

## **Effects of rapid or slow body weight reduction on intramuscular protein degradation pathways during equivalent weight loss on rats**

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Running head: Rapid and slow weight loss and muscle atrophy

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1 **Summary**

2 The purpose of this study was to compare the effects of short-term fasting-induced rapid  
3 weight loss with those of slower but equivalent body weight loss induced by daily calorie  
4 restriction on muscle protein degradation pathways and muscle protein content. Male  
5 Fischer rats were subjected to either 30% calorie restriction for 2 wk to slowly decrease  
6 body weight (Slow) or 3-day fasting to rapidly decrease body weight by a comparable level  
7 of that of the Slow group (Rapid). The final body weights were about 15% lower in both the  
8 Slow and Rapid groups than in the Con group ( $p < 0.001$ ). The total protein content and wet  
9 weight of fast-twitch plantaris muscle, but not slow-twitch soleus muscle, were significantly  
10 lower in the Rapid group compared with the control rats fed ad libitum. Substantial  
11 increases in the expression ratio of autophagosomal membrane proteins (LC3-II/-I ratio)  
12 and polyubiquitinated protein concentration, used as biomarkers of autophagy-lysosome  
13 and ubiquitin-proteasome activities, respectively, were observed in the plantaris muscle of  
14 the Rapid group. Moreover, the LC3-II/-I ratio and polyubiquitinated protein concentration  
15 were negatively correlated with the total protein content and wet weight of plantaris muscle.  
16 These results suggest that short-term fasting-induced rapid body weight loss activates  
17 autophagy-lysosome and ubiquitin-proteasome systems more strongly than calorie  
18 restriction-induced slower weight reduction, resulting in muscular atrophy in fast-twitch  
19 muscle.

20

21 **Key words:** skeletal muscle, fasting, calorie restriction, autophagy-lysosome,  
22 ubiquitin-proteasome

23

24

## 25 **Introduction**

26 Many athletes restrict their caloric intake to improve their force-to-mass ratio, to  
27 achieve a certain body mass category, or for aesthetic reasons. In particular, athletes in  
28 weight-classified sports such as wrestling and boxing usually lose body weight rapidly  
29 before competitions (Choma *et al.* 1998, Reljic *et al.* 2013). The rapid weight loss, also  
30 known as “weight cutting”, typically involves several-day fasting until the targeted weight is  
31 met. However, fasting is a recognized stimulus of skeletal muscle atrophy (Jagoe *et al.*  
32 2002), which results in a significant loss of lean body mass that compromises exercise  
33 performance. Muscle atrophy occurs when rate of protein degradation exceeds that of  
34 protein synthesis. There are two major protein degradation pathways in skeletal muscle.  
35 One, the ubiquitin-proteasome pathway, plays a major role in selective protein degradation  
36 and serves as the primary degradation route for most short-lived proteins (Rock *et al.*  
37 1994). The other, the autophagy-lysosome pathway, is an intracellular bulk degradation  
38 system that is responsible for the degradation of most long-lived proteins, as well as some  
39 organelles (Mortimore and Pösö 1987). Both proteolytic pathways become activated during  
40 fasting to maintain amino acid pools, leading to muscle atrophy (Mitch and Goldberg 1996,  
41 Bujak *et al.* 2015).

42 An alternative dietary weight-loss approach practiced by athletes is daily calorie  
43 restriction, which results in slower body weight loss compared with fasting. Many Japanese  
44 bodybuilders empirically believe that the slower body weight loss induced by daily calorie  
45 restriction has less atrophic effects on skeletal muscle than the fasting-induced rapid  
46 weight loss and therefore adopt the slower body weight-loss strategy before competitions.  
47 However, it remains unclear which type of body weight loss more strongly activates the  
48 autophagy-lysosome and ubiquitin-proteasome pathways and induces muscle atrophy

49 when body weight is reduced to the same extent, because no study has directly compared  
50 the effects of rapid vs. slow body weight reduction on the major protein degradation  
51 pathways and on protein content in skeletal muscle. Thus, the purpose of this study was to  
52 directly compare the effects of rapid or slow body weight loss on the autophagy-lysosome  
53 and ubiquitin-proteasome pathways and on protein content in rat skeletal muscle during an  
54 equivalent weight loss.

55

## 56 **Methods**

### 57 **Animal treatment**

58 Nineteen-week-old male Fischer-344 rats were obtained from Japan SLC  
59 (Shizuoka, Japan) and individually housed under a 12:12-h light:dark cycle (light  
60 09.00-21.00 h) in an air-conditioned room (23°C). Rats were given a standard laboratory  
61 diet ad libitum (CE-2; CLEA Japan, Tokyo, Japan) and water and acclimated to the housing  
62 facility for 1 wk.

63 After the acclimation period, the rats were divided into three groups, matched for  
64 body weight: one group continued to receive the standard diet ad libitum for the entire  
65 14-day experimental period (Con; n = 5); a second group received the standard diet equal  
66 to 70% of the average amount of food eaten by the Con group during the 14 days to  
67 decrease their body weight slowly (Slow; n = 5); the third group was fed the standard diet  
68 ad libitum for 11 days and fasted thereafter for the last 3 days of the study period to rapidly  
69 decrease their body weight to a comparable extent as that of the Slow group (Rapid; n = 5).  
70 All rats were allowed to drink water freely during the 14-day dietary intervention. Body  
71 weight and food intake were recorded daily during the dietary intervention.

72 At the end of the dietary intervention, fast-twitch plantaris, extensor digitorum

73 longus (EDL), and slow-twitch soleus muscles were quickly and carefully dissected out  
74 under anesthesia with isoflurane immediately after the 12-h dark period during which rats  
75 eat most food. The muscle samples were weighed, quickly frozen in liquid N<sub>2</sub>, and stored at  
76 –80°C until analysis. After the blood samples were collected from the heart,  
77 intra-abdominal fat (sum of the epididymal, mesenteric, and retroperitoneal fat pads) was  
78 removed and weighed. The experimental protocols were approved by the Animal  
79 Experimental Committee of The University of Tokyo.

80

### 81 **Muscle homogenization**

82 Frozen plantaris and soleus muscles were homogenized in ice-cold  
83 Radio-Immuno Precipitation Assay (RIPA) lysis buffer (EMD Millipore, Temecula, CA, USA)  
84 containing 50 mM Tris-HCl, pH 7.4, 150 mM NaCl, 0.25% deoxycholic acid, 1% NP-40, 1  
85 mM ethylenediaminetetraacetic acid (EDTA), protease inhibitor cocktail (SIGMA-Aldrich, St.  
86 Louis, MO, USA), and phosphatase inhibitors (PhosSTOP; Roche, Basel, Switzerland).  
87 The homogenates were frozen and thawed three times to disrupt intracellular organelles  
88 and rotated end-over-end at 4°C for 60 min to solubilize the protein. Total protein content  
89 per muscle was measured with a bicinchoninic acid (BCA) protein assay kit (Pierce,  
90 Rockford, IL, USA). Homogenized samples were then centrifuged at 700 × g for 5 min at  
91 4°C and the supernatants were harvested.

92

### 93 **Western blotting**

94 Protein concentrations of the supernatant harvested as described above were  
95 measured with the BCA protein assay kit. Samples were prepared in Laemmli sample  
96 buffer (Wako Pure Chemical, Osaka, Japan) and heated for 5 min in a heating block at

97 95°C. Equal amounts of sample protein were subjected to sodium dodecyl  
98 sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) (7.5% or 15% resolving gels) and  
99 then transferred to polyvinylidene difluoride (PVDF) membranes at 200 mA for 90 min. After  
100 transfer, membranes were blocked for 1 h at room temperature in Tris-buffered saline  
101 (TBS) with 0.1% Tween 20 (TBS-T; 20 mM Tris base, 137 mM NaCl, pH 7.6) supplemented  
102 with 5% (w/v) nonfat powdered milk or 5% (w/v) bovine serum albumin. Membranes were  
103 incubated overnight at 4°C with the primary antibody diluted 1:1000 in TBS-T containing  
104 5% bovine serum albumin. The primary antibodies used were anti-microtubule-associated  
105 protein light chain 3 (LC3) (Medical & Biological Laboratories, Nagoya, Japan),  
106 anti-phospho-p70S6K (Cell Signaling Technology, Danvers, MA, USA), and  
107 anti-phospho-Akt (Ser473) (Cell Signaling Technology). After the incubation with primary  
108 antibody, membranes were incubated for 1 h at room temperature with secondary  
109 antibodies (goat anti-rabbit IgG or goat anti-mouse IgG, Jackson ImmunoResearch  
110 Laboratories, West Grove, PA, USA) diluted 1:5000 in TBS-T containing 1% nonfat  
111 powdered milk. Bands were visualized by enhanced chemiluminescence (ECL) reagent  
112 (GE Healthcare Life Sciences, Piscataway, NJ, USA) and quantified by Image Studio  
113 (LI-COR, Lincoln, NE, USA). The membranes were stained with Ponceau (Sigma-Aldrich)  
114 to verify equal loading of protein across lanes.

115

#### 116 **Polyubiquitinated protein concentration analysis**

117 The supernatants of the plantaris and soleus muscle homogenates were also  
118 used for the measurement of polyubiquitinated protein concentrations. Polyubiquitinated  
119 protein concentrations were measured with an enzyme-linked immunospecific assay  
120 (ELISA) kit according to the manufacturer's instructions (Cyclex Poly-Ubiquitinated Protein

121 ELISA Kit; Medical & Biological Laboratories).

122

### 123 **Serum glucose and insulin concentrations**

124 Serum glucose and insulin concentrations were determined with the Glucose C2

125 Test Wako kit (Wako Pure Chemical) and Rat Insulin ELISA Kit (Merckodia AB, Uppsala,

126 Sweden), respectively.

127

### 128 **Muscle glycogen concentration**

129 For the measurement of the muscle glycogen concentration, EDL muscles were

130 homogenized with 0.3 M perchloric acid. The glycogen concentration was determined by

131 the enzymatic methods of Lowry and Passonneau after acid hydrolysis (Lowry and

132 Passonneau 1972).

133

### 134 **Statistical analysis**

135 All data are presented as means  $\pm$  SEM. Statistical analysis was performed by

136 Welch's ANOVA and Bonferroni correction for post-hoc analysis (Social Survey Research

137 Information Co., Ltd., Tokyo, Japan). We performed least-squares regression analyses to

138 examine relationships between variables. Statistical significance was defined as  $p < 0.05$ .

139

## 140 **Results**

### 141 **Body weight, total intra-abdominal fat weight, and total food intake**

142 Changes in body weights during the 2-wk dietary intervention are shown in Fig.1.

143 During the intervention period, daily calorie restriction in the Slow group for 2 wk and 3-day

144 fasting in the Rapid group caused a substantial reduction in body weight. The body weight

145 in the Slow group became significantly different from the Con and Rapid groups at day 3  
146 ( $p < 0.05$ ). In addition, significant body weight reduction in the Slow group from day 0 was  
147 observed at day 3. The body weight in the Rapid group became significantly different from  
148 the Con group at day 12 (1 day after the onset of fasting). The final body weights were  
149 about 15% lower in both the Slow and Rapid groups than in the Con group ( $p < 0.001$ )  
150 (Table 1). Total intra-abdominal fat weights were also significantly lower in both the Slow  
151 and Rapid groups than in the Con group ( $p < 0.001$ ), with no significant differences between  
152 the Slow and Rapid groups (Table 1).

153 Total food intake during the 2-wk experimental period was significantly lower in the  
154 Slow and Rapid groups than in the Con group ( $p < 0.001$ ; Table 1). Furthermore, total food  
155 intake was significantly lower in the Slow group than in the Rapid group ( $p < 0.05$ ).

156

### 157 **Serum glucose and muscle glycogen concentration**

158 At the completion of the 14-day dietary intervention, there was no significant  
159 difference in serum glucose concentration among the three groups (Table 1). Although  
160 there was no significant difference in the glycogen concentration of EDL muscle between  
161 the Con and Slow groups, the muscle glycogen concentration was significantly lower in the  
162 Rapid group than in the Con and Slow groups ( $p < 0.001$ ; Table 1).

163

### 164 **Muscle wet weight and muscle total protein content**

165 There were no significant differences in muscle wet weight and total protein  
166 content of the soleus muscle among the three groups (Table 1). Although the wet weight  
167 and total protein content of the plantaris muscle did not differ between the Con and Slow  
168 groups, the muscle weight and total protein content in the plantaris muscle were

169 significantly lower in the Rapid group than in the Con group ( $p < 0.05$ ; Table 1).

170

### 171 **Autophagy-lysosome activity**

172 The microtubule-associated protein LC3 is now widely used to monitor the  
173 autophagy-lysosome system. The cytosolic form of LC3 (LC3-I) conjugates with  
174 phosphatidylethanolamine to form the LC3-phosphatidylethanolamine conjugate (LC3-II),  
175 which is recruited to autophagosomal membranes (Mizushima and Yoshimori 2007).  
176 Because the amount of LC3-II is correlated with the extent of autophagosome formation  
177 and an increased LC3-II/LC3-I ratio is representative of accelerated autophagy-lysosome  
178 activity (Lee *et al.* 2014), we determined the expression levels of LC3-I and LC3-II and  
179 used the LC3-II/LC3-I ratio as a marker of autophagy-lysosome activity.

180 In both plantaris and soleus muscles, LC3-II/LC3-I ratios were significantly higher  
181 in the Slow group than in the Con group ( $p < 0.01$ ; Fig. 2-A and -B). Further increases in  
182 LC3-II/LC3-I ratios were observed in the plantaris and soleus muscles of the Rapid group  
183 ( $p < 0.001$  vs. the Con and Slow groups; Fig. 2-A and -B). In the plantaris muscle but not the  
184 soleus muscle, the LC3-II/LC3-I ratio was significantly and negatively associated with the  
185 muscle wet weight ( $p < 0.01$ ) and muscle protein content ( $p < 0.05$ ; Fig. 3-A and -C).

186

### 187 **Polyubiquitinated protein concentration**

188 Intracellular proteins are marked with a polyubiquitin chain, after which they are  
189 degraded to peptides and free ubiquitin by the 26S proteasome (Goldberg 2003). In the  
190 present study, we used the polyubiquitinated protein concentration as a marker of  
191 ubiquitin-proteasome pathway activity. The polyubiquitinated protein concentrations of the  
192 soleus muscle did not significantly differ among the three groups (Fig. 2-D). In contrast, in

193 the plantaris muscle, they were significantly higher in the Slow and Rapid groups than in  
194 the Con group (Con vs. Slow:  $p < 0.01$ ; Con vs. Rapid:  $p < 0.001$ ; Fig.2-C). Moreover, the  
195 polyubiquitinated protein concentration was higher in the Rapid group than in the Slow  
196 group ( $p < 0.05$ ; Fig. 2-C). The polyubiquitinated protein concentrations were significantly  
197 and negatively associated with the muscle wet weight ( $p < 0.01$ ) and muscle protein content  
198 ( $p < 0.05$ ) of the plantaris muscle (Fig. 3-B and -D).

199

### 200 **Protein synthesis pathway**

201 Although mechanistic target of rapamycin (mTOR) is a master regulator of muscle  
202 protein synthesis (Wulfschleger *et al.* 2006), the phosphorylation status of mTOR  
203 (phospho-mTOR) does not necessarily reflect mTOR activity (Eliasson *et al.* 2006, Fujita *et*  
204 *al.* 2007, Miyazaki *et al.* 2011). Many recent studies have instead evaluated the  
205 phosphorylation of p70S6K (phospho-p70S6K), a downstream target of mTORC1, as a  
206 biomarker of mTOR activity (Jacinto and Hall 2003, Tamura *et al.* 2014). Although both the  
207 Slow and Rapid groups tended to have lower phospho-p70S6K content in the plantaris and  
208 soleus muscles than the Con group, the difference was not statistically significant due to a  
209 considerable variability in phospho-p70S6K levels (Fig. 2-E and -F).

210

### 211 **Serum insulin concentration and phosphorylated-Akt content in skeletal muscle**

212 The insulin–Akt axis has strong inhibitory effects on both autophagy-lysosome  
213 and ubiquitin-proteasome pathways in skeletal muscle (Price *et al.* 1996, Mitch *et al.* 1999,  
214 Lee *et al.* 2004, Satchek *et al.* 2004, Stitt *et al.* 2004, Wang *et al.* 2006). Here, the serum  
215 insulin concentration was significantly lower in both the Slow and Rapid groups than in the  
216 Con group ( $p < 0.001$ ; Table 1). In addition, the serum insulin concentration was significantly

217 lower in the Rapid group than in the Slow group ( $p < 0.01$ ; Table 1). The levels of  
218 phospho-Akt, which is the active form of Akt, in the plantaris and soleus muscles were  
219 significantly lower in the Rapid group than in the Con and Slow groups, with no significant  
220 differences between the Con and Slow groups (Con vs. Rapid:  $p < 0.01$ ; Slow vs. Rapid in  
221 plantaris muscle:  $p < 0.01$ ; Slow vs. Rapid in soleus muscle:  $p < 0.05$ ; Fig. 2-G and -H). The  
222 serum insulin concentration significantly and negatively correlated with the LC3-II/LC3-I  
223 ratio and polyubiquitinated protein concentration in the plantaris muscle ( $p < 0.01$ ; Fig. 4-A  
224 and -B). In addition, the phospho-Akt content was significantly and negatively correlated  
225 with the LC3-II/LC3-I ratio and polyubiquitinated protein concentration in the plantaris  
226 muscle ( $p < 0.05$ ; Fig. 4-C and -D).

227

## 228 **Discussion**

229 A severe energy deficit during body weight loss causes significant reductions in  
230 skeletal muscle and body fat masses. To our knowledge, this is the first study to directly  
231 compare the effects of rapid and slow weight reductions, which result in acute and gradual  
232 energy deficits, respectively, on protein degradation pathways and protein content in  
233 skeletal muscle during an equivalent weight loss in rats. We found that the rapid weight  
234 loss induced by the 3-day fast potently activated both autophagy-lysosome and  
235 ubiquitin-proteasome pathways (Fig. 2). This fast resulted in significant reductions in the  
236 total protein content and wet weight of the fast-twitch plantaris muscle (Table 1), although  
237 both weight-loss methods decreased rat body weight and total intra-abdominal fat mass to  
238 a similar extent (Table 1 and Fig. 1).

239 Muscle atrophy occurs when protein degradation rates exceed protein synthesis  
240 rates. Although the phospho-p70S6K contents of the plantaris and soleus muscles

241 appeared to be lower in both the Slow and Rapid groups than in the Con group, the  
242 difference was not statistically significant (Fig. 2-E and -F). In addition, the  
243 phospho-p70S6K contents of the plantaris muscle were almost identical in both weight-loss  
244 groups. It is therefore unlikely that fasting-induced atrophy in the plantaris muscle of the  
245 Rapid group was due to a diminished protein synthesis rate, although we did not directly  
246 evaluate the muscle protein synthesis rate.

247           Even though the total food intake during the 14-day dietary intervention was  
248 significantly higher in the Rapid group than in the Slow group, the muscle glycogen  
249 concentration was substantially lower in the Rapid group, suggesting that only the 3-day  
250 fast resulted in a severe energy deficit in muscle cells. A severe energy deficit induced by  
251 fasting and starvation activates autophagy-mediated protein degradation (Mizushima *et al.*  
252 2004). In this study, we found that the LC3-II/LC3-I ratio, frequently used as a biomarker of  
253 autophagy-lysosome activity (Lee *et al.* 2014), was significantly higher in the plantaris  
254 muscle of the Rapid group than in that of both the Slow and Con groups (Fig. 2). This  
255 finding indicates that the autophagy-lysosome pathway was more potently activated in the  
256 Rapid group than in the Slow group, despite the equivalent weight loss. In addition, the  
257 LC3-II/LC3-I ratio was significantly and negatively correlated with the total protein content  
258 and muscle wet weight in the plantaris muscle (Fig. 3-A and -C). Based on these results, it  
259 is plausible that the higher autophagy-lysosome activity was responsible for the rapid  
260 weight loss-induced atrophy in the plantaris muscle.

261           Our results also indicate that another protein degradation pathway, the  
262 ubiquitin-proteasome system, might be involved in the muscular atrophy in the Rapid group.  
263 In the ubiquitin-proteasome system, proteins are targeted for degradation by the 26S  
264 proteasome through covalent attachment of a chain of ubiquitin molecules (Goldberg 2003).

265 We determined the polyubiquitinated protein concentration in skeletal muscle and found  
266 that the plantaris muscle in the Rapid group had a significantly higher polyubiquitinated  
267 protein concentration (Fig.2-C), as well as LC3-II/LC3-I ratio, than the Con and Slow  
268 groups. Similar to the LC3-II/LC3-I ratio, the polyubiquitinated protein concentration was  
269 significantly and negatively associated with total protein content and muscle wet weight in  
270 the plantaris muscle (Fig. 3-B and -D). These results provide strong evidence that the two  
271 major proteolytic pathways are more strongly activated during rapid body weight loss than  
272 during slow body weight reduction, resulting in a significant loss of total protein content and  
273 wet weight of fast-twitch muscle.

274           Insulin and its downstream effector Akt strongly inhibit both autophagy-lysosome  
275 and ubiquitin-proteasome pathways in skeletal muscle (Price *et al.* 1996, Mitch *et al.*1999,  
276 Lee *et al.* 2004, Satchek *et al.* 2004, Stitt *et al.* 2004, Wang *et al.*2006). Whereas the  
277 serum insulin concentration and phospho-Akt content of the plantaris muscle were almost  
278 negligible in the Rapid group, the Slow group rats maintained a serum insulin concentration  
279 and phospho-Akt content similar to that of the ad libitum-fed Con group rats (Table 1 and  
280 Fig. 2-G). In addition, the serum insulin concentration and phospho-Akt content were  
281 negatively correlated with the LC3-II/LC3-I ratio and polyubiquitinated protein concentration  
282 in the plantaris muscle (Fig. 4). Taken together, it is likely that the slow body weight  
283 reduction induced by daily calorie restriction can partially prevent muscular atrophy during  
284 weight loss, at least in part by maintaining the serum insulin level and its signaling pathway  
285 in skeletal muscle. Because it has been well documented that blood insulin level  
286 substantially decreases even after overnight fasting, we could not rule out the possibility  
287 that the higher proteolytic activities observed in the muscles of the Rapid group might be  
288 reflecting an acute effect of fasting (acute insulin deficiency) rather than chronic effects.

289 However, Ogata *et al.* (2010) reported that LC3-II content in rat skeletal muscle did not  
290 increase in response to 1-day fasting, whereas 3-day fasting induced significant and large  
291 increase in muscle LC3-II content. It is therefore plausible that higher proteolytic activities  
292 in the Rapid group resulted from chronic and accumulated effects of 3-day fasting, but not  
293 acute effect. Future studies are required to measure the proteolytic activities in the Con and  
294 Slow group in the fasting condition, or in the Rapid group after a few hours feeding in order  
295 to assess the chronic adaptations and differentiate them from potential acute effects.

296 As shown in Table 1, the soleus muscle, unlike the plantaris muscle, did not show  
297 any atrophic changes in response to either the rapid or slow body weight reduction. Our  
298 results support a previous finding that the degree of fasting-induced atrophy is greater in  
299 fast-twitch muscle than slow-twitch muscle (Li and Goldberg 1976, Frayn and Maycock  
300 1979). Ogata *et al.* (2010) reported that a fasting-induced increase in LC3-II expression  
301 was notably greater in rat fast-twitch plantaris muscle than in slow-twitch soleus muscle.  
302 Consistent with these results, we observed that the magnitudes of the increase in the  
303 LC3-II/LC3-I ratio after rapid and slow body weight reductions appeared to be relatively  
304 lower in soleus than plantaris muscle (Fig. 2-A and -B), providing further evidence that the  
305 autophagy pathway is preferentially induced in fast-twitch muscle in an energy deficient  
306 state. Another major finding of the present study was that the polyubiquitinated protein  
307 concentration was markedly higher in the soleus muscle than in the plantaris muscle under  
308 basal conditions (the Con group) and that it did not increase in response to fasting and  
309 daily calorie restriction (Fig. 2-C and -D). The blunted responses in the  
310 ubiquitin-proteasome and autophagy-lysosome systems might be associated with the  
311 atrophy resistance of soleus muscle to a severe energy deficit.

312 This study has several limitations. First, muscle strength and exercise capacity

313 after the rapid or slow weight loss were not assessed in this study. Thus, we could not  
314 clarify which weight loss strategy is effective in improving exercise performance although  
315 slow weight loss induced by calorie restriction could maintain muscle mass. Second, we  
316 did not evaluate the effects of fasting or calorie restriction in combination with exercise on  
317 muscle protein content and muscle weight. The results obtained in this study cannot be  
318 directly extrapolated to athletic population, who engage in exercise training. Future  
319 extensive studies are required to examine the combined effects of exercise and dietary  
320 interventions on muscle functions as well as muscle mass in order to elucidate whether  
321 exercise training can prevent muscle atrophy induced by weight loss.

322

## 323 **Conclusion**

324 During an equivalent weight loss, the rapid weight loss induced by short-term  
325 fasting more strongly activates autophagy-lysosome and ubiquitin-proteasome pathways  
326 than a slow body weight reduction induced by daily calorie restriction, resulting in muscular  
327 atrophy in fast-twitch plantaris muscle but not in soleus muscle.

328

## 329 **Conflict of interest**

330 The authors declare no conflict of interest.

331

## 332 **Acknowledgements**

333 This work was supported by JSPS KAKENHI Grant Numbers JP15K01615,  
334 JP25750330, and JP16J10555 (to S.T.).

335 **References**

- 336 BUJAK AL, CRANE JD, LALLY JS, FORD RJ, KANG SJ, REBALKA IA, GREEN AE, KEMP BE,  
337 HAWKE TJ, SCHERTZER JD, STEINBERG GR: AMPK Activation of Muscle Autophagy  
338 Prevents Fasting-Induced Hypoglycemia and Myopathy during Aging. *Cell Metab* **21**:  
339 883–890, 2015.
- 340 CHOMA CW, SFORZO GA, KELLER BA: Impact of rapid weight loss on cognitive function in  
341 collegiate wrestlers. *Med Sci Sports Exerc* **30**: 746–749, 1998.
- 342 ELIASSON J, ELFEGOUN T, NILSSON J, KÖHNKE R, EKBLÖM B, Blomstrand E: Maximal  
343 lengthening contractions increase p70S6 kinase phosphorylation in human skeletal muscle  
344 in the absence of nutritional supply. *Am J Physiol Endocrinol Metab* **291**: E1197–205,  
345 2006.
- 346 FRAYN KN, MAYCOCK PF: Regulation of protein metabolism by a physiological concentration  
347 of insulin in mouse soleus and extensor digitorum longus muscles. Effects of starvation  
348 and scald injury. *Biochem J* **184**: 323–30, 1979.
- 349 FUJITA S, ABE T, DRUMMOND MJ, CADENAS JG, DREYER HC, SATO Y, VOLPI E,  
350 RASMUSSEN BB: Blood flow restriction during low-intensity resistance exercise increases  
351 S6K1 phosphorylation and muscle protein synthesis. *J Appl Physiol* **103**: 903–10, 2007.
- 352 GOLDBERG AL: Protein degradation and protection against misfolded or damaged proteins.

353 *Nature* **426**: 895–9, 2003.

354 JACINTO E, HALL MN: TOR signalling in bugs, brain and brawn. *Nat Rev Mol Cell Biol* **4**:  
355 117–126, 2003.

356 JAGOE RT, LECKER SH, GOMES M, GOLDBERG AL: Patterns of gene expression in  
357 atrophying skeletal muscles: response to food deprivation. *FASEB J* **16**: 1697–712, 2002.

358 LEE JH, LEE JH, JIN M, HAN SD, CHON GR, KIM IH, KIM S, KIM SY, CHOI SB, NOH YH: Diet  
359 control to achieve euglycemia induces significant loss of heart and liver weight via  
360 increased autophagy compared with ad libitum diet in diabetic rats. *Exp Mol Med* **46**: e111,  
361 2014.

362 LEE SW, DAI G, HU Z, WANG X, DU J, MITCH WE: Regulation of muscle protein degradation:  
363 coordinated control of apoptotic and ubiquitin-proteasome systems by phosphatidylinositol  
364 3 kinase. *J Am Soc Nephrol* **15**: 1537–45, 2004.

365 LI JB, GOLDBERG AL: Effects of food deprivation on protein synthesis and degradation in rat  
366 skeletal muscles. *Am J Physiol* **231**: 441–8, 1976.

367 LOWRY OH, PASSONEAU JV: A Flexible System of Enzymatic Analysis. Academic Press;  
368 New York, 1972

369 MITCH WE, BAILEY JL, WANG X, JURKOVITZ C, NEWBY D, PRICE SR: Evaluation of  
370 signals activating ubiquitin-proteasome proteolysis in a model of muscle wasting. *Am J*

371 *Physiol* **276**: C1132–8, 1999.

372 MITCH WE, GOLDBERG AL: Mechanisms of muscle wasting. The role of the  
373 ubiquitin-proteasome pathway. *N Engl J Med* **335**: 1897–905, 1996.

374 MIYAZAKI M, MCCARTHY JJ, FEDELE MJ, ESSER KA: Early activation of mTORC1 signalling  
375 in response to mechanical overload is independent of phosphoinositide 3-kinase/Akt  
376 signalling. *J Physiol* **589**: 1831–46, 2011.

377 MIZUSHIMA N, YAMAMOTO A, MATSUI M, YOSHIMORI T, OHSUMI Y: In vivo analysis of  
378 autophagy in response to nutrient starvation using transgenic mice expressing a  
379 fluorescent autophagosome marker. *Mol Biol Cell* **15**: 1101–11, 2004.

380 MIZUSHIMA N, YOSHIMORI T: How to interpret LC3 immunoblotting. *Autophagy* **3**: 542–5,  
381 2007.

382 MORTIMORE GE, PÖSÖ AR: Intracellular protein catabolism and its control during nutrient  
383 deprivation and supply. *Annu Rev Nutr* **7**: 539–64, 1987.

384 OGATA T, OISHI Y, HIGUCHI M, MURAOKA I: Fasting-related autophagic response in slow-  
385 and fast-twitch skeletal muscle. *Biochem Biophys Res Commun.* **394**: 136–40, 2010.

386 PRICE SR, BAILEY JL, WANG X, JURKOVITZ C, ENGLAND BK, DING X, PHILLIPS LS,  
387 MITCH WE: Muscle wasting in insulinopenic rats results from activation of the  
388 ATP-dependent, ubiquitin-proteasome proteolytic pathway by a mechanism including gene

389 transcription. *J Clin Invest* **98**: 1703–8, 1996.

390 RELJIC D, HÄSSLER E, JOST J, FRIEDMANN-BETTE B: Rapid weight loss and the body fluid  
391 balance and hemoglobin mass of elite amateur boxers. *J Athl Train* **48**: 109–17, 2013.

392 ROCK KL, GRAMM C, ROTHSTEIN L, CLARK K, STEIN R, DICK L, HWANG D, GOLDBERG  
393 AL: Inhibitors of the proteasome block the degradation of most cell proteins and the  
394 generation of peptides presented on MHC class I molecules. *Cell* **78**: 761–71, 1994.

395 SACHECK JM, OHTSUKA A, MCLARY SC, GOLDBERG A: IGF-I stimulates muscle growth by  
396 suppressing protein breakdown and expression of atrophy-related ubiquitin ligases,  
397 atrogin-1 and MuRF1. *Am J Physiol Endocrinol Metab* **287**: E591–601, 2004.

398 STITT TN, DRUJAN D, CLARKE BA, PANARO F, TIMOFEYVA Y, KLINE WO, GONZALEZ M,  
399 YANCOPOULOS GD, GLASS DJ: The IGF-1/PI3K/Akt pathway prevents expression of  
400 muscle atrophy-induced ubiquitin ligases by inhibiting FOXO transcription factors. *Mol Cell*  
401 **14**: 395–403, 2004.

402 TAMURA Y, MATSUNAGA Y, MASUDA H, TAKAHASHI Y, TAKAHASHI Y, TERADA S,  
403 HOSHINO D, HATTA H: Postexercise whole body heat stress additively enhances  
404 endurance training-induced mitochondrial adaptations in mouse skeletal muscle. *Am J*  
405 *Physiol Regul Integr Comp Physiol* **307**: R931–43, 2014.

406 WANG X, HU Z, HU J, DU J, MITCH WE: Insulin resistance accelerates muscle protein

407 degradation: Activation of the ubiquitin-proteasome pathway by defects in muscle cell  
408 signaling. *Endocrinology* **147**: 4160–8, 2006.

409 WULLSCHLEGER S, LOEWITH R, HALL MN: TOR signaling in growth and metabolism. *Cell*  
410 **124**: 471–84, 2006

411

## 412 **Figure legends**

413 Figure 1. Changes in the body weights of rats during a 14-day dietary intervention. Con, ad  
414 libitum-fed control group; Slow, daily calorie restriction-induced slow body weight-loss  
415 group; Rapid, fasting-induced rapid weight-loss group. Values are means  $\pm$  SEM. <sup>#</sup>p<0.05,  
416 <sup>##</sup>p<0.01 vs Con and Rapid, respectively; \* p<0.001 vs Con; <sup>†</sup> p<0.05, <sup>††</sup> p<0.01 vs day 0  
417 in Slow group, respectively.

418

419 Figure 2. Effects of rapid or slow body weight loss on the LC3-II/-I ratio, polyubiquitinated  
420 protein concentration, phospho-p70S6K content, and phospho-Akt content in rat plantaris  
421 (A, C, E, and G) and soleus (B, D, F, and H) muscles. Con, ad libitum-fed control group;  
422 Slow, daily calorie restriction-induced slow body weight-loss group; Rapid, fasting-induced  
423 rapid weight-loss group; Values are means  $\pm$  SEM. \*\* and \*\*\* indicate significant  
424 differences from the values obtained in the Con group at p<0.01 and p<0.001, respectively.  
425 <sup>§</sup>, <sup>§§</sup>, and <sup>§§§</sup> indicate significant differences from the values obtained in the Slow group at  
426 p<0.05, p<0.01, and p<0.001, respectively.

427

428 Figure 3. Correlations between the LC3-II/-I ratio (A and C) and the polyubiquitinated  
429 protein concentration (B and D) and the total protein content and wet weight of rat plantaris

430 muscle. Con, ad libitum-fed control group; Slow, daily calorie restriction-induced slow body  
431 weight-loss group; Rapid, fasting-induced rapid weight-loss group.

432 Figure 4. Correlations between the serum insulin concentration or phospho-Akt content  
433 and LC3-II/-I ratio (A and C) or polyubiquitinated protein concentration (B and D) of rat  
434 plantaris muscle. Con, ad libitum-fed control group; Slow, daily calorie restriction-induced  
435 slow body weight-loss group; Rapid, fasting-induced rapid weight-loss group.

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Table 1. Body weight, intra-abdominal fat weight, total food intake, muscle wet weight, total protein content, serum glucose, insulin and muscle glycogen concentrations in rats.

	Con	Slow	Rapid
Initial body weight (g)	303 ± 5	304 ± 5	304 ± 4 <sup>***</sup>
Final body weight (g)	321 ± 6	270 ± 2 <sup>***</sup>	265 ± 3 <sup>***</sup>
Intra-abdominal fat weight (g)	13.7 ± 0.3	8.5 ± 0.5 <sup>***</sup>	9.2 ± 0.9 <sup>***</sup>
Food intake (g)	255 ± 7	170 ± 1 <sup>***</sup>	193 ± 3 <sup>***,§</sup>
Plantaris muscle wet weight (mg)	287 ± 9	267 ± 6	251 ± 6 <sup>*</sup>
Plantaris muscle protein content (mg)	56 ± 3	54 ± 1	49 ± 1 <sup>*</sup>
Soleus muscle wet weight (mg)	114 ± 3	103 ± 6	105 ± 2
Soleus muscle protein content (mg)	20 ± 1	18 ± 1	19 ± 1
Serum glucose (mmol/L)	12.4 ± 0.5	13.2 ± 0.3	11.1 ± 0.7
Serum insulin (µg/L)	7.6 ± 0.6	3.8 ± 0.4 <sup>***</sup>	1.0 ± 0.4 <sup>***,§§</sup>
Muscle glycogen concentration (µmol/g wet tissue)	47.1 ± 1.2	41.8 ± 1.6	27.7 ± 1.7 <sup>***,§§§</sup>

445 Values are means ± SEM, n=5. \* and \*\*\* indicate significant differences from the values  
 446 obtained in the Con group at p<0.05 and p<0.001, respectively. §, §§ and §§§ indicate  
 447 significant differences from the values obtained in the Slow group at p<0.05, p<0.01 and  
 448 p<0.001, respectively.

Figure 1

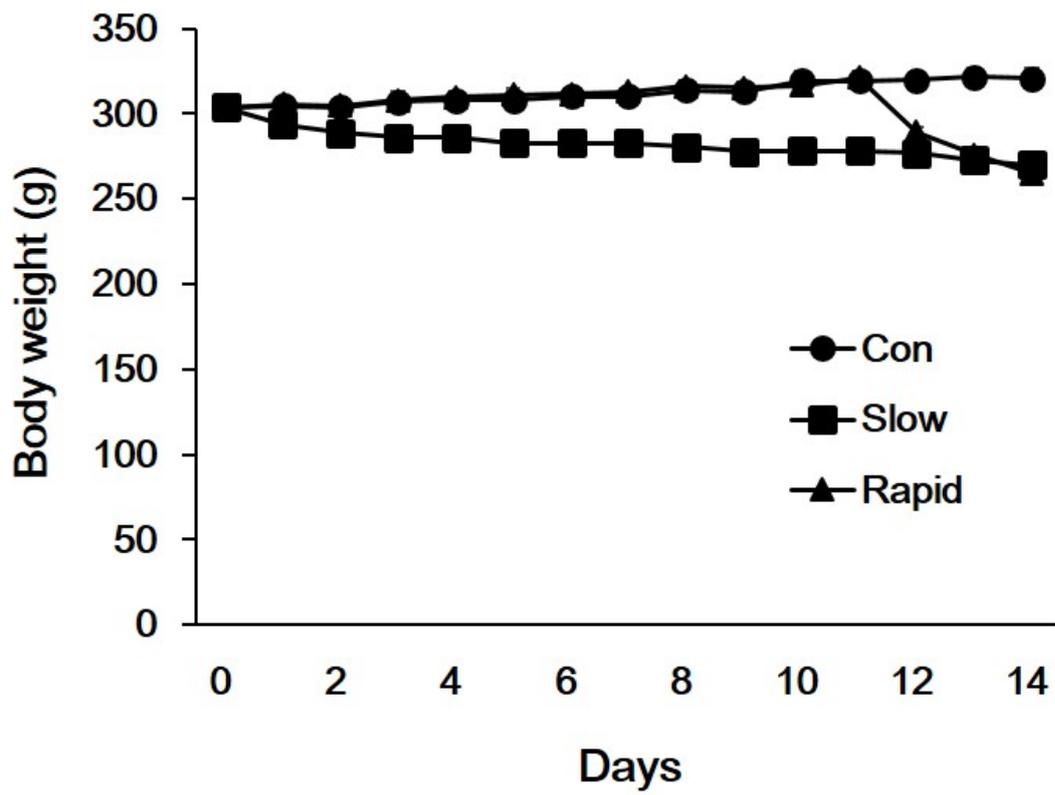


Figure 2

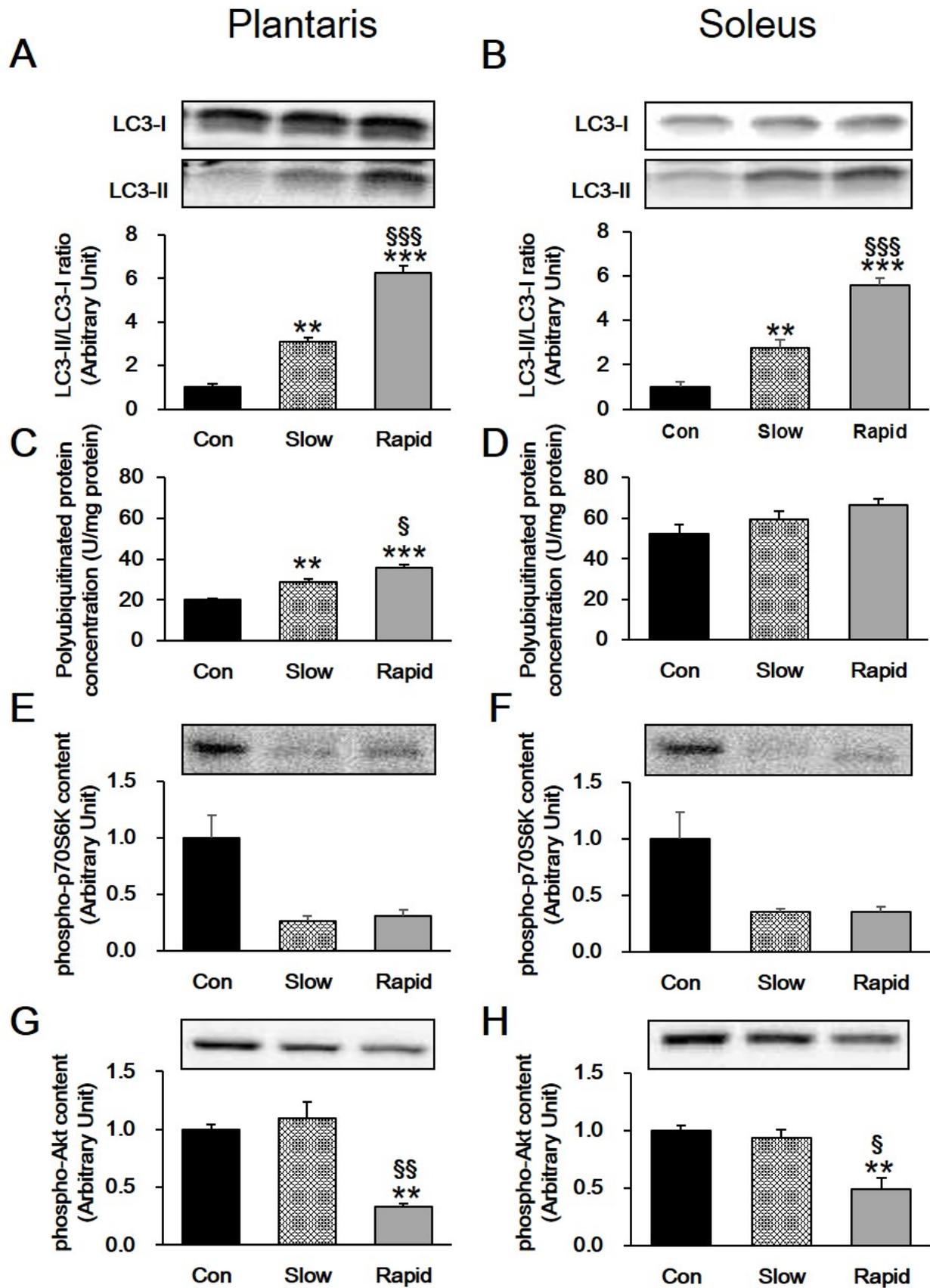


Figure 3

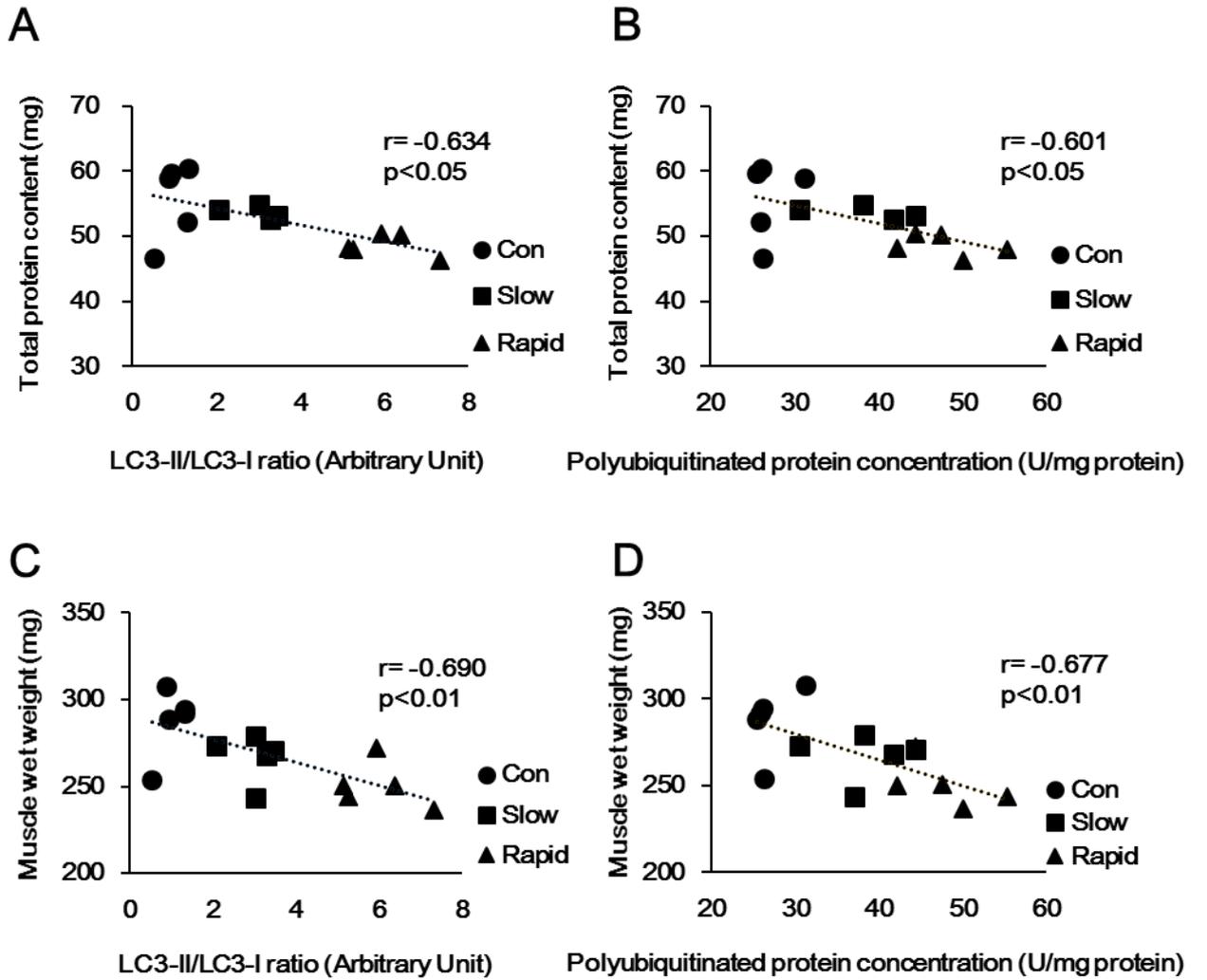


Figure 4

