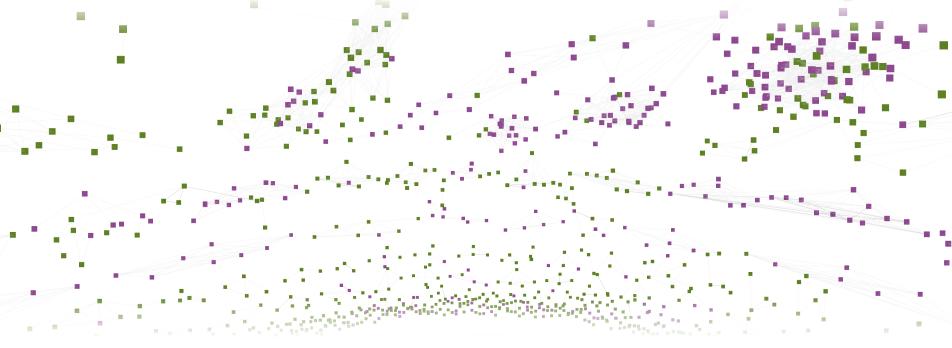
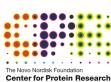
Network biology





Lars Juhl Jensen jensenlab.org



what will you learn?

STRING database

network creation

enrichment analysis

FAVA

co-expression networks

Cytoscape tool

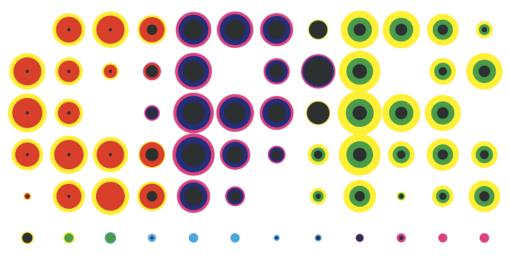
STRING integration

network visualization

who are we?

Lars Juhl Jensen

group leader 2009–



The Novo Nordisk Foundation

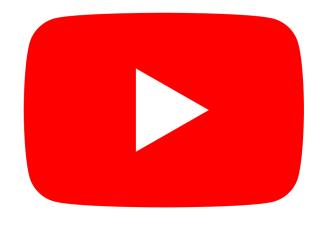
Center for Protein Research

cofounder 2009–2022



intomics

YouTuber 2020–



youtube.com/larsjuhljensen

Mikaela Koutrouli

PhD student

single-cell transcriptomics

Nadezhda Doncheva

assistant professor

omics data analysis

network analysis

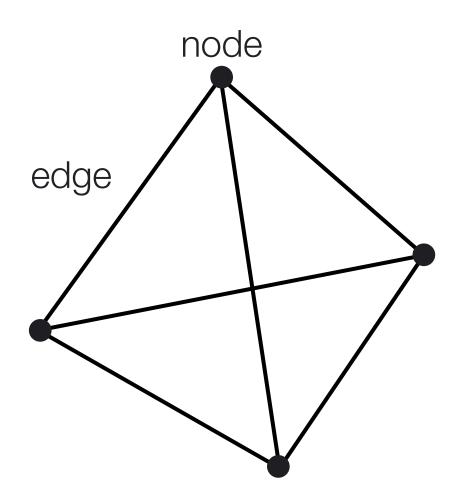
why networks?

entities or concepts

understand their interplay

useful abstraction

lends itself to visualization



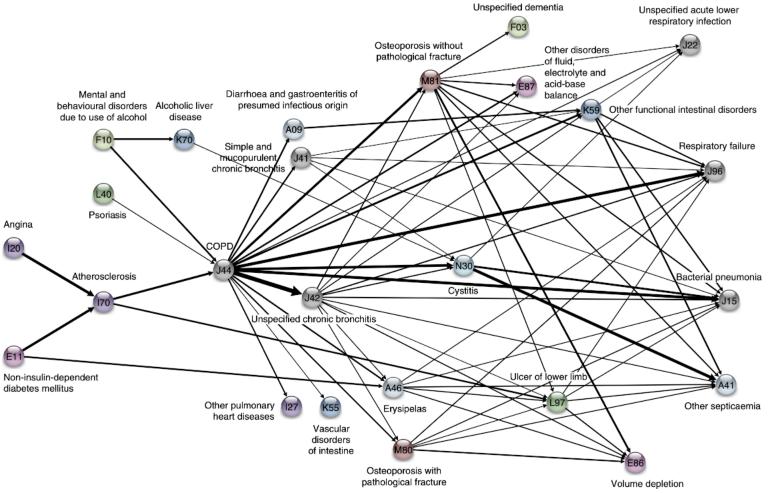
nodes

things to be linked

edges

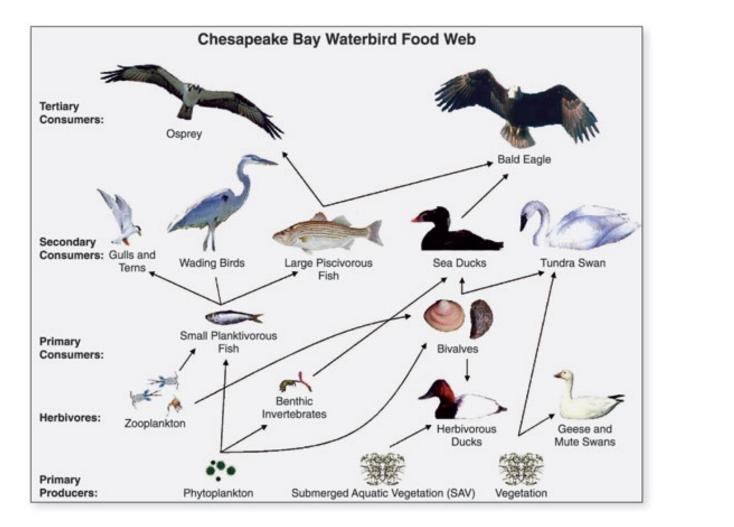
links between things

disease networks



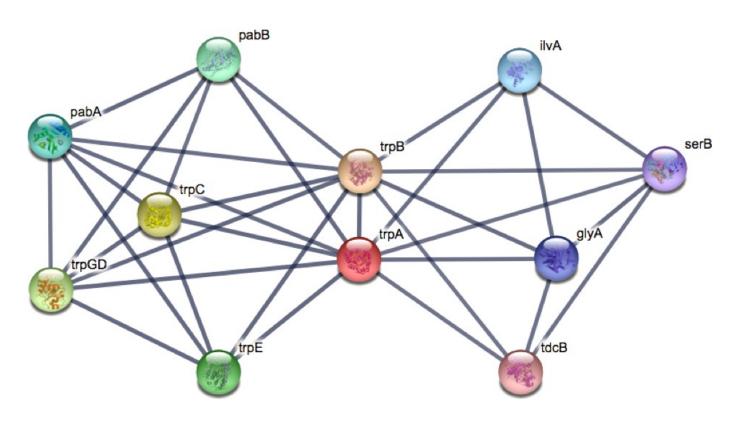
Jensen et al., Nature Communications, 2014

food webs



protein networks

The STRING database



14,094 genomes

67.6 million proteins

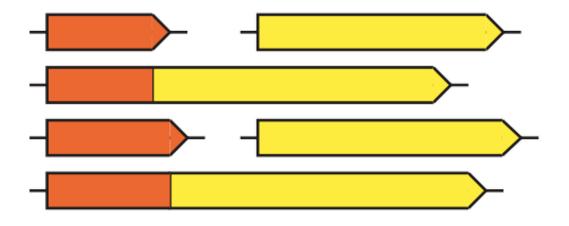
physical interactions

functional associations

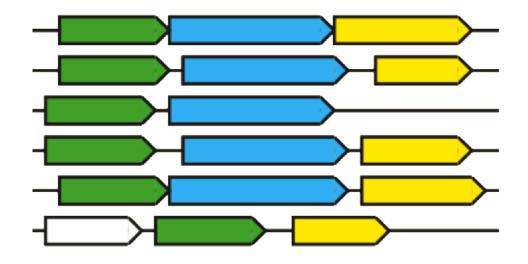
data integration

genomic context

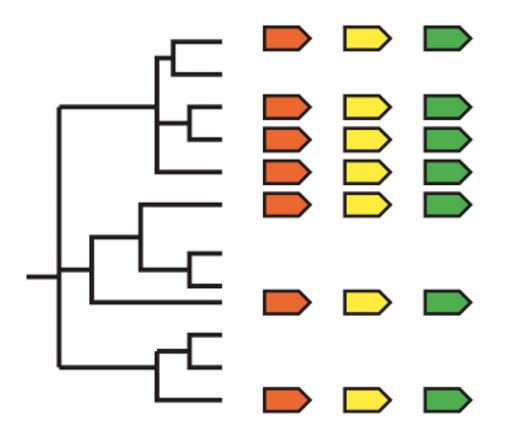
gene fusion



gene neighborhood



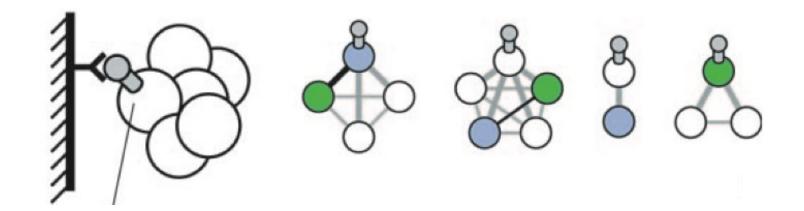
phylogenetic profiles



experimental data

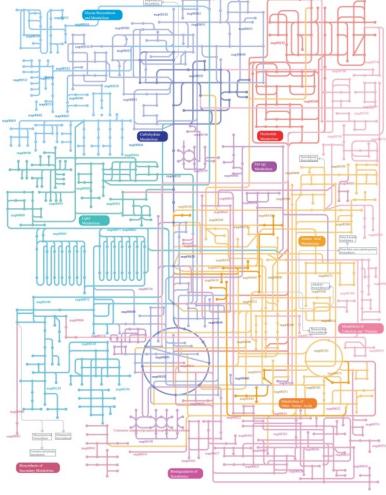
gene coexpression

interaction experiments



curated knowledge

molecular pathways



Kanehisa et al., Nucleic Acids Research, 2008

many databases

different formats

different names

varying quality

not comparable

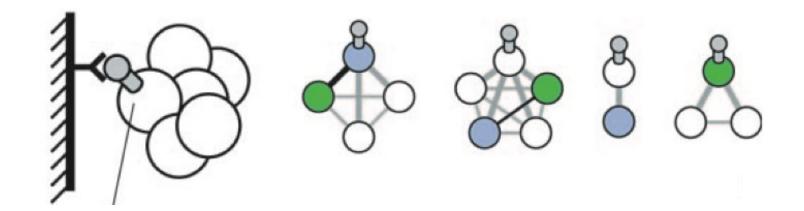
not same species

hard work

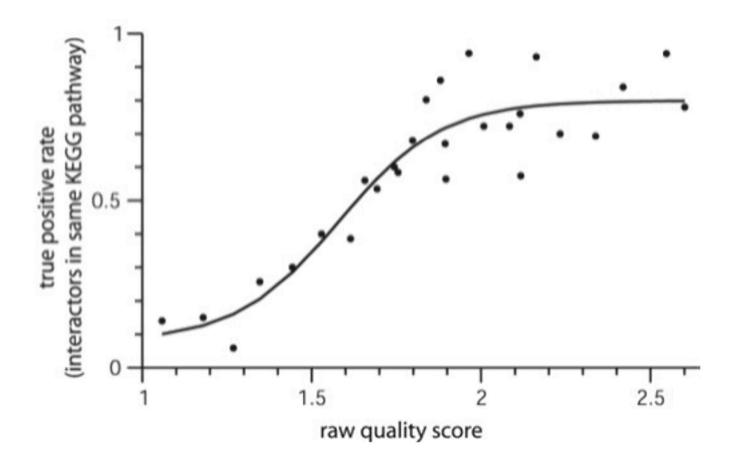
parsers

mapping files

scoring schemes



score calibration



von Mering et al., Nucleic Acids Res., 2005

transfer by orthology

missing most of the data

text mining



>10 km

too much to read

computer

named entity recognition

dictionary

cyclin dependent kinase 1

CDK1

block list

SDS

co-mentions

counting

distance-weighted score

consensus view

no relation type

deep learning

pre-trained transformers

huge unlabeled corpora

finetune for specific task

physical protein interactions

practical session



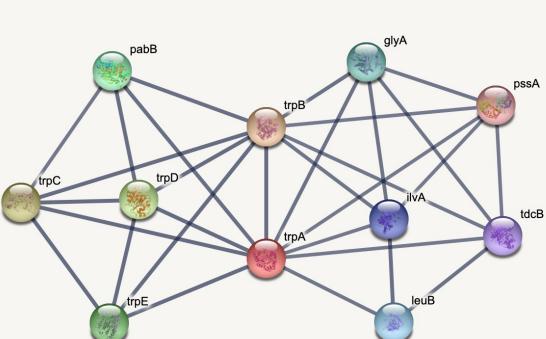
Download

LOGIN

Help

REGISTER

My Data



Search

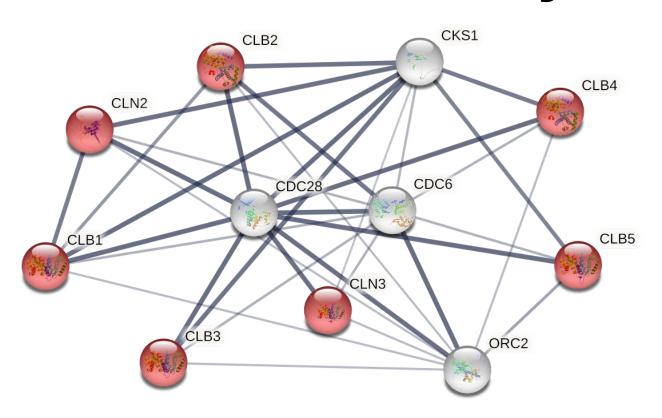
query the database

inspect the evidence

exercise 1

jensenlab.org/training/string/

Enrichment analysis



characterize a gene list

classes of genes

overrepresented

associated with the study

overrepresentation

omics study

regulated proteins

mitochondria enriched?

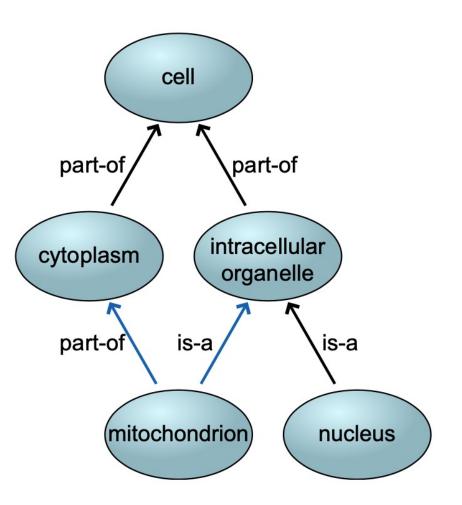
	Mitochondrial	Not mitochondrial	
Regulated	a	b	
Not regulated	С	d	

-

Fisher's exact test

p-value

Gene Ontology



systematically test all

many gene sets

Gene Ontology

pathways

protein domains

UniProt keywords

subcellular localizations

tissues

associate diseases

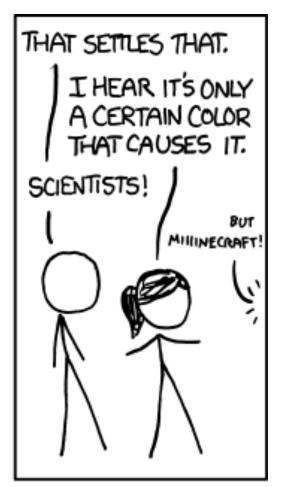
local network clusters

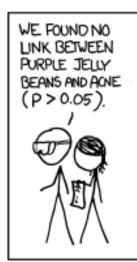
publications

multiple testing



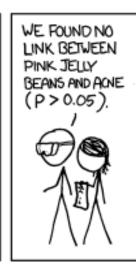




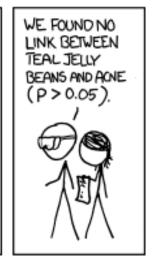


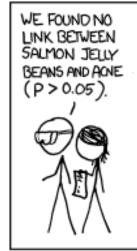


WE FOUND NO



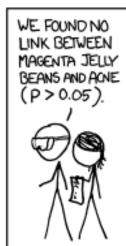


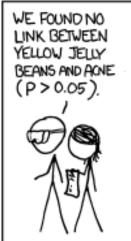




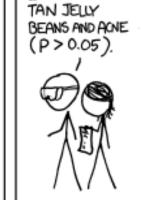






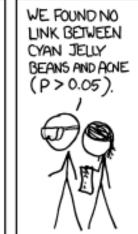




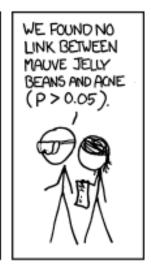


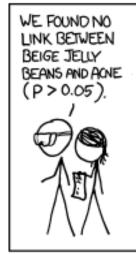
WE FOUND NO

LINK BETWEEN



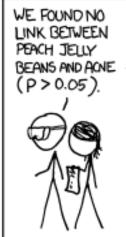


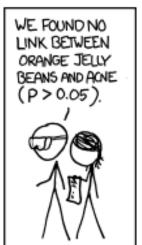


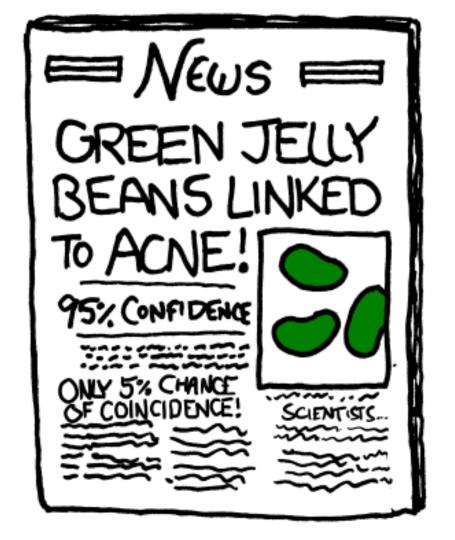












enrichment analysis

thousands of tests

crucial to correct

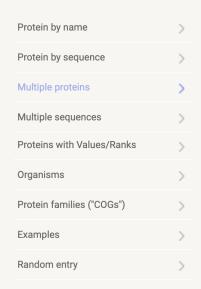
Bonferroni correction

false discovery rate

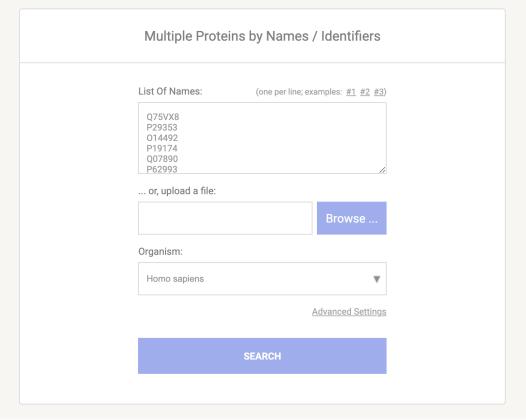
practical session

proteomics study





SEARCH



protein network

change query parameters

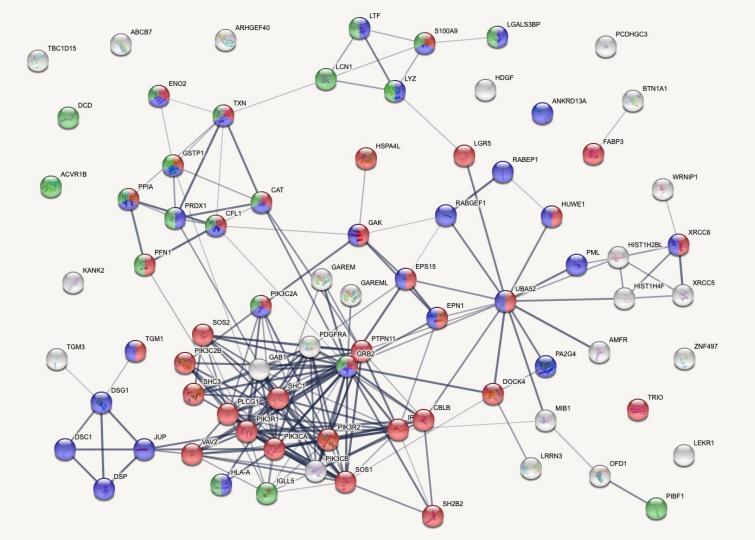
functional enrichment

>	KEGG Pathways			
pathway	description	count in network	strength	false discovery rate
hsa01521	EGFR tyrosine kinase inhibitor resistance	12 of 78	1.59	6.17e-13
hsa04012	ErbB signaling pathway	12 of 83	1.56	6.17e-13
hsa04722	Neurotrophin signaling pathway	13 of 114	1.46	6.17e-13
hsa04650	Natural killer cell mediated cytotoxicity	13 of 121	1.43	6.57e-13
hsa05214	Glioma	11 of 72	1.58	2.17e-12
				(more)
>	Reactome Pathways			
pathway	description	count in network	strength	false discovery rate
HSA-8853659	RET signaling	12 of 40	1.88	2.00e-14
HSA-168249	Innate Immune System	29 of 1025	0.85	1 44e-13

>	Reactome Pathways			
pathway	description	count in network	strength	false discovery rate
HSA-8853659	RET signaling	12 of 40	1.88	2.00e-14
HSA-168249	Innate Immune System	29 of 1025	0.85	1.44e-13
HSA-177929	Signaling by EGFR	11 of 50	1.74	3.83e-12
HSA-9006335	Signaling by Erythropoietin	9 of 25	1.96	3.20e-11
HSA-2424491	DAP12 signaling	9 of 28	1.91	6.02e-11
				(more)

WikiPathways ≜ false discovery rate pathway description count in network strength EGF/EGFR signaling pathway 15 of 162 1.37 4.14e-13 WP437 Insulin signaling 15 of 159 1.37 4.14e-13 WP481 EGFR tyrosine kinase inhibitor resistance WP4806 12 of 83 1.56 1.56e-12 WP673 ErbB signaling pathway 12 of 90 1.52 2.82e-12 WP2037 Prolactin signaling pathway 11 of 76 1.56 1.43e-11 (more ...)

show on network



exercise 2

jensenlab.org/training/string/