



The Effect of Various Types of Exercise on Gut Microbiota: A Systematic Review



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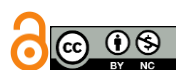
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ABSTRACT

Objective: Gut microbiota composition plays a pivotal role in health and emerging evidence supports the ability of exercise training programs to alter the gut microbiota's composition and function, which could counteract dysbiosis and the effects of chronic diseases. This study aims to investigate the effectiveness of different exercise methods on the composition of intestinal microbiota.

Methods: We searched for published peer-reviewed articles in PubMed and Google Scholar databases up until January 2023. We searched using the terms "exercise", "education", "microbiome" and "microbiota".

Results: By electronic search until January 2023 in the databases with keywords made in Mesh by mentioning Title/Abstract, several 1814 articles were collected. By repeating and performing filtering at each stage, 15 clinical trials were finally left in the study. The results showed varying degrees of efficacy and high inter-individual variations. In conclusion, the baseline microbiome profile was shown to have a decisive role in microbiome responsiveness to training intervention and training dose and duration seem to be a determining factor in all exercise modalities.

Conclusion: In general, can be said that exercise can balance GM. More importantly, exercise is proposed to present a stressor to the gut that stimulates beneficial adaptations and improves long-term gut barrier flexibility over time through regular physical activity. It seems that the GM changes caused by aerobic exercise are reversible after returning to a sedentary lifestyle. Therefore, it is recommended that exercise initially causes a disrupts in GM that, with continued exercise and thus adaptation, revenues to the pre-exercise state.

Keywords: Gut Microbiota, Exercise, Aerobic Exercise, Resistance Training, Combined Training

1. Introduction

G

ut microbiota (GM) composition can influence human metabolism and has been shown to influence metabolic health and decrease the risk of metabolic diseases (1).

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Exercise is increasingly reported to positively influence GM by improving the diversity and abundance of health-promoting species, and altering its functional capacity through increasing metabolites (2).

However, the physiologic adaptations to different types of exercise such as resistance and aerobic training are distinctly different and they elicit several pathways in our body (3). Because of the differences in metabolic pathways and the fuel source utilized in these exercise modalities, their impact on the human GM might be different (4). For example, the contribution of anaerobic energy in resistance exercises is predominant (5) and it has been shown that high-intensity anaerobic exercises produce lactate which enters the gut lumen through circulation and provides a selective advantage for lactate-utilizing species (6). Moreover, aerobic exercise specifically enhances cardiorespiratory fitness which is associated with greater diversity in GM (3).

GM can be effective in regulating energy metabolism, hydration, inflammatory response, and oxidative stress (7). Exercise can affect the GM due to its metabolic and immune effects. Exercise-induced changes in GM may depend on the type of exercise (8). The effect of physical activity on GM has been investigated in many studies, most of which are animal models that have investigated the effect of a specific exercise with a food combination (7, 9, 10). This study is a pulmonary systemic investigation that examines the effect of various exercise interventions on GM in human populations.

2. Methods and Materials

2.1 Type of Study

The present study is a secondary study of review type. How to conduct the study is shown in Figure 1.

2.2 Search Methodology

A PubMed search using the terms “exercise”, “training”, “microbiome” and “microbiota” was conducted for articles published until January 2023. The secondary search was extended to the Google Scholar database. To ensure that all relevant literature was included, the reference lists of identified articles were also searched, and relevant papers were identified.

2.3 Search strategy

In this study, we classified exercises into three classes (aerobic, resistance (strength) training, or combined aerobic and resistance training) and their effects on GM characteristics (alpha diversity, beta diversity, changes in microbial composition and abundance, and metabolites produced by specific species). Then, the keywords used in this search were combined with the medical subject profiles (Mesh) and with the abstract and title (Table 1).

Table 1. Descriptive Findings of Research Variables Before and After COVID-19

Search strategy		
Keyword	Aerobic, Resistance (strength) Training, Combined Aerobic, Exercise and Resistance Training	microbiome, microbiota, alpha diversity, beta diversity, changes in microbial composition and abundance, and metabolites produced by specific species
Search terms	(Aerobic OR Resistance Training OR Combined Aerobic OR Exercise OR Resistance Training) AND (microbiota OR microbiome OR alpha diversity OR beta diversity OR changes in microbial composition OR metabolites produced by specific species)	

2.4 Inclusion Criteria

In this study, only longitudinal studies were examined.

2.5 Exclusion criteria

Studies, where intervention participants received specific medications, supplements, or diets (other than exercise intervention) were excluded (11).

2.6 Quality Assessment

To evaluate the quality of the articles, the Jadad scale checklist was used due to its ability to score quantitatively (12).

3. Results

By electronic search until January 2023 in the databases with keywords made in Mesh by mentioning Title/Abstract, several 1814 articles were collected. By repeating and performing filtering at each stage, 15 clinical trials were

finally left in the study (Figure 1). A summary of the characteristics and findings of the included studies are presented in separate tables in each section (Table 2).

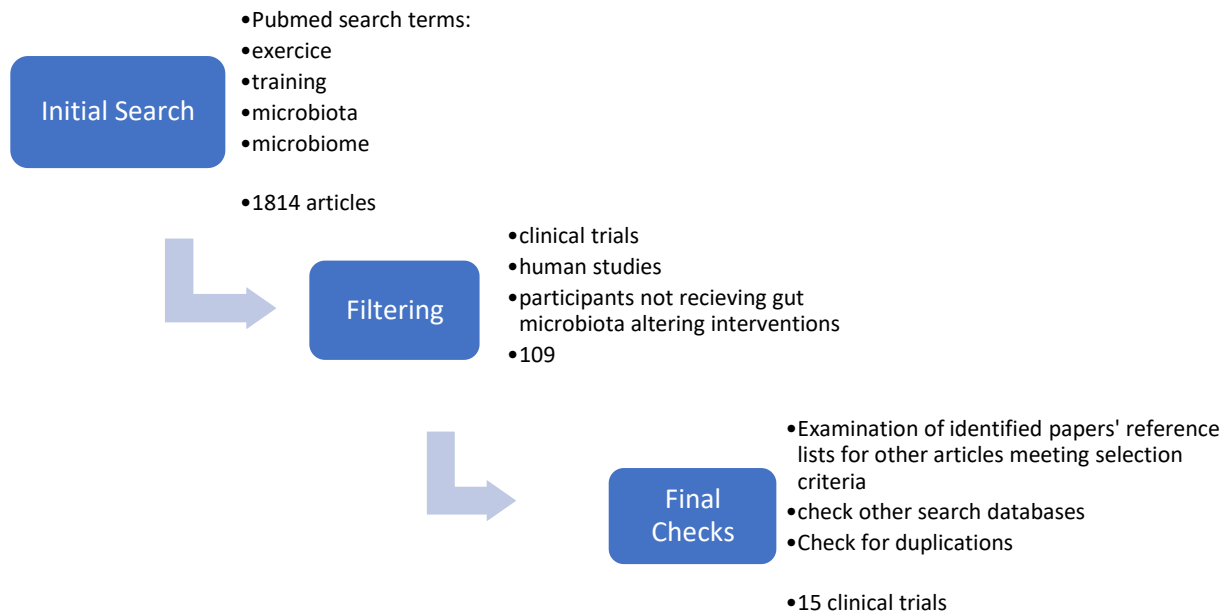


Figure 1. Steps for including articles in the review

Table 2. Summary of characteristics of reviewed studies

Author	Country	Sample size	Participants/	Intervention	Outcome	Follow-up period	Conclusion
Wiley Barton (13)	Ireland	86	Athletes/ sedentary individuals	AE	GM	12 weeks	A significant difference in fecal microbiota between athletes and sedentary individuals cannot indicate the effect of exercise.
Emiko Morita (2)	Japan	32	Elderly Women	AE	GM	12 weeks	Aerobic exercise increases intestinal bacteroids in healthy elderly women.
Rocío Quiroga (8)	Spain	39	Childhood obesity	CT	Proteobacteria phylum/Gammaproteobacteria class	12 weeks	Exercise training can be considered as an efficient non-pharmacological treatment, reducing inflammatory signaling pathways caused by obesity in children through microbiota modulation.
Owen Cronin (14)	Ireland	90	Sedentary Adults	Exercise	GM	8 weeks	Little changes in gut microbial composition and function were observed following increased physical activity.
Yan Liu (10)	China	39	Diabetic men	Exercise	GM	12 weeks	Exercise increases GT in diabetics

M. Dupuit (15)	France	17	Women with Overweight	HIIT*	GT	12 weeks	HIIT has a positive effect on gut GT in postmenopausal women.
Ryan P (16)	San Francisco	37	Healthy Participants	Exercise	GT	7 days	There is a relationship between the number and proportion of GT and cardiorespiratory fitness.
K. MOTIANI (8)	Turku	26	Sedentary Subjects	MIC†/SIT‡	GT	2 weeks	Exercise improves the intestinal microbial profile.
Ayane S (3)	Brazil	24	Healthy Men	AE	GT	10 weeks	Moderate intensity AE improves VO2 and boosts men's GT.
Hirokazu Taniguchi (17)	Japanese	33	Elderly Men	STE§	GT	5 weeks	STE has little effect on GT in the elderly.
Timo Kern (4)	Denmark	88	Overweight or Obesity	BIKE**/MOD††/ VIG‡‡	GT	6-month	Exercise produces little change in GT.
Sabrina Donati Zeppa (18)	Italy	17	seventeen healthy male	HIIT	GT	9 weeks	HIIT moves the number of GT in the gut towards healthier GT.
Elizabeth A (19)	New Zealand	29	overweight Men	HIIT	GT	3 weeks	Short-term HIIT has no effect on GT composition
Fei Zhong (20)	China	14	Older Women	Exercise	GT	8-week	Exercise can increase the frequency of GT
Kristine M (21)	USA	22	Older, Sedentary Adults	RT	GT	24week	Performing exercise interventions is associated with changes in GT

Findings of the included studies are presented in three sections: 1. Aerobic Exercise, 2. Resistance Training, and 3. Combined Training.

3.1 Aerobic Exercise Training (AE)

Most of the studies included in our review had AE as one of their primary educational interventions. AE has effects in several ways such as improving endurance and cardiopulmonary fitness and improving body composition through increasing lean body mass and decreasing body fat percentage (22). Cardiorespiratory fitness (VO2 max and maximum power) showed a positive correlation with the characteristics of GM such as increased bacterial diversity and butyrate-producing bacteria (10). VO2max has been shown to predict more than 20% of the variance of an individual's relative gut bacteria and is positively correlated with GM diversity (3). However, the results of longitudinal studies in this area are varied. For example, a moderate-intensity AE intervention has shown changes in GM composition, without improvements in cardiorespiratory fitness parameters (8). While a similar intervention by

Resende et al (3) improved VO2peak and affected GM composition in non-obese men, it did not alter alpha or beta variability.

Overall, it seems that exercise-induced changes in GM might not be a result of improved VO2 max and cardiorespiratory fitness and other physiological changes that occur during AE. Instead, improved cardiorespiratory fitness and improved diversity of GM composition might sometimes co-occur as a mutual outcome of exercise and no causality relationship might exist between VO2max and GM composition. Probably the effect of this type of AE depends on the GM composition and domestic characteristics of the analyzed subject; in particular on the BMI and age of the subject studied (3).

3.1.1 Richness and Diversity

The majority of AE interventions induced no changes in α -diversity. In the studies reviewed, only Kern et al. observed a 5% increase in the Shannon index in the vigorous-intensity cycling group after 3 months (4). Most AE interventions that we reviewed had employed a

* High-Intensity Interval Training

† Moderate-Intensity Continuous Training

‡ Sprint Interval Training

§ Short-Term Endurance

** Active Commuting by Bike

†† Leisure-Time Exercise of Moderate

‡‡ Vigorous Intensity

frequency of 3 training sessions per week, but Kern et al. intervention was performed 5 times per week (4). However, Morita et al. (2) which implemented daily training sessions for 3 months did not observe the same results. It is also noteworthy that Morita's study participants were healthy elderly over 66 years old, while Kern's study population aged between 20–45 years old, which might be a determining factor for the different results. In Kern et al. study, three intervention groups with different intensities were compared: moderate, leisure, and vigorous; and only vigorous AE resulted in changes in alpha diversity. Interventions with moderate intensity AE did not induce changes in alpha diversity (23). Moreover, when comparing a 2-week high-intensity and moderate-intensity AE training, no differences in microbiome richness and diversity were reported (8). Therefore, it can be inferred that a low to moderate-intensity AE or a short-term vigorous AE might not be enough to alter alpha diversity. Regarding AE effects on β -diversity, the data have been heterogeneous even among the participants of one intervention group with the same intervention protocol (3). These varied results can be due to confounding factors such as age, diet, fitness level, BMI, and overall, factors that determine the baseline GM profile.

3.1.2 Differences in Baseline GM Profile

Large interindividual differences in human GM have been observed even in healthy populations. Moreover, it is well recognized that there is great individual variability in microbiome change patterns in response to exercise training (24), even between individuals of the same group (25). For example, some studies (10) have shown that exercise non-responders or low responders have an attenuated microbiome response to exercise training. In Munukka's study (23) only half of the subjects' gut microbiome responded to PA significantly. Moreover, in lean participants, GM modulations were significantly associated with body composition changes, whereas in obese participants these changes were associated with changes in VO_2 max. In Allen's study, five butyrate-producing bacterial genera and consequently, fecal concentrations of SCFAs (acetate and butyrate) were increased in lean individuals (26). In contrast, obese subjects showed a decrease in the abundance of *Faecalibacterium* spp. and an increase in *Bacteroides* species. In another study (19), 3 weeks of AE intervention showed that the *Subdoligranulum* genus was increased in lean subjects, while it decreased in obese

subjects. Allen et al. showed a marked improvement in VO_2 max in both lean and obese subjects. The basal GM profile plays a decisive role in the regulation of physiological adaptations and exercise-induced changes in GM composition, the effect of exercise on GM strongly depends on the basic profile of intestinal microbiota as well as the type and intensity of exercise (26).

3.1.3 GM changes in response to AE training

Due to variations in AE protocols (intensity, duration, and frequency) and because of the interpersonal variations between subjects, the influence of AE training on GM has been very diverse. For instance, Taniguchi et al. (17), showed that a five-week AE program in the elderly had modest effects on GM composition. In this study no changes in the GM's richness and diversity were reported; however, the relative abundance of *Clostridium difficile* and *Oscillospira* genus were decreased and increased, respectively, even if the changes in *Oscillospira* abundance was no longer significant after adjusting for dietary changes that had occurred. Furthermore, in the study by Motiani et al. (8) high and moderate intensity AE has been shown to decrease the ratio of Firmicutes/Bacteroidetes, mainly due to the increase in the relative abundance of Bacteroidetes phyla. Moreover, the *Clostridium* genus and *Blautia* genus were decreased in both training modes (8).

3.1.4 GM alterations after AE training are reversible

Allen et al. have shown that GM changes after an AE intervention are reversed once the sedentary lifestyle is resumed (26). Notably, some studies have shown that even without going back to a sedentary lifestyle and while the intervention is still ongoing, the initial GM alterations tend to reverse and diminish (4). Inconsistent, in a study by Bycura et al., GM composition changed after the first few weeks of the exercise program, but after the completion of the intervention, individuals' GM composition was no longer significantly different from the baseline point (6).

3.2 Resistance Training (RT)

In RT, the contribution of anaerobic energy is predominant and it has been shown that high-intensity anaerobic exercises produce lactate which enters the gut lumen through circulation and provides a selective advantage for lactate-utilizing bacterial species (1). Moreover, it has been shown that changes in GM due to

exercise are associated with modifications in body composition, especially lean mass (26). Animal study have also shown that RT can affect GM and alter its composition (27) however, human intervention studies on RT and GM are scarce, with various protocols and doses of intervention that will be discussed in the following text.

3.3 Combined Training (CT)

Greater plasticity of the gut microbiome in young makes them more responsive to the effect of exercise on microbial composition, above all at the phylum level (28). In obese children (aged 7-12 years old), Quiroga et al. (29) have shown that combined exercise training reduces the abundance of obesity-associated bacteria, and changes the microbiota toward a non-obese profile, according to Allen (26) and Donati Zeppa et al. (18) results that showed a shift towards healthier microbiome profile after AE training.

Training interventions are shown to induce changes in habitual diet, toward healthier dietary patterns (30) and it could be quite challenging to maintain the habitual diet during a training intervention. For example, Donati Zeppa and colleagues (18) have shown that despite instructions to maintain their usual diets, participants showed significant increases in protein, carbohydrate, fiber, and vitamin C intake. These changes were recorded using consistent daily records, which no other study has applied. Since dietary changes drastically influence the GM composition, not all the changes in longitudinal studies are linked to exercise training and the results should be interpreted with this fact in mind. In another study (6), even with subjects reporting no major changes in their dietary pattern, significant increases in body weight of half of the participants were observed, which could indicate an unintended increase in energy intake.

A further influential factor to be considered is the protocol of training. In CT interventions, it is suggested that the modality, frequency, and duration of the selected aerobic training, influence the effects of resistance training. E.g., running, but not cycling, seems to decrease hypertrophy and strength alterations through RT (31). Even the sequence of training modes (AE or RT) in a mixed exercise intervention affects its efficacy (31).

4. Conclusion

In general, can be said that exercise can balance GM. More importantly, exercise is proposed to present a stressor to the gut that stimulates beneficial adaptations and improves

long-term gut barrier flexibility over time through regular physical activity. It seems that the GM changes caused by aerobic exercise are reversible after returning to a sedentary lifestyle. Therefore, it is recommended that exercise initially causes a disruption in GM that, with continued exercise and thus adaptation, revenues to the pre-exercise state. The disorder returns. Therefore, GM differences between athletes and non-athletes may be the result of long-term lifestyle differences, including diet and exercise training.

4.1 Limitation

This study is a Systematic Review study, and due to the variety of exercise interventions, there was a large heterogeneity of the studied population, which did not allow us to perform a meta-analysis.

4.2 Suggestion

It is suggested to carry out meta-analysis studies separately for each type of sport and population (Athletes/sedentary individuals) in order to make a better decision about the results of exercise in GM.

Authors' Contributions

M.S.M. was responsible for the study concept and design, analysis and interpretation of data, critical revision of the manuscript for important intellectual content, statistical analysis, and study supervision. M.R.R. contributed to the acquisition of data, drafting of the manuscript, and provided administrative, technical, and material support. Both authors have actively participated in the research process and approved the final version of the manuscript.

Declaration

In order to correct and improve the academic writing of our paper, we have used the language model ChatGPT.

Transparency Statement

Data are available for research purposes upon reasonable request to the corresponding author.

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Declaration of Interest

The authors report no conflict of interest.

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Ethics Considerations

Ethical standards, including fair use of information and proper citation, were strictly adhered to, maintaining the integrity of the review.

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