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fasting glucose concentration, insulin concentration, insulin resistance, homeostasis model assessment (homa), and matsuda index were evaluated at baseline and after the intervention. the matsuda index was calculated as 10,000 / (10,000 - fasting insulin) x (fasting glucose/fasting insulin). the matsuda index is a measure of insulin sensitivity, and a higher index indicates a better insulin sensitivity and beta-cell function. adipose tissue samples were collected from the sat of each individual at baseline and after the intervention and stored in liquid nitrogen or 10% neutral buffered formalin for histological analysis. adipocyte size was determined in sections stained with hematoxylin and eosin using a macroscopic assessment of the average adipocyte size based on size frequency distribution. cell size was measured with the image-pro plus software (media cybernetics, usa). the volume of each adipocyte was estimated by the software using a point grid on the background and the sections were annotated by a single observer (m.e.) who was blinded to the participant and treatment groups. two whole abdominal sat biopsies were removed from each participant at baseline and after the intervention. the tissues were minced in liquid nitrogen, homogenized in ice-cold protein extraction buffer and centrifuged at 11,000 rpm for 15 minutes. the supernatant was collected and stored at -80°c until the analyses of protein expression levels of ampk and pampk using western blotting. users are able to choose the software version according to their needs. they can also customize the options of their data acquisition and analysis such as the number of wells, cycles and temperature. furthermore, they can change the experimental settings of real-time pcr for amplification, melt curve analysis, and pcr product detection. they can also define the gel electrophoresis settings and the data export.

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if you do not have software to properly analyze the data, and if you have not performed the analysis already, the best thing that you can do is to contact the company that produced the data. tell them what happened. if they are willing to help, they will then tell you what you need to do to get the data into the software that you have. if you do not have software to properly analyze the data, and if you have not performed the analysis already, the best thing that you can do is to contact the company that produced the data. this part of the internet is provided to assist individuals in making a decision on which of the products to acquire and which ones they wish to refrain from acquiring. based on our experience we've created an unique testing procedure that enables us to tell whether a given software is, in fact, adware, and if so, which of the programs is adware. the purpose of using this service is to help you avoid getting adware on your computer if you see adware listed below, then you need to remove it as soon as possible, as it can have a negative impact on your computer's efficiency and security. fairview is a group of windows applications and software development tools designed to extract relevant information from large databases and archives quickly and efficiently. fairview can be used as an automatic way to search through many thousands of distinct document types, and we hope you will find it useful. for more information, please visit www.fairview.com . data were analyzed using graphpad prism 5 (graphpad software inc., ca). descriptive statistics were used to summarize the data, with data expressed as mean ± sd and analyzed by the student's t test or by one-way analysis of variance (anova) with bonferroni correction. linear regression analysis was used to test the relationship between variables. correlations were assessed by spearman's correlation coefficient. the significance level was set at p 5ec8ef588b

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