



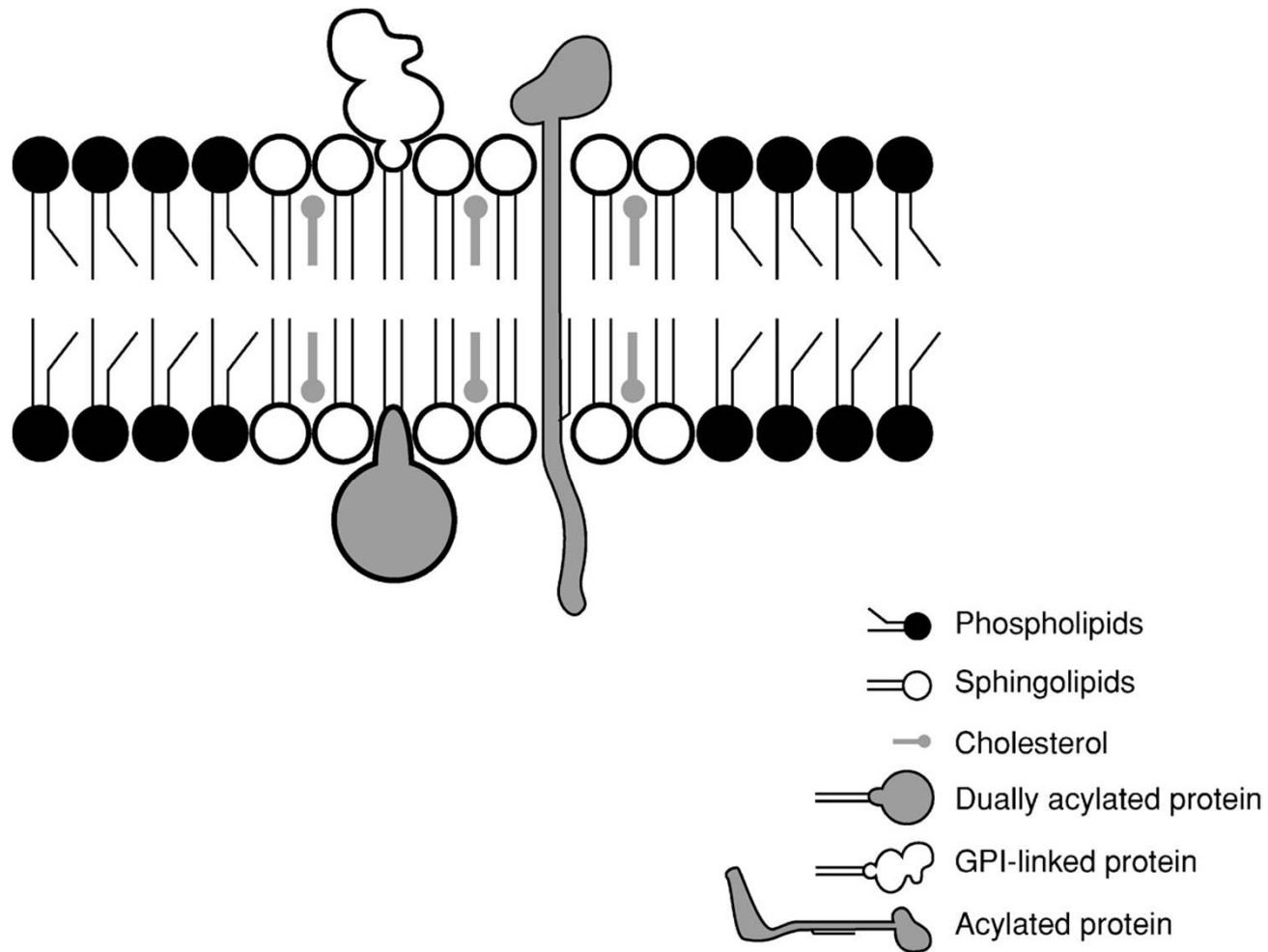
**E3 ligase TRIM72 negatively regulates
myogenesis by IRS-1 ubiquitination**

Young-Gyu Ko

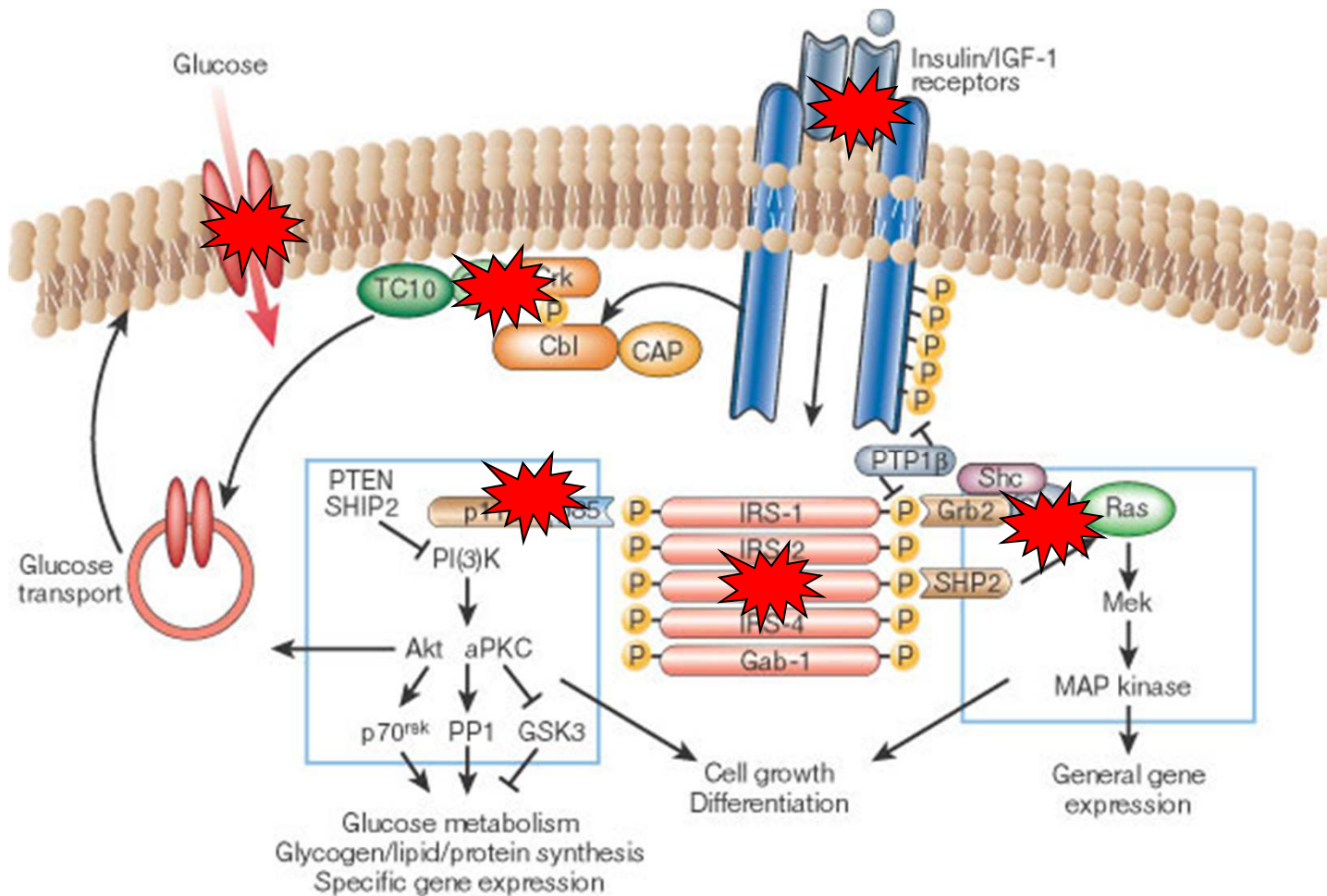
**College of Life Science and Biotechnology
Korea University**

MyHC immunofluorescence

Lipid rafts are plasma membrane compartments composed of cholesterol and glycosphingolipids.

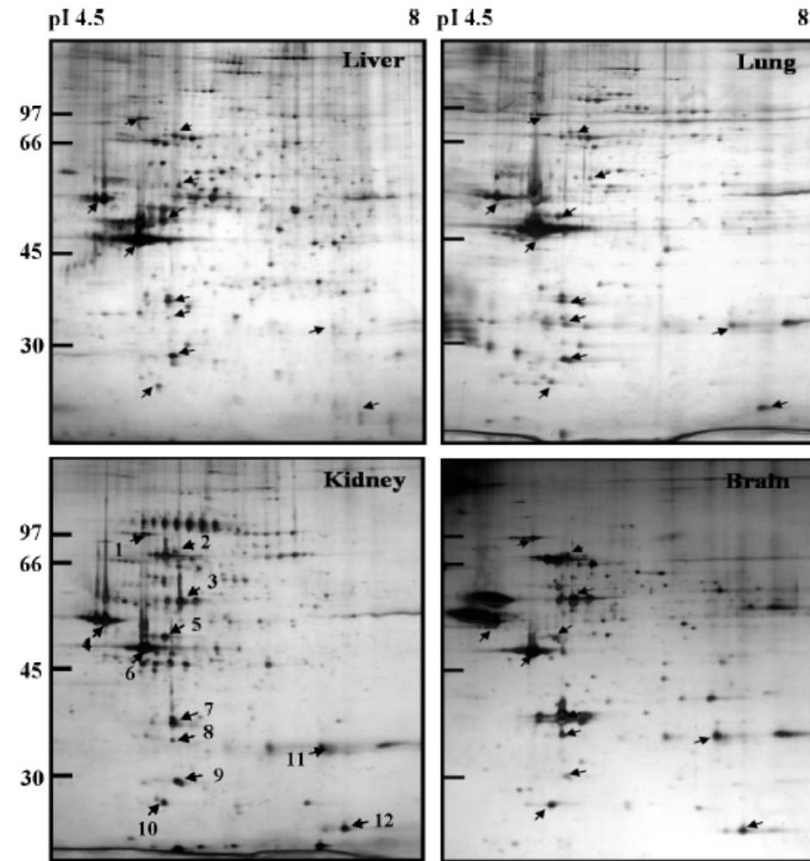
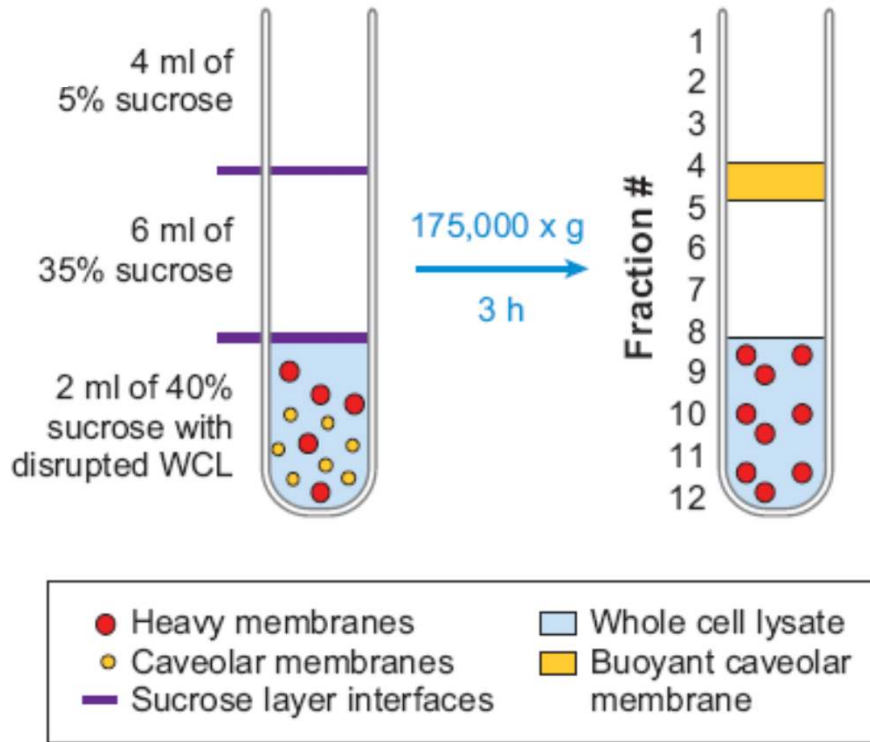


Insulin action through lipid rafts



★ Signaling molecules found in lipid rafts; IR, IRS, Grb2, Sos, Ras, PI-3-K, TC10, Cbl, CAP, and GLUT4

Identification of novel signaling molecules by lipid raft proteome

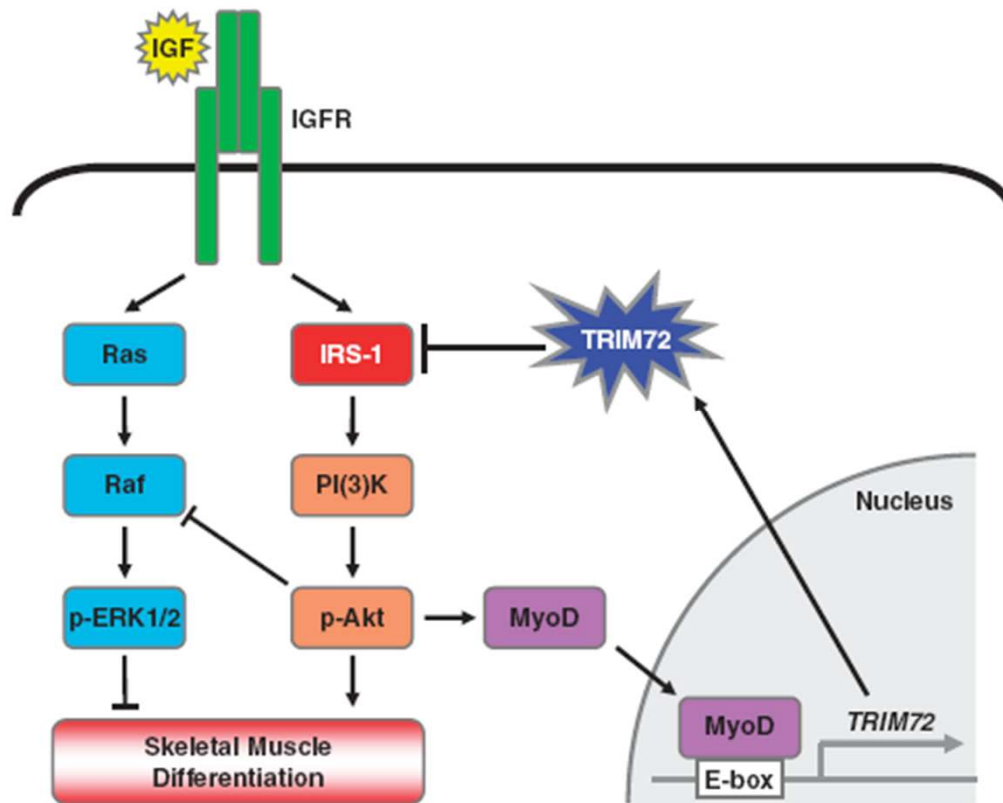


지질래프트는 신호전달물질을 농축하고 있어서 새로운 신호전달물질의 동정에 매우 유리

Functional proteomics of lipid rafts



Functional proteomics of lipid rafts



Lipid raft proteome papers in our lab

TRIM72, gC1qR, surface OXPHOS

JBC, 2011; **a paper of the week**

Expert Rev. Proteomics, 2010

CDD, 2010

BBRC, 2010

Proteomics, 2010; **cover paper**

Proteomics, 2009; **issue paper**

Proteomics, 2006; **cover paper, 51 citations**

Diabetologia, 2006; **82 citations**

EMM, 2004; **66 citations**

Proteomics, 2004; 30 citations

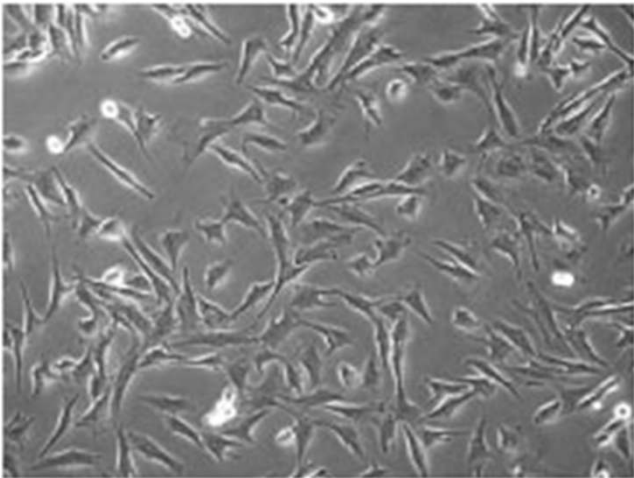
Proteomics, 2004; **108 citations**

The function of TRIM72

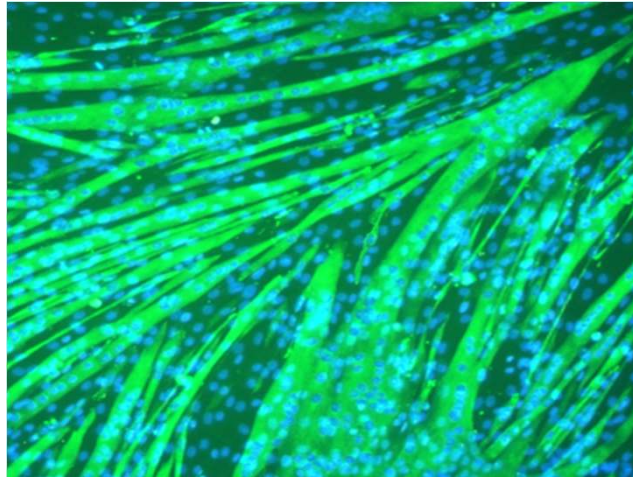
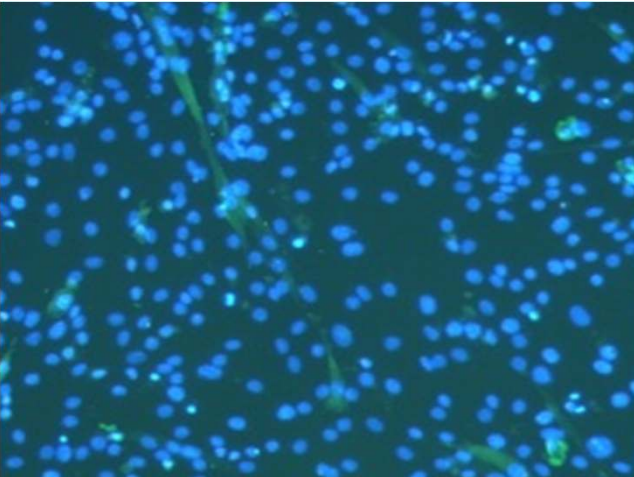
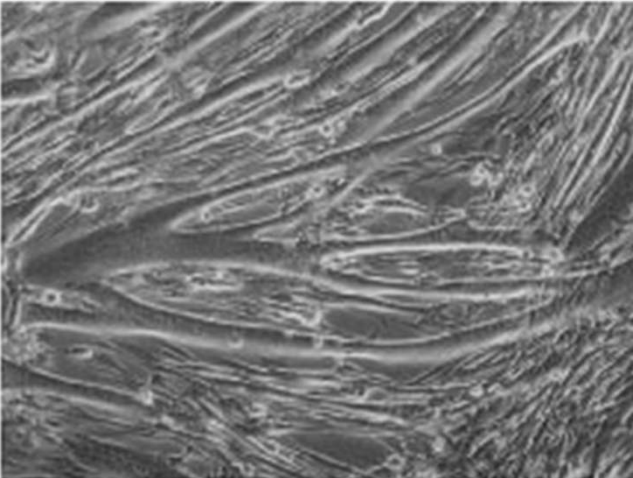
Cell Death Differ, 2010

C2C12 Myogenesis

Myoblasts



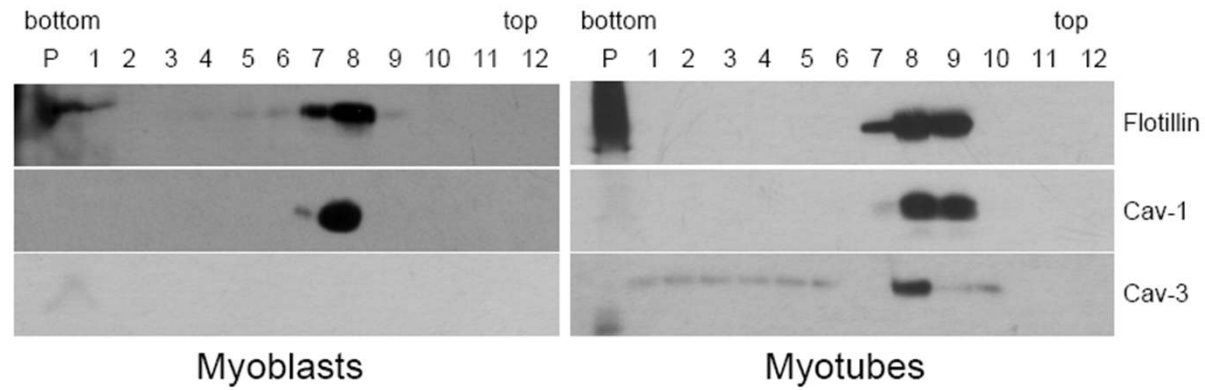
Myotubes



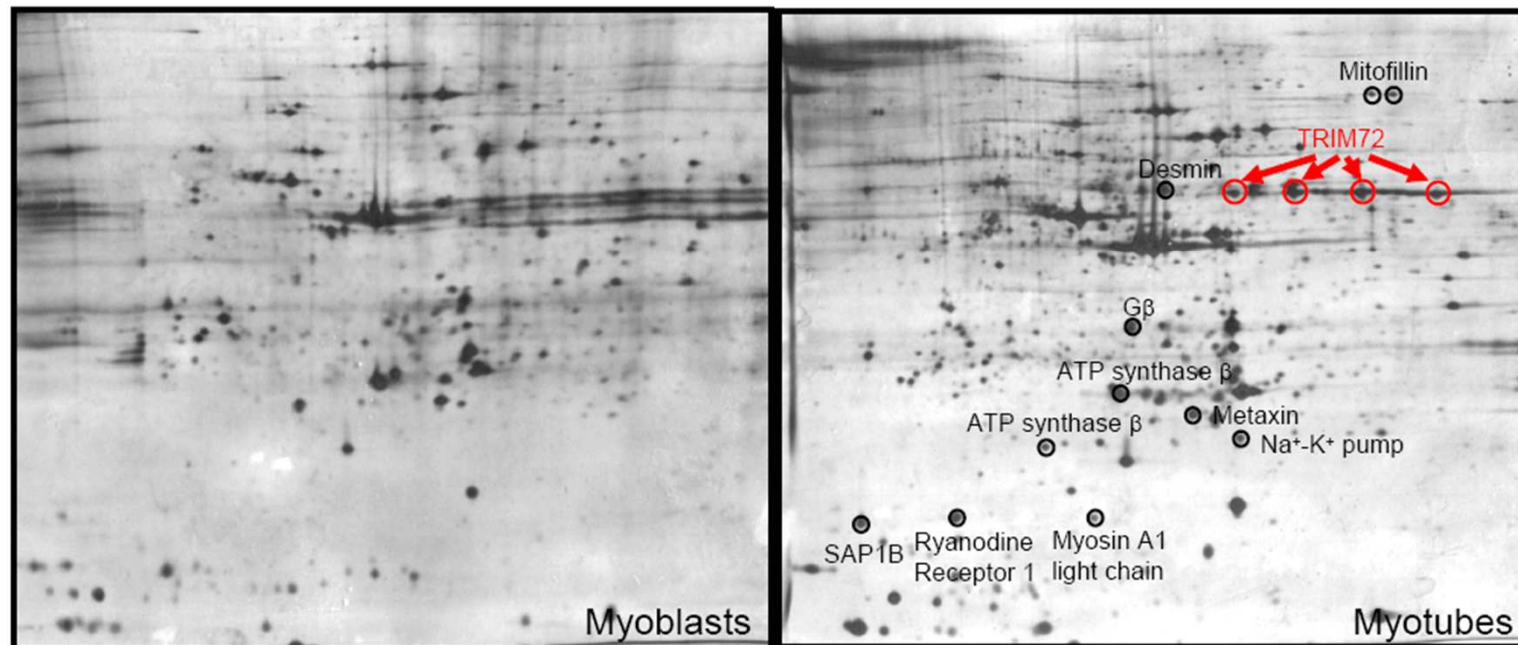
Myosin Heavy Chain

TRIM72 is found in the lipid rafts of C2C12 myotubes.

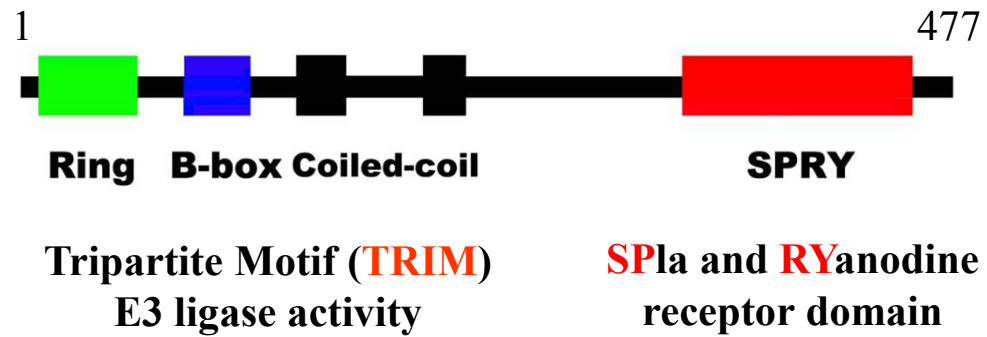
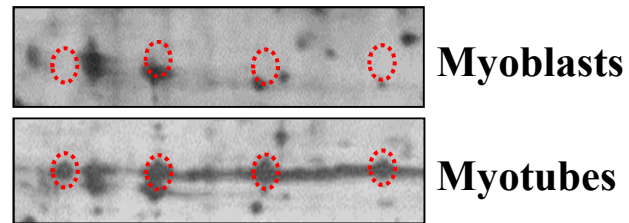
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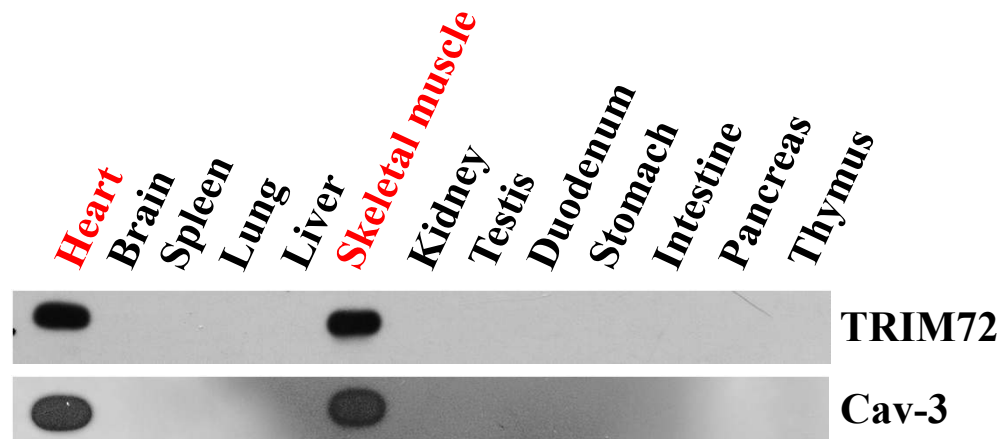
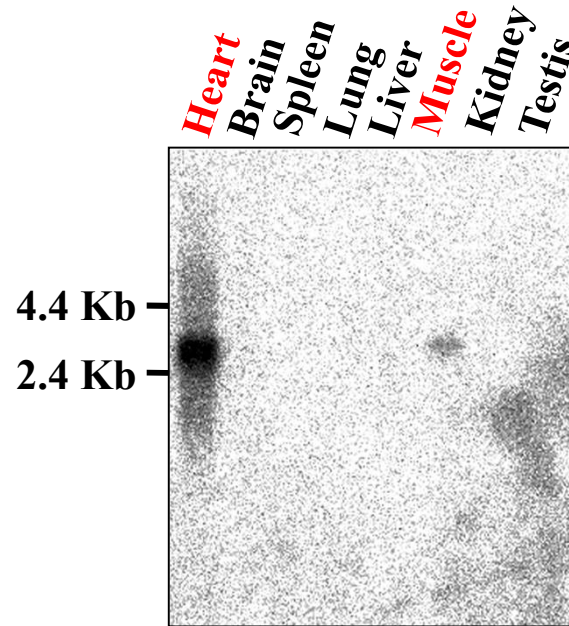
B



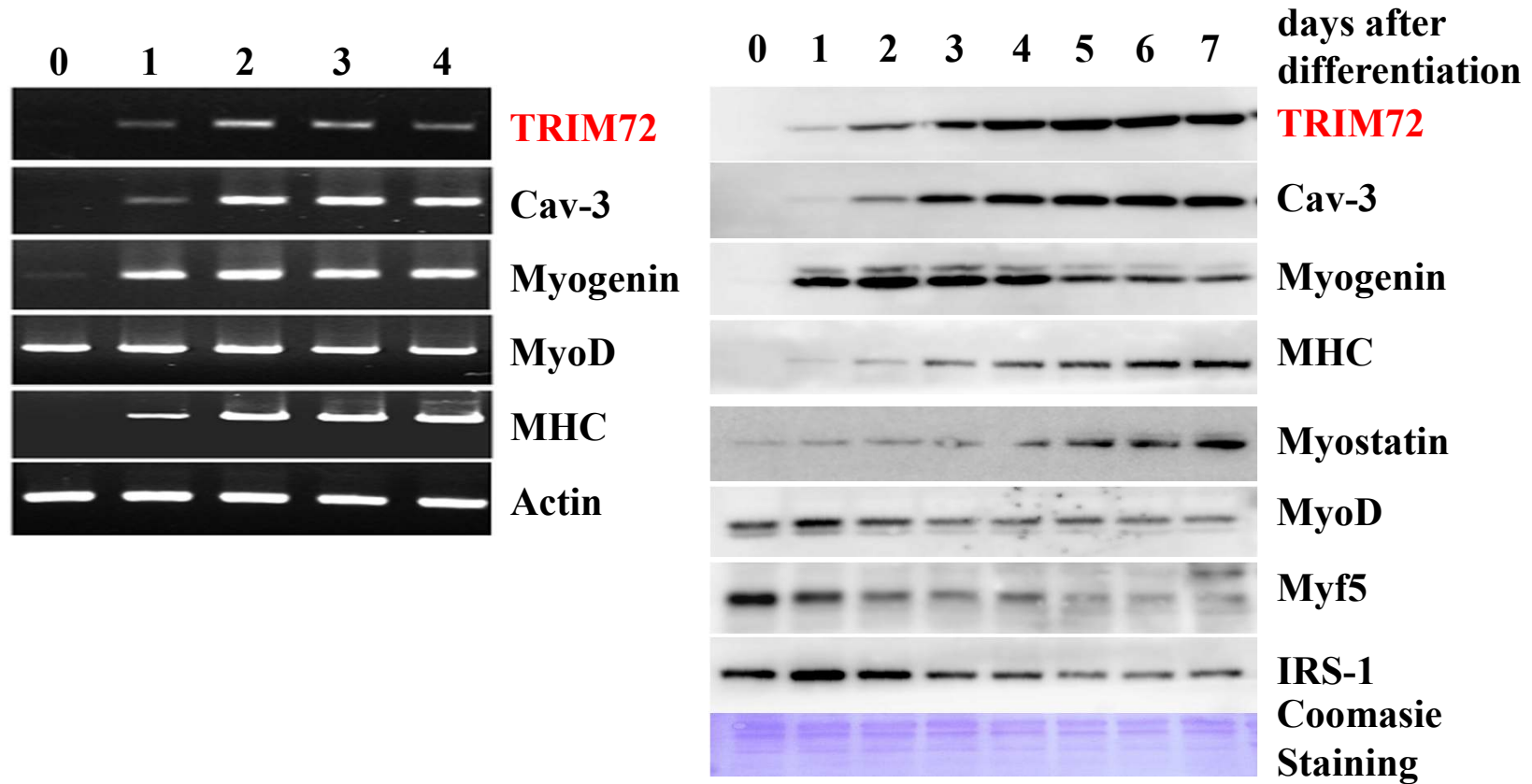
TRIM72 gene analysis



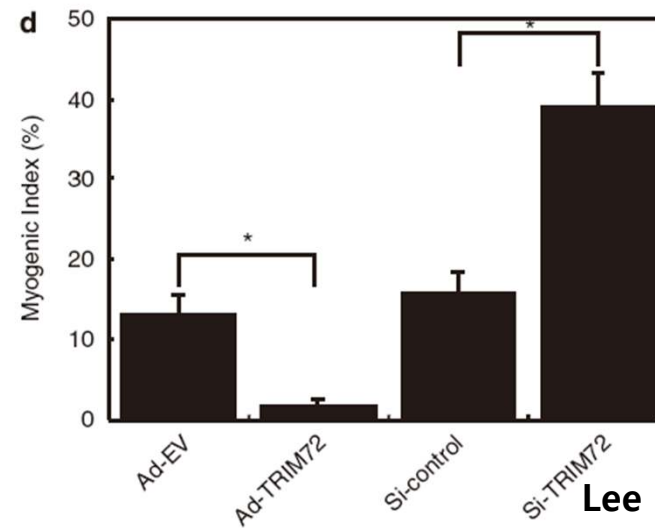
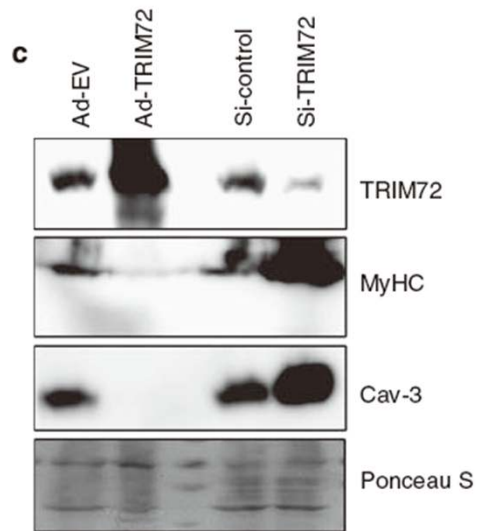
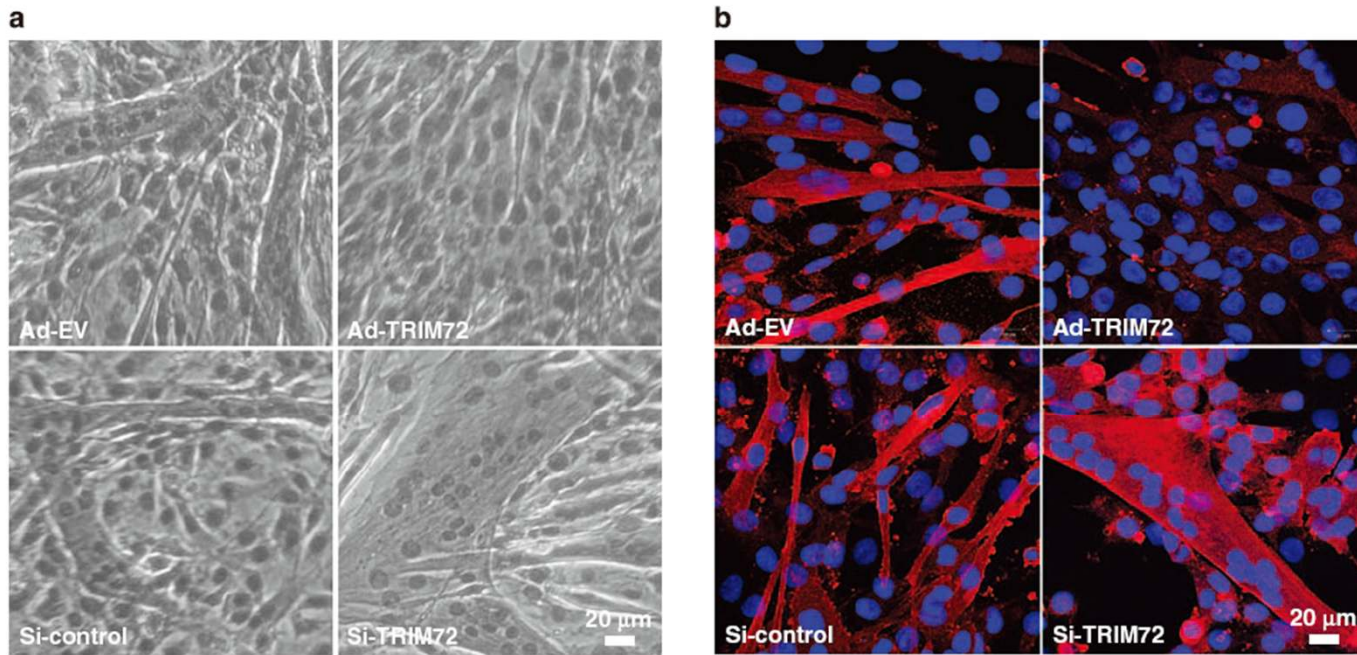
TRIM72 is specifically expressed in skeletal and cardiac muscle



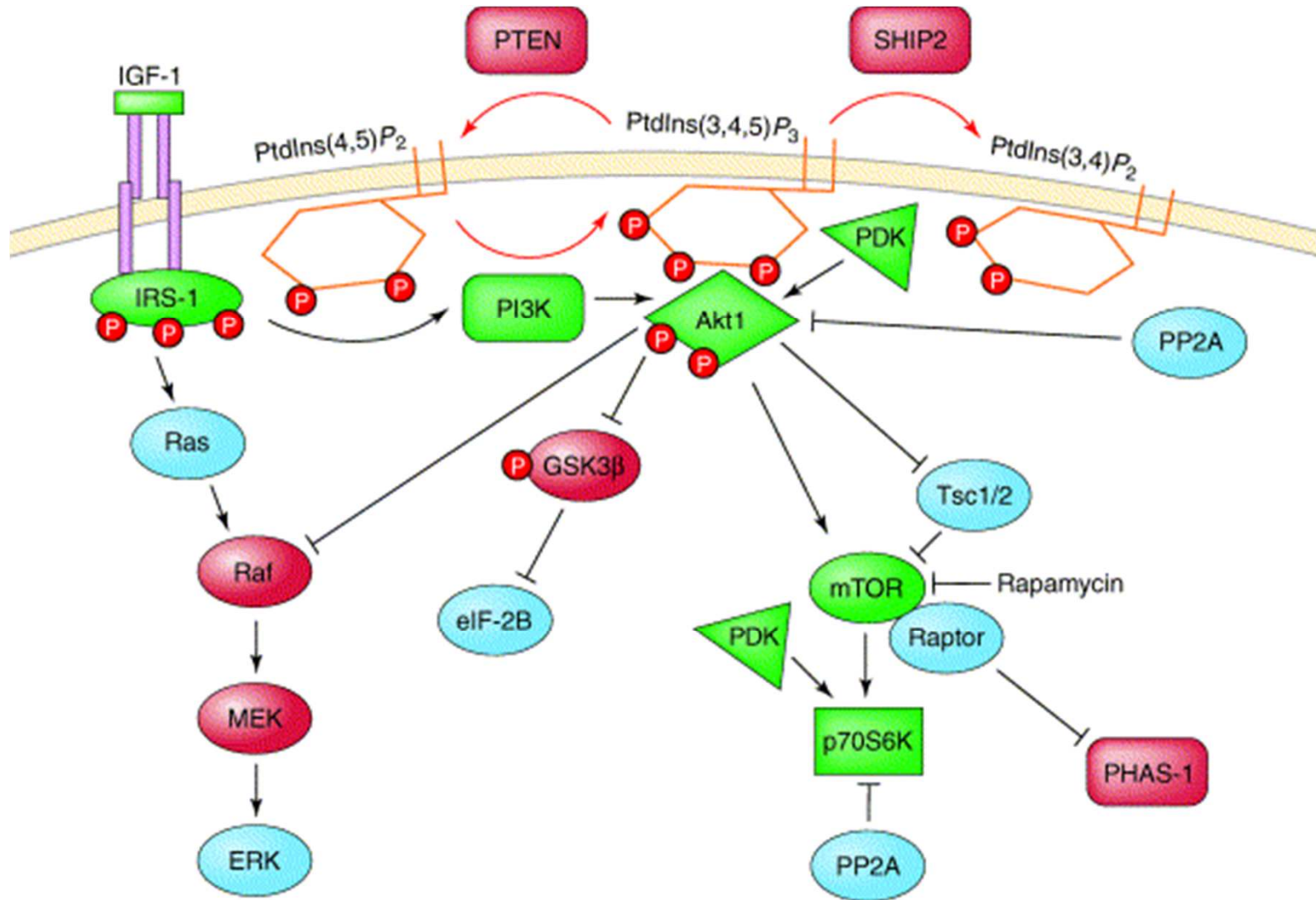
TRIM72 is highly expressed during C2C12 myogenesis



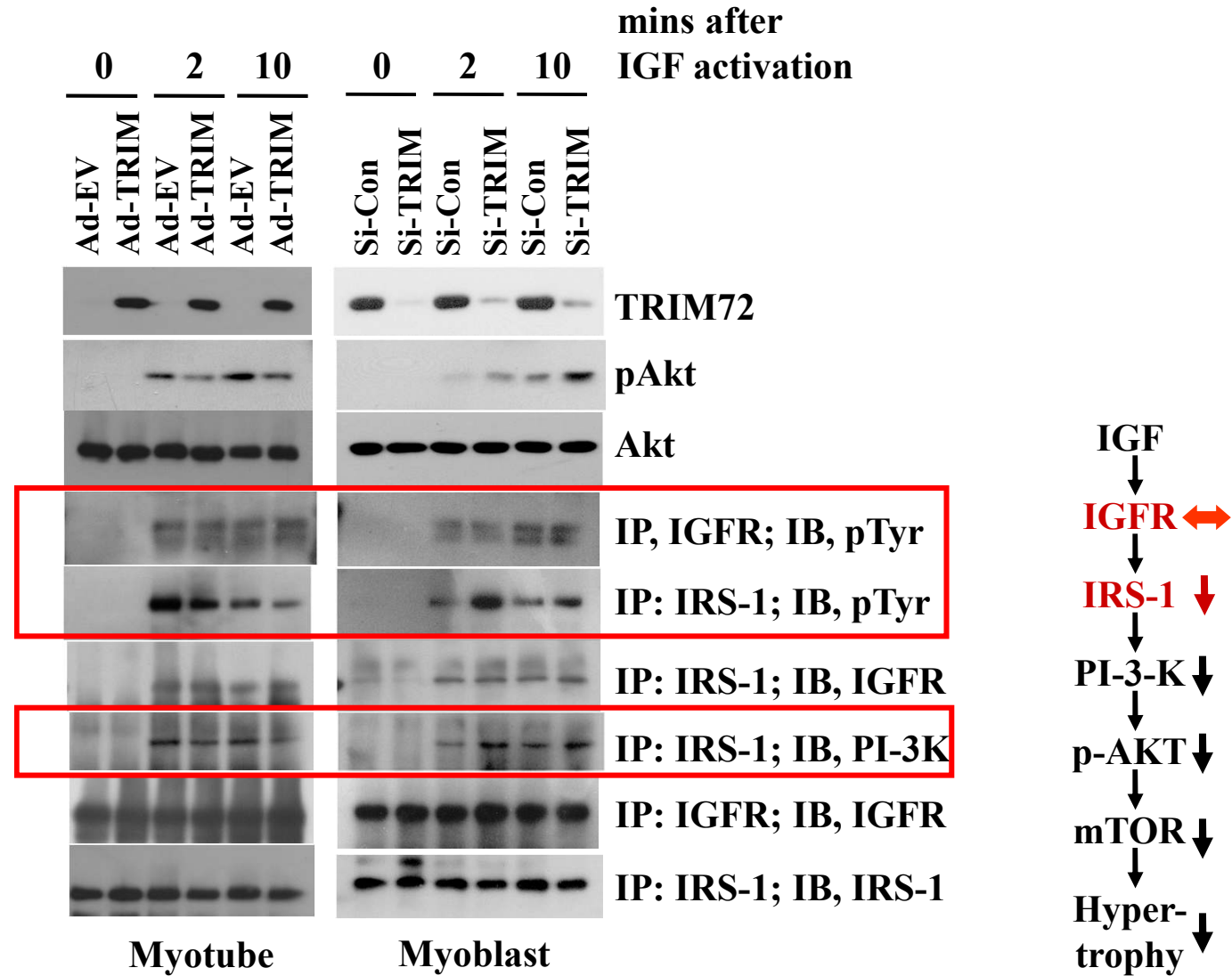
TRIM72 is a myogenesis inhibitor



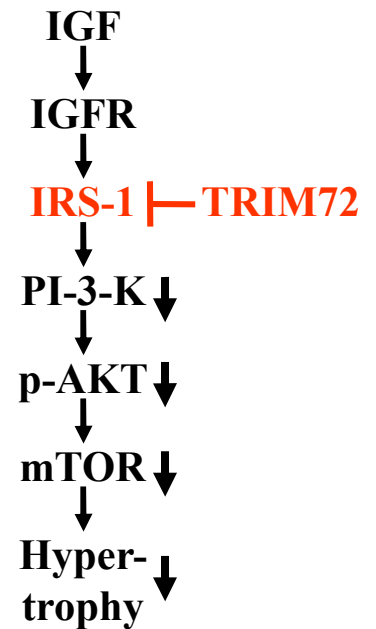
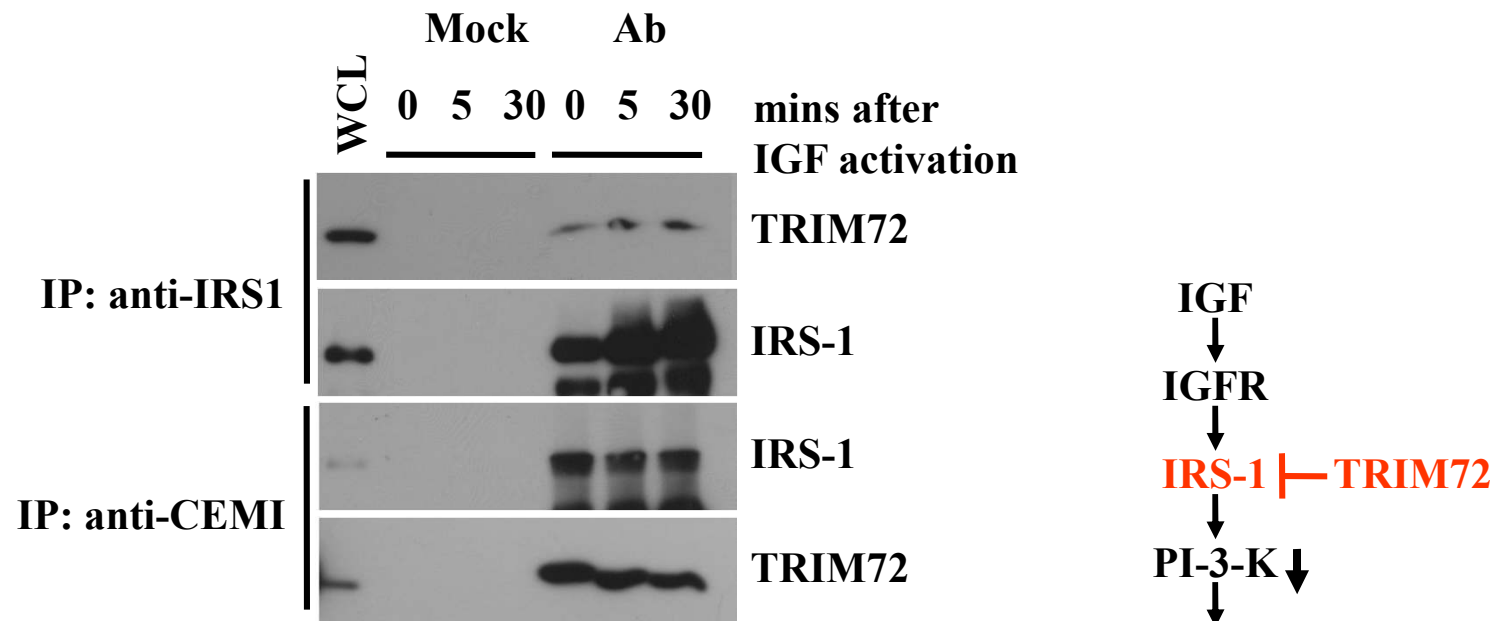
What is a molecular target of TRIM72 in IGF signaling?



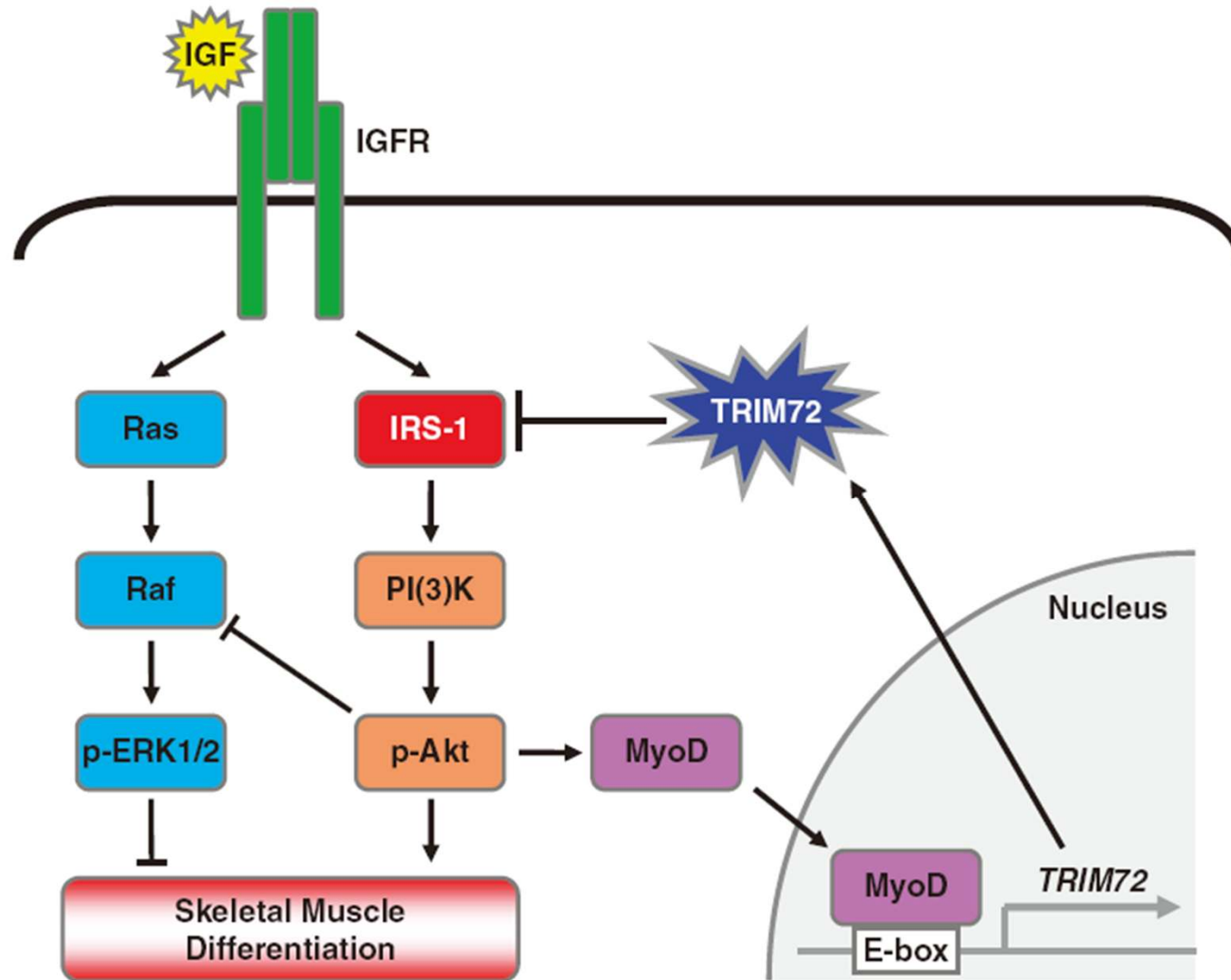
TRIM72 supresses IRS-1 phosphorylation by IGF-1.



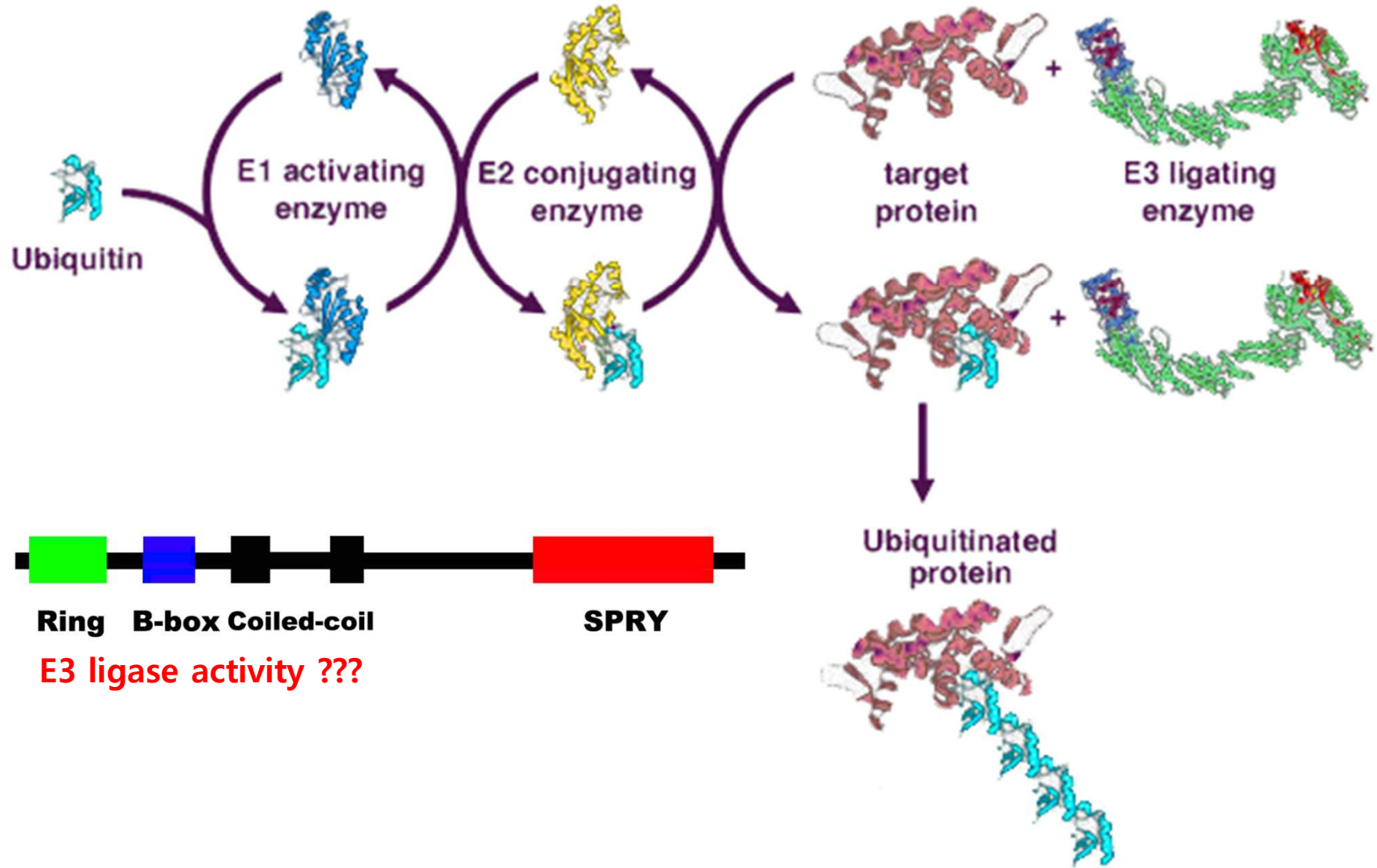
Molecular association of TRIM72 with IRS-1



TRIM72 negatively regulates myogenesis via targeting IRS-1.



Protein Ubiquitination



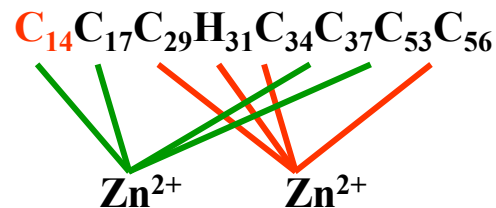
Ring Finger domain of TRIM72

Consensus Sequence of Ring Finger Domain, Zn²⁺ binding site

CX₂CX₍₉₋₃₉₎CX₍₁₋₃₎HX₍₂₋₃₎C/HX₂CX₍₄₋₄₈₎CX₂C

TRIM72 sequence

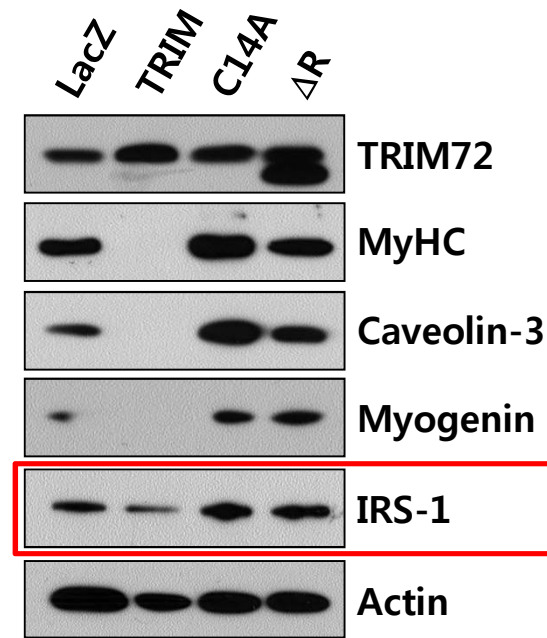
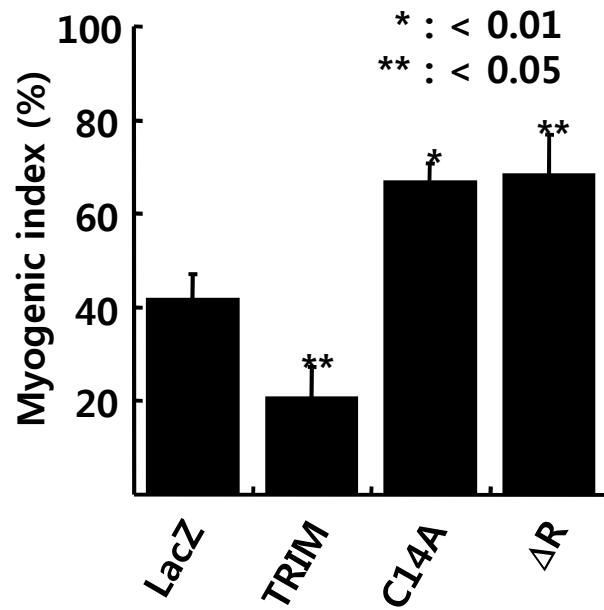
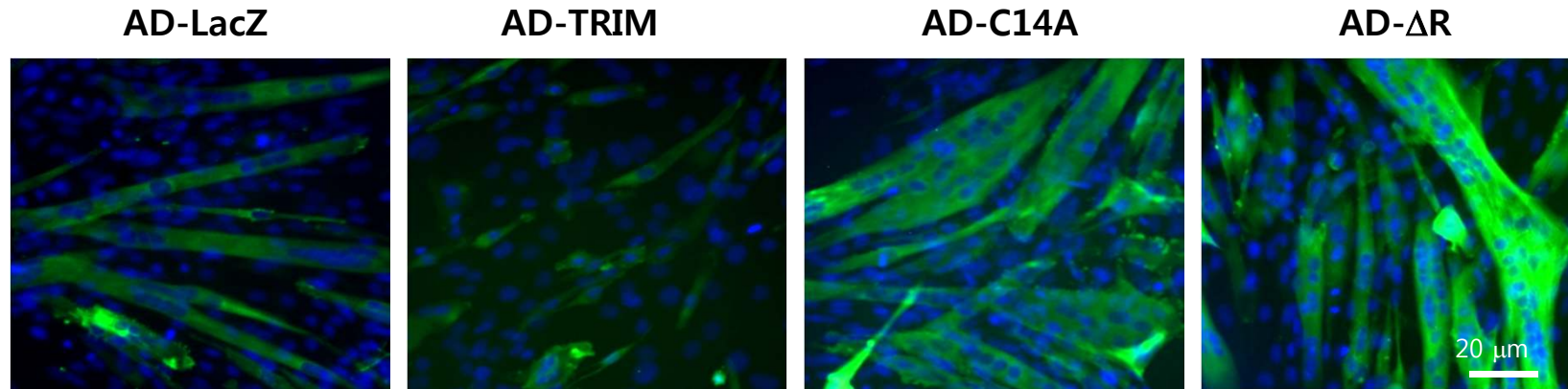
MSAAPGLLHQELS**C**PL**C**LQLFDAPVTAE**C**G**H**S**F****C**RA**C**LGRVAGEPAADGTVL**C**
P**C**QAPTRPQALSTNLQLARLVEGLAQVPQGHCEEHLDP
LSIYCEQDRALVCGV
CASLGSHRGHRLLPAAEAHARLKTQLPQQKLQLQEACMRKEKSVAVLEHQLVV
EETVRQFRGAVGEQLGKMRFVFLAALEGS
LDCEAERVGEAGVALRRELGSLNS
YLEQLRQMEKVLEEVADKPQTEFLMKYCLVTSRLQKILAESPPPARLDIQLPIISD
DFKFQVWRKMFRALMPALEELTFDPSSAHPSLVVSSSGRRVECSEQKAPPAGED
PRQFDKAVAVVAHQQLSEGEHYWEVDVGDKPRWALGVIAAEAPRRGRLHAVPS
QGLWLLGLREGKILEAHVEAKEPRALRSPERRPTRIGLYLSFGDGVLSFYDASD
ADALVPLFAFHERLPRPVYPPFDVCWHDKGKNAQPLLLVGPEGAEA



TRIM72 C14A mutant ?

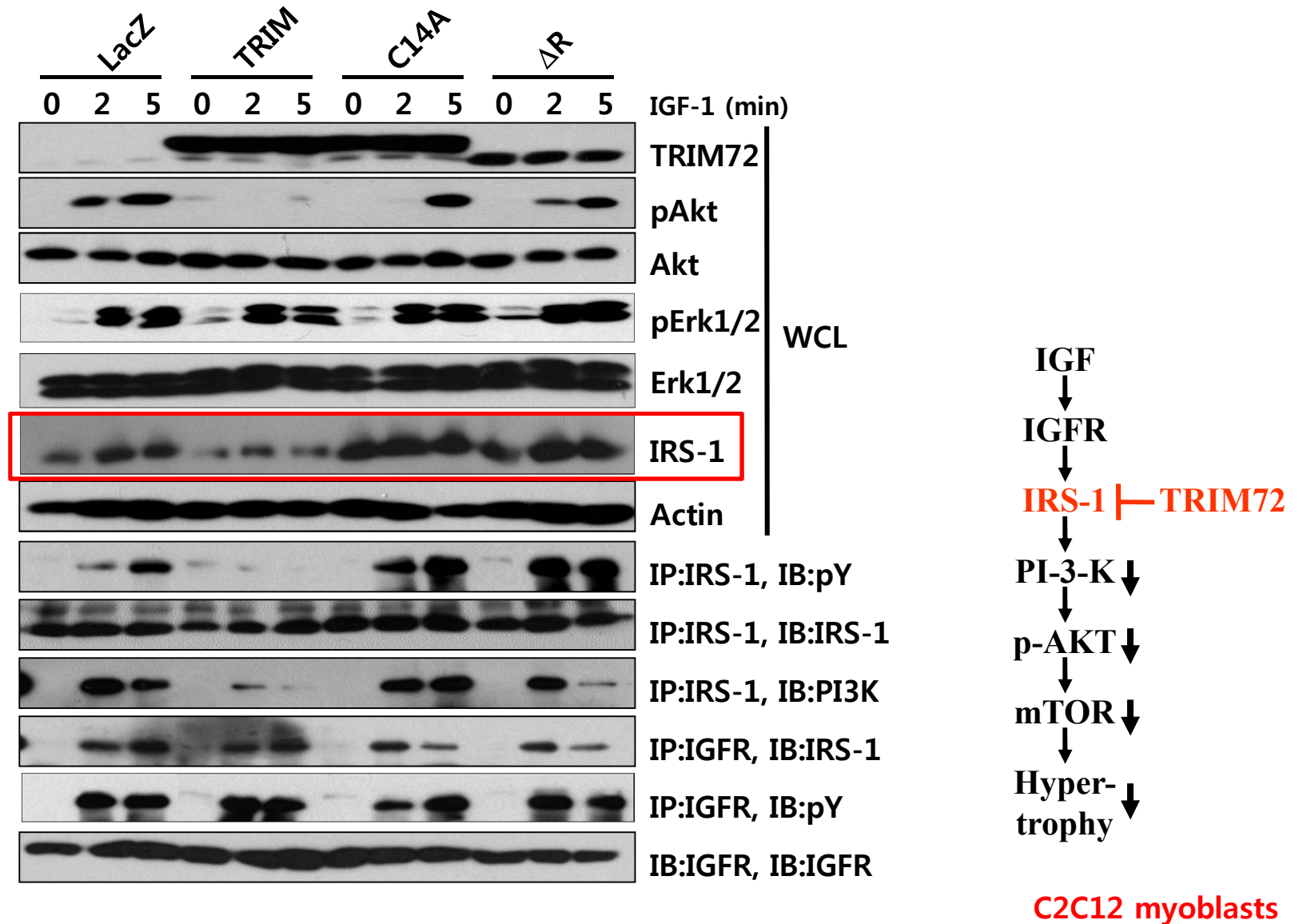
TRIM72 ΔRING mutant ?

RING domain of TRIM72 is essential for the negative regulation of myogenesis.

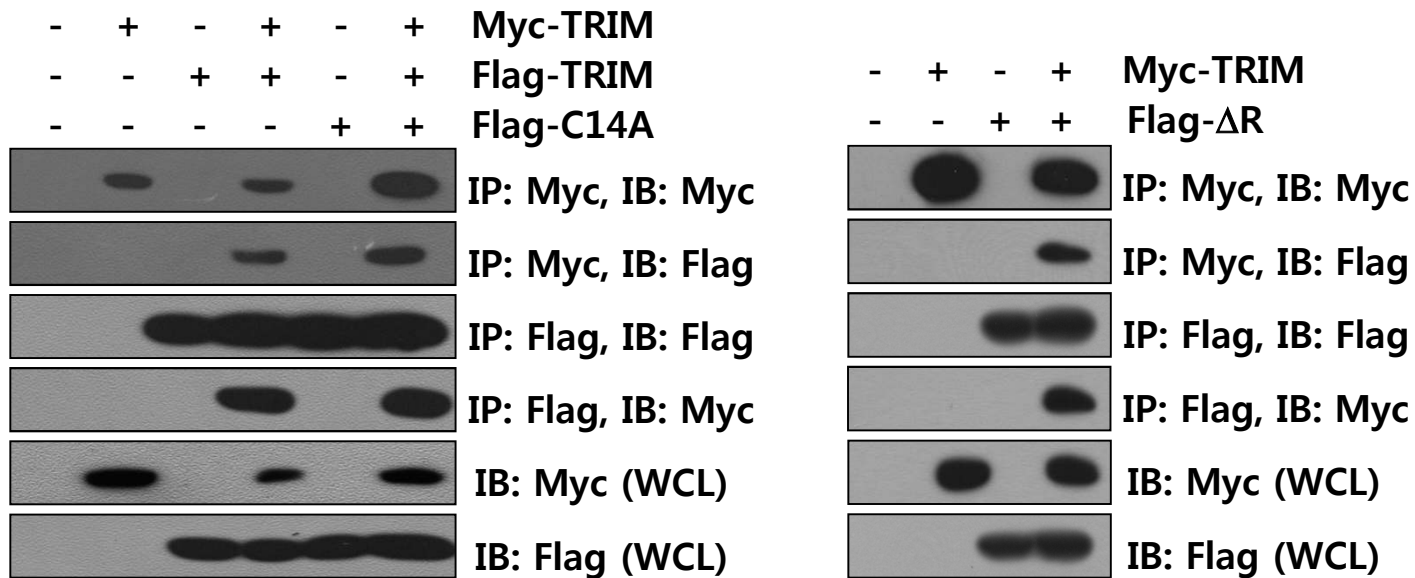


C2C12 myotubes

RING domain of TRIM72 is required for the negative regulation of IGF signaling.



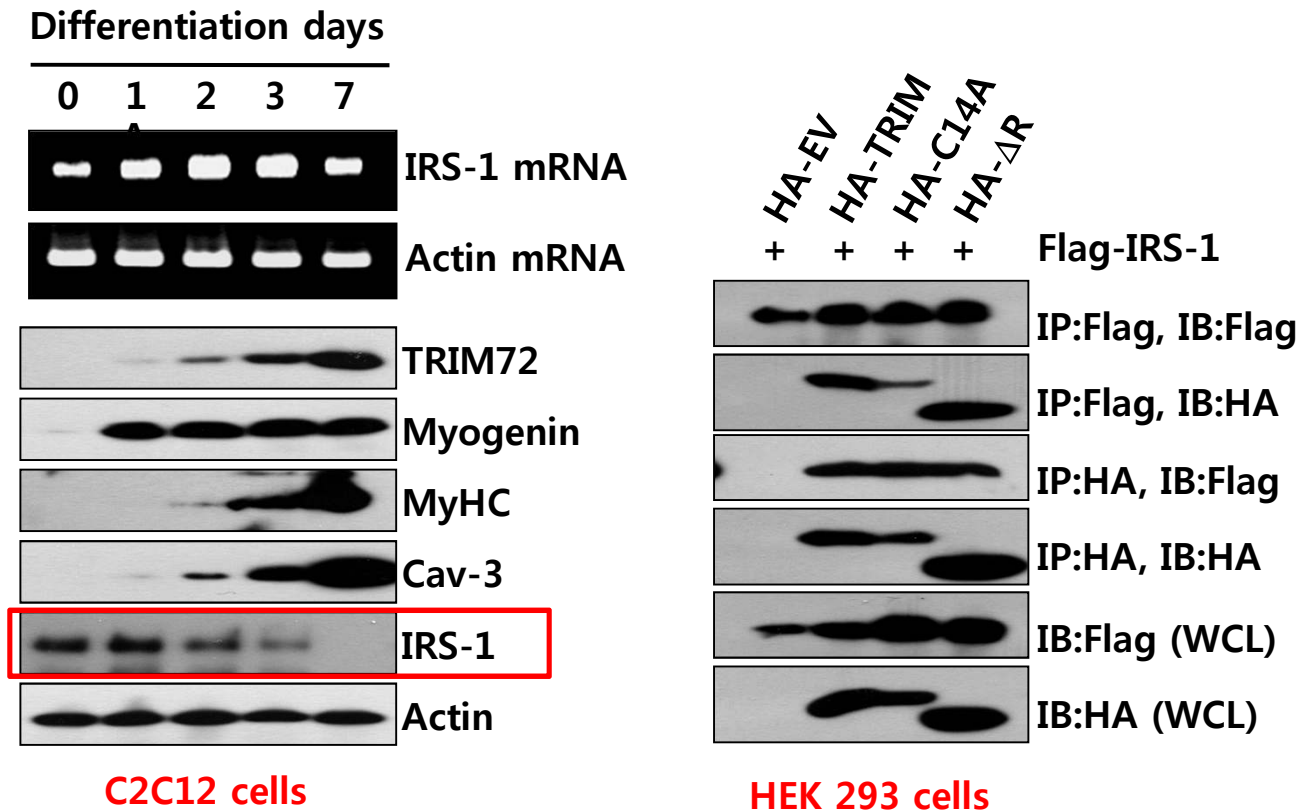
RING domain-lacking TRIM72 mutants work as dominant negative forms of endogenous TRIM72.



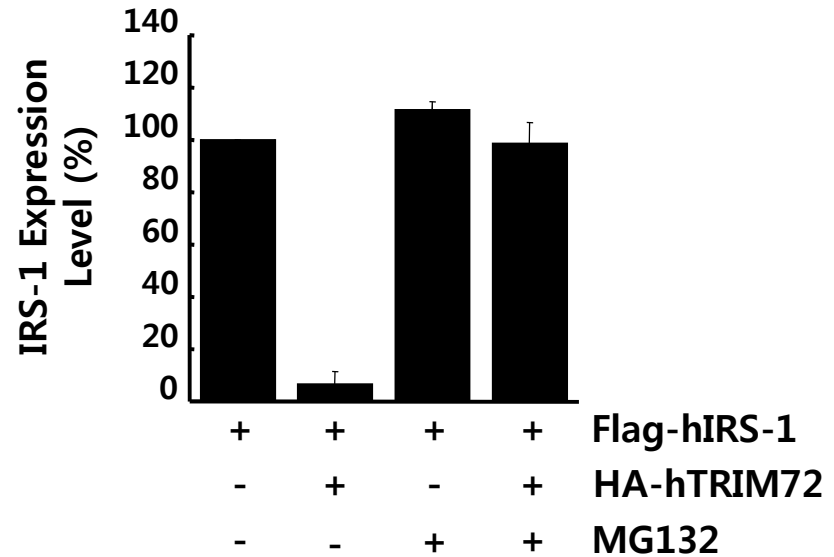
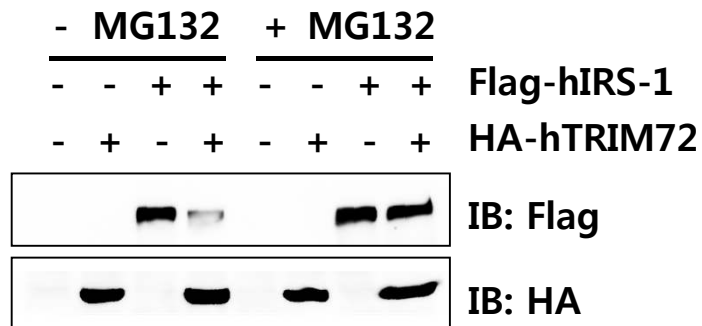
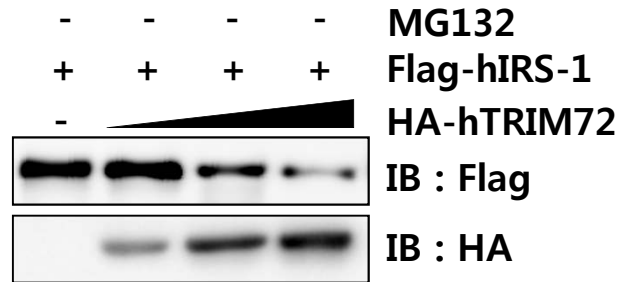
TRIM72 oligomerization

HEK 293 cells

RING domain does not prevent binding of TRIM72 to IRS-1

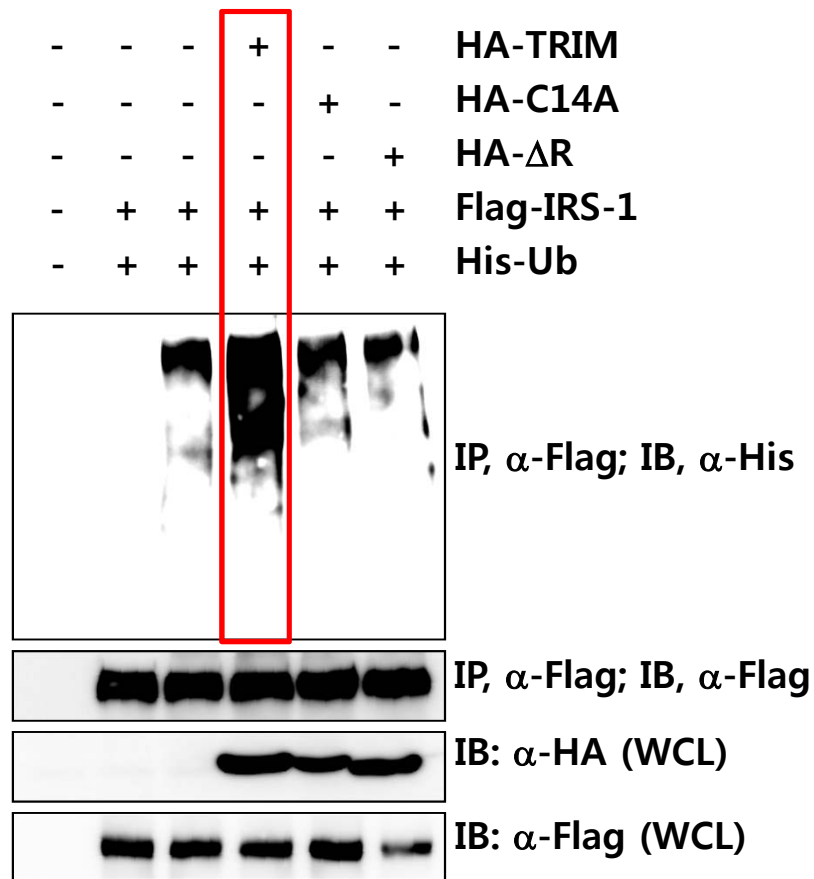


RING domain of TRIM72 is required for IRS-1 degradation.



HEK 293 cells

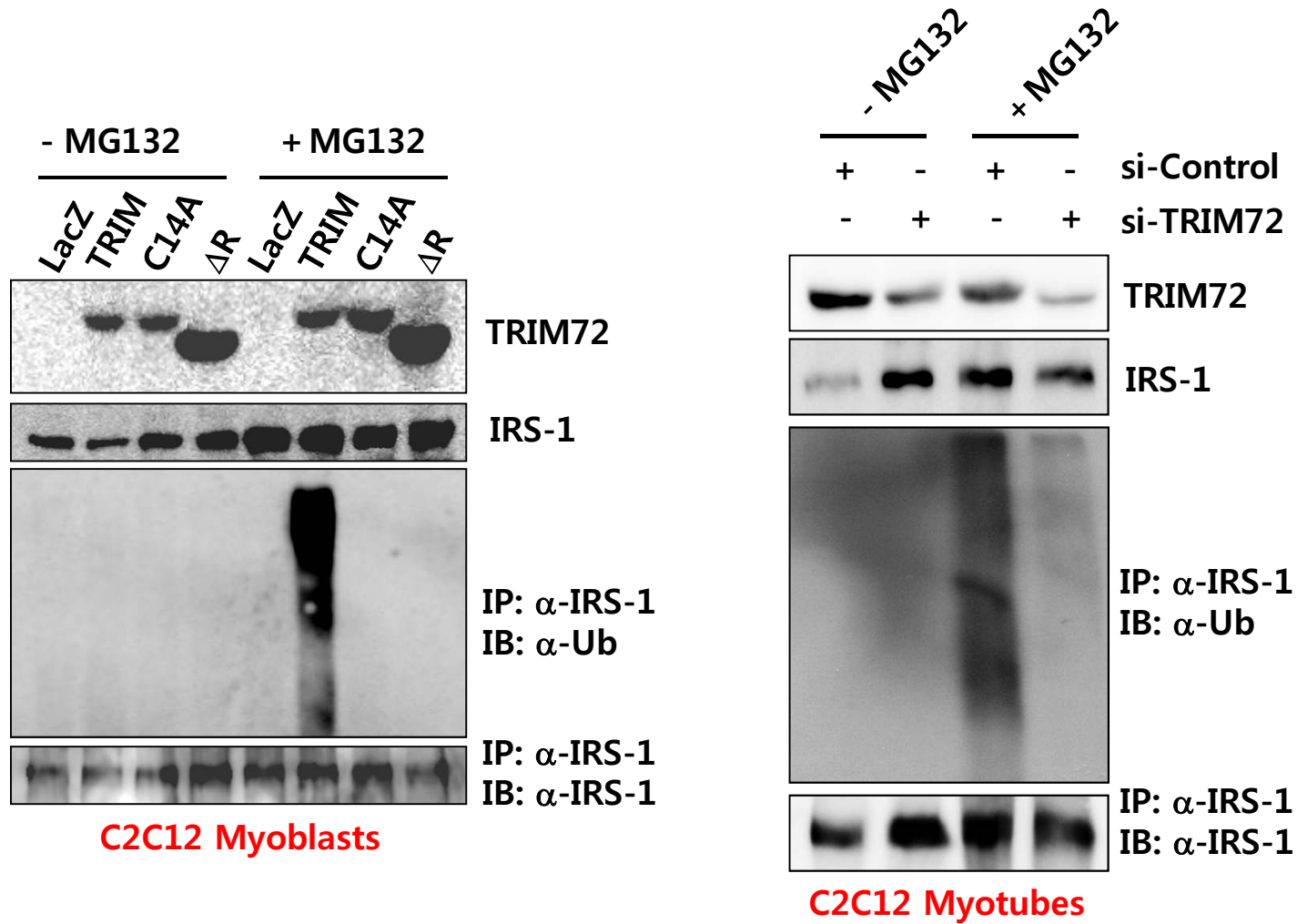
RING domain of TRIM72 is required for IRS-1 ubiquitination.



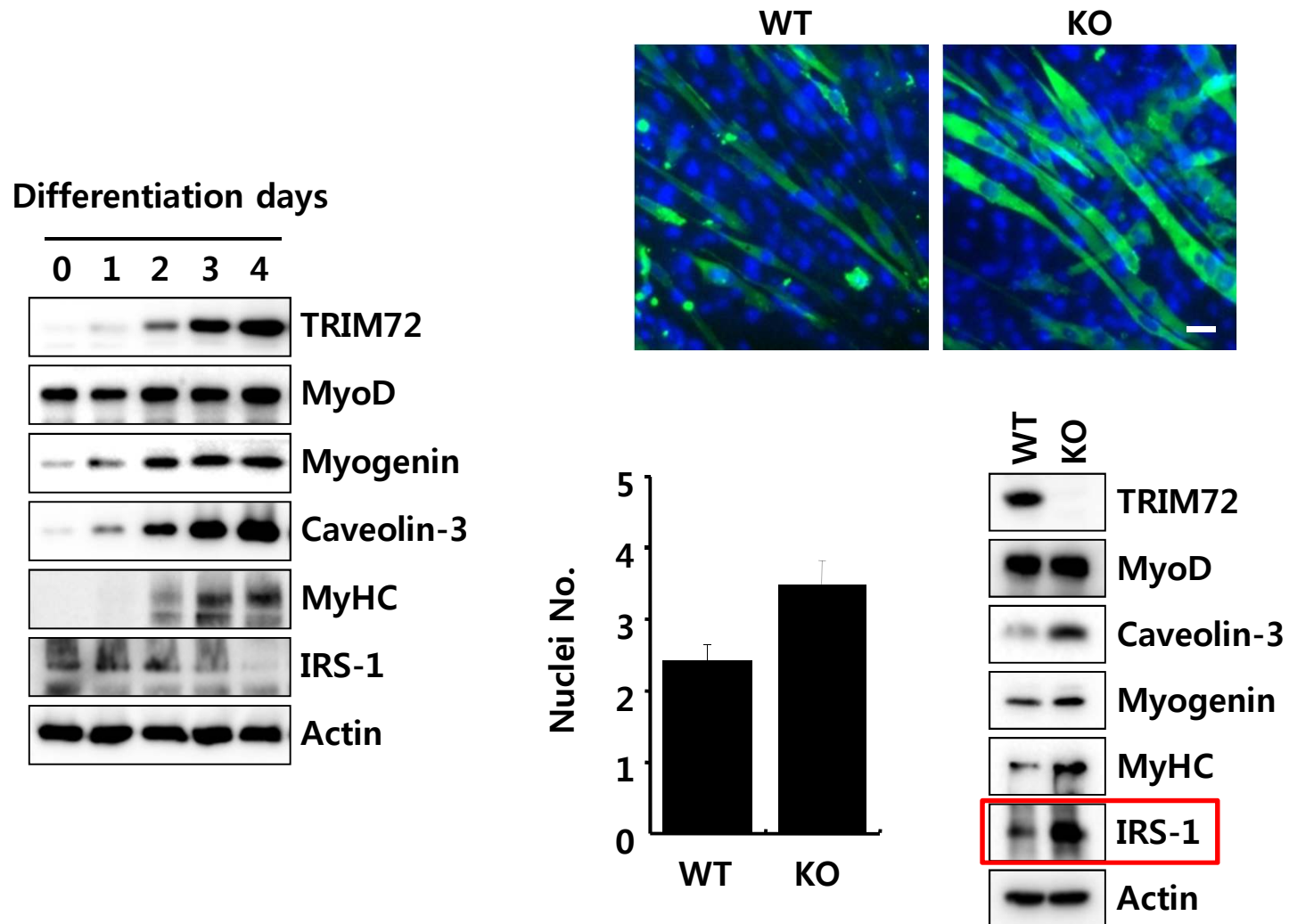
In the presence of MG132

HEK 293 cells

RING domain of TRIM72 is required for IRS-1 ubiquitination.

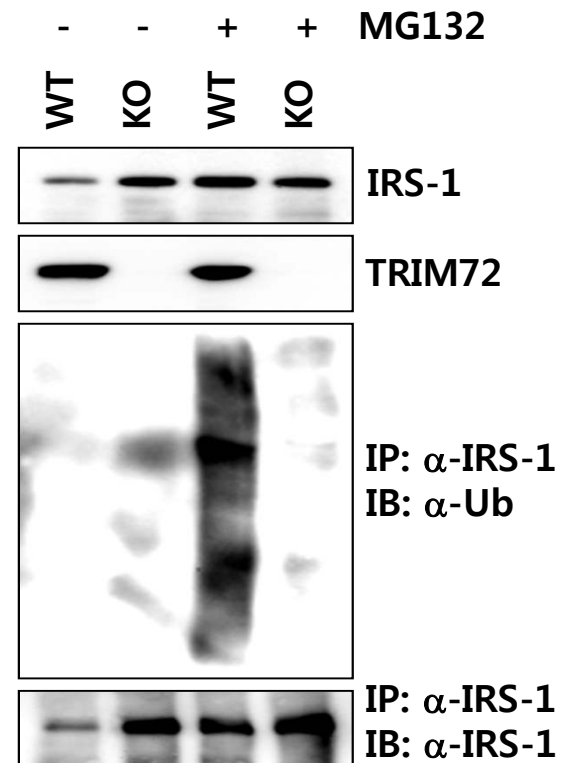


TRIM72 disruption enhances MyoD-driven myogenesis in MEFs.



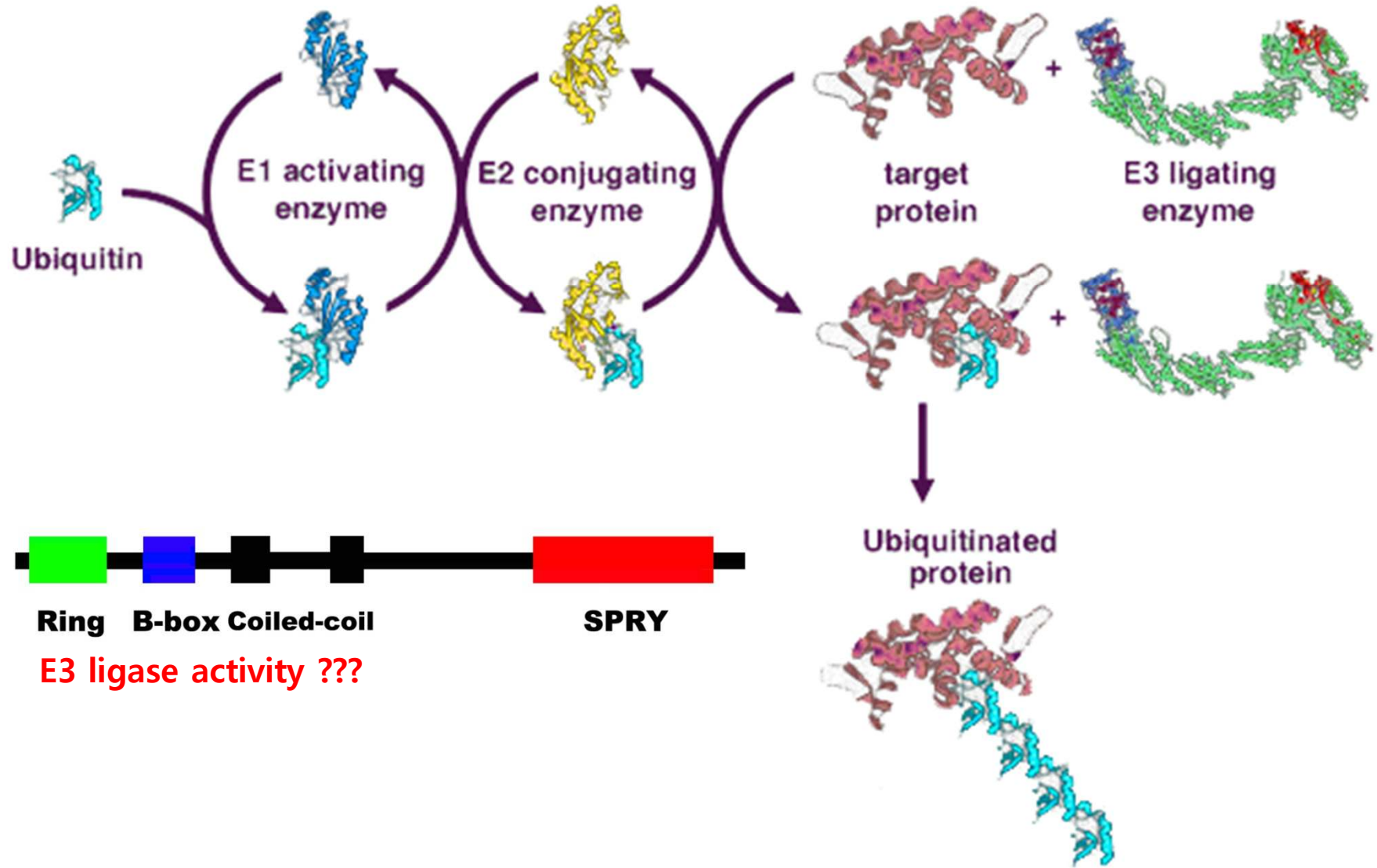
MyoD-driven myotubes from TRIM72^{+/+} and ^{-/-} MEFs

**TRIM72 disruption abolishes IRS-1 ubiquitination
in MyoD-driven myotubes of MEFs.**

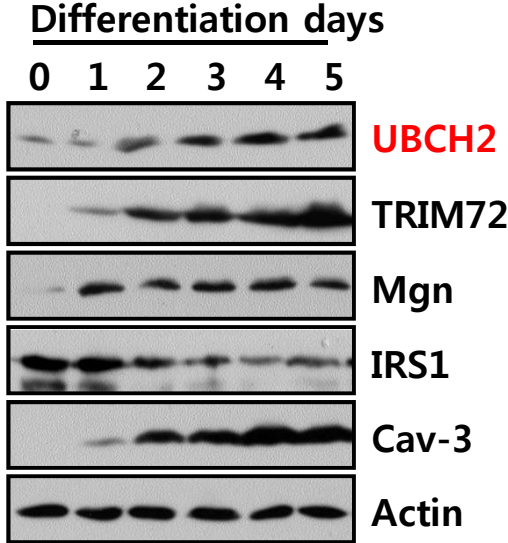
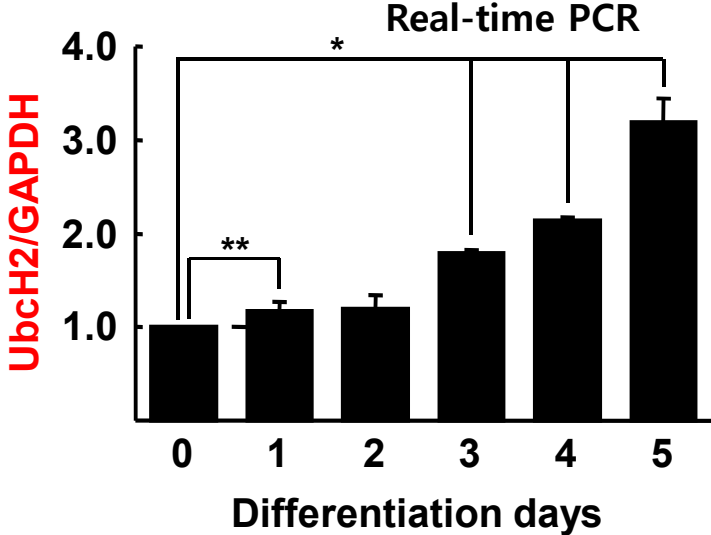
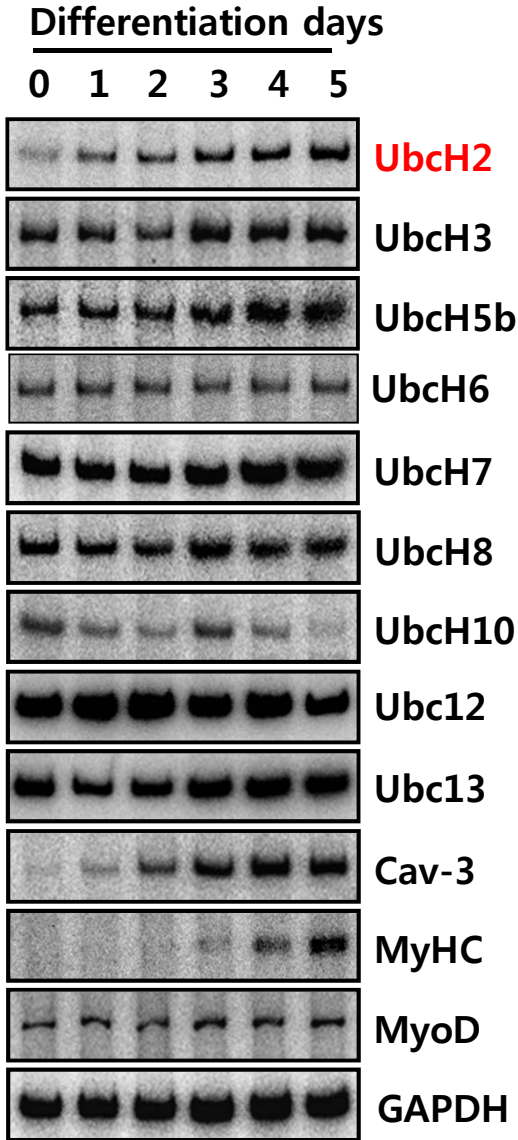


MyoD-driven myotubes from TRIM72^{+/+} and ^{-/-} MEFs

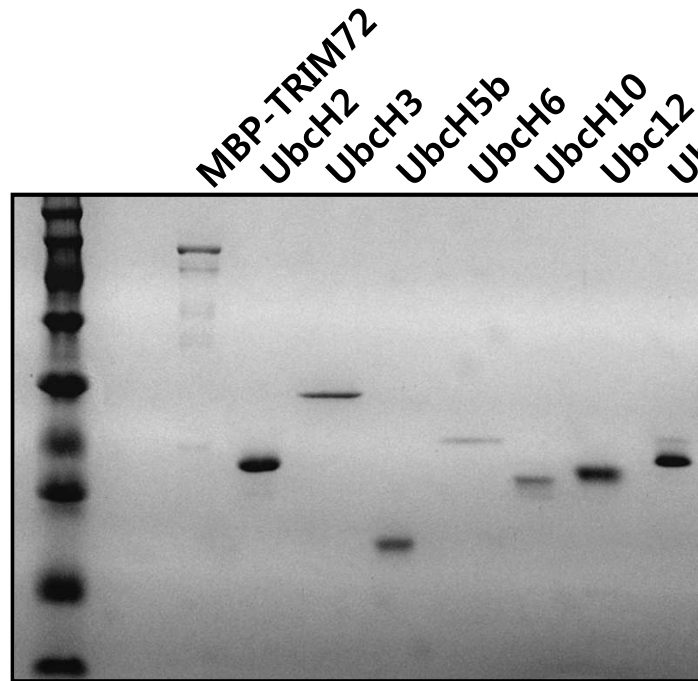
Protein Ubiquitination



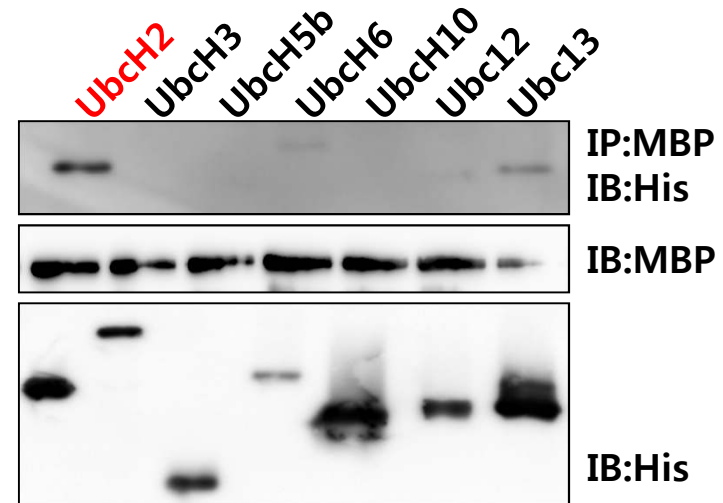
Ubch2 mRNA and protein are increased during C2C12 myogenesis



Molecular association of TRIM72 with UbcH2

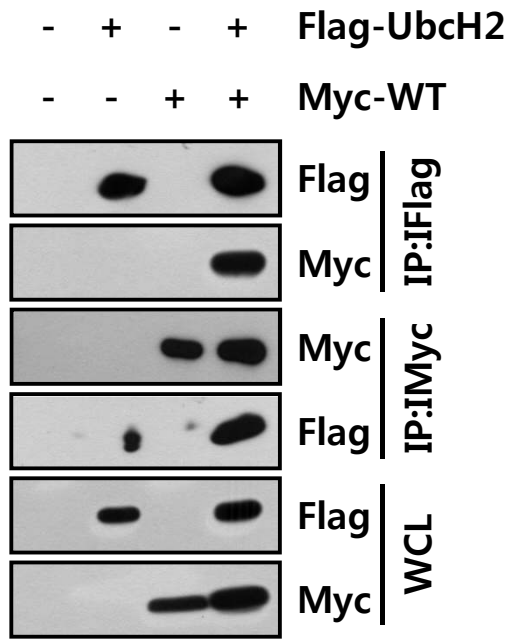


Purified TRIM72 and E2 enzymes

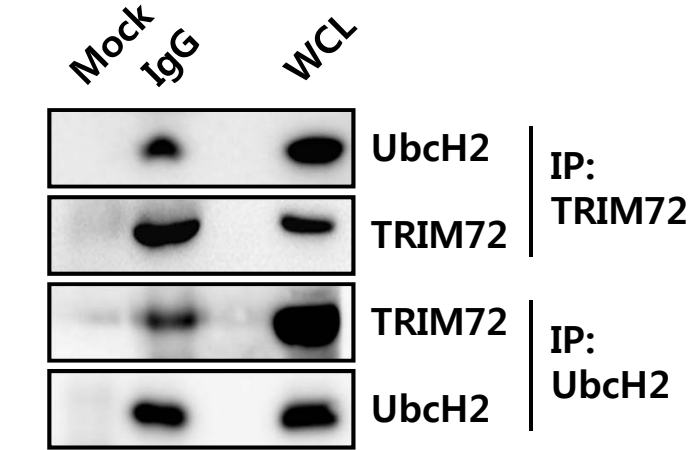


In vitro binding assay

Molecular association of TRIM72 with Ubch2

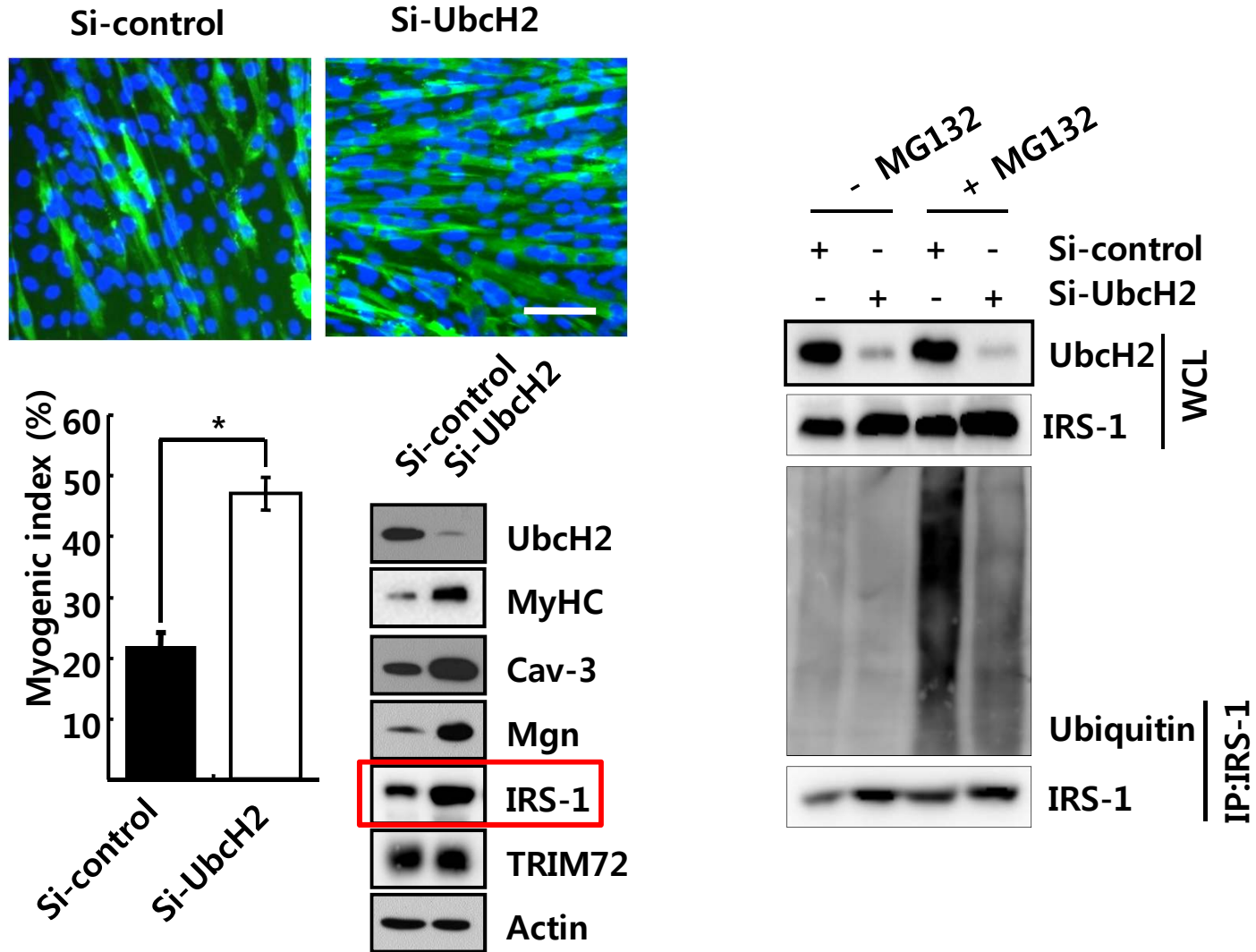


Exogenous IP
in **293 cells**



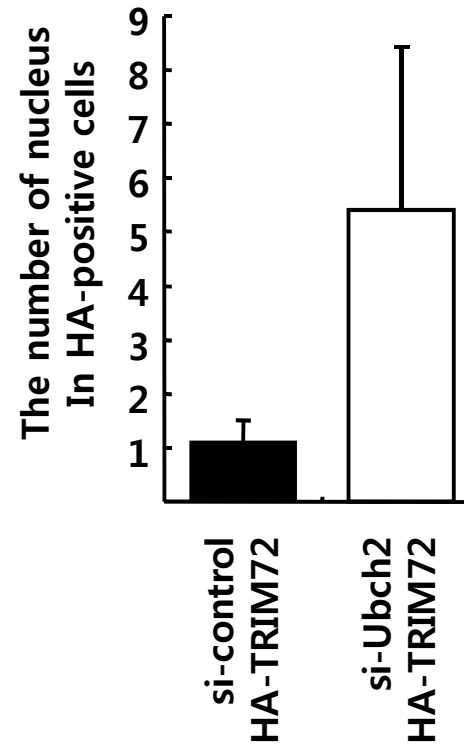
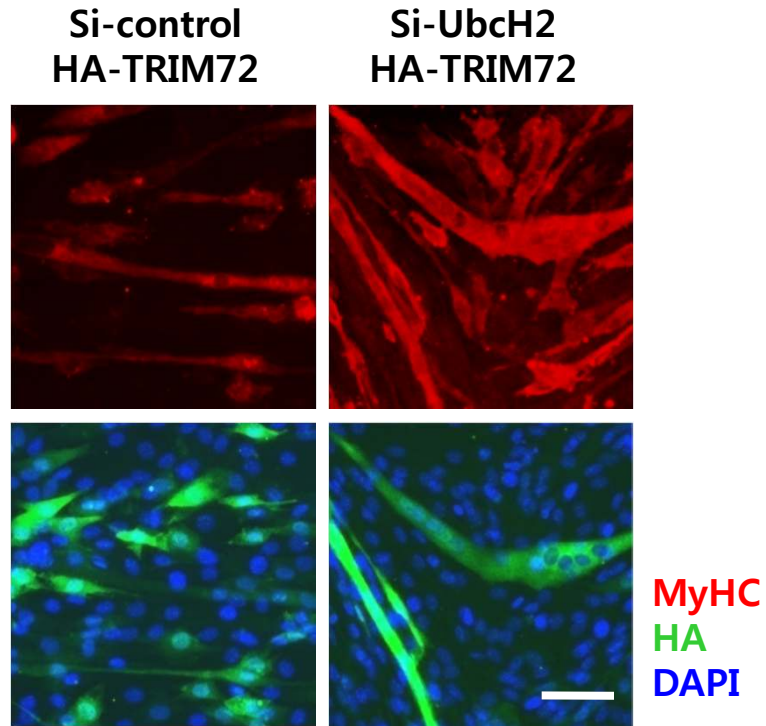
Endogenous IP
in **C2C12 myotubes**

UbcH2 knockdown enhances myogenesis by inhibiting IRS-1 ubiquitination.



C2C12 myotubes

TRIM72 inhibition of myogenesis is released by Ubch2 knockdown

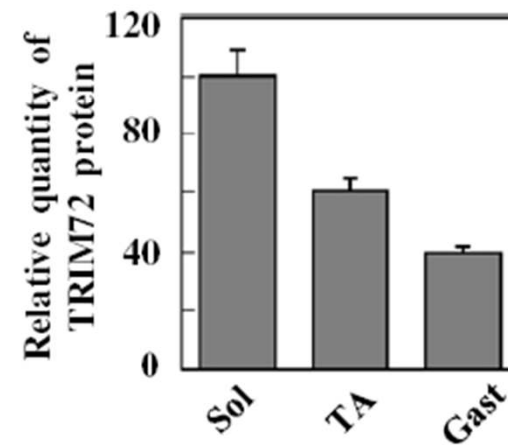
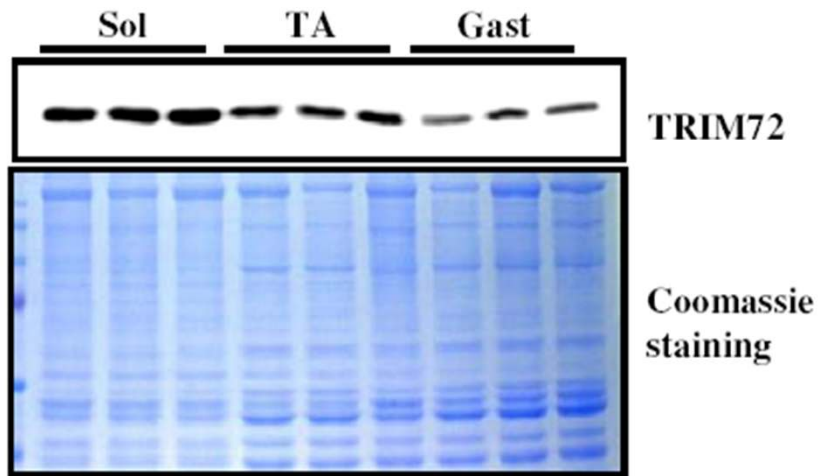


C2C12 myotubes

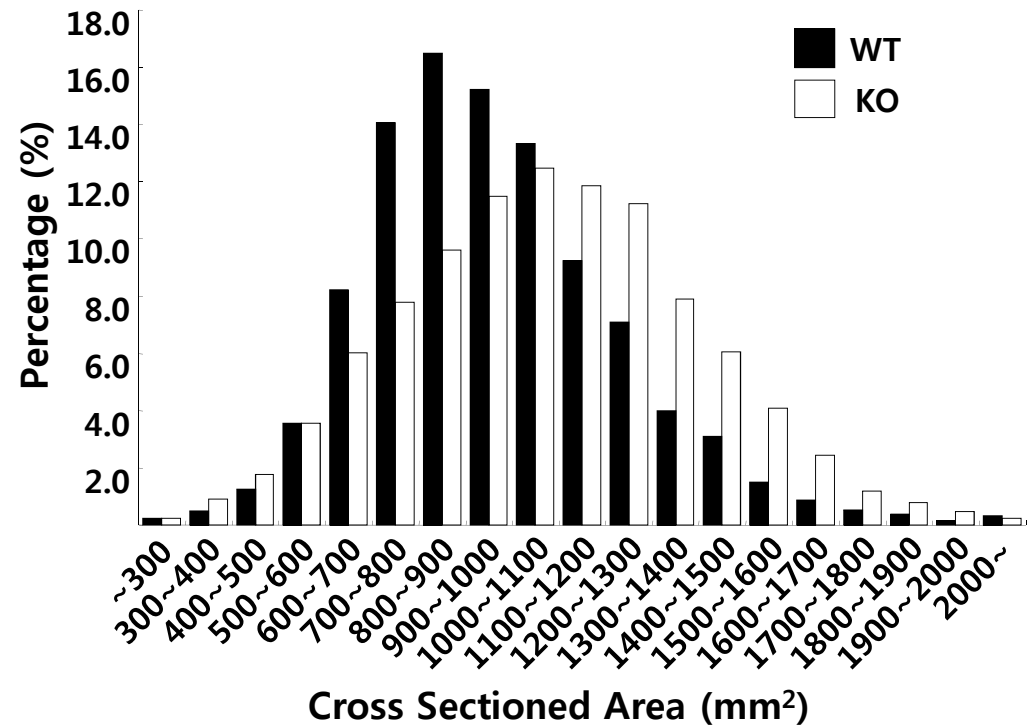
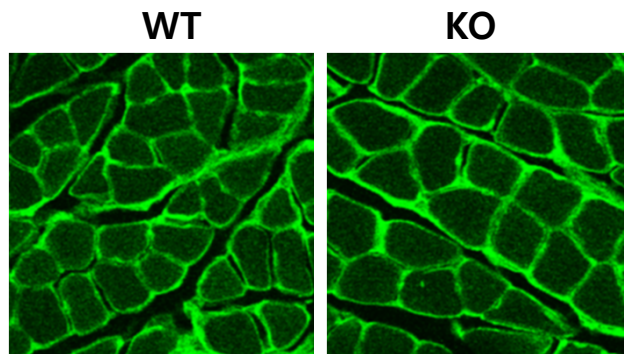
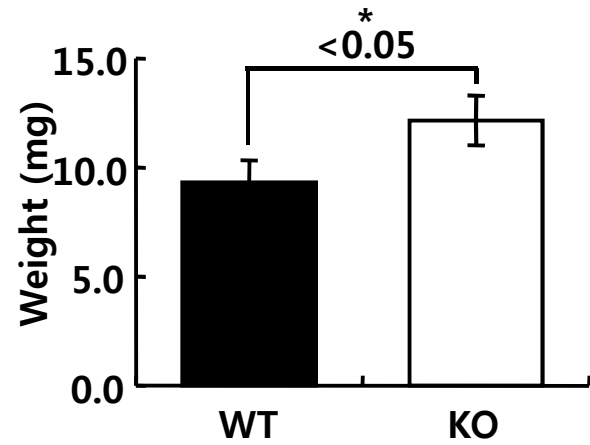
TRIM72 is highly expressed in soleus muscle.



	soleus	gastrocnemius
Color	Red muscle	White muscle
Size	Smaller	Larger
Fatigue	Resistant	Sensitive
TRIM72	High	Low

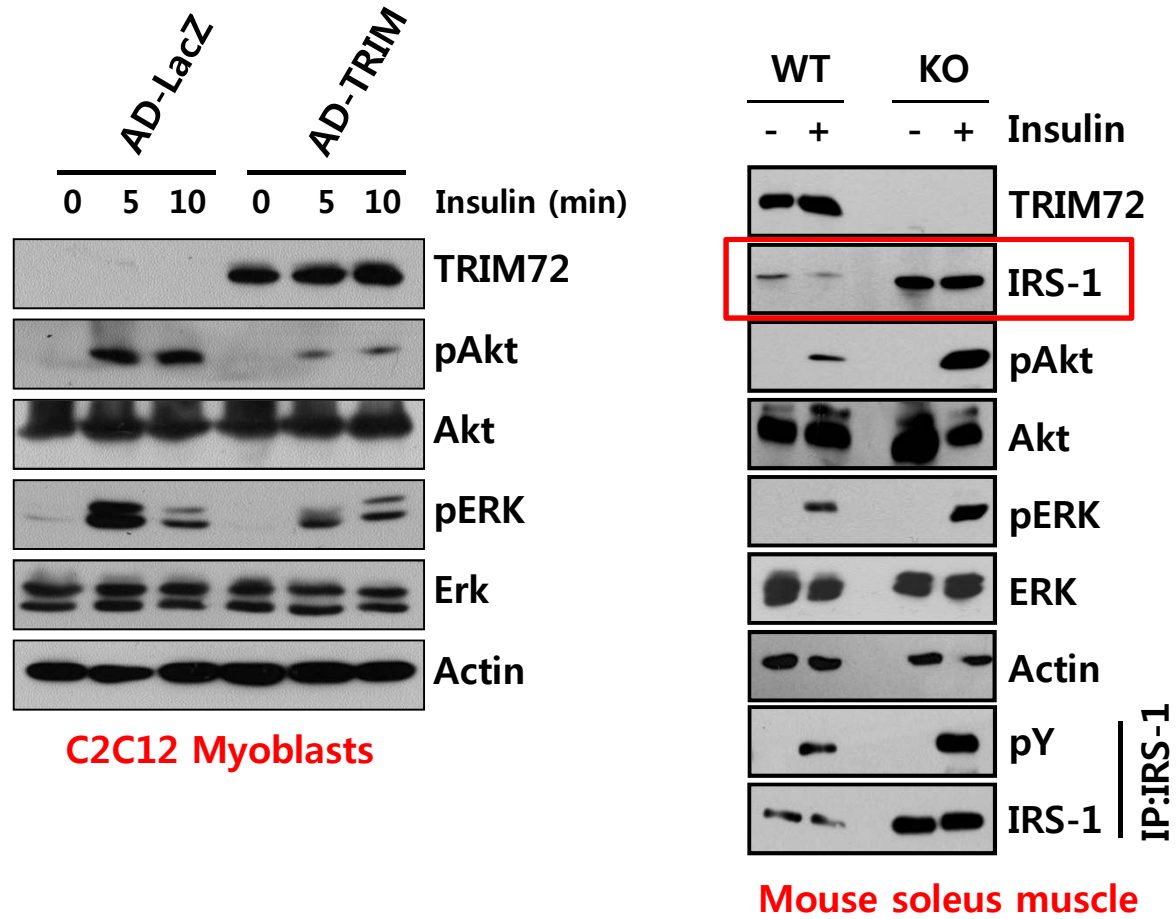


Skeletal muscle size is increased in TRIM72-disrupted mice.

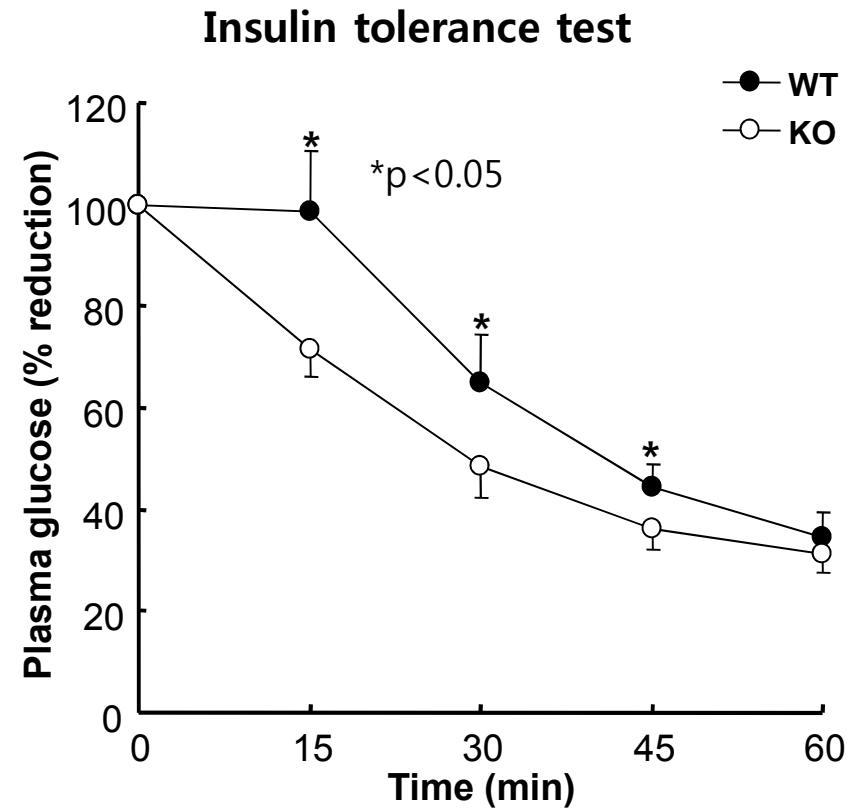
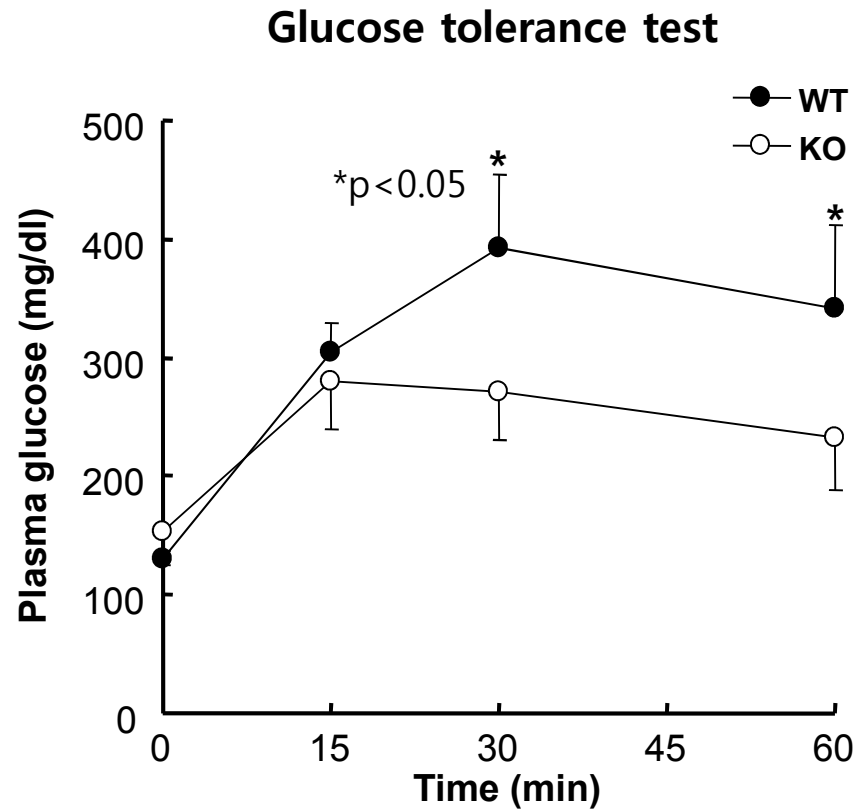


Mouse soleus muscle

IRS-1 expression level is elevated in TRIM-disrupted skeletal muscle

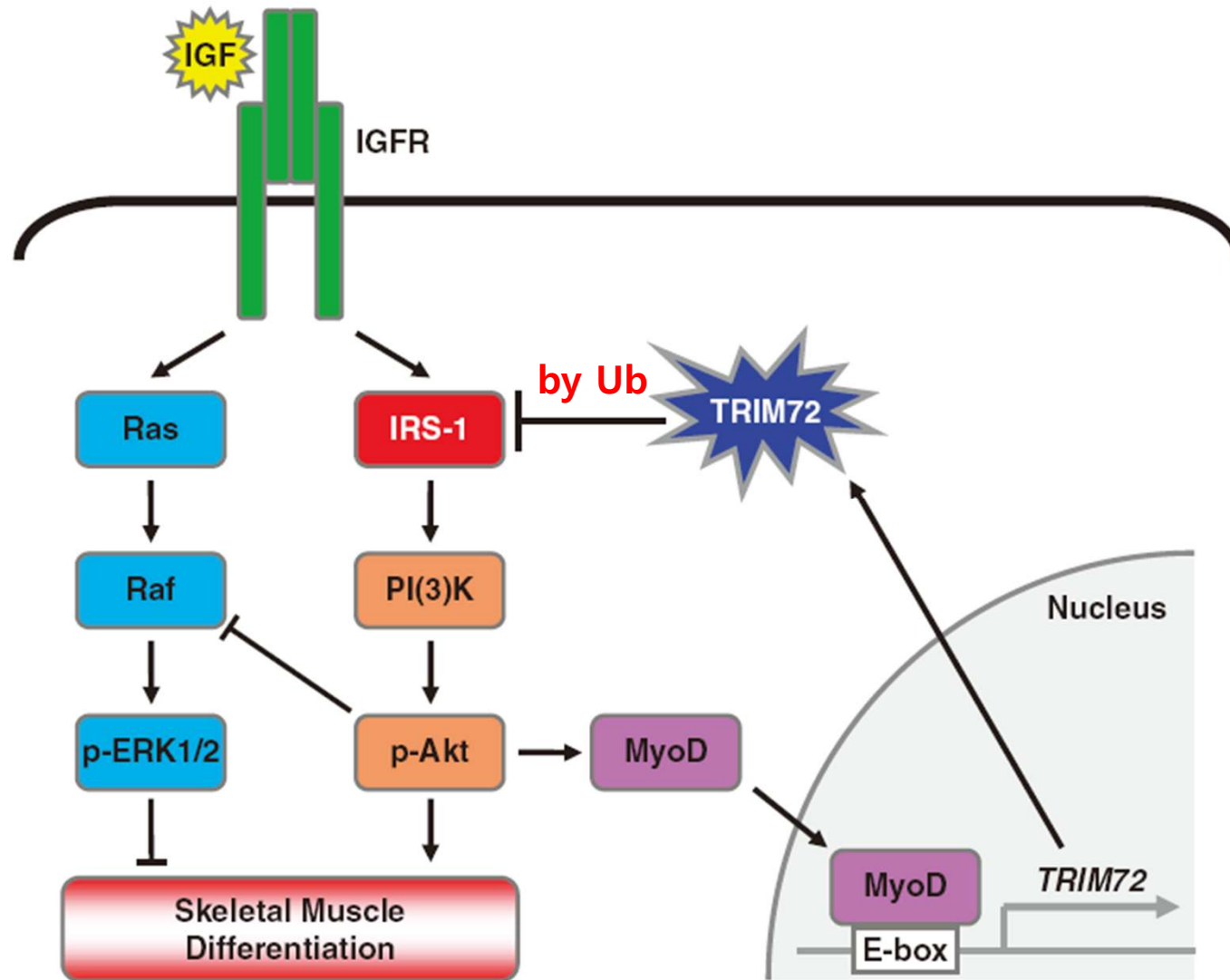


TRIM72 is a therapeutic target for type 2 diabetes



High fat diet-fed mice

E3 ligase TRIM72 negatively regulates myogenesis by IRS-1 ubiquitination.



Suggestion; TRIM72 is a negative muscle size regulator.



How do organs and whole organisms know when to stop growing?

A person's right and left legs almost always end up the same length, and the hearts of mice and elephants each fit the proper rib cage. **How genes set limits on cell size and number continues to mystify.**



JUPITER IMAGES

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Dr. Chang-Seok Lee

Dr. Young-Mi Ham

Miss Na-Rae Lee

Ms. Nga Nguyen

Mr. Jing Hong

Mr. June-Seop Lee

Mr. Jeong-Woo Lee

Mr. Dong-Min Yoo