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### A role for Raptor phosphorylation in the mechanical activation of TOR signaling

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#### ABSTRACT

The activation of mTOR signaling is necessary for mechanically-induced changes in skeletal muscle mass, but the 22 mechanisms that regulate the mechanical activation of mTOR signaling remain poorly defined. In this study, we 23 set out to determine if changes in the phosphorylation of Raptor contribute to the mechanical activation of mTOR. 24 To accomplish this goal, mouse skeletal muscles were subjected to mechanical stimulation via a bout of eccentric 25 contractions (EC). Using mass spectrometry and Western blot analysis, we found that ECs induced an increase in 26 Raptor S696, T706, and S863 phosphorylation, and this effect was not inhibited by rapamycin. This observation 27 suggested that changes in Raptor phosphorylation might be an upstream event in the pathway through which 28 mechanical stimuli activate mTOR. To test this, we employed a phospho-defective mutant of Raptor (S696A/ 29 T706A/S863A) and found that the EC-induced activation of mTOR signaling was significantly blunted in muscles 30 expressing this mutant. Furthermore, mutation of the three phosphorylation sites altered the interactions of Raptor with PRAS40 and p70<sup>S6k</sup>, and it also prevented the EC-induced dissociation of Raptor from p70<sup>S6k</sup>. Combined, 32 these results suggest that changes in the phosphorylation of Raptor play an important role in the pathway 33 through which mechanical stimuli activate mTOR signaling. 34

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#### 40 1. Introduction

Skeletal muscles make up the largest proportion of the body's mass,
and they are not only the motors that drive locomotion, but they also
play a crucial role in whole body metabolism [1,2]. Accordingly, it has
been well recognized that the maintenance of skeletal muscle mass contributes significantly to disease prevention and issues associated with
the quality of life [3].

Skeletal muscle mass is known to be regulated by a variety of stimuli,
one of which is mechanical loading. For instance, chronic mechanical
loading induces an increase in muscle mass, while chronic mechanical
unloading results in a loss of muscle mass [4]. Although it is well recognized that mechanical loads play a role in the regulation of muscle mass,
the molecular mechanisms that regulate these changes remain poorly
defined [5].

0898-6568/\$ – see front matter © 2013 Elsevier Inc. All rights reserved. http://dx.doi.org/10.1016/j.cellsig.2013.11.009 One molecule that has been widely implicated in the regulation 54 of skeletal muscle mass is the Serine/Threonine kinase called the 55 mammalian [or mechanistic] Target of Rapamycin (mTOR). For ex- 56 ample, previous studies have shown that mechanically-induced 57 muscle hypertrophy is associated with increases in mTOR signaling, 58 while atrophy resulting from mechanical unloading is associated with 59 decreased mTOR signaling [6,7]. Furthermore, recent studies have pro- 60 vided genetic evidence which indicates that the activation of mTOR is 61 sufficient to induce skeletal muscle hypertrophy, and that signaling 62 through mTOR is necessary for a mechanically-induced hypertrophic re- 63 sponse [8,9]. 64

mTOR exists in at least two distinct multi-protein complexes called 65 mTORC1 and mTORC2 and signaling by mTORC1, but not mTORC2, 66 can be inhibited by the drug rapamycin [10]. This is important because 67 rapamycin has been shown to inhibit mechanical load-induced in- 68 creases in mTOR-dependent signaling events, protein synthesis, and 69 growth [9,11,12]. Thus, it has been widely concluded that signaling by 70 mTORC1 is critical for mechanically-induced changes in muscle mass. 71

The mTORC1 complex contains mTOR and other accessory proteins 72 such as Raptor and PRAS40 [13–16]. Previous studies have shown that 73 Raptor, which binds directly to mTOR, acts as a scaffold protein and is 74 important in the regulation of mTOR-dependent signaling events. For 75 example, knockdown of Raptor using RNAi results in a reduction of 76 cell size *in-vitro*. Furthermore, it has been demonstrated that skeletal 77 muscle specific ablation of Raptor *in-vivo* results in the loss of muscle 78 mass, and it also prevents mechanical load-induced hypertrophy 79

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*Abbreviations:* EC, eccentric contraction; GβL, G-protein β-subunit-like protein; GFP, green fluorescent protein; JNK, c-jun n-terminal kinase; MAPK, mitogen-activated protein kinase; mLST8, lethal with sec13 protein 8; mTOR, mammalian [or mechanistic] target of rapamycin; mTORC1, mTOR complex 1; mTORC2, mTOR complex 2; p38, p38 mitogen-activated protein kinase; P70<sup>56k</sup>, ribosomal S6 kinase 1; PI3K, phosphotidylinositol-3-kinase; PRAS40, proline-rich Akt substrate of 40 kDa; Raptor, regulatory associated protein of mTOR; TA, tibialis anterior muscle.

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## **ARTICLE IN PRESS**

J.W. Frey et al. / Cellular Signalling xxx (2013) xxx-xxx

[17–19]. Hence, it seems apparent that Raptor plays an important role in
 the regulation of mTOR signaling and muscle mass, but a complete un derstanding of the mechanisms through which Raptor regulates these
 processes is only beginning to be established.

Changes in the phosphorylation state of Raptor appear to be one 84 mechanism through which Raptor can regulate mTOR signaling. For ex-85 ample, alterations in Raptor phosphorylation have been implicated in 86 87 the regulation of mTOR signaling that occurs in response to insulin, os-88 motic stress and energetic stress [20-23]. It has also been shown that 89 signaling through the Ras/MAPK pathway can induce changes in 90 mTOR signaling by promoting alterations in the phosphorylation of Raptor [24,25]. Combined, these studies have provided strong evidence 91to support the conclusion that mTOR signaling can be regulated through 92alterations in the phosphorylation state of Raptor. However, to date, the 93 potential role of Raptor phosphorylation in the regulation of mTOR by 94 mechanical stimuli has not been explored. Thus, the goal of this study 95 was to determine if changes in the phosphorylation of Raptor contribute 96 97 to the mechanical activation of mTOR signaling.

#### 98 2. Experimental procedures

#### 99 2.1. Materials

Primary antibodies, anti-total p70<sup>S6k</sup> (#2708), phospho Serine/ 100 Threonine – Proline (S\*/T\*–P) (#9391), phospho Akt substrate 101 (RxRxxS\*/T\*) (#10001), total Raptor (#2280), phospho Raptor (\$792) 102 (#2083), total mTOR (#2972) and PRAS40 (#2691) were purchased 103 104 from Cell Signaling Technology (Danvers, MA, USA). The primary antibody anti-phospho p70<sup>S6k</sup>(T389) (#sc-11759-R) was purchased from 105 Santa Cruz Biotechnology (Santa Cruz, CA, USA). Phospho-specific Rap-106 tor antibodies (S696, T706, S859, S863, and S877) have been previously 107 108 described [20] and were kindly provided by Dr. Diane Fingar (University of Michigan Medical School, Ann Arbor, Michigan, USA). The peroxidase 109 labeled anti-rabbit IgG (H + L) secondary antibody (#PI-1000) was 110 purchased from Vector Laboratories Inc. (Burlingame, CA, USA). Perox-111 112 idase labeled anti-rabbit IgG (Light Chain Specific) (#211-032-171) and anti-mouse IgG (Light Chain Specific) (#115-035-174) antibodies were 113114 purchased from Jackson ImmnunoResearch Laboratories Inc. (West Grove, PA, USA). 115

#### 116 2.2. Animals

C57BL6 mice (Jackson Laboratories, Bar Harbor, MA, USA) 8-10 117 weeks of age were randomly assigned to different experimental 118 groups. Before all surgical procedures, the mice were anaesthetized 119 120 with an intraperitoneal injection of ketamine (100 mg/kg) and xylazine 121 (10 mg/kg). After tissue extraction, the mice were sacrificed by cervical 122dislocation. All animals were housed in a room maintained with a 12-12 h light-dark cycle and received food and water ad libitum. The 123Institutional Animal Care and Use Committee at the University of 124 Wisconsin-Madison approved all of the methods employed in this 125study. 126

#### 127 2.3. Plasmid constructs and purification

The GFP and GST-p70<sup>S6k</sup> plasmid constructs have been previously 128described [8]. Wild type FLAG-Raptor and FLAG-Raptor (S696A, 129 T706A, S863A) have been previously described [21] and were kindly 130 provided by Dr. Sung Ho Ryu (Pohang University of Science and 131 Technology, Pohang, Kyungbuk, Korea). Myc-p70<sup>S6k</sup> was generated 132by using the QuickChange II Site-Directed Mutagenesis Kit (Agilent 133 Technologies, Santa Clara, CA, USA) to insert a stop codon before 134the region that encodes for the GST tag on the Myc-p70<sup>S6k</sup>-GST plas-135mid that has been previously described [8]. All plasmid DNA was 136 137 grown in DH5 $\alpha$  E. Coli, purified with an Endofree plasmid kit (Qiagen, Valencia, CA) and resuspended in sterile phosphate buff- 138 ered saline (PBS). 139

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#### 2.4. Skeletal muscle transfection (electroporation)

As previously described [8], a small incision was made through the 141 skin covering the TA muscle and then a 27-gauge needle was used to in- 142 ject 12 µl of plasmid DNA solution containing 7.5–25 µg of GFP, wild 143 type FLAG-Raptor (WT) or mutant FLAG-Raptor (S696A, T706A, 144 S863A) (3A) and, when indicated, 2 µg of GST- or Myc-tagged p70<sup>S6k</sup>. 145 After the injections, electric pulses were applied through two stainless 146 steel pin electrodes (1-cm gap; Harvard Apparatus, Holliston, MA, 147 USA) laid on top of the proximal and distal myotendinous junctions. 148 Eight 20 ms square-wave electric pulses at a frequency of 1 Hz were de- 149 livered with an ECM 830 electroporation unit (BTX; Harvard Apparatus) 150 at a field strength of 160 V/cm. After the electroporation procedure, the 151 incision was closed with Vetbond surgical glue (3 M Animal Care Prod- 152 ucts, St. Paul, MN, USA). Following the surgery, mice were given an 153 intraperitoneal injection of buprenorphine (0.05  $\mu$ g/g) dissolved in 154 100 µl of PBS. 155

#### 2.5. Eccentric contractions and insulin injections

The model previously described by O'Neil et al. (2009) was used to 157 induce eccentric contractions (EC) in the TA muscle [26]. Specifically, 158 electrodes were placed on the sciatic nerve of the right leg and contractions were elicited by stimulating the sciatic nerve with an SD9E Grass 160 stimulator (Grass Instruments, Quincy, MA, USA) at 100 Hz, 4–8 V 161 pulse, for 10 sets of 6 contractions. Each contraction lasted 3 s and 162 was followed by a 10 s rest period, and a 1 min rest period was provided 163 between each set. The left (control) TA muscle and the right (EC) TA 164 muscle were collected 40 min after the last set of contractions. When 165 indicated, mice were also given an intraperitoneal injection of 20 U/kg 166 Novolog® insulin (Novo Nordisk, Princeton, NJ, USA) dissolved in 167 100 µl of PBS at 30 min prior to the collection. 168

#### 2.6. Rapamycin injections

Rapamycin was purchased from LC laboratories (Woburn, MA, USA) 170 and was dissolved in DMSO to generate a 5  $\mu$ g/µl stock solution. 171 Rapamycin was administered at a dose of 2.0 mg/kg and dissolved in 172 a final volume of 200 µl PBS. For the vehicle condition, an equal amount 173 of DMSO was dissolved in a final volume of 200 µl PBS. The vehicle or 174 rapamycin solutions were administered via an intraperitoneal injection 175 100 min prior to the initiation of the EC protocol. 176

#### 2.7. Sample preparation for immunoprecipitations and Western blot 177 analysis 178

Upon collection, TA muscles were either immediately homogenized 179 or frozen in liquid nitrogen. In some cases, the samples were 180 homogenized with a Polytron for 20 s in ice-cold buffer A (40 mM 181 Tris (pH 7.5), 1 mM EDTA, 5 mM EGTA, 0.5% Triton X-100, 25 mM 182 β-glycerophosphate, 25 mM NaF, 1 mM Na3VO4, 10 mg/ml leupeptin, 183 and 1 mM PMSF) and the whole homogenate was used for further anal- 184 ysis. In other cases (e.g. preservation of the mTORC1 complex) the sam- 185 ples were homogenized with a Polytron for 20 s in ice-cold CHAPS lysis 186 buffer (40 mM HEPES (pH. 7.4), 2 mM EDTA, 0.3% CHAPS, 10 mM sodi- 187 um pyrophosphate, 10 mM  $\beta$ -glycerophosphate, and 1 tablet of EDTA- 188 free protease inhibitors (Roche) per 25 ml) and the whole homogenate 189 was used for further analysis. Sample protein concentration was deter- 190 mined with a DC protein assay kit (Bio-Rad, Hercules, CA, USA), and un- 191 less otherwise noted (e.g. immunoprecipitation), equivalent amounts 192 of protein from each sample were dissolved in 2× Laemmli buffer, heat- 193 ed to 100 °C for 5 min and then subjected to Western blot analysis as 194 described below. 195

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