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## Dietary wild bitter gourd displays selective androgen receptor modulator like activity and improves the muscle decline of orchidectomized mice†

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Loss of skeletal muscle mass and strength is often associated with disability and poor quality of life. Selective Androgen Receptor Modulators (SARMs) are under development as potential treatment. This study aims at examining the potential of wild bitter gourd (BG) as a SARM and its effects on the muscle decline induced by orchietomy. In the cell-based androgen receptor (AR) transactivation assay, the BGP extract showed weak agonistic and antagonistic activities, resembling those of some SARMs. Male C57BL/6J mice were sham-operated (Sham group) or castrated (Cast groups) and fed a modified AIN-93G high sucrose diet supplemented without (Cast group) or with 5% lyophilized BG powder (Cast + BGP) or with testosterone propionate (7 mg TP per kg diet, Cast + TP) for 23 weeks. In contrast to the Cast + TP group, the BGP supplementation did not affect the serum testosterone concentration, and prostate and seminal vesicle mass. Both TP and BGP supplementation increased the weight of androgen responsive muscles, bulbocavernosus (BC) and levator ani (LA) ( $p < 0.05$ ). The grip strength and the performance on a rotarod of the Cast + BGP group were comparable to those of the Cast + TP group ( $p > 0.05$ ). The number of succinate dehydrogenase (SDH)-positive fibers of the Cast + BGP group was not significantly different from that of the Sham and Cast + TP groups ( $p > 0.05$ ). The BGP supplementation up-regulated the *Pgc1 $\alpha$* , *Ucp2* or *Ucp3* gene expressions in skeletal muscles of castrated mice ( $p < 0.05$ ). BGP showed some characteristics of the SARM and might improve skeletal muscle function through the up-regulation of mitochondrial biogenic genes and oxidative capacity, and ameliorated the castration-induced decline of skeletal muscle function in mice.

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### 1. Introduction

Loss of skeletal muscle mass and strength, resulting from ageing or co-morbidity of various diseases, is often associated with disability and poor quality of life.<sup>1</sup> Sarcopenia, a gradual loss of skeletal muscle mass and function with age, and cachexia, weight loss due to an underlying illness, are the two most common muscle wasting disorders and represent important syndromes impacting quality and quantity of life in the geriatric population.<sup>2</sup> Besides exercise and nutrition, the management of these conditions also includes hormones/signaling molecules known to enhance muscle development.<sup>2</sup> For

example, testosterone, a primary androgenic hormone in men, exerts a broad range of male physiological functions, and has a potent anabolic effect on skeletal muscles.<sup>3</sup> Experimental androgen deprivation using either castration or androgen receptor knockout (ARKO) animal models typically results in a rapid loss of muscle mass and decreased muscle strength.<sup>4,5</sup> Interestingly, not all muscles respond to androgen deprivation in the same way in these animal studies. The bulbocavernosus (BC) and levator ani (LA), skeletal muscles that attach to the base of the penis, have been found to be highly responsive to androgen.<sup>6</sup> In contrast, limb muscles, such as triceps brachii (TRI) and extensor digitorum longus (EDL), are relatively less responsive.

Epidemiologic studies have demonstrated a correlation among the bioavailable testosterone concentration, fat-free mass and muscle strength.<sup>7,8</sup> Hormone replacement therapy with androgens improved muscle function in both castrated animals<sup>9</sup> and hypogonadal individuals.<sup>10,11</sup> However, the uncertainty of the side effects associated with testosterone replacement therapy has limited its wide applications.<sup>12</sup>

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Currently, selective androgen receptor modulators (SARMs) are under development for the treatment of sarcopenia<sup>13</sup> or cancer cachexia.<sup>14</sup> SARMs form a novel class of androgen receptor ligands which provide anabolic benefits in the absence of androgenic effects on sexual organs.<sup>15</sup> The development of SARMs is intended to offer anabolic effects in muscle and bone, and they might also revert other castration-related adverse effects such as those in adipose tissues. As SARMs have an optimal effect on myo-anabolic activity but less of growth effects on prostate and other secondary sexual organs than traditional androgen therapy, they are promising as a new class of function promoting anabolic therapy for multiple clinical indications.<sup>16,17</sup>

Androgen deficiency contributes to mitochondrial dysfunction and disruption of normal cellular function, including production of ATP and promoting cell death.<sup>18</sup> Mitochondrial dysfunction adversely affected the quality and quantity of skeletal muscles and was closely related to the loss of skeletal muscle mass with ageing<sup>19</sup> and cachexia.<sup>20</sup> Peroxisome proliferator-activated receptor gamma coactivator 1-alpha (PGC1 $\alpha$ ) plays a crucial role in maintaining muscle metabolic function and controls numerous genes that affect muscle morphology and physiological functions.<sup>21</sup> The expression levels of PGC1 $\alpha$  in the gastrocnemius skeletal muscle is up-regulated by supplementation of supra-physiological testosterone and down-regulated by AR knockout.<sup>22</sup> The roles of PGC1 $\alpha$  in enhancing mitochondria biogenesis<sup>23</sup> and suppressing gene expressions of forkhead transcription factors 3 (*FoxO3*), atrophy-specific atrogen-1 (*Fbxo32*), and Muscle RIN-finger protein-1 (*MuRF-1* or *Trim63*) in skeletal muscles<sup>24</sup> have made it another molecular target for developing therapeutics for muscle wasting.

Bitter melon (*Momordica charantia*, BG) is a common tropical vegetable belonging to the Cucurbitaceae family. It has traditionally been used in herbal medicine. Scientific evidence for its biological effects,<sup>25</sup> especially for metabolic disorders<sup>26</sup> have been addressed. BG supplementation improved insulin sensitivity in the skeletal muscles.<sup>27,28</sup> Our previous study demonstrated that BG supplementation significantly up-regulated the PGC1 $\alpha$  mRNA level in the gastrocnemius muscle of mice.<sup>29</sup> Recently, Hsiao *et al.* reported that 4-week supplementation of BG enhanced muscle strength and reduced fatigue in male ICR mice.<sup>30</sup> Bioactive compounds isolated from BG can activate nuclear receptors, including peroxisome proliferator activated receptors (PPARs)<sup>31</sup> and estrogen receptors (ERs).<sup>32</sup> In this study, we examined the BG extract and index compounds for the transactivation of the androgen receptor (AR) in a cell-based assay. The orchidectomized mouse model was then used to evaluate the SARM-like characteristics of BG.

## 2. Materials and methods

### 2.1 Preparation of wild bitter melon powder and experimental diets

The strain 2381 of wild bitter melon bred and provided by Hualien District Agricultural Research and Extension Station

was used. Whole fresh fruits (with seeds) were sliced, freeze dried and ground to produce the bitter melon powder (BGP). The high sucrose (HS) diet was modified from the AIN-93G<sup>33</sup> diets in which carbohydrate was provided by 50% sucrose and 12.95% cornstarch. The BGP diet was formulated by incorporating 5% (w/w) of BGP into the HS diet. Testosterone propionate (TP) (Sigma-Aldrich, St Louis, MO, USA) was incorporated into the HS diet (7 mg TP per kg diet, TP diet) as a positive control. The composition of experimental diets is listed in ESI Table 1.†

### 2.2 Animals

Six-week old male C57BL/6J mice were purchased from the National Laboratory Animal Centre (Taipei, Taiwan) and housed individually in steel wire cages in an animal room with controlled temperature and a light-dark cycle. Mice were acclimatized for 2 weeks and then randomly assigned to castration or sham surgery. After a 1-week surgical recovery period, they were further assigned to four experimental groups: sham-operated mice fed the HS diet (Sham,  $n = 7$ ), castrated mice fed the HS diet (Cast,  $n = 8$ ), castrated mice fed the BGP diet (Cast + BGP,  $n = 8$ ) and castrated mice fed the TP diet (Cast + TP,  $n = 7$ ). The animal study was approved by the Institutional Animal Care and Use Committee (IACUC) of National Taiwan University (No. NTU-105-EL-00009).

### 2.3 Castration and sham-operated surgery

Surgery was performed under sterile conditions. Mice were anesthetized by an *i.p.* injection of ketamine (100 mg kg<sup>-1</sup>) and xylazine (20 mg kg<sup>-1</sup>). Then, both testes were gently pushed from the abdomen into the scrotum and a 5 mm incision was made on the midline of the scrotum. Another smaller incision was made on the parietal tunica, and then one of the testes was carefully pulled out through the incision till the spermatic cord was exposed. The spermatic cord was clamped with a hemostat and ligated above, and the spermatic cord below the hemostat through the testis was dissected away. The remaining tissue was placed back into the scrotum. This procedure was repeated on the contralateral testis. The scrotal incision was closed with sutures. The sham-operated animals were anesthetized and incisions were made. Their testes were gently pulled out and then placed back into the scrotum, the incisions were then closed with sutures. The absence of testes in every castrated mouse was confirmed at necropsy after 23 weeks of feeding.

### 2.4 Serum testosterone level analysis

Blood samples were collected from the retro-orbital sinus under light anesthesia. The collected blood samples were centrifuged twice at 12 000g, 4 °C, for 20 min. Serum samples were isolated for measuring serum testosterone concentrations using the Testosterone ELISA kit (Cayman Chemical Company, Ann Arbor, Michigan, USA) following the manufacturer's instructions.

## 2.5 Muscle strength and performance tests

The grip strength and rotarod performance of the tested mice were examined during the experimental period. To minimize variability, each test was conducted in the evening of the same day and in the same environment by the same operator.

To measure the forelimb grip strength, a mouse was allowed to grab a metal wire grid with its forelimb and then pulled backwards in the horizontal plane with increasing force until it released the grid. The maximal force (in gram) was recorded using an automatic force transducer (Muromachi, Japan). The test consisted of two sessions with at least 30 seconds of rest between the two sessions. In each session, each mouse was tested five times with 5 second intervals. The mean maximal force was taken as the grip strength.

A 5-lane accelerating rotarod (TSE Systems, Bad Homburg, Germany) was used for measuring muscle endurance and motor abilities. Before the test, mice were trained at a relatively slow speed (12 rpm) 3 times with 3 min at each time and at least 10 min intervals on each day. The training was repeated for four consecutive days. On the test days, mice were placed on a static rod, and the motor was turned on and accelerated from 5 to 40 rpm over 5 min until all mice fell off the rod. Each mouse was tested six times with at least 30 min interval between tests. The mean duration times (in second) the mouse remained on the rod were recorded as the latency to fall.

## 2.6 Blood and tissue sample collection

After 23 weeks of feeding, mice were fasted overnight and killed by CO<sub>2</sub> asphyxiation. Blood samples were withdrawn through cardiac puncture. Liver, seminal vesicle (SV)/prostate, testis, epididymis, adrenal gland and skeletal muscle tissue including gastrocnemius (GAS), soleus (SOL), tibialis anterior (TA), extensor digitorum longus (EDL), triceps brachii (TRI), levator ani (LA) and bulbocavernosus (BC) were excised, weighed and immediately frozen in liquid nitrogen and stored at -80 °C for mRNA analysis. For histology analysis, muscle tissues were fixed by overnight immersion in 30% sucrose/PBS at 4 °C. Muscle tissues were then embedded in an optimal cutting temperature compound (OCT, Sakura Finetek USA, Torrance, CA, USA) in dry-ice-cooled 2-methylbutane (Sigma) and stored at -80 °C until preparation of cryosections.

## 2.7 Histological analysis

Cryosections were cut from the mid-belly region of TA, GAS and TRI muscles on a cryostat (Leica 3050 S, Leica GmbH, Germany) at -23 °C to -25 °C. The sections were then placed on poly-L-lysine coated glass slides (Sigma) and stored at -20 °C until the analysis of succinate dehydrogenase (SDH) activity within a month. For SDH staining, slides were soaked in PBS at room temperature for 10 min and then dried for 5 min before incubation in an SDH working solution, which consisted of 0.24 M disodium succinate (Sigma), 1 mM 1-methoxyphenazine methosulphate (mPMS, Sigma), 1.5 mM nitroblue tetrazolium (NBT, Sigma) and azide/EDTA/PO<sub>4</sub> buffer (including 0.02 M sodium azide, 0.125 M disodium EDTA,

0.3 M NaH<sub>2</sub>PO<sub>4</sub> and 0.3 M Na<sub>2</sub>HPO<sub>4</sub> in deionized water, pH = 7.4) at room temperature for 15 min. Slides were then washed in deionized water to stop the enzymatic reaction and air-dried for 10 min in the dark. Slides were mounted with mounting medium (Leica) and stored at -20 °C. For the evaluation of the muscle fiber cross-section area (CSA), hematoxylin and eosin (H&E) staining was performed on muscle cryosections. Briefly, the slides were fixed in 4% paraformaldehyde overnight and then H&E stained following the standard protocol.<sup>34</sup> Digital photographs were taken from each section under a microscope (Leica DM1000LED), which was connected to a computerized imaging system. All images were analyzed by manually quantifying the SDH intensity and fiber size with the Image J software (NIH, U.S. National Institutes of Health, Bethesda, MD, USA). In SDH distribution and intensity analyses, all images were converted to a grayscale for histogram analysis with Image J. The intensities of the gray-scale images were expressed on a scale of 0–255 (0: black, 255: white). The number of myofibers with various intensities were quantified and expressed in a distribution graph.

## 2.8 RNA extraction and quantification of mRNA

Total RNA was isolated from various muscles by using the TRIZOL reagent (Invitrogen, Carlsbad, CA, USA) and an RNA miniprep kit (ZYMO Research, Irvine, CA, USA) following the manufacturer's instruction. Total RNA (2 µg) was reverse transcribed to cDNA in a mixture containing 10× RT (reverse transcription) buffer, 100 mM dNTP mixture buffer, 10× RT random primer and MultiScribe™ RTase (Invitrogen, Carlsbad, CA, USA). The polymerase chain reaction (PCR) was performed in a final volume of 25 µL containing 10 µL cDNA, 12.5 µL TaqMan® Gene Expression Master Mix, 1.25 µL probe/primer assay mix and water. PCR primers and probes were purchased from Applied Biosystems (Carlsbad, CA, USA; ESI Table 2†). The PCR and fluorogenic analysis was performed using a CFX connect™ real-time PCR Detection System (Bio-Rad Laboratories, Berkeley, CA, USA). Amplification conditions were 2 min at 50 °C, 10 min at 95 °C then 40 cycles of 15 s at 95 °C and 1 min at 60 °C. Gene expression was quantified using the relative (ΔΔCT) method with normalization to the levels of glyceraldehyde 3-phosphate dehydrogenase (*Gapdh*) mRNA expression.

## 2.9 Transactivation assay for AR activity

BGP was extracted with ethyl acetate (1 : 20, w/v) with stirring at room temperature overnight. The ethyl acetate extract (EAE) of BGP was obtained by filtration and evaporation to remove the solvent. The extraction rates of the EAE was 9.4%. This BGP extract was dissolved in a minimal amount of absolute ethanol and diluted with cell culture medium to concentrations that did not affect cell growth or viability.

Androgen receptor transactivation activity was examined using a cell-based transactivation assay. Briefly, plasmids including pcDNA3-AR, pRL-TK (renilla luciferase plasmid, normalization vector) and 3× ARE Δ56-c-fos GL3 (reporter gene vector), provided by Dr Chih-Pin Chuu (Institute of Cellular and System Medicine, National Health Research Institutes, Miaoli, Taiwan), were transiently co-transfected into CHO-K1

cells (American Type Culture Collection; Manassas, VA, USA) using Lipofectamine™ 2000 (Invitrogen). The transfected cells were treated with varying concentrations of the BGP extract, conjugated linoleic acid (*cis*-9, *trans*-11-CLA or *trans*-10, *cis*-12-CLA) or 9*c*,11*t*,13*t*-CLN ( $\alpha$ -eleostearic acid) in the absence or presence of 5 $\alpha$ -dihydrotestosterone (5 $\alpha$ -DHT, androgen receptor agonist; Wako, Japan) for 24 h. Luciferase activity was then measured using a Dual-Glo® Luciferase kit (Promega, Madison, Wisconsin, USA). 9*c*,11*t*,13*t*-CLN is one of the active compounds in BG that can activate PPARs and can be converted to 9*c*,11*t*-CLA in rats and mice.<sup>35–37</sup> The conjugated fatty acids were purchased from Cayman Chemical Co. (Ann Arbor, MI, USA). Samples were dissolved in absolute ethanol and diluted to concentrations that did not affect the cell viability. Folds of activation were calculated by taking the vehicle-treated cells as 1.0.

## 2.10 Statistical analysis

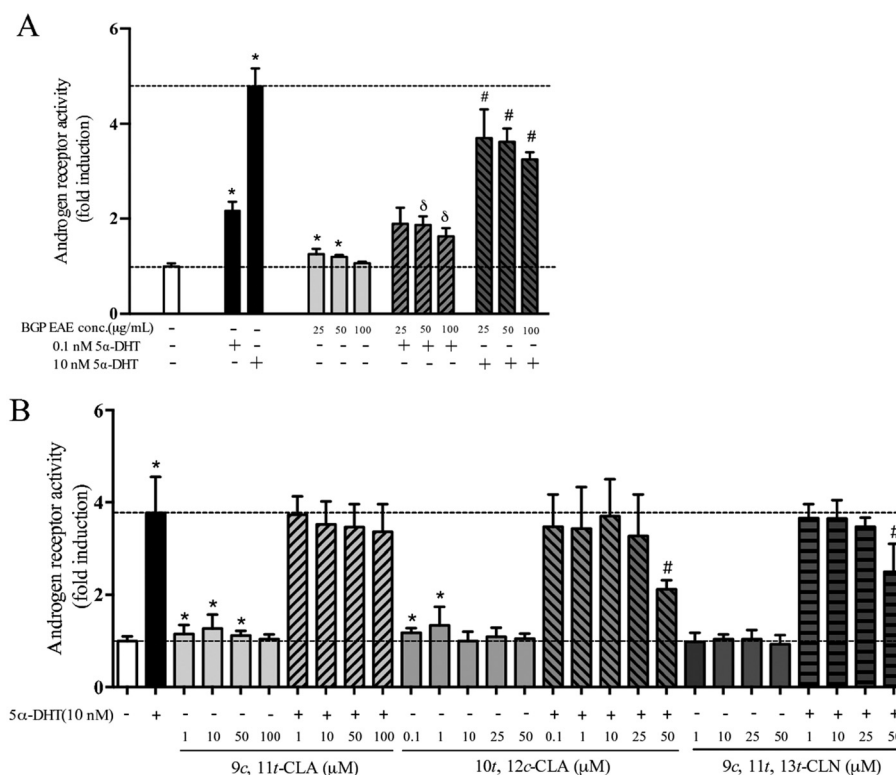
Data are presented as means  $\pm$  SD. Data were checked for the adherence to normal distribution. Those that did not were transformed to log, reciprocal or root square values. The significance of differences was analyzed by one-way ANOVA. Where there was a significant effect ( $p < 0.05$ ), a Duncan's *post-hoc* test was performed for comparison among groups. In some cases, a two-sample Student's *t* test was further performed to compare

between the supplemented Cast group and the Cast group. The Pearson correlation analysis was used to examine the association between muscle mass and gene expression levels. All analyses were conducted using the SPSS, version 21.0 (SPSS, Chicago, IL).  $P$  value  $< 0.05$  was considered significant.

## 3 Results

### 3.1 The ethyl acetate extract of BGP weakly modulates the AR activation activity in the cell-based transactivation assay

In the cell-based AR transactivation assay, the dose response was tested in the concentration ranges that did not show cytotoxicity, 25–100  $\mu\text{g mL}^{-1}$  for the BGP extract, 1–100  $\mu\text{M}$  for 9*c*,11*t*-CLA, 1–50  $\mu\text{M}$  for 9*c*,11*t*,13*t*-CLN, and 0.1–50  $\mu\text{M}$  for 10*t*,12*c*-CLA. The BGP extract (50  $\mu\text{g mL}^{-1}$ ) increased 5.3% of the AR transactivation but suppressed 26.2% and 30.9% of the AR activation when co-treated with 0.1 nM or 10 nM 5 $\alpha$ -DHT, respectively ( $p < 0.05$ , Fig. 1A). Similarly, 9*c*,11*t*-CLA (10  $\mu\text{M}$ ) and 10*t*,12*c*-CLA (1  $\mu\text{M}$ ) increase 27% and 34% of AR transactivation, respectively ( $p < 0.05$ , Fig. 1B). 9*c*,11*t*,13*t*-CLN (50  $\mu\text{M}$ ) and 10*t*,12*c*-CLA (50  $\mu\text{M}$ ) suppressed 36% and 46% of the AR activation when co-treated with 10 nM 5 $\alpha$ -DHT, respectively ( $p < 0.05$ , Fig. 1B).



**Fig. 1** Transactivation assay of an AR using a BGP ethyl acetate extract (EAE) and conjugated fatty acids with or without 5 $\alpha$ -DHT. CHO-K1 cells were co-transfected with pcDNA3-AR, 3 $\times$  ARE  $\Delta$ 56-c-fos GL3 and pRL-TK (renilla luciferase control plasmid). Four hours after the transfection, cells were treated with 5 $\alpha$ -DHT or various concentrations of test samples for 24 h and then harvested. Dual luciferase activities were measured and firefly values were normalized with Renilla values and then calculated as the fold of activation by taking the vehicle-treated cells as 1.0. Values are means  $\pm$  SD of at least three separate experiments. 5 $\alpha$ -Dihydrotestosterone (5 $\alpha$ -DHT) is the positive control for the AR. Data were analyzed by Student's *t* test. \* denotes significant difference from the vehicle (veh).  $\delta$  and # denote significant difference from 0.1 nM DHT and 10 nM DHT, respectively.



### 3.2 Dietary BGP increased the mass of androgen responsive muscle but not SV/prostate in castrated mice

At the beginning of the feeding study, all mice had similar body weights ( $23.5 \pm 0.6$  g). After 23 weeks of feeding, the Cast + BGP group had significantly lower final body weight compared to the Sham and Cast groups ( $p < 0.05$ ). At necropsy, the absolute weights of androgen responsive tissues/organs including epididymis, SV/prostate, and LA and BC muscles of the Cast group were significantly lower than those of the Sham group ( $p < 0.05$ ) (Table 1). Compared to the Cast group, the Cast + TP group had a significantly higher mass of SV/prostate as well as LA, BC and quadriceps ( $p < 0.05$ ). In the Cast + BGP group, weights of LA and BC were also higher than those of the Cast group ( $p < 0.05$ ). However, the mass of the SV/prostate was comparable to that of the Cast group ( $p > 0.05$ ). Similar results were observed when these tissue weights were calcu-

lated as % of the body weight except that the relative weight of quadriceps and GAS of the Cast + BGP group was comparable to those of the Cast + TP group ( $p > 0.05$ ) (Table 1). Thus, the castration-induced mass reduction of the highly androgen responsive muscles<sup>38</sup> was restored (LA) or partially restored (BC) by both TP and BGP treatment, but BGP did not affect the weights of prostate and seminal vesicles, in contrast to the TP treatment.

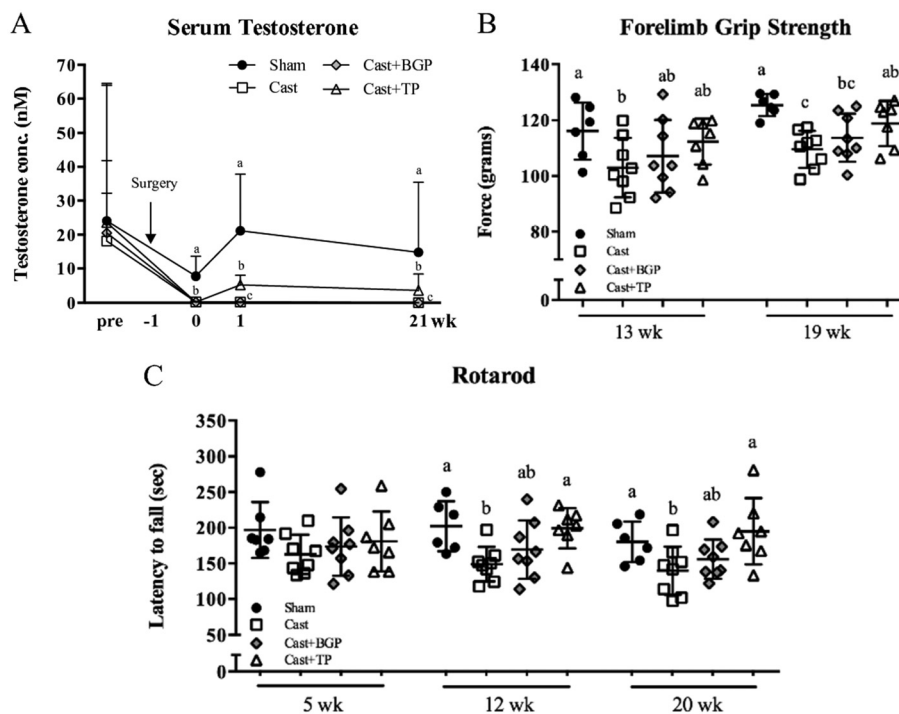
### 3.3 BGP did not affect the testosterone concentration but ameliorated the castration-induced decline of muscle strength and performance in mice

As shown in Fig. 2, all 4 groups have similar serum testosterone levels before surgery. After castration, serum testosterone declined rapidly as reported.<sup>39</sup> BGP supplementation did not affect the serum testosterone level in castrated mice. In con-

**Table 1** The final body weight and organ/tissue weight of sham or castrated mice fed test diets for 23 weeks

Group	Sham	Cast	Cast + BGP	Cast + TP
Final body weight (g)	$37.9 \pm 3.6^a$	$34.7 \pm 2.4^b$	$30.7 \pm 2.1^c$	$32.3 \pm 3.6^{bc}$
Absolute organ/tissue weight (g)				
	Sham	Cast	Cast + BGP	Cast + TP
<b>Organ/tissue weight</b>				
Liver	$1.532 \pm 0.241^a$	$1.484 \pm 0.114^{ab}$	$1.284 \pm 0.092^c$	$1.324 \pm 0.137^{bc}$
Testis	$0.196 \pm 0.010^b$	N/A	N/A	N/A
Epididymis	$0.123 \pm 0.022^a$	$0.009 \pm 0.025^b$	$0.003 \pm 0.008^b$	$0.004 \pm 0.007^b$
SV <sup>a</sup> /prostate	$0.519 \pm 0.039^a$	$0.004 \pm 0.004^c$	$0.005 \pm 0.006^c$	$0.133 \pm 0.021^b$
Adrenal gland	$0.006 \pm 0.002^b$	$0.009 \pm 0.002^a$	$0.008 \pm 0.002^a$	$0.008 \pm 0.002^{ab}$
<b>Skeletal muscles</b>				
LA <sup>a</sup>	$0.078 \pm 0.017^a$	$0.038 \pm 0.014^b$	$0.065 \pm 0.013^a$	$0.071 \pm 0.028^a$
BC <sup>a</sup>	$0.098 \pm 0.007^a$	$0.015 \pm 0.004^d$	$0.024 \pm 0.009^c$	$0.042 \pm 0.009^b$
Triceps brachii	$0.253 \pm 0.014^a$	$0.217 \pm 0.030^{ab}$	$0.209 \pm 0.013^b$	$0.226 \pm 0.023^{ab}$
Quadriceps	$0.411 \pm 0.033^a$	$0.377 \pm 0.021^b$	$0.400 \pm 0.024^{ab}$	$0.427 \pm 0.035^a$
Gastrocnemius	$0.345 \pm 0.021^a$	$0.328 \pm 0.017^a$	$0.303 \pm 0.011^b$	$0.333 \pm 0.022^a$
Soleus	$0.021 \pm 0.003$	$0.024 \pm 0.003$	$0.024 \pm 0.003$	$0.023 \pm 0.004$
Tibialis anterior	$0.098 \pm 0.017$	$0.095 \pm 0.020$	$0.095 \pm 0.013$	$0.098 \pm 0.014$
Relative organ/tissue weight (% body weight)				
	Sham	Cast	Cast + BGP	Cast + TP
<b>Organ/tissue</b>				
Liver	$4.022 \pm 0.284$	$4.285 \pm 0.367$	$4.197 \pm 0.272$	$4.113 \pm 0.347$
Testis	$0.522 \pm 0.064$	N/A	N/A	N/A
Epididymis	$0.327 \pm 0.072^a$	$0.023 \pm 0.066^b$	$0.009 \pm 0.024^b$	$0.011 \pm 0.020^b$
SV <sup>a</sup> /prostate	$1.373 \pm 0.096^a$	$0.010 \pm 0.012^b$	$0.017 \pm 0.021^b$	$0.409 \pm 0.042^{ab}$
Adrenal gland	$0.016 \pm 0.006^b$	$0.026 \pm 0.006^a$	$0.027 \pm 0.007^a$	$0.025 \pm 0.005^a$
<b>Skeletal muscles</b>				
LA <sup>a</sup>	$0.208 \pm 0.055^a$	$0.112 \pm 0.046^b$	$0.214 \pm 0.046^a$	$0.218 \pm 0.070^a$
BC <sup>a</sup>	$0.262 \pm 0.041^a$	$0.042 \pm 0.011^d$	$0.078 \pm 0.031^c$	$0.130 \pm 0.021^b$
Triceps brachii	$0.673 \pm 0.078$	$0.626 \pm 0.084$	$0.684 \pm 0.067$	$0.709 \pm 0.123$
Quadriceps	$1.091 \pm 0.127^b$	$1.094 \pm 0.126^b$	$1.310 \pm 0.096^a$	$1.330 \pm 0.110^a$
Gastrocnemius	$0.914 \pm 0.049^c$	$0.947 \pm 0.045^{bc}$	$0.990 \pm 0.042^{ab}$	$1.038 \pm 0.088^a$
Soleus	$0.057 \pm 0.008^b$	$0.069 \pm 0.008^a$	$0.079 \pm 0.011^a$	$0.072 \pm 0.009^a$
Tibialis anterior	$0.259 \pm 0.050$	$0.277 \pm 0.067$	$0.312 \pm 0.047$	$0.307 \pm 0.058$
EDL <sup>a</sup>	$0.056 \pm 0.011$	$0.065 \pm 0.010$	$0.068 \pm 0.004$	$0.68 \pm 0.017$

<sup>a</sup> SV: seminal vesicle, LA: levator ani muscle, BC: bulbocavernosus muscle, EDL: extensor digitorum longus. Mice were sham-operated or orchidectomized and fed test diets for 23 weeks. The relative organ weight is expressed as the percentage of the body weight. The values represent mean  $\pm$  SD. Data were analyzed by one-way ANOVA and Duncan's multiple comparison test. Values not sharing the same superscripted letter are significantly different at  $p < 0.05$ .



**Fig. 2** Serum testosterone concentration and muscle function tests of sham or castrated mice supplemented with BGP or TP. (A) Serum testosterone concentration; (B) average forelimb grip strength in grams; (C) latency to fall in seconds in the accelerating rotarod test (5–40 rpm/5 min). In (B) and (C), each data point is the mean value of each mouse and the group means and SD are also shown. Values are means  $\pm$  SD. Data were analyzed by one-way ANOVA and Duncan's multiple range test. Data of serum testosterone levels were transformed to log values to conform to normality before statistical analysis. Values not sharing the same superscript letters are significantly different at  $p < 0.05$ .

trast, the Cast + TP group had significantly higher serum testosterone levels compared to the Cast and Cast + BGP groups ( $p < 0.05$ ). However, the levels of serum testosterone of the Cast + TP group was still lower than that of the Sham group ( $p < 0.05$ ) (Fig. 2A).

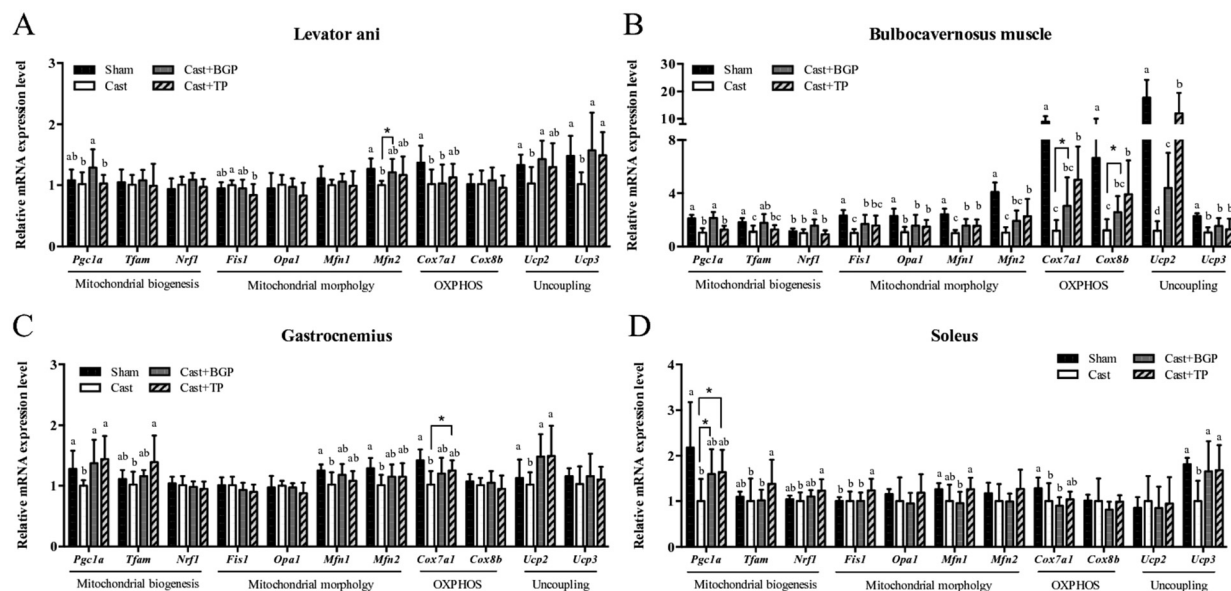
Muscle strength and motor coordination performance were measured during the experimental period. After feeding test diets for 19 weeks, the forelimb grip strength showed an 11% reduction in the Cast group compared to that of the Sham group ( $p < 0.05$ ), while TP treatment restored it (Fig. 2B). In the Cast + BGP group, the grip strength was comparable to that of the Cast + TP group ( $p > 0.05$ ) (Fig. 2B). After feeding test diets for 12 weeks, the Sham and Cast + TP groups showed significantly longer duration times on the rotarod than the Cast group ( $p < 0.05$ ) (Fig. 2C). At 20 weeks, the duration time on the rotarod showed a 19% reduction in the Cast group compared to that in the Sham group, while TP supplementation restored it to the level of the Sham group. The performance of the Cast + BGP group on the rotarod was also comparable to that of the Cast + TP group ( $p > 0.05$ ).

### 3.4 BGP increased the expression of *Pgc1 $\alpha$* and improved oxidative capacity in skeletal muscles

The mRNA expression levels of *Pgc1 $\alpha$*  and its target genes in some muscles are shown in Fig. 3. Remarkably, the Cast group showed significantly lower mRNA expression levels of most

genes related to mitochondrial biogenesis, morphology and oxidative phosphorylation (OXPHOS) in the androgen responsive BC muscle. Similarly, in GAS (consisting of >50% fast-twitch muscle fibers) and SOL (fast-twitch) muscles, the mRNA expressions of *Pgc1 $\alpha$*  and some of its target genes associated with mitochondrial function were also lower in the Cast group compared to those of the Sham group (Fig. 3C and D). Interestingly, the Cast + BGP group of mice had up-regulated mRNA of *Pgc1 $\alpha$*  and its target genes in the BC muscle ( $p < 0.05$ ). In addition, the mRNA expressions of *Pgc1 $\alpha$* , *Ucp2* and *Ucp3* were also higher in the LA muscle of the Cast + BGP group ( $p < 0.05$ ). Furthermore, the expression of *Pgc1 $\alpha$* , *Ucp2* or *Ucp3* in the hindlimb muscles of the BGP group was also higher than those of the Cast group ( $p < 0.05$ ).

We next examined the distribution and intensity of SDH, a marker of oxidative capacity, using SDH staining, in TA, GAS and TRI muscle sections. As shown in Fig. 4, the muscle fibers in the Sham group stained more darkly than those of the Cast group. Compared to the Sham group, the number of SDH-positive fibers of the Cast group decreased 31% ( $p < 0.05$ ) and 12% in TA and GAS muscle, respectively. On the other hand, the number of SDH-positive fibers in the TA and GAS of the Cast + BGP group was comparable to those of the Sham and Cast + TP groups ( $p > 0.05$ ) (Fig. 4). In contrast, there was no difference in percent SDH-positive fibers in TRI muscle among groups, which is relatively less responsive to androgen deprivation.



**Fig. 3** The mRNA expression levels of genes related to mitochondrial functions in various muscles of sham or castrated mice supplemented with BGP or TP. The mRNA of genes related to mitochondrial biogenesis, morphology, oxidative phosphorylation and uncoupling in highly androgen-responsive levator ani (A) and bulbocavernosus muscles (B) or the hind limb skeletal muscles gastrocnemius (C) and soleus (D) were analyzed. Values are means  $\pm$  SD. Data are transformed into log or reciprocal if they are not normally distributed before the statistical analysis. The significance of differences were analyzed by one-way ANOVA and Duncan's multiple range test. Values not sharing the same superscript letters are significantly different at  $p < 0.05$ . In some cases, a two-sample Student's *t* test was further performed for the comparison with the Cast group. \* denotes significant difference from the Cast group ( $p < 0.05$ ).

### 3.5 BGP did not change the fiber size distribution and most of the myogenesis-related gene expression levels in castrated mice

As shown in Fig. 5, the mRNA expressions of myogenic factor 6 (*Myf6*), insulin-like growth factor-1 (*Igf1*), eukaryotic translation initiation factor 3, subunit F (*eIF3f*) and myostatin (*Mstn*) in the BC muscle of the Cast group were lower than those of the Sham group ( $p < 0.05$ ). The BGP + Cast group had significantly higher mRNA expressions of *Mf6* and *Igf1* in the BC muscle than the Cast group ( $p < 0.05$ ).

Since it is believed that, at the single fiber level, the maximal force is directly related to its CSA, we also examined the distributions of the myofiber CSA of TA, GAS and TRI muscles. As shown in Fig. 6, the castrated mice tended to have a smaller CSA and their mean CSA was significantly lower than that of the Sham group ( $p < 0.05$ ). However, the size distribution and mean CSA of Cast + BGP and Cast + TP groups were comparable to those of the Cast group. Supplementation did not significantly change the CSA distribution in castrated mice (Fig. 6). The size distribution of these muscle fibers in castrated mice tended to shift toward the left when compared to the Sham group, and the BGP and TP supplementation did not restore the distribution of the fiber size to the control levels.

### 3.6 The mRNA expression of *Igf1*, *Ucp2* and *Pgc1a* positively correlated with the relative weight of androgen responsive muscles

The correlations between the muscle mass and mRNA expressions of several genes in the highly androgen responsive

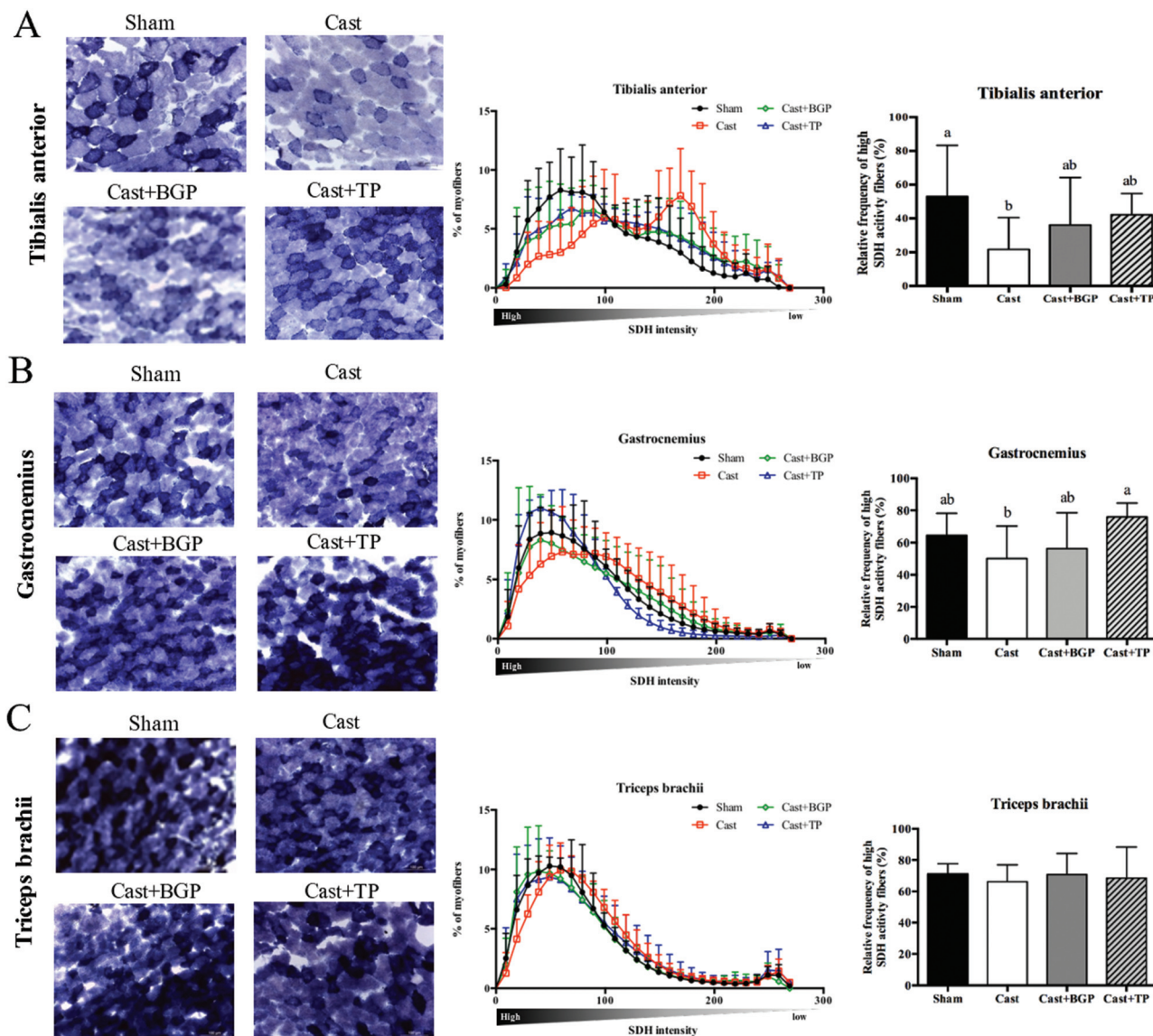
LA and BC muscles are shown in Table 2. Interestingly, the expressions of *Igf1*, *Ucp2* and *Pgc1a* mRNA levels were positively and significantly correlated with the mass in both muscles ( $p < 0.05$ ). In addition, the mRNA expressions of *Ucp3*, cytochrome c oxidase subunit VIIa 1 (*Cox7a1*) and cytochrome c oxidase subunit VIIIb (*Cox8b*) also positively correlated with the relative weight of the BC muscle. Remarkably, the mRNA expressions of most genes listed are positively and significantly correlated with each other in the BC muscle, implying an association in the expressions of these genes.

## 4 Discussion

The *in vivo* efficacy of a SARM is usually tested by the Hershberger assay, in which the weights of LA muscle, and prostate and seminal vesicles of treated castrated rats are compared to those of castrated and sham-operated rats.<sup>16</sup> Using a similar approach in a mouse model, this study provides evidence that BGP shows SARM-like properties. In addition, our cell-based AR transactivation assay further demonstrated that a BGP EA extract weakly modulated AR activation activities similar to those of some SARMS in development.<sup>40,41</sup>

In our animal study, data of the serum testosterone concentration, weights of androgen responsive tissues, including prostate/seminal vesicles, and LA and BC muscles, as well as grip strength and rotarod performance tests confirmed the androgen deficient status of the Cast mice. Supplementation of TP restored these parameters in a non-selective manner. In contrast,





**Fig. 4** Histochemical staining of succinate dehydrogenase activity in various limb muscles of sham or castrated mice supplemented with BGP or TP. SDH activity was observed from stained frozen sections of tibialis anterior (A), gastrocnemius (B) and triceps brachii (C) muscles. The staining intensity was analyzed with Image J and plotted as intensity distribution diagrams and high SDH activity fibers were quantified in TA, GAS and TRI muscles of each group. Values in the bar graph are means  $\pm$  SD. Data were analyzed by one-way ANOVA and Duncan's multiple range test. Values not sharing the same letter are significantly different at  $p < 0.05$ . Scale bar: 100  $\mu$ m.

supplementation of BGP to castrated mice did not affect the serum testosterone and seminal vesicle/prostate weights, but improved the androgen responsive LA and BC muscle weight as well as the grip strength and rotarod performance, suggesting a tissue selective androgen receptor modulation activity.

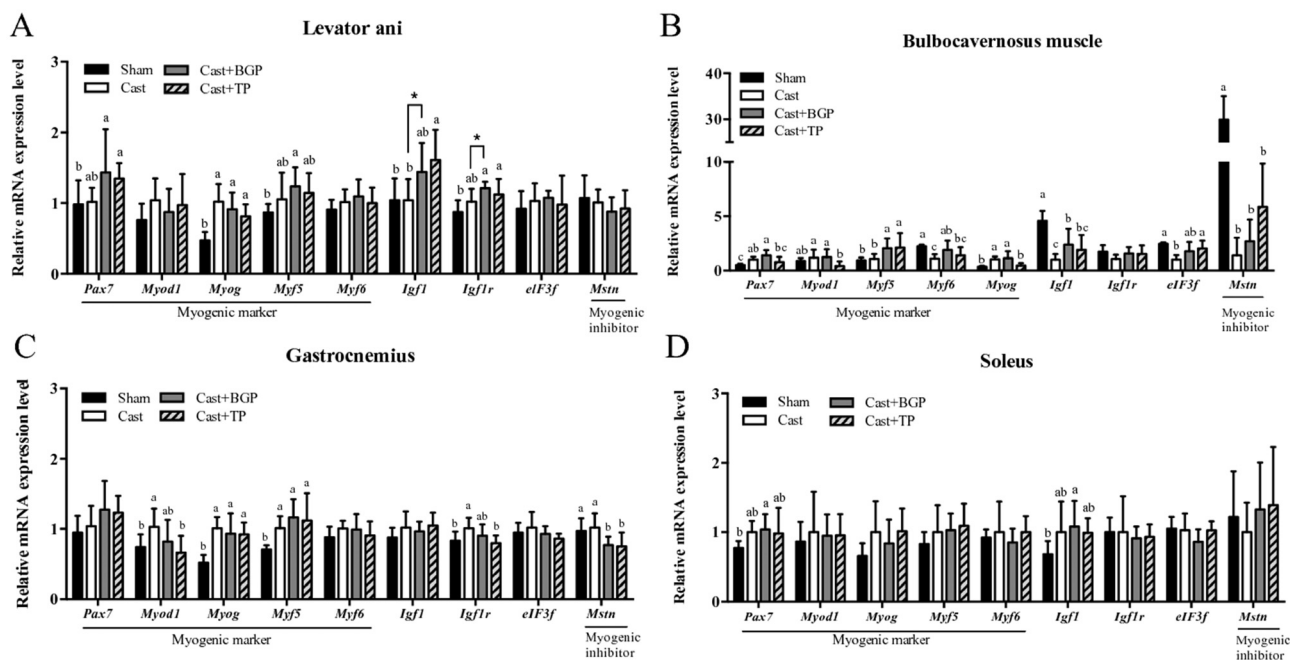
Although the SV/prostate and BC muscle weights of the Cast + TP group were significantly higher than those of the Cast group, these values were still significantly lower than those of the Sham group (Table 1). These differences between the Sham and Cast + TP groups coincide with the difference in the serum testosterone concentrations between these two groups (Fig. 2A). These results suggest that the androgen status of the Cast + TP group in this study was lower than the

normal physiological condition. Interestingly, the weight of the LA muscle of the Cast + TP group was comparable to that of the Sham group, suggesting that a lower androgen threshold is required to support this muscle.

Except for LA and BC, the remaining skeletal muscles weighed in this study hardly corresponded to the androgen status, agreeing with previous reports.<sup>6,38</sup> However, our results of the SDH staining (Fig. 4) showed that the oxidative capacity of TA and GAS muscles was affected by the androgen status. Results of the two functional tests, grip strength and rotarod performance, corresponded to the androgen status as well.

The CSA is a commonly used indicator of muscle fiber hypertrophy. The anabolic effects of testosterone have been





**Fig. 5** The mRNA expressions of genes related to myogenesis in various muscles of sham or castrated mice supplemented with BGP or TP. The mRNA expression levels of genes related to myogenesis in the highly androgen-responsive levator ani (A) and bulbocavernosus muscles (B) or the hind limb skeletal muscles gastrocnemius (C) and soleus (D) were analyzed. Values are means  $\pm$  SD. Data were transformed into log or reciprocal if they are not normally distributed before the statistical analysis. The significance of differences were analyzed by one-way ANOVA and Duncan's multiple range test. Values not sharing the same superscript letters are significantly different at  $p < 0.05$ . In some cases, a two-sample Student's *t* test was further performed for a comparison with the Cast group. \* denotes significant difference from the Cast group ( $p < 0.05$ ).

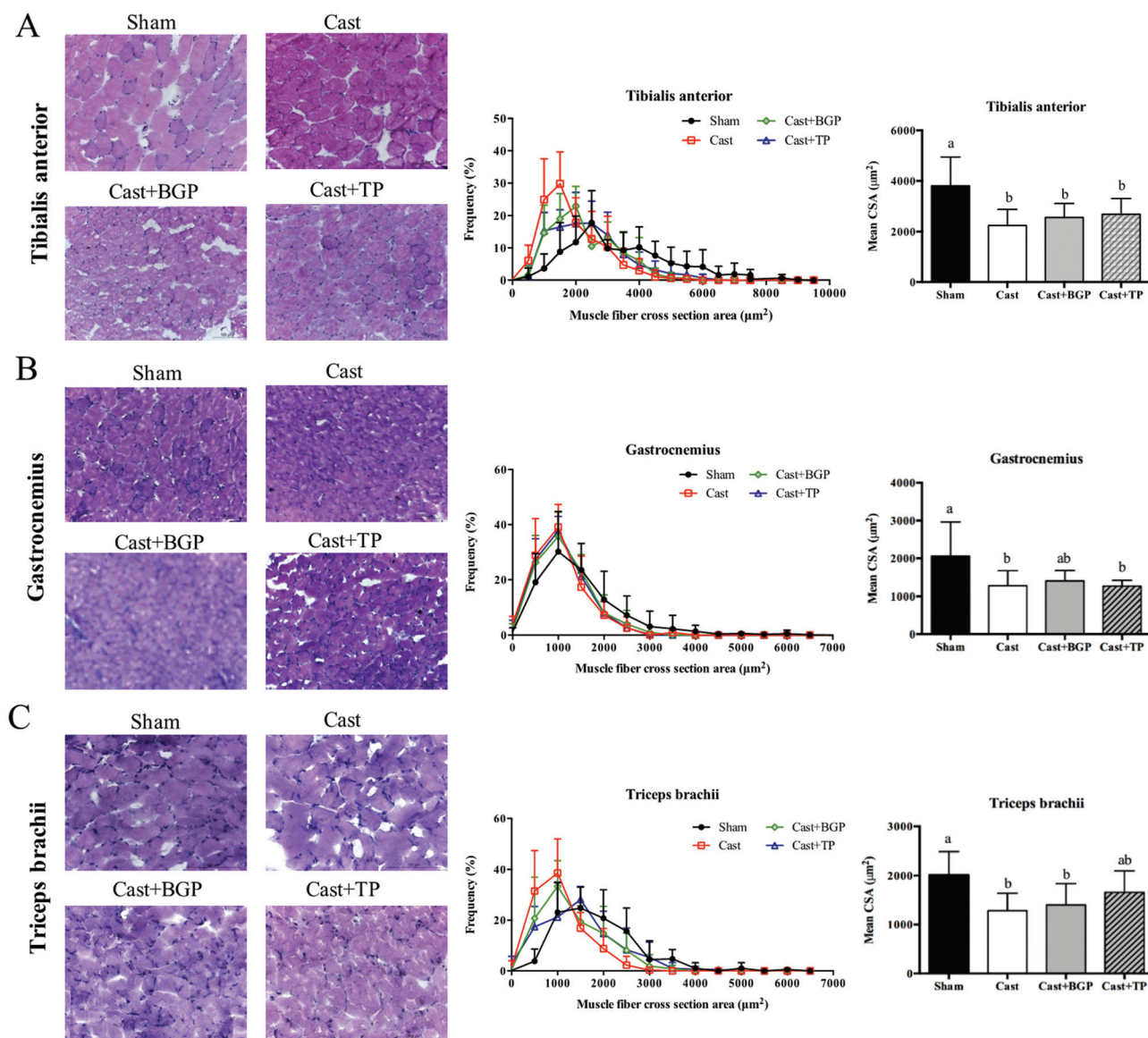
shown to increase the muscle weight and CSA in mice<sup>42</sup> or humans.<sup>43</sup> In this study, castrated mice showed a greater number of small size fibers in TA, GAS and TRI muscles as expected. However, the BGP and TP treatment did not change the mean CSA in castrated mice. Previous studies also showed that the anabolic effect of testosterone on muscle is dose dependent.<sup>44</sup> Serum testosterone levels equal to or higher than those of a young adult are required to show the increased volume of some muscles.<sup>44</sup> In addition, fiber-hypertrophy in mice was shown to be induced by supra-physiological testosterone supplementation for 8 weeks.<sup>45</sup> The TP and BGP treatment in this study did not change the mean CSA in the castrated mice which might be related to the lower-than normal physiological androgen status.

Among the 4 muscles measured, the mRNA expressions of genes related to mitochondria of the BC muscle corresponded most pronouncedly to the androgen status, especially those of oxidative phosphorylation and *Ucp2* (Fig. 2). These results agree with the data of BC weight which was supported by the positive and significant correlations of BC weight and mRNA expressions of *Ucp2*, *Ucp3*, *Pgc1 $\alpha$* , *Cox7a*, and *Cox8b* (Table 2), again, suggesting that this androgen responsive muscle tissue might be more sensitive to the androgen status. In this regard, BGP supplementation significantly elevated the BC weight of castrated mice, but the BC weight of the Cast + BGP group was still significantly lower than that of the Cast + TP group. It is thus speculated that the androgen status of the Cast + BGP

group might be even lower than that of the Cast + TP group. The weak agonistic activity of the BG extract in the AR transactivation assay also supports this speculation. This may account for the insignificant differences between the performance in the functional tests and SDH staining of TA and GAS muscles between the Cast + BGP group and the Cast group. Nevertheless, there were also no significant differences of these parameters between the Cast + BGP and Cast + TP groups, suggesting that BGP showed a trend of effects similar to TP.

Testosterone up-regulates *Igf1* gene expression through the androgen response elements (AREs) within the upstream promoter region of the IGF1 gene.<sup>46</sup> Interestingly, the *Igf1* mRNA expression level in the BC muscle corresponded to the androgen status in this study (Fig. 5B) and was positively correlated with the BC weight (Table 2). The BGP supplementation also significantly up-regulated *Igf1* mRNA expression to a level comparable to TP supplementation. Similarly, the mRNA of *Ucp3* expression in the LA, BC and SOL muscles, and *Ucp2* in the LA, BC and GAS muscles, corresponded to the androgen status as well. The BGP supplementation also up-regulated *Ucp3* and/or *Ucp2* expressions in these muscles, to levels comparable to those of the Sham and Cast + TP groups, except *Ucp2* in the BC which was significantly lower than that of the Cast + TP group.

PGC1 $\alpha$  is well recognized for its role in mitochondrial biogenesis and respiration through an induction of nuclear respiratory factor NRF1, NRF2 and UCP2 gene expressions<sup>47,48</sup>



**Fig. 6** The cross-section area (CSA) of skeletal muscle fibers in various limb muscles of sham or castrated mice supplemented with BGP or TP sections from dry ice-cooled isopentane frozen tibialis anterior (A), gastrocnemius (B) and triceps brachii (C) muscles were fixed in 4% paraformaldehyde in PBS and stained with hematoxylin and eosin. Histogram distribution of the myofiber cross-section area (CSA) (A, B and C) and the average CSA (D) was analyzed with Image J. Values are means  $\pm$  SD. Data were analyzed by one-way ANOVA and Duncan's multiple range test. Values not sharing the same letter are significantly different at  $p < 0.05$ . Scale bar: 100  $\mu\text{m}$ .

and has been demonstrated to spare muscle mass and improve endurance in atrophic muscles.<sup>49</sup> Skeletal muscles PGC1 $\alpha$  and PPAR $\beta/\delta$  have been proposed to be key regulators of exercise performance. Overexpression of PGC1 $\alpha$  specifically in skeletal muscles increased the muscle oxidative capacity and the proportions of type I and IIa fibers in transgenic mice.<sup>50,51</sup> In contrast, the muscle-specific knock-out of the *Pgc1 $\alpha$*  gene in mice exhibited a reduction in the exercise performance and muscle respiratory capacity compared to those of wild type control mice.<sup>52</sup> In addition, testosterone replacement up-regulated *Pgc1 $\alpha$*  and *Ucp3* expression in the TRI muscle of castrated mice.<sup>53</sup> In this study, castration lowered the mRNA expressions of *Pgc1 $\alpha$*  in BC, GAS and SOL muscles.

The BGP supplementation increased *Pgc1 $\alpha$*  mRNA in all these four muscles. In contrast, TP supplementation only restored these to some extent in GAS and SOL, but not in BC muscles. The up-regulation of *Pgc1 $\alpha$*  mRNA in muscles of the Cast + BGP group might thus be, at least in part, from mechanisms other than the androgen receptor.

Emerging evidence suggests that UCP2 plays a positive physiological role by regulating mitochondrial biogenesis, substrate utilization and ROS elimination,<sup>54</sup> thereby providing possibly anti-ageing effects.<sup>55</sup> We also observed up-regulations of the *Ucp2* mRNA in LA, BC and GAS muscles of the BGP and TP treated castrated mice. Overexpression of *Pgc1 $\alpha$*  in C2C12 myotubes has been shown to up-regulate UCP2

**Table 2** The correlation coefficients (*r*) among the relative weight of LA and BC muscles and the level of mRNA expression

	Relative weight (% of BW)	<i>Ucp2</i>	<i>Ucp3</i>	<i>Pgc1a</i>	<i>Cox7a1</i>	<i>Cox8b</i>	<i>Igf1</i>	<i>Igf1r</i>
<b>LA muscle</b>								
Relative weight (% of BW)	1							
<i>Ucp2</i>	0.437*	1						
<i>Ucp3</i>	0.189	0.327	1					
<i>Pgc1a</i>	0.378*	0.454*	0.309	1				
<i>Cox7a1</i>	-0.267	0.120	0.477**	0.061	1			
<i>Cox8b</i>	-0.434*	-0.152	0.324	0.243	0.574**	1		
<i>Igf1</i>	0.516**	0.095	-0.079	0.304	-0.320	-0.26	1	
<i>Igf1r</i>	-0.110	0.126	0.309	0.407*	0.129	0.444*	0.182	1
<b>BC muscle</b>								
Relative weight (% of BW)	1							
<i>Ucp2</i>	0.889**	1						
<i>Ucp3</i>	0.608**	0.597**	1					
<i>Pgc1a</i>	0.494*	0.419*	0.730**	1				
<i>Cox7a1</i>	0.836**	0.773**	0.663**	0.451*	1			
<i>Cox8b</i>	0.689**	0.668**	0.620**	0.441*	0.883**	1		
<i>Igf1</i>	0.818**	0.688**	0.617**	0.595**	0.826**	0.770**	1	
<i>Igf1r</i>	0.345	0.528**	0.683**	0.561**	0.356	0.355	0.317	1

Data were analyzed by Pearson's correlation. \* denotes significant correlation,  $p < 0.05$ . \*\* denotes significant correlation,  $p < 0.01$ . LA: levator ani muscle, BC: bulbocavernosus muscle.

expression.<sup>48</sup> But the pattern of UCP2 expression in response to various treatments in this study were not completely parallel to that of the *Pgc1a* expression in all 4 muscles tested, suggesting that other mechanisms might be involved. Computational BLAST was thus conducted to preliminarily search for possible AREs in the promoter region of the mouse UCP2 gene (ESI Table 3†). Further studies are required to explore whether these putative ARE sites are functional.

The fiber size and oxidative capacity of muscles are known to be determined from the balance among myofibrillar protein synthesis, mitochondrial biosynthesis and mitochondrial degradation.<sup>56</sup> The inverse relationship between the striated muscle fiber size and its oxidative capacity, the so-called “fiber type-fiber size paradox”, is interpreted in part from the competition between signaling pathways for the synthesis of myofibrillar proteins and proteins associated with oxidative metabolism, such as increased mitochondrial biogenesis *via* AMP-activated protein kinase (AMPK).<sup>56</sup> AMPK controls several processes *via* the transcriptional co-activator PGC1 $\alpha$  and promotes the slow, oxidative phenotype in skeletal muscles.<sup>57</sup> Previous studies have clarified another role of AMPK in inhibiting protein synthesis and opposing actions of mTOR and AMPK on muscle homeostasis.<sup>58</sup> For example, AMPK activation suppresses the translation of myofibrillar proteins by inhibiting the mTOR-p70S6K pathway *via* activation of TSC-2 and by phosphorylation and inactivation of eEF-2. This interaction indicates that AMPK, in addition to being a stimulus of biosynthesis of mitochondria, is also an inhibitor of protein synthesis.<sup>59,60</sup> These findings showed that mTOR and AMPK are cross-balancing pathways under physiologic conditions. In this regard, some cucurbitane-type triterpenoids isolated from BG were shown to activate AMPK in adipocytes and myocytes.<sup>61,62</sup> Furthermore, AMPK can increase gene expressions of mitochondria biogenesis through directly phosphorylating the

PGC1 $\alpha$  protein. The phosphorylated PGC1 $\alpha$  protein is required for the PGC1 $\alpha$  dependent induction of the PGC1 $\alpha$  promoter.<sup>63</sup> We previously showed that *Pgc1a* mRNA is up-regulated in muscles of mice fed the BGP diet.<sup>29</sup> It is therefore speculated that the BGP supplementation might improve the oxidative capacity of muscle fibers through the activation of the AMPK pathway, but these fibers remained in relatively small sizes.

We have identified various biologically active compounds from the BG. Those that can activate PPAR $\alpha$  and  $\gamma$  include 9c,11t,13t-CLN,<sup>64</sup> some cucurbitane-type triterpenoids,<sup>29</sup> phytol and lutein.<sup>65</sup> Some of the cucurbitane-type triterpenoids have also been shown to activate estrogen receptors<sup>32</sup> and stimulate glucagon-like peptide 1 secretion in an enteroendocrine cell line.<sup>66</sup> Some cucurbitane-type triterpenoids have been shown to activate AMPK.<sup>61,62</sup> Therefore, multiple compounds in this common tropical vegetable may act on various molecular targets and lead to beneficial health effects. In this study, we tested 9c,11t,13t-CLN and its metabolite in rodents, 9c,11t-CLA<sup>35-37</sup> and its isomer 10t-12c CLA,<sup>67</sup> along with the BGP extract, in the AR transactivation assay. The weak agonistic and antagonistic activity of the BGP extract can also be observed in these conjugated fatty acids. These data imply that 9c,11t,13t-CLN and its metabolite 9c,11t-CLA may be one of the bioactive compounds in BGP. The possibility of the contribution of other active compounds cannot be excluded.

Bitter melon is a common vegetable in the tropical area. In our previous study, ~5 g day<sup>-1</sup> freeze-dried bitter melon powder was supplemented to patients with the metabolic syndrome for 3 months in a preliminary clinical trial. This level is equivalent to 1% (wt/wt) BGP in the diet for mice study, as the common Taiwanese diet has been estimated to be 450–500 g d<sup>-1</sup> in the dry weight basis. Results of this trial demonstrated not only the safety of the supplementation, but also the activity of BG in ameliorating the metabolic syndrome.<sup>68</sup> On the other



hand, ~5 g of BGP is equivalent to ~100 g of fresh bitter melon which is a serving of vegetable. Dietary guidelines recommended at least 3 servings of vegetable per day.

In conclusion, this study demonstrated that BGP supplementation may ameliorate castration-induced decline of grip strength, exercise performance, and muscle mass and up-regulate *Pgc1α*, *Ucp2* and mitochondrial biogenic genes in some muscles, but did not increase the mass of the prostate in orchidectomized mice. *In vitro*, the BGP extract weakly modulated the AR transactivation activity.

## Abbreviations

5α-DHT	5α-Dihydrotestosterone
AR	Androgen receptor
BC	Bulbocavernosus
BGP	Bitter melon powder
Cast group	Castrated mice fed a high sucrose diet
Cast + BGP group	Castrated mice fed a high sucrose diet supplemented with 5% lyophilized BGP
Cast + TP group	Castrated mice fed a high sucrose diet supplemented with TP
CSA	Cross-section area
GAS	Gastrocnemius
LA	Levator ani
PGC1α	Peroxisome proliferator-activated receptor gamma coactivator 1 alpha
SARM	Selective androgen receptor modulator
SDH	Succinate dehydrogenase
Sham group	Sham-operated mice fed the high sucrose diet in which carbohydrate was provided by 50% sucrose and 12.95% cornstarch
SOL	Soleus
SV	Seminal vesicle
TA	Tibialis anterior
TP	Testosterone propionate
TRI	Triceps brachii
UCP2/3	Uncoupling protein 2/3

## Conflicts of interest

There are no conflicts of interest to declare.

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