

# Contrarily to whey and high protein diets, dietary free leucine supplementation cannot reverse the lack of recovery of muscle mass after prolonged immobilization during ageing

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## Key points

- During ageing, there is a lack of recovery of muscle mass following immobilization.
- We showed, in old rats, an 'anabolic resistance' of muscle protein synthesis to food intake during immobilization and only a slight increase of protein synthesis during the recovery, which explain a poor muscle nitrogen balance that is insufficient to induce a muscle mass gain.
- A supplementation with free leucine, an essential amino acid known to stimulate muscle protein metabolism, was efficient in inducing a greater anabolism but failed to induce muscle mass recovery.
- This discrepancy was explained by a 'desynchronization' between the leucine signal and amino acids coming from dietary protein digestion.
- An induction of a larger increase and a longer availability of amino acids in the postprandial state with rich-protein leucine (i.e. whey) and high protein diets were efficient in inducing a muscle mass recovery after immobilization.

**Abstract** During ageing, immobilization periods increase and are partially responsible of sarcopaenia by inducing a muscle atrophy which is hardly recovered from. Immobilization-induced atrophy is due to an increase of muscle apoptotic and proteolytic processes and decreased protein synthesis. Moreover, previous data suggested that the lack of muscle mass recovery might be due to a defect in protein synthesis response during rehabilitation. This study was conducted to explore protein synthesis during reloading and leucine supplementation effect as a nutritional strategy for muscle recovery. Old rats (22–24 months old) were subjected to unilateral hindlimb casting for 8 days (I8) and allowed to recover for 10–40 days (R10–R40). They were fed a casein ( $\pm$ leucine) diet during the recovery. Immobilized gastrocnemius muscles atrophied by 20%, and did not recover even at R40. Amount of polyubiquitinated conjugates and chymotrypsin- and trypsin-like activities of the 26S proteasome increased. These changes paralleled an 'anabolic resistance' of the protein synthesis at the postprandial state (decrease of protein synthesis, P-S6 and P-4E-BP1). During the recovery, proteasome activities remained elevated until R10 before complete normalization and protein synthesis was slightly increased. With free leucine supplementation during recovery, if proteasome activities were normalized earlier and protein synthesis was higher during the whole recovery, it nevertheless failed in muscle mass gain. This discrepancy could be due to a 'desynchronization' between the leucine signal and the availability of amino acids coming from casein digestion. Thus, when

supplemented with leucine-rich proteins (i.e. whey) and high protein diets, animals partially recovered the muscle mass loss.

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**Abbreviations:** ASR, absolute synthesis rate; CAS, casein; LEU, leucine; PA, postabsorptive; PP, postprandial.

## Introduction

Sarcopaenia is the progressive loss of muscle mass and strength associated with normal ageing (Rosenberg 1989). This phenomenon is a highly predictive factor of frailty, of limited mobility, of increased susceptibility to injury and of impaired recovery (Harris 1997). Many mechanisms have been proposed to explain sarcopaenia. Besides chronic and progressive loss, periods of immobilization or acute inactivity which increase with age may contribute by themselves to muscle atrophy. Thus, the unloading-induced atrophy due to limb immobilization, chronic bed-rest or lack of physical inactivity impairs muscle mass and functions in the elderly (Bar-Shai *et al.* 2005). However, the effect of disuse by itself on skeletal muscle in elderly individuals has not been extensively investigated and the subsequent recovery ability has been even less studied despite the fact that we and others have shown an impaired recovery in old animals and elderly humans after immobilization-induced muscle atrophy (Chakravarthy *et al.* 2000; Suetta *et al.* 2009; Magne *et al.* 2011). These periods of uncompleted recoveries repeated over time may contribute to a significant muscle mass loss and then worsen sarcopaenia, and this phenomenon has been named the 'acute catabolic crisis' model (English & Paddon-Jones, 2010).

At a cellular level, muscle atrophy following immobilization results from an imbalance of protein turnover as the maintenance of skeletal muscle mass depends on the overall balance between rates of protein synthesis and protein breakdown. In adults, protein synthesis has been reported to be reduced at the post-absorptive state (Gibson *et al.* 1987) and muscle protein breakdown increased (Loughna *et al.* 1986; Taillandier *et al.* 1993, 1996; Lawler 2003; Krawiec *et al.* 2005; Vazeille *et al.* 2008) leading to the muscle protein loss. Muscle mass recovery after unloading is also dependent on the generation of a sustained positive nitrogen balance which results from normalization of changes in protein metabolism associated with an increased protein synthesis, decreased proteolysis, or simultaneous changes in both processes. Thus, during recovery in adults, muscle protein synthesis was increased (Booth 1982) and muscle protein breakdown was down-regulated, normalized or increased according to the model of immobilization used (Tilgnaç *et al.* 2002; Taillandier *et al.* 2003; Minnaard *et al.* 2005; Vazeille *et al.* 2008), which allowed a muscle mass

recovery over time after reloading. We have shown that during ageing, mechanisms of casting-induced atrophy were similar to those observed in adult animals, i.e. an increased ubiquitin-proteasome-dependent proteolysis and apoptosis in skeletal muscle (Magne *et al.* 2011). However, we recorded a lack of muscle mass recovery, which was not due to a sustained activation of these pathways after reloading as they were rapidly normalized during the recovery period. If muscle protein synthesis seems to be reduced after an immobilization period in elderly people (Kortebein *et al.* 2007), no data on muscle protein synthesis during recovery are available. Our previous work strongly suggests that protein synthesis could be impaired during the recovery period (Magne *et al.* 2011) and we have hypothesized that this alteration may explain the absence of a positive nitrogen balance and subsequently the lack of recovery of muscle mass we observed during ageing. If protein synthesis is indeed impaired during reloading, finding ways to stimulate it may be essential.

It is now well established that feeding is a robust stimulator of muscle protein synthesis (Kimball *et al.* 2000; Bohe *et al.* 2003; Moore *et al.* 2009). However, it is also well known that the stimulating effect of food intake on muscle anabolism is already less efficient in both old animals and elderly humans and this without any pathologies or physical inactivity (reviewed in Balage & Dardevet, 2010). These alterations of muscle protein synthesis response have been partially explained by the presence of a low grade inflammation and/or oxidative stress development with age (Marzani *et al.* 2008; Rieu *et al.* 2009). Interestingly, we and others have previously shown that immobilization generated a further increase in the intramuscular inflammatory and oxidative stress markers (Kondo *et al.* 1991; Biolo *et al.* 2008; Magne *et al.* 2011), which could further alter protein synthesis response to food intake and then contribute to the total lack of muscle mass recovery in the elderly after reloading. Then, during the recovery period, the postprandial phase should play a critical role since it is during this period that the major anabolic factors regarding protein metabolism are elevated.

Amino acids and particularly leucine play an important role in the stimulation of postprandial muscle protein synthesis (Anthony *et al.* 1999, 2001; Koopman *et al.* 2005). Leucine is known as a 'nutrient signal' as it reduces muscle protein breakdown and further stimulates muscle

protein synthesis (Buse & Reid, 1975; Frexes-Steed *et al.* 1992; Crozier *et al.* 2005) particularly during ageing (Dardevet *et al.* 2002; Rieu *et al.* 2003, 2006, 2007; Combaret *et al.* 2005; Katsanos *et al.* 2006). The 'leucine signal' stimulates the mammalian target of rapamycin (mTOR) pathway and 70 kDa ribosomal protein S6 kinase (p70S6 kinase) activity, and enhances eIF4E-binding protein (4EBP1) phosphorylation and the association of eukaryotic initiation factor 4E (eIF4E) with eukaryotic initiation factor 4G (eIF4G) both *in vitro* and *in vivo* (Anthony *et al.* 2000; Dardevet *et al.* 2000; Bolster *et al.* 2004). These signalling pathways are known to become resistant to amino acid (i.e. leucine) stimulation in the case of chronic inflammation or oxidative stress (Lang *et al.* 2002; Marzani *et al.* 2008; Balage *et al.* 2009), phenomena that occur during immobilization in aged muscle (Magne *et al.* 2011). We hypothesized that following immobilization, a sustained resistance of muscle protein synthesis to leucine persisted and that increasing leucine intake could have a beneficial effect on muscle mass recovery by making the nitrogen balance positive after reloading. To our knowledge, no study has yet tested such supplementation in aged muscle recovering from immobilization. The aims of this study were to (1) determine if muscle protein synthesis response to food intake is altered during atrophy following immobilization and remains resistant to food intake stimulation during the subsequent recovery, and (2) test the effect of a leucine supplementation on muscle mass recovery by measuring muscle protein synthesis, proteolysis and signalling proteins associated with these pathways in both the postabsorptive and postprandial states.

## Methods

### Animals and experimental design

All procedures were performed in accordance with institutional guidelines on animal experimentation in France and comply with the policies and regulations of *The Journal of Physiology* given by Drummond (2009). Male Wistar rats aged 22–24 months were housed individually under controlled environmental conditions (room temperature 22°C; 12 h light–dark cycle, light period starting at 08.00 h), fed *ad libitum* a standard 13% casein diet (Table 1) and given free access to water.

After a 3 week adaptation period, 143 rats were anaesthetized with isoflurane inhalation and subjected to unilateral hindlimb cast immobilization with an Orfit-soft plaque (Gibaud, France) for 8 days (I8). The foot was positioned in plantar extension to induce maximal atrophy of the gastrocnemius muscle (Goldspink, 1977; Booth, 1982; Krawiec *et al.* 2005; Vazeille *et al.* 2008; Magne *et al.* 2011). For muscle recovery, casts were removed and animals were allowed to recover for 10 (R10), 20 (R20),

**Table 1. Composition of the standard and the experimental diets**

	Standard diet	Control and experimental diets	
	CAS	CAS + ALA	CAS + LEU
Ingredients			
Casein	166	166	166
L-Cystine	1.8	1.8	1.8
Alanine <sup>1</sup>	—	59	—
Leucine <sup>2</sup>	—	—	44.5
Valine <sup>2</sup>	—	—	5.1
Isoleucine	—	—	9.8
Colza oil	30	30	30
Peanut oil	27	27	27
Sunflower oil	3	3	3
Cellulose	35	35	35
Saccharose	100	100	100
lactose	134	134	134
Wheat flour	458.2	399.2	398.8
Mineral mixture	35	35	35
AIN93			
Vitamins mixture	10	10	10
AIN93			
given during	adaptation immobilization	recovery	recovery

Quantities are expressed in g/kg dry matter.

<sup>1</sup>Alanine was included in the control CAS+ALA diet to render the diets isonitrogenous. This amino acid has no effect on muscle protein metabolism.

<sup>2</sup>Valine and isoleucine were included in the CAS+LEU diet to prevent the fall of their plasma concentrations induced by leucine supplementation. CAS: casein; CAS+ALA: casein + alanine; CAS+LEU: casein + leucine.

30 (R30) or 40 (R40) days. All rats were fed the standard 13% casein diet during immobilization. Then half of the animals were fed a control diet (casein (CAS) + alanine (ALA)) and constituted the control (CON) group; the other half were fed a 4.45% leucine-supplemented diet (CAS + leucine (LEU)) and were the LEU group (Table 1). The CAS + LEU diet was supplemented with leucine to increase plasma leucine concentration. To prevent the fall of plasma valine and isoleucine concentrations induced by leucine supplementation, the CAS + LEU diet was also supplemented with appropriate amounts of these amino acids Rieu *et al.* 2003). Alanine, an amino acid that has no effect on muscle protein metabolism, was included in the control CAS + ALA diet to render the diets isonitrogenous.

Casted rats reduced their food intake during the immobilization period. Therefore, 39 control non-casted rats were pair-fed (PF) to the casted group (18 at I8 and 21 at R20). Seventeen rats were studied as a reference point before the immobilization period (I0).

Before immobilization (I0) and at the end of the immobilization (I8) or recovery periods (R10, R20, R30,

R40), animals were killed under pentobarbital sodium anaesthesia (50 mg kg<sup>-1</sup> i.p.). On the morning of each time point studied, half of the rats in each group were not fed, so that they were in a postabsorptive state. These rats were designed 'PA' (PA CON and PA LEU). The others ate their respective diet (i.e. CAS + ALA or CAS + LEU) for 1 h and then were in the postprandial state; they were named 'PP' (PP CON and PP LEU). Non-casted pair-fed rats were also studied in the PA and the PP states using the same procedure (PA standard diet and PP standard diet).

### Measurements of *in vivo* protein synthesis

Protein synthesis rates were measured using the flooding-dose method. Each rat was injected intravenously with [1-<sup>13</sup>C]valine (99%) (150 μmol (100 g body weight)<sup>-1</sup>), 40 min before killing (i.e. 110–140 min after the beginning of the experimental diets), to flood the precursor pool with [1-<sup>13</sup>C]valine as previously described (Mosoni *et al.* 1996). Rats were then killed under pentobarbital sodium anaesthesia (50 mg kg<sup>-1</sup> i.p.). Blood was withdrawn from the aorta, and hindlimb gastrocnemius muscles were carefully dissected, weighed and frozen in liquid nitrogen.

Free and bound valine enrichments were determined as follows. Muscles were powdered in liquid nitrogen in a ball mill (Dangoumeau, Prolabo, Paris, France). A 200 mg aliquot of frozen muscle powder was homogenized in 2 ml of 10% trichloroacetic acid (TCA). Homogenates were centrifuged (10,000 g, 15 min, 4°C) and supernatants, containing free amino acids, were desalted by cation-exchange chromatography (AG 50X8, 100–200 mesh, H<sup>+</sup> form, Bio-Rad, Richmond, CA, USA) in minidisposal columns. Valine and other amino acids were eluted with 4 mol l<sup>-1</sup> NH<sub>4</sub>OH. After evaporation of NH<sub>4</sub>OH under vacuum, free amino acids were resuspended in 0.01 mol l<sup>-1</sup> HCl for enrichment measurements. TCA-insoluble materials were washed 3 times in 4 volumes of cold 10% TCA and once in 4 volumes of 0.2 mol l<sup>-1</sup> perchloric acid (PCA). Resultant pellets were resuspended in 0.3 mol l<sup>-1</sup> NaOH and incubated at 37°C for 1 h. Protein concentration was determined using the bicinchoninic procedure. Proteins were precipitated with 20% PCA overnight at 4°C, and samples centrifuged (10,000 g, 5 min, 4°C). The protein pellet was hydrolysed in 6 mol l<sup>-1</sup> HCl at 110°C for 24 h. HCl was removed by evaporation and amino acids purified by cation-exchange chromatography as described above. Measurement of free valine enrichment was done as its *t*-butyldimethylsilyl derivative by gas chromatography electron impact mass spectrometry, using a gas chromatograph coupled to an organic mass spectrometer quadrupole (GC-M; Hewlett-Packard 5971A (Hewlett-Packard Co., Palo Alto, CA, USA)). Enrichment of [1-<sup>13</sup>C]valine into muscle proteins was measured as its *N*-acetyl-propyl derivatives by gas chromatography–combustion–isotope ratio mass

spectrometry (GC-c-IRMS 'Isoprime' (Elementar France, Lyon, France)).

### Calculations

The absolute synthesis rate (ASR) was calculated from the product of the protein fractional synthesis rate (FSR) and the protein content of the tissue and expressed in mg day<sup>-1</sup>. FSR (in % day<sup>-1</sup>) was calculated from the formula:  $FSR = S_b \times 100/S_a \times t$ , where  $S_b$  is the enrichment at time  $t$  (minus natural basal enrichment of protein) of the protein-bound valine,  $t$  is the incorporation time in days, and  $S_a$  is the mean enrichment of free tissues valine between time 0 and  $t$ . The mean  $S_a$  enrichment was the  $S_a (t_{1/2})$  value calculated from the linear regression obtained in tissue between time 0 and time  $t$ .

### Plasma amino acid measurements

Plasma amino acid concentrations were determined for each group at I0, I8, R20 and R40. Plasma amino acids were purified, i.e. 500 μl of plasma was added to 125 μl of sulfosalicylic acid solution (1 mol l<sup>-1</sup> in ethanol with 0.5 mol l<sup>-1</sup> thiodiglycol) previously completely evaporated. Norleucine (100 μl) was added as an internal standard. Amino acid concentrations were determined using an automated amino acid analyser with BTC 2410 resin (Biotronic LC 3000, Roucaire, Velizy, France).

### Measurement of proteasome activities

In this pathway, two distinct steps are depicted: (1) the ubiquitination of proteins and (2) their degradation by the 26S proteasome. We first investigated the chymotrypsin- and the trypsin-like activity of the proteasome.

One hundred and fifty milligrams of gastrocnemius muscle powder from pair-fed, non-immobilized and immobilized muscles at each time point was homogenized in 10 volumes of an ice-cold buffer containing 50 mM Tris-Cl (pH 7.5), 5 mM MgCl<sub>2</sub>, 250 mM sucrose, 1 mM dithiothreitol (DTT), 10 nM adenosine triphosphate (ATP) and protease inhibitors (10 μg ml<sup>-1</sup> antipain, 10 μg ml<sup>-1</sup> leupeptin, 10 μg ml<sup>-1</sup> aprotinin, 10 μg ml<sup>-1</sup> pepstatin A, 20 μM phenylmethanesulphonyl fluoride (PMSF)).

Briefly, extracts were centrifuged at 10,000 g for 20 min at 4°C. Supernatants were then centrifuged at 150,000 g for 30 min at 4°C and resulting supernatants were finally centrifuged at 150,000 g for 2.5 h at 4°C. The resulting protein pellets were resuspended in 150 μl of a buffer containing 20% glycerol, 5 mM MgCl<sub>2</sub> and 50 mM Tris-Cl (pH 7.5) (Buffer A). Protein concentration was determined on these resuspended pellets (Bio-Rad assay). The peptidase activities of the proteasome (chymotrypsin-like and trypsin-like activities) were determined by measuring

the hydrolysis of the fluorogenic substrates succinyl-Leu-Leu-Val-Tyr-7-amido-4-methylcoumarin (LLVY-AMC) (Sigma, USA) and the Boc-Leu-Arg-Arg-7-amido-4-methylcoumarin (LRR-AMC) (Enzo Life Sciences, Exeter, UK), respectively. To measure the proteasome chymotrypsin-like and trypsin-like activities, 15  $\mu\text{g}$  of proteins from the resuspended pellets diluted in 15  $\mu\text{l}$  of Buffer A were added to 60  $\mu\text{l}$  of medium containing 50 mM Tris-Cl (pH 7.5), 11.25 mM  $\text{MgCl}_2$ , 1.25 mM DTT, 0.01 U apyrase and 300  $\mu\text{M}$  LLVY-AMC or 800  $\mu\text{M}$  LRR-AMC. Pilot experiments were performed with or without inhibitors of the chymotrypsin-like activity (MG132, Affiniti) to ensure that the activities were totally inhibited. The trypsin-like activity was measured with and without the specific inhibitor lactacystin (lactacystin, 100  $\mu\text{M}$ , Sigma). Both activities were determined by measuring the accumulation of the fluorogenic cleavage product (amido-4-methylcoumarin; AMC) using a luminescence spectrometer FLX800 (Biotek, USA) during 45 min at 380 nm excitation wavelength and 440 nm emission wavelength. The time course for the accumulation of AMC after hydrolysis of the substrate was analysed by linear regression to calculate activities, i.e. the slopes of best fit of accumulation of AMC vs. time.

### Analysis of muscle protein synthesis and proteolysis signalling pathways by Western blotting

Powder (300 mg) of gastrocnemius muscles was homogenized in 10 volumes of a buffer containing 1 mM dithiothreitol (DTT), 0.1 mM phenylmethanesulphonyl fluoride (PMSF), 1 mM benzamidine and 0.5 mM sodium vanadate. Extracts were then centrifuged at 9,500 rpm for 12 min at 4°C. Aliquots of supernatants were diluted in sample buffer, boiled for 5 min, and stored at -20°C until protein immunoblot analyses. Equal amounts of proteins were separated by SDS-PAGE and transferred to PVDF membranes (GE Healthcare, Orsay, France).

### Muscle proteasome proteolysis pathway

To explore the first step of the ubiquitin-proteasome-dependant proteolysis (tagging of proteins by ubiquitin before their recognition by the proteasome) the anti-ubiquitinated proteins antibody, which recognizes poly-ubiquitin chains (Millipore, USA), was used at 1:2000 dilution. Twenty-five micrograms of proteins was separated on 7% acrylamide gels. FoxO3a is a transcription factor involved in atrogene transcription. When phosphorylated, its nuclear translocation is impossible. Thus, the ratio between FoxO3a and its phosphorylated form traduces an inhibition of atrogene transcription. The abundance of transcription factor FoxO3a and its phosphorylated form phospho-FoxO3a (Ser253) was determined using appropriate antibodies

(Cell Signaling Technology, Inc., Danvers, MA, USA) at 1:1000 and 1:1000 dilution, respectively. Fifty micrograms of proteins was separated on 7.5% acrylamide gels.

### Muscle protein synthesis pathway

Immunoblotting was performed using appropriate antibodies: S6 and phospho-S6 (Ser235/236) (Cell Signaling Technology) at 1:6000 and 1:6000 dilution, respectively. To determine the amount of S6, 25  $\mu\text{g}$  of proteins was separated on 15% acrylamide gels, and 30  $\mu\text{g}$  of proteins was separated on 12% acrylamide gels to quantify amount of phospho-S6. The amount of total 4EBP1 ( $\alpha$ ,  $\beta$  and  $\gamma$  forms) was determined on 50  $\mu\text{g}$  of proteins separated on 15% acrylamide gels, using an antibody (Cell Signaling Technology) at 1:4000 dilution.

Signals were detected using the ECL+ detection kit (GE Healthcare, France) after exposition onto radiographic film (Hyperfilm ECL, GE Healthcare) and quantified by densitometry using the ImageJ software. Signals were normalized against the signal obtained with a pool of samples used as reference.

### Index of muscle functionality

Muscles are composed of different fibre types which have different metabolic and contractile properties. As myosin is the most abundant contractile protein, muscle functionality is linked to the muscle myosin content. Myosin content was determined on total protein extracts according to the method described by Mizunoya *et al.* (2008). Briefly, 5  $\mu\text{g}$  of proteins was separated on 40% acrylamide-2% acrylamide-bisacrylamide gel during 15 h at 140 V. Gels were then coloured with Coomassie Blue and quantified with ImageQuant software (GE Healthcare).

### Pilot experiment: whey diet and high protein diet supplementations

To improve our nutritional strategy after the free leucine supplementation, a pilot experiment was carried out. Forty-one animals were subjected to cast immobilization for 8 days and then allowed to recover during 20 (R20) and 40 (R40) days. Eight animals were killed at I8 and the others were divided in two groups: the first one ( $n = 16$ ) received a soluble milk protein diet and the second one ( $n = 17$ ) received a high protein diet. Seven animals were used as control before casting (I0). All animals were studied in the postprandial state.

Soluble milk proteins diet was composed of (in  $\text{g kg}^{-1}$  diet): 144 whey (Prolacta, Lactalis ingredients), 5.7 proline, 30 colza oil, 3 sunflower oil, 27 peanut oil, 35 cellulose, 35 AIN93 mineral mix, 10 AIN93 vitamin mix, 2.5 cholin, 100 saccharose, 126 lactose and 482 wheat flour. High protein diet was composed of (in  $\text{g kg}^{-1}$  diet): 156

casein (Lactalis ingredients), 144 whey (source = ProLacta, Lactalis ingredients), 30 colza oil, 3 sunflower oil, 27 peanut oil, 35 cellulose, 35 AIN93 mineral mix, 10 AIN93 vitamin mix, 2.5 cholin, 100 saccharose, 124 lactose and 334 wheat flour.

Muscle protein synthesis was measured with the flooding dose method as described previously.

### Statistical analysis

All data are expressed as means  $\pm$  SEM. Food intake and body weight comparisons were assessed using repeated measures analysis of variance (StatView statistical software package, v. 5, SAS Institute, Cary, NC, USA). Other measurements were analysed using a two-way ANOVA (time, diet). When significant differences were detected by ANOVA, *post hoc* comparisons between groups were made using Fisher's PLSD test. Significance was defined at the  $P < 0.05$  level.

## Results

### Parameters of non-casted pair-fed animals

Food intake of the pair-fed group perfectly matched the one of the casted group during immobilization ( $11.29 \pm 0.06$  g day<sup>-1</sup> at I8) and recovery ( $17.86 \pm 1.09$  g day<sup>-1</sup> at R20). The body weight of pair-fed animals decreased slightly during the experimental protocol ( $-4.4\%$  at I8 and  $10.1\%$  at R20). However, gastrocnemius muscle mass from these rats was stable at each point measured ( $2.498 \pm 0.038$  g at R20 vs.  $2.553 \pm 0.042$  g at I8). Except the amount of polyUb conjugates which slightly increased at I8 compared to I0 ( $+30\%$ ,  $P < 0.05$  using a Student's *t* test I8 vs. I0), before returning to I0 value, all the others parameters did not change from I0 to the end of the recovery period (data not shown).

### Animal characteristics

Food intake was  $16.89 \pm 0.42$  g day<sup>-1</sup> (I0) and decreased by 65% the first day of cast-immobilization (Fig. 1A). Then, animals increased their food intake up to I8 ( $11.19 \pm 0.29$  g day<sup>-1</sup>). The nutritional intervention started the first day of the recovery period (R1). Food intake was similar in the two groups during the whole recovery period (Fig. 1A).

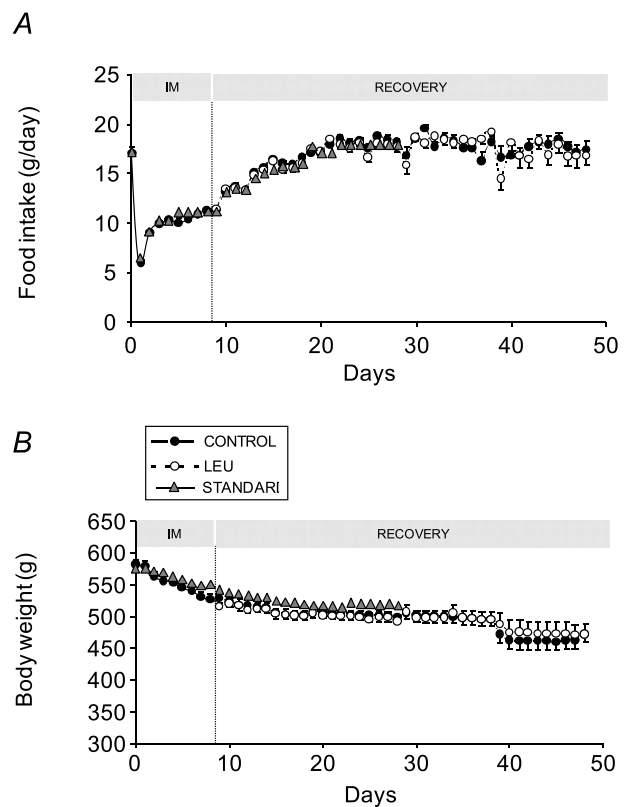
Casted rats exhibited a slight decrease of body weight during immobilization ( $-8.6\%$  at I8, not significantly different from the pair-fed group). The decrease of body weight continued during the recovery period whatever the experimental diet given ( $-18.8$  and  $-18.9\%$  in CON and LEU group respectively at I8) (Fig. 1B).

### Plasma amino acid concentrations

In rats, postprandial total amino acid concentrations and essential amino acid concentrations were not affected by the immobilization ( $1900.5 \pm 145.0$  and  $1995.0 \pm 144.8$   $\mu\text{mol l}^{-1}$  for total amino acid concentrations at I0 and I8, respectively) (Table 2). The nutritional intervention had no effect on total and essential amino acid concentration after 20 (R20) and 40 (R40) days of recovery. During this period, only plasma free leucine differed between the two diets. Indeed, leucine was markedly higher in the LEU group ( $\sim \times 2.5$  and  $\times 3.5$  at R20 and R40, respectively,  $P < 0.05$  LEU group vs. CON group).

### Effect of the leucine supplemented-diet on ubiquitin-proteasome-dependent pathways

Figure 2A shows an increased amount of polyUb conjugates at I8 ( $+70\%$ ,  $P < 0.05$  vs. I0) which normalized as soon as R10. No effect of the diet was observed during the recovery period. Both chymotrypsin- (Fig. 2B) and trypsin-like activities (Fig. 2C) were increased at I8 ( $+42$



**Figure 1. Food intake (A) and body weight (B) of casted and non-casted PF rats**

Food intake and body weight of casted and non-casted rats are similar during the whole experiment between the CON and the LEU groups and the STANDARD diet group. IM: immobilization period. Data are means  $\pm$  SEM.

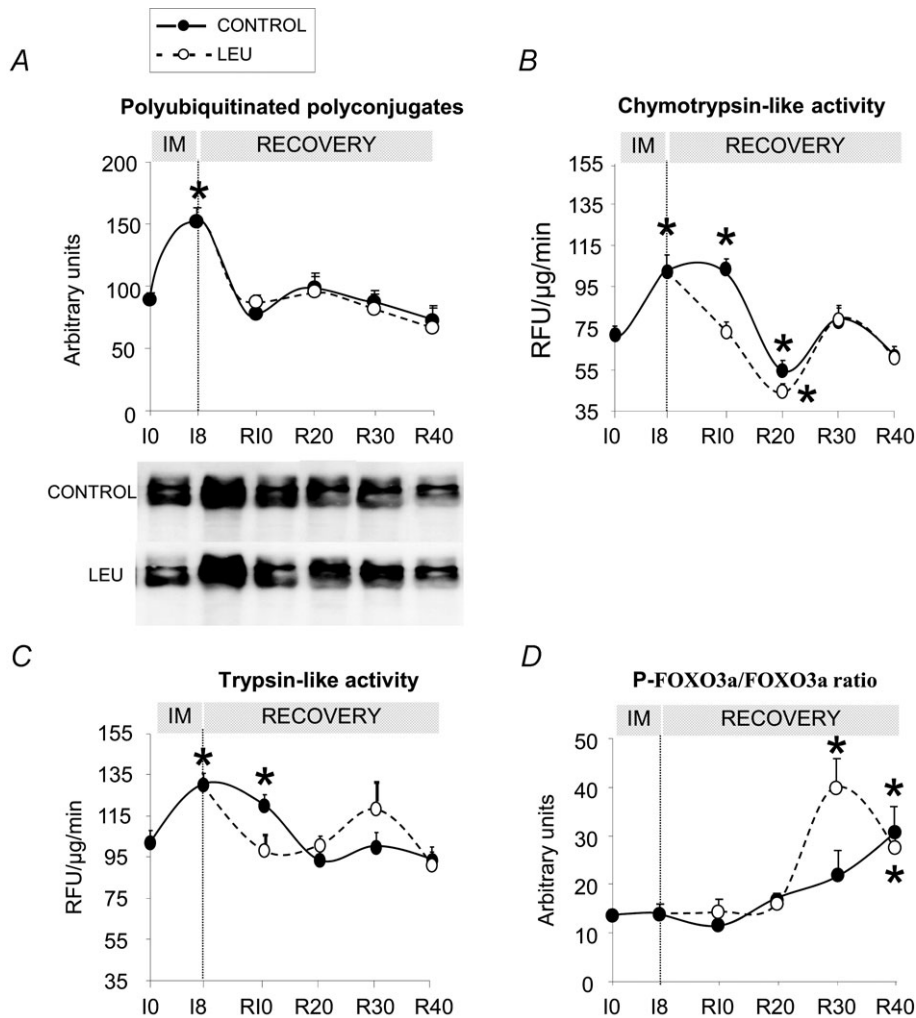
**Table 2. Plasma amino acid concentration in rats chronically fed casein or leucine in the postprandial state**

	I0	I8	R20 CAS	R20 LEU	R40 CAS	R40 LEU
Essential amino acids	859.5 ± 70.6	885.7 ± 52.2	766.0 ± 40.7	830.3 ± 66.1	722.8 ± 120.6	863.1 ± 66.8
Leucine	62.6 ± 2.7	78.6 ± 6.0	87.7 ± 7.1	211.7 ± 19.3*	60.8 ± 4.0	214.3 ± 27.0*
Total amino acids	1900.5 ± 145.0	1995.0 ± 144.5	1792.2 ± 95.5	1771.5 ± 173.4	1569.4 ± 208.6	1849.6 ± 131.8

Amino acid concentrations are presented at the postprandial before (I0) and after (I8) the immobilization period and also during the recovery period (R20 and R40; 10 and 40 days of recovery). Values explained in  $\mu\text{mol/L}$  are means  $\pm$  SEM. CAS: casein; LEU: leucine. \*  $P < 0.05$ , LEU vs. CAS.

and +27%, respectively,  $P < 0.05$  vs. I0). Both activities were still elevated at R10 within the CON group but were largely decreased with the leucine supplementation

( $\sim 18\%$  and  $\sim 30\%$  for chymotrypsin- and trypsin-like activity, respectively,  $P < 0.05$ , CON group vs. LEU group). Activities were then similar between the two



**Figure 2. Ubiquitin–proteasome-dependent proteolysis in immobilized gastrocnemius muscles before and after the free leucine supplementation.** A, accumulation of polyubiquitinated polyconjugates was assessed on 25  $\mu\text{g}$  of proteins by immunoblotting using an antibody that recognizes polyubiquitin chains. B and C, the chymotrypsin-like activity (B) and trypsin-like activity (C) of the proteasome were measured by using the fluorogenic substrate succinyl-LLVY-AMC and Boc-LRR-AMC as indicated in Methods. Data are expressed in relative fluorescence units ( $\text{RFU } \mu\text{g}^{-1} \text{ min}^{-1}$ ). D, phospho-FoxO3a/FoxO3a ratio traduces an anti-proteolytic potential. FoxO3a and its phosphorylated form phospho-FoxO3a (Ser253) were determined using appropriate antibodies on 50  $\mu\text{g}$  of proteins. IM: immobilization period; I0: before immobilization; I8: 8 days of casting; R10 to R40: 10 to 40 days of recovery. Data are means  $\pm$  SEM. As no effect of the meal was observed, postabsorptive and postprandial values are pooled. \*  $P < 0.05$ , vs. I0. Data are means  $\pm$  SEM.

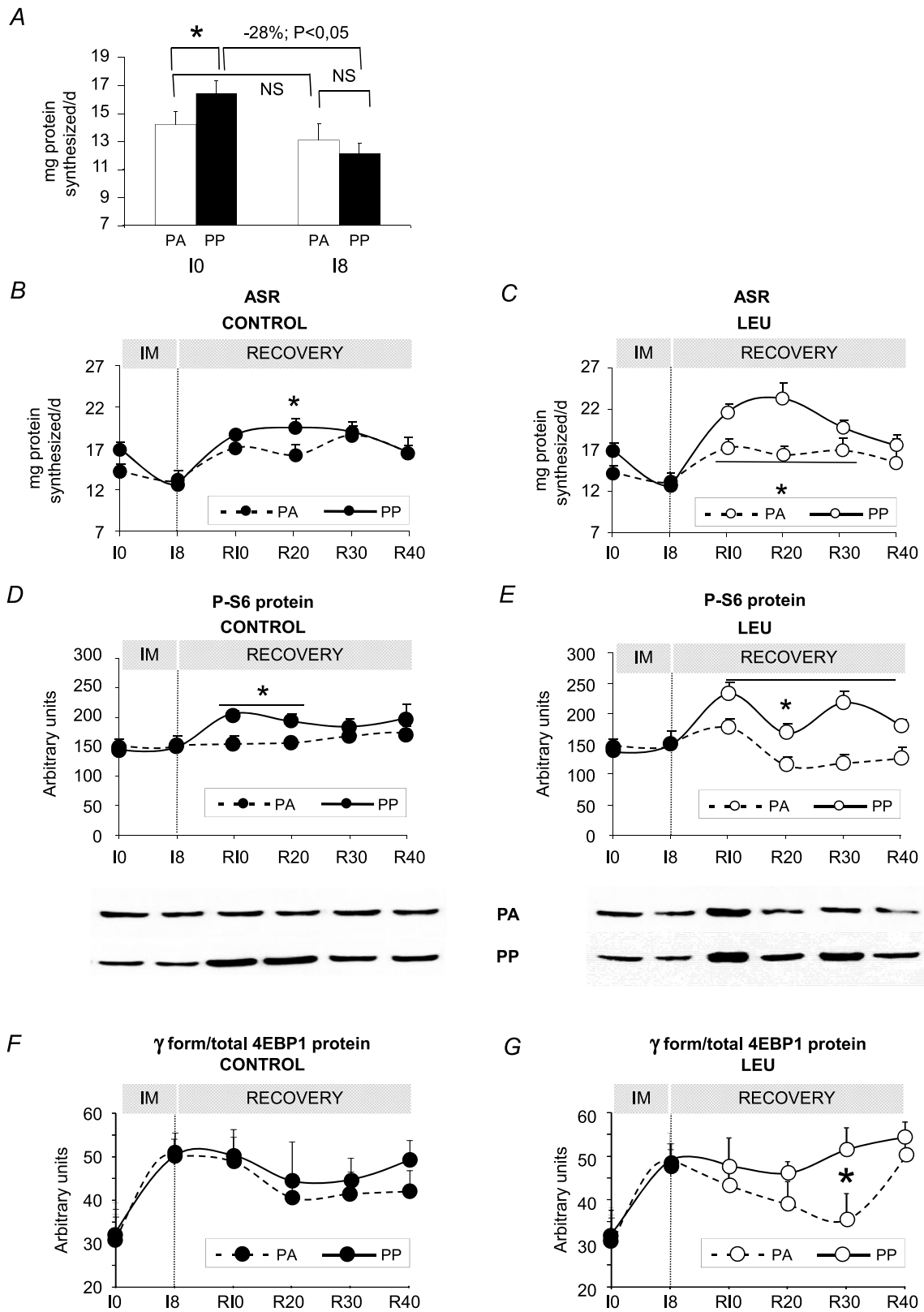


Figure 3. Muscle protein synthesis and protein pathway in immobilized gastrocnemius muscles before and after the free leucine supplementation



groups. phospho-FoxO3a/FoxO3a ratio was not different between the two diets until R20. At R30, the ratio was largely increased in the LEU group (+80%,  $P < 0.05$  vs. CON group) (Fig. 2D).

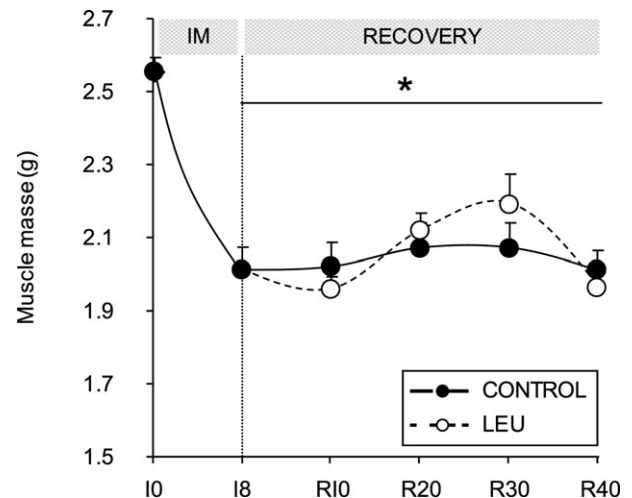
### Effect of the leucine supplemented diet on protein synthesis and protein synthesis pathway

At I0, before immobilization, protein synthesis increased slightly but significantly by 15.5% after food intake ( $P < 0.05$  PA vs. PP). However at I8, the stimulatory effect of food intake disappeared as protein synthesis was not different between the PA and the PP states ( $13.10 \pm 1.14$  and  $12.13 \pm 0.72$  mg day<sup>-1</sup> in the PA and the PP states respectively) (Fig. 3A) demonstrating an anabolic resistance during immobilization. Indeed, immobilization did not induce any significant modifications on the ASR in the PA state but largely decreased protein synthesis in the PP state (-28% at I8,  $P < 0.05$  vs. I0) (Fig. 3A). The stimulatory effect of food intake was only detectable at R20 in the CON group (Fig. 3B), whereas, this effect was present from R10 to R30 in the LEU group (Fig. 3C) ( $P < 0.05$ , PA vs. PP).

The amount of protein S6 phosphorylated is normally increased with the PA/PP transition reflecting an activation of the protein synthesis pathway. In the CON group this increase was only present at R10 and R20 (+30% and +24% respectively,  $P < 0.05$  PA vs. PP) (Fig. 3D). In the LEU group this increase was present during the whole recovery period, from R10 to R40 with a peak at R30 (+30%,  $P < 0.05$  PA vs. PP) (Fig. 3E). Similarly, the leucine supplementation induced a higher amount of  $\gamma$  form/total 4EBP1 protein (+44%,  $P < 0.05$  PA vs. PP) and no effect was observed with the control diet (Fig. 3F and G).

### Gastrocnemius muscle mass during immobilization and recovery

Immobilized gastrocnemius muscles atrophied by 20% at I8 ( $2.553 \pm 0.042$  g at I0 and  $2.013 \pm 0.059$  g at I8,  $P < 0.05$ ) (Fig. 4). The leucine supplementation, started after reloading and did not have any effect on muscle mass as muscle mass recovery was absent and muscle mass were



**Figure 4. Muscle mass of gastrocnemius after immobilization and after free leucine supplementation**

In immobilized muscles, muscle mass decreased after an 8 day immobilization period. The nutritional supplementation started after cast removal but muscle mass was not different between the CON group and the LEU group at R40 and never returned to its pre-immobilization value. IM: immobilization period; I0: before immobilization; I8: 8 days of casting; R10 to R40: 10 to 40 days of recovery. \* $P < 0.05$ , vs. I0. No significant differences between CON and LEU groups were recorded. Data are means  $\pm$  SEM.

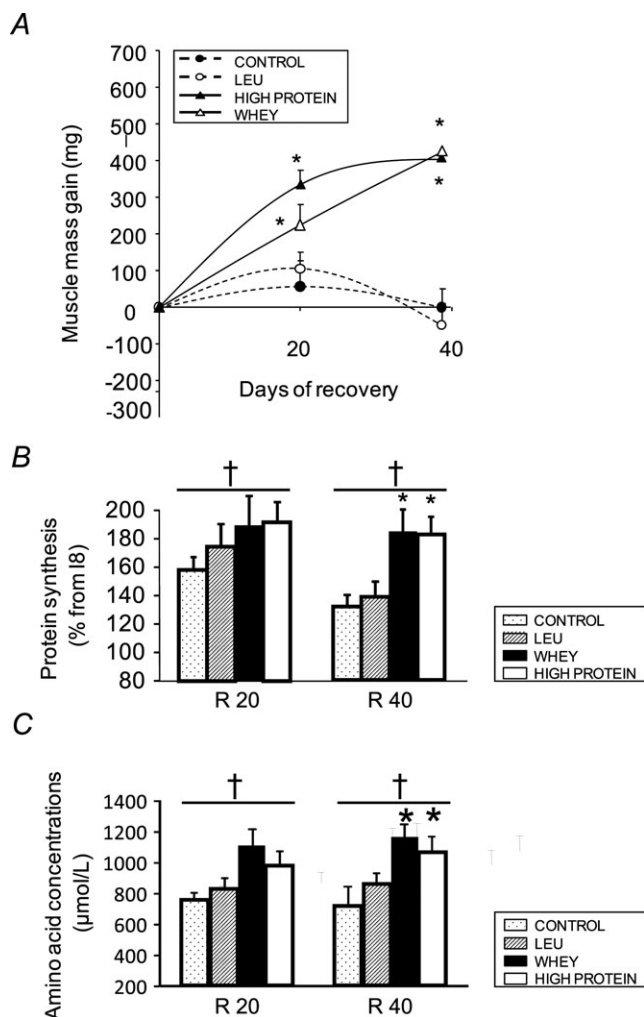
similar at R40 between the two groups ( $2.012 \pm 0.053$  g and  $1.963 \pm 0.059$  g in the CON group and the LEU group, respectively).

### Effect of whey diet and high protein diet supplementations on muscle mass recovery

Figure 5A shows the muscle mass gain following immobilization. As mentioned before, no muscle mass gain was obtained after 40 days in the CON or the LEU groups. However, within the WHEY and HIGH PROTEIN groups, a progressive muscle mass recovery appeared as soon as 20 days after cast removal, to achieve  $\sim 400$  mg at R40. At R20, the muscle mass gain was  $\sim 300$  mg within the HIGH PROTEIN group and only  $\sim 200$  within the WHEY group ( $P < 0.05$  vs. I8). The increase of postprandial muscle protein synthesis (% from I8) (Fig. 5B) was increased at R20 without any difference

A, muscle protein synthesis at I0 and I8 in the postabsorptive and the postprandial states for each time is expressed as the absolute synthesis rate (ASR), i.e. the amount of proteins synthesized in mg day<sup>-1</sup>. B and C, ASR during immobilization and recovery in the postabsorptive (B) and the postprandial states (C). D and E, protein S6 phosphorylation in gastrocnemius for the CON group (D) and the LEU group (E) in the postabsorptive and the postprandial states. The amount of protein S6 phosphorylated was assessed by immunoblotting on 30  $\mu$ g of proteins. F and G, amount of protein 4EBP1 expressed in arbitrary units as the ratio  $\gamma$  form/total forms in the CON group (F) and LEU group (G). IM: immobilization period; I0: before immobilization; I8: 8 days of casting; R10 to R40: 10 to 40 days of recovery; PA: postabsorptive state; PP: postprandial state. \* $P < 0.05$ , PA vs. PP. Data are means  $\pm$  SEM.

between the four diets. However, muscle protein synthesis at R40 was largely higher with WHEY and HIGH PROTEIN diets ( $\sim +30$ – $40\%$ ,  $P < 0.05$  vs. CON or LEU, Student's  $t$  test). Similarly, essential amino acid concentrations were greater at R40 with WHEY diet and HIGH PROTEIN diet ( $\sim +25$ – $70\%$ ,  $P < 0.05$  vs. CON or LEU) (Fig. 5C).



**Figure 5. Effect of the whey and high protein diets on muscle mass gain, muscle protein synthesis and amino acid concentrations**

A, muscle mass gain after the nutritional intervention is presented as the gain versus I8, in mg of muscle. Four diets were tested during the recovery period: a CON diet (13% casein), a LEU diet (13% casein + 4.45% Leucine), a WHEY diet (13% whey, i.e. leucine-rich and fast digested protein diet), and a HIGH PROTEIN diet (13% casein + 13% whey). \* $P < 0.05$ , vs. I8, Student's  $t$  test. Data are means  $\pm$  SEM. B, muscle protein synthesis is expressed as the percentage of the absolute synthesis rate (ASR), i.e. the amount of proteins synthesized in  $\text{mg day}^{-1}$ , at I8. C, plasma amino acid concentrations expressed in  $\mu\text{mol l}^{-1}$  were determined on animals fed each experimental diet at R20 and R40. I8: 8 days of casting; R20 and R40: 10 and 40 days of recovery; LEU: leucine. \* $P < 0.05$ , vs. CON and LEU; † $P < 0.05$ , vs. I8.

## Muscle fibre composition

Table 3 presents the percentage of each major fibre types in gastrocnemius muscles before immobilization, after immobilization and during recovery. There was no effect of immobilization on the composition of muscle fibre types and no diet induced modification of the percentage of fibre types.

## Discussion

In this study, we have demonstrated that during ageing, muscle protein synthesis was depressed in the postprandial state by the period of disuse and only normalized during reloading. As proteolysis was also only normalized during the recovery period, the resultant nitrogen balance was then not positive enough for muscle protein gain, hence contributing to the age-related incomplete muscle mass recoveries. We also demonstrated that a dietary free leucine supplementation had no effect on muscle mass gain despite its positive anabolic effect on muscle protein metabolism during the first 20 days of the recovery period, contrarily to high protein or whey protein diets.

Sarcopaenia is the progressive loss of muscle mass and strength related to normal ageing (Rosenberg, 1989; Cruz-Jentoft *et al.* 2010). One of the mechanisms leading to this muscle mass loss could be the impaired muscle mass recovery following acute catabolic states observed in aged animals (Dardevet *et al.* 1995; Mosoni *et al.* 1995; Chakravarthy *et al.* 2000; Magne *et al.* 2011) or elderly humans (Suetta *et al.* 2009). This was recently named 'the catabolic crisis model' by English & Paddon-Jones (2010). At a cellular level, we have previously shown that an 8 day immobilization period by casting induced an increase of muscle proteolysis, i.e. activation of the ubiquitin-proteasome-dependent pathway (Magne *et al.* 2011). We have completed these observations by showing in the present study a large depression of protein synthesis in the postprandial state. This alteration could explain a large imbalance of protein metabolism, leading therefore to protein loss and ultimately muscle mass loss. During reloading we observed a normalization of protein breakdown and only a slight increase of protein synthesis above those recorded before immobilization. These phenomena may result in an insufficient positive nitrogen balance that is normally required to recover muscle mass. Immobilized muscles exhibited a decrease in responsiveness of muscle protein synthesis to anabolic stimuli (i.e. amino acids or insulin) (Phillips *et al.* 2009). If this impaired response of protein synthesis has already been observed during immobilization in the adult (Ferrando *et al.* 1996; Paddon-Jones *et al.* 2006; de Boer *et al.* 2007; Glover *et al.* 2008), this is the first study to demonstrate this 'anabolic resistance' during ageing and reloading. We and others have observed in the immobilized muscle

**Table 3. Effect of immobilization and diets on muscle fibres composition**

		Type IIA	Type IIX	Type IIB	Type I
I0		0.2 ± 0.2	38.1 ± 1.5	57.7 ± 1.8	3.9 ± 0.6
I8		0.4 ± 0.3	37.8 ± 1.5	58.5 ± 2.0	2.8 ± 0.9
R20	CON	ND	42.9 ± 2.2	53.0 ± 2.4	4.4 ± 0.9
	LEU	ND	41.4 ± 2.2	56.5 ± 1.8	3.8 ± 0.8
	WHEY	ND	41.4 ± 2.7	53.2 ± 3.3	5.3 ± 0.9
	HIGH PROTEIN	ND	35.2 ± 1.4	60.7 ± 1.7	4.1 ± 0.9
R40	CON	ND	41.2 ± 3.1	54.1 ± 3.3	4.7 ± 0.5
	LEU	ND	38.6 ± 2.5	58.1 ± 3.0	3.3 ± 0.7
	WHEY	ND	38.3 ± 0.7	56.1 ± 2.1	5.6 ± 1.4
	HIGH PROTEIN	ND	36.7 ± 2.3	58.8 ± 2.1	4.5 ± 0.2

Muscle fibres composition has been assessed by electrophoresis. The composition in the 4 major types (type I, type IIA, type IIX, type IIB) is presented in % of total fibres content at I0, I8, R20 and R40 for the 4 different diets tested. CONTROL diet = 13% casein, LEU diet = (13% casein + 4.45% Leucine), WHEY diet = 13% whey, HIGH PROTEIN diet = (13% casein + 13% whey), I8: 8 days of casting; R10 and R40: 10 and 40 days of recovery. ND: non detectable. Data are means ± SEM.

the presence of an inflammation (Zarzhevsky *et al.* 2001; Magne *et al.* 2011) which may be responsible for this alteration of the protein synthesis pathway. Indeed, inflammation is known to be deleterious for protein synthesis during ageing (Lang *et al.* 2002; Balage *et al.* 2009) and the immobilization-induced inflammation may worsen this phenomenon, hence contributing to altered protein synthesis.

To generate the positive nitrogen balance required for protein accretion during recovery, a further increased protein synthesis, decreased proteolysis, or simultaneous changes in both processes is required. If exercise is the best way to induce these modifications to initiate muscle mass gain, it is not always possible particularly after prolonged immobilization periods in elderly people. Thus, as protein intake is a robust stimulator of muscle protein synthesis (Kimball *et al.* 2000; Bohe *et al.* 2003; Moore *et al.* 2009), we focused our work on this aspect. Amino acids and particularly leucine have been shown to be efficient in the stimulation of postprandial muscle protein synthesis (Anthony *et al.* 1999, 2001; Koopman *et al.* 2005). In adults, some studies have used a mix of branched amino acids during the immobilization period (Stein *et al.* 1999; Paddon-Jones *et al.* 2004; Trappe *et al.* 2007), but few studies have evaluated the isolated effect of leucine supplementation on muscle mass recovery after disuse situations. In 2010, Baptista *et al.* supplemented diets with free leucine and showed that these diets given during immobilization could attenuate the muscle mass atrophy (Baptista *et al.*, 2010). Up to now, no study has explored its potential effect during the reloading in adult or aged animals or humans. This study was conducted to test the effect of a free leucine supplementation which is

also known to stimulate protein synthesis by reactivating specific kinases implicated in pathways involved in the regulation of translation initiation, particularly during ageing (Dardevet *et al.* 2002; Rieu *et al.* 2003, 2006, 2007; Combaret *et al.* 2005; Katsanos *et al.* 2006). Its action on muscle protein metabolism by stimulating the m-TOR pathway has been well demonstrated (Anthony *et al.* 2000; Dardevet *et al.* 2000; Bolster *et al.* 2004) and these leucine effects could be maintained when administered chronically (Lynch *et al.* 2002; Rieu *et al.* 2003). In our study, a free leucine supplemented diet given during rehabilitation had a positive effect on the muscle protein synthesis. Indeed, free leucine supplementation induced an increased amount of phosphorylation of S6 and 4EBP1 proteins, but also a decrease of chymotrypsin- and trypsin-like activities of the 26S proteasome associated to a phospho-FoxO3a/FoxO3a ratio in favour of anti-proteolytic process. This resulted in a higher protein synthesis at the postprandial state with leucine and a proteolysis normalized earlier. However, surprisingly, it failed in the muscle mass recovery.

Our measures of muscle protein synthesis were done at a single time point postprandially and confirmed that our free leucine supplementation was efficient 150–180 min after food intake. As muscle mass gain was absent, we hypothesized that the stimulation of protein synthesis was nevertheless not sustained enough to translate into a significant muscle protein accretion. We postulated that free leucine, as it is absorbed rapidly, induced an anabolic 'leucine signal' for protein synthesis before there was sufficient availability of amino acids coming from dietary protein digestion. Therefore, the duration or the intensity of the protein synthesis could have been insufficient. This

desynchronization between the 'leucine signal' and the availability of substrates for protein synthesis could explain the insufficient positive nitrogen balance and the lack of protein accretion when free leucine is used.

To verify this hypothesis we improved our nutritional strategy by resynchronizing both the 'leucine signal' and the amino acid availability. The strategy was (1) to change the quality of proteins to use rich-leucine and fast digested proteins to induce simultaneously the 'leucine signal' and a large increase of amino acid concentration and (2) to increase the quantity of proteins to provide amino acids for a longer time during the postprandial state. We then used soluble milk proteins (i.e. rich-leucine and fast digested proteins) and high protein diet (i.e. casein and whey proteins), respectively. Indeed, soluble milk proteins are known to be digested faster than casein, i.e. they have a rapid absorption kinetic, but also high leucine content (Boirie *et al.* 1997; Dangin *et al.* 2001, 2003). Whey proteins have already demonstrated their positive effect in re-stimulating muscle protein accretion in defective protein synthesis stimulation during ageing (Pennings *et al.* 2011). A high protein diet, composed of one-half casein and the other half of soluble milk proteins, combined the properties of whey (i.e. rich-leucine and fast digested proteins) and mass effect (high quantity of amino acids for a long time; Boirie *et al.* 1997). We have shown in the present study that these two diets had a positive impact on muscle mass gain, inducing a gain of ~60% of the total muscle mass loss. These two diets induced a greater protein synthesis compared to casein or free leucine-supplemented diets, associated to higher amino acid concentrations. These observations confirm data presented in the recent review of Breen & Phillips (2011), which suggest that protein-based interventions could be a good way to induce hypertrophy, particularly during ageing.

As sarcopaenia and immobilization affect muscle mass but also muscle functionality, we proposed to evaluate this functionality by measuring the muscle fibres composition. We did not record any modification of muscle fibre composition with immobilization. This could be explained by (1) a low composition of slow fibres, which are mainly affected by immobilization, and (2) a short period of (8 days), as some authors demonstrated that 10–21 days of casting are required to induce a fibre-type switch in rats (Pattison *et al.* 2003; Coutinho *et al.* 2004). Moreover, primary effects of immobilization could be a reduction of cross-sectional area (Tomanek & Lund 1974; Nicks *et al.* 1989; Berg *et al.* 1997) before modifications of fibre type composition.

In summary, we have demonstrated here for the first time that the lack of recovery after immobilization-induced atrophy during ageing is due to an 'anabolic resistance' of protein synthesis to amino acids during rehabilitation. Our nutritional

study showed that chronic leucine supplementation, despite a greater postprandial protein synthesis response, failed in muscle mass gain. Furthermore, this work provides an insight into the mechanisms involved in this discrepancy: our data strongly suggest the presence of a desynchronization between the stimulatory 'leucine signal' and the availability of amino acid substrates for protein synthesis. Our results highlight a novel approach to induce muscle mass recovery following atrophy in the elderly by giving soluble milk protein or high protein diets.

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### Author contributions

H.M., I.S.A. and D.D. contributed to the conception and design of the experiments. H.M., I.S.A., C.M., M-A.P., L.C., D.R. and D.D. participated to the collection, analysis and interpretation of the data and revised the manuscript critically for important intellectual content. All authors drafted the manuscript and approved the final version for publication. All experiments were performed in the Unité de Nutrition Humaine from the Institut National de la Recherche Agronomique (Saint Genès Champanelle, France).

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