The Effect of Exhaustive Exercise on Plasma Metabolic Profiles of Male and Female Rats

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Abstract
The objective of the study was to evaluate the alteration in biochemical composition and gender difference within exhaustive exercise in male and female rats using a metabolomics strategy. Sixty male and female rats were randomly assigned to control, exhaustive exercise and one-week recovery groups, respectively. The metabolic profiles of plasma were investigated by gas chromatograph-mass spectrometry (GC-MS) and data further underwent orthogonal partial least-squares (OPLS) analysis. The current study found that gender was a significant determinant of the effects of exhaustive exercise on the cortisol, blood urea nitrogen, creatine kinase, and the ratio of reduced glutathione to oxidized glutathione, whereas, no significant interaction effects between gender and exhaustive exercise were found on the levels of testosterone, malonaldehyde, reduced glutathione, oxidized glutathione and lactic dehydrogenase. In male rats, the altered metabolites within exhaustive exercise included increased tricarboxylic acid cycle intermediates (citric acid, fumaric acid, butanedioic acid), branch-chain amino acids (valine, leucine), fatty acids and metabolite (oleic acid, linoleic acid, 3-hydroxybutyric acid), phosphate and decreased glucose, lactic acid, serine, and glutamic acid. In female rats, the levels of fatty acids and metabolite (linoleic acid, oleic acid, arachidonic acid, 3-hydroxybutyric acid), amino acids (valine, leucine, glutamic acid, 5-oxo-proline, methionine, ornithine), other metabolites urea, myo-inositol and phosphate were increased. The results indicated that exhaustive exercise increased the rates of energy metabolism, glucose metabolism, amino acid catabolism and fatty acid metabolism in male rats, whereas, female rats showed an increased propensity to oxidize lipid and conserve carbohydrate and protein metabolism against physical stress. Disordered urea cycle and inositol metabolism also occurred in female rats with exhaustive exercise. Exhaustive exercise affected the balance of hormone adjustment and caused oxidative stress, subsequent cell membrane damage both in male and female rats. A significant gender-related difference in the metabolic profiles was also found between male and female rats within exhaustive exercise.

Key words: Exhaustive exercise, physiology, metabolomics, plasma, gender difference.

Introduction
Physical exercise is one of the major modulators of systemic metabolism. It increases the rate of metabolic processes and modulates the levels of different metabolites. A persistent, high exercise volume and intensity with limited recovery periods could lead to declined performance, and subsequent accelerated fatigability (Margonis et al., 2007). Various biological markers, hormonal markers and immunological markers have been used to evaluate the physical status within exhaustive exercise (Aguiló et al., 2005; Ferreira et al., 2018; Sarikaya et al., 2017; Zheng et al., 2018). However, these conventional methods, which involved measuring the concentrations of only a few biochemistry parameters or a few target metabolites in test samples, could not sensitively reflect all physiological differences and would be unable to explain the interaction between varieties of the metabolites (Taysi et al., 2008).

Metabolites are the final biological products of the system, which means that investigating the changes of these endogenous compounds can verify the response endpoint of biological substances (Gao et al., 2014). Metabolomics strategy is a useful analytic platform to determine endogenous metabolites and assess the global and dynamic metabolic responses of living system (Nicholson and Lindon, 2008). Recently, metabolomics has attracted increasing interest in the field of sports medicine and shown great potential for monitoring the changes of physiological state. Huang et al. (2010) used metabolomics to determine the biochemical variations in male rats’ liver with exhaustive and endurance exercises. Kume et al. (2015), using plasma metabolome analysis, identified potential biomarkers of fatigue in male rats. Miao et al. (2018) investigated the mechanism of the anti-fatigue effect of Danggui Buxue Tang on fatigue of male mice induced by forced swimming, and identified 14 metabolites providing evidence for the anti-fatigue effect. Jang et al. (2018) determined changes in urinary metabolic profiles and metabolomics markers in muscle damage following eccentric exercise in men and women, which clarify the metabolic response within eccentric exercise-induced muscle damage and gender-dependent patterns. By measurement of the endogenous metabolites, metabolomics investigations have the potential to distinguish the change of metabolic profiling and to identify the biomarkers associated with exercise performance, fatigue or exercise-induced disorders (Ra et al., 2014; Yan et al., 2009).

Although there have been several studies that examined the metabolic response to exercise, there is a lack of information on the gender-related differences in the metabolic response to exhaustive exercise. Therefore, in this study, the metabolomics study was used to distinguish the variations in metabolite profiles and determine biomarkers changed during exhaustive exercise and recovery in male and female rats.
Methods

Rat blood sampling
Sixty eight-week-old adult male and female Sprague-Dawley rats were randomly subdivided into control, exhaustive exercise and recovery groups consisting of 10 rats each. Before the commencement of the experiment, rats from exhaustive exercise and recovery group were familiarized to adaptive swimming training for 10 min/day for 3 days. Weight-loaded, forced swimming was performed according to the methodology described previously (Xu et al., 2013) with some modifications. The rats swim with a load of aluminum sheets that weighed 5% of their body weight and were attached to their tails. Rats swim individually until exhaustion once a day for 10 consecutive days; exhaustion was reached when the rat was unable to constantly keep its nose out of water and its nose remained below the water surface for 10 seconds (Ma et al., 2017). The recovery group was given a 7-day period to refresh after the exhaustive exercise period, while the rats in the control group were left in cages without swimming. The water temperature was controlled around 32-36 °C. The animals were housed at a controlled ambient temperature of 22-25 °C with 55-65% relative humidity and a 12 h light/12 h dark cycle (lights on at 7:00 AM) and were given food and water ad libitum at the Laboratory Animal Center in Shanghai University of Traditional Chinese Medicine, Shanghai, China. All animal procedures were performed in accordance with Institutional Animal Care and Use Committee guidelines of Shanghai University of Traditional Chinese Medicine. To investigate the metabolites generated in the system, several blood collections from each animal were collected immediately at day 5 and day 10 of swimming, as well as day 3 and day 7 post recovery.

Biochemical assay
Serum testosterone (T) and cortisol (CORT) were analyzed using specific assays strictly according to the manufacturer’s instructions (Nanjing Jiancheng CO., China) and measured by the microplate reader (Biotek, USA). The malondialdehyde (MDA) assay kit (TBA method) was used to detect the MDA level as a marker of lipid peroxidation and the kit was bought from Nanjing Jiancheng CO., China. Blood urea nitrogen (BUN), creatine kinase (CK) and lactate dehydrogenase (LDH) were measured using an automated biochemistry analyzer (Hitachi, Japan). Plasma reduced glutathione (GSH) and oxidized glutathione (GSSG) were evaluated as markers of oxidative stress using specific assay kits from Nanjing Jiancheng CO., China and measured by the microplate reader (Biotek, USA).

Reagents
4-Chlorophenyl-alanine, heptadecanoic acid, methoxyamine and N,O-bis(trimethylsilyl)-trifluoroacetamide (BSTFA) were purchased from Sigma-Aldrich (Sigma-Aldrich, USA). Methanol (HPLC Grade) and chloroform were obtained from Thermo Fisher (Thermo Fisher, USA). Pyridine was purchased from China National Pharmaceutical Group Corporation (China National Pharmaceutical Group Corporation, China). Pure water was produced by a Milli-Q purification system (Millipore, USA).

Sample preparation for GC-MS analysis
Plasma samples were treated with chemical derivatization following our previously published procedure (Liao et al., 2012). Each plasma sample (100 μL) was added with two internal standard solutions (10 μL of 4-chlorophenyl-alanine in water, 0.3 mg/mL; 10 μL of heptadecanoic acid in methanol, 1 mg/mL) and with 300 μL methanol/chloroform (3:1) to extract the metabolites. After vortex mixing for 30 s, the mixtures were incubated at -20 °C for 10 min and centrifuged at 12,000 rpm for 10 min. A 300 μL supernatant aliquot was transferred into a GC vial and evaporated to dryness under N2 at 30 °C. Methoxyamine (80 μL) in pyridine (15 mg/mL) was added to the dried residue and vortex mixed for 1 min. The methoximation reaction was carried out for 90 min while rocking in an air-shaker at 30 °C, followed by trimethylsilyl for 60 min by adding 80 μL BSTFA at 70 °C. At last, the solution was vortex mixed 30 s and cooled into room temperature for GC-MS analysis.

Metabolite analysis by GC-MS
Each 1 μL derivatized sample was injected onto a HP-5MS capillary column (30 m × 250 μm inner diameter, 0.25 μm film thickness, Agilent J&W Scientific, USA) on an Agilent 7890A GC/5975C MSD (Agilent J&W Scientific, USA). Helium was used as the carrier gas through the column with a constant flow rate of 1.0 mL/min. The sample was injected at 270 °C in splitless mode. The optimized GC-MS gradient temperature programming was selected following our previous experiment (Liao et al., 2012): the GC oven was started at 80 °C for 2 min, then the temperature was increased step-wise, starting at 10 °C/min to 140 °C, 4 °C/min to 210 °C, 10 °C/min to 240 °C, 25 °C/min to 290 °C and then maintained at 290 °C for 3 min. The ion source temperature and the quadrupole temperature were set at 230 °C and 150 °C, respectively. The mass data were acquired in scan (m/z 30-600) mode at a rate of 20 spectra/s with electron impact ionization (70 eV). The solvent delay time was set to 5.0 min.

Data analysis
The data from the GC-MS were converted into CDF formats, and the data were processed by the XCMS toolbox (http://metlin.scripps.edu/download/) to carry out baseline correction, peak deconvolution and alignment using XCMS’s default settings. The result (CSV file) was exported into Microsoft Excel (Microsoft Inc., USA) where normalization was performed. The resulting data were analyzed in the SIMCA-P 11.0 Software (Umetrics, Umea, Sweden) for multivariate statistical analysis. The supervised orthogonal partial least-squares (OPLS) were employed to process the acquired data and to identify the general separation and cluster. Then the differential variables were selected based on a threshold of variable importance in the projection (VIP) value (VIP>1.0) from OPLS model. Subsequently, those variables were validated at a univariate level using the nonparametric Wilcoxon-Mann-Whitney test by SPSS 18.0 (SPSS, Chicago, IL, USA) with the p-value set at 0.05 (Liao et al., 2012). The corresponding fold change showed how these selected metabolites varied between groups in male and female rats. Additionally,
compounds were identified by searching in NIST 2011 database. The Kyoto Encyclopedia of Genes and Genomes (KEGG) and Human Metabolome Database (HMDB) were used to give the biochemical interpretation of changed metabolites.

The biochemical parameters data were reported as mean ± Standard Deviations (SD). Statistical analysis was performed using SPSS 18.0 (SPSS Inc. Chicago, IL, USA). A two-way ANOVA was performed to examine the main effects of gender and exhaustive exercise and interaction on the measured variables. When a significant main effect or interaction was detected, data were subsequently analyzed by use of a post-hoc Bonferroni test. The mean differences in biochemical parameters levels between genders from control, exhaustive exercise and recovery group were also examined by running tests for simple main effects. Statistical significance was set at a p < 0.05.

Results

Effects of gender and exhaustive exercise on serum biochemical parameters

A two-way ANOVA was conducted to examine the effect of the gender and exhaustive exercise on T, CORT, BUN, MDA, GSH, GSSG, CK, and LDH; and the results are shown in Figure 1. The T levels were significantly affected by gender (F = 269.28, p = 0.00) and exhaustive exercise (F = 19.61, p = 0.00). However, no significant gender × exhaustive exercise interaction (F = 2.28, p = 0.12) was found for T levels, which indicated the male and female rats exhibited similar patterns of response for T levels. Subsequent post hoc analyses revealed significantly lower values for T in exhaustive exercise group compared to control group. In addition, a significant increase of T levels was observed from exhaustive exercise to recovery in male and female rats. There was a significant interaction effect between gender and exhaustive exercise on CORT level (F=5.89, p=0.008). Analyses for simple main effects indicated that the CORT levels were greater for the male rats than the female rats in the control, exhaustive exercise and recovery groups. The post hoc analyses revealed the CORT increased from control to exhaustive exercise, and returned to basal levels at recovery for the male and female rats. There was a significant interaction effect between gender and exhaustive exercise for the BUN levels (F = 5.90, p = 0.008). Analyses for simple main effects, female rats had higher BUN levels than the male in recovery group, however, no significant difference was found between male and female rats either in control or exhaustive exercise group. The post hoc analyses showed a significant increase of BUN levels from control to exhaustive exercise in male and female rats. From exhaustive exercise to recovery, a significant decrease of BUN values was observed in male rats whereas no changes were found in female rats.

The MDA levels were significantly affected by gender (F = 173.47, p = 0.00) and exhaustive exercise (F = 12.94, p = 0.00), while no significant interaction effect was found between gender and exhaustive exercise for MDA levels (F = 0.58, p = 0.57). The post hoc analyses revealed that the MDA levels were significantly higher in exhaustive exercise group when compared to the control in male and female rats. Further, the MDA levels significantly decreased from exhaustive exercise to recovery group in male and female rats, and back to basal values. The values of GSH, GSSG and GSH/GSSG were significantly affected by gender (GSH: F = 87.78, p = 0.00; GSSG: F = 69.75, p = 0.00; GSH/GSSG: F = 80.44, p = 0.00) and exhaustive exercise (GSH: F = 198.66, p = 0.00; GSSG: F = 241.91, p = 0.00; GSH/GSSG: F = 150.54, p = 0.00). No significant interaction effect between gender and exhaustive exercise in GSH and GSSG were obtained (GSH: F = 0.78, p = 0.47; GSSG: F = 1.80, p = 0.19), while a significant interaction (gender × exercise) was found for the GSH/GSSG (F = 5.91, p = 0.008). Subsequent post hoc analyses revealed that exhaustive exercise decreased GSH levels and increased GSSG levels in male and female rats and their ratio decreased after exercise in both groups. At recovery, the GSH and GSSG levels returned to basal values. In addition, male rats had higher levels of GSSG, as well as lower levels of GSH and GSH/GSSG than female rats in all three groups.

There was a significant interaction (gender × exercise) effect for the CK level (F = 19.87, p = 0.00). Analyses for simple main effects, male rats had higher CK levels than female rats in all three groups. The post hoc analyses revealed that the CK levels were significantly higher in exhaustive exercise and recovery group when compared to the control for the male and female rats. In addition, male rats had higher levels of CK at recovery compared to exhaustive exercise. The LDH level was significantly affected by gender (F = 6.408, p = 0.017) and exhaustive exercise (F = 76.97, p = 0.00), but no significant interaction effect between gender and exhaustive exercise (F = 0.79, p = 0.46) was found. The post hoc analyses revealed that the LDH levels increased from control to exhaustive exercise group for male and female rats. At recovery, however, LDH levels were still greater than the control. The LDH values in female rats were lower than male rats in the recovery group, but no significant differences were found in either the control or exhaustive exercise group.

Metabolomics differences between male and female rats

The data matrix was introduced into SIMCA-P 11.0 Software for multivariate statistical analysis. The OPLS model was employed to process the acquired data to distinguish the separation between groups of observation and to understand which variables contribute to the class-separating. Figure 2A illustrated the score plots of the male and female rats from control, exhaustive-exercise (10-day swimming) and recovery groups (7-day recovery). The X-axis (t1) and Y-axis (t2) indicated the first principle component and second principle component. One point represents one observation sample and the closer the points, the greater the similarity of the samples’ metabolites composition; vice versa, the greater the distance between points, the greater difference in composition. The four principal-component model explained 90.2% R²Y and predicted 82.8% Q²Y of the data according to the cross-validation. This model also explained 65.8% of the variables (R²X).
Gender-based difference within exhaustive exercise

Figure 1. Effects of gender and exhaustive exercise on the levels of T, CORT, BUN, MDA, GSH, GSSG, CK and LDH in male and female rats. The data were analyzed by a two-way ANOVA with post-hoc Bonferroni test and statistical significance was set at a $p<0.05$. Values are shown as mean ± SD (n=10). * $p<0.05$: significant change compared with the control group in male or female rats. # $p<0.05$: significant change compared with the exhaustive exercise group in male or female rats. § $p<0.05$: significant change between male and female rats from control, exhaustive exercise and recovery group. S-d10: 10-day swimming (exhaustive exercise group); R-d7: 7 days recovery.

According to the metabolic patterns and the value of $R^2$ and $Q^2$, a distinct cluster of metabolites in male and female rats undergoing control group, exhaustive exercise group and one-week recovery group was found. The observation samples for the control, exhaustive exercise and one-week recovery group from male rats are distributed in the upper part of the Figure 2A and those for female rats are in the lower part, indicating that the model can differentiate between male and female rats at control, exhaustive exercise and one-week recovery stages. Additionally, both in male and female rats, the samples from the exhaustive exercise group are clearly separated from the control group, while those for recovery group are closer to the control group.

To further evaluate the endogenous variations between the male and female rats, the loading plot (Figure 2B) was employed to display the specific metabolites that differentiated the male and female within the control, exhaustive exercise and recovery groups. Each point represents one variable. The differentially expressed metabolites between male and female rats in those three groups were distinguished based on the VIP values (VIP>1.0). Then, univariate statistical analysis, Wilcoxon-Mann-Whitney test (non-normally distributed data), was performed on those variations to evaluate their significance. Male control rats exhibited notably lower levels of citric acid, 3-hydroxybutyric acid and oleic acid as compared with female control rats. Within exhaustive exercise, male rats had
higher levels of citric acid, valine, leucine, linoleic acid, oleic acid, 3-hydroxybutyric acid and phosphate than those in female rats. Compared with female recovery group, male rats exhibited significant elevation of citric acid, valine, leucine, oleic acid, 3-hydroxybutyric acid and phosphate. The results are presented in Table 1.

**Figure 2.** The score plot of OPLS analysis to compare the metabolome of male and female rats from control, exhaustive exercise and recovery groups (n=10 per group). A: The score plot showed distinct clustering of metabolites with gender and exhaustive exercise periods. Metabolic patterns were grouped by different colored points. One point represents one sample. B: The loading plot was created to display the specific metabolites positively correlated with score plot. Each point presents one variable (metabolite). The bigger red triangle (▲) indicates statistically significant difference in variables that differentiated these groups. S-d10: 10-day swimming (exhaustive exercise group); R-d7: 7-day recovery.

**Table 1.** Altered plasma metabolites between male and female rats in the control, exhaustive exercise and recovery groups (n=10).

<table>
<thead>
<tr>
<th>Compounds</th>
<th>RT(min)</th>
<th>VIP</th>
<th>MC/FC FC</th>
<th>MS-d10/FS-d10 FC</th>
<th>MR-d7/FR-d7 FC</th>
</tr>
</thead>
<tbody>
<tr>
<td>Amino acid</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Valine</td>
<td>7.68</td>
<td>1.57</td>
<td>2.80</td>
<td>&lt;0.001</td>
<td>2.80</td>
</tr>
<tr>
<td>Leucine</td>
<td>8.48</td>
<td>1.63</td>
<td>2.80</td>
<td>&lt;0.001</td>
<td>2.11</td>
</tr>
<tr>
<td>Fatty acid</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Linoleic acid</td>
<td>27.65</td>
<td>1.62</td>
<td>2.35</td>
<td>0.002</td>
<td></td>
</tr>
<tr>
<td>Oleic acid</td>
<td>27.73</td>
<td>1.84</td>
<td>0.46</td>
<td>0.005</td>
<td>2.29</td>
</tr>
<tr>
<td>Organic acid</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>3-Hydroxybutyric acid</td>
<td>6.88</td>
<td>2.14</td>
<td>0.45</td>
<td>0.004</td>
<td>2.80</td>
</tr>
<tr>
<td>Citric acid</td>
<td>19.91</td>
<td>1.78</td>
<td>0.37</td>
<td>&lt;0.001</td>
<td>2.64</td>
</tr>
<tr>
<td>Other metabolites</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Phosphate</td>
<td>8.65</td>
<td>1.61</td>
<td>2.80</td>
<td>&lt;0.001</td>
<td>2.80</td>
</tr>
</tbody>
</table>

RT: The retention time of each compound from GC-MS; VIP: Variable importance in the project; FC: Fold change. Mean ranks were calculated using the nonparametric Wilcoxon-Mann-Whitney test by SPSS 18.0, and FC was obtained by comparing those metabolites’ mean ranks between male and female rats in control, exhaustive exercise and recovery group. FC with a value >1 indicates a relatively higher level present in male rats as compared to the female rats, while a value <1 means a relatively lower level in male rats as compared to the female rats. S-d10: 10 days swimming group (exhaustive exercise group); R-d7: 7 days recovery group; p: Correlation coefficient from Mann-Whitney test. p<0.05 means the change is significant.
Gender-based difference within exhaustive exercise

Figure 3. OPLS score plots of male rats (A) and female rats (B) over the whole exercise program (n=10). Both for the male and female rats, clear movement of data points along with exhaustive exercise period and recovery period was shown, indicating that exhaustive exercise had significant impact on plasma metabolome, and that one-week recovery could attenuate the exhaustive exercise induced metabolic perturbation in rats. Metabolic patterns were grouped by different colored points, one point represents one sample.

Metabolic profiles of male and females induced by exhaustive exercise

The OPLS model was established to analyze metabolic profiles of male and female rats induced by exhaustive exercise. Figure 3 (A, B) illustrates metabolomics movements of the control, 5-day swimming, 10-day swimming, 3-day recovery and 7-day recovery samples from male and female rats, respectively (Male: R²X = 0.472, R²Y = 0.932, Q²Y = 0.811; female: R²X = 0.324, R²Y = 0.820, Q²Y = 0.727). The observation samples for the control and 7-day recovery are located on the left of the Figure 3, whereas those for the 5-day swimming and 10-day swimming are on the right; this plot shows that the metabolic profile of the 10-day swimming (exhaustive exercise group) is the furthest one from the control. As the number of post exercise resting days increased, the metabolic profiles moved closer to the control group. Since clear movement of data points along with exhaustive exercise and recovery periods has been shown, it indicated that exhaustive exercise had significant impact on plasma metabolome both in male and female rats.

Endogenous variations contributing to the classification were determined in loading plots. In male rats, a total of 13 variations were selected over the exercise periods (Table 2). Exhaustive exercise decreased the levels of lactic acid, glucose, serine, glutamic acid, and increased the levels of critic acid, fumaric acid, butanedioic acid, valine, leucine, oleic acid, linoleic acid, 3-hydroxybutyric acid, and phosphate in male rats. Among the variations, 7 metabolites in the recovery group still showed significant differences compared with the control, indicating that not all the metabolites were normalized within the one-week recovery period.

In female rats, 13 variations were identified (Table 2). Exhaustive exercise significantly increased the levels of valine, leucine, methionine, 5-oxo-proline, glutamic acid, ornithine, linoleic acid, oleic acid, arachidonic acid, 3-hydroxybutyric acid, as well as urea, phosphate, and
myo-inositol. Within the one-week recovery, the levels of the amino acids valine, leucine, methionine, 5-oxo-proline, glutamic acid and ornithine were still higher in comparison to the control group. The fatty acids linoleic acid, oleic acid and arachidonic acid diminished within the one-week recovery period and showed no significant differences compared with the control.

Table 2. Altered plasma metabolites in the control, exhaustive exercise and recovery groups for male and female rats (n=10).

<table>
<thead>
<tr>
<th>Compounds</th>
<th>RT (min)</th>
<th>VIP</th>
<th>S-d10/Control</th>
<th>R-d7/Control</th>
<th>R-d7/S-d10</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>FC (Male)</td>
<td>p</td>
<td>FC (Male)</td>
</tr>
<tr>
<td>Valine</td>
<td>7.68</td>
<td>1.43</td>
<td>1.88</td>
<td>0.016</td>
<td>2.62</td>
</tr>
<tr>
<td>Serine</td>
<td>8.25</td>
<td>1.29</td>
<td>0.51</td>
<td>0.010</td>
<td>5.80</td>
</tr>
<tr>
<td>Leucine</td>
<td>8.48</td>
<td>1.67</td>
<td>2.09</td>
<td>0.005</td>
<td>2.82</td>
</tr>
<tr>
<td>Methionine</td>
<td>12.88</td>
<td>1.45</td>
<td>2.75</td>
<td>&lt;0.001</td>
<td>1.69</td>
</tr>
<tr>
<td>5-oxo-proline</td>
<td>12.93</td>
<td>1.71</td>
<td>2.82</td>
<td>&lt;0.001</td>
<td>2.82</td>
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<tr>
<td>Glutamic acid</td>
<td>15.05</td>
<td>1.25</td>
<td>0.41</td>
<td>&lt;0.001</td>
<td>2.56</td>
</tr>
<tr>
<td>Ornithine</td>
<td>19.65</td>
<td>1.34</td>
<td>2.82</td>
<td>&lt;0.001</td>
<td>2.82</td>
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<tr>
<td>Lactic acid</td>
<td>5.56</td>
<td>1.68</td>
<td>0.35</td>
<td>&lt;0.001</td>
<td>0.35</td>
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<tr>
<td>3-Hydroxybutyric acid</td>
<td>6.58</td>
<td>1.96</td>
<td>2.22</td>
<td>&lt;0.001</td>
<td>2.62</td>
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<tr>
<td>Fumaric acid</td>
<td>9.55</td>
<td>1.16</td>
<td>2.18</td>
<td>0.003</td>
<td>2.56</td>
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<td>Butanedioic acid</td>
<td>12.37</td>
<td>1.26</td>
<td>2.50</td>
<td>&lt;0.001</td>
<td>0.47</td>
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<td>Citric acid</td>
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<td>1.81</td>
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<td>Linoelic acid</td>
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<td>Oleic acid</td>
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<td>2.23</td>
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<td>Arachidonic acid</td>
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<td>D(+)-Glucose</td>
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<td>Urea</td>
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<tr>
<td>Phosphate</td>
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<td>1.18</td>
<td>2.28</td>
<td>0.002</td>
<td>2.00</td>
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<td>Myo-inositol</td>
<td>26.17</td>
<td>1.31</td>
<td>1.80</td>
<td>0.023</td>
<td>2.18</td>
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</table>

Discussion

The current study demonstrated significant gender-related differences in biochemical parameters and metabolic responses to exhaustive exercise. It was found that the gender was a significant determinant of the effects of exhaustive exercise on the CORT, BUN, CK, and the ratio of GSH to GSSG. The levels of T, LDH, MDA, GSH and GSSG were affected by gender and exhaustive exercise, however, no significant interaction effects were found. Exhaustive exercise increased the rates of energy metabolism, glucose metabolism, amino acid catabolism and fatty acid metabolism in male rats, whereas, female rats showed an increased propensity to oxidize lipid and conserve carbohydrate and protein against physical exercise.

Biochemical parameters

It has been proven that endogenous hormones are essential for physiological adoptions during exercise and influence the recovery phase by modulating anabolic and catabolic processes (Urhausen et al., 1995). In the current study, serum CORT, which was considered as a catabolic hormone (Elloumi et al., 2003), was significantly increased both in male and female rats within exhaustive exercise period, whereas the levels of T (which was considered as anabolic hormone) (Elloumi et al., 2003) decreased in male and female rats, indicating that exhaustive exercise affected the balance of hormone adjustment (Lac and Maso, 2004). Within the recovery period, the CORT and T levels returned to base values in both male and female rats. The basal levels of urea derived from nitrogen metabolism are reported to be amplified by intensive exercise (Haralambie and Berg, 1976). In the current study, the BUN levels in male and female rats were significantly increased by exhaustive exercise, suggesting an increased breakdown of nitrogen-containing compounds, in line with a reported rise in blood serum urea after heavy exercise (Haralambie and Berg, 1976). Within the recovery period, the BUN levels returned to base values in male rats, whereas no significant change was found in female rats compared to exhaustive exercise group, which was considered as to be an indicator of insufficient regeneration with recovery period.

Regarding the fact that GSH/GSSG in blood plasma stand for clinical measure of oxidative stress (Johns, 2006), reduction in plasma GSH and GSH/GSSG ration causes a shift in redox balance towards a more oxidizing environ-
ment. In other words, significant increase in GSSG and further decrease in the GSH/GSSG immediately after exhaustive exercise probably indicate elevation in free-radical production (Aguiló et al., 2005). In the current study, as compared with the control group, significant increases in GSSG level, as well as significant decreases in GSH level and GSH/GSSG were found within exhaustive exercise in plasma from male and female rats, reflecting exercise-induced oxidative stress. MDA is a byproduct of oxidation of polyunsaturated fatty acid which has been used as a marker of lipid peroxidation (Powers et al., 2016; Vollaard et al., 2005). In the present study, MDA level was increased within exhaustive exercise and gender showed similar alteration, which also imply that exercise-induced oxidative stress occurred both in male and female rats. Our results are consistent with the research by Goldfarb et al. (2007). In addition, female rats exhibited higher levels of GSH and GSH/GSSG than male rats in all three groups. Similarly, some studies have found higher levels of antioxidant molecules or lower levels of oxidative molecules in the blood in females than in males (Balog et al. 2006, Goldfarb et al. 2007). Another experimental study in rats suggested that the mitochondria from females exhibited higher antioxidant capacity than in males at rest (Borrás et al., 2003). Our data also revealed that female rats had lower levels of MDA than male rats in all three groups. Several studies have reported that the MDA levels in women are lower than in men at rest (Bloomer et al., 2008). These differences between males and females have been attributed to estrogen action (Borrás et al., 2003). Taken together, the present study does support the findings of Goldfarb et al. (2007) that females are slightly better protected from oxidative stress than males. However, male and female rats showed similar changes to the oxidative stress markers undergoing the exhaustive exercise.

For the most part, increased levels of tissue enzymes in serum have been treated as indirect indicators of increased cell permeability caused by tissue membrane damage (Kanter et al., 1988). In the current study, post-exercise CK and LDH values were significantly greater than the control both in male and female rats, suggesting that exhaustive exercise might cause cell membrane damage. Additionally, persistently elevated CK and LDH levels within recovery period in male and female rats were suggestive of over-exercise. Besides these effects, male rats had higher CK and LDH levels at recovery than female rats, which is consistent with the intriguing work of Sewright et al. (2008), who have reported that male rats showed a large CK response 7 d after exercise than female rats.

**Plasma metabolomics**

Exhaustive exercise significantly increased the tricarboxylic acid cycle (TCA) intermediates levels including citric acid, fumaric acid and butanedioic acid in male rats. However, no changes in TCA intermediates levels were found in female rats. It is generally accepted that energy expenditure will be elevated with increasing exercise volume and intensity, and the TCA cycle will be activated accordingly to oxidize and produce more ATP for the body (Peake et al., 2014), with a concomitant, marked increase in plasma TCA intermediates (Gibala et al., 1988). Additionally, it has been known that carbohydrate oxidation is the primary ATP-producing system during exercise and subsequent to maintain high levels of TCA intermediates (Bowtell et al., 2007). However, since there were no significant changes of TCA intermediates in female rats, females might rely less on carbohydrate utilization and TCA cycling during exercise. In addition, although male rats had lower levels of citric acid than females in the control group, male levels became higher than in females within exhaustive exercise, which also suggested that males might rely more on the TCA cycle to provide energy for the body. During exhaustive exercise, glucose is an important energy source supplying most part of the oxidative energy production (Hargreaves, 2015; Jensen and Richter, 2012). In the current study, the level of glucose was decreased in male rats undergoing exhaustive exercise, which suggested greatly increased energy expenditure. Moreover, the decrease in lactate content indicated a reduction in the whole body rate of glycosis, which was in agreement with previous report in which lowered lactate levels were found in mice doing the wheel running exercise (Monleón et al., 2014).

Generally, the increased levels of free fatty acids and ketone bodies suggest that lipolysis is involved (Romijn et al., 1993; Turcotte et al., 1992). Free fatty acids represent a major energy source supporting whole body energy flux in men and women (Friedlander et al., 1998). 3-Hydroxybutyrate is an important byproduct produced by oxidation of fatty acids in the liver and kidney (Liao et al., 2012). Current results reveal that the levels of fatty acids including oleic acid, linoleic acid and metabolite 3-hydroxybutyric acid were increased in male and female rats within exhaustive exercise, indicating enhanced lipolysis. In addition, these data show that male rats had lower levels of 3-hydroxybutyric acid and oleic acid than females in control group, but those two variations together with linoleic acid were significantly elevated and became higher than in female rats within exhaustive exercise, which suggest that female rats might oxidize more fatty acids for energy consumption during exhaustive exercise. Results of another study (Henderson et al., 2007) also suggest that fatty acids comprise a greater proportion of energy source in women than in men for a given relative work intensity. In our study, no significant changes of carbohydrate and TCA intermediates were found in female rats within exhaustive exercise, indicating that females might have an increased propensity to oxidize lipid and conserve carbohydrate against physical stress. It has been reported that the relative abundance of ovarian hormones in women might alter the regulation of metabolic pathways to favor fat oxidation and conserve carbohydrate (D’Eon and Braun, 2002; Ruby and Robergs, 1994). Other studies indicated that elevated estrogen and progesterone in women altered sensitivity to catecholamines and enhanced lipolysis and lipid utilization, consequently, carbohydrate utilization decreased (Braun and Horton, 2001; Horton et al., 1998). During the recovery period, the same trends were found in declining levels of those fatty acids for male and female rats.

Valine and leucine, the branched-chain amino acids
(BCAA\textsubscript{5}) participating in blood glucose regulation, were found to be significantly increased in male and female rats within exhaustive exercise. It has been recognized that skeletal muscle is a major site of BCAAs utilization (She et al., 2010). As a result of the reduced availability of the muscular glycogen during exercise, there were higher blood levels of BCAAs on account of increased proteolysis in skeletal muscle (Kume et al., 2015). The other amino acids serine and glutamic acid were significantly decreased in male rats, which are in agreement with the results of Dohm et al. (1981). They found significantly decreased levels of alanine, glutamate, and glutamine in plasma following exercise. In contrast, plasma concentrations of the BCAAs were generally elevated by exercise. These data were, thus, quite similar to those of Kasperek (1989).

The results suggest an increased metabolic use of these amino acids with exercise. Our data also show that male rats have higher valine and leucine levels than female rats in the exhaustive exercise group, whereas no significant differences are found between male control rats and female control rats for the levels of valine and leucine, which suggests that male rats might use more proteolysis to supply energy. Tarnopolsky et al. (1990) found that the lower insulin and higher epinephrine levels in males might partially explain the greater glycogenolysis and protein catabolism observed in the exercise group.

As mentioned before, plasma GSH couple GSH/GSSG is a critically important redox biomarkers (Seifi-Skishahr et al. 2016). Our data suggest that exhaustive exercise-induced oxidative stress were observed in both male and female rats. GSH, a tripeptide composed of glutamate, cysteine, and glycine, is a major non-enzymatic endogenous antioxidant (Ji, 2002). In the present study, elevated level of glutamate was detected in female rats within exhaustive exercise. It has been known that the cystine/glutamate antiporter is a plasma membrane-bound protein critically involved in glutamatergic transmission. Upon oxidative stress, it starts to pump out glutamate in exchange for cysteine, which is essential for glutathione synthesis (Lu, 2013). The increase in the influx of cysteine during increased glutathione synthesis results in increased export of glutamate to extracellular compartment and to the circulation, which ultimately results in the increase of extracellular glutamate levels (Tapiero et al., 2002; Anderson, 1998). Additionally, the levels of 5-oxoproline, methionine and arachidonic acid were also significantly elevated in female rats within exhaustive exercise, suggesting that exhaustive exercise induced diverse responses to oxidative stress in female rats. In our study, exhaustive exercise also increased the levels of urea and ornithine in female rats, which suggested that the urea cycle might be elevated in activity within exercise. Myo-inositol is the primary biologically active form of inositol which could improve glucose tolerance by insulin sensitivity (Croze et al., 2013). The increased levels of myo-inositol in female rats indicated the disorder of inositol metabolism caused by exhaustive exercise.

Actually, selected body fluid is important in detection of oxidative stress. GSH is found in all extracellular biological fluids and here we chose plasma as an extracellular fluid circulating between body cells and exchanging oxidative biomarkers. However, according to the study from Seifi-Skishahr et al. (2016), they found that the changes in erythrocyte GSH/GSSG are not parallel with plasma GSH/GSSG, which suggest that plasma redox biomarker might not accurately reflect tissue redox status compared with erythrocyte biomarkers. One of limitations of the current study is lack of data on erythrocyte oxidative stress biomarkers after exhaustive exercise which is suggested for future studies.

**Conclusion**

It is concluded that exhaustive exercise affects hormone balance and causes oxidative stress, and subsequent cell membrane damage both in male and female rats. Females are slightly better protected from oxidative stress than males. However, undergoing exhaustive exercise, male and female rats showed similar changes to the oxidative stress markers. Significant gender-related differences in metabolic profiles in male and female rats within exhaustive exercise occur, with increased rates of TCA cycling, glucose metabolism, amino acid catabolism and fatty acid metabolism in male rats, whereas, female rats might have an increased propensity to oxidize lipid and conserve carbohydrate and protein metabolism against physical stress. Disordered urea cycle and inositol metabolism also occur in females with exhaustive exercise. The one-week recovery period attenuated many, but not all, of the metabolic perturbation by exhaustive exercise in rats.

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**References**


Key points

- Exhaustive exercise affected hormonal balance and caused oxidative stress and subsequent cell membrane damage both in male and female rats.
- Female rats exhibited higher antioxidant capacity at rest, but showed similar changes to the oxidative stress markers as male rats with exhaustive exercise.
- A significant gender-based difference in the metabolic profiles was found in male and female rats within exhaustive exercise.
- Exhaustive exercise increased the rates of the TCA cycling, glucose metabolism, amino acid catabolism and fatty acid metabolism in male rats, whereas, female rats demonstrated an increase propensity to lipid utilization and conserve carbohydrate and proteolytic metabolism.

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