

8:7

Hormonal and metabolic substrate status in response to exercise in men of different phenotype

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Abstract

Objectives: The present study verified the effect of moderate-to-high-intensity aerobic exercise on the endocrine response profile and adipose tissue in young healthy men with different phenotype characteristics.

Design: Eighteen men were divided into three experimental groups with defined body components and specific physical fitness: Endurance phenotype – EP (n = 6; low body mass; low fat content; aerobic endurance trained), Athletic phenotype – AP (n = 6; high body mass; low fat content, resistance trained), Obesity phenotype – OP (n = 6; high body mass; high fat content, untrained).

Methods: The participants performed an progressive exercise protocol on a treadmill (30% VO_{2max} , 50% VO_{2max} , 70% VO_{2max}), separated by 45 s of passive rest for blood collection.

Results: Plasma glucose oxidation increased in relation to exercise intensity, but to a greater extent in the AP group. The free fatty acids' plasma level decreased with a rise in exercise intensity, but with different kinetics in particular phenotypes. Plasma growth hormone increased after the cessation of exercise and was significantly higher in all groups 45 min into recovery compared to resting values. Plasma insulin decreased during exercise in all groups, but in the OP, the decrease was blunted.

Conclusions: The results indicate that the rate of lipolysis, hormonal and metabolic response to aerobic exercise depends on the individuals' phenotype. Thus, exercise type, duration and intensity have to be strictly individualized in relation to phenotype in order to reach optimal metabolic benefits.

Key Words

- ▶ free fatty acid metabolism
- endocrine response
- lipolytic hormones
- ▶ phenotype
- exercise metabolism

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Introduction

Somatotypes, which to some extent express genetic determinism, are significantly associated with the level of physical fitness (1, 2), and as a consequence with the content of adipose tissue (1). Adipocytes are highly metabolically active cells releasing adipokines and adipocytokines which affect the processes in neighboring and distant cells by autocrine, paracrine and endocrine fashion (3). The rate of release of the above

mentioned adipocyte-derived factors can be modified by environmental changes, among which physical activity and dietary habits play a vital role.

Adipose tissue (AT) does not display homogenous properties, and its physiological role depends on the location in the body (4, 5). Considering body metabolism, the white adipose tissue (WAT) also referred to as subcutaneous or visceral fat (intraabdominal: mesenteric,





Exercise metabolism in different phenotypes

8:7

perirenal and epididymal depots) serves as a primary energy store, and lipolysis in WAT is a crucial process for whole-body energy homeostasis. Physical exercise is recognized as a strong environmental stimulus of lipolysis in WAT, being a significant preventive factor against the so-called non-communicable diseases (2, 5). Most exercise protocols designed to induce AT loss have focused on regular steady-state aerobic exercise (6). In response to moderate-intensity aerobic exercise (~30-60% VO_{2max}) whole-body lipolysis is increased two- to threefold compared with resting values (6, 7, 8). When exercise intensity increases (above ~50% VO_{2max} for the general population), lipid oxidation starts to decrease and drops dramatically once the intensity exceeds 80% of VO_{2max} (6, 9). On the other hand, some authors suggest that higher exercise intensity (>70% VO_{2max} , HIIE) has the potential to be an economical and effective form of reducing body fat (10).

Exercise reveals the influence on the rate of AT lipolysis via stimulation of release of both lipolytic and antilipolytic hormones in an exercise intensitydependent manner (9). Endocrine exercise response depends on body mass, fat mass content and subjects' training status (11). It implies a possible role of phenotype as a factor modifying AT lipolysis response to hormonal action. In line with such an assumption are studies investigating fat oxidation during exercise in obese and lean subjects albeit mechanisms responsible for the above mentioned response are not fully recognized and provide conflicting data (12). For example, the plasma concentrations of non-esterified fatty acids (NEFAs) are not significantly different in high-intensity compared to low-intensity aerobic exercise (13). Moreover, postexercise circulating NEFA and fat oxidation remain in stimulation for several hours (13), which indicates the essential contribution of AT in resting post-exercise metabolism. Thus, the ability to alternate between storage and breakdown of AT is crucial for a healthy substrate metabolism in humans (14).

Most of the above mentioned study outcomes have been obtained during steady-state endurance exercises. However, there remains a paucity of evidence on how continuous running with low, moderate, or high intensity during a single continuous bout of exercise affect plasma hormonal and metabolic substrate status among individuals with different phenotypes. To approach this issue, we enrolled subjects with endurance, athletic and obese phenotypes and compared their plasma hormonal and metabolic substrate status.

Materials and methods

The study included 18, healthy trained and untrained men (mean age 32 ± 5.4 years). The participants were divided into three experimental groups with defined body components and level of physical fitness. The inclusion criteria into particular experimental groups were age, body mass, fat content, physical fitness and type of training (Table 1). Exclusion criteria on the other hand included infectious and inflammatory diseases, the use of food supplements or any systemic pharmacological treatment within 2 months prior to the study. All subjects kept their dietary habits constant during the study period. The endurance phenotype – EP group (n=6; mean body mass (BM) 73.8±7.6kg; fat content (FAT%) 11.6±2.7%; VO_{2max} - 56.8±4.3 mL/min/kg, aerobic endurance trained), the athletic phenotype – AP group (n=6; mean body mass (BM) 115 ± 7.0 kg; fat content (FAT%) 11.0 ± 1.7 %, VO_{2max} 43.2 ± 5.2 mL/min/kg, resistance trained), the obesity phenotype – OP group (n=6; mean body mass (BM) 132.5±3.6kg; fat content (FAT%) 30.0±7.7%, VO_{2max} -32.8±4.7 mL/min/kg, untrained).

The experiment lasted 10 days. The study consisted of three visits to the laboratory on the 1st, 3rd and 10th day of investigation (Fig. 1).

Visits to the laboratory began at 08:00h. On the 1st day all participants underwent medical examinations, anthropometric and somatic measurements to exclude any contraindications to the exercise protocol and to classify each subject into one of the three groups. On the 3rd day of experiment, progressive ramp test was performed on a treadmill to determine VO_{2max} and successive loads for the exercise test protocol. Energy intake, as well as macro and micronutrient intakes of all subjects, were determined by the 24-h nutrition recall. Basal metabolic rate (BMR), as well as energy expenditure at rest (REE) and during exercise (EE), were evaluated using the indirect calorimetry method. The participants were informed about the risks and benefits of the study and provided their written consent before the beginning of the experiment. The study received the approval of the Bioethics Committee at the Academy of Physical Education in Katowice, Poland (7/215).

The measurements of body mass were performed on a medical scale with a precision of 0.1 kg. Body composition was evaluated in the morning, between 08:00 and 08:30 h. The day before, the participants had the last meal at 20:00 h. They reported to the laboratory after an overnight fast, refraining from exercise for 48 h. Body composition





816

8:7

Variables	Obesity phenotype (<i>n</i> = 6)	Athletic phenotype (n = 6)	Endurance phenotype (<i>n</i> = 6)
Age (years)	31 ± 3.2	32 ± 2.8	33±3.0
Height (cm)	194.5 ± 2.12 ^a	186.3 ± 4.9	177.3 ± 5.03
BMI (kg/m ²)	34.3 ± 3.2	33.2 ± 2.6	22.9 ± 2.3
BM (kg)	132.5 ± 3.6	115 ± 7.00	73.8 ± 7.6 ^{bc}
FM (%)	30 ± 7.7 ^a c	11 ± 1.72	11.5 ± 2.17
FFM (kg)	92.15 ± 0.7	101.83 ± 5.8	64.9 ± 5.02^{bc}
BMR (kcal)	2363 ± 11.31	2555.7 ± 91.7	1772 ± 108.2 ^{bc}
TDEE (kcal)	3054 ± 169	3174 ± 207	2640 ± 525
DEI (kcal)	1853.5 ± 297 kcal	3839 ± 520 ^{ba}	2451 ± 240
Carb (%) of diet energy	37.5 ± 10	52 ± 7	39±16
Fats (%) of diet energy	38.5 ± 0.7	21 ± 8	30 ± 10
Pro (%) of diet energy	25 ± 9	27 ± 1	31 ± 5

 Table 1
 Basic anthropometric characteristics of the study participants.

Data are expressed as mean \pm s.p., n = 6.

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^aSignificant difference compared to EP (P < 0.05); ^bSignificant difference compared to OP (P < 0.05); ^cSignificant difference compared to AP (P < 0.05). BM (kg), total body mass; BMI (kg/m²), body mass index; BMR (kcal), basal metabolic rate; DEI, daily energy intake; FFM (kg), fat free mass; FM (%), fat mass; TDEE (kcal), total daily energy expenditure.

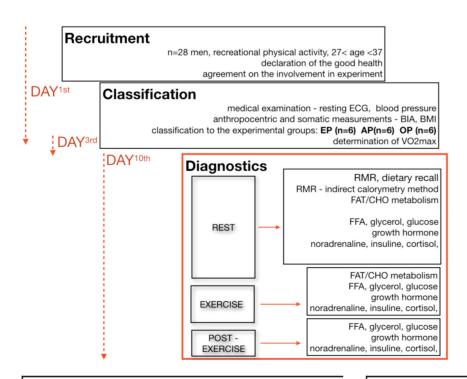
was evaluated using the electrical impedance technique (Inbody 720, Biospace Co., Japan).

The ramp VO_{2max} test was performed on a treadmill (Pulsar; H/P Cosmos, Nussdorf-Traunstein, Germany), starting at a speed of 6 km/h, which was increased linearly (1 km/h/1 min) until volitional exhaustion. During the test, heart rate (HR), oxygen uptake (VO₂), expired carbon dioxide (CO₂), minute ventilation (VE), breath frequency (BF) and respiratory exchange ratio (RER) were measured continuously using the MetaLyzer 3B-2R spiroergometer (Cortex, Leipzig, Germany) in the breath-by-breath mode. The following two criteria were used to determine

 VO_{2max} : (a) a plateau in VO_2 despite an increase in running speed, (b) RER >1.10.

Resting energy expenditure was measured immediately after waking at 08:00 h. Measurements were made with the subject in a supine position on a comfortable mattress. RER was measured for 30 min using the rest test procedure (Metalyzer). The gasses collected during a period of 20 min were analyzed by the MetaSoft® software.

On the 10th day of the experiment, at 09:00h, after an overnight fast, the participants performed a 35-min progressive treadmill exercise protocol at speeds corresponding to 30, 50 and 70% of their personal







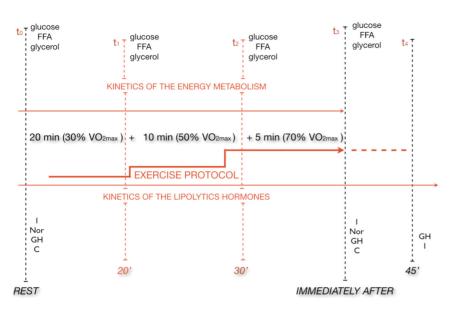
J Chycki *et al.*

Endocrine

Exercise metabolism in different phenotypes

8:7

817





 VO_{2max} (20-min walk at 30% VO_{2max} , 10-min jog at 50% VO_{2max} , 5-min run at 70% VO_{2max}) separated by 45s of passive rest for blood collection. Blood samples were obtained before (t0), during (20' (t1), 30' (t2) the exercise protocol, immediately after (t3), and 45 min into recovery (t4)) (Fig. 2).

At each of the five time points (t0, t1, t2, t3, t4), venous blood samples were collected to determine growth hormone (GH) - t0, t3, t4 noradrenaline (NA) - t0, t3, insulin (I) - t0, t3, t4 cortisol (C) - t0, t3, glucose (G) - t0, t1, t2, t3, free fatty acids (FFA) – t0, t1, t2, t3, and glycerol (GC) – t0, t1, t2, t3. All hormone concentrations as well as FFA and glycerol concentrations were evaluated in blood serum. On the other hand, glucose concentration was evaluated in blood plasma collected for EDTA with NaF. An enzymatic assay kit was used to measure glucose (bioMérieux SA, Marcy l'Etoile, France), and the colorimetric analysis was applied to measure FFA (FA115; Randox, Crumlin, United Kingdom) and glycerol (GY105; Randox). C, GH and I concentrations were measured in duplicate using EDTA plasma and immunoassay kits customized on an automated analyzer (Cobas e411; Roche Diagnostics). NA was measured once by the highperformance liquid chromatography method, what was justified by the recommendations of the producers of the biochemical assays. The intra-assay coefficient of variation was 6.5% for NA, for cortisol 2.2%, for GH 2.3% and 4.6% for insulin.

Statistical analysis

Shapiro–Wilk, Levene's and Mauchly's tests were used in order to verify the normality, homogeneity and sphericity

of the sample's data variances, respectively. All data are presented as mean \pm s.D. Dynamic changes of considered variables were analyzed using time series and fixed base indexes. Verifications of the differences between groups in considered variables were carried out using multivariate ANOVA. Statistical significance was set at *P*<0.05. All statistical analyses were performed using Statistica software and Microsoft Excel.

Results

All participants completed the described testing protocol. The procedure was carried out in identical environmental conditions with the air temperature of $19.4\pm0.4^{\circ}$ C and humidity of $58\pm0.5\%$ (Carl Roth hydrometer – Germany).

The dynamics of RER, glucose, glycerol and FFA concentrations in particular phenotypes (OP, AP and EP) are presented in Table 2A, B, C and D).

Analysis of time series (fixed base indexes) revealed significant changes of RER values (%), for all time periods from rest to the 35th min of the exercise protocol in all studied groups (Table 2).

Additionally multivariate ANOVA revealed significant differences between groups for RER, glucose, glycerol and FFA (Table 2A, B, C and D).

Analysis of time series (fixed base indexes) revealed significant changes of RER (in the 30th and 35th min for OP, AP and for EP), G (in the 30th min for OP; 20th min for AP and 35th min for EP), GC (in the 30th and 35th min for OP; 35th min for AP and 30th min for EP) and FFA (in the 35th min for OP; 30th and 35th min for AP; in the 30th and 35th min for EP) values expressed in %, for





8:7

Table 2Results of time series – fixed base indexes.

	Obesity phenotype		Athletic phenotype		Endurance phenotype	
		Fixed base		Fixed base		Fixed base
Time	Value	indexes (%)	Value	indexes (%)	Value	indexes (%)
A. Respiratory excha	nge ratio					
Rest	0.76	Base	0.92	Base	0.75	Base
20 min	0.77	4.73	0.94	1.45	0.78	3.21
30 min	0.81	10.81 ^a	1.02	10.14 ^b	0.87	18.35℃
35 min	0.88	18.24 ^a	1.08	15.94 ^b	0.92	24.77 ^c
		Fixed base		Fixed base		Fixed base
Time	Value (mmol/L)	indexes (%)	Value (mmol/L)	indexes (%)	Value (mmol/L)	indexes (%)
B. Glucose						
Rest	4.93	base	4.21	base	4.66	base
20 min	4.71	-4.9	3.99	- 18.7 ^b	4.61	-6.1
30 min	4.27	-13.9	4.44	-10.5	4.93	0.1 ^c
35 min	4.82	-1.9	4.71	-4.6	5.4	10 ^c
		Fixed base		Fixed base		Fixed base
Time	Value (µmol/L)	indexes (%)	Value (µmol/L)	indexes (%)	Value (µmol/L)	indexes (%)
C. Glycerol						
Rest	61.56	base	40.61	base	33.40	base
20 min	91.55	48.7	65.23	6	54.1 9	-12
30 min	136.06	121 ^d	73.82	19.9	95.49	55.1°
35 min	163.11	164.9 ^d	94.45	53.4 ^b	91.41	48.5
		Fixed base		Fixed base		Fixed base
Time	Value (mmol/L)	indexes (%)	Value (mmol/L)	indexes (%)	Value (mmol/L)	indexes (%)
D. Free fatty acids						
Rest	0.58	base	0.43	base	0.32	base
20 min	0.51	-13.2	0.56	4.3	0.44	24.1
30 min	0.48	-18.4	0.35	- 40.4 ^b	0.35	– 39.1 ℃
35 min	0.35	- 40 ^d	0.33	- 42.9 ^b	0.38	-35.5°

 ^{a}P < 0.01 significant difference compared to AP and EP; ^{b}P < 0.01 significant difference compared to OP and EP; ^{c}P < 0.01 significant difference compared to OP and AP; ^{d}P < 0.01 significant difference compared to AP.

all time points above the 29th min of exercise (Table 2A, B, C and D).

while I values were significantly lower after 35 min of exercise (Table 3A and B).

GH values after 35 min of exercise and 45 min of recovery were significantly higher in comparison with basal level in all phenotype groups. Similarly, NA values after 35 min of exercise were significantly higher in comparison with basal level in all examined groups, Analysis of time series – fixed base indexes, revealed significant changes of NA (in the 35th min of exercise and 45th min of recovery for OP, AP and EP), GH (in the 35th min of exercise for OP; in the 35th min of exercise and 45th min of recovery for AP; as well as for the 35th min

 Table 3
 Results of time series – fixed base indexes.

	Obesity phenotype		Athletic phenotype		Endurance phenotype	
Time	Value (ng/mL)	Fixed base indexes (%)	Value (ng/mL)	Fixed base indexes (%)	Value (ng/mL)	Fixed base indexes (%)
A. GH						
Rest	0.14	100	0.49	100	0.51	100
35′	2.51	1652.3ª	9.03	1756.1 ^b	10.87	2030.7 ^c
45′res	0.30	108.1	5.03	934.2 ^b	3.92	667.97 ^c
Time	Value (µIU/mL)	Fixed base indexes (%)	Value (µIU/mL)	Fixed base indexes (%)	Value (µIU/mL)	Fixed base indexes (%)
B. Insulin						
Rest	17.00	100	3.97	100	5.02	100
35′	8.67	50.98	1.98	50	2.95	58.80
45′res	13.67	80.39ª	2.62	65.97 ^b	3.78	75.42°

 $^{a}P < 0.01$ significant difference compared to AP and EP; $^{b}P < 0.01$ significant difference compared to OP and EP; $^{c}P < 0.01$ significant difference compared to OP and AP.



Exercise metabolism in different phenotypes

8:7

of exercise and 45th min of recovery for EP) and I (in the 45th min of recovery for OP, AP and EP) values in %, for all time points from rest to the 45th min of recovery.

Endocrine

Multivariate ANOVA showed, statistically significant differences (P<0.05) between all groups for GH, I and NA values at all time points of measurement, while there were no statistically significant differences between groups in cortisol values (P>0.05).

Discussion

In this study, we identified three phenotypes, i.e. EP, AP, OP using body mass, fat content, physical fitness and type of training as independent variables to investigate selected plasma hormonal and metabolic substrate status in response to continuous exercise of low, moderate and high-intensity in healthy men. These phenotypes can be attributed to different lifestyles and related to non-genetic risk (obesity) or protective (physical activity) factors of non-communicable diseases (15, 16, 17, 18).

The body content of subcutaneous AT is generally attributed to environmental factors (19). This tissue is a key energy storage site in humans which dominates in postprandial conditions (20). One of the primary functions of this tissue is the release of NEFA which are exported to peripheral tissues, mostly skeletal muscles, where they can be used as a substrate for aerobic metabolism. The contribution of NEFA to the total oxidative metabolism is dependent on a variety of factors, among which exercise intensity and duration, as well as dietary and training status play a pivotal role. In this study, regardless of these items, we explored this issue by selecting individuals of different phenotypes who were characterized by different body mass, body fat content and fat distribution, different levels of fat free mass and physical fitness (Table 1).

Fat oxidation during identical relative exercise intensity was repeatedly found to be higher in trained compared with untrained individuals (21). Our analysis has revealed that dynamics of circulating NEFA discloses a biphasic pattern in both athletic and endurance phenotypes, being more pronounced in the second group of individuals. After a 20-min walk at 30% VO_{2max} the level of NEFA achieved peak values, and next decreased during the run at 50% VO_{2max} in both phenotypes, when NEFA oxidation reaches a maximal rate (6). These results are in agreement with the view that lipid metabolism is dependent on the intensity of exercise, and in line with such an assumption is also a further lack of a drop in FFA level despite the increase in exercise intensity (70% VO_{2max}). It was shown that plasma NEFA uptake and oxidation was limited during high-intensity exercise (22). Under these circumstances intramuscular TG hydrolysis may at least partially substitute reduced accessibility of NEFA from plasma by adrenaline (23) and contraction (24) stimulation of hormone-sensitive lipase (HSL) within muscle cells.

It seems that the efficiency of fat oxidation at muscle cell level is the most likely controlling factor, which can be modified by substrate availability before the appearance of leveling off (about 85% of VO_{2max}) in net NEFA (predominantly long chain) uptake. This was confirmed in our study by a constant rise of plasma glycerol levels and non-parallel changes in plasma NEFA during particular exercise intensities in AP and EP individuals. A linear drop in plasma NEFA in the obese phenotype suggests that this mechanism likely operates in a different manner in this group of subjects, and may be attributed to their dietary status, which was characterized by a significantly higher energy intake from fats (Table 1). The driving force of the FFA flux from vascular space to muscle cells is assumed to be the concentration gradient between these compartments, which could be kept upregulated (higher plasma NEFA level) in obese phenotype individuals within all ranges of exercise intensities. The view that delivery from vascular space can play an important role in fat metabolism of the above mentioned subjects also appeared from RER values which were lower both at 30 and 50% $\mathrm{VO}_{\mathrm{2max}}$ in these subjects in comparison to other investigated groups, indicating their higher reliance on lipids as a source of energy supply. On the other hand, the observed changes in RER in the athletic phenotype indicate the highest glucose contribution to oxidative metabolism. Since the dynamics of insulin concentration had a similar pattern in all phenotypes, the differences in glucose concentration between phenotypes may reflect the activation of different regulatory mechanisms at the muscle cellular level including NEFA re-esterification. Among regulatory factors that may play a major role in this phenomenon, one can point to differences in muscle insulin sensitivity, different activation of contraction-induced insulin signaling pathways and impact of substrate and exercise on muscle malonyl-CoA concentration. Physiological catecholamines, GH and insulin are primary controllers of both AT lipolysis and glucose metabolism. Catecholamines and GH have a stimulatory effect on AT metabolism, while insulin inhibits lipolysis. There is significant evidence for a hormone-coordinated regulation of AT TG hydrolysis during prolonged exercise (25). A linear increase in plasma NA and GH and a drop in plasma I observed in our study



J Chycki *et al.*

820

8:7

indicate that the above mentioned hormone regulatory mechanisms expressed this general physiological pattern in all investigated phenotypes, and this mechanism seems not to be affected by phenotype.

Endocrine

Exercise is a major stimulus of GH secretion. GH becomes an important regulator of lipolysis during prolonged fasting, where lipolysis is stimulated together with increased GH and decreased insulin levels (21, 26). Previous studies showed that circulating GH concentration can be regulated by several factors including food intake (27). It could thus be argued that a slower decrease in GH plasma concentration seen in individuals with athletic phenotype over the recovery period could be attributed to their nutritional status and content of amino acids in diet.

It has been established that adipocytes contribute to a lesser extent to whole-body insulin stimulated glucose disposal in normal individuals compared with skeletal muscle. Consequently, the primary and most evident site of insulin action should be observed in individuals with a higher FFM. This variable reached the highest values both in athletic and surprisingly obesity phenotypes. However, the later phenotype is also characterized by the highest content of FM. Because of similar exercise induced drops in insulin levels in all groups, changes in body composition could be considered as another factor taking part in different exercise induced adaptations between athletic and obesity phenotypes. These results suggest that individuals with obesity phenotype are more susceptible to insulin resistance, what has been confirmed in obese individuals (28).

Among known catecholamines, A belongs to the best investigated in term of its lipolytic effect. However, NA was detected in the interstitial fluid of adipose tissue at concentrations similar to those observed in plasma (29). This suggests that NA impacts AT lipolysis, similarly to A, and can be expressed via a dual mechanism: direct action on adrenergic receptors located in adipose cell membranes (30), and indirectly, by regulation of blood flow to AT (31). Many studies are not fully conclusive, as some authors found enhanced whole-body lipolysis during exercise in trained compared with untrained subjects (21, 32), while others did not find these differences (33). Such an association may be expected in face of results revealing that variation in catecholamine pathway genes contribute to the inter-individual variability in plasma NA and A concentrations, both at rest and after recovery from exercise (33).

In summary, the present study provides more data about the metabolic and hormonal responses to low and moderate intensity aerobic exercise. Our findings highlight the benefits of targeted training protocols dependent on phenotype and dietary habits. The results of this research can contribute to the individualization of training protocols, which will optimize the lipolytic effect. Combining phenotype and metabolism with a range of counter-regulatory lipolytic hormones provides a more complete perspective on cross-talk between the endocrine system and exercise metabolism. It has been shown experimentally that inhibition of whole-body lipolysis improves whole-body insulin sensitivity and metabolic profile of subjects. Studies investigating the combined effect of lipolysis inhibition and different types of physical activity may bring new information into this field of study. The results indicate that the rate of lipolysis, hormonal and metabolic response to aerobic exercise to some extent depends on the individuals' phenotype. Thus, exercise type, duration and intensity have to be strictly individualized in relation to phenotype in order to reach optimal metabolic benefits. The observed changes in RMR and RER indicate the highest glucose contribution to oxidative metabolism in the athletic phenotype, suggesting the importance of FFM in the regulation of lipolysis and energy expenditure. For people with a high content of lean body mass and significant glycogen stores, performing high-intensity exercise $(>70\% \text{ VO}_{2\text{max}})$ seems to be a rational training strategy for improving lipolysis. Moreover, in overweight and obese people, two mechanisms regulating exercise metabolism and especially lipolysis are significantly blunted. First of all post-exercise insulin decrease is significantly lower and the efflux of GH and noradrenaline is much smaller than in resistance and endurance trained subjects. Thus, for people with obesity using lower intensity aerobic efforts (50% VO_{2max}) seems most appropriate for improving fat metabolism and inducing favorable changes in body composition.

Declaration of interest

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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8:7

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