

Just - Evotec Biologics

Multi-Omic Characterization of CHO media and lysate during the production of biotherapeutics



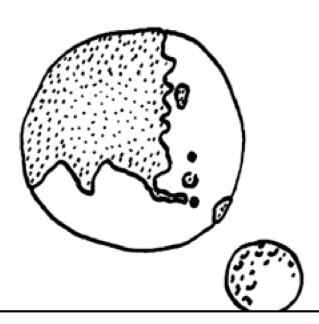
Agenda

Just - Evotec Biologics - Why Omics?

Inducible Cell line and Experimental Design

Results

Conclusion





Just - Evotec Biologics is a technology design company

Integrated data support to harness machine learning

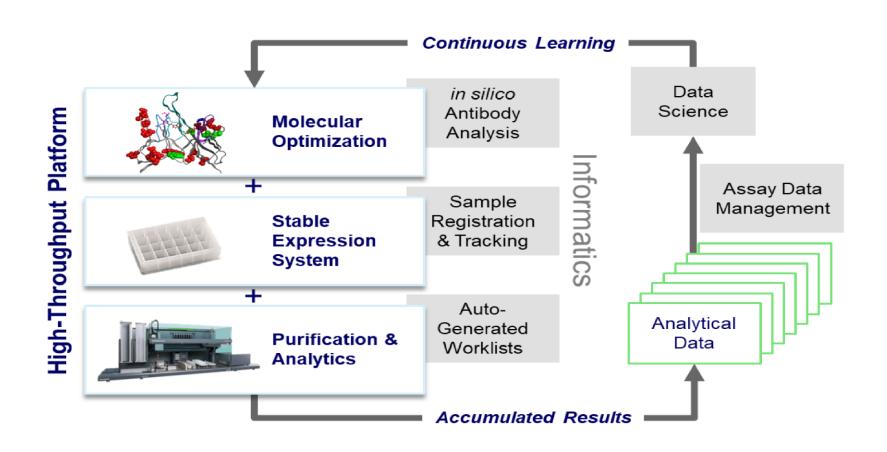
the Just. mission
design and apply
innovative technologies
to dramatically expand
global access
to biotherapeutics

Superior Sequence Molecule optimized for **MOLECULE DISCOVERY J.DISCOVERY Discovery** developability **MOLECULE DESIGN** J.MD[™] **J.DESIGN** MANUFACTURING DESIGN J.POD° Cell line, **PROCESS & PRODUCT DESIGN** Clinical and process and JP3° formulation Commercial Manufacturing development



Platform optimization requires large data input

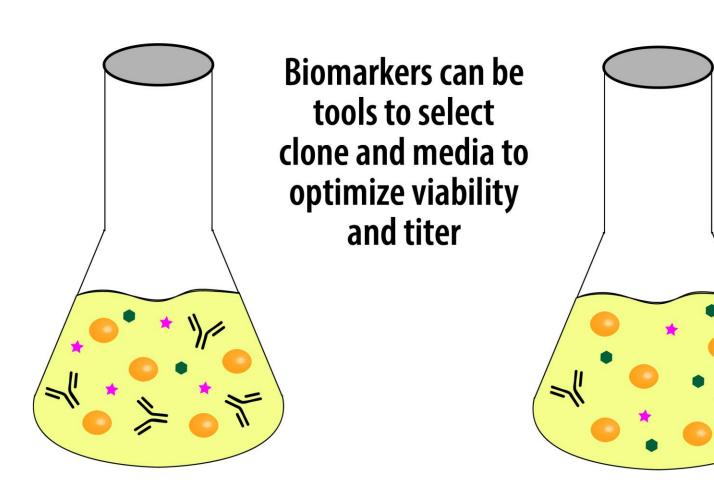
Analytical data drives platform improvement





Cell metabolism is perturbed by high expression of biologics

Biomarkers of high expression allow for optimization across platform





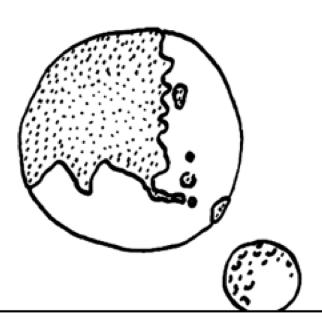
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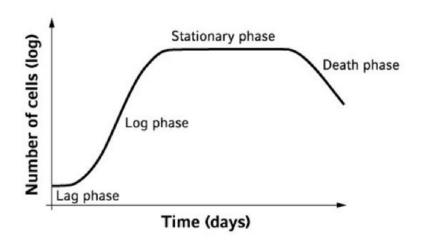
Conclusion



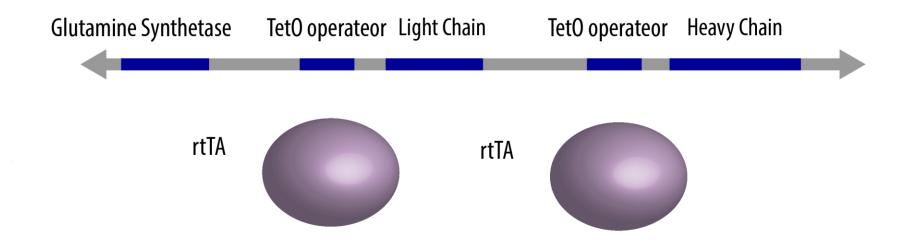


Inducible Cell Line allows seperation of growth and production phases

6F5 Inducible Cell Line



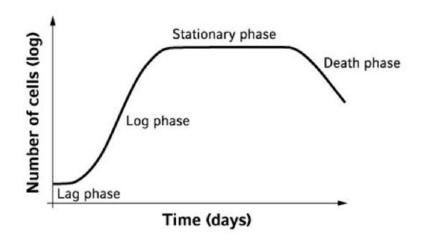
Selecting pools without addition of doxycycline allows cells to recover more effectively without needing to produce protein



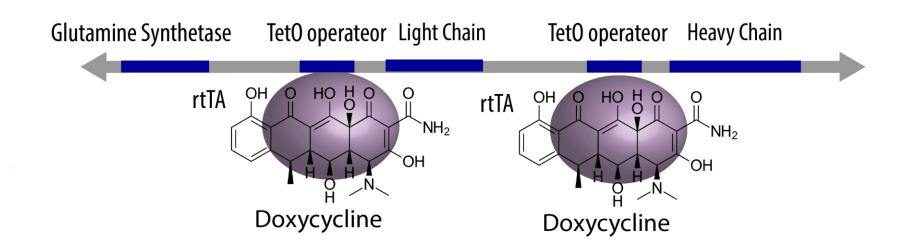


Inducible Cell Line allows seperation of growth and production phases

6F5 Inducible Cell Line



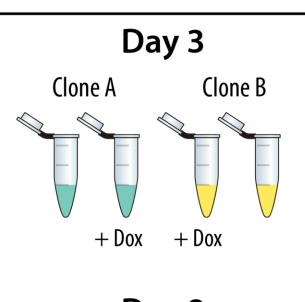
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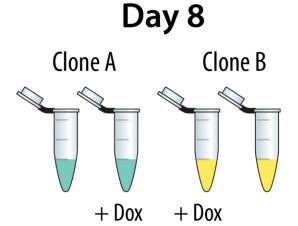


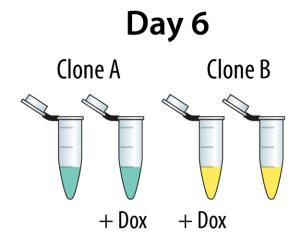


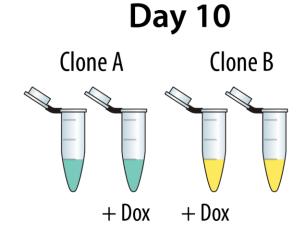
Experimental Design

16 Media and cell pellet samples collected for proteomic and metabolomic analysis





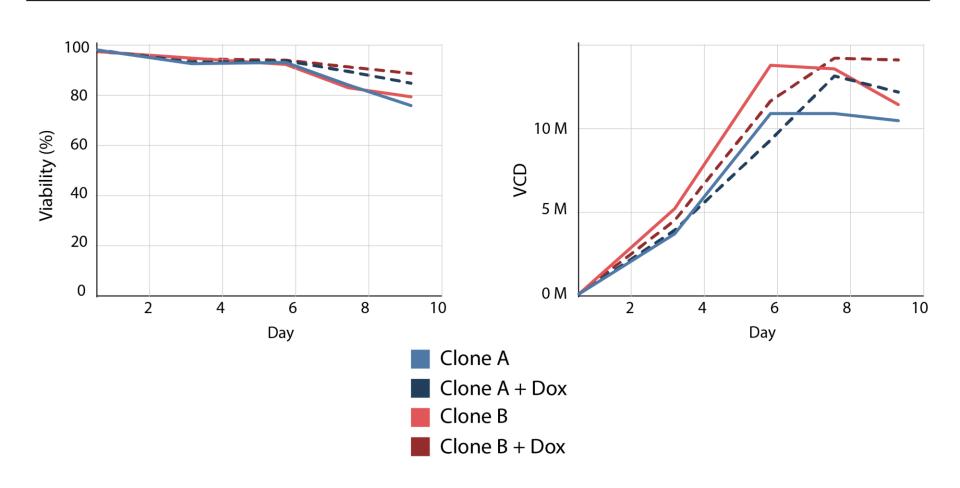






Cell culture performance

Less viability and slightly faster growth without doxycyline

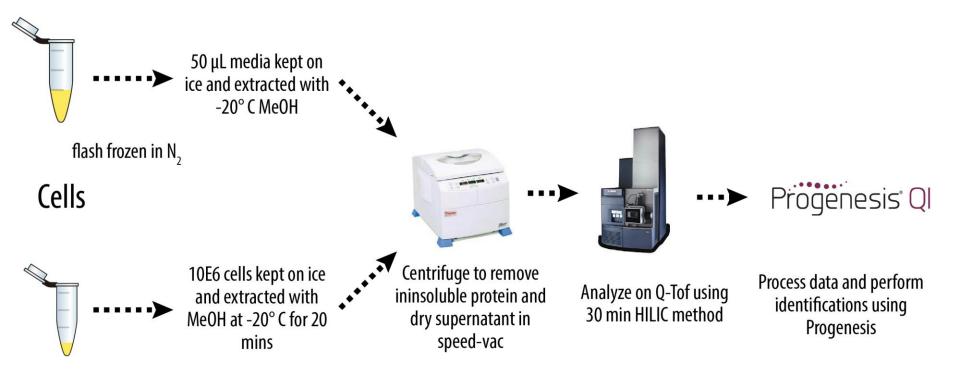




Metabolomics Sample Prep

Media and Cell Pellets

Media

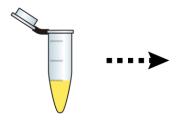


washed in PBS, flash frozen in N₂



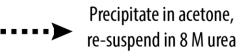
Proteomics Sample Prep

Media



flash frozen in N₂

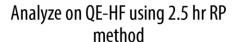
Automated ProA purification of all samples, flow through collected for analysis





BCA of all samples,







De-salt with HLB cartridges and dry in speed vac



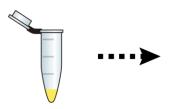
Reduce, alkylate, dilute to 50 mM Tris and digest overnight with trypsin



Proteomics Sample Prep

Cell Pellets

Cells



1E6 cells re-suspended in 8 M Urea/7 M GnHCl, DNA dispersed using probe sonicator, centrifuge and transfer supernatant

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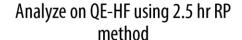
BCA of all samples, aliquot 100 µg

Reduce and alkylate samples, precipitate protein in cold EtOH

Cells washed in PBS, flash frozen in N₂









De-salt with HLB cartridges and dry in speed vac



Re-suspend dried cell pellets in 8 M Urea, dilute with 50 mM Tris, digest with trypsin overnight



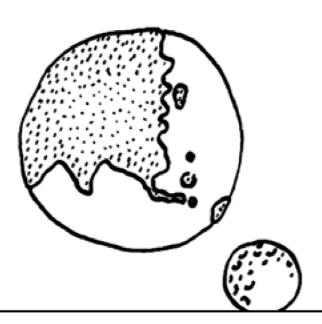
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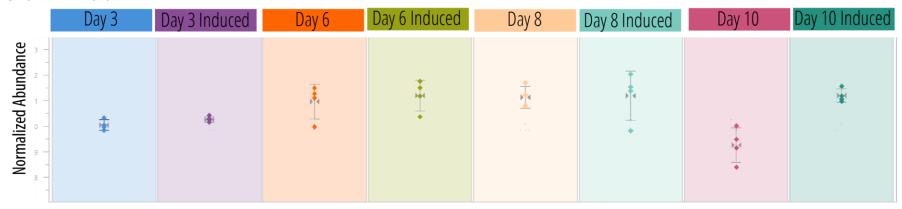


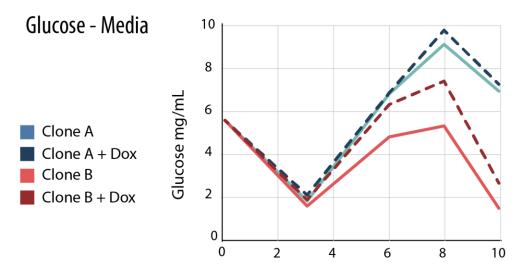


Glucose

Doxycycline appears to decrease overall consumption of glucose

Glucose - Media





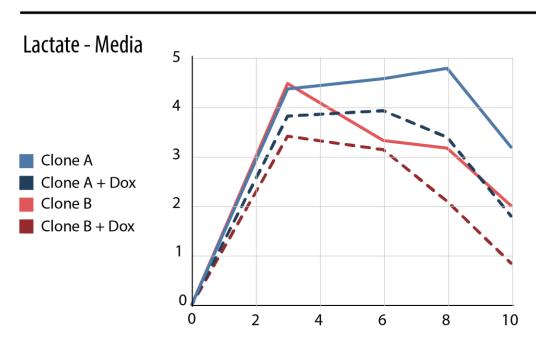
Day

- Cells treated with doxycycline typically grow slower consuming less glucose and take longer to reach carrying capacity
- Hypothesize that this is due to increased energy being put toward antibody production



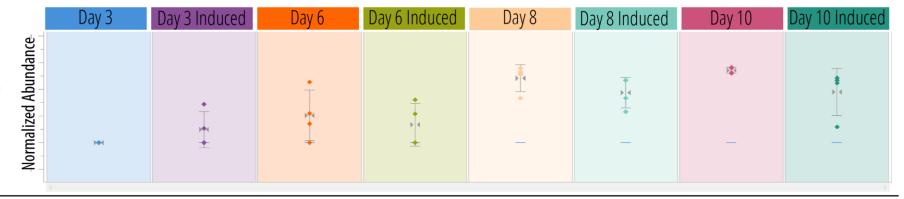
Lactate

Shift to less lactate production, more lactate consumption



- Lower lactate could also maybe explain growth/viability differences
- This shift could also be explained by differences in GS expression, possibly through a push in the TCA-aKG-Glutamate cycle

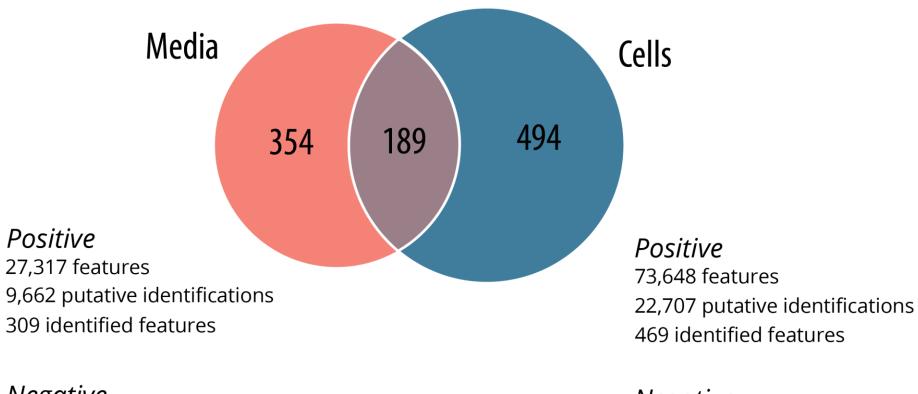
Lactate - Cells





Over 200 metabolites identified from each sample set

Metabolomics results



Negative

13,144 features3,658 putative identifications279 identified features

Negative

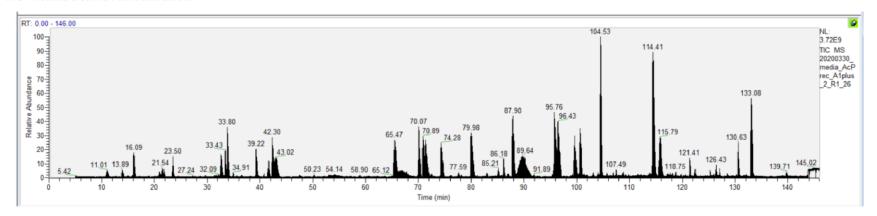
9,096 features2,730 putative identifications343 identified features



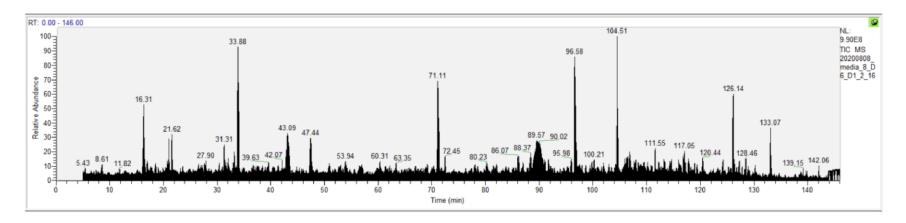
Antibody removal with ProA improved media analysis

More low abundant proteins are visible in flow through

TIC - Media before ProA filtration



TIC - Media after ProA filtration

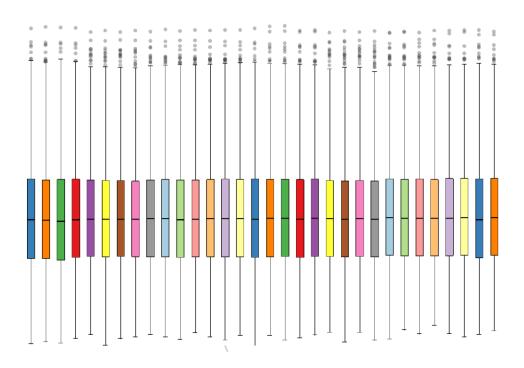




Over 3,500 proteins identified in cells and 1,500 from media

Proteomics results

All samples were normalized to equivalent abundances



3,953 proteins identified with FDR < 1%

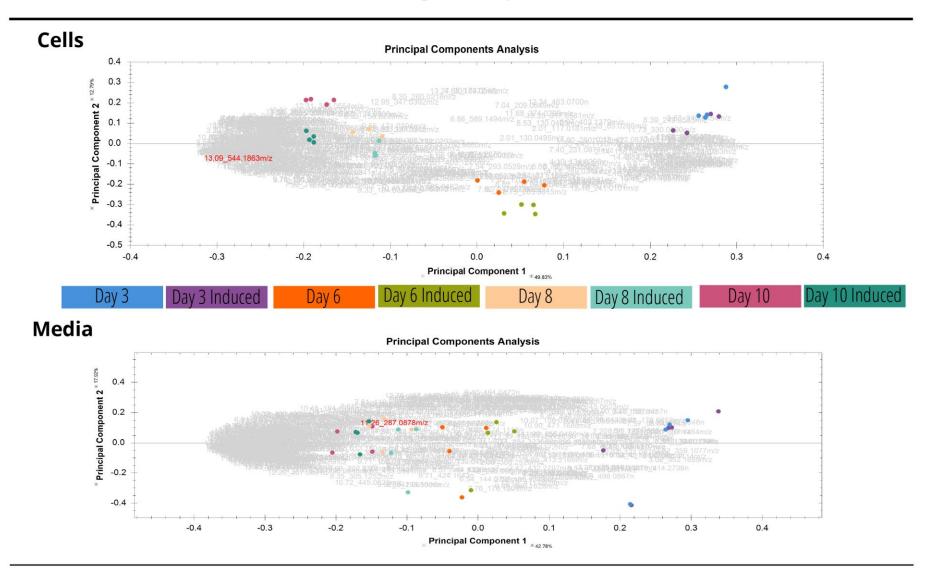
1,679 proteins identified with FDR < 1%

Raw data searched against database of mouse proteins using Proteome Discoverer



Cells and media show seperation by time in PCA

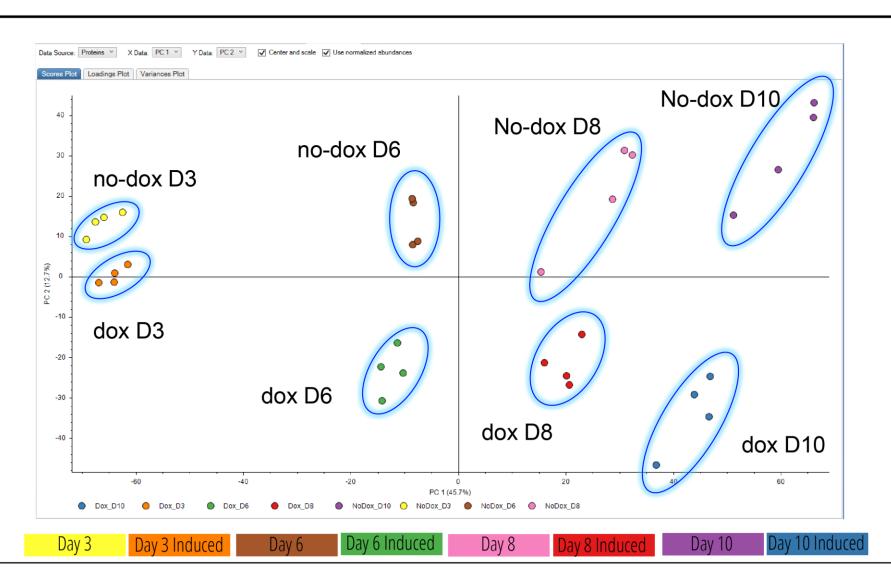
Cells show better separation by condition in metabolomics





Distinctive grouping by category seen in PCA analysis

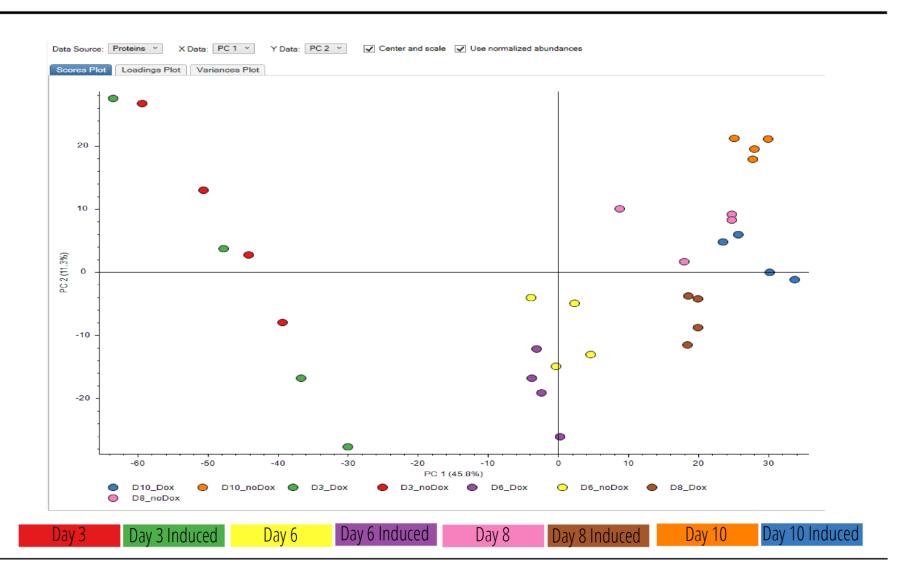
Proteomics Results - Cells





Media shows seperation by time in PCA analysis

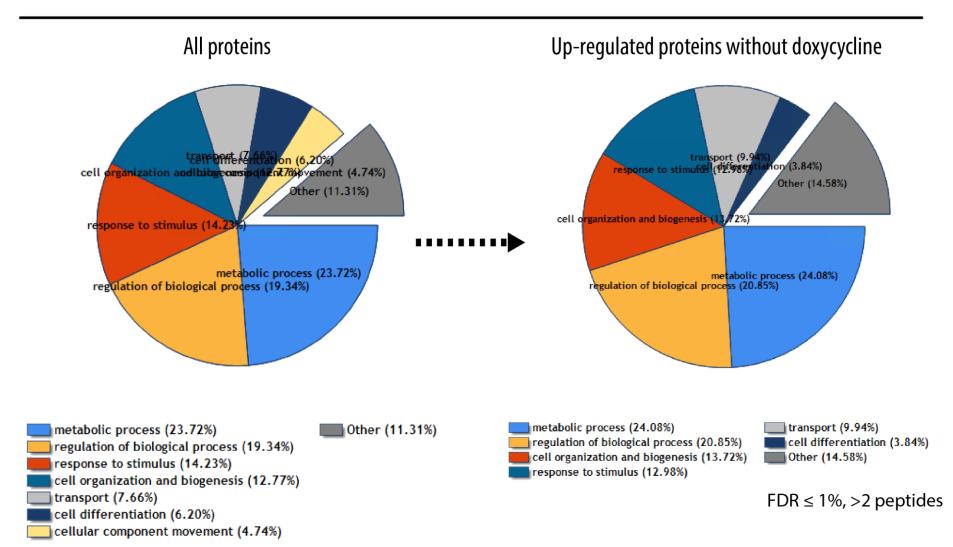
Proteomics Results - Meida





Fewer proteins associated with transport, more with differentiation in untreated cells

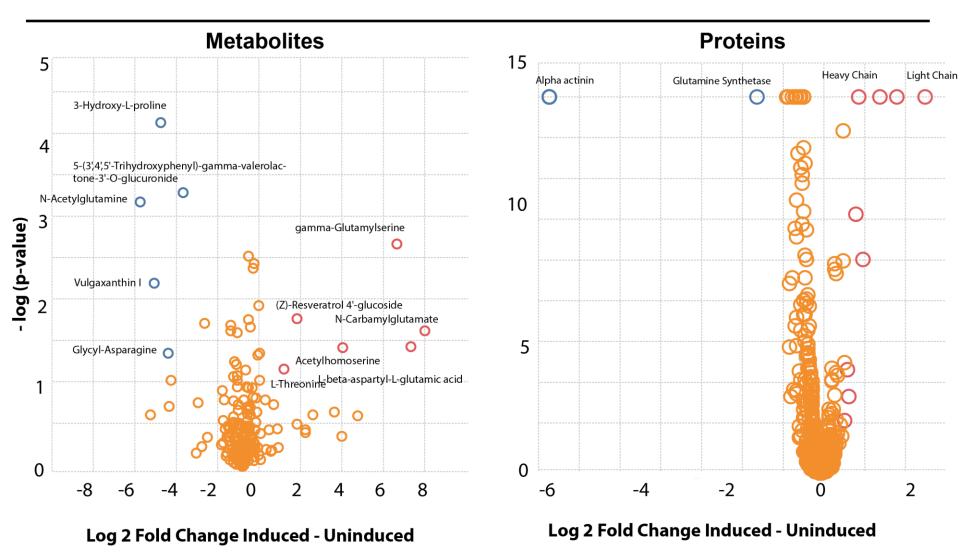
GO enrighment of KEGG pathways





Induced and uninduced samples similar at day 3

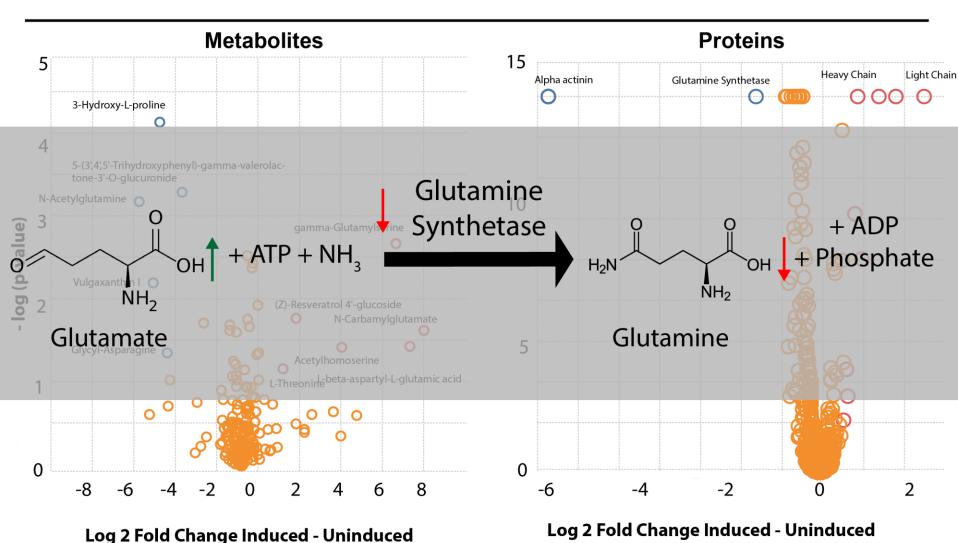
Metabolomics and Proteomics results: Day 3 Cells





Induced and uninduced samples similar at day 3

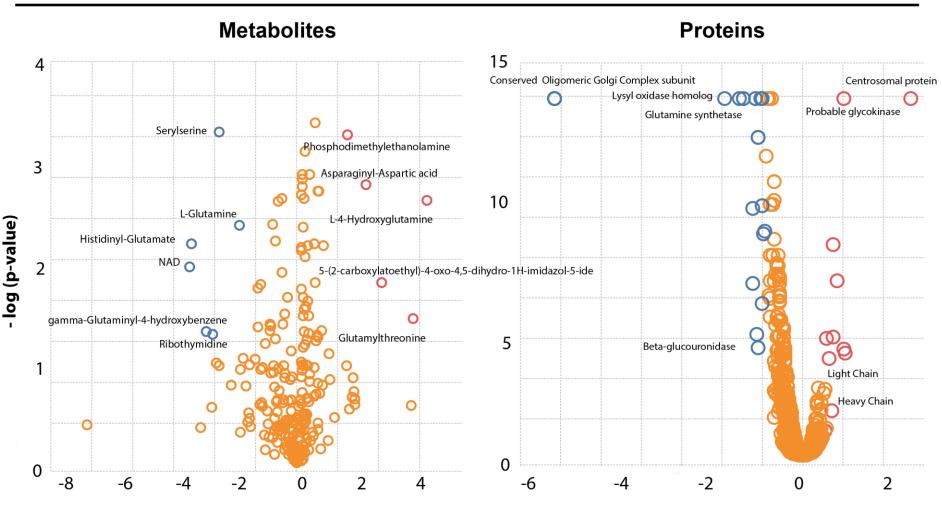
Metabolomics and Proteomics results: Day 3 Cells





Decreases seen in Glutamine and NAD

Metabolomics and Proteomics results: Day 6 Cells



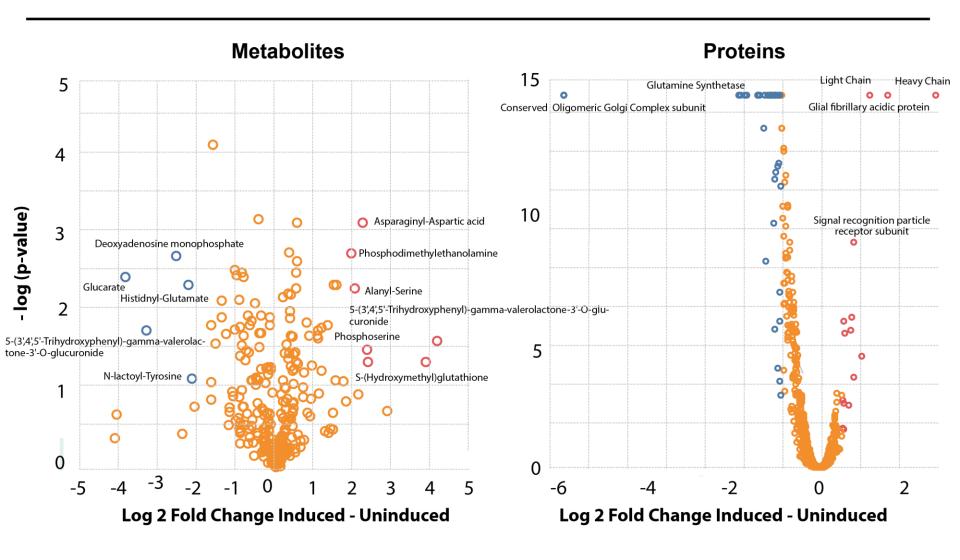
Log 2 Fold Change Induced - Uninduced

Log 2 Fold Change Induced - Uninduced



Increasing number of proteins down-regulated

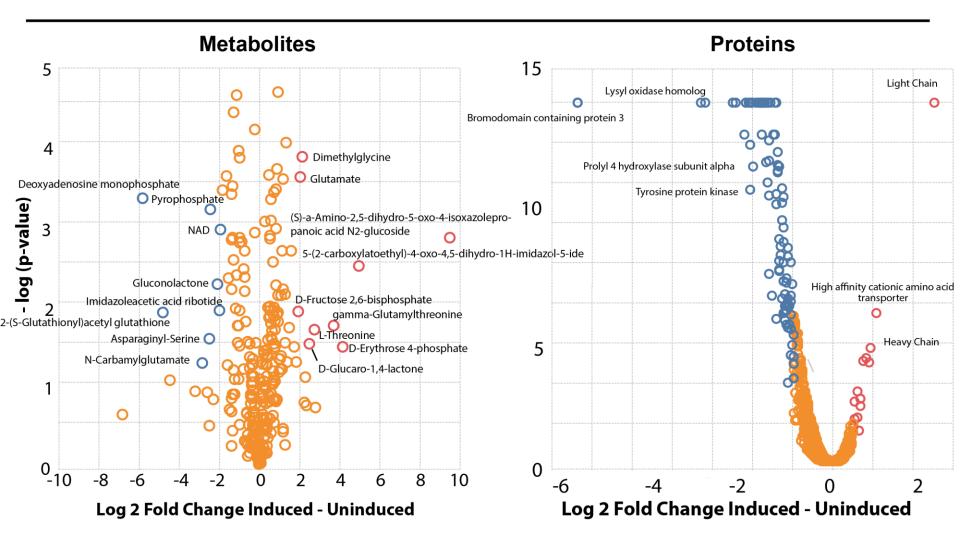
Metabolomics and Proteomics results: Day 8 Cells





Large number of proteins down-regulated by day 10

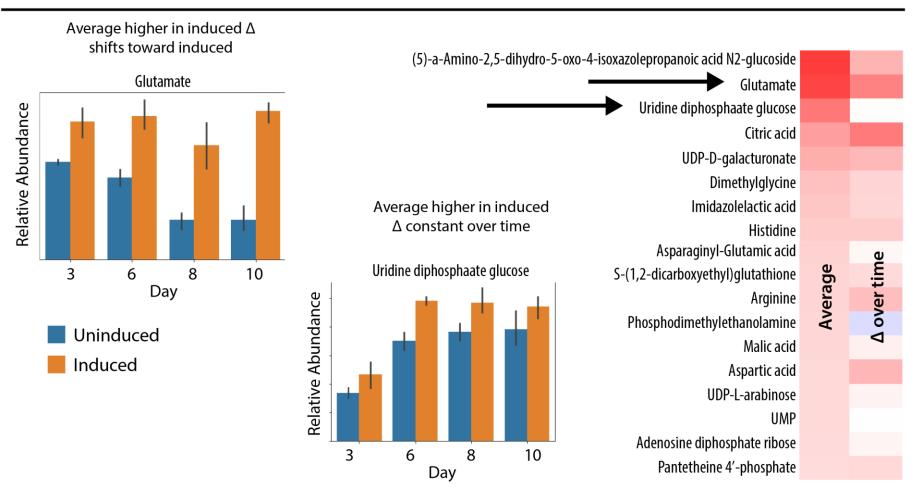
Metabolomics and Proteomics results: Day 10 Cells





Metabolomics - Cells: Effect of Induction

Species increased by induction



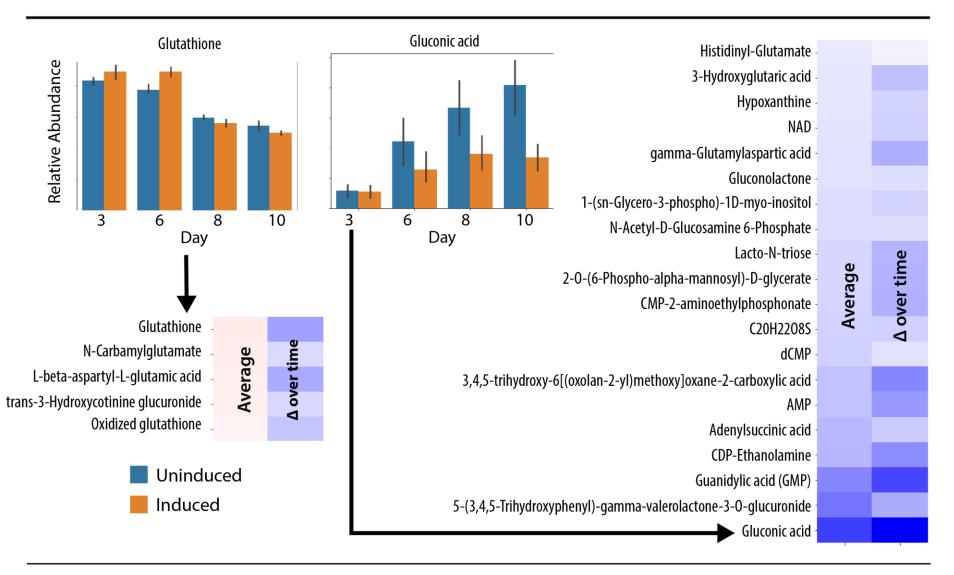
ASCA - ANOVA Stimultaneous Component Analysis

Smilde et al, "ANOVA-simultaneous component analysis (ASCA): a new tool for analyzing designed metabolomics data", Bioinformatics, 2005. Zwanenburg et al., "ANOVA-principal component analysis and ANOVA-simultaneous component analysis: a comparison". J. Chemometrics, 2011



Metabolomics - Cells: Effect of Induction

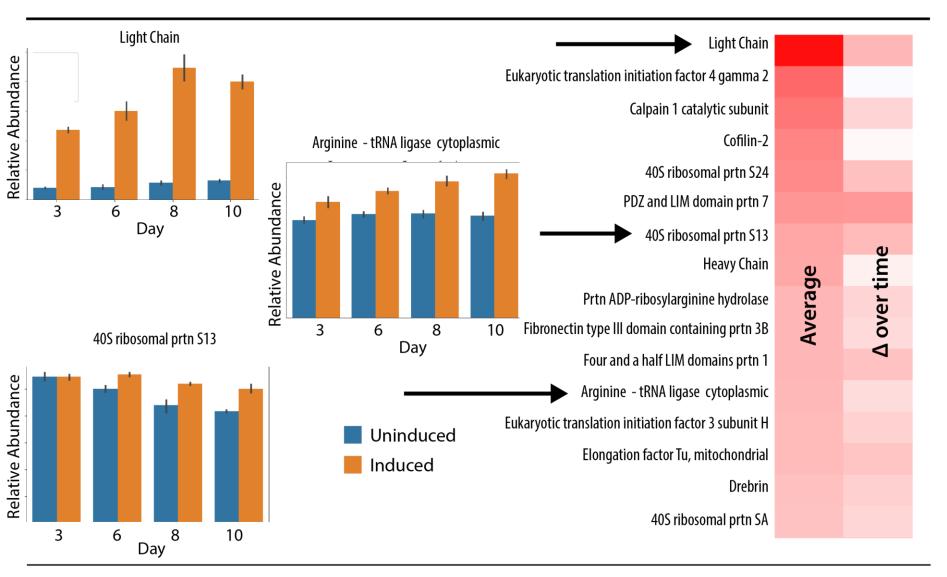
Species decreased by induction





Proteins - Cells: Effect of Induction

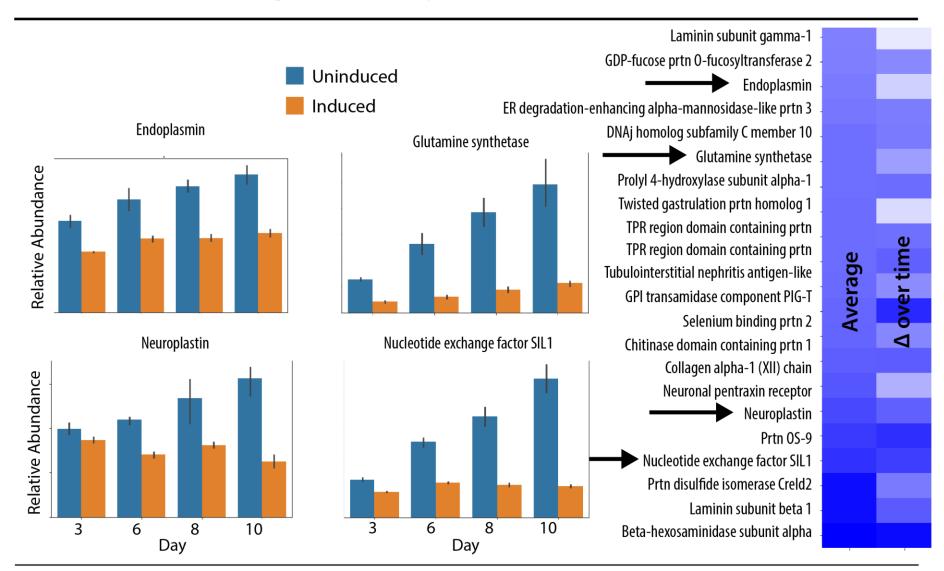
Species increased by induction





Proteins - Cells: Effect of Induction

Species decreased by induction





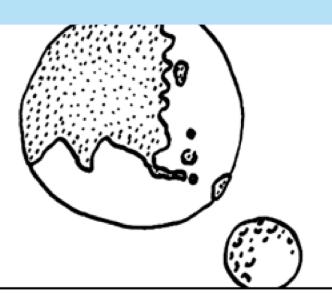
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Summary

- Omics can provide a snapshot of cell physiology at different conditions
- An inducible cell line exhibits slightly slower growth and decreased glucose consumption while producing protein
- Over 3,900 protein and 650 metabolites were identified in cells
- Over 1,679 proteins and 500 metabolites were identified in media
- Significant shifts are seen in protein expression and metabolism are seen during protein production
- Glutamine synthetase downregulated as part of induction system, corresponding metabolites affected
- Ribosomes up-regulated in induced state, predicted to be due to high demand for protein production



Acknowledgments

Just-Evotec Biologics

Yuko Ogata Gabe Stancu Jeremy Shaver Jeff Meyer