

Supplemental data

Tables

Table I Sequence of oligopeptides used as substrates with corresponding protein names.

Oligopeptide sequence	Protein name
[H] <u>SY</u>SMDHF RWGRPVGRRRRPVRVYP[OH]*	ACTH (aa 138-161) (P01189)
[H] <u>SE</u>SSSKS RWGRPVGRRRRPVRVYP[OH] *	High mobility group protein A1 (P17096)
[H] <u>MLG</u>PEGG RWGRPVGRRRRPVRVYP [OH]*	hnRNP F; (P52597)
[H] <u>MLD</u>PEGG RWGRPVGRRRRPVRVYP [OH]*	hnRNP F; G3D mutant
[H] <u>MLL</u>PEGG RWGRPVGRRRRPVRVYP [OH]*	hnRNP F; G3L mutant
[H] <u>MLR</u>PEGG RWGRPVGRRRRPVRVYP [OH]*	hnRNP F; G3R mutant
[H] <u>MIG</u>PEGG RWGRPVGRRRRPVRVYP [OH]*	hnRNP F; L2I mutant
[H] <u>MFG</u>PEGG RWGRPVGRRRRPVRVYP [OH]*	hnRNP F; L2F mutant
[H] <u>MLGTE</u>GG RWGRPVGRRRRPVRVYP[OH]*	hnRNP H; (P31943)
[H] <u>MLALIS</u>R RWGRPVGRRRRPVRVYP [OH]*	hARL8b; (Q9NVJ2)
[H] <u>MLGTG</u>PA RWGRPVGRRRRPVRVYP [OH]*	mTOR; (P42345)
[H] <u>MKEE</u>VKG RWGRPVGRRRRPVRVYP[OH]*	Kinesin KIF4A (Q95239)
[H] <u>MDEL</u>FPL RWGRPVGRRRRPVRVYP [OH]*	NF- κ B/p65 (Q04206)
[H] <u>DDIA</u>AL RWGRPVGRRRRPVRVYP[OH]*	β -actin (NP_001092)

The custom made peptides were obtained from BioGenes or Sigma Genosys.

* Bold indicates amino acids found in corresponding protein while underlines indicate the residues used to identify the peptides in the text.

Table II Primer sequences used to mutate GST-hNaa50p

hNaa50p mutants	Sequence 5' – 3'
R84A	GTCTGGCACCTTACGCAAGGCTAGGAATAGG
Y124F	GTCGGCAATTGACTTCTTCAGGAAGTTTGGCTTTGAG
K34R	CCAGTCAGCTACAATGACAGGTTCTACAAGGATGTGCTG
K37R	GACAAGTTCTACAGGGATGTGCTGGAGGTTGGC
K140R	CAAAGAAGA ACTACTATAGGAGGATAGAGCCCGC

Table III Potential human *in vivo* substrates for hNaa50p mediated acetylation

Protein name	Protein ID code	N-terminal protein sequence
EF-hand calcium-binding domain-containing protein 2	Q5VUJ9	<u>MLGP</u> GQVRLRPR
Protein kinase-like protein SgK071	Q8NE28	<u>MLGP</u> GSNRRRPT
Putative uncharacterized protein C1orf180	Q8NAE3	<u>MLGP</u> NAQVTVVV
Dihydrodipicolinate synthase-like, mitochondrial precursor	Q86XE5	<u>MLGP</u> QVWSSVRQ
Synaptopodin	Q8N3V7	<u>MLGP</u> HLPPLA
Taste receptor type 1 member 3 precursor (Sweet taste receptor T1R3)	Q7RTX0	<u>MLGP</u> AVLGLSLW
Alkaline phosphatase, placental type precursor (PLAP-1)	P05187	<u>MLGP</u> CMLLLLLL
Olfactory receptor 52I2	Q8NH67	<u>MLGP</u> AYNHTMET
Olfactory receptor 52I1 (Olfactory receptor OR11-13)	Q8NGK6	<u>MLGP</u> AYNHTMET
Heterogeneous nuclear ribonucleoprotein F (hnRNP F)	P52597	<u>MLGP</u> EGGEFV
Calcium-dependent secretion activator 1 (CAPS-1)	Q9ULU8	<u>MLDP</u> SSSEESD
Calcium-dependent secretion activator 2 (CAPS-2)	Q86UW7	<u>MLDP</u> SSSEESD
TIMELESS-interacting protein	Q9BVW5	<u>MLEP</u> QENGVIDL
Putative forkhead box protein ENSP00000381211	A8MTJ6	<u>MLSP</u> ERRDQPRS
Transmembrane protein C9orf144B	Q6ZU69	<u>MLSP</u> TFVLWEVG
RUN and SH3 domain-containing protein 1 (Nesca)	Q9BVN2	<u>MLSP</u> QRALLCNL
Armadillo repeat-containing protein 2	Q8NEN0	<u>MLSP</u> NDKMLGKL
Rho GTPase-activating protein 22	Q7Z5H3	<u>MLSP</u> KIRQARRA
Zinc finger protein 671	Q8TAW3	<u>MLSP</u> VSRDASDA
Atlastin-3	Q6DD88	<u>MLSP</u> QRVAAAAS
p53 and DNA damage-regulated protein 1	Q9NUG6	<u>MLSP</u> EAERVRLRY
Olfactory receptor 5M10 (Olfactory receptor OR11-207)	Q6IEU7	<u>MLSP</u> NHTIVTEF
Sulfotransferase family cytosolic 1B member 1	O43704	<u>MLSP</u> KDILRKDL
Uncharacterized protein C11orf49	Q9H6J7	<u>MLSP</u> ERLALPDY
Uncharacterized protein MGC35361	Q8N0U4	<u>MLAP</u> CSGWELGC
Thrombospondin-4 precursor	P35443	<u>MLAP</u> RGAAVLLL
Leucine-rich repeat and transmembrane domain protein 2 precursor	Q8N967	<u>MLAP</u> GSSPGQRG
Kanadapin (Lung cancer oncogene 3 protein - HLC-3)	Q9BWU0	<u>MLAP</u> LRNAPGRE
Uncharacterized protein FLJ37543 precursor	Q2M2E5	<u>MLAP</u> LFLCCLRN
Putative transcript Y 12 protein	Q9BZ98	<u>MIDP</u> ETRHKAFL
Retrotransposon-like protein 1	A6NKG5	<u>MIEP</u> SEDSFETM
Chloride anion exchanger (Solute carrier family 26 member 3)	P40879	<u>MIEP</u> FGNQYIVA
C-type lectin domain family 2 member A	Q6UVW9	<u>MINP</u> ELRDGRAD
Zinc finger protein 221	Q9UK13	<u>MISP</u> SLELLHSG
TRAF3-interacting modulator (TRAF3-interacting protein 3)	Q9Y228	<u>MISP</u> DPRPSPGL
Alkaline phosphatase, tissue-nonspecific isozyme precursor	P05186	<u>MISP</u> FLVLAIGT
Beta-2-glycoprotein 1 precursor (Beta-2-glycoprotein I)	P02749	<u>MISP</u> VLILFSSF
Putative BMP-2-inducible kinase-like protein	Q5H9B9	<u>MIAP</u> SPKSSEEE
Zinc finger BED domain-containing protein 5	Q49AG3	<u>MIAP</u> LLCILSYN

Supplemental Figures

Figure S1. MALDI-MS spectra of immunoprecipitated Xpress-hNaa50p from transfected HeLa cells.

Xpress-hNaa50p was isolated from transfected HeLa cells by immunoprecipitation with anti-Xpress antibody. **A.** MALDI-MS spectra (mass range 2.100- 2.200) showing a peptide with m/z of 2173.2 corresponding to acetylated lysine 34. **B.** MALDI-MS spectra (mass range 1.190- 2.050) showing a peptide with m/z of 1987.1 corresponding to acetylated lysine 140.

Figure S2. MALDI-MS and MS/MS spectra of autoacetylated GST-hNaa50p.

A. MALDI-MS spectrum of Lys-C produced peptides of recombinant GST-hNaa50p. Monoisotopic peaks are labelled with their respective m/z ratio. The +42 Da MALDI-molecular ions of m/z 1552.769, 1986.980 and 2173.072, are indicated with **B**, **C** and **D**, respectively, which also refer to the MS/MS spectra shown in the figure. The identities of the acetylated lysines present in the peptides are indicated. **B**, **C** and **D**. Tandem mass spectra of the m/z 1552.769, 1986.980 and 2173.072 ions, respectively, indicate the observed fragmentation pattern and the sequence ion assignments. Predicted ions (b and y) are shown at top of the figures. The acetylated lysines are displayed as 'acK' in the amino acid sequences.

Figure S3. Knockdown of human NatA with siRNA targeting hNAA15p also reduces protein levels of hNaa50p.

WB analyses using anti-hNaa50p, anti-hNaa15p and anti-tubulin antibodies. si-non-targeting represent cell lysate from HeLa cells transfected with a non-targeting siRNA, lane 2: Cell lysate from HeLa cells transfected with si-*NAA15*, lane 3: Cell lysate from HeLa cells transfected with si-*hNAA50*.

Figure S1

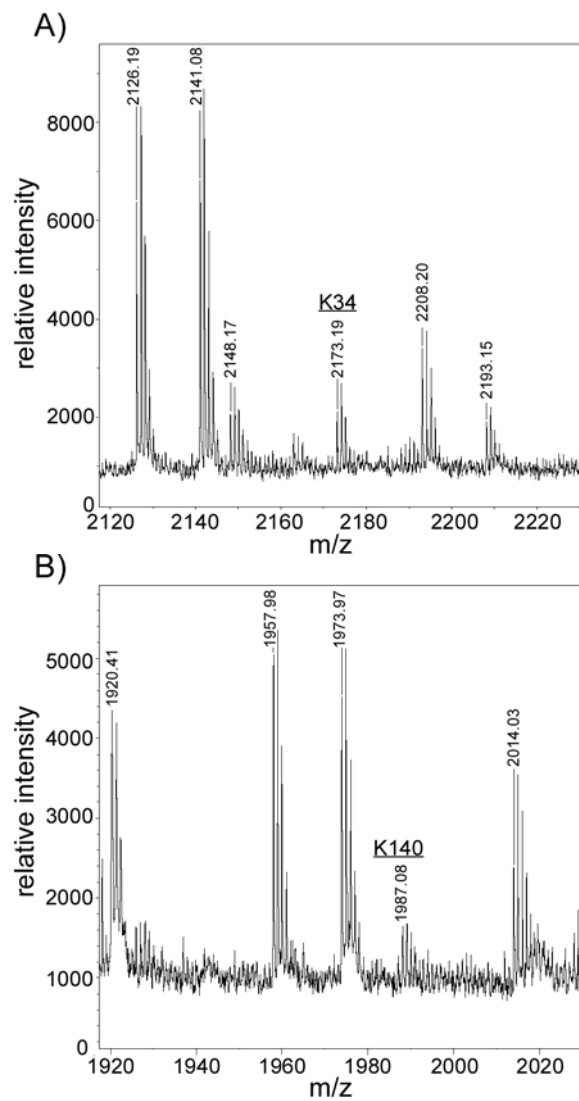
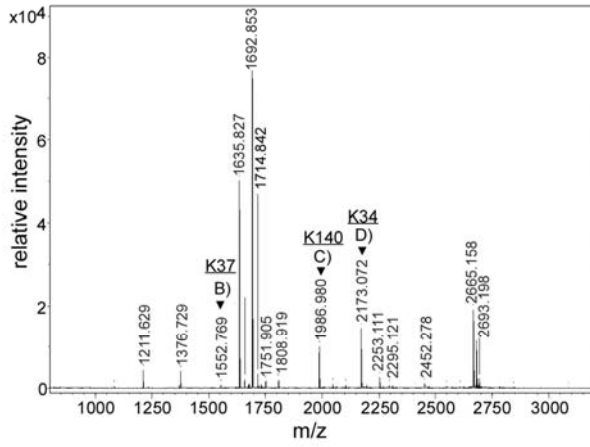
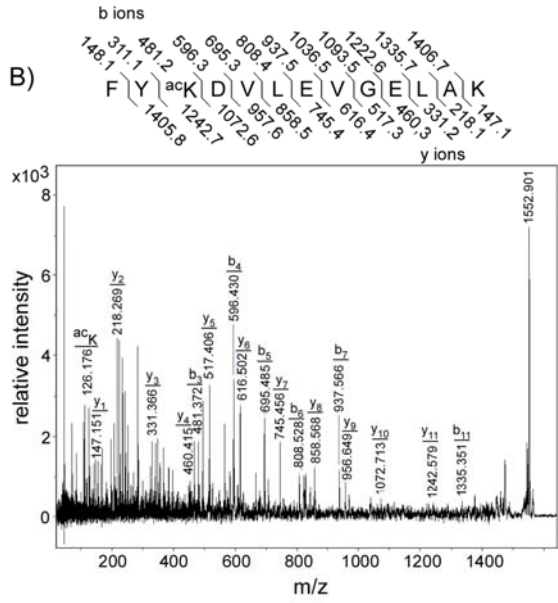


Figure S2

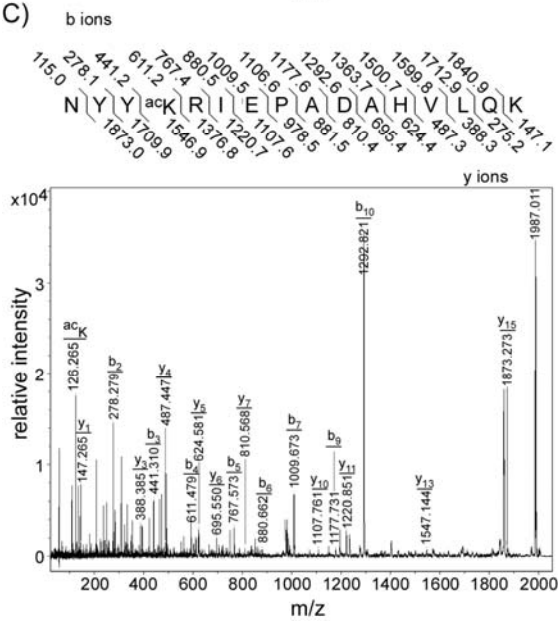
A)



B)



C)



D)

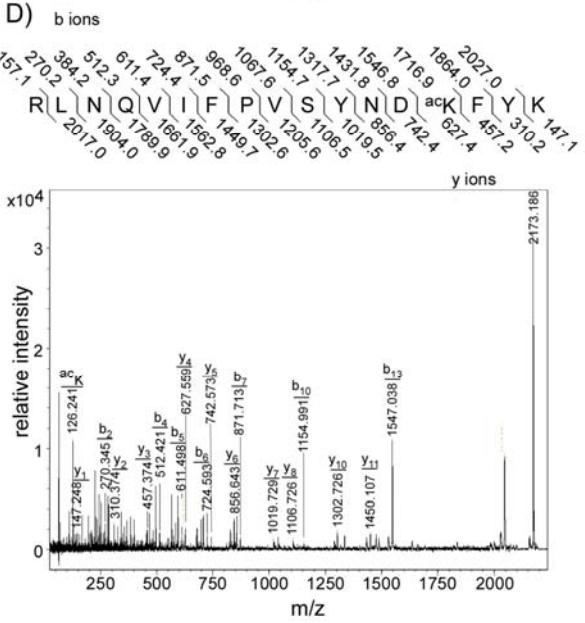


Figure S3

