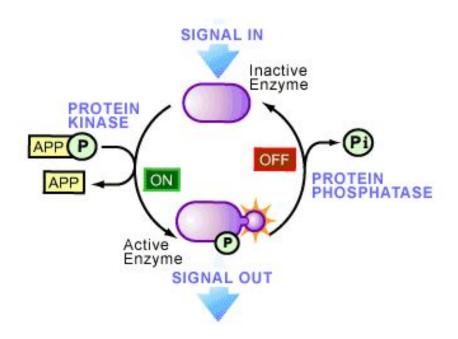
Cell Type Deconvolution by Phosphoproteomics

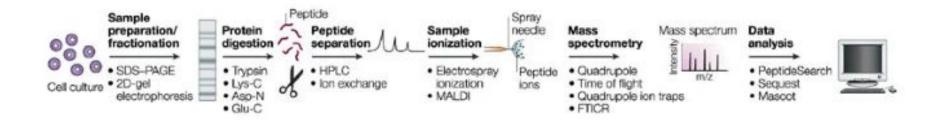
Background

- Signal transduction
- Protein phosphorylation
 - Kinases
 - Phosphatases
 - Adapter proteins
- Cancer



Background

- Phosphoproteomics
- Mass spectrometry



Project Overview

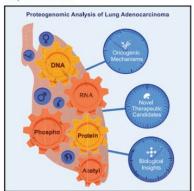
- Identify cell types in tumor tissue samples based on bulk phosphoproteomic data
- Data from a 2020 study in Cell
- Started by doing correlation analysis



Resource

Proteogenomic Characterization Reveals Therapeutic Vulnerabilities in Lung Adenocarcinoma

Graphical Abstract



Highlights

- Comprehensive LUAD proteogenomics exposes multi-omic clusters and immune subtypes
- Phosphoproteomics identifies candidate ALK-fusion diagnostic markers and targets
- Candidate drug targets: PTPN11 (EGFR), SOS1 (KRAS), neutrophil degranulation (STK11)
- Phospho and acetyl modifications denote tumor-specific markers and druggable proteins

Authore

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In Brief

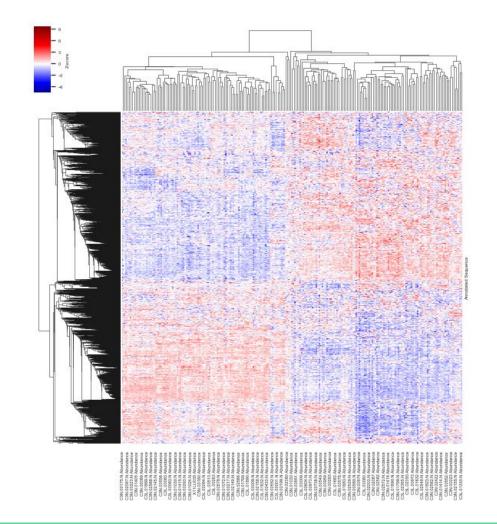
Comprehensive proteogenomic characterization of lung adenocarcinomas and paired normal adjacent tissues from patients of diverse smoking status and country of origin yields insights into cancer taxonomy, oncogenesis, and immune response; offers novel candidate biomarkers and therapeutic targets; and provides a community resource for further discovery.





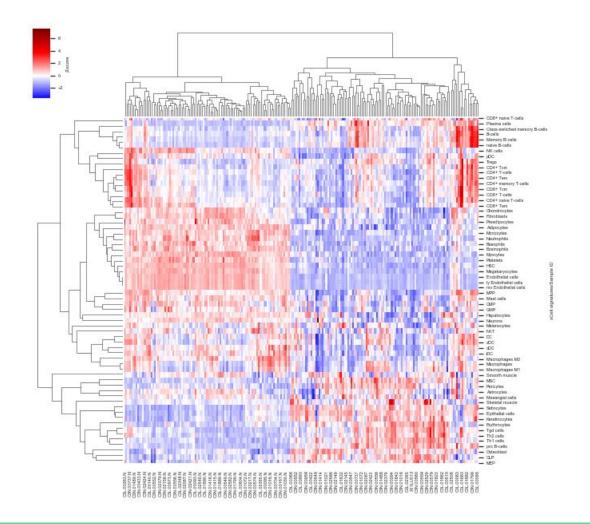
Phosphopeptide vs. Sample Heatmap

- 5256 peptides
- 211 samples

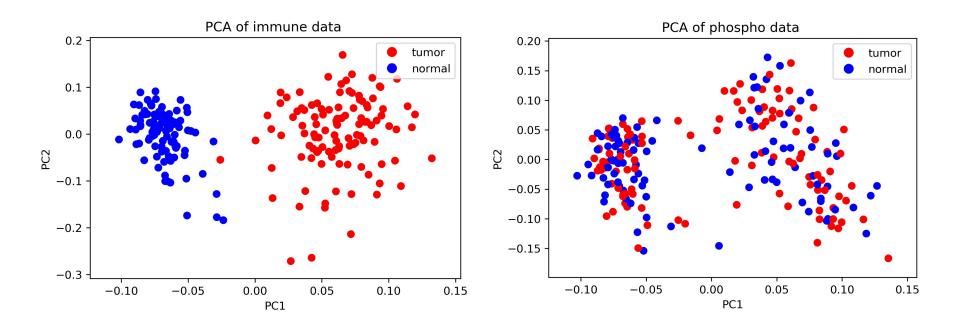


Immune Cell vs. Sample Heatmap

- 64 immune cells
- 211 samples

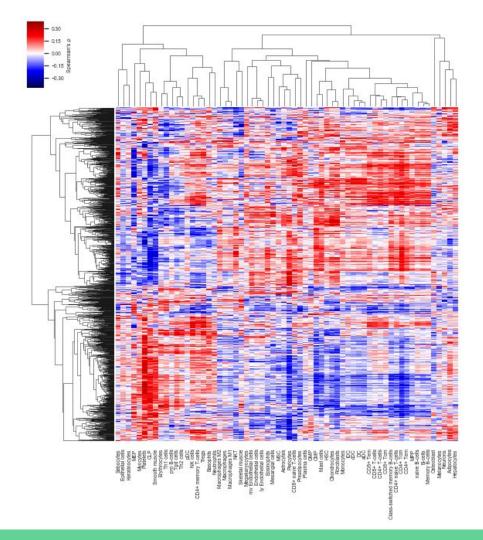


Principal Component Analysis (PCA)

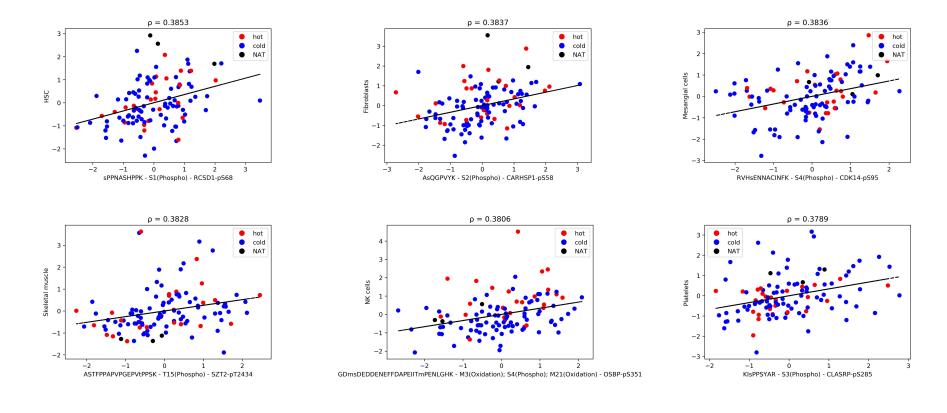


Phosphopeptide vs. Immune Cell Heatmap

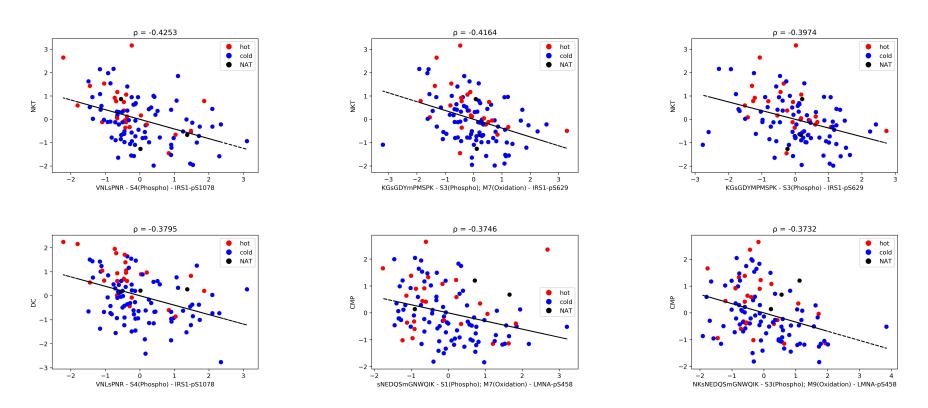
- 1584 peptides
- 64 immune cells



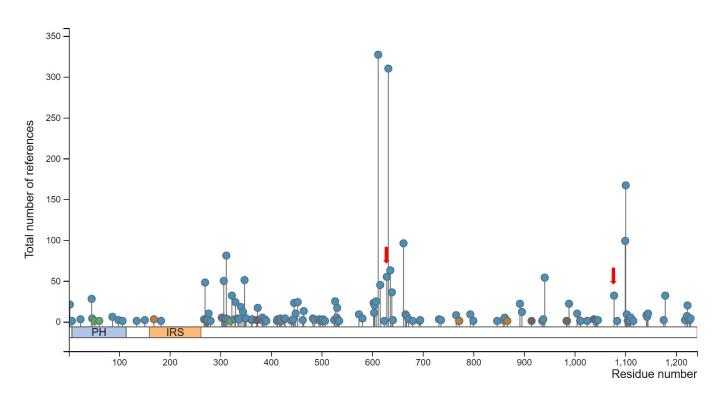
Most Correlated Peptides and Immune Cells



Least Correlated Peptides and Immune Cells



Insulin Receptor Substrate 1 (IRS1)



Future Work

- Biomarker analysis
 - Found matches between sample peptides and biomarkers
 - ['NES', 'VIM', 'NRP1', 'SPN', 'RORA', 'S100A9', 'NFKB1', 'STAT1', 'C5AR1', 'CD44', 'GP1BB', 'KIT', 'STAT3', 'RUNX1', 'ALDOC', 'PECAM1', 'PTPRC', 'AQP4', 'PEA15', 'ITGA4', 'FOXO1', 'NDRG2', 'MSR1', 'VPS54']
- Predictions based on signatures
- Other datasets

