

Creative Proteomics

Cutting-edge Bioinformatics Solutions for Proteomics, Metabolomics, & Proteins

- Experienced Staff: Ph.D. team has an average of 10+ years' hands-on experience.
- Custom-tailored & Modularized Services: Analytical solutions customized for your research.

♦ High-quality: Validated process to ensure reproducibility of your results.



Featured Bioinformatics Services at Creative Proteomics

Equipped with advanced computing instruments and software, Creative Proteomics can provide cutting-edge bioinformatics services to accommodate a wide range of needs from basic to complex.

Bioinformatics for Proteomics

- ♦ Functional Annotation & Enrichment Analysis: GO, KEGG, domain annotation and enrichment, *etc.*
- ♦ Cluster Analysis: Hierarchical clustering, K-means clustering, STEM analysis, and others
- ♦ Network analysis: PPI analysis, IPA analysis, Gene co-expression network analysis
- Subcellular localization: A significant part of bioinformatics-based prediction of protein function and genome annotation

Bioinformatics for Metabolomics

- ♦ Univariate analysis: Volcano plot, T-test analysis, Fold change analysis, *etc*.
- ♦ Multivariate analysis: PCA, PLS-DA, OPLS-DA
- ♦ Clustering analysis: Heatmap, K-means, SOM
- ♦ Identification of potential biomarkers
- Correlation network of differential metabolites

Bioinformatics for Proteins

- ♦ Protein site prediction: Amino acid composition, Signal peptide prediction, Protein motif prediction, etc.
- Protein structure prediction: Homology modeling, Molecular docking, Structure features analysis, Secondary structure prediction, *etc*.
- ♦ **Protein evolution Analysis:** Sequence-based analysis, Structure-based analysis, Genome-based analysis

