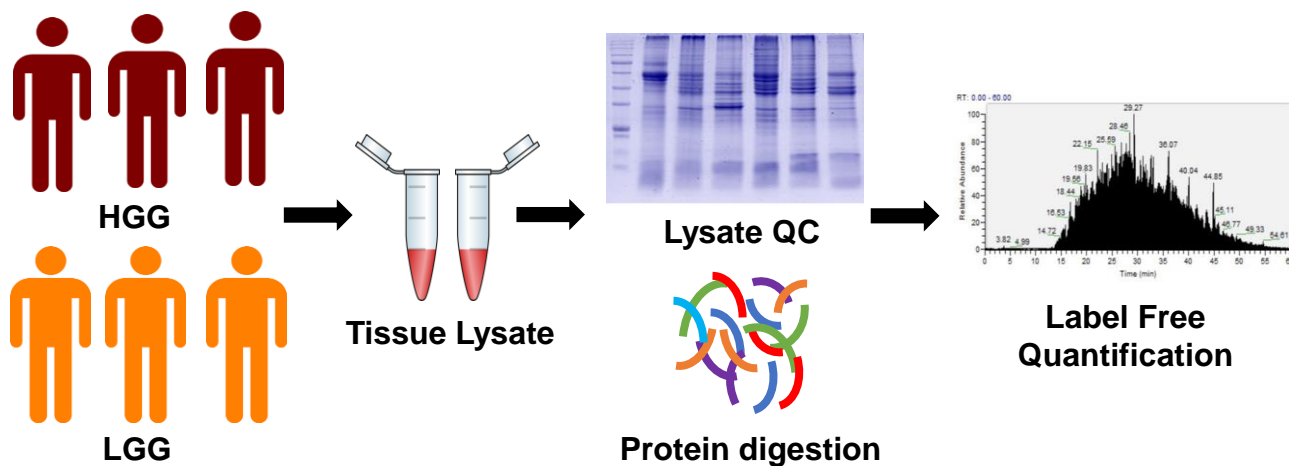
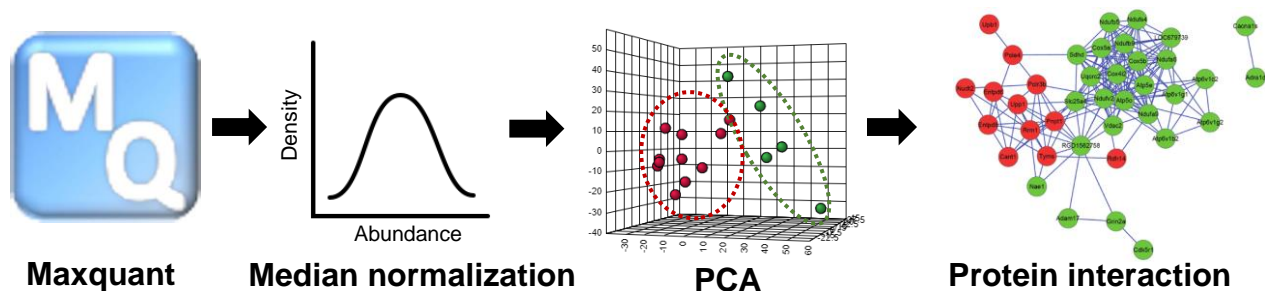


A. Sample collection and processing for MS analysis



B. Data normalization and Statistical data analysis



C. Validation using multiple reaction monitoring



D. *In-silico* Molecular Docking of Target proteins

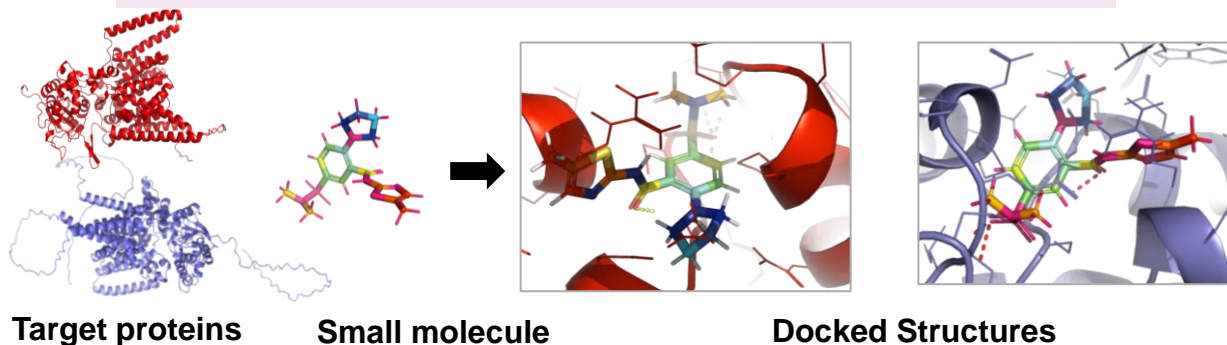


Figure 1