Pyrroloquinoline quinone attenuates cachexia-induced muscle atrophy via suppression of reactive oxygen species

Tongtong Xu^{1,2#}, Xiaoming Yang^{1#}, Changyue Wu², Jiaying Qiu¹, Qingqing Fang¹, Lingbin Wang¹, Shu Yu¹, Hualin Sun¹

¹Laboratory of Neuroregeneration, Jiangsu Clinical Medicine Center of Tissue Engineering and Nerve Injury Repair, Co-Innovation Center of Neuroregeneration, ²School of Medicine, Nantong University, Nantong 226001, China

Contributions: (I) Conception and design: H Sun; (II) Administrative support: H Sun, X Yang; (III) Provision of study materials or patients: T Xu, X Yang, C Wu, J Qiu, Q Fang, L Wang, S Yu; (IV) Collection and assembly of data: T Xu, X Yang, C Wu; (V) Data analysis and interpretation: T Xu, X Yang, C Wu, S Yu; (VI) Manuscript writing: All authors; (VII) Final approval of manuscript: All authors.

[#]These authors contributed equally to this work.

Correspondence to: Dr. Hualin Sun. 19 Qixiu Road, Jiangsu Key Laboratory of Neuroregeneration, Nantong University, Nantong 226001, China. Email: sunhl@ntu.edu.cn.

Background: Cachexia, a wasting syndrome, is most commonly observed in individuals with advanced cancer including lung cancer, esophageal cancer, liver cancer, etc. The characteristic sign of cachexia is muscle atrophy. To date, effective countermeasures have been still deficiency to alleviate muscle atrophy. Reactive oxygen species (ROS) are important regulators of muscle atrophy. Therefore, the effects of a naturally antioxidant, pyrroloquinoline quinone (PQQ), were explored on muscle atrophy induced by cachexia in the present study.

Methods: Tumor necrosis factor- α (TNF- α) induced C2C12 myotubes atrophy model was constructed. The atrophied C2C12 myotubes were dealt with the presence or absence of N-acetyl-L-cysteine (NAC, an antioxidant for ROS abolition) (5 mM) or PQQ (80 μ M) for 24 hours. ROS content was determined by dichlorodihydrofluorescein diacetate (DCFH-DA) staining. The diameter of myotubes was analyzed by myosin heavy chain (MHC) staining. The protein levels of MHC, muscle atrophy F-box (MAFbx) and muscle RING finger-1 (MuRF-1) in each group were observed by Western blotting.

Results: First, ROS generation was enhanced in C2C12 myotubes treated with TNF- α . NAC treatments significantly avoided the reduction in the diameter of C2C12 myotubes, and concomitantly increased MHC levels, and decreased ROS contents, MuRF1 and MAFbx levels. These data suggested that the increased ROS induced by TNF- α might play a central role in muscle wasting. PQQ (a naturally occurring antioxidant) administration inhibited C2C12 myotubes atrophy induced by TNF- α , as evidenced by the increase of the diameter of C2C12 myotubes, together with increased MHC levels and decreased ROS, MAFbx and MuRF-1 levels.

Conclusions: PQQ resists atrophic effect dependent on, at least in part, decreased ROS in skeletal muscle treated with TNF- α .

Keywords: Pyrroloquinoline quinone (PQQ); muscle atrophy; oxidative stress

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Introduction

Cachexia, a wasting syndrome, is most commonly observed in the majority of patients with advanced cancer including lung cancer, liver cancer, gastric cancer, renal cancer, etc. (1-3).

The hallmark of cancer associated cachexia is muscle wasting (4). Cachexia is characterized by inexorable muscle atrophy that largely affects patient prognosis and contributes significantly to cancer morbidity and mortality (3,5). To date, effective countermeasures have been still deficiency to alleviate muscle wasting. Inflammatory mediators [tumor necrosis factor- α (TNF- α), interleukin-1 and interleukin-6] are key contributors to the development of cancer cachexia, which is the driving forces of skeletal muscle atrophy caused by cancer cachexia (6,7). TNF- α was often used to induce a model of muscle atrophy in cancer cachexia (8-10). Therefore, to elucidate the molecular mechanism and develop targeted therapies of cachexia associated muscle atrophy is of significant importance.

The balance between protein synthesis and degradation is crucial to preserve muscle healthy. Skeletal muscle atrophy occurs when protein degradation exceeds protein synthesis (11). Skeletal muscle atrophy is closely related to a reduced myofibrillar protein which results from both increased proteolysis and decreased protein synthesis (2). The ubiquitin-proteasome pathway (UPS) and the autophagy-lysosomal pathway (ALP) are two main intracellular proteolysis systems (12,13). The UPS is well recognized to be involved in mediating muscle atrophy, and is a key factor of muscle atrophy under various physiopathological situations (14,15). muscle RING finger-1 (MuRF1) and muscle atrophy F-box (MAFbx) are two muscle-specific E3 ubiquitin ligases of the UPS (16). Our previous study has shown that MuRF1 and MAFbx displayed significant up-regulation during skeletal muscle atrophy under various pathophysiological conditions (17-20). Likewise, MuRF1 and MAFbx displayed an upregulation in muscle atrophy caused by cancer cachexia and cerebral ischemia (16,21,22).

Reactive oxygen species (ROS) generation is at relatively low level in skeletal muscle under physiological conditions (23). While elevated ROS led to increased protein degradation through increasing the expression of key atrophic factors such as MuRF1 and MAFbx (24). Our previous study had also shown that the gene expression of the positive regulators for ROS production were up-regulated and the negative regulators for ROS production were down-regulated in denervation induced muscle atrophy (20). Although growing evidence has suggested that high levels of ROS might play a crucial role in muscle wasting, and effective targeted therapeutic strategies are still deficient for muscle atrophy. The therapeutic strategy of skeletal muscle wasting with natural compounds has recently received more and more attention. Pyrroloquinoline quinone (PQQ) is a naturally occurring redox cofactor, and it is known to protect cells from oxidative stress (25). Nonetheless, the effects of PQQ on skeletal muscle atrophy caused by cachexia have not yet been clearly addressed. Therefore, the aim of this study is to determine the protective effect of PQQ on muscle atrophy.

In this work, we first detected ROS production and analyzed whether ROS were central for cachexiainduced muscle atrophy. Furthermore, the effects of PQQ administration on cachexia-induced skeletal muscle atrophy were estimated. PQQ administration could resist muscle atrophy, as evidenced by increased muscle fiber crosssectional area (CSA) or diameter of myotubes, together with increased myosin heavy chain (MHC) levels and decreased ROS, MAFbx and MuRF-1 levels.

Methods

Cell culture and treatments

Mouse skeletal muscle C2C12 cells were purchased from the ATCC (American Type Culture Collection), and were cultured in high-glucose DMEM (Dulbecco's modified Eagle's medium) containing with 10% fetal bovine serum (FBS) at 37 °C in a humidified atmosphere with 5% CO₂. Mouse skeletal muscle C2C12 cells were differentiated in C2C12 differentiation medium containing DMEM, antibiotics and 2% horse serum (Invitrogen, Waltham, MA, USA) for 7 days (19). For antioxidant treatment, 10 ng/mL recombinant mouse TNF- α (R&D Systems, USA) was added and incubated with the differentiated C2C12 myotubes in the presence or absence of N-acetyl-L-cysteine (NAC) (5 mM) or PQQ (Sigma-Aldrich, 80 µM) for 24 hours. Finally, the C2C12 myotubes were examined by morphometric or biochemical assays.

Western blotting analysis

Western blotting analysis was carried out as described previously (26). For the protein extraction, C2C12 myotubes were homogenized in the radio immunoprecipitation assay buffer and the extraction was quantified with Bio-Rad Protein Assay Kit (Bio-Rad, Hercules, CA, USA). Then, total proteins were subjected to SDS-PAGE (polyacrylamide gel electrophoresis) and transferred onto polyvinylidene difluoride (PVDF) membranes (Millipore, USA). Following, the PVDF membranes were blocked with 5% nonfat dry milk diluted in Tris-buffer saline (TBS), and incubated with primary antibodies: mouse anti-MHC (1:3,000) (R&D Systems, USA), rabbit anti-MAFbx (1:3,000) (Abcam, USA), rabbit anti-MuRF-1 (1:3,000) and mouse anti-tubulin (1:3,000) (Santa Cruz, CA, USA) at 4 °C overnight. Next morning, the PVDF membranes were incubated with the appropriate secondary antibody. The immunoreactions were performed by enhanced chemiluminescence (Thermo Scientific, USA) and the intensity values were obtained for further normalization against the internal reference.

Myotube diameter measurement and quantification

The diameter of myotubes was analyzed by MHC staining. Briefly, C2C12 cells were seeded and differentiated on the glass coverslips. Then C2C12 myotubes were fixed and incubated with mouse anti-MHC (1:3,000) (R&D Systems, USA) for 1 hour. When the antibody incubation fluid was removed, the slips were incubated with affinity-purified Alexa Fluor dye-conjugated goat anti-mouse antibody (1:100) (Life Technologies, USA) for 30 min. The slips were photographed with the fluorescence microscopy. The diameter of myotubes was evaluated by the Image J software (27).

Determination of ROS

The ROS content is measured by using dichlorodihydrofluorescein diacetate (DCFH-DA) staining (Sigma-Aldrich, USA). DCFH-DA is widely used techniques for directly measuring the ROS content (28). When DCFH-DA is oxidized by ROS, it would be changed into the fluorescent compound dichlorofluorescein (DCF). Briefly, C2C12 myotubes were washed with PBS and fresh DMEM without phenol red, and incubated with 10 µM of ROS sensitive dve DCFH-DA for 30 min in the dark at room temperature. Then, the C2C12 myotubes stained with DCFH-DA were immediately analyzed, and the ROS content was assessed by the increase in the fluorescence intensity of DCF. The fluorescence intensity of DCF was measured at an excitation wavelength of 488 nm and an emission wavelength of 519 nm. Mean fluorescence intensities were obtained by histogram statistics.

Statistical analysis

All data were expressed as mean \pm SEM from at least three independent experiments. Statistical methods including Student's *t*-test and one-way ANOVA were used to analyze the differences between groups. All statistical analyses were performed with the SPSS statistical software version 17.0 (SPSS Inc. Chicago, IL, USA). P values <0.05 were considered statistically significant.

Results

ROS generation is enhanced in TNF-a induced C2C12 myotubes atrophy

The ROS content was measured by using DCFH-DA staining in C2C12 myotubes in the presence or absence of TNF- α treatment. Results showed that the relative fluorescence intensity in C2C12 myotubes exposed to TNF- α was higher than that in the C2C12 myotubes without TNF- α treatment (P<0.05) (*Figure 1A*), which indicated that the ROS generation was enhanced in TNF- α induced C2C12 myotubes atrophy, and also demonstrated that ROS generation might play a central role in TNF- α induced C2C12 myotubes atrophy.

ROS are crucial for skeletal muscle atrophy induced by TNF-a

Next, the effect of increased ROS on the atrophic response induced by cachexia was determined by incubating myotubes with NAC, an antioxidant for ROS abolition, and assessing the diameter of myotubes. NAC treatments significantly relieved the decrease in the diameter of myotubes (*Figure 1B*,*C*,*D*,*E*) and the increase in ROS content (*Figure 1F*). Following, the influences of NAC on the MHC, E3-ubiquitin ligases MAFbx and MuRF-1 levels were examined by Western blotting. Treatments with 5 mM of NAC avoided the reduction of the expression of MHC, and also inhibited the increase of MAFbx and MuRF-1 levels (*Figure 2*). These results suggested that the increased ROS caused by TNF- α might play a critical role in muscle wasting.

PQQ prevents ROS generation and muscle atrophy induced by TNF-a

In order to explore the effects of PQQ, a natural antioxidant, on the muscle atrophic effect induced by



Figure 1 NAC treatments significantly alleviated the reduction in the diameter of C2C12 myotubes induced by TNF- α through inhibiting ROS. (A) TNF- α induces increased ROS levels in C2C12 myotubes; (B-E) NAC treatments significantly alleviated the reduction in the diameter of C2C12 myotubes. C2C12 myotubes were incubated with TNF- α in the presence or absence of NAC (5 mM) for 24 h. Then, immunodetection was performed to determine the presence of MHC in C2C12 myotubes, and the value of diameter of C2C12 myotubes was expressed as the mean ± SEM from three independent experiments; (F) NAC could inhibit ROS caused by TNF- α treatment. DCF fluorescence intensities in C2C12 myotubes suffered from TNF- α treatment. NC: C2C12 myotubes without any treatment; TNF- α : C2C12 myotubes treated with TNF- α and NAC simultaneously. *, P<0.05 versus NC; *, P<0.05 versus TNF- α . ROS, reactive oxygen species; NAC, N-acetyl-L-cysteine; MHC, myosin heavy chain; DCF, dichlorofluorescein.

TNF- α , muscle atrophy and ROS generation were analyzed by incubating myotubes with PQQ, and evaluating the diameter of C2C12 myotubes and the ROS generation. The diameter of C2C12 myotubes was determined through MHC staining and quantified (*Figure 3A*, *B*, *C*, *D*). The results showed that PQQ treatments significantly alleviated the reduction in the diameter of C2C12 myotubes and the increase of ROS levels induced by TNF- α (*Figure 3E*).

The expression level of MAFbx, MuRF-1 and MHC, which are dysregulated during muscle atrophy, were assessed by Western blotting. The expression level of MHC displayed remarkable decrease, and MAFbx and MuRF-1 displayed dramatically increase in C2C12 myotubes treated with TNF- α . PQQ treatments significantly prevented the decrease of MHC and the increase of MAFbx and MuRF-1 levels (*Figure 4*).

Discussion

Cachexia, a devastating, multifactorial and often irreversible

syndrome, is most commonly observed in individuals with advanced cancer (1,2,29). Cachexia leads to substantial weight loss, primarily from loss of skeletal muscle and body fat. The characteristic sign of cachexia is muscle wasting, driven by inflammatory response, which significantly affects patient prognosis and increases mortality (5,30). To date, effective countermeasures have been still deficiency to alleviate muscle atrophy. It is well-known that inflammatory cytokines such as IL-1 beta, TNF-a, IL-6 and interferongamma have been involved in cancer cachectic response (31). Inflammatory mediators are key contributors to cancer cachexia, and they are the driver of skeletal muscle atrophy in cancer cachexia (6,32,33). Therefore, blocking inflammatory response was considered to be an effective treatment to prevent skeletal muscle atrophy under cachectic conditions. Cachexia is more common in malignant tumor, such as non-small cell cancer (34). TNF- α displayed an elevated expression level in sarcopenia and cancer cachexia and the elevated TNF- α is remarkably correlated with muscle wasting (35). Tsubouchi et al.



Figure 2 NAC alleviated the decrease of MHC levels and the increase of MuRF-1 and MAFbx levels caused by TNF- α treatment. (A) MHC, MuRF-1 and MAFbx expression in C2C12 myotubes were determined by Western blotting; (B) the relative expression of MHC, MuRF-1 and MAFbx in C2C12 myotubes. NC: C2C12 myotubes without any treatment; TNF- α : C2C12 myotubes treated with TNF- α only; TNF- α + NAC: C2C12 myotubes treated with TNF- α and NAC simultaneously. *, P<0.05 versus NC; [#], P<0.05 versus TNF- α . NAC, N-acetyl-L-cysteine; MHC, myosin heavy chain; MAFbx, muscle atrophy F-box; MuRF1, muscle RING finger-1.



Figure 3 PQQ treatments significantly avoided the reduction in the diameter of C2C12 myotubes induced by TNF- α through inhibiting ROS. (A-D) C2C12 myotubes were incubated with TNF- α for 24 h in the presence or absence of PQQ (80 µM). Then, immunodetection was performed to investigate the presence of MHC in C2C12 myotubes, and the value of the diameter of C2C12 myotubes was expressed as the mean ± SEM from three independent experiments; (E) PQQ could inhibit ROS caused by TNF- α treatment. DCF fluorescence intensities in C2C12 myotubes suffered from TNF- α treatment; NC: C2C12 myotubes without any treatment; TNF- α : C2C12 myotubes treated with TNF- α and PQQ simultaneously. *, P<0.05 versus NC; [#], P<0.05 versus TNF- α . PQQ, pyrroloquinoline quinone; ROS, reactive oxygen species; DCF, dichlorofluorescein.

suggested that inhibiting the inductions of TNF- α could ameliorate body weight loss in the mouse model of lung adenocarcinoma (36). Now, TNF- α is often used to induce cachexia associated C2C12 myotubes atrophy model (8-10). Skeletal muscle atrophy is closely related to a reduced myofibrillar protein which results from both increased proteolysis and decreased protein synthesis, as evidenced by the reduced muscle fiber CSA and diameter (2). Normally,



Figure 4 PQQ alleviated the decrease of MHC levels and the increase of MuRF-1 and MAFbx levels caused by TNF- α treatment. (A) MHC, MuRF-1 and MAFbx expression in C2C12 myotubes were determined by Western blotting; (B) the relative expression of MHC, MuRF-1 and MAFbx in C2C12 myotubes. NC: C2C12 myotubes without any treatment; TNF- α : C2C12 myotubes treated with TNF- α only; TNF- α + PQQ: C2C12 myotubes treated with TNF- α and PQQ simultaneously. *, P<0.05 versus NC; [#], P<0.05 versus TNF- α . PQQ, pyrroloquinoline quinone; MHC, myosin heavy chain; MAFbx, muscle atrophy F-box.

ROS content was at relatively low levels in skeletal muscle fibers under physiological conditions (23). Whereas, the increased ROS caused the increase in protein breakdown through upregulating the atrophy related factors such as MuRF1 and MAFbx (23,24). Although growing evidence has suggested that high levels of ROS might play a crucial role in muscle wasting, effective targeting therapies are still deficient for muscle atrophy.

Regarding this, the current study was the first to report that ROS generation was enhanced in atrophied skeletal muscle induced by cachexia, and increased ROS might play an important role in skeletal muscle atrophy induced by cachexia. Then, the function of increased ROS on the atrophic response was examined. NAC, free radical scavenger, was incubated with C2C12 myotubes treated with TNF-α. These results showed that NAC treatments significantly inhibited the reduction in the diameter of myotubes, as well as the increase in ROS production. Concomitantly, the decrease of the expression level of MHC and the increase of MAFbx and MuRF-1 levels were attenuated. These results were consistent with the previous reports that free radical scavenger NAC efficiently decreased the ROS levels and restored the expression level of MHC (27,37). Our data suggested that the increased ROS caused by cachexia played a central role in skeletal muscle atrophy.

PQQ is a water-soluble naturally occurring antioxidant component that has a strong anti-oxidant capacity. It not only serves as a mediator participating in redox reactions in the mitochondrial respiratory chain but also serves as a critical scavenger participating in scavenging ROS (38).

Pretreated rats with PQQ could observably decrease ROS generation after ICH, which would be probably due to its antioxidant properties (39). PQQ could also significantly enhance the activities of superoxide dismutase, catalase and glutathione peroxidase (40). Therefore, in this study, we elucidated the potential of PQQ to ameliorate the muscle atrophy caused by cachexia. Our data demonstrated that the protective effect of PQQ on muscle atrophy was similar to that of NAC. PQQ treatments significantly alleviated the reduction in the diameter of myotubes and the increase of ROS content caused by cachexia, as well as preventing the decrease of MHC levels and the increase of MAFbx and MuRF-1 levels. Previous study indicated that PQQ could activate PGC-1a (41). PGC-1a expression could be useful to preserve muscle mass in functional denervation or aging (42). Overexpression of PGC-1a not only could block the TWEAK-induced atrophy program but also decrease the expression level of Fn14 in the skeletal muscle suffered from denervation (42). At the same time, TWEAK may activate the nuclear factor-kB (NF-kB) signaling pathway and the proteolytic pathways such as UPS and the ALP to induce muscle proteolysis (43). According to these, we might speculate that PQQ prevented the increase of MAFbx and MuRF-1 levels to relieve muscle atrophy by activating PGC-1 α , and then blocking TWEAK. These data might be responsible for the explanation for PQQ preventing muscle atrophy. However, the exact molecular mechanism is still unclear.

Therefore, investigations into the signaling pathways that connect PQQ to cachexia induced muscle atrophy is still to be an important topic for future research, which might lead to the development of biological targets for therapeutic intervention to protect against skeletal muscle atrophy induced by cachexia.

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Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

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