Day to Day and Leg to Leg Variation in Gene Expression

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ABSTRACT

BACKGROUND: Skeletal muscle biopsies are extensively used in research to determine the effects of exercise. Usually, sequential biopsies are taken before and after exercise in order to determine the influence of exercise on gene expression. Often it is not clear if the results are due to exercise or as a result of damage cause by the biopsy to the muscle. Previous studies have proven that multiple biopsies in the same leg cause stress to the muscle and such stress leads to inflammation and other response pathways that alter the gene expression. PURPOSE: The purpose of this study is to determine the day to day variance in skeletal muscle gene expression in the same leg and between legs to establish an experimental design that eliminates artifacts and reduces the number of biopsies needed for a study. METHODS: Eight participants had a muscle biopsy taken from the vastus lateralis on three separate occasions separated by 7 days. Before the biopsies were taken, all the subjects were instructed to not exercise and keep the same diet one day prior to the biopsy. Two biopsies were taken from one leg and a third biopsy was taken from the other. This will allow comparisons between and within legs on different days. A total of 24 muscle biopsies were analyzed to test the expression of genes using real-time PCR. Genes related to mitochondrial development were ERRa, GABPA, MEF2a, NRF1, PPARG, PGC1a, SIRT1, TFAM, VEGF, and FNDCS and stable reference genes B2M, CYC, GAPDH and RPS18. Gene expression was expressed using the 2^ΔΔCT and 2^ΔCT method. The stable reference control condition for 2^ΔΔCT was defined as the first biopsy. A one-way repeated measures analysis of variance (ANOVA) was used to determine statistical significance. All data is presented as mean ± standard error.

RESULTS

Eight healthy college age volunteers had a muscle biopsy taken from the vastus lateralis on three separate occasions separated by 7 days. Before the biopsies were taken, all the subjects were instructed to not exercise and keep the same diet one day prior to the biopsy. Two biopsies were taken from one leg and a third biopsy was taken from the other. This will allow comparisons between and within legs on different days. A total of 24 muscle biopsies were analyzed to test the expression of genes using real-time PCR. Genes related to mitochondrial development were ERRa, GABPA, MEF2a, NRF1, PPARG, PGC1a, SIRT1, TFAM, VEGF, and FNDCS and stable reference genes B2M, CYC, GAPDH and RPS18. Gene expression was expressed using the 2^ΔΔCT and 2^ΔCT method. The stable reference control condition for 2^ΔΔCT was defined as the first biopsy. A one-way repeated measures analysis of variance (ANOVA) was used to determine statistical significance. All data is presented as mean ± standard error.

CONCLUSION AND FUTURE DIRECTION

- Genes related to mitochondrial biogenesis are not statistically different between legs or between days except for FNDCS which was different between biopsy 1 and 2 but not different than biopsy 3.
- Two of the four stable reference genes were stable between biopsies and two were higher on biopsy 3 than the other biopsies. However, by using the geometric mean of all four reference genes these differences disappear.
- While many genes are statistically not different between legs and between days, researchers need to determine the amount of error that is acceptable in each study design.
- 12 more subjects are currently being analyzed to expand the sample size.
- These data will aid researchers utilizing the muscle biopsy technique in the best study design in terms of the appropriate timing of the biopsy in relation to subsequent biopsies.

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