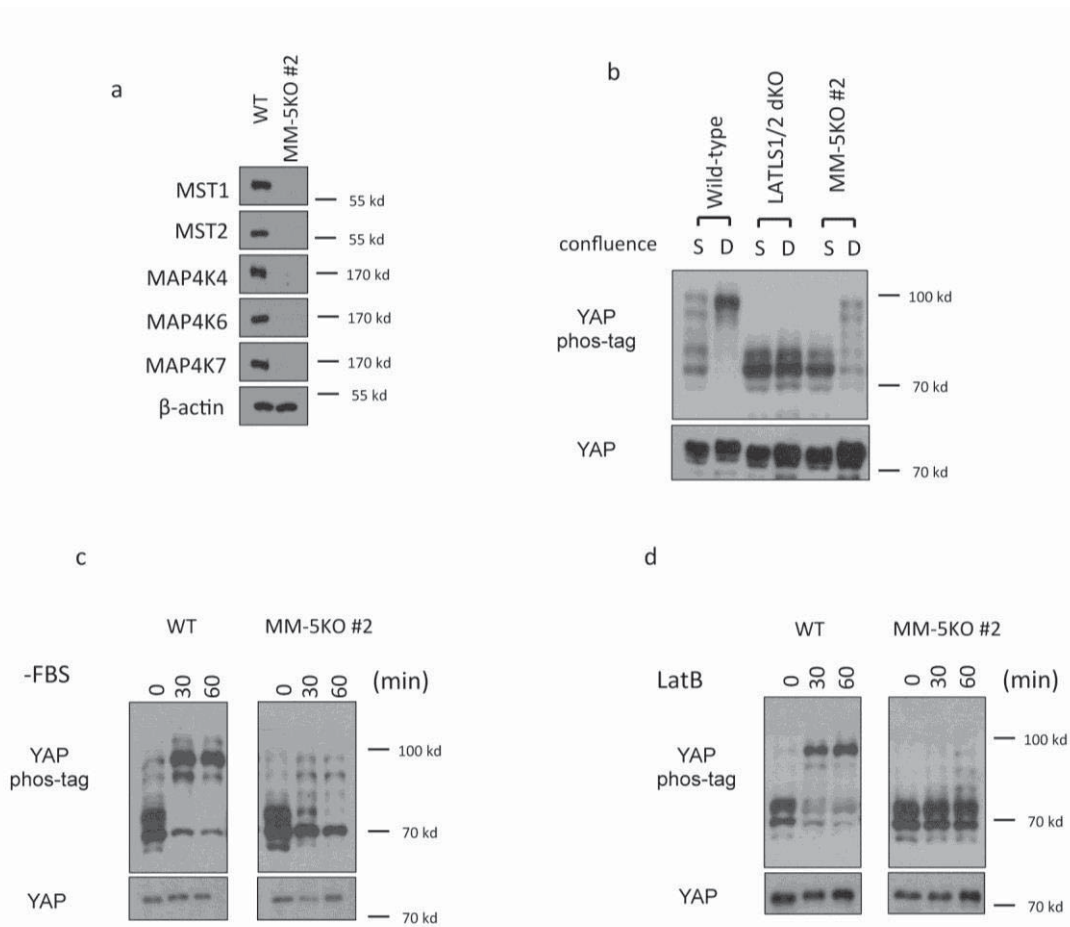
Supplementary fig. 6.



Supplementary fig. 6. Related to Figure 5. Deletion of MAP4K4 in MST1/2-dKO cells decreases and delays YAP phosphorylation induced by various signals.

- (a) Deletion of MAP4K4 in MST1/2-dKO HEK293A cells.
- (b) Decreased YAP phosphorylation from serum deprivation by deletion of MAP4K4 in MST1/2 dKO cells.
- (c) Deletion of MAP4K4 further lowers the responsiveness of YAP phosphorylation to LatB in MST1/2 dKO cells.
- (d) Deletion of MAP4K4 antagonizes contact inhibition-induced YAP phosphorylation and TAZ degradation.

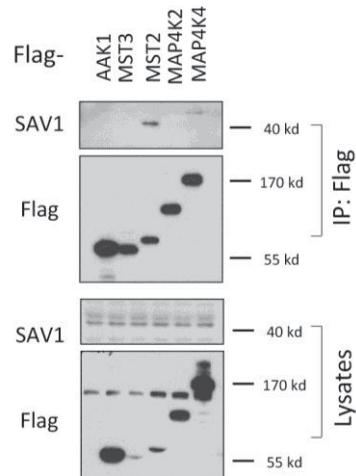
Supplementary fig. 7.



Supplementary fig. 7. Related to Figure 5. Another clone with deletion of MST1/2 and MAP4K4/6/7 shows similar responses to various upstream signals.

- A separate clone with deletion of MST1/2 and MAP4K4/6/7 using different sgRNAs for each of MAP4K4/6/7 in HEK293A cells.
- Contact inhibition-induced LATS and YAP phosphorylation is significantly compromised in the second clone of MM-5KO HEK293A cells. The experiments were performed along with the samples in Figure 5b in the same phos-tag gel. Therefore, the wild-type control cells were omitted.
- Serum deprivation-induced LATS and YAP phosphorylation is largely blocked in MM-5KO HEK293A cells.
- Actin depolymerization-induced LATS but not YAP phosphorylation is largely blocked in MM-5KO HEK293A cells.

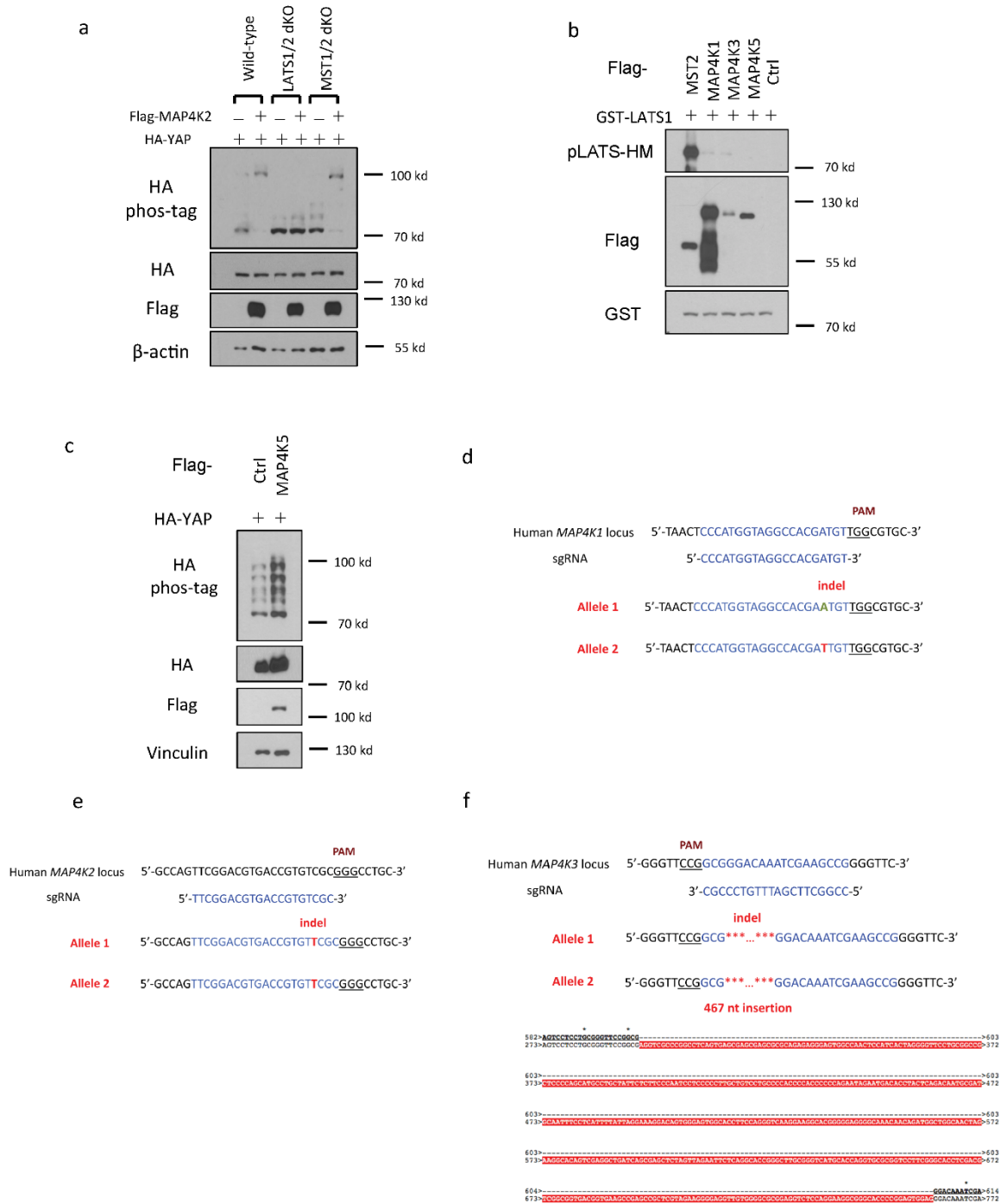
Supplementary fig. 8.



Supplementary fig. 8. Related to Figure 6. MAP4K2/4 does not interact with SAV1.

Flag-tagged kinase constructs were transfected into HEK293A cells. Cell lysate was used for immunoprecipitation with anti-Flag antibody. The immunoprecipitates were subjected to Western blot analyses with SAV1 antibody.

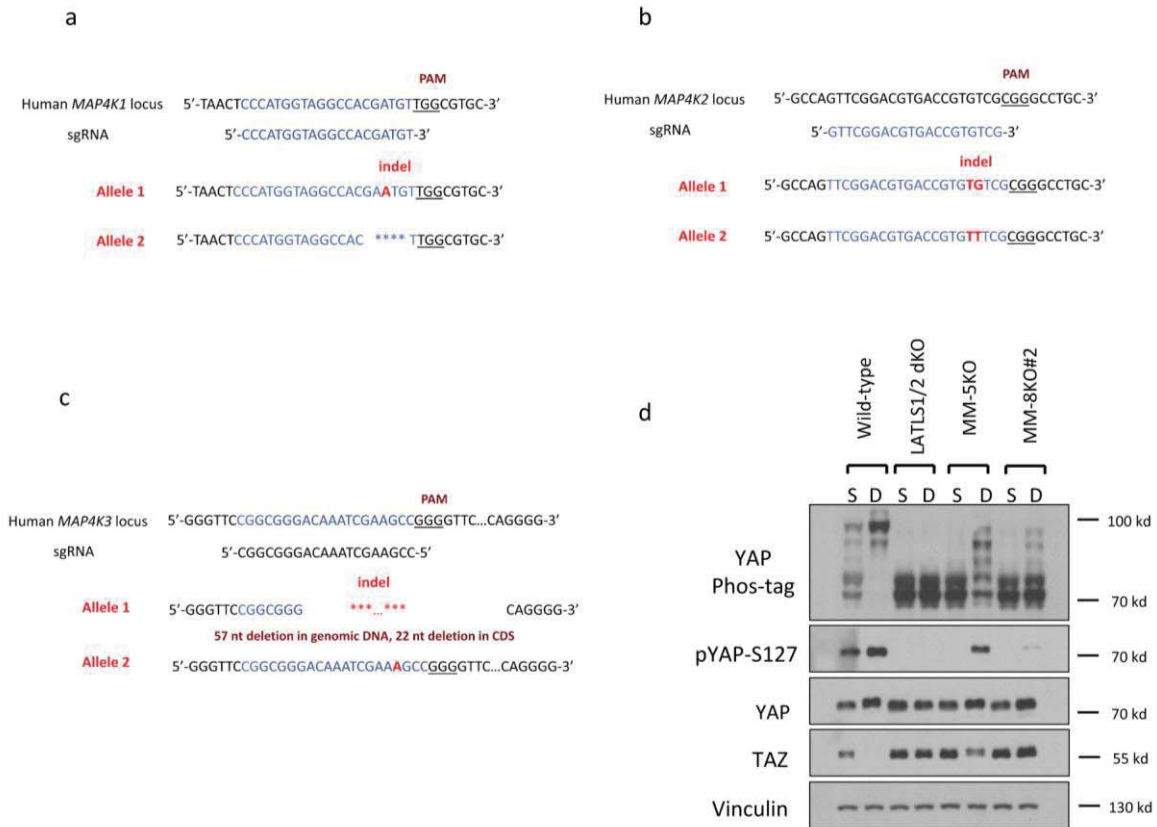
Supplementary fig. 9.



Supplementary fig. 9. Related to Figure 7. MAP4K1/2/3 are also kinases for LATS and induce YAP phosphorylation.

- (a) MAP4K2 induces YAP phosphorylation in a LATS1/2 dependent but MST1/2 independent manner.
- (b) MAP4K1 and MAP4K3 show very weak kinase activities toward LATS.
- (c) MAP4K5 is not able to induce YAP phosphorylation.
- (d) The DNA mutation MAP4K1 in the clone of the MST1/2 MAP4K1/2/3/4/6/7 8 gene knockout (MM-8KO) cells shown in Figure S7c.
- (e) The DNA mutation MAP4K2 in the clone of the MST1/2 MAP4K1/2/3/4/6/7 8 gene knockout cells shown in Figure S7c.
- (f) The DNA mutation MAP4K3 in the clone of the MST1/2 MAP4K1/2/3/4/6/7 8 gene knockout cells shown in Figure S7c.

Supplementary fig. 10.

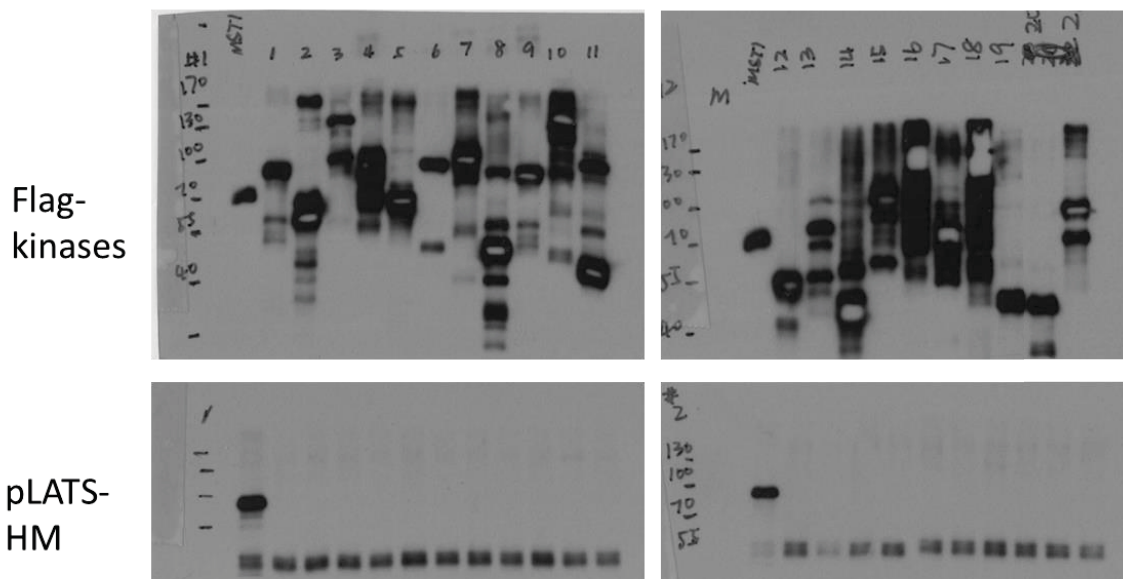


Supplementary fig. 10. Related to Figure 7. A second clone with deletion of *MST1/2* and *MAP4K1/2/3/4/6/7* showed further decreased YAP phosphorylation.

- The DNA mutation *MAP4K1* in another clone of the *MST1/2* *MAP4K1/2/3/4/6/7* 8 gene knockout (MM-8KO) cells.
- The DNA mutation *MAP4K2* in the second clone of the *MST1/2* *MAP4K1/2/3/4/6/7* 8 gene knockout cells.
- The DNA mutation *MAP4K3* in the second clone of the *MST1/2* *MAP4K1/2/3/4/6/7* 8 gene knockout cells.
- YAP phosphorylation is further decreased by deletion of *MAP4K1/2/3* in MM-5KO cells as shown in the second clone of MM-8KO cells.

Supplementary fig. 11.

Group1: 1-21

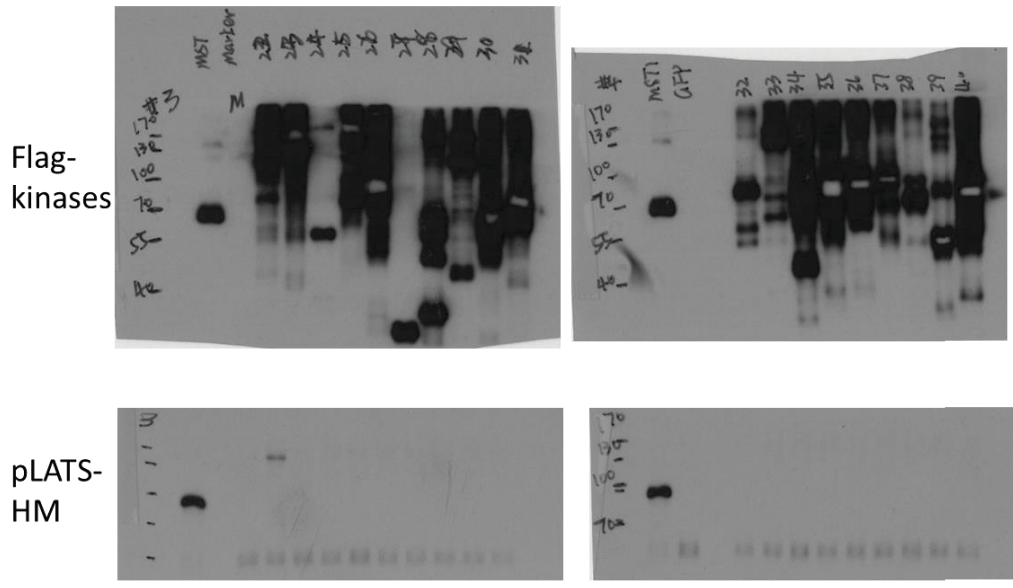


1	ARAF	12	PSKH2
2	AKT3 (isoform 2)	13	PINK1
3	PKN2	14	MAPK7 (isoform 1)
4	AMHR2	15	TNK1
5	FYN	16	AXL
6	RIPK1	17	TNK2
7	LIMK2	18	FGFR3
8	CDK9	19	CDC2L5
9	ARAF	20	PFTK2
10	TIE1	21	ITK
11	CDC2L1		

Supplementary fig. 11-18. Related to Figure 2. Identification of MST1/2-independent LATS-activating kinases.

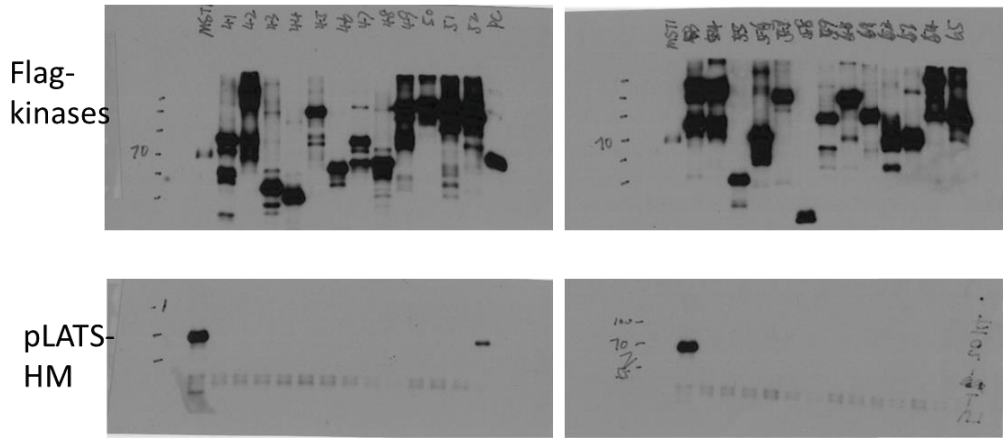
A human kinome library of kinase-expressing constructs is used for expression in HEK293A cells and purification of kinases by immunoprecipitation. The purified kinases are applied to the truncated LATS1 recombinant proteins for in vitro kinase assays. The phosphorylation of LATS1 is detected by immunoblotting with an antibody targeting the hydrophobic motif of the LATS1 kinase domain (pLATS1-T1079).

Group1: 22-40



- | | | | |
|----|-------------------|----|-------------------|
| 22 | DSTYK (isoform 2) | 32 | JAK3 |
| 23 | FGFR1 | 33 | NPR2 |
| 24 | PSKH1 | 34 | MOS |
| 25 | INSRR | 35 | HCK |
| 26 | IRAK2 | 36 | NEK11 (isoform 3) |
| 27 | RAGE | 37 | GRK4 |
| 28 | PRKACB | 38 | FASTK |
| 29 | FER | 39 | MYLK4 |
| 30 | STK11 | 40 | DYRK4 |
| 31 | TESK2 | | |

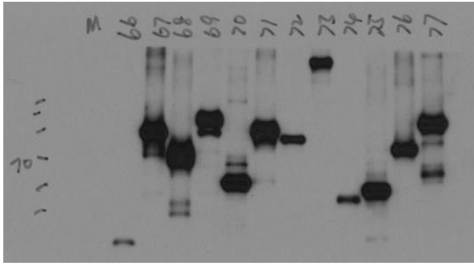
Group1: 41-65



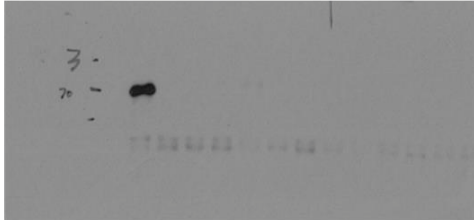
- | | | | |
|----|--------|----|---------------------|
| 41 | RAF1 | 53 | ABL1 |
| 42 | FLT4 | 54 | DDR2 |
| 43 | TSSK2 | 55 | ALS2CR7 |
| 44 | PRKY | 56 | MATK |
| 45 | KSR2 | 57 | MAST2 |
| 46 | FASTK | 58 | ULK3 |
| 47 | PRKAA1 | 59 | EIF2AK4 (isoform 3) |
| 48 | WNK1 | 60 | MYLK3 |
| 49 | EPHB6 | 61 | DCLK2 |
| 50 | EPHB1 | 62 | ACVR1C |
| 51 | NTRK1 | 63 | FRK |
| 52 | RIPK5 | 64 | MYO3B |
| | | 65 | DYRK1B |

Group1: 66-77

Flag-kinases



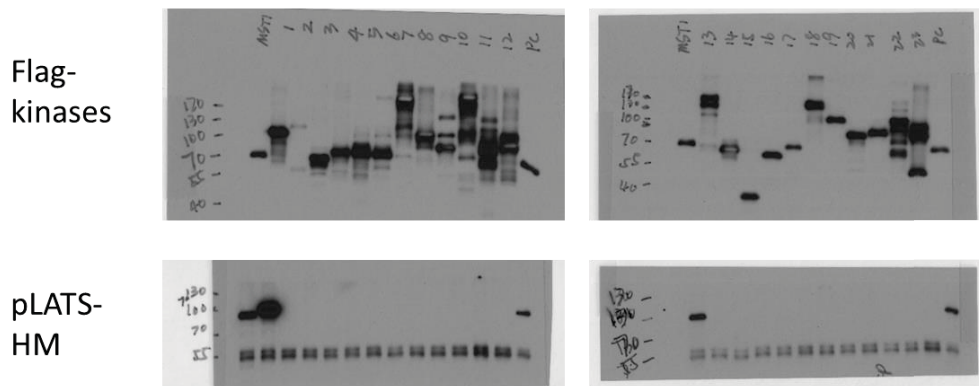
pLATS-HM



- | | |
|----|-----------------|
| 66 | ICK (isoform 2) |
| 67 | IRAK2 |
| 68 | ACVR1B |
| 69 | PTK2 |
| 70 | PRKX |
| 71 | GRK6B |
| 72 | PRKR |
| 73 | WNK4 |
| 74 | CDK7 |
| 75 | STYK1 |
| 76 | CLK3 (isoform1) |
| 77 | NEK8 |

Supplementary fig. 12.

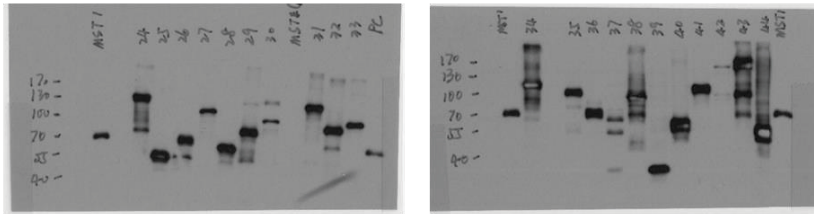
Group 2: 1-23



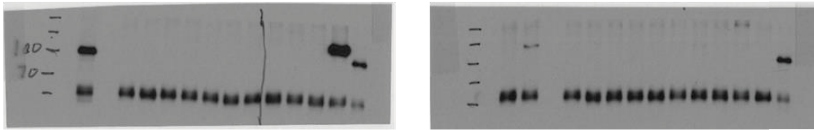
1	PRKCB	13	NTRK3
2	MERTK	14	GSK3A
3	PCTK3	15	TSSK6
4	LCK	16	MAP2K7 (isoform 4)
5	EPHB6	17	CLK3 (isoform 3)
6	LYN	18	EPHA2
7	ERBB2	19	TNNI3K
8	HIPK4	20	CAMKK1 (isoform 2)
9	GRK7	21	TESK1
10	ALK	22	DCAMKL2
11	HCK	23	PDIK1L
12	ACVR2B		

Group 2: 24-44

Flag-kinases



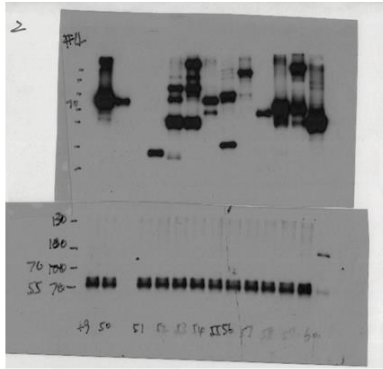
pLATS-HM



24	DDR1	34	PRKCE
25	SGK2	35	PLK2
26	SGK1	36	RIOK
27	GSG2	37	PRKACB
28	ILK	38	IKBKE
29	SRPK3	39	CDK10 (isoform 4)
30	EIF2AK4	40	TSSK1B
31	PKN1	41	NUAK2
32	BTK	42	PIK3R4
33	PRKG2	43	CSF1R
		44	STK32C (isoform 2)

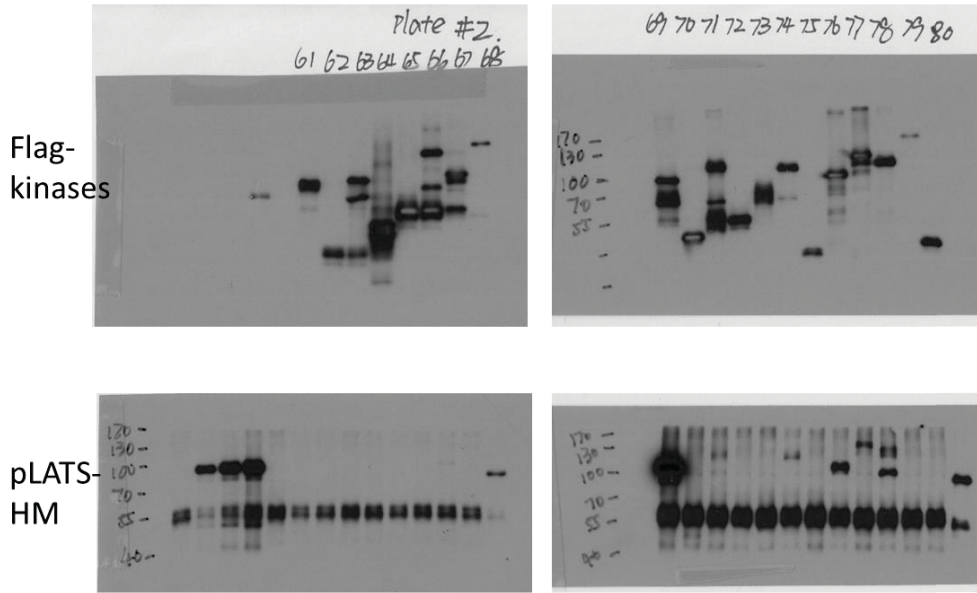
Group 2: 49-60

Flag-kinases



- 49 CAMKV
- 50 MAP3K12
- 51 MAPK15
- 52 NEK2
- 53 ERBB3
- 54 ITK
- 55 CSNK1A1
- 56 TYK2
- 57 CDK8
- 58 BLK
- 59 RET
- 60 CSNK1D (isoform 2)

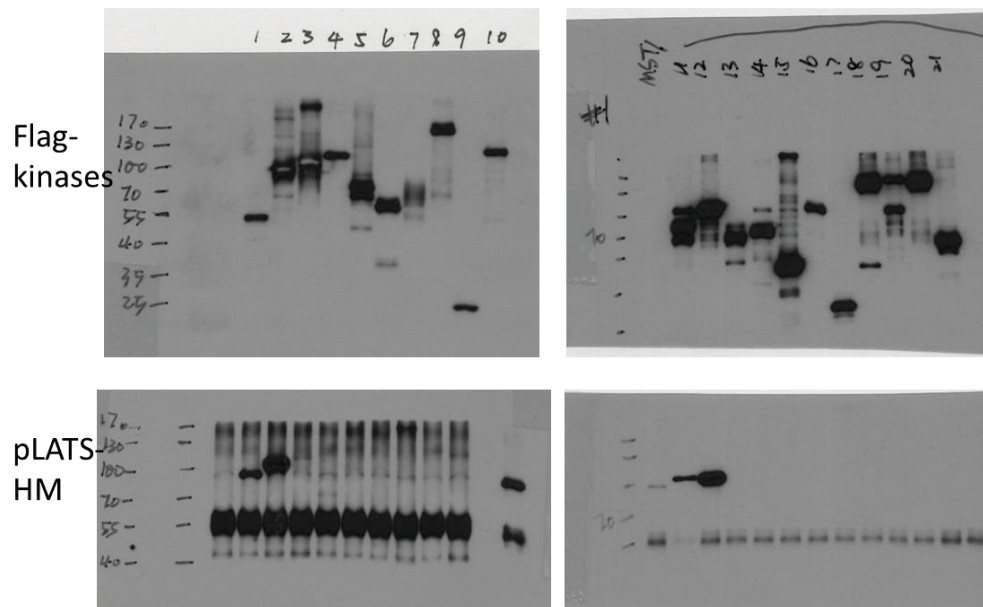
Group 2: 61-80



61	BMPR1A	69	PRKCQ
62	NRBP2 (isoform 2)	70	CDK6
63	LYN	71	MUSK
64	PIM3	72	CSNK1E
65	CSNK1D (isoform 1)	73	LCK
66	STK33	74	KSR
67	RPS6KA2	75	TP53RK
68	ALPK1	76	PRKCG
		77	PTK2B
		78	MAP4K2
		79	DKFZp761P0423 (?)
		80	CDK3

Supplementary fig. 13.

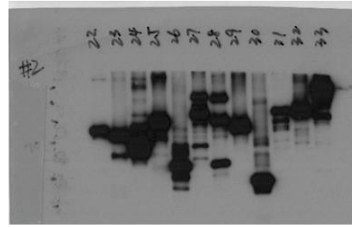
Group 3: 1-21



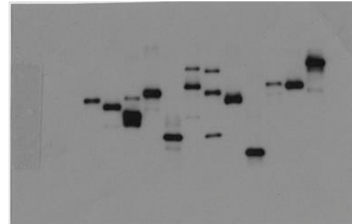
1	LOC653052(?)	11	ACVRL1
2	PRKCI	12	PRKCB1
3	PRKCH	13	SGK
4	FES	14	RPS6KL1
5	RPS6KA4	15	EPHA10
6	PTK6	16	TBK1
7	TGFBR2	17	CCRK
8	RET	18	TYRO3
9	RIPK3	19	JAK1
10	LOC91807	20	TYRO3
		21	CAMK1G

Group 3: 22-33

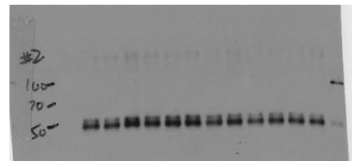
Flag-kinases



Flag-kinases
Short exposure



pLATS-HM



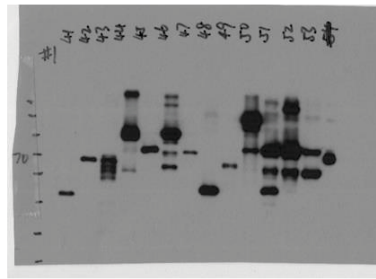
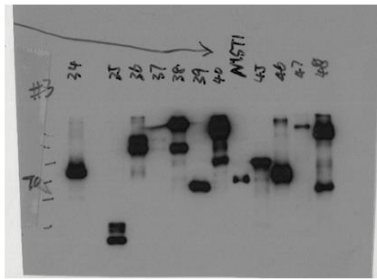
- 22 RPS6KA5 (isoform 2)
- 23 ADCK1 (isoform 2)
- 24 TSSK1
- 25 LIMK1
- 26 PRKACG
- 27 EPHB1
- 28 DSTYK (isoform 1)
- 29 FGR
- 30 CDK10 (isoform 3)
- 31 PAK6
- 32 PRKCZ
- 33 ROR2

Group 2: 45-48

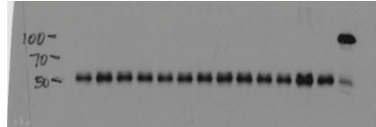
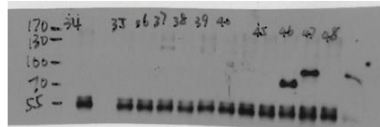
Group 3: 34-40

Group 3: 41-53

Flag-kinases



pLATS-HM



Group 2

45 MYLK2
46 RIPK2
47 MAP4K4
48 FLT3

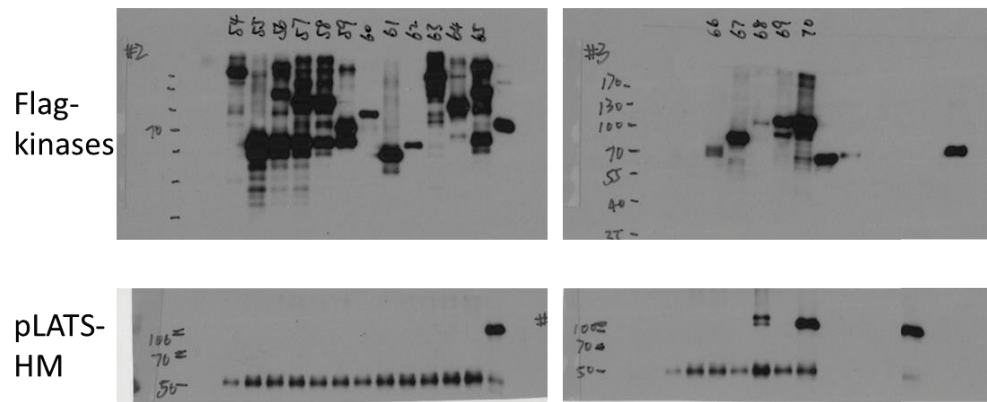
Group 3

34 IRAK3
35 TSSK4 (isoform 3)
36 EPHA1
37 STK32A
38 IGF1R
39 DAPK3
40 PDGFRB

Group 3

41 CDK4
42 EPHA3
43 CAMK2B (isoform 2)
44 BMX
45 PRKAA2
46 PFTK1
47 DYRK2 (isoform 2)
48 AURKC
49 MGC42105
50 NTRK2
51 CSNK1A1L
52 AKT1
53 UHMK1

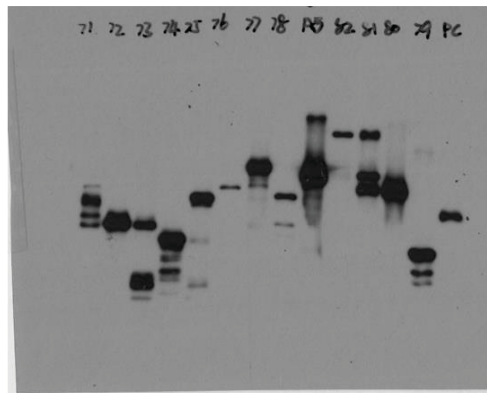
Group 3: 54-70



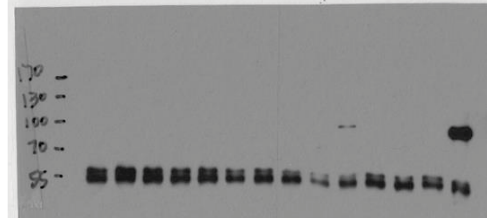
- 54 PASK
- 55 PDPK1 (isoform 4)
- 56 PHKG2
- 57 EPHA4
- 58 ADRBK1
- 59 ZAK (isoform 2, or beta)
- 60 PAK4 (isoform 1)
- 61 LOC91461
- 62 STRADB
- 63 DAPK1
- 64 PLK3
- 65 PDGFRA
- 66 TGFBR1
- 67 SGK3
- 68 EIF2AK1
- 69 MAP3K7 (isoform 1A)
- 70 PRKCA

Group 3: 71-82

Flag-kinases



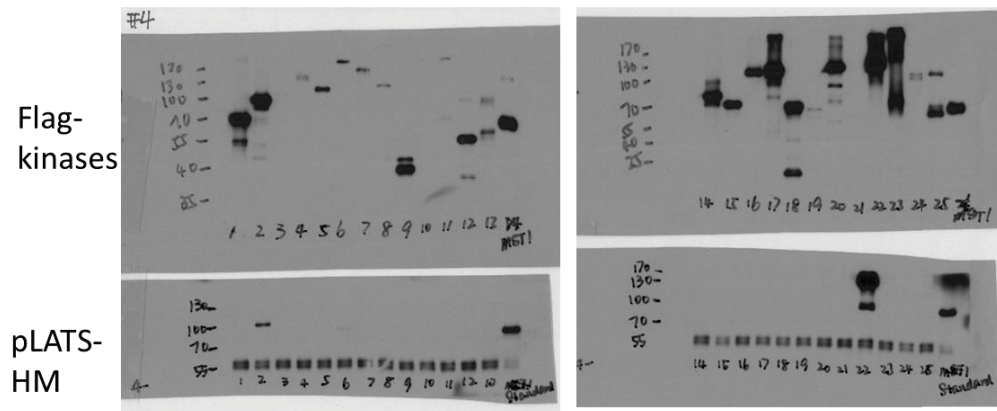
pLATS-HM



- | | |
|----|-----------|
| 71 | BMPR1B |
| 72 | PTK6 |
| 73 | LOC340371 |
| 74 | TRIB3 |
| 75 | STK38L |
| 76 | ULK4 |
| 77 | SIK1 |
| 78 | PCK2 |
| 79 | AURKB |
| 80 | GRK6A |
| 81 | EGFR |
| 82 | MINK1 |

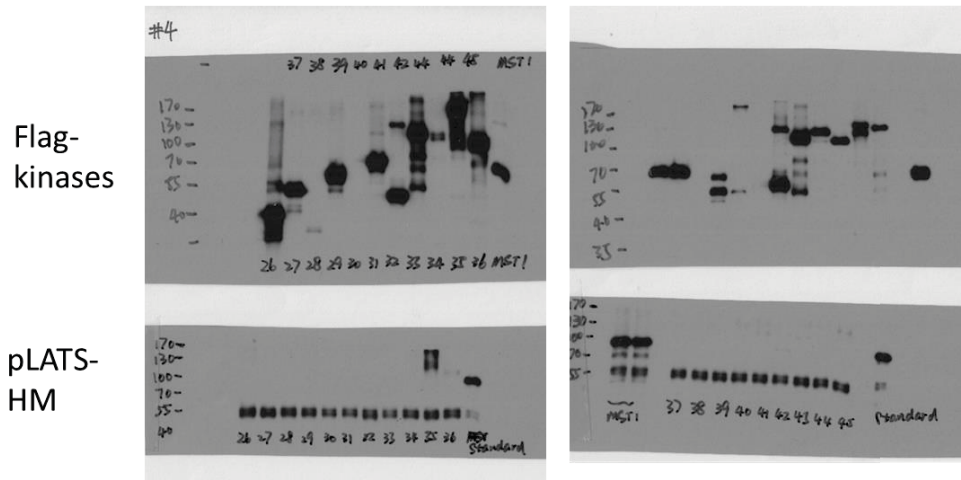
Supplementary fig. 14.

Group 4: 1-25



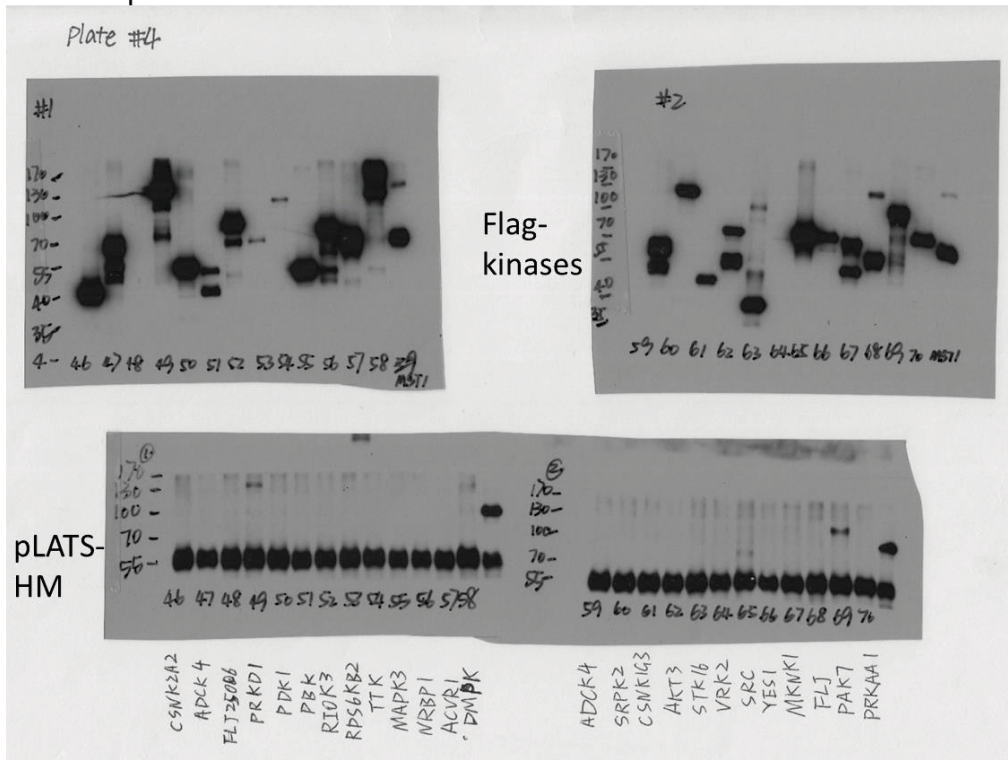
1	AKT2	14	DYRK2 (isoform 1)
2	RPS6KA1	15	PAK2
3	HIPK2	16	MAP4K3
4	ULK2	17	PRKD2
5	SRPK1	18	STK38
6	SLK (isoform 2)	19	NEK10
7	BMPR2	20	PTK2
8	PLK4	21	BRAF
9	BCKDK	22	NEK9
10	FASTKD1	23	CLK2
11	LMTK2	24	MGC16169
12	AURKA	25	CHEK1
13	ACVR2A		

Group 4: 26-45



- 26 PIM1 (isoform 2)
- 27 MAPKAPK2
- 28 TSSK3
- 29 VRK1
- 30 CDKL4
- 31 CDC2L6
- 32 DAPK2
- 33 MAP4K1 (isoform 2)
- 34 SNRK
- 35 ERBB4
- 36 BMX
- 37 FASTK (isoform 2)
- 38 ALPK2
- 39 DYRK3 (isoform 2, short)
- 40 ERN1
- 41 RPS6KC1
- 42 TAOK3
- 43 MAP4K5
- 44 MAST1
- 45 BUB1B

Group 4: 46-70

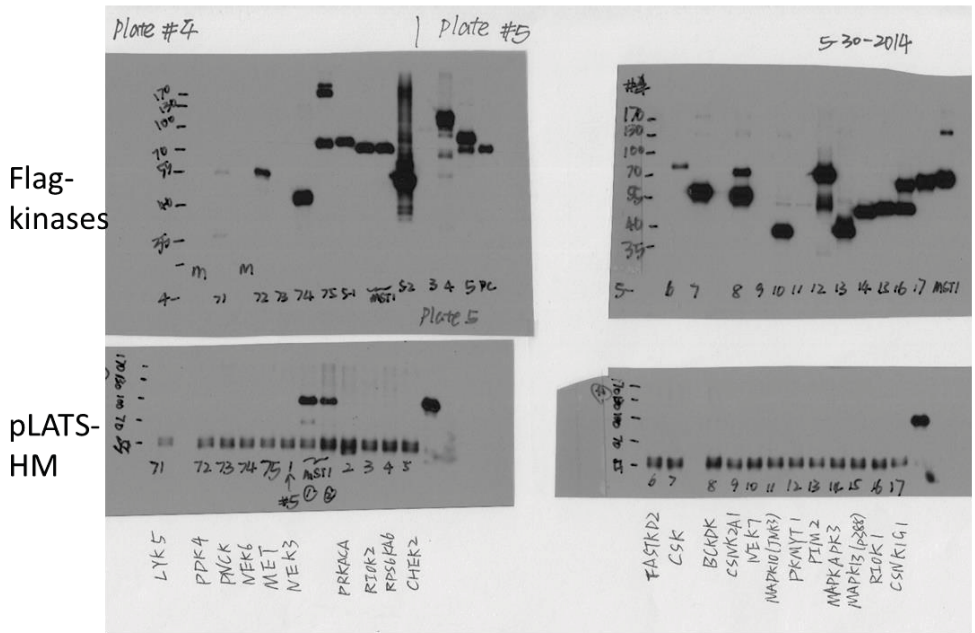


46	CSNK2A2	59	ADCK4 (isoform 2)
47	ADCK4 (isoform 1)	60	SRPK2
48	FLJ25006	61	CSNK1G3 (isoform 3 or 4)
49	PRKD1	62	AKT3 (isoform 1)
50	PDK1	63	STK16
51	PBK	64	VRK2
52	RIOK3	65	SRC
53	RPS6KB2	66	YES1
54	TTK	67	MKNK1
55	MAPK3	68	FLJ23356
56	NRBP1	69	PAK7
57	ACVR1	70	PRKAA1
58	DMPK		

Supplementary fig. 15.

Group 4: 71-75

Group 5: 1-17



Group 4

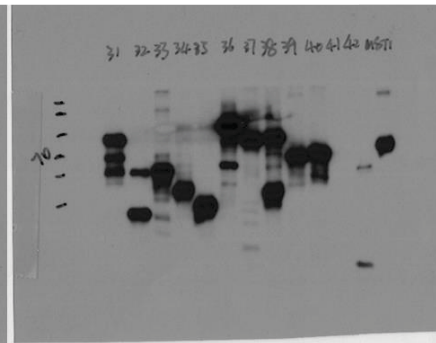
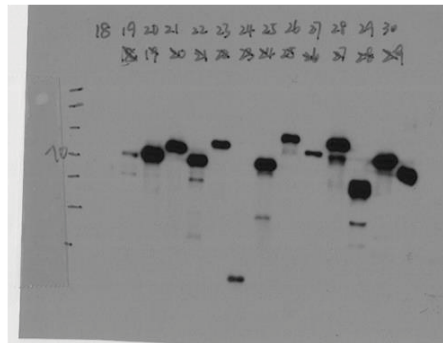
71 LYK5
72 PDK4
73 PNCK
74 NEK6
75 MET

Group 5

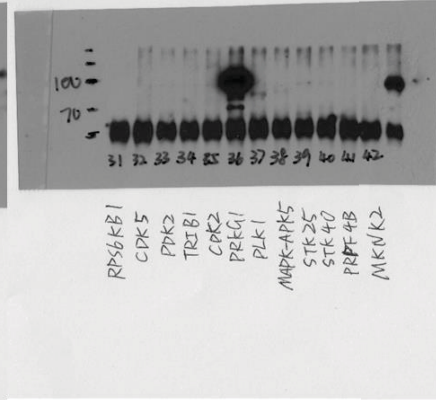
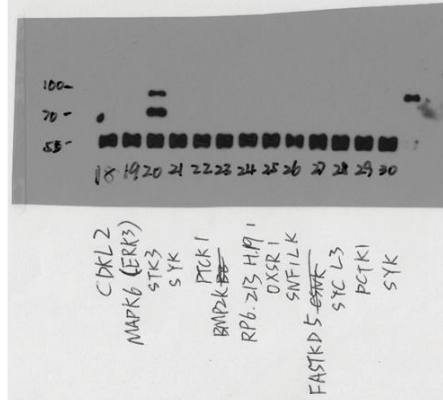
1 MAPK14
2 NEK3
3 PRKACA
4 RIOK2
5 RPS6KA6
6 CHEK2
7 FASTKD2
8 CSK
9 BCKDK
10 CSNK2A1
11 NEK7
12 MAPK10
13 PKMYT1
14 PIM2
15 MAPKAPK3
16 MAPK13
17 RIOK1

Group 5: 18-42

Flag-kinases

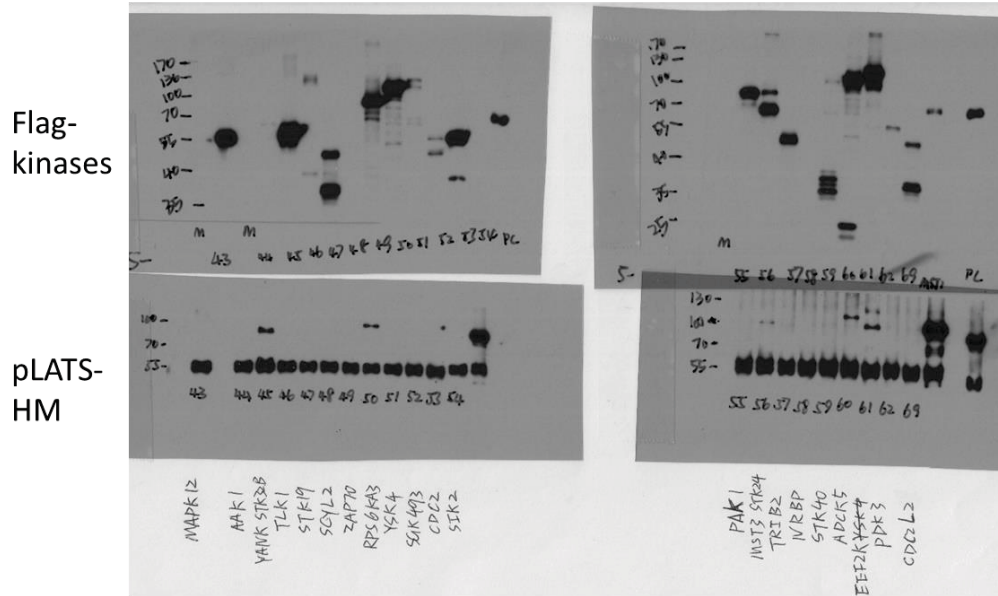


pLATS-HM



18	CSNK1G1	31	RPS6KB1
19	MAPK6	32	CDK5
20	STK3	33	PDK2
21	SYK	34	TRIB1
22	PCTK1	35	CDK2
23	BMP2K (isoform 2)	36	PRKG1
24	RP6-213H19.1	37	PLK1
25	OXSR1	38	MAPKAPK5 (isoform 2)
26	SNF1LK	39	STK25
27	FASTKD5	40	STK40
28	SCYL3 (isoform 2)	41	PRPF4B
29	PCTK1	42	MKNK2
30	SYK		

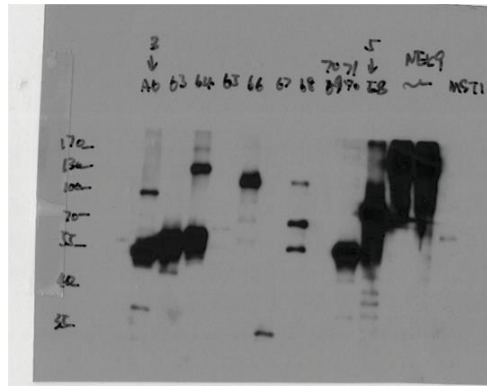
Group 5: 43-62, 69



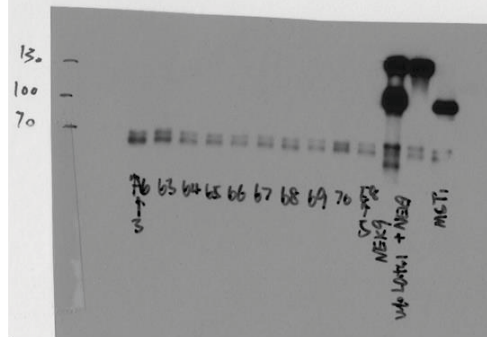
43	MAPK12	54	SIK2
44	AAK1	55	PAK1
45	STK32B	56	STK24 (isoform A)
46	TLK1	57	TRIB2
47	STK19	58	NRBP
48	SCYL2	59	STK40
49	ZAP70	60	ADCK5
50	RPS6KA3	61	EEF2K
51	YSK4 (isoform 6)	62	PDK3
52	SGK493	69	CDC2L2
53	CDC2		

Group 5: 63-68; 70,71

Flag-kinases



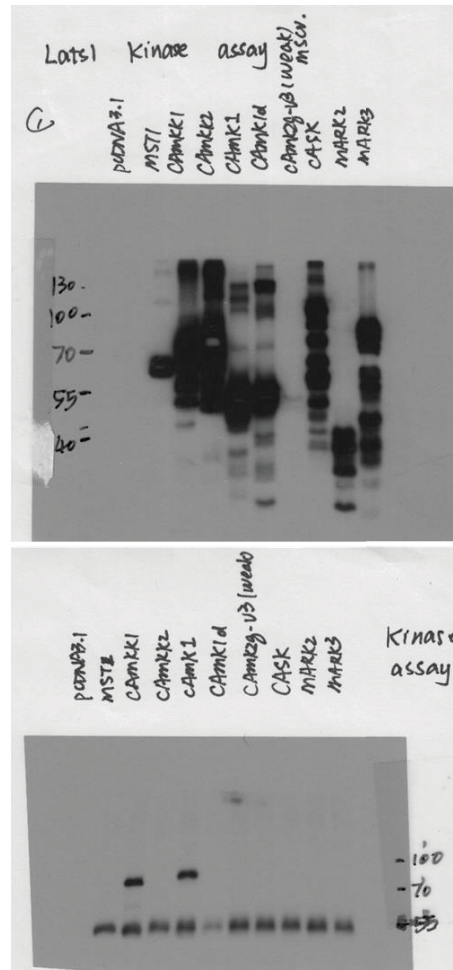
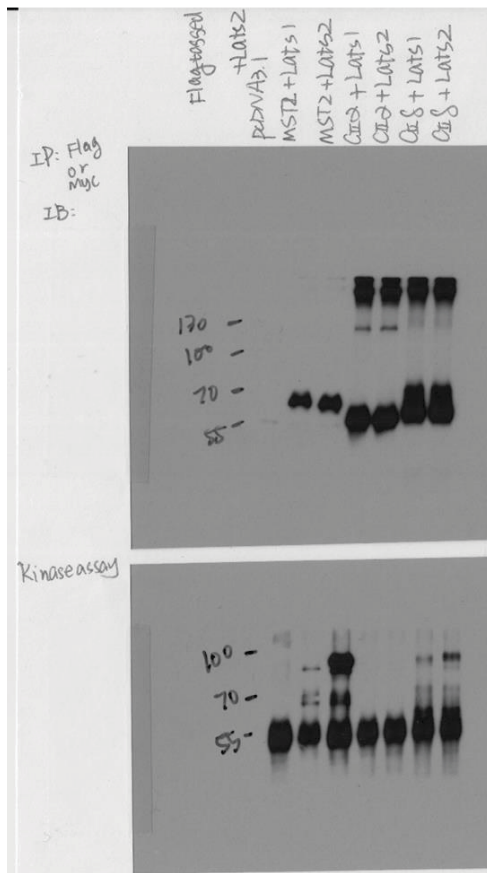
pLATS-HM



- | | |
|----|--------------------|
| 63 | PXK |
| 64 | SGK196 |
| 65 | EPHA7 |
| 66 | TLK2 |
| 67 | STRADA |
| 68 | STK19 (isoform 2) |
| 70 | BRSK2 (isoform 3?) |
| 71 | PAK4 (isoform 2) |

Supplementary fig. 16.

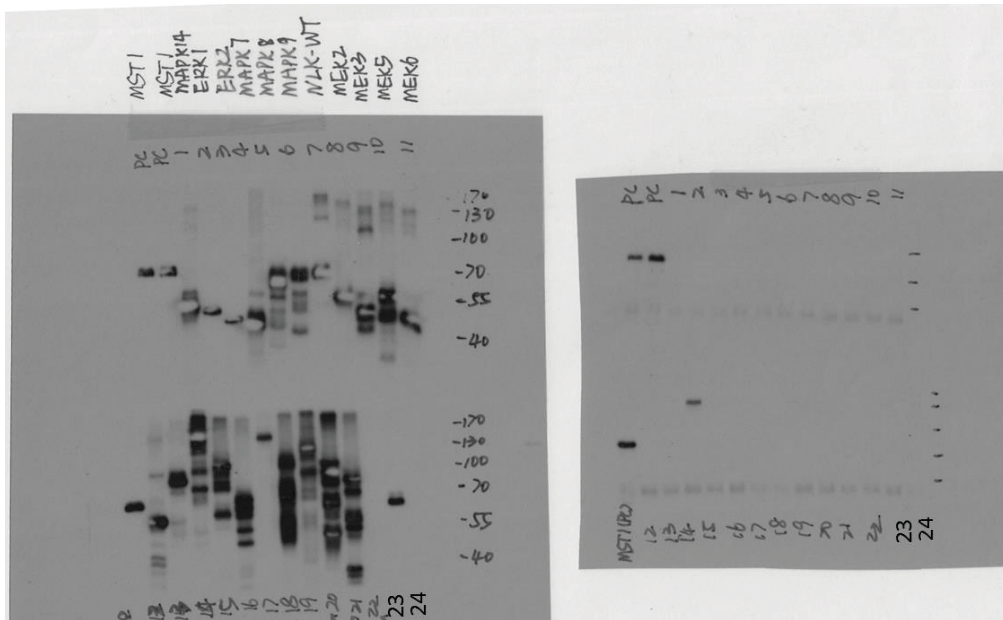
Group 6: CAMK and CAMKK



- CAMKIIA
- CAMKIID
- CAMKK1
- CAMKK2
- CAMK1
- CAMK1d
- CAMK2g-T287D
- CASK
- MARK2
- MARK3

Supplementary fig. 17.

Group 7: MAPKs

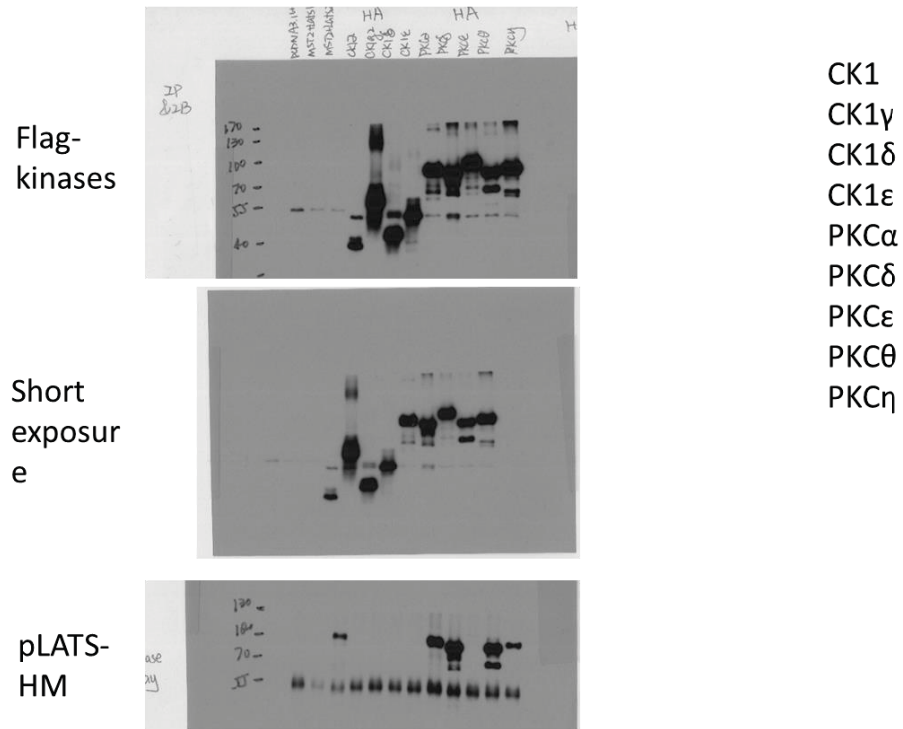


1. MAPK14-p38a
2. hERK1
3. rErk2
4. MAPK7-ERK5
5. MAPK8
6. MAPK9
7. NLK
8. MAP2K2
9. MAP2K3
10. MAP2K5
11. MAP2K6

12. MAP2K7
13. MAP3K2
14. MAP3K5
15. MAP3K6
16. MAP3K8
17. MAP3K9
18. MAP3K13
19. MAP3K14
20. MAP3K15
21. Raf-1
22. pWZL-CAMK1g
23. pWZL-CAMK2b
24. pWZL-CAMK4

Supplementary fig. 18.

Group 8: CKs and PKCs



Note that the molecular weight of pLATS-HM signals is actually the same with that of the PKC kinases.

Supplementary fig. 19. Uncropped immunoblots.

Fig 1A.

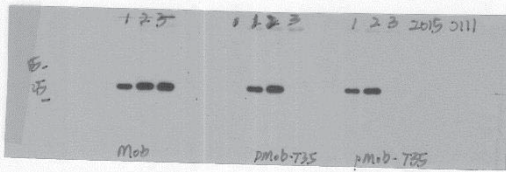
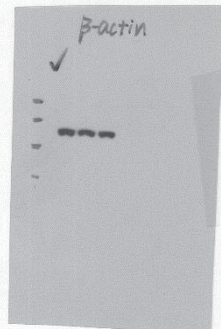
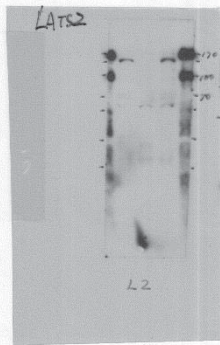
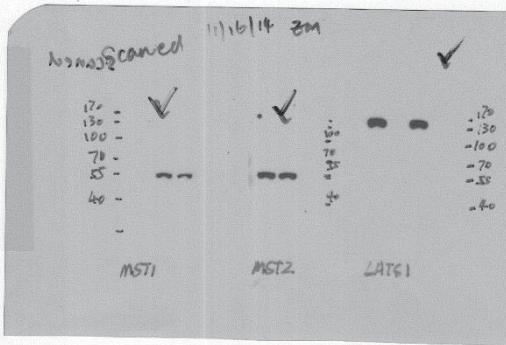


Figure 1B

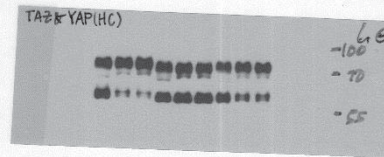
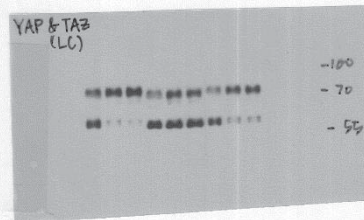
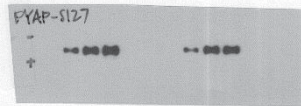
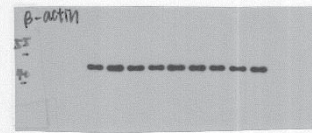
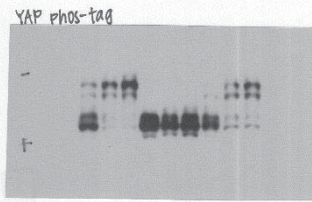


Fig. 1c.

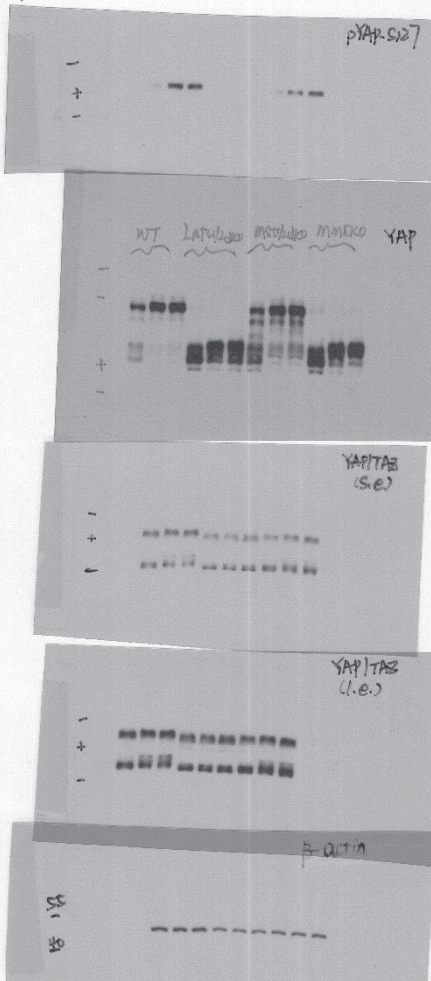
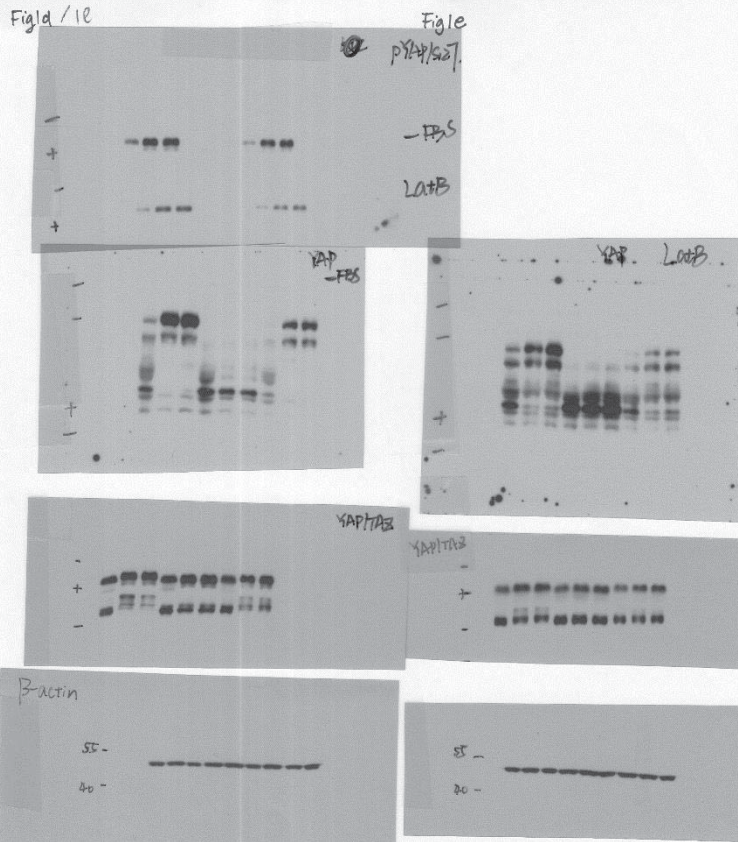
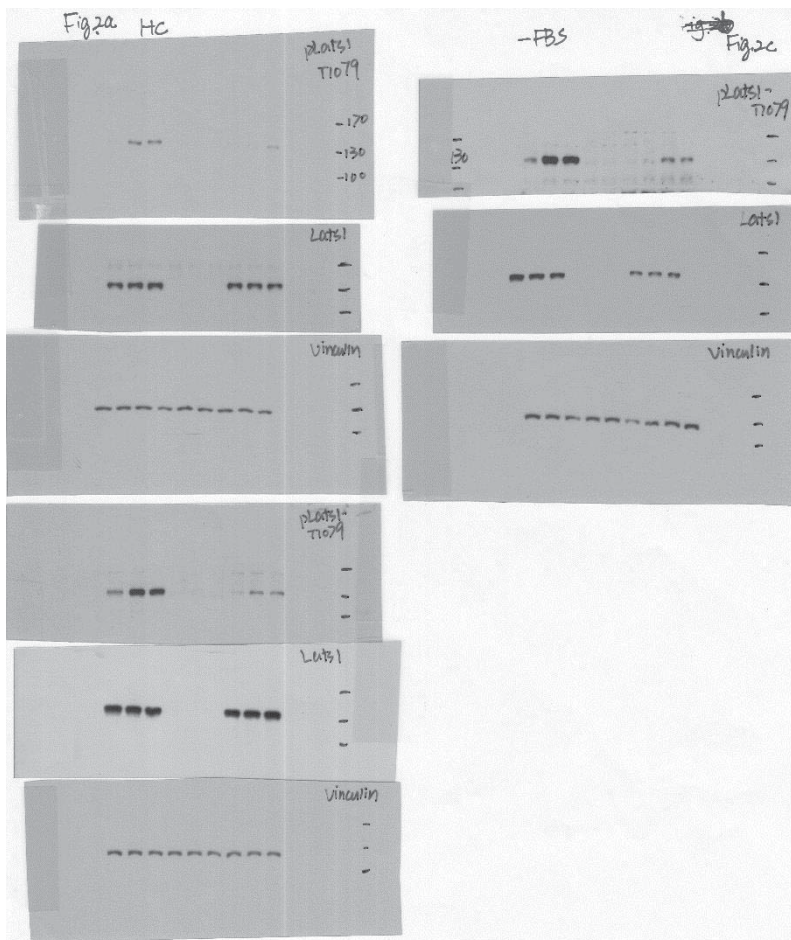


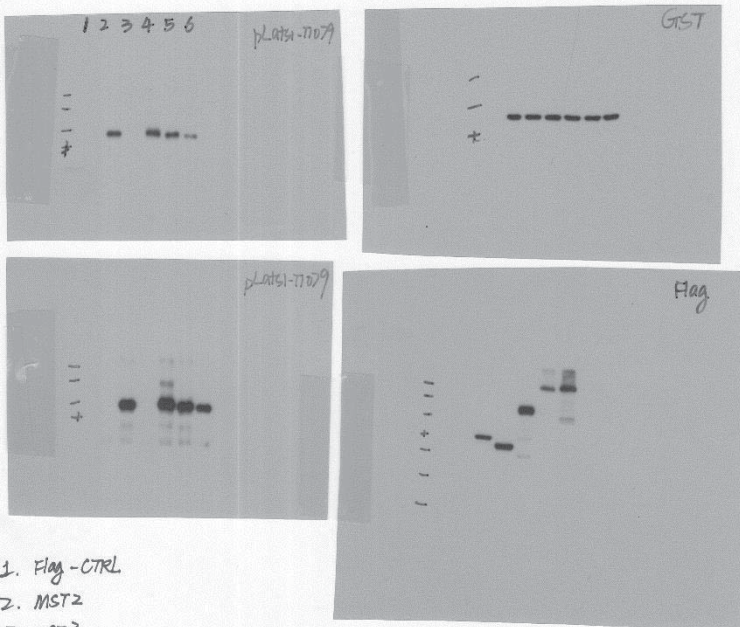
Fig. 1d / 1e





2DG Fig. 2b

Fig. 2f



1. Flag-CTRL
2. MST2
3. MST3
4. MAP4K2
5. MAP4K4
6. MAP4K6

Fig. 2g

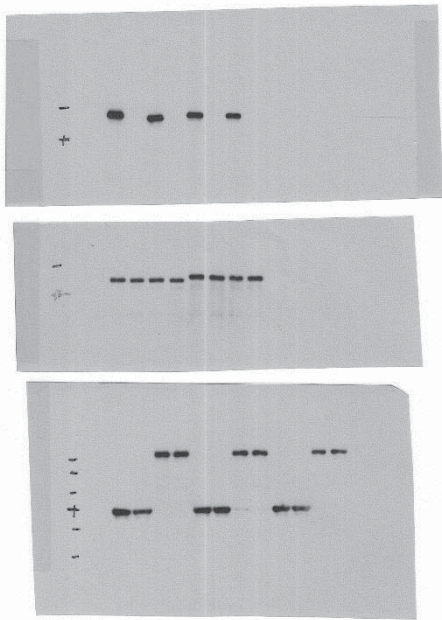


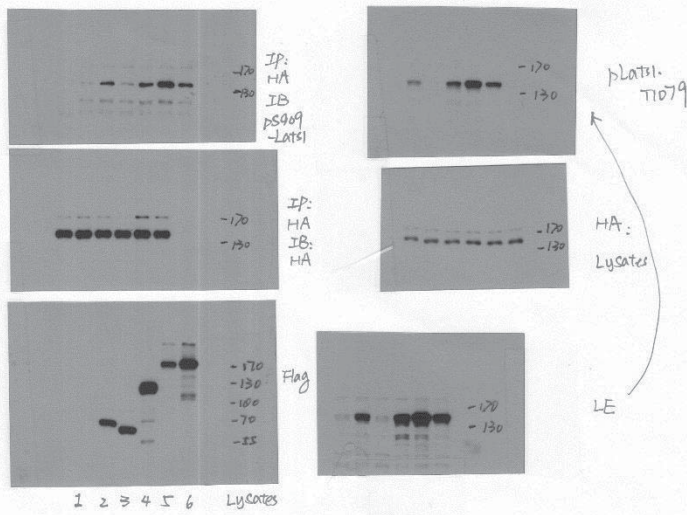
Fig. 2g

plats1-11079

GST

Flag

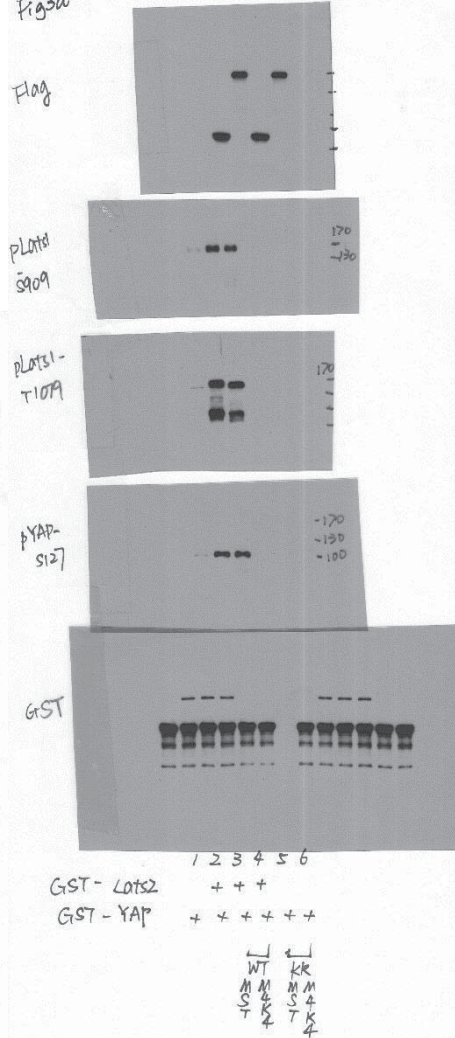
Fig. 2h



1. N-Flag
2. MST2
3. MST3
4. 4K2
5. 4K4
6. 4K6

12/3/2014

Fig. 3a



GST - LATS2 + + +
 GST - YAP + + + + +

WT KK
 M S
 S T K T
 K K

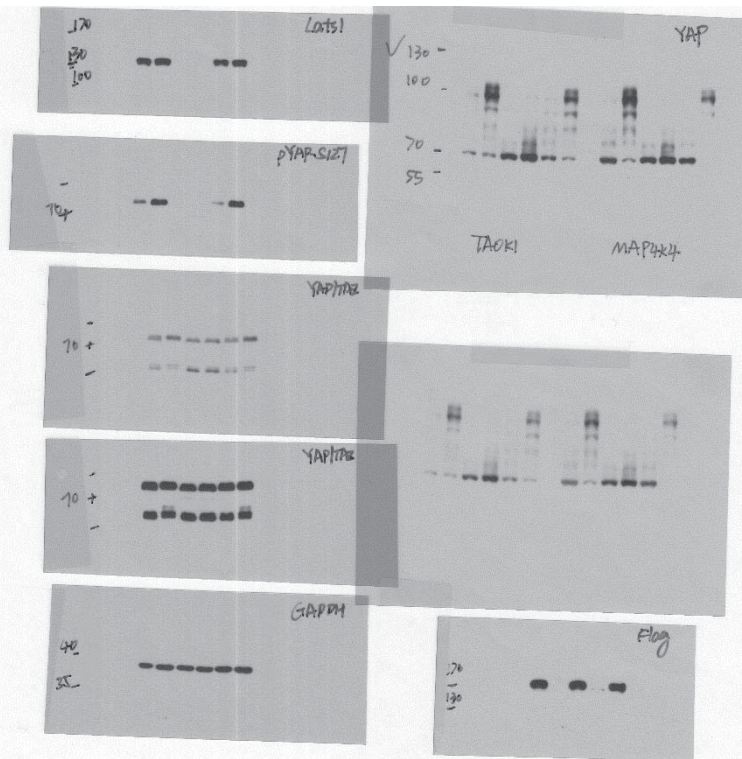
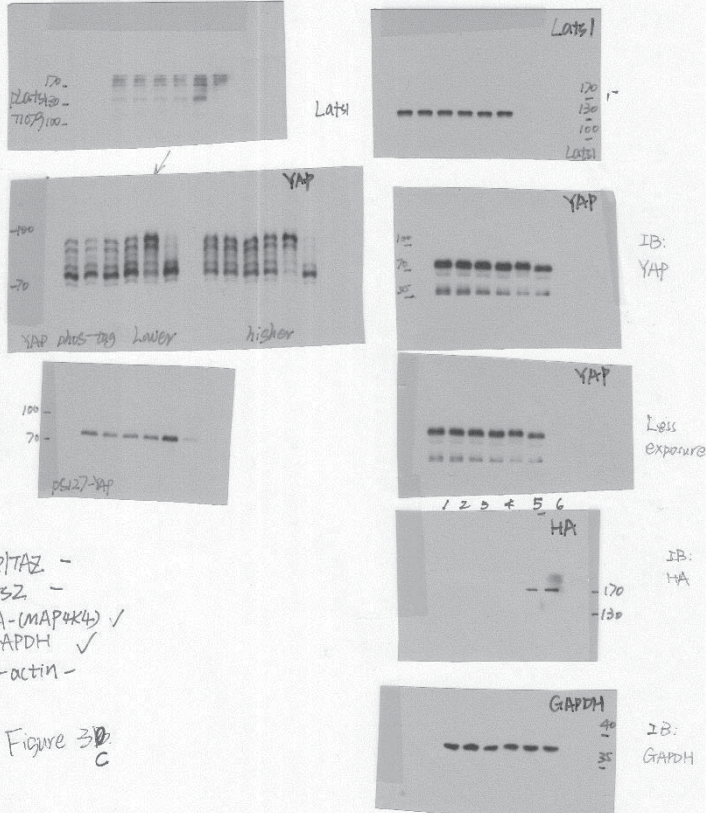


Fig 3c
 Dox - - - + + +
 CON WT KR CON WT KR HA

10/7/2014



YAP172 -
 Lats2 -
 HA-(MAP4K4) ✓
 GAPDH ✓
 β-actin -

Figure 3c

Fig 3d

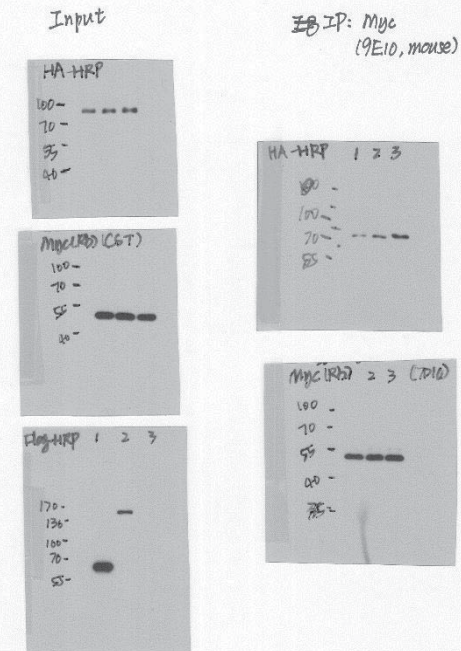


Fig. 4a

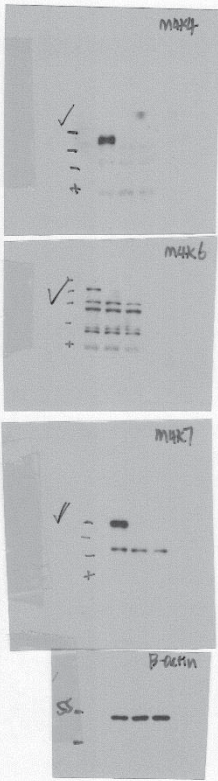


Fig. 4b

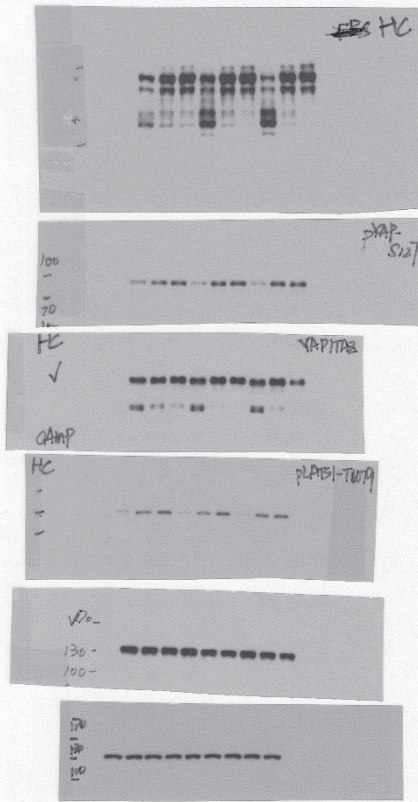


Fig. 4c

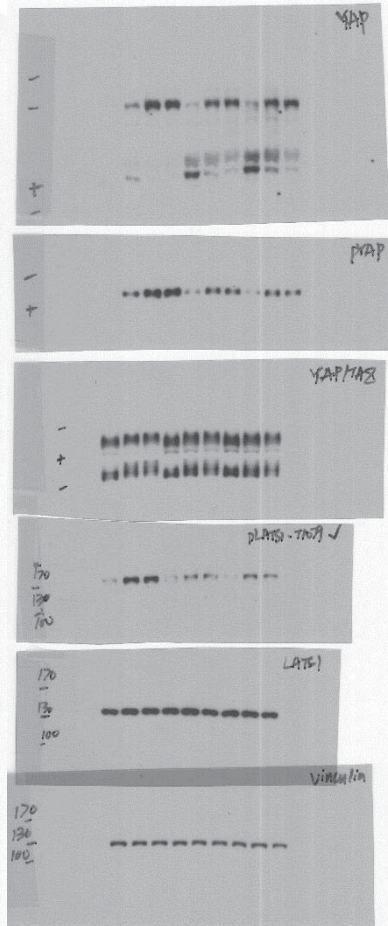


Fig. 4d

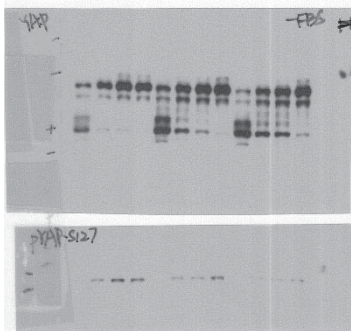


Fig. 4e

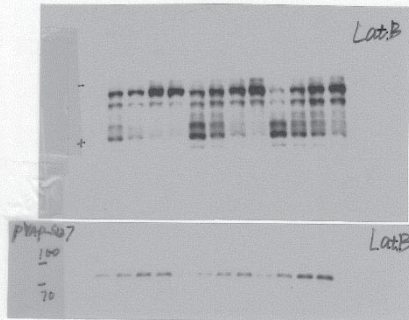


Figure 5

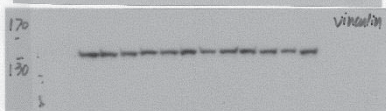
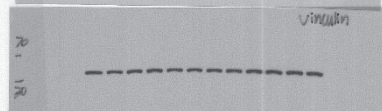
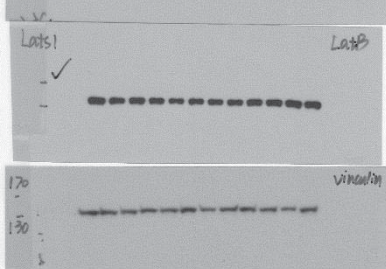
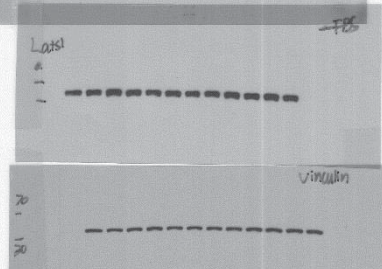
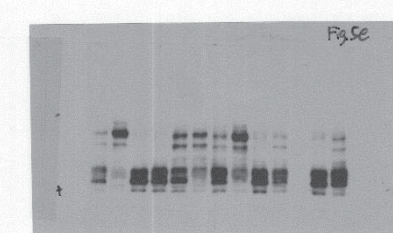
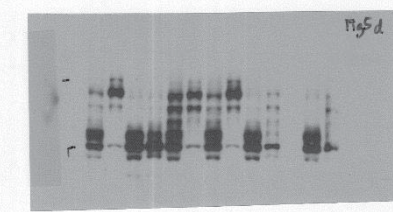
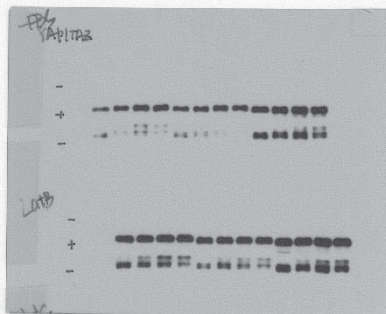
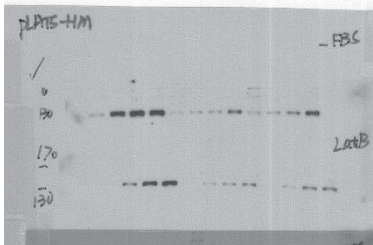
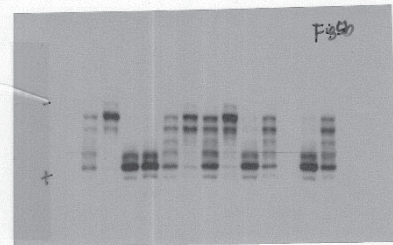


Fig 5b.

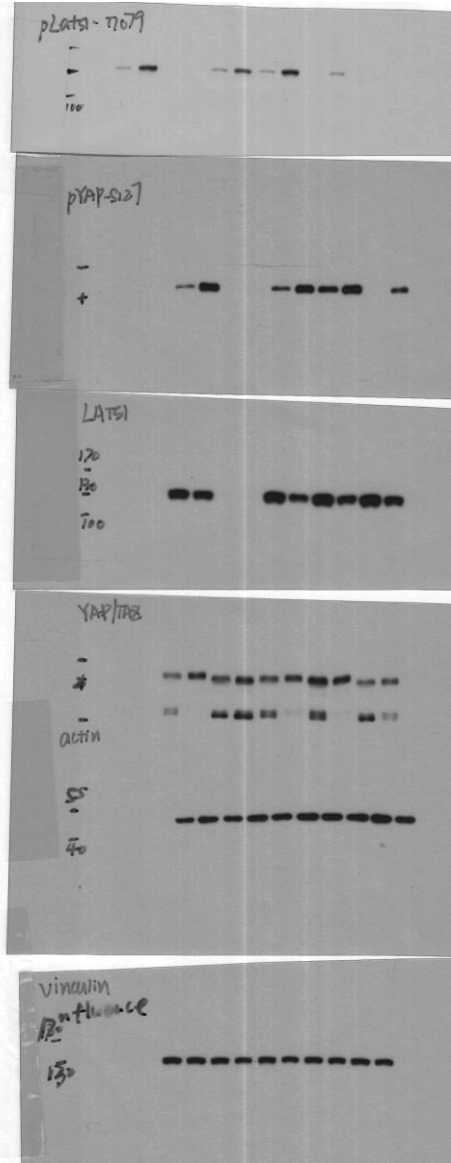
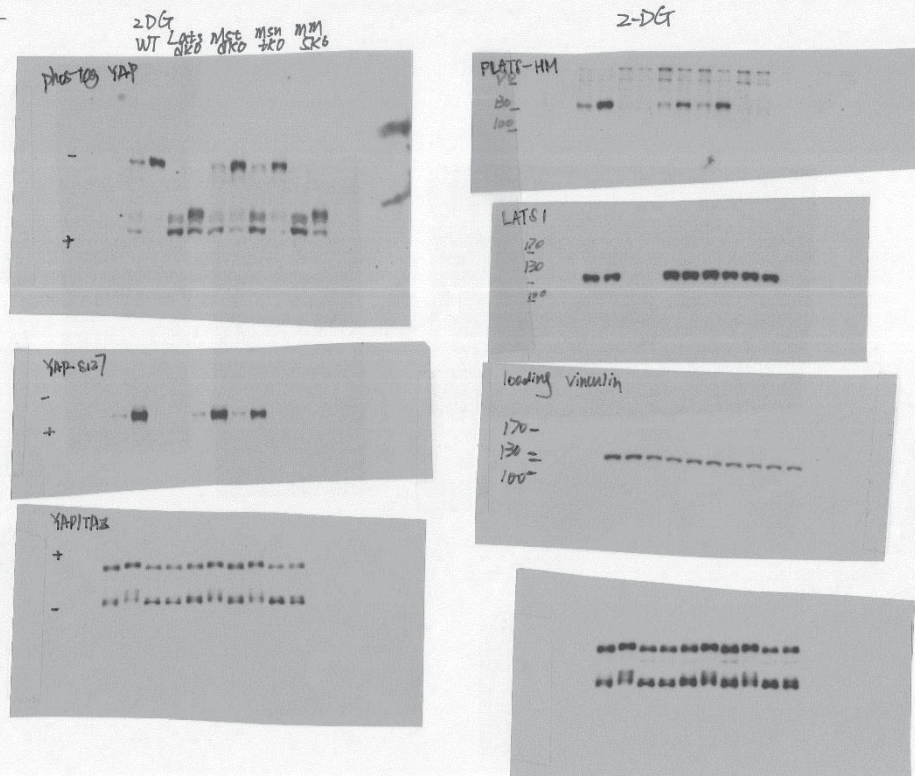


Fig 5c



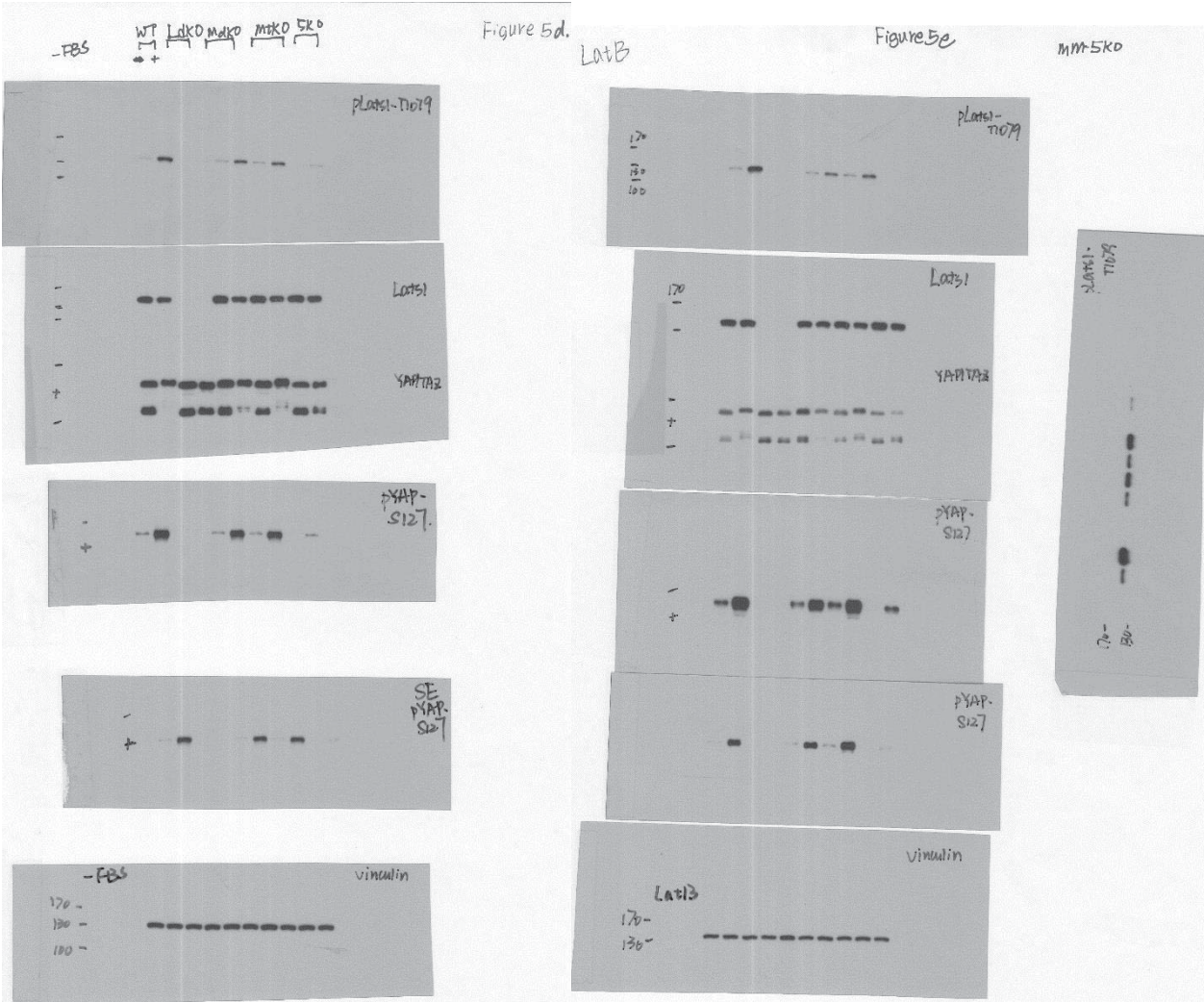


Fig. 6c & Fig. S3d

7/2/2015

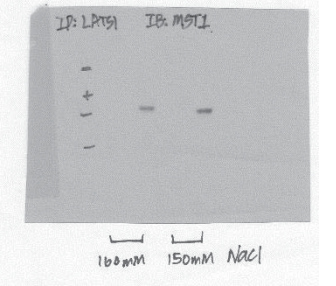
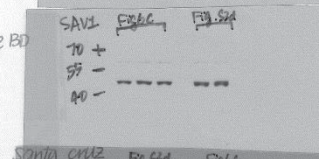
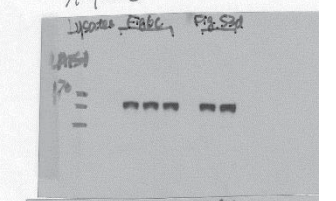
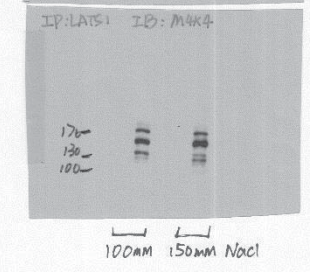
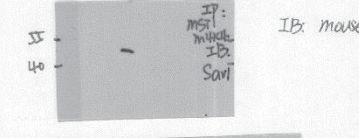
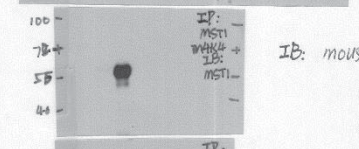
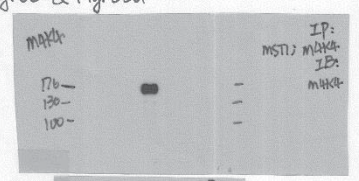
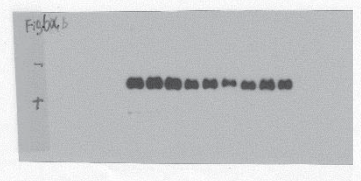
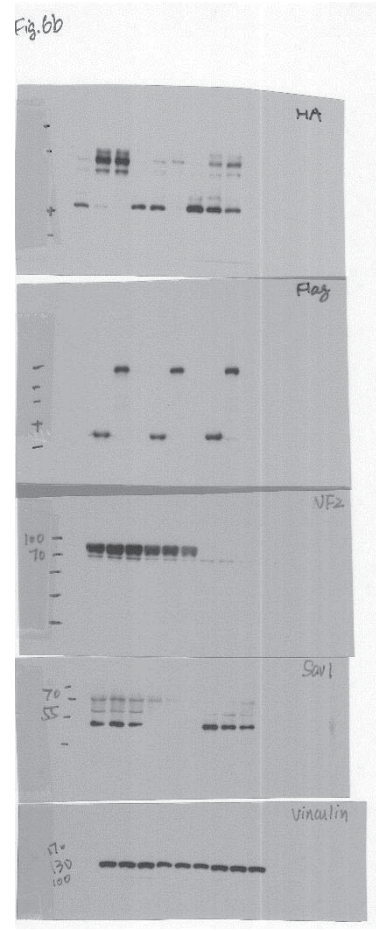


Fig. 6a

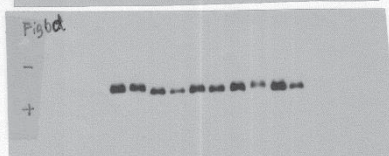
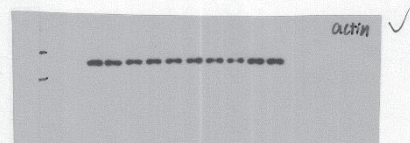
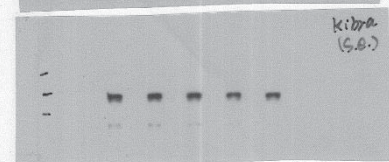
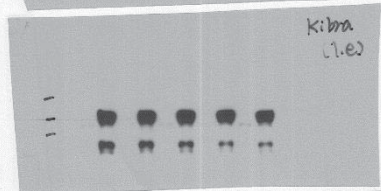
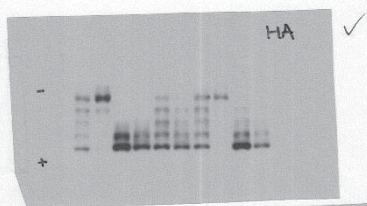


Fig. 6b

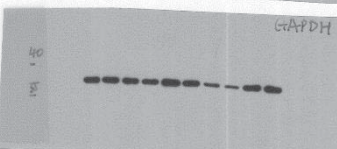
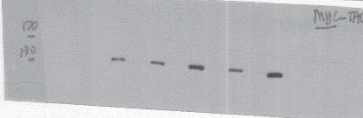
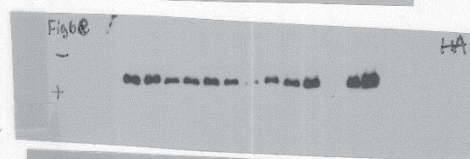
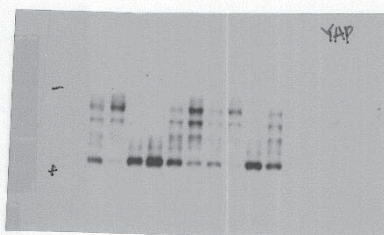


Fig. 7a

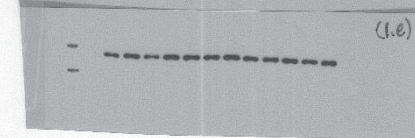
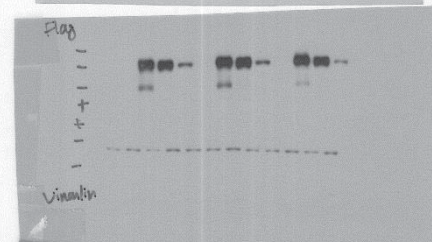
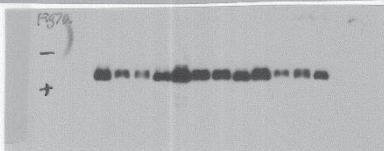
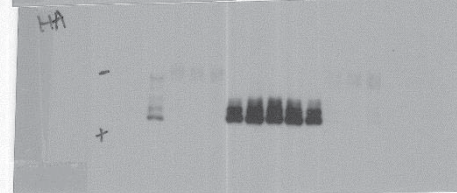
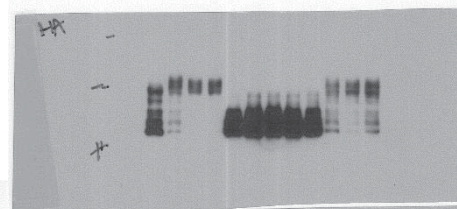
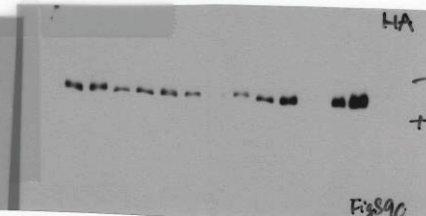
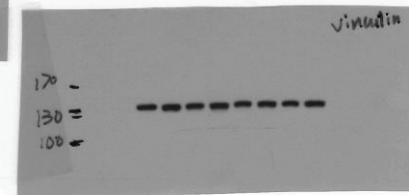
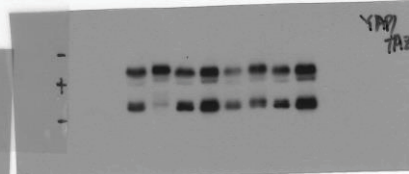
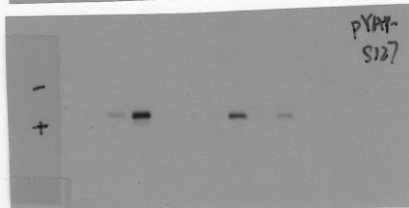
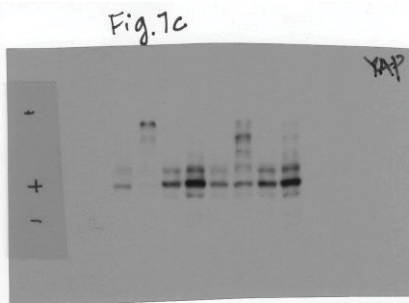
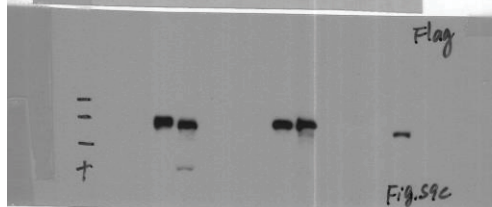
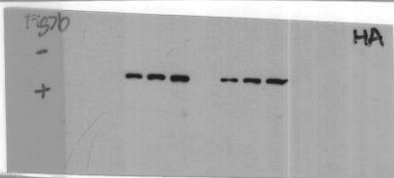
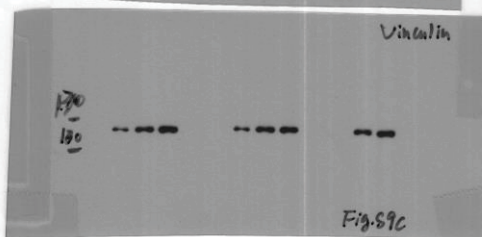
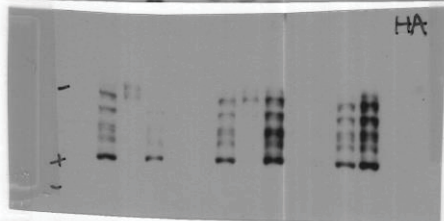
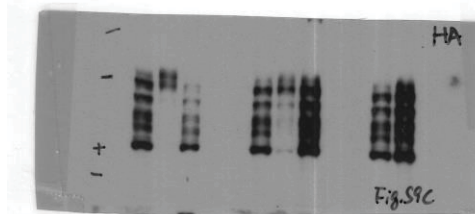


Fig. 7b & Fig. 9c.



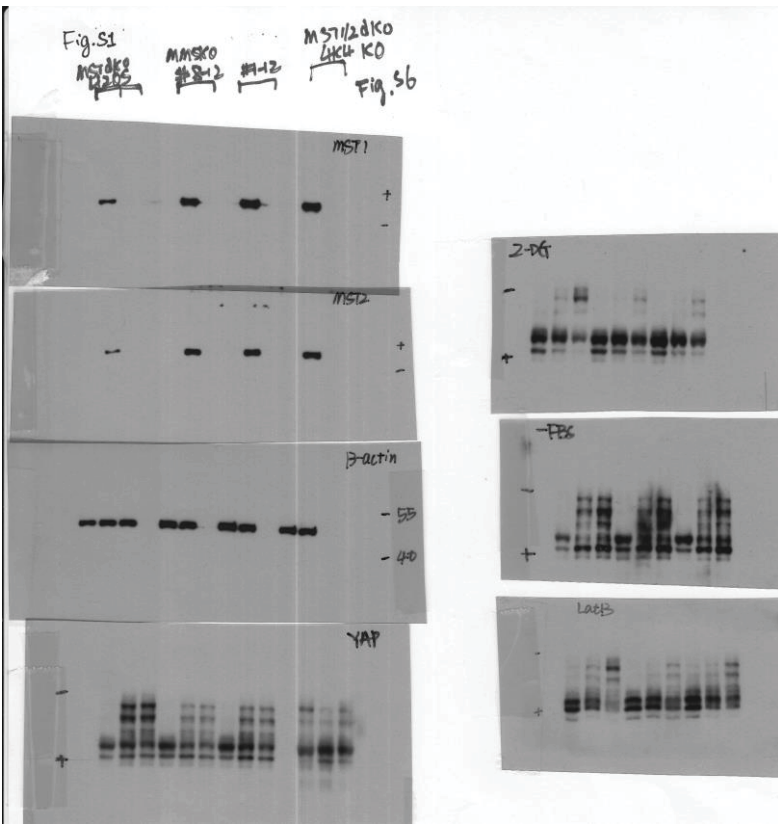


Fig.S1

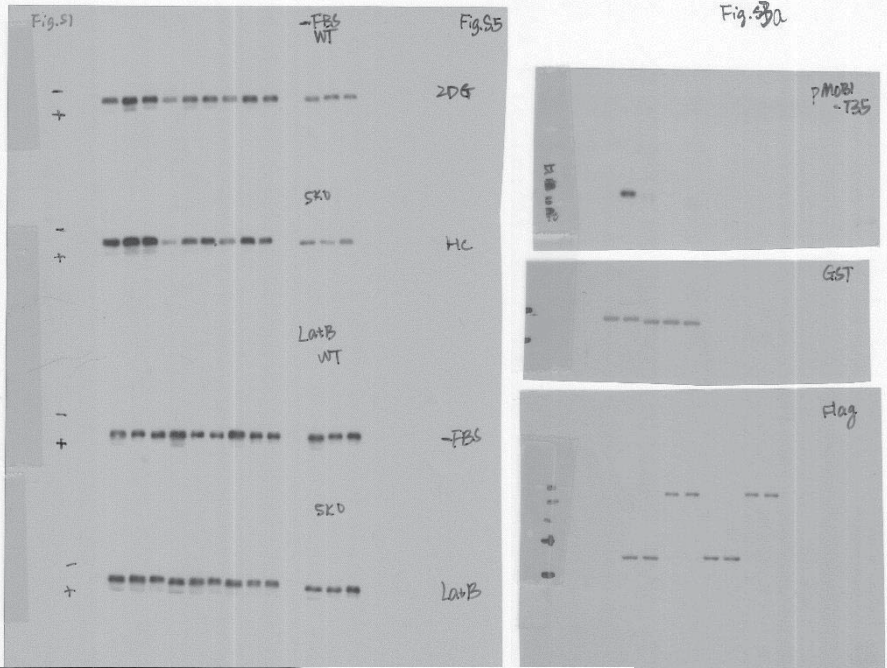
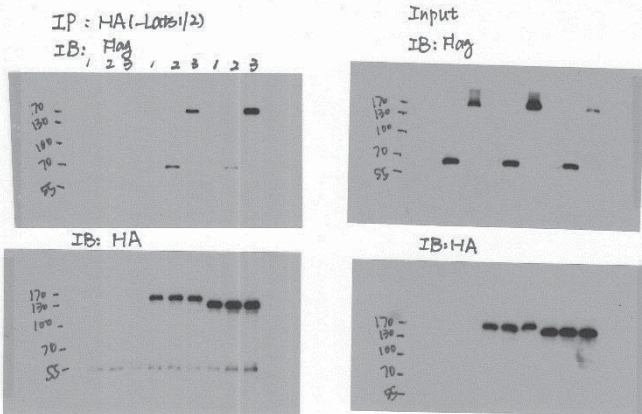


Fig S3b



1. PRK7- N-Flag
2. MST2-Flag
3. MAP4K4-Flag

PCNA-SHA Lats1-HA Lats2-HA

Fig. S3c

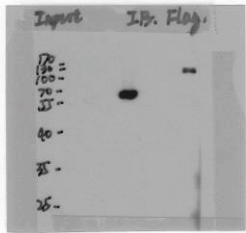
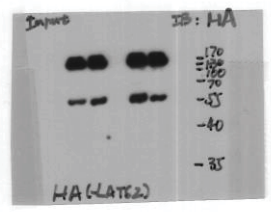
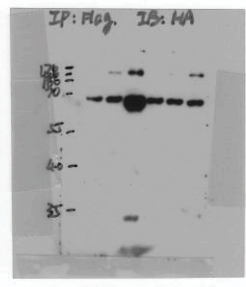
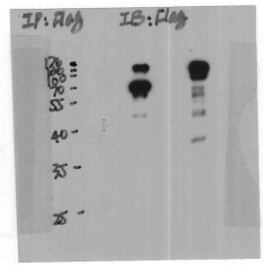
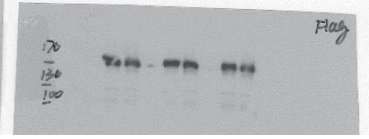
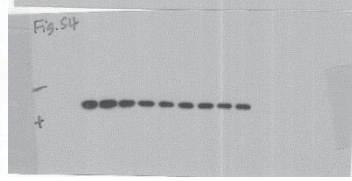
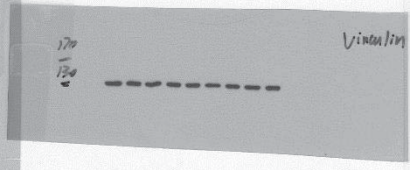
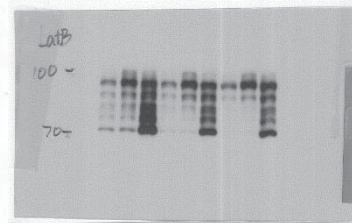
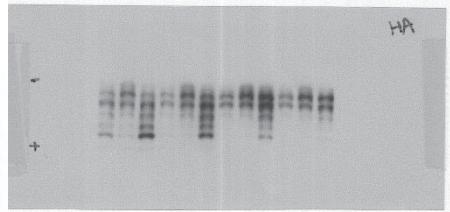
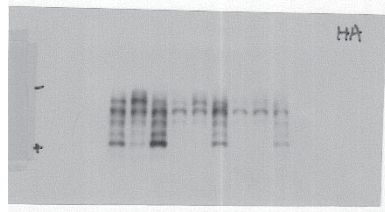


Fig. S4



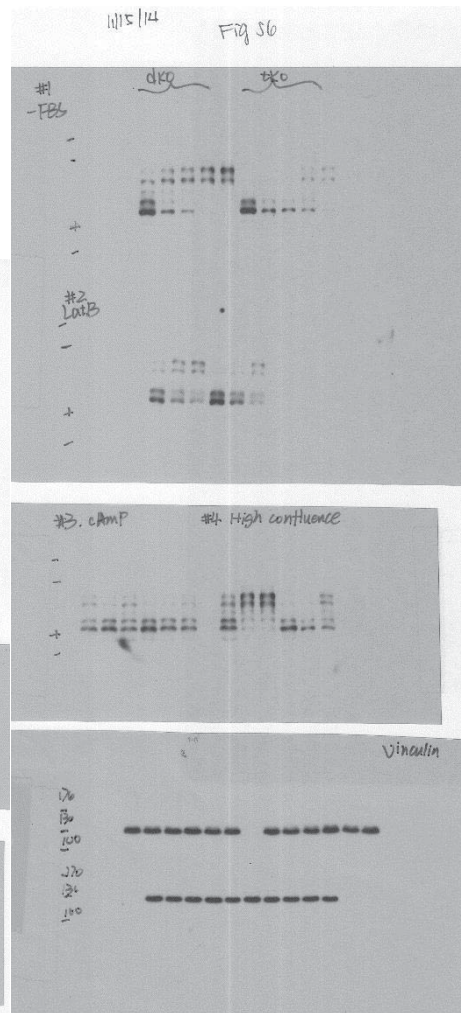
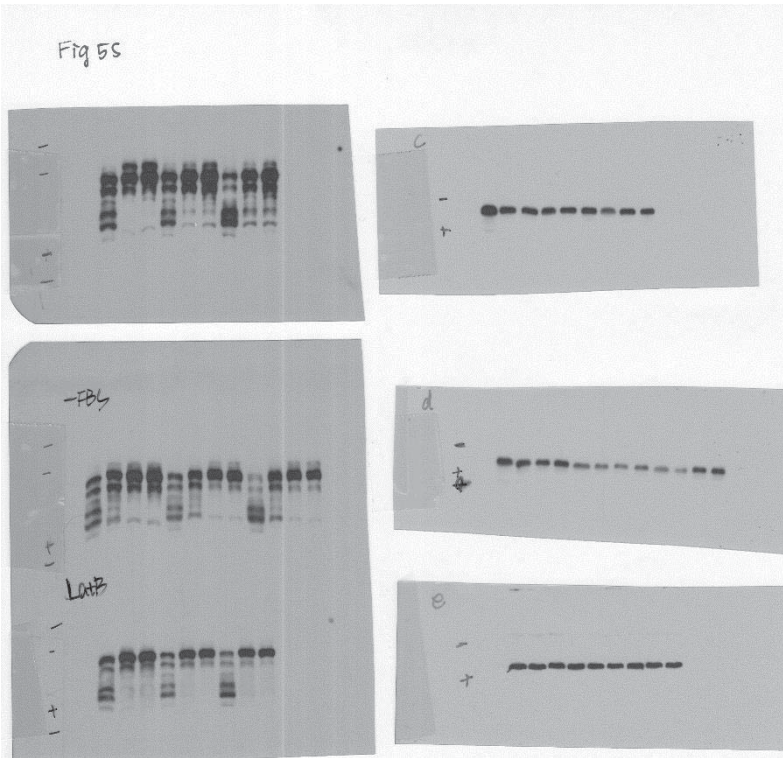


Fig 56

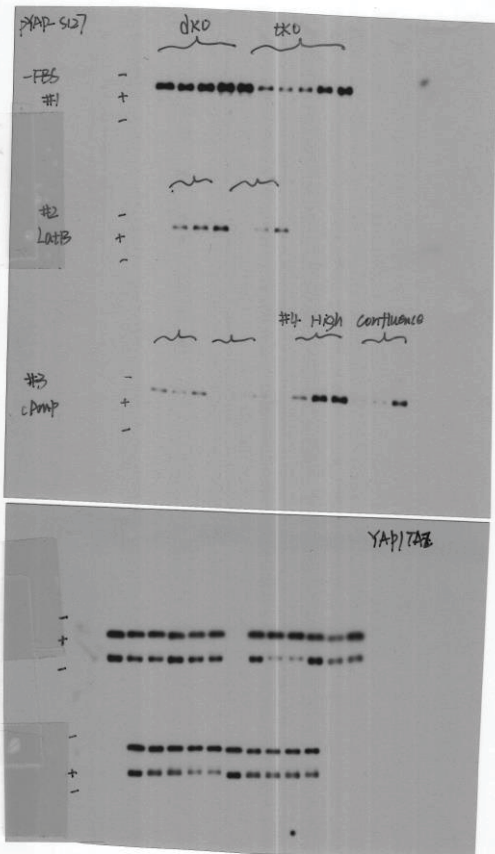


Fig 57

Fig. 57a

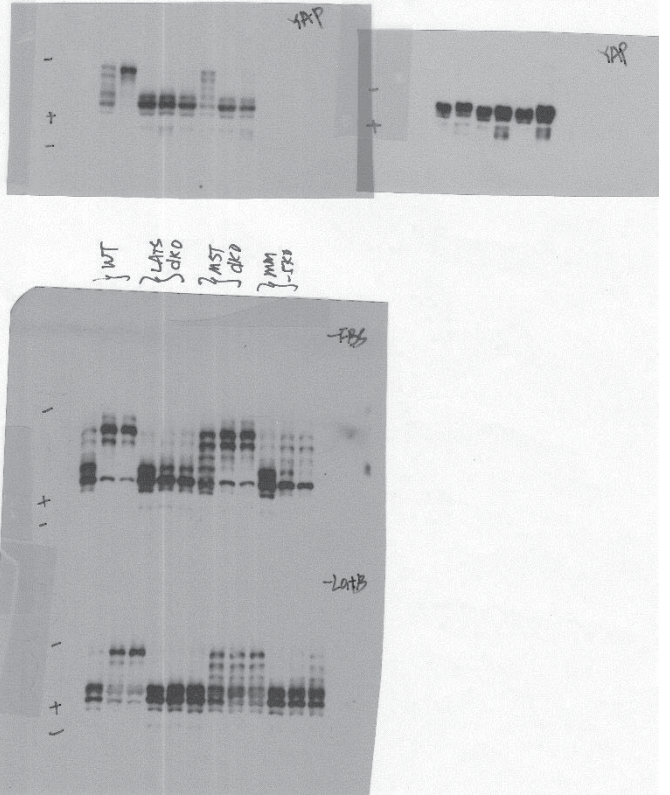


Fig. S9a/b

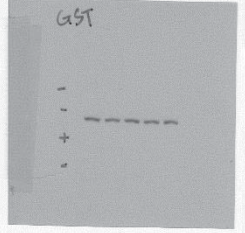
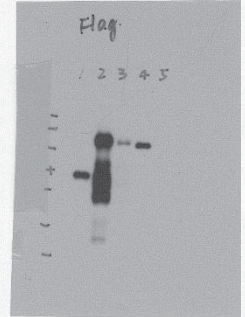
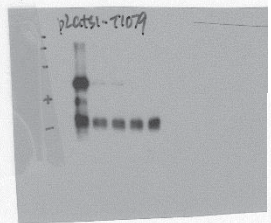
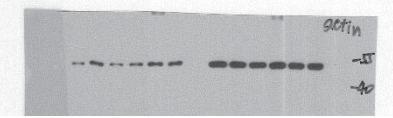
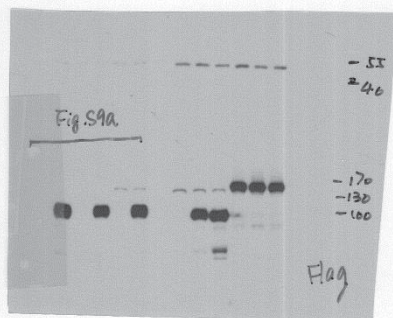
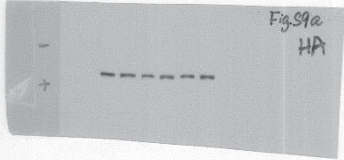
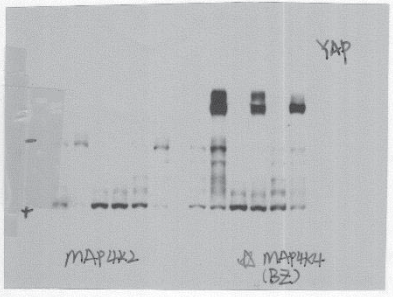
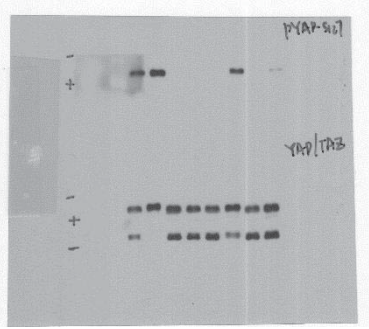
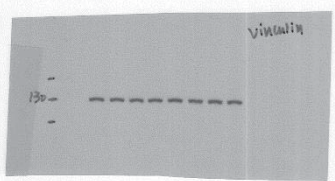
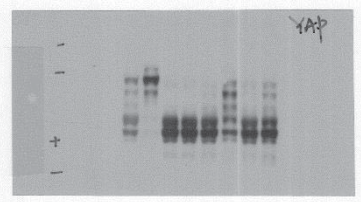


Fig. S10



Supplementary table. 1.

List of sgRNA used for CRISPR

Cell type	Clone	Gene	sgRNAs
LATS1/2 dKO HEK293A		LATS1	CGTGCAGCTCTCCGCTCTAA
		LATS2	TACGCTGGCACC GTAGCCCT
MST1/2 dKO HEK293A		MST1	ATACACCGAGATATCAAGGC
		MST2	AGTACTCCATAACAATCCAG
MST1/2 dKO U2OS	#1	MST1	AGCTTTGTATACGCTGCCAT
		MST2	TTTAATTGCGACA ACTTGAC
	#2	MST1	ATACACCGAGATATCAAGGC
		MST2	AGTACTCCATAACAATCCAG
SAV1 KO HEK293A		SAV1	TCCAGGAGGAAGTCCTTCTC
NF2 KO HEK293A		NF2	GTCCATGGTGACGATCCTCA
MAP4K4 KO HEK293A	#1	MAP4K4	CAGGACATGATGACCAACTC
	#2	MAP4K4	GGGCGGAGAAATACGTTTCAT
MAP4K4/6/7 tKO HEK293A	#1	MAP4K4	GGGCGGAGAAATACGTTTCAT
		MAP4K6	CGGACAGGTCGATGTCGTCC
		MAP4K7	CGACTCCCCGGCTCGAAGCC
	#2	MAP4K4	GGGCGGAGAAATACGTTTCAT
		MAP4K6	AGGGTCGGCATGTCAAGACG
		MAP4K7	CGACTCCCCGGCTCGAAGCC
MST1/2-MAP4K4/6/7 5KO	#1	MST1	ATACACCGAGATATCAAGGC
		MST2	AGTACTCCATAACAATCCAG
		MAP4K4	GGGCGGAGAAATACGTTTCAT
		MAP4K6	CGGCAATGGAACCTACGGAC
		MAP4K7	CGACTCCCCGGCTCGAAGCC
	#2	MST1	ATACACCGAGATATCAAGGC
		MST2	AGTACTCCATAACAATCCAG
		MAP4K4	CAGGACATGATGACCAACTC
		MAP4K6	AGGGTCGGCATGTCAAGACG
		MAP4K7	TTCATCCAGGCTTCGAGCCG

Supplementary table 2.

List of antibodies

Gene	Vendor	Catalog #	Used for CRISPR	Dilution for Western blot
LATS1	Cell signaling	3477	Yes	1:2000
LATS2	Cell signaling	5888	Yes	1:1000
	Abcam	ab70565	Yes	1:1000
MST1	Cell signaling	3682	Yes	1:2000
	BD Biosciences	611052	Yes	1:2000
MST2	Abcam	ab52641	Yes	1:2000
MAP4K4	Cell signaling	5146	Yes	1:1000
	Bethy	A301-502A	Yes	1:2000
MAP4K6	Bethy	A302-191A	Yes	1:2000
	Abcam	ab86385	Yes	1:1000
MAP4K7	Santa Cruz	ab95887	Yes	1:1000
	Bethy	A310-985A	Yes	1:1000
NF2	Cell signaling	12888	Yes	1:2000
SAV1	Cell signaling	13301	Yes	1:2000
	Santa Cruz	sc-374366		1:1000
pLATS1-T1079 (HM)	Cell signaling	8654		1:1000
pLATS1-S909 (AL)	Cell signaling	9157		1:1000
YAP	Abcam	ab52771		1:2000
YAP/TAZ	Santa Cruz	sc-101199		1:1000
pYAP-S127	Cell signaling	4911		1:2000
MOB1	Cell signaling	13730		1:1000
pMOB1-T35	Cell signaling	8699		1:1000
Actin	Sigma	A5441		1:5000
Vinculin	Sigma	V9131		1:5000
Flag-HRP	Sigma	A8592		1:5000
Flag	Sigma	F1804		1:2000
HA-HRP	Cell signaling	2999		1:5000
HA	Covance	MMS-101P		1:2000
GST	Sigma	SAB4200237		1:2000
	Cell signaling	2625		1:1000
Myc	Santa Cruz	sc-40		1:250
	Cell signaling	2278		1:1000
KIBRA	Cell signaling	8774		1:2000
GAPDH	Cell signaling	5174		1:2000