Effect of Hazelnut Oil on Muscle Cell Signalling and Differentiation

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Abstract: Nuts-enriched diets were shown to bear beneficial effects for human’s health. Among nuts, hazelnut plays a major role in human nutrition and health because of its unique fatty acid composition (predominantly MUFA), fat soluble bioactives (tocopherols and phytosterols), vitamins (vitamin E), essential minerals (selenium), essential amino acids, antioxidant phenolics (caffeic acid), dietary fiber (soluble form), and bioactive phytochemicals. The current study was designed to explore the cellular effects of two particular hazelnut strains (Ordu and Tonda).

Four hazelnut oils were obtained from 2 common strains (Ordu hazelnut oil, Ordu cuticle oil, Tonda “gentile” hazelnut oil, Tonda “gentile” cuticle oil). The metabolic and nutritional effects of the four hazelnut oils were assessed using an in vitro model of mouse myoblasts, identifying the intracellular mechanisms involved in muscle differentiation and in the modulation of specific muscle genes.

We demonstrated that hazelnut oils induced morphological changes in neo-formed myotubes increasing myotubes size. In particular, the diversified effects of the hazelnuts and cuticle oils on muscle fibres shape (on length and diameter respectively) determine a diversified pattern of action on elongation or hypertrophy of the muscle fibres. Furthermore, hazelnut oils regulate different pathways associated with myoblasts growth and development, stimulate signal transduction, and activate cell commitment and differentiation.

The present results provide evidence that hazelnut oils may affect skeletal muscle growth and differentiation, constituting the proof of principle for the future development of novel foods and integrators.

Key words: hazelnut oils, skeletal muscle differentiation, novel foods, integrators

1 Introduction

Muscle is not only the major site of metabolic activity but it is the largest protein reservoir, providing a source of amino acids to be used for energy production during periods of food deprivation. Several studies have suggested that the loss of skeletal muscle may have serious health consequences. In contrast, physical activity, particularly resistance exercise, can generate large increases in skeletal muscle mass.

During the skeletal muscle differentiation, mononucleated myoblasts withdraw from the cell cycle and fuse into multinucleated myotubes, the basic cellular structures of muscle. Myogenesis is regulated by Myogenic Transcription Factors: MyoD, Myf5, Myogenin, Muscle Regulatory Factor 6 (Myf6), and Myocyte specific Enhancer Factor 2 (MEF2) regulating the expression of muscle-specific genes, such as myosin heavy chain (MyHC) and creatine kinase.

Numerous reports previously addressed the intracellular pathways, downstream of hormonal (insulin) and nutritional signals, involved in skeletal muscle protein synthesis during differentiation and hypertrophy, revealing two major pathways by which these signals are transmitted: (1) the IRS1/phoshatidylinositol 3 kinase (PI3K)/AKT/p70 S6K pathway and (2) the Ras/Raf/mitogen-activated protein kinase (MAPK) pathway. PI3-Kinase activation is involved in cellular anti-apoptotic processes via AKT activation, which is sufficient to induce hypertrophy in vivo and regulates the number, size and survival of mature muscle cells. Downstream, the p70 S6 kinase phosphorylation starts translation process. A strong positive relationship between activation of p70 S6 kinase and the long-term increase in muscle mass has been observed in rat skeletal muscle.
muscle following resistance training. In the second pathway, the activation of ERKs is necessary for the expression of muscle specific genes and phospho-ERK2 is essential for the formation of mature muscle fibres.

Skeletal muscle is intrinsically a stimulus-responsive tissue not only sensitive to the insulin stimulation, but also to the mechanical stimulus. Increased mechanical stretching is a powerful spur toward protein synthesis in the skeletal muscle. The PI3K/AKT/p70 S6 kinase and ERKs pathways, the same pathways stimulated by insulin, have been indicated to be of importance to the control of skeletal muscle mass, especially under mechanical stimulation.

Free fatty acids are the muscle preferred energy fuel in endurance exercises. Their accumulation into myofibrils was described in marathon runners and endurance athletes. Therefore, there is increasing evidence to suggest that fatty acids play a major role in regulating metabolism through effects on gene expression. Previous works have demonstrated that fatty acids may play an important role in regulating skeletal muscle growth.

The most studied source of unsaturated fatty acids is olive oil with its 80% content in oleic acid. Hazelnut oil has also a similarly high content of oleic acid, but it also contains several anti-oxidants and additional saturated and unsaturated fatty acids. Hazelnut oil is used much less in Mediterranean diet prescriptions, although its use is increasing in recent years.

The current study was designed to explore the cellular effects of four particular hazelnut oils (Ordu hazelnut oil, Ordu cuticle oil, Tonda gentile hazelnut oil, Tonda gentile cuticle oil) using an in vitro model of mouse myoblasts, assessing the intracellular mechanisms involved in muscle differentiation and the modulation of muscle specific genes. Hazelnut plays a major role in human nutrition and health because of its unique fatty acid composition (predominating MUFA), fat soluble bioactives (tocopherols and phytosterols), vitamins (vitamin E), essential minerals (selenium), essential amino acids, antioxidant phenolic (caffeic acid), dietary fibre (soluble), and phytochemicals.

### 2 Experimental

#### 2.1 Materials

Anti-GAPDH (FL-335), anti p-ERK (E-4), anti p-p70 S6 kinase (Thr 421/Ser 424)-R, anti-MyoD (C-20), anti-Myf5 (C-20), anti-Myf6 (C-19), anti-Myogenin (D-10) and anti-MyHC (H-300), monoclonal or polyclonal primary antibodies, the peroxidase-conjugated secondary antibodies and rhodamine-conjugated secondary antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, U.S.A.). All other reagents were purchased from Sigma Chem. Co. (St. Louis, MO, U.S.A.). Mouse C2C12 myoblasts cells were purchased from the European Collection of Animal Cell Cultures (ECACC).

#### 2.2 Cuticle oil extraction

The hazelnut skin oil extractions were performed with a NATEX extractor (Natex, Extraction Co2 Pilot Unit 5.0L–Terniz, Austria). With NATEX extractor was possible to work in dynamic phase where the flux of CO2 was in continuous. FOOD 2 Carbon dioxide was used as solvent. A sample of about 1300 g has been loaded into the extraction reactor.

The start of extraction was set to achieve temperature and pressure of work and after a period of stabilization parameters. In the dynamic phase of supercritical CO2 extraction was observed the following conditions:
- Extractor pressure 300 bar
- Temperature 50°C
- Flow CO2 Kg/h 30, Tot solvent 120 kg

After that, a cycle with reduced extraction time, pres-
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or temperature was conducted.

Extraction yield was about 27% for Turkey hazelnut and 26% for Piedmont hazelnut.

2.3 Kernel Oil extraction

Hazelnut kernel oil extractions were performed with IR heating BROVIND pilot plant. Sample were first roasted, then ground in a disc mill. 350 g aliquots have been pressed using hydraulic press. Leaving oil temperature was about 41°C.

Yields have been calculated by difference:

\[
\text{Yield} = \left( \frac{(P_e - P_s)}{P_e} \right) \times 100
\]

where

\[ P_e = \text{initial weight sample (loaded into the press)} \]
\[ P_s = \text{dry sample weight (drained from the press)} \]

Extraction yield was about 63% for Turkey hazelnut (Ordu hazelnut oil and Ordu cuticle oil) and 67% for Piedmont hazelnut (Tonda gentile hazelnut oil and Tonda gentile cuticle oil). The main properties of oils used in this work are reported in Table 1.

2.4 Experimental protocol

C2C12 myoblasts were cultured at 37°C in humidified 5% CO₂ atmosphere in a growth medium (GM) containing DMEM supplemented with 20% (v/v) FBS (fetal bovine serum), 1% penicillin–streptomycin, and 1% L-glutamine. Cell differentiation was initiated by placing 70% confluent cells cultures in DMEM supplemented with 1% HS (horse serum) and antibiotics (DM). DM was changed every 24 h.

For the proliferating myoblasts experiments, the cells were grown to 40-50% confluence in GM, for the myoblasts experiments the cells were cultured until 70-80% confluence in GM, while for the myotube experiments C2C12 cells were differentiated in DM for 72 h (Fig. 1a). The cells used as control were maintained respectively in GM (C0) or in DM (C0).

Five other groups of cells were placed in medium with the selective addition respectively of Ordu hazelnut oil (OHO; 10%) or of Ordu cuticle oil (OCO; 10%), of Tonda gentile hazelnut oil (TGHO; 10%) or of Tonda gentile cuticle oil (TGCO; 10%), of commercial olive oil (OLIVE; 10%). Cells were lysed at 30, 60 and 240 min after the stimuli addition. For Immunofluorescence analysis, C2C12 differentiated myocytes were stimulated for 4 h.

2.5 RT-PCR array analysis

Gene expression levels were assessed by RT2 -PCR Arrays System. In general, RT2-PCR Array plates produced by SABiosciences (SABiosciences Corporation, Frederick, MD 21703 USA) were designed to analyze simultaneously a panel of genes. We decided to study the following gene expression: MyoD, Myf5, Myf6, Myogenin, Myoatin, MyHC, p21, Rb, cyclin D and cyclin B1. GAPDH/Actin housekeeping genes were included as endogenous normalization controls. Total RNA was isolated from C2C12 cells using the RNeasy Plus Mini Quiagen kit (Quiagen GmbH, Germany) according to the manufacturer’s instructions. Total RNA (1 µg) was reverse transcribed using RT2 First Strand Kit (SABiosciences Corporation, Frederick, MD 21703 USA). The reverse transcripts were used as templates for analysis of gene expression level using RT2-PCR Arrays plates according to the manufacturer’s instructions. Each sample was run in triplicate. The expression level of the housekeeping genes chosen for normalization in the threshold cycle (ΔCt) for each experimental condition and then the fold-change (ΔΔCt) for each gene from treatment group compared to the control group was calculated. If the ΔΔCt is greater than 1, then the result may be reported as a fold up-regulation. If the ΔΔCt is less than 1, then the result may be reported as a fold down-regulation.

2.6 Immunoblotting analysis

C2C12 extracts were prepared by lising the cells in RIPA buffer at 4°C. Aliquots of 30 µg supernatant proteins were separated with SDS-polyacrylamide gel electrophoreses (SDS-PAGE) and electrophoretically transferred to nitrocellulose membranes (Protran, Whatman Schleicher & Schuell) using a Bio-Rad Mini Trans-Blot system. Blots were blocked and then incubated with respective primary antibodies and then with species-specific secondary antibodies. To confirm equal protein loading per sample, we used antibody anti-GAPDH. Immunoreactive bands were visualized by an enhanced chemiluminescence method (Amersham Pharmacia Biotech, Piscataway, NJ, USA) and quantified by densitometric analysis using the Scion image software (Scion corporation, Frederick, MD, USA). Data were then converted into fold-changes (FC) of the controls.

2.7 Immunofluorescence analysis

Proliferating myoblasts or differentiated myotubes were fixed in prepared 4% paraformaldehyde and then, maintained in PBS. The resulted cells were permeabilized with 0.2% Triton X-100 and blocked with PBS containing 1% bovine serum albumin (BSA). Then, cells were incubated with primary antibodies. To detected primary antibody binding sites, cells were washed in PBS and followed by incubation with specific antibodies rhodamine-conjugated. Coverslips with cells were mounted with Moviol and observed using Nikon Eclipse 50i fluorescence microscopy and images were captured using Nis-Elements D 4.00 software.

2.8 Statistical analysis

All experiments were performed three times. The data are presented as the mean ± standard deviation and statistical comparisons were performed with specific statistical packages (Prism v 7.00 GraphPad Software, San Diego, CA, USA). Statistically significant differences were determined using ANOVA test followed by Tukey’s multiple comparison
3 Results

In the experiments, we tested the ability of both hazelnut oils (OHO-OOC / TGHO-TGCO) and of a common commercial extra-virgin olive oil (OLIVE) to stimulate the same pathways in myoblast un-differentiated cells.

3.1 Effects of hazelnut oils during proliferative phase

In the proliferative phase, we investigated whether hazelnut and olive oils were able to induce protein synthesis by the activation of p70 S6 kinase. OCO (Fig. 1A) or TGHO (Fig. 1B) treatments caused a significant p70 S6 kinase phosphorylation increase (30, 60 and 240 min), while OHO (Fig. 1A), TGCO (Fig. 1B) or OLIVE oil did not modify p70 S6 activity.

Results were considered statistically significant when $p \leq 0.05$. 

Fig. 1 During proliferative phase, OCO (A) or TGHO (B) promoted p70 S6 kinase activation, while OHO, OCO (A) or TGHO (B) increased ERK1/2 phosphorylation at 240 min. Representative immunoblots are shown. Tukey post test: § $p \leq 0.001$ vs GM40 confluence.
Moreover, we analyzed the Ras/MEK/ERK cascade involved in mitosis and cell proliferation. We determined the levels of ERK1 and ERK2 phosphorylation\(^7\); at 240 min, ERK1 and ERK2 were activated by OCO, OHO (Fig. 1A) and TGOH oils (Fig. 1B). These results suggest how in myoblast cells hazelnut oils could positively induce myocyte commitment in respect to the OLIVE.

To confirm this preliminary evidence, we investigated MyoD and Myf5 protein expression in C2C12 myoblast cells by Immunofluorescence analysis\(^3\) after 4 h of oils treatment: our data suggested that OCO and TGOH induced the first morphologic changes to commit myoblasts in myocyte cells (Fig. 2).

### 3.2 Effects of hazelnut oils during differentiation phase

Since myocytes differentiation requires cell cycle withdrawal\(^2, 3\), we studied the gene expression of cell cycle regulators and myogenic regulatory factors (MRFs) MyoD, Myf5, Myf6 and Myogenin in myoblast cells after 4 (Fig. 3A) and 48 h (Fig. 3B) of oils treatment. As reported in Experimental section, if the \(\Delta\Delta C_t\) is greater than one, then the result can be reported as a fold up-regulation. If the \(\Delta\Delta C_t\) is less than one, then the result can be reported as a fold down-regulation.

We observed the increase of key cell cycle regulators and of MRFs gene expression after 4 (Fig. 3A) and 48 h (Fig. 3B) of hazelnut oils supplementation. In contrast, the expression of these MRFs was not detected in cells treated with OLIVE.

These results indicate that treatments with hazelnuts oils promoted the gene expression of MRFs\(^7\) and skeletal muscle specific protein MyHC not only in the early phase (Fig. 3A) but also during differentiation (Fig. 3B). To confirm this observation and clarify the degree of myogenic differentiation induced by oils stimulation, we studied the effects on protein synthesis by Western Blot and Immunofluorescence analysis.

Subsequently, we examined the effect of oils during the
differentiation phase. After 24 h of differentiation in presence of OLIVE oils, most of cells showed a markedly reduced vitality at 48 h, the total C2C12 cells were died. We did not observe any toxic effects in presence of hazelnut oils.

First, we studied the activation of the principal protein kinases involved in protein synthesis and muscle differentiation\(^2\). As shown in Fig. 4\(^A\), all hazelnut oils induced p70 S6 activation (Figs. 4A, 4B).

ERK1/2 play an important role during muscle differentiation and hypertrophy\(^10\). Hazelnut and cuticle supplements increased ERK1/2 activation at all time-points (Figs. 4A, 4B). These results indicated that hazelnut oils could modulate specific signaling pathways involved in myogenic factors synthesis.

In fact, OCO, TGHO or TGCO treatment increased MyoD protein level (Figs. 5A, 5B) and all hazelnut oils increased Myogenin protein levels in respect with untreated C2C12 (DM). We determined MyHC synthesis: at the end of differentiation, MyHC levels were the same in the negative control (DM) and in treated cells with OCO, OHO or TGCO (Fig. 5A), while TGHO increased MyHC protein levels (Fig. 5B).

3.3 Effects of hazelnut oils on post-differentiation hypertrophy

To clarify hazelnut oils action on hypertrophy process, neo-formed myotubes were treated for 30, 60 and 240 min...
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with the five oils after 72 h differentiation.

We observed a strong increase of p70 S6 kinase phosphorylation after 30 min OHO or TGCO stimuli (Figs. 6A, 6B). In contrast, OLIVE oil significantly decreased p70 S6 kinase phosphorylation level. Furthermore, we observed a weak modulation of ERKs pathway (Fig. 6). At 240 min, OCO and TGCO oils promoted ERKs phosphorylation (Figs. 6A, 6B).

We examined neo-formed myotubes morphologic features. Images of MyHC-positive myotubes, detected by Immunofluorescence after 4 h, showed that OLIVE oil negatively influenced myotubes formation. In contrast, cells stimulated with OHO showed an increase of the diameter respect DM, while neo-formed myotubes stimulated with OCO or TGCO were longer compared to DM (Fig. 7A).

This evidence was further confirmed by Myf6 Immunofluorescence assay (Fig. 7B): OCO, TGHO and TGCO oils showed a major number of Myf6 positive myotubes.

Taken together, these results indicate that hazelnuts oils can induce the activation of signaling pathways involved in muscle differentiation and hypertrophy and, in contrast to

Fig. 4 After 48 h of differentiation in presence of OLIVE oil, the total C2C12 cells were died. All hazelnut oils increased p70 S6 kinase and ERK1/2 activation at all time point (A, B). Representative immunoblots are shown. Tukey post test: ^ p ≤ 0.002, § p ≤ 0.001 vs DM 100% confluence.
common commercial OLIVE, can promote the different phases of muscle differentiation process (Table 2).

4 Discussion
The skeletal muscle mass is closely related to the number of cells which trigger the myogenic program and are able to undergo hypertrophy at later stages of muscle development\(^{5,9}\). The nutritional and humoral factors controlling skeletal muscle mass and function acting during development deserve particular attention. In fact, muscle hypotrophy/atrophy (sarcopenia) is an important clinical problem, negatively impacting human health. Chronic inactivity (such as prolonged bed rest), accelerates the aging process\(^{21}\). Poor nutrition and physical inactivity are the 2nd...
cause of mortality (next to cigarette smoking) in humans (Center of Diseases Control, Atlanta, GA). Currently, dietary recommendations and physical exercise counselling are the prevailing means to counteract muscle hypotrophy/atrophy.

Among dietary nutrients, amino acids were known since decades to stimulate protein anabolism and to mimic insulin effect activating intra-cellular signals into several cell types\(^{22}\). Overall, the effect of amino acids is comparable to the one of high protein diet, although branched-chain amino acids seem to have the highest insulin-mimetic and trophic activity in skeletal cells\(^{23, 24}\). Similar effects of high-fat diets and single fatty acids integration on muscle accretion are much less known. Chanseaume and others\(^{25}\) demonstrated that an increase muscle mixed and mitochondrial protein synthesis rates after a high-fat or high-sucrose diet. Very recently, Rincón-Cervera and others\(^{26}\) showed that supplementation with extra-virgin olive oil

Fig. 6 In neo-formed myotubes, OHO or TGCO induced a strong increase of p70 S6 kinase phosphorylation after 30 min of stimulation. OLIVE oil significantly decreased p70 S6 phosphorylation level (A, B). At 240 min, OCO and TGCO oils promoted ERKs phosphorylation (A, B). Representative immunoblots are shown. Tukey post test: § p ≤ 0.001 vs DM\textsubscript{post-diff}.
prevents hepatic oxidative stress and reduction of desaturation capacity in mice fed a high-fat diet. We presently report the effects of hazelnut oils on intracellular pathways and myotubes differentiation in a model of murine myoblasts. This is the first study to describe the effects of hazelnut oil on Myogenic Regulatory Factors (MRFs) expression during differentiation in C2C12 skeletal muscle cells. We presently show that hazelnut oil integration is able to stimulate myogenic phenotype acquisition and differentiation process through MRFs protein expression in C2C12 in vitro model.

Concerning the mechanism(s), our data suggest that the activation of ERKs pathways and the strong activation of kinases involved in the protein synthetic pathway (AKT and p70) appears to be responsible of an enhanced differentiation process with a selective stimulation in the proliferative (Fig. 1) differentiation (Fig. 4) and post-differentiation (Fig. 6) phases. This is a novel mechanism, not common to other "healthy" oils as olive oil. In fact, olive oil, used as control, showed a scarce or absent effect on proliferation, differentiation and post-differentiation phase of myoblasts. A previous study described the reduction of fat mass with an increase of muscle mass following the assumption of 12 table green olives/day for 30 days. We presently use commercial standard olive oil as control, whilst Accardi et al used extra-virgin olive oil. Moreover, the experimental set-up (whole-body vs. murine cell models) used in the 2 studies was significantly different. However, the differential effect on muscle cell metabolism/homeostasis between hazelnut oil and olive oil is noteworthy and may be explained as follows: 1) hazelnut oil has a higher content of oleic acid with respect to standard olive oil; 2) the polyphenols content and total antioxidant power is definitely higher in hazelnut oil with respect to standard olive oil. Since the PREDIMED trial showed that dietary polyphenols trigger nitric oxide (NO) production, it is conceivable that a more pronounced NO release, besides having the well-known vasodilatory effect, might also contribute to cell proliferation as previously demonstrated in cardiomyoblasts.

Del Rio and others previously reported on the 3-fold higher anti-oxidant power of cuticle with respect to the body in hazelnuts. Therefore, we tested also oils derived only by cuticle of hazelnut on cellular signals as well as cell differentiation and proliferative activity. All four hazelnut oils induced morphological changes in neo-formed myotubes increasing myotubes size. Nonetheless, OCO and TGHO have the stronger effect in activating p70 S6 kinase and pERK2 at 60 and 240 min of the proliferative phase. This indicates the 2 hazelnut oils as potential candidates to be used as novel foods or to develop and test food integrators with the specific target of stimulating protein anabolism. For instance, poorly-controlled diabetes mellitus is characterized by increased protein catabolism, reduced

Fig. 7 In neo-formed myotubes, OHO cells showed an increase of the diameter with respect to the control DM. The cells stimulated with OCO or TGCO showed a rise in the length in MyHC positive myotubes compared with the un-stimulated differentiated cells (DM) (A). OCO, TGHO and TGCO treatments showed a major number of Myf6 positive myotubes (B).
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5 Conclusion

Studies are required to specifically address this issue in our hazelnut oils strains and extraction methodologies in muscle mass and muscle fibers strength. The present in vitro data constitute a proof of principle for testing specific hazelnut oils strains and extraction methodologies (mainly OCO and TGHO) as potential food integrators to counteract protein catabolism in diabetic patients.

An important issue to be considered is whether the effect of OCO and TGHO may be mediated by epigenetic modifications induced by a hazelnut oil enriched diet over the decades. In a previous study, Surra and others demonstrated that, dietary nut supplementation delays the development of atherosclerotic lesions in female apoE-deficient mice, increasing the mRNA expression of genes involved in non-HDL cholesterol synthesis. Additional studies are required to specifically address this issue in our experimental setting.

Table 2 Table resuming the effects of the 5 oils tested in this study during: a) Proliferative stage, b) Differentiation phase, and c) Post-differentiation hypertrophy. ▲, increase; ▼, decrease; =, no effects vs Control.

|                | Proliferative stage | Differentiation phase | Post-differentiation hypertrophy |
|----------------|---------------------|-----------------------|---------------------------------|
|                | pp70 pERK Ilyocyte Commitment | pp70 pERK Myogenic factors synthesis | pp70 pERK Myogenic factors synthesis |
| OLIVE          | ▼ ▼ ▼ | Cell Death            | ▼ ▼ ▼ ▼ ▼ ▼ ▼ |
| OHO            | ▼ ▲ ▼ | ▲ ▲ ▼ ▼ ▼ | ▼ ▼ ▼ ▼ ▼ |
| OCO            | ▼ ▼ ▼ ▼ | ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ | ▼ ▼ ▼ ▼ ▼ |
| TGHO           | ▼ ▼ ▼ ▼ ▼ | ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ | ▼ ▼ ▼ ▼ ▼ |
| TGCO           | ▼ ▼ ▼ ▼ ▼ | ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ ▼ | ▼ ▼ ▼ ▼ ▼ |

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