

## FUNCTIONAL PROTEOMICS%0A

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[Functional genomics reveals that Clostridium ...](#)  
Functional classification of *C. difficile* Spo0A regulated genes. Enriched functional gene classes of genes (a) downregulated or (b) upregulated in the *C. difficile* spo0A mutant. The number of genes present in each functional class based on RNAseq (grey) and proteomics (black) analysis.

[Functional genomics reveals that Clostridium difficile ...](#)  
Background. *Clostridium difficile* is an anaerobic, Gram-positive bacterium that can reside as a commensal within the intestinal microbiota of healthy individuals or cause life-threatening antibiotic-associated diarrhea in immunocompromised hosts.

[pSITE Vectors for Stable Integration or Transient ...](#)  
Plant functional proteomics research is increasingly dependent upon vectors that facilitate high-throughput gene cloning and expression of fusions to autofluorescent proteins.

[Proteomics/Protein Primary Structure/Alternative Splicing ...](#)  
Alternative Splicing is a major source of protein diversity in living organisms. It has been estimated that at least 30% of all genes in the human genome are alternatively spliced and this number continues to expand. It was originally thought that the number of alternatively spliced genes accounted

[Structural genomics of protein phosphatases](#)  
The impact of various structural genomics efforts on our structural and functional understanding of the human protein phosphatases and protein phosphatases from a wide range of biomedically-relevant pathogens is already apparent. Current coverage of the human protein phosphatome is ~45%, with the promise of many more structures to come within the next few years. These data will provide

[Metabolic Maturation during Muscle Stem Cell ...](#)  
Furthermore, our study highlights the benefits of genetic approaches to unveil functions of miRNAs in vivo, in particular when studying miRNAs that are highly abundant (miR-1), serve overlapping functions (miR-1-1 and miR-1-2), or represent a functional unit (miR-1 and miR-133a).

[BBA - Proteins and Proteomics](#)  
The functional diversity of IDPs/IDPRs may be attributed to their potential promiscuity in binding to different structurally unrelated partners in living cells [2].

[Research Sabbatical to Explore Molecular and Functional ...](#)

research sabbatical to explore molecular and functional genomics approaches for foodborne pathogen investigations

**Department of Molecular Biology**

Research in Molecular Biology at Princeton addresses biological questions in five broad areas.