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Supplemental Information

Inc-Rps4I-encoded peptide RPS4XL regulates

RPS6 phosphorylation and inhibits the

proliferation of PASMCs caused by hypoxia

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Supplemental material Figure S1

-5'3' Frame 1

TPLVLWEGEGRVSACSSAMARGPKKHLKRVAAPRHWMLDKLTGVFAPRPSAGPHRLRECLPLAIFLRNRLKYA LTGDEVKKICMQRLIKVDGKVRTDVAYPAGFMDVISIDKSGENFRLVYDTKGRFAVHRITPEEAKYKLCKVRK VFVGTKGIPHLVTHDARTIRYPDPLIKVNDTVQISLDSGKITDAIKFDTGNLCMVTGGANLGRIGVITNRERH PGSFDVVHVKDANGNGFATRLSNIFVIGKGNKPWISLPRGKGIRLTIAEERDKRLAAKQSG-SGL-GTLRGRG VYVCLYLKVF-AAC-IKKESK

-5'3' Frame 2-

LRLCSGKEKGVFPLAPPPWLVVPRNT-SVWRPHATGCWTN-PVCSRPVRLPARTACGSACRSPSS-GTGSSTP -PATR-RRSACSGSLRSTARSEPMWPTQLASWMSSASTRAARTSASSTTPRAASRYTASRPRRPSTSCAR-GK SSWVPRASRTSSRTTRVPSATLTRSSRSTTPCRSR-TAEKSPMPSSLIPATCAW-PEAPTWVASASSPTASAT PAPLTWST-KMPTAMASLPASPTSS-SGRATSRGSPFPGEKESASPLPKRETRGWRPSRAGEVDSEGHCGEGA FTFVYT-KFFKQHVELKKKAS

5'3' Frame 3-

SACALGRRRACFRLLLRHGSWSQETPEACGGPTPLDAGQTDRCVRAPSVCRPAPPAGVPAARHLPEEQAQVRP DRRRGEEDLHAAAH-GRRQGQNRCGLPSWLHGCHQHRQERRELPPRLRHQGPLRGTPHHARGGQVQAVQGEES LRGYQGHPAPRHARRAYHPLP-PAHQGQRHRADLARQRKNHRCHQV-YRQPVHGNRRRQPGSHRRHHQPRAPP RLL-RGPRERCQRQWLRYPPLQHLRDREGQQAVDLPSPGKRNPPHHCRRERQEAGGQAERVKWTLRDIAGKGR LRLFILKSFLSSMLN-KRKQA

Supplemental figure 1. Rps4l's open reading frame and coding sequence were

detected by ExPASy.

Figure S2



Supplemental figure 2. Overexpression efficiency of the plasmid were used. All values are represented as the mean \pm SEM (**p < 0.01, and ***p < 0.001; n \geq 3).





Supplemental figure 3. Mass spectrum of the RPS4XL. (A) Mass spectrum of RPS4XL amino acids 76-94. (B) Mass spectrum of RPS4XL amino acids 156-168. (C) Mass spectrum of RPS4XL amino acids 169-191. (D) Mass spectrum of RPS4XL amino acids 212-221.

Figure S4



Supplemental figure 4. Detection of the specificity of anti-RPS4XL antibody. (A) Western blot analysis of expression of RPS4XL detected by the antibody in PASMCs treated with lvRps41 or NC. (B) Western blot analysis of expression of RPS4XL detected by the antibody in PASMCs treated with si-Rps41 or si-NC.





Supplemental figure 5. Expression of Rps4l in different organs of overexpression transgenic mice. (A) Agarose gel electrophoresis of the expression of Rps4l in the heart, liver, spleen, lung and kidney of WT and Rps4lTg mice. (B) qPCR analysis of the overexpression efficiency of Rps4l in the heart, liver, spleen, lung and kidney of Rps4l mice. All values are represented as the mean \pm SEM (*p<0.05 and NS, no significance; $n \ge 3$).



Supplemental figure 6. RPS4XL does not regulate hypoxia-induced inhibition of apoptosis in PASMCs. (A) WB analysis of cleaved caspase-3 in hypoxic and control PASMCs transfected with lvRps4l, ORFmut, or NC. (B) WB analysis of cleaved caspase-9 in hypoxic and control PASMCs transfected with lvRps4l, ORFmut, or NC. All values are represented as the mean \pm SEM (*p<0.05, **p < 0.01 and NS, no significance; $n \ge 3$).





The si-3 was used in this study. (A) Interference efficiency of RPS6 in mRNA level. (B) Interference efficiency of RPS6 in protein level. Si-3 was used for the study. All values are represented as the mean \pm SEM (***p < 0.001 and NS, no significance; n \geq 3).

Figure S8









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Hyp+ Hyp+ si-NC 10µg/ml+ si-NC

Nor+ si-NC

CyclinA

CyclinD

CyclinE

CDK1

CDK2

CDK4

Hyp+ 10µg/ml+ si-RPS6

54kDa

31kDa

53kDa

33kDa

33kDa

34kDa





1.0

0.5

В

PCNA

β-actin

36kDa

43kDa

Hyp+ lv-mut+ si-RPS6

NS

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NS

NS

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OIHO

NS

proliferation by inhibiting RPS6 phosphorylation. (A) MTT assay in hypoxic and control PASMCs co-transfected with lvRps4l, ORFmut, or lv-NC and si-RPS6 or si-NC. (B) WB analysis of PCNA in hypoxic and control PASMCs co-transfected with lvRps4l, ORFmut, or lv-NC and si-RPS6 or si-NC. (C) WB analysis of cyclin A, cyclin D, cyclin E, CDK1, CDK2, and CDK4 in hypoxic and control PASMCs co-transfected with si-RPS6 or si-NC. (D) MTT assay in hypoxic and control PASMCs simultaneously add 10ug/ml exogenous peptide RPS4XL and si-NC or si-RPS6. (E) WB analysis of PCNA in hypoxic and control PASMCs simultaneously add 10ug/ml exogenous peptide RPS4XL and si-NC or si-RPS6. (E) WB analysis of PCNA in hypoxic and control PASMCs simultaneously add 10ug/ml exogenous peptide RPS4XL and si-NC or si-RPS6. (F) WB analysis of cyclin A, cyclin D, cyclin E, CDK1, CDK2, and CDK4 in hypoxic and control PASMCs simultaneously add 10ug/ml exogenous peptide RPS4XL and si-NC or si-RPS6. All values are represented as the mean \pm SEM (*p < 0.05, **p < 0.01, ***p < 0.001 and NS, no significance; n \geq 3). Nor, normoxia; Hyp, hypoxia.

Figure S9



Supplemental figures 9. The peptide RPS4XL is conserved in humans and inhibits the proliferation of human pulmonary artery smooth muscle cells induced by hypoxia. (A) Bioinformatics alignment of the homologous sequences of Rps41-encoded peptide RPS4XL in human and mouse. (B) WB analysis of RPS4XL in hypoxic, control human PASMCs and pulmonary hypertension patient PASMCs. (C) MTT assay in hypoxic and control human PASMCs treated with RPS4XL with concentrations of 5 ug/ml, 7.5ug/ml or 10 ug/ml. (D) WB analysis of PCNA. (E) WB analysis of cyclin A, cyclin D, cyclin E, CDK1, CDK2, and CDK4 in hypoxic and control human PASMCs treated with RPS4XL with concentrations of 5 ug/ml, 7.5ug/ml or 10 ug/ml. All values are represented as the mean \pm SEM (*p<0.05, **p<0.01 and NS, no significance; n \geq 3). Nor, normoxia; Hyp, hypoxia.





Supplemental figure 10. The efficiency of AAV9 overexpresses Rps4l, mutates Rps4l and interferes with of RPS6. (A) qPCR for the efficiency of AAV9 overexpression of Rps4l and mut-Rps4l. (B) qPCR for the efficiency of AAV9 interference with RPS6. All values are represented as the mean \pm SEM (*p<0.05, **p<0.01 and ***p < 0.001; n \geq 3).





Supplemental figure 11. RPS4XL blocks hypoxia-induced pulmonary hypertension in vivo through RPS6. (A-E) (A) RV systolic pressure (RVSP) (B) Right ventricular (RV)/left ventricular (LV)+S weight ratio (C) PAAT (D) PAVTI and (E) HE staining (Scale bar = 200 μ m) of hypoxic mouse model infected with AAV9-lv-NC, AAV9-lv-Rps4l, AAV9-lv-mut and AAV9-si-NC, AAV9-si-RPS6. All values are represented as the mean \pm SEM (*p < 0.05, **p < 0.01, ***p < 0.001 NS, and no significance; n \geq 3). Nor, normoxia; Hyp, hypoxia.