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DETERMINANTS OF INTRA-INDIVIDUAL VARIATION IN ADAPTABILITY TO RESISTANCE TRAINING OF DIFFERENT VOLUMES WITH SPECIAL REFERENCE TO SKELETAL MUSCLE PHENOTYPES

Determinants of intra-individual variation in adaptability to resistance training of different volumes with special reference to skeletal muscle phenotypes

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THESIS FOR DOCTORAL DEGREE (Ph.D.)

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Abstract

The preface pretty much says it all.

Second paragraph of abstract starts here.

List of scientific papers

I. **Hammarström D**, Øfsteng S, Koll L, Hanestadhaugen M, Hollan I, Apró W, Blomstrand E, Rønnestad B, Ellefsen S Benefits of higher resistance-training volume are related to ribosome biogenesis. The *Journal of physiology.* 2020;598(3):543-65.

II. Khan Y, **Hammarström D**, Rønnestad B, Ellefsen S, Ahmad R Increased biological relevance of transcriptome analyses in human skeletal muscle using a model-specific pipeline. *Submitted.*

III. **Hammarström D**, Øfsteng S, Koll L, Jacobsen N, Flobergseter K, Rønnestad B, Ellefsen S Ribosome accumulation during early phase resistance training. *Manuscript*

IV. **Hammarström D**, Ellefsen S. generefer: A R package for unbiased selection of reference genes for qPCR in repeated measures designs. *Manuscript*

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1. Introduction

Skeletal muscle health is essential for physical independence. In a lifespan perspective, measures of muscle mass and/or strength are inversely associated with mortality (1–6) and disability (7). Besides adverse associations between of low muscle mass and strength and clinical conditions, muscle weakness also accounts for increased health care costs in patient populations (8,9). The intercept between muscle mass, muscle function and health status is interrelated with variables such as age and primary illness or injury (10). This highlights that interventions designed to increase muscle mass and strength are likely to prevent adverse health outcomes across the lifespan. A higher level of muscle mass and functional capacity would counteract the effects of muscle loss due to illness, age or inactivity.

Although a large degree of the observed variations in lean mass and strength are attributed to genetic components (11,12), environmental factors also contribute, leaving a window of opportunity to increase muscle mass and functional capacity. Among factors affecting muscle mass and functioning are nutrition and pharmacological agents. However, physical activity and specifically systematic resistance training of sufficient volume, intensity and frequency provides a stimulus that promote morphological and functional changes to the human neuromuscular system without adverse side-effects. Irrespective of age, resistance training generally leads to increased muscle mass and strength (13,14) and is considered safe when performed in a well organized manner (14,15).

Resistance training can be modulated indefinitely through combined variations of training variables such as frequency, intensity and volume (16,17). Well designed training prescriptions should incorporate information about the current state and goals of the trainee to maximize the potential outcome of the training program (16–18). Training volume has received particular attention in the scientific community for many reasons. Evidence suggests that exercise volume affects selected molecular determinants of muscle hypertrophy in a dose-dependent manner (19–21). Such effects are believed to facilitate long-term training effects as training programs with higher volume generally result in higher gains in muscle mass and strength

with little evidence of differences between age groups or participants with different training backgrounds (22–24).

A consequence of a more extensive training program is the increased time required to complete such a program. As time constraints has been reported as a limiting factor for engaging in physical activity (25) some merit can be given to arguments against guidlines suggesting higher volume in resistance training prescription (18,26). From an individual perspective, training prescription that balances time-requirement with efficacy presumably increases the likelihood of participation in physical activity (25). From a more general perspective, increased knowledge about mechanisms governing responses to physical training could improve training prescription also for individuals and populations that experience attenuated benefit of resistance training (27). The overreaching goal of the present thesis is to contribute to understanding individualized training loads. To this end, training volume was used to study the effects of variable training stimulus in within-participant models of exercise-training.

2. Background

2.1 Exercise training variables affecting training outcomes

2.2 Exercise volume

2.2.1 Meta-analysis of exercise volume

2.3 Molecular determinants of training-induced muscle hypertrophy

2.3.1 Protein synthesis

A single exercise session promotes protein synthesis Positive net protein balance in responce to exercise

Inhibition of RNA synthesis restrict protein synthesis

Indicated in Goldspink 1977 and 1976 RNA reflects ribosomal availability

Protein synthesis is proportional to RNA content

Increase loading leads to increased RNA

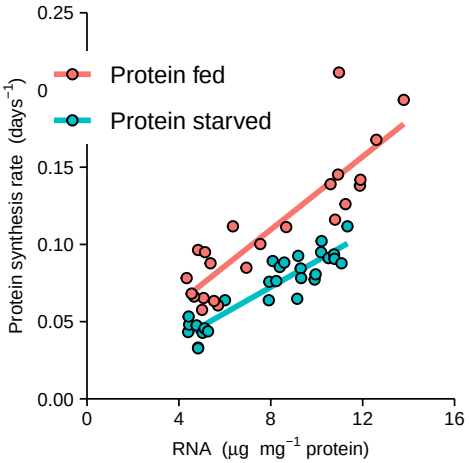


Figure 2.1: Data from Millward et al. 1973. Group A were fed a diet containing protein, group B were starved or fed a diet not containing protein.

2.3.2 The mammalian target of rapamycin (mTOR) and translational efficiency

The mammalian target of rapamycin (mTOR) is a large serine-threonine protein kinase wich in complex with oyher regulatory proteins forms a signaling hub responsible for responses to environmental cues such as nutrients and mechanical stress.

mTOR has several phosphorylation sites

Phosphorylation of Ser2448 is mediated by S6K1 to reduce mTOR activity in a negative feedback loop .

Ser2448 is phosphorylated by S6K1, changes in nutrient avaliability modifies S6K1 and Ser2448, Ser2448 phosphorylation is abolished when S6K1 is depleted

When the C-terminal is deleted, mTOR gets constitutively active

2.4 Ribsome biogenesisand muscle growth

Crossland et al. noted that using IGF-1 induces specific growth related effects in muscle cells. These may be different than the effects induced by sreum as in Stec et al.

Brook 2016 Stec 2016 Nakada 2016 Figueriedo 2015

Millward 1973 correlation between RNA and protein synthesis

2.4.1 Ribsome biogenesis

Transcription of ribsomal RNA (rRNA)

2.5 Transcriptional activity related to muscle hypertrophy

2.5.1 Methods for studying transcriptional regulation

3. Aims

The primary aim of this thesis was to relate the adaptive response to resistance training with low- and moderate-volume to skeletal-muscle characteristics in previously untrained individuals. The key question was whether manipulation of exercise-volume will have diverse effects in different individuals related to muscular intrinsic characteristics. A further aim was to characterize exercise-volume dependence and time course profiles of molecular mechanism thought to control resistance training-induced muscle growth. Based on these aims, the objectives of the present thesis were;

- to relate skeletal muscle and systemic characteristics to benefit of moderate- compared to low-volume resistance training;
- To determine volume-dependence in molecular networks related to muscle growth and remodelling in response to mechanical stress
- To determine a time course of markers related to ribosome biogenesis in the early phase of resistance training.

4. Methods

4.1 Study participants, protocols and training interventions

Study I was designed to examine effects of low- and moderate-volume on responses to acute exercise and long-term training within participants. Forty-one healthy individuals were recruited and 34 of these completed at least 85% of the prescribed sessions and were thus included in subsequent data analyses. Reasons for not completing the trial included injury not related to the study ($n = 1$), pain or discomfort during exercises ($n = 5$) and non-adherence to the study protocol. There were no differences in characteristics between participants included in or excluded from data analysis in Study I. Study II was designed to study the effects of resistance training *per se* and effects of variable volume on selected markers related to ribosome biogenesis. Participants were therefore recruited to a training group ($n = 11$) and a non-training control group ($n = 8$). Eligible for participation were young (Study I 18-40; Study II 18-35), non-smoking men and women. Exclusion criteria included a training history of more than one weekly session during the last 12 (Study I) or six (Study II) months leading up to the study. Participants were also screened for intolerance to local anesthetic, current or previous injuries affecting their ability to perform resistance training, self-reported symptoms or history of disease, intake of medication or supplements with known effects on adaptations to training. Participant characteristics for both studies are shown in Table 4.1. Each training session started with a light standardized warm-up (5 min ergometer cycling and 10 repetitions each of push-ups, sit-ups, back-extensions and squats). Before each exercise in the main program, one set of 10 repetitions were performed in the specific exercise with approximately 50% of 1RM.

Both studies were fully or partially performed as within-participant studies as each participant had their legs assigned to different training conditions (not including the control group in Study II). Allocation was performed after enrollment where each participant had their legs randomized to either low- or moderate volume

Table 4.1: Participant characteristics

		Sex	Age (years)	Stature (cm)	Mass (kg)	Fat mass (%)	Lean mass (%)
Study I	Included	Female	22.0 (1.3)	168 (7)	64.4 (10.4)	34.1 (5.6)	64.3 (6.2)
		Male	23.6 (4.1)	183 (6)	75.8 (10.7)	20.4 (6.0)	79.3 (5.0)
	Excluded	Female	22.9 (1.6)	166 (8)	64.6 (9.7)	28.8 (8.7)	68.6 (9.1)
		Male	24.3 (1.5)	189 (5)	88.2 (22.4)	24.3 (15.3)	76.8 (12.7)
Study II	Training	Female	23.4 (2.9)	168 (8)	64.0 (9.2)	30.8 (7.1)	65.5 (6.8)
		Male	25.7 (5.8)	177 (3)	77.5 (8.0)	25.3 (3.9)	71.3 (2.4)
	Control	Female	24.1 (3.5)	166 (4)	63.8 (0.6)	30.5 (6.4)	66.3 (5.2)
		Male	25.5 (5.5)	182 (5)	76.5 (7.7)	18.2 (5.1)	78.7 (4.2)

Data are means and (SD)

(Study I), or variable or constant volume (Study II).

In Study I, the low-volume protocol consisted of a single set of each exercise and the moderate-volume consisted of three sets per exercise. Three unilateral leg exercises were used (leg press, leg curl and knee extension). The moderate volume-leg commenced all sessions and the low volume-leg performed a single set of each exercise in the rest between second and third set of the moderate volume training protocol.

In Study II, only unilateral knee-extension was performed in an effort to concentrate the stimulus to the quadriceps muscles. The constant-volume leg performed six sets of 10RM throughout the study and variable leg performed six sets in session one to four, three sets in session five to eight and nine sets in session nine to twelve with same intensity (10RM).

4.1.1 Ethical considerations

Both studies were approved by the local ethics committee Lillehammer University College/Inland Norway University of Applied Sciences and the Norwegian Centre for Research Data. In accordance with the *Declaration of Helsinki*(28) the studies were pre-registered in publicly accessible databases (Study I, ClinicalTrials.gov Identifier: NCT02179307; Study II, `https://osf.io/wa96y`). Participants were informed of the study design, potential risks and sources of discomfort prior to giving their informed consent.

4.2 Measures of muscle mass

In Study I muscle mass was measured by magnetic resonance imaging (MRI) and dual energy X-ray absorptiometry (DXA) prior to and after the intervention. Both MRI and DXA measurements were completed during the same visit to the laboratory. Participants were instructed to refrain from strenuous physical activity during the last 48 h leading up to the measurements. The post-training measurements were completed at least 48 h after the last strength testing session. Participants were asked to refrain from food consumption during 2 h leading up to the measurements.

MRI images were obtained from the mid-thigh and analyzed by the same investigator blinded for time (pre- and post-training) and condition (low- and moderate-volume). Multiple images were used to estimate the cross-sectional area of the extensor muscles at the same distance from the knee-joint.

See figure

Dallin et al. recently estimated the (29)

4.3 Muscle strength assessments

Muscle strength was with

4.4 Blood variables

4.5 Muscle tissue sampling and preparations for downstream analyses

Muscle samples were obtained under local anesthesia (Study I, Xylocaine, $10\,\mathrm{mg\,ml^{-1}}$ with adrenalin $5\,\mu\mathrm{g\,ml^{-1}}$, AstraZeneca, Oslo, Norway; Study II, Lidocaine Mylan, $10\,\mathrm{mg\,ml^{-1}}$, Mylan Ireland Ltd, Ireland) with a fine needle (12-14 gauge; Universal-plus, Medax, Italy) operated with a spring-loaded instrument (Bard Magnum, Bard Norway AS, Norway). Sampling was performed as previously described (30), with modifications. Anestheisa was injected in the subcutanous tissue with care taken not to inject anesthesia into the muscle itself. Following a short period (5 min) the effect of the anesthesia was confirmed using an injection needle. Following pilot experiments we decided not to use an insertion cannula as described in (30) as the biopsy needle itself could be used to puncture the skin

and muscle fascia. This also resulted in less discomfort. Several passes through the same skin puncture was made to obtain sufficient material for downstream analyses. A smaller needle (14 vs. 12 gauge) was used to further minimized discomfort in Study II where more biopsies were sampled over a shorter time span, with exception from when material was used for immunhistochemistry. The first biopsy was sampled at one third of the distance between the patella to the *anterior superior iliac spinae* with subsequent biopsies sampled ∼2 cm proximal to previous samples. In Study II samples obtained more than one week apart were sampled with closer proximity and distally from previous samples but never at previous sampling sites.

The microbiopsy technique produces smaller samples compared to other biopsy techniques (31), and thus requires several passes to produce sufficient material for multiple downstream experiments. However, reports confirms that the microbiopsy technique is comparable to the traditionally used Bergström technique in several measures of muscle characteristics at the same time as being well tolerated (30,32). Any reported differences in fiber type distributions between sampling techniques have been suggested relating to differences in sampling depth (32,33).
For determination of fiber type distributions, a threshold of 200-300 fibers has been suggested as a suitable sample size per specimen as more fibers does not reduce the variation between dupliacte samples (34). In Study I one or several pieces of muscle (total weight ∼15 mg) were chosen per sampling for analysis of fiber type distributions (described in detail below). The total number of fibers were counted from these specimens (Figure ref fig). Using an average of fibers from the first sampling time point the between leg coefficient of variation was determined to 14% for Type I fibers and 11.3 for type II fibers. The between leg variation in Type I fibers is similar to what has been previously reported [Blomstrand Ekblom]

Latest paper on variability between samples from the same leg and due to number of fibers counted Appl Physiol Nutr Metab . 2020 Apr;45(4):368-375. doi: 10.1139/apnm-2019-0263. Epub 2020 Mar 24.

The within-leg was similar to between leg comparison in our samples. This might highlight sampling depth variation in microbiopsy technique in immunohistochemistry ?

To calculate variation in proportions

`https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3875499/`

```
Using ',' as decimal and '.' as grouping mark. Use read_delim() for more control

Parsed with column specification:
```

```
cols(
  subject = col_character(),
  multiple = col_character(),
  single = col_character(),
  sex = col_character(),
  include = col_character()
)


Joining, by = c("subject", "leg")
Joining, by = c("subject", "leg")
Joining, by = c("subject", "leg")
```

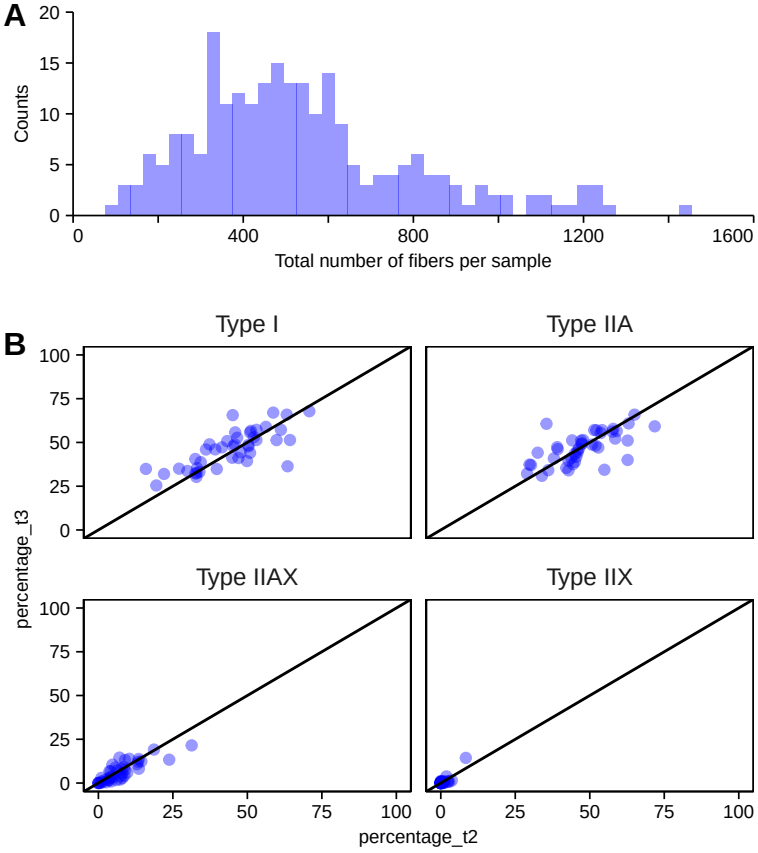


Figure 4.1: Number of fibres in immunohistochemistry analyses

4.6 Gene expression analysis

4.7 Determination of protein abundance

4.8 Statistics and data analysis

TO DO:

- For methods discussion, compare product length, efficiencies and ct values in relation to RQI-values. See Fleige 2006 for reference.

4.9 Gene expression analysis

4.9.1 Normalization

- An external reference gene was added at a constant amount in Trizol preps
- A normalization factor was used to express relative target gene abundance per-weight tissue.
- In qPCR the linearised expression (effectivety ^cq) was used to express the fraction of external reference per total RNA.
- In RNA-seq the external reference gene was sequenced and counts were used to express external RNA as a fraction of total RNA.
- In both cases the normalization factor was calculated as mw * counts.

A simulation to see that this is equivalent to tissue used in prep when no measurement errors exists.

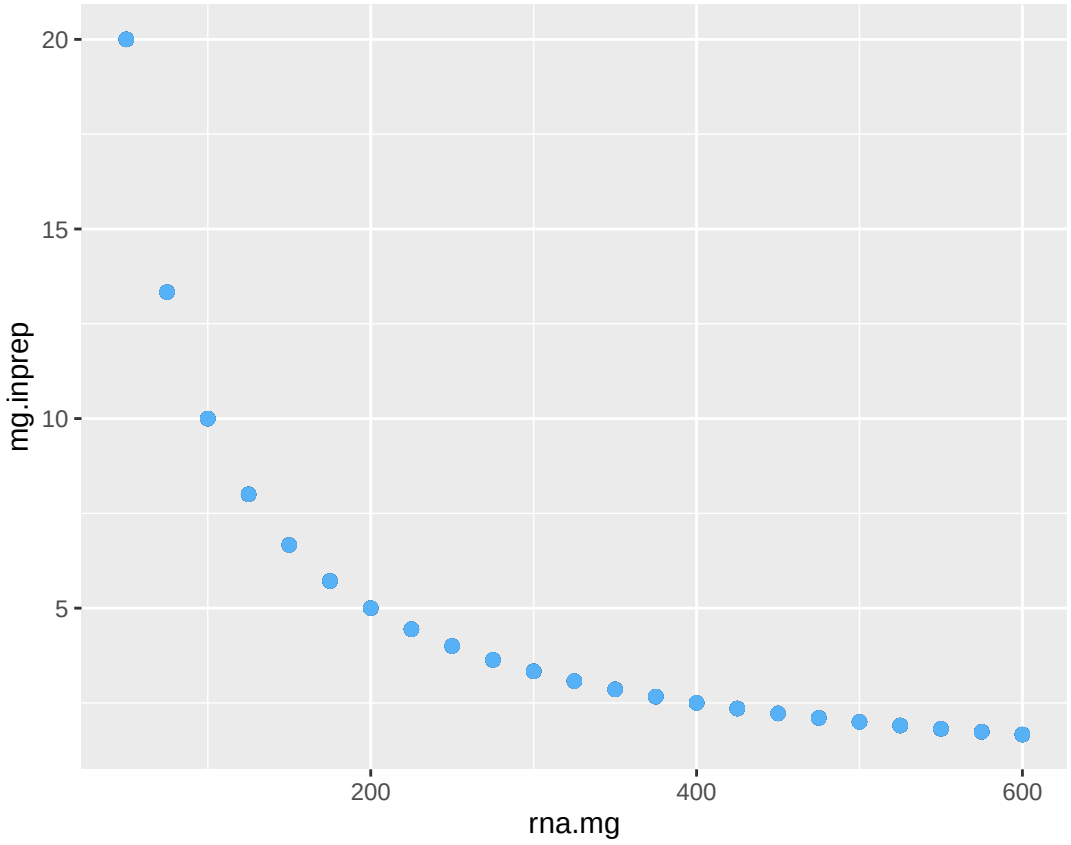
```
library(tidyverse)

expand_grid(mg = seq(from = 5, to = 100, by = 5),
            rna.mg = seq(from = 250, to = 600, by = 25),
            ext = 0.04) %>%
  mutate(tot.rna = mg * rna.mg,
         ext.frac = ext / (ext + tot.rna),
         mg.inprep = 1000 / ((ext + tot.rna) / mg),
         nf = ext.frac * mg)
```

```
# A tibble: 300 x 7
      mg rna.mg   ext tot.rna  ext.frac mg.inprep        nf
```

```
   <dbl>  <dbl> <dbl>   <dbl>     <dbl>     <dbl>     <dbl>
 1     5    250  0.04    1250 0.0000320      4.00 0.000160
 2     5    275  0.04    1375 0.0000291      3.64 0.000145
 3     5    300  0.04    1500 0.0000267      3.33 0.000133
 4     5    325  0.04    1625 0.0000246      3.08 0.000123
 5     5    350  0.04    1750 0.0000229      2.86 0.000114
 6     5    375  0.04    1875 0.0000213      2.67 0.000107
 7     5    400  0.04    2000 0.0000200      2.50 0.000100
 8     5    425  0.04    2125 0.0000188      2.35 0.0000941
 9     5    450  0.04    2250 0.0000178      2.22 0.0000889
10     5    475  0.04    2375 0.0000168      2.11 0.0000842
# ... with 290 more rows
```

```
expand_grid(mg = seq(from = 5, to = 100, by = 5),
            rna.mg = seq(from = 50, to = 600, by = 25),
            ext = 0.04) %>%
  mutate(tot.rna = mg * rna.mg,
         ext.frac = ext / (ext + tot.rna),
         mg.inprep = 1000 / ((ext + tot.rna) / mg),
         nf = ext.frac * mg) %>%
         ggplot(aes(rna.mg, mg.inprep,  color = mg)) + geom_point(size = 2)
```



4.10 Training protocols

A full body protocol was used in study I including

5. Results and Discussion

5.1 Effects of different training volume on changes in muscle size and function

Average within participant differences in responses to low- and moderate volume were consistent across measures of muscle hypertrophy and strength gains. Taken together these indicate that MOD was more efficient than LOW in increasing muscle size and strength (Figure 5.1).

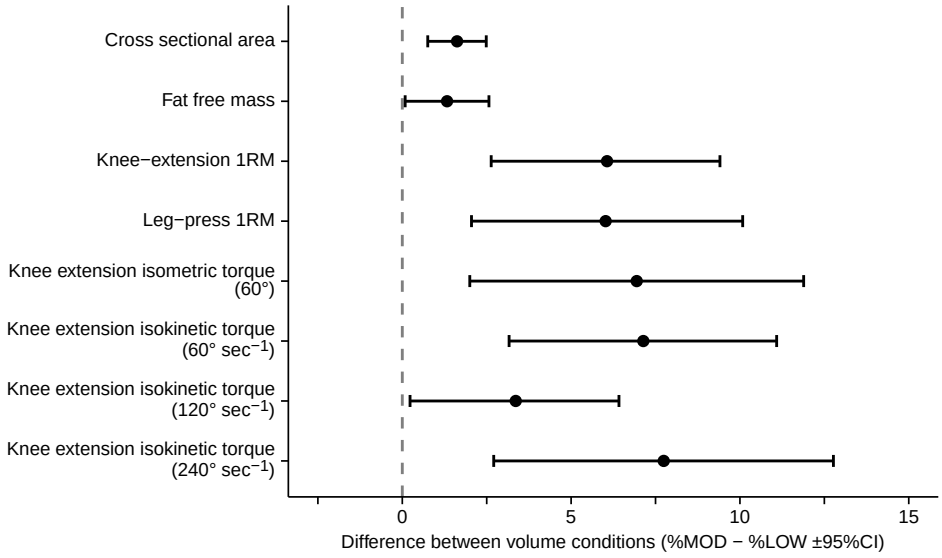


Figure 5.1: Differences in training induced relative changes in muscle mass and strength measures. Estimates are derived from ANCOVA models controling for baseline values and sex.

6. Discussion

Conclusion

If we don't want Conclusion to have a chapter number next to it, we can add the `{-}` attribute.

More info

And here's some other random info: the first paragraph after a chapter title or section head *shouldn't be* indented, because indents are to tell the reader that you're starting a new paragraph. Since that's obvious after a chapter or section title, proper typesetting doesn't add an indent there.

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