NMR-based metabolomic analysis for the effects of alanyl-glutamine supplementation on C2C12 myoblasts injured by energy deprivation

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Figure Legends

- **Figure S1.** A representative 1D ¹H NMR spectrum of Ala-Gln dissolved in D₂O recorded on 850 MHz NMR spectrometer at 298K.
- **Figure S2.** A representative 2D ¹H-¹³C HSQC spectrum of aqueous extracts derived from C2C12 myoblast cells recorded on 850 MHz NMR spectrometer at 298K in PBS (pH 7.4).
- **Figure S3.** A selected region (0.8-2.8 ppm) of a representative 2D ¹H-¹H TOCSY spectrum of aqueous extracts derived from C2C12 cells recorded at 298K in PBS (pH 7.4). Resonance assignments are labeled.
- **Figure S4.** A selected region (2.0-4.6 ppm) of a representative 2D ¹H-¹H TOCSY spectrum of aqueous extracts derived from C2C12 cells recorded at 298K in PBS (pH 7.4). Resonance assignments are labeled.
- **Figure S5.** A selected region (5.5-9.5 ppm) of a representative 2D ¹H-¹H TOCSY spectrum of aqueous extracts derived from C2C12 cells recorded at 298K in PBS (pH 7.4). Resonance assignments are labeled.

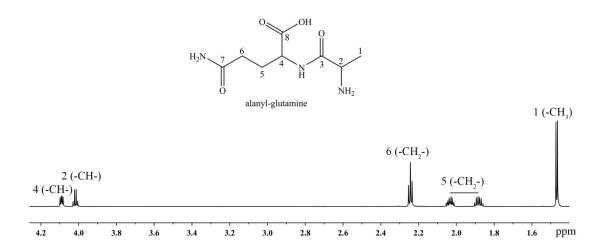


Fig. S1 A representative 1D 1 H NMR spectrum of Ala-Gln dissolved in D₂O recorded on 850 MHz NMR spectrometer at 298K.

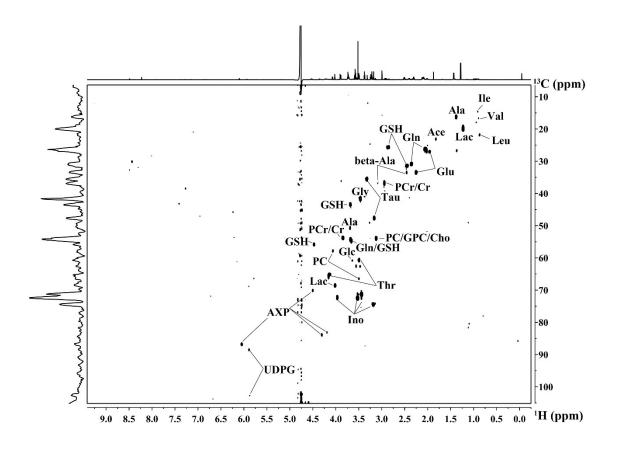


Fig. S2 A representative 2D ¹H-¹³C HSQC spectrum of aqueous extracts derived from C2C12 myoblast cells recorded on 850 MHz NMR spectrometer at 298K in PBS (pH 7.4).

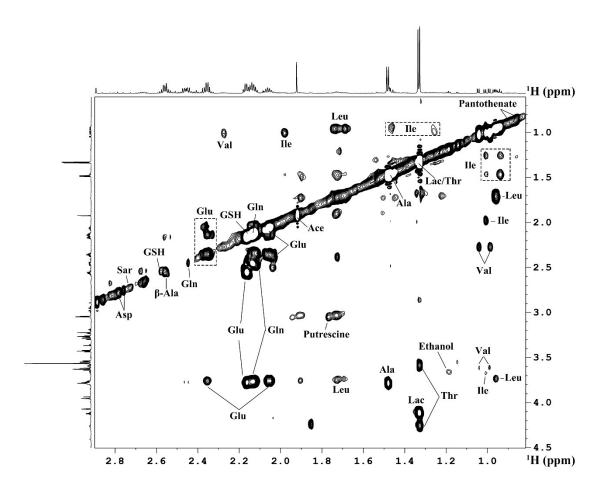


Fig. S3 A selected region (0.8-2.8 ppm) of a representative 2D ¹H-¹H TOCSY spectrum of aqueous extracts derived from C2C12 cells recorded at 298K in PBS (pH 7.4). Resonance assignments are labeled.

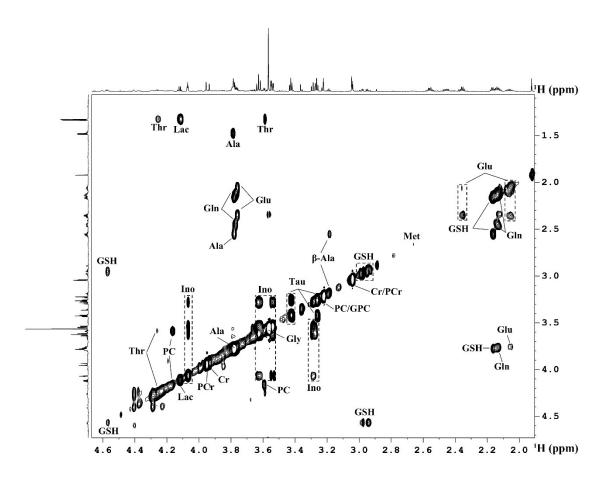


Fig. S4 A selected region (2.0-4.6 ppm) of a representative 2D ¹H-¹H TOCSY spectrum of aqueous extracts derived from C2C12 cells recorded at 298K in PBS (pH 7.4). Resonance assignments are labeled.

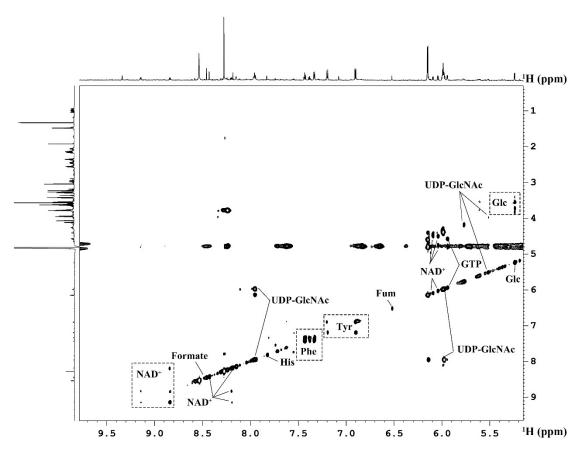


Fig. S5 A selected region (5.5-9.5 ppm) of a representative 2D ¹H-¹H TOCSY spectrum of aqueous extracts derived from C2C12 cells recorded at 298K in PBS (pH 7.4). Resonance assignments are labeled.