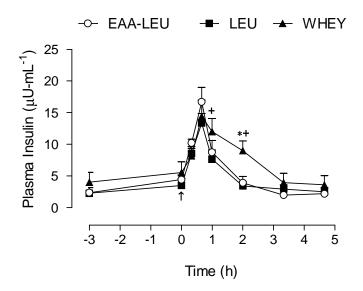
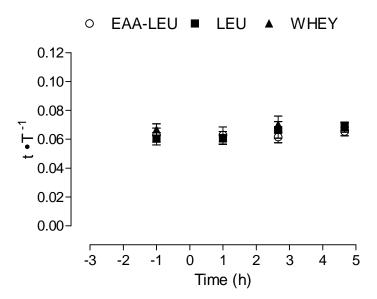
Online Supplemental Material Figure 1.



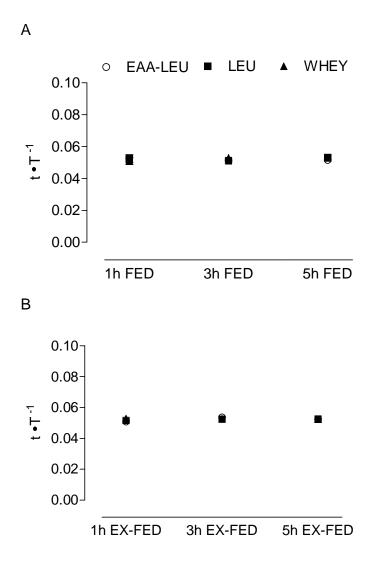
Supplemental Figure 1. Mean (\pm SEM) plasma insulin concentration (μ U-mL⁻¹) following EAA-LEU, LEU, and WHEY treatments. Upward arrow indicates time of treatment administration. *Significantly greater than EAA-LEU (P < 0.05); +Significantly greater than LEU (P < 0.05).

Online Supplemental Material Figure 2.



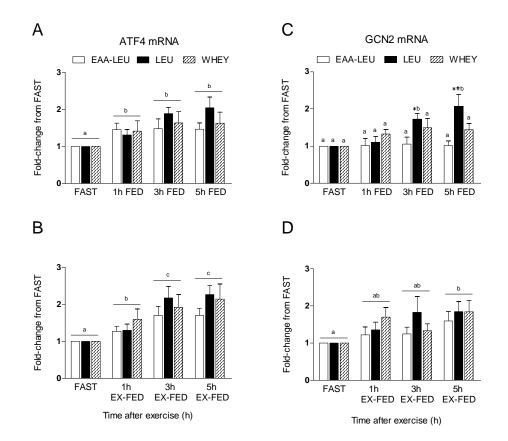
Supplemental Figure 2. Mean (\pm SEM) plasma free phenylalanine enrichments (tracer-to-tracee ratio t•T⁻¹) over time. Data were analyzed using a 2-factor (treatment × time) repeated measures ANOVA and linear regression. There were no differences between treatments (P = 0.66) or across time (P = 0.34). The slope of plasma free phenylalanine enrichment by time was not different from zero for any treatment group (EAA-LEU P = 0.95; LEU P = 0.11; WHEY P = 0.40).

Online Supplemental Material Figure 3.



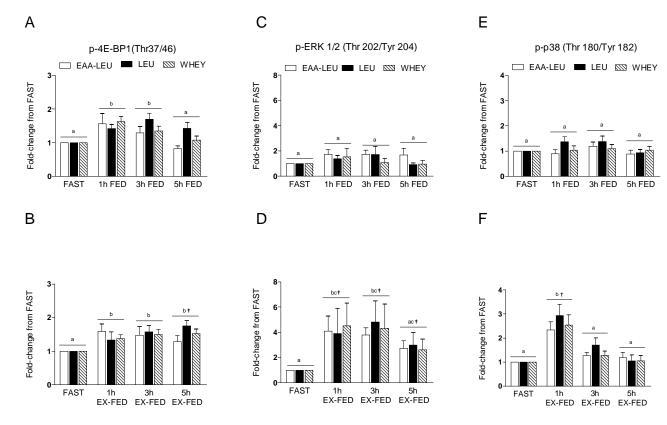
Supplemental Figure 3. Mean (\pm SEM) intracellular free phenylalanine enrichments (tracer-to-tracee ratio – t•T⁻¹) in both FED (A) and EX-FED (B) conditions. Conditions (i.e. FED and EX-FED) were analyzed separately using a 2-factor (treatment \times time) repeated measures ANOVA (FED: time P = 0.92; treatment, P = 0.90. EX-FED: time P = 0.30; treatment P = 0.88). Condition effects were analyzed separately at 1 (P = 0.90), 3 (P = 0.42), and 5h (P = 0.98) post-exercise recovery using a 2-factor (treatment \times condition) ANOVA. Data were also analyzed using linear regression for the difference of the linear regression slope from zero (EAA-LEU FED P = 0.77, EX-FED P = 0.41; LEU FED P = 0.84, EX-FED P = 0.68; WHEY FED P = 0.56, EX-FED P = 0.84).

Online Supplemental Material Figure 4.



Supplemental Figure 4. Mean (\pm SEM) mRNA expression of ATF4 (A and B) and GCN2 (C and D) (expressed as fold-difference from FAST) at 1, 3, and 5 post-exercise recovery in both FED and EX-FED conditions following EAA-LEU, LEU, and WHEY treatments. Times with different letters are significantly different from each other within that treatment and condition. *Significantly greater than EAA-LEU within that time and condition (P < 0.05); ‡Significantly greater than WHEY within that time and condition (P < 0.05).

Online Supplemental Material Figure 5



Supplemental Figure 5. Mean (\pm SEM) phosphorylation status of 4E-BP1^{Thr 37/46} (A and B), ERK 1/2^{Thr 202/Tyr 204} (C and D) and p38^{Thr 180/Tyr 182} (E and F) (expressed as fold-difference from FAST) at 1, 3, and 5 post-exercise recovery in both FED and EX-FED conditions following EAA-LEU, LEU, and WHEY treatments. Times with different letters are significantly different from eachother within that treatment and condition. †Significantly greater than EX-FED condition at that time-point; all P < 0.05.

Online Supplemental Material Figure 6

Supplemental Figure 6. Representative blot images for p-Akt Ser473 , p-mTOR Ser2448 , p-p70S6k Thr389 , p-4E-BP1 $^{Thr 37/46}$, p-ERK $1/2^{Thr202/Tyr204}$, p-p38 $^{Thr180/Tyr182}$, and α -tubulin at FAST, 1h EX-FED, 1h FED, 3h EX-FED; 3h FED, 5h EX-FED, and 5h FED following EAA-LEU, LEU, and WHEY treatments. Image contrast adjusted to improve clarity.

