

**Table S1. List of high-confidence interactors identified by affinity purification and mass spectrometry analysis. Related to Figure 1**

<b>Protein name</b>	<b>Accession number</b>	<b>Molecular weight (kDa)</b>
NPM1	P06748	32
SET	Q01105	33
PPM1G/PP2C $\gamma$	O15355	59
hnRNP-F	P52597	46
MePCE/BCDIN3	Q7L2J0	74
DDX21	Q9NR30	87
Sart3/Tip110	Q15020	110
DHX57	Q6P158	155

## SUPPLEMENTAL EXPERIMENTAL PROCEDURES

### Materials

5,6-dichloro-1- $\beta$ -ribofuranosylbenzimidazole (DRB) (Sigma, D1916) was used at a final concentration of 100  $\mu$ M for 4 hrs. TNF- $\alpha$  (Sigma, T6674) was used at final concentration of 50 ng/ml for 1 hr.

### Cell culture and treatments

For the heat shock experiment, HeLa cells were heat shocked at 42°C for 2 hrs. To induce the inflammatory pathway, HeLa cells were treated with TNF- $\alpha$  (50 ng/ml) for 1 hr.

### RNAi and rescue assay

For the rescue assay (**Figure S5**), HeLa cells were transfected with 100 ng of an empty pcDNA4/TO (-) or pcDNA4/TO-PPM1G:S (PPM1G) plasmid DNA using PolyJet (SignaGen). One day post-transfection, cells were re-transfected with the indicated siRNAs. Two days post-siRNA transfection, cells were induced with TNF- $\alpha$  and used for RNA extraction and gene expression analysis by RT-qPCR and Western Blot.

### UV crosslinking of RNA-protein complexes *in vivo*

293T cells were seeded at  $2 \times 10^6$  cells into a 10-cm<sup>2</sup> plate and transfected with 5  $\mu$ g of a Strep-tagged Tat plasmid 24 hrs post seeding. For the experiment in **Figure S1A**, cells were irradiated with 125 mJ/cm<sup>2</sup> using a Spectrolinker XL-1500 UV crosslinker and harvested and lysed using denaturing lysis buffer (150 mM NaCl, 20 mM Tris-HCl pH = 7.0, 5% Glycerol, 1.5 mM MgCl<sub>2</sub>, 1 mM DTT, 1% NP-40, 0.25% Sodium Deoxycholate and 0.1 % SDS) and sonicated using the Bioruptor water bath Sonicator (Diagenode) with 8 cycles (30 sec on 30 sec off). Tat complexes

were affinity purified using Strep-tactin Superflow Resin (IBA Life Sciences) and washed using denaturing washing buffer (250 mM NaCl, 20 mM Tris-HCl pH = 7.0, 5% Glycerol, 1.5 mM MgCl<sub>2</sub>, 1 mM DTT, 0.05% NP-40, 0.1 % Sodium Deoxycholate, 0.1 % SDS). Co-immunoprecipitating RNAs were extracted using Proteinase K digestion [0.5 mg/mL Proteinase K, 20mM Tris-HCl pH = 7.0, 5 mM EDTA, 0.5% SDS, 300 pg/μl *in vitro* transcribed β-globin RNA (used as precipitation control, PC)] (Conrad, 2008) followed by Phenol:Chloroform:Isoamyl alcohol (25:24:1) extraction and ethanol precipitation. The immunoprecipitated material was analyzed by Northern blot using a 7SK probe encompassing stem-loop I made with primers P15 and P16 (**Table S4**) and labeled using a PCR DIG DNA labeling kit (Roche).

**Table S2. Plasmids used in this study**

<b>Gene</b>	<b>Vector / Tag</b>	<b>Restriction sites</b>
Tat	pcDNA-4TO / Strep	BamHI - XhoI
Tat	pcDNA-4TO / Flag	BamHI - XhoI
Larp7	pcDNA-4TO / Strep	BamHI - XhoI
Larp7	pcDNA-4TO / Flag	BamHI - XhoI
Sart3	pcDNA-4TO / Strep	HindIII - XhoI
Sart3	pcDNA-4TO / Flag	HindIII - XhoI
CycT1	pcDNA-4TO / Strep	HindIII - EcoRI
CycT1	pcDNA-4TO / Flag	HindIII - EcoRI
Cdk9	pcDNA-4TO / Strep	HindIII - XhoI
Cdk9	pcDNA-4TO / Flag	HindIII - XhoI
MePCE	pcDNA-4TO / Strep	HindIII - XhoI
MePCE	pcDNA-4TO / Flag	HindIII - XhoI
DDX21	pcDNA-4TO / Strep	BamHI - XhoI
DDX21	pcDNA-4TO / Flag	BamHI - XhoI
NPM1	pcDNA-4TO / Strep	BamHI - XhoI
NPM1	pcDNA-4TO / Flag	BamHI - XhoI
hnRNP-F	pcDNA-4TO / Strep	BamHI - XhoI
hnRNP-F	pcDNA-4TO / Flag	BamHI - XhoI
SET	pcDNA-4TO / Flag	NotI - XhoI
GFP	pcDNA-4TO / Strep	BamHI - XhoI
GFP	pcDNA-4TO / Flag	XhoI - ApaI
PPM1G	pcDNA-4TO / Strep	BamHI - NotI
PPM1G	pcDNA-4TO / Flag	BamHI - NotI

**Table S3. siRNAs used in RNAi assays**

<b>Gene</b>	<b>Flexitube siRNA product name</b>	<b>Catalog number</b>
Cdk9	Hs_CDK9_5	SI00605066
MePCE	Hs_BCDIN3_1	SI02778265
hnRNP-F	Hs_HNRPF_1	SI00300461
DHX57	Hs_DHX57_5	SI02757839
NPM1	Hs_NPM1_1	SI00300979
PPM1G	Hs_PPM1G_6	SI02658684
PPM1G*	Hs_PPM1G_10	SI05016627
PPM1G*	Hs_PPM1G_11	SI05016634
PPM1A	HS_PPM1A_5	SI02659258
SLC25A5	Hs_SLC25A5_5	SI02654358
CycT1	Hs_CCNT1_1	SI00024073
Larp7	Hs_HDCMA18P_2	SI00434980
SET	Hs_SET_5	SI03021291
DDX21	Hs_DDX21_7	SI04311412
Sart3	Hs_SART3_7	SI04357304
RelA	Hs_RELA_5	SI00301672
Negative control	AllStars Negative Control	SI1027280

\* siRNA oligos used in rescue experiments because they target the 3' non-coding region.

**Table S4. DNA oligonucleotides used in this study**

<b>Gene</b>	<b>Primer number/Sequence (5'–3')</b>	<b>Purpose / Assay</b>
<b>MePCE</b>	<b>507/TGAAGCCAGAGCAGTTCAGTTCCTA</b> <b>508/TACACAGGACGCTGGAAGCCTTTA</b>	RNAi validation
<b>Larp7</b>	<b>511/TCCGGGATACTTTGGCAGCAATCT</b> <b>512/AGGATCTCGAGTTTCCAGCAGTGT</b>	RNAi validation
<b>Cdk9</b>	<b>517/GTGTTTCGACTTCTGCGAGCATGAC</b> <b>518/CTATGCAGGATCTTGTCTGTGG</b>	RNAi validation
<b>SLC25A5</b>	<b>519/GCAGCTGATGTGGGTAAAGCTGG</b> <b>520/CCGGAAGCATTCCCTTTGCAGT</b>	RNAi validation
<b>NPM1</b>	<b>525/GTTGTGAACTAAAGGCCGACAAAG</b> <b>526/TGTGCAACTCATCCTTTGCACCAG</b>	RNAi validation
<b>hnRNP-F</b>	<b>527/TGTATTGGTCTCCTGCTCCTAGA</b> <b>528/AGGACTGGTTTCTGTTGCTACC</b>	RNAi validation
<b>DDX21</b>	<b>531/CTGGGTGTTTGCTTTGATGTACC</b> <b>532/AGTTCTGGTTGCTCTGTGG</b>	RNAi validation
<b>DHX57</b>	<b>537/CCGCTGTTTGGATCCTGCTCT</b> <b>538/GCTGCCATCCCTTATACGCTTG</b>	RNAi validation
<b>SET</b>	<b>539/AATATAACAACTCCGCCAACCC</b> <b>540/CTGGTCAAATAATGCAGTGCCTC</b>	RNAi validation
<b>PPM1G</b>	<b>543/AAAATGGCAACAGCGACAAG</b> <b>544/CACCTCATACCCACTGCTA</b>	RNAi validation
<b>Sart3</b>	<b>549/CTTTACTCGTGCCCTTGGAGTAT</b> <b>550/CCGAGCTTTCTGCATGTTATTG</b>	RNAi validation
<b>CycT1</b>	<b>551/GCAGACTTTAGGCTTTGAACTAAC</b> <b>552/TACTGCAGGCTAAATGTGGT</b>	RNAi validation
<b>7SK RNA</b>	<b>15/GGATGTGAGGGCGATCTG</b> <b>16/GGAGCGGTGAGGGAGGAAG</b>	RIP assay, Northern blot
<b>U6 snRNA</b>	<b>9/CTCGCTTCGGCAGCACATATAC</b> <b>10/GGAACGCTTCACGAATTTGCGTG</b>	RIP assay
<b>β-actin</b>	<b>389/CCCCCGGGCCGTCTTCCCCTC</b> <b>390/TGAGGATGCCTCTCTTGCTCTG</b>	Internal control RNAi
<b>gapdh</b>	<b>239/CCCTGTGCTCAACCAGT</b> <b>240/CTCACCTTGACACAAGCC</b>	Internal control RNAi
<b>c-Myc</b>	<b>860/ACAGCTACGGAActCTTGTGCGTA</b> <b>861/CAGCCAAGTTGTGAGTTGCATT</b>	Internal control RNAi
<b>Rpl19</b>	<b>354/ATCGATCGCCACATGTATCA</b> <b>355/GCGTGCTTCCTTGGTCTTAG</b>	Internal control RNAi
<b>Hsp90</b>	<b>838/CCTTCTATTTGTCCCACG</b> <b>839/ATCCTCCGAGTCTACCAC</b>	Heat shock response
<b>Hsp70</b>	<b>842/ACCCGCATCCCCAAGGTGCAG</b> <b>843/TCAGGGCAGTCATCACGCCTC</b>	Heat shock response
<b>IL-8</b>	<b>245/GGGCCATCAGTTGCAAATC</b> <b>246/TTCTTCCGGTGGTTTCTTC</b>	Inflammatory response ChIP (Promoter)
<b>IL-8</b>	<b>282/GCCATAAAGTCAAATTTAGCTGGAA</b> <b>283/GTGCTTCCACATGTCCTCACA</b>	Inflammatory response ChIP (Gene body)
<b>IκBα</b>	<b>255/GATCCGCCAGGTGAAGGG</b> <b>256/GCAATTTCTGGCTGGTTGG</b>	Inflammatory response

**Table S5. Antibodies used in IF, IP, ChIP and Western blot assays**

<b>Protein/Tag</b>	<b>Source (Company)</b>	<b>Catalog Number</b>
CycT1	Santa Cruz Biotechnologies	sc-10750
Cdk9	Santa Cruz Biotechnologies	sc-484
Phospho-Cdk9 (Thr186)	Cell Signaling	2549
STREP-Tag II	Novagen	71591
FLAG M2	Sigma	F1804
MePCE/BCDIN3 (N-17)	Santa Cruz Biotechnologies	sc-82542
Larp7/PIP7S	Sigma	AV40847
Hexim1	Abcam	ab25388
RNA Polymerase II CTD	Active Motif	102660
RNA Polymerase II S5P-CTD	Abcam	ab5131
RNA Polymerase II S2P-CTD	Abcam	ab5095
Normal Rabbit IgG	Santa Cruz Biotechnologies	sc-2027
$\beta$ -actin (C4)	Santa Cruz Biotechnologies	sc-47778
PPM1G/PP2C $\gamma$	Santa Cruz Biotechnologies	sc-136320
PPM1G/PP2C $\gamma$	Abcam	ab70794
SART3 (C3)	GeneTex	GTX107684
p65/NF- $\kappa$ B/RelA (C-20)	Santa Cruz Biotechnologies	sc-372X
HIV-1 p24 (AG3.0)	AIDS Research and Reference Reagent Program	4121
GST	Santa Cruz Biotechnologies	sc-33613

## **SUPPLEMENTAL REFERENCES**

Conrad, N.K. (2008). Chapter 15. Co-immunoprecipitation techniques for assessing RNA-protein interactions in vivo. *Methods Enzymol.*, 317-342.

Li, Q., Price, J.P., Byers, S.A., Cheng, D., Peng, J., and Price, D.H. (2005). Analysis of the large inactive P-TEFb complex indicates that it contains one 7SK molecule, a dimer of HEXIM1 or HEXIM2, and two P-TEFb molecules containing Cdk9 phosphorylated at threonine 186. *J. Biol. Chem.* *280*, 28819-28826.