

# *Network Biology and Evolution of Human Genetic Diseases*



Sanguk Kim  
POSTECH

# Outline

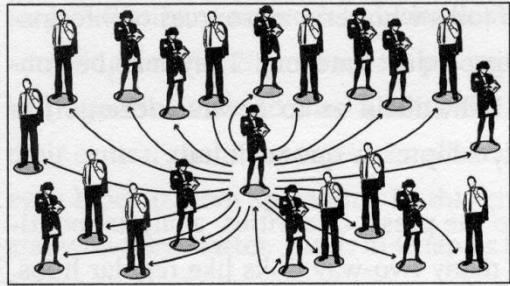
- *Introduction of Network Biology and Medicine*

- *Network Distance & Localization* → *Disease Comorbidity* *Nature Mol Sys Biol.* 2011  
*Human disease evolution* *Nature Scientific Reports* 2012  
*Mitochondrial protein network* *Nature Scientific Reports* 2013

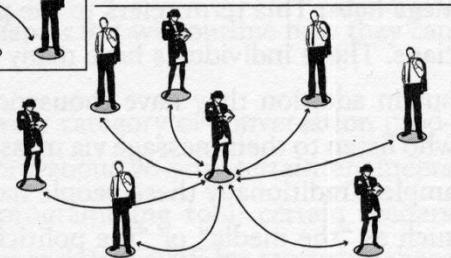
- *Network Clustering* → *Cancer* *PLoS Comp. Biol.* 2011

- *Network Rewiring and Evolution* → *Gene essentiality changes* *Nature Scientific Reports* 2012  
*Neuronal Disease* *PLoS Genetics* 2012

1. Spatial and functional organization of mitochondrial protein network. *Nature Scientific Reports* 2013 3:1403.
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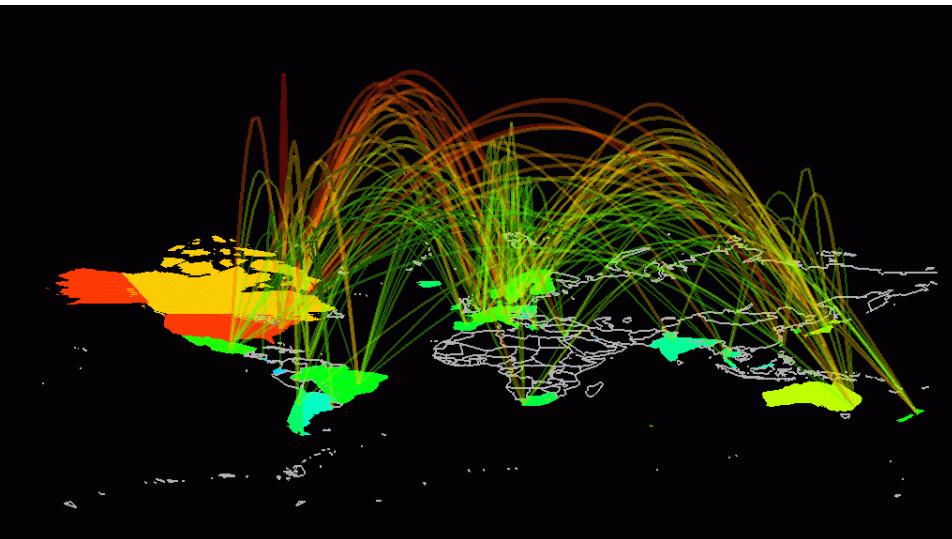
**Mega-Hub.** An MTV veejay spreads the word to thousands or millions of people through one-way links.



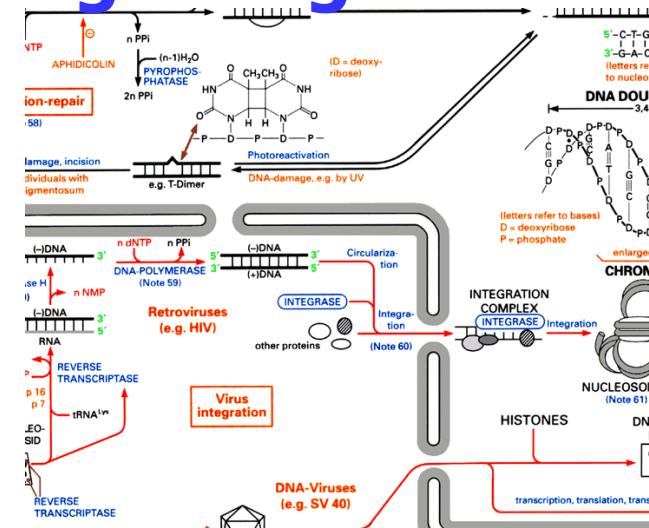
**Hub.** This undergraduate has spread the word to seven other people through two-way links.

# Society

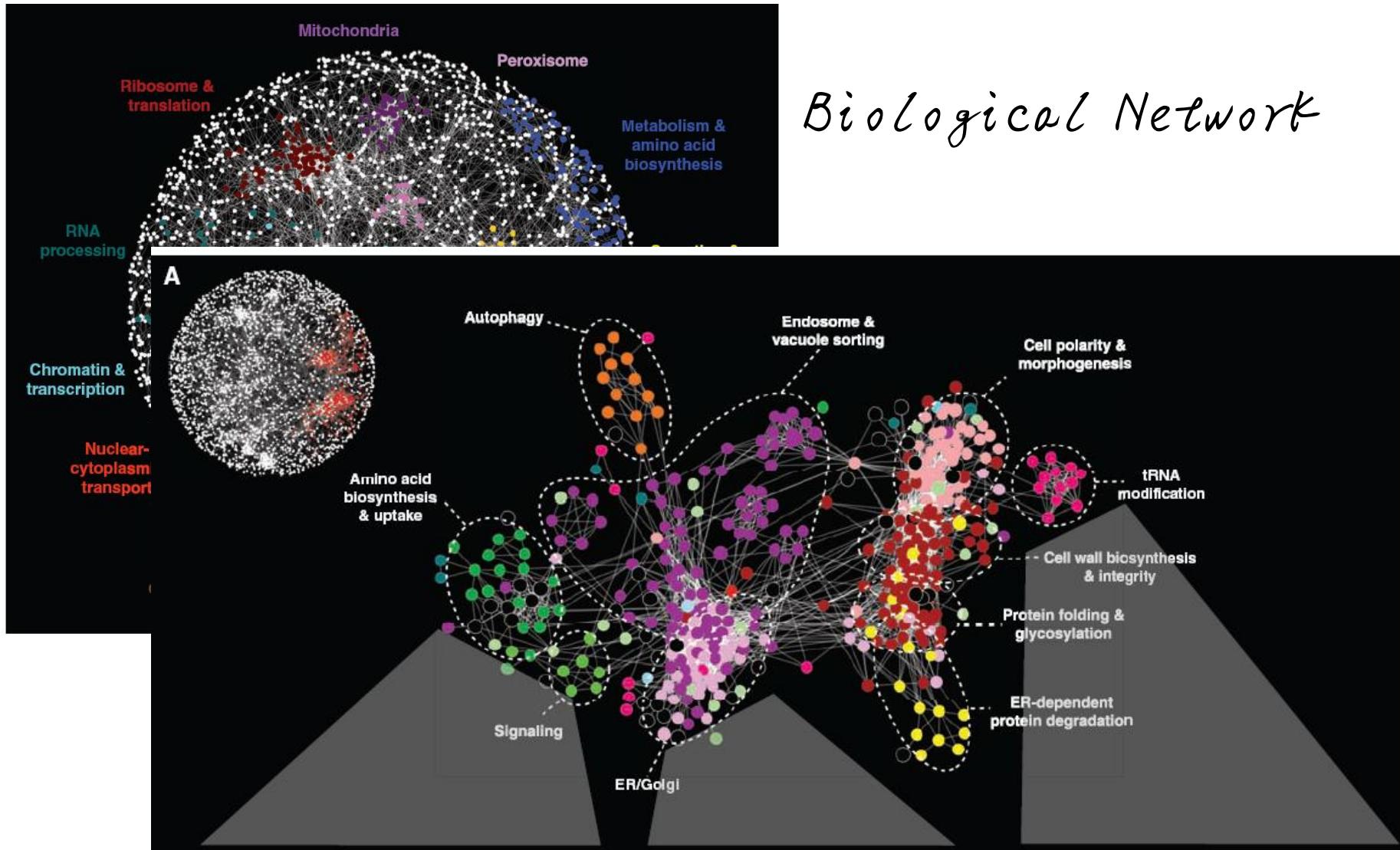
# Internet



# Biological signaling network



# Biological Network

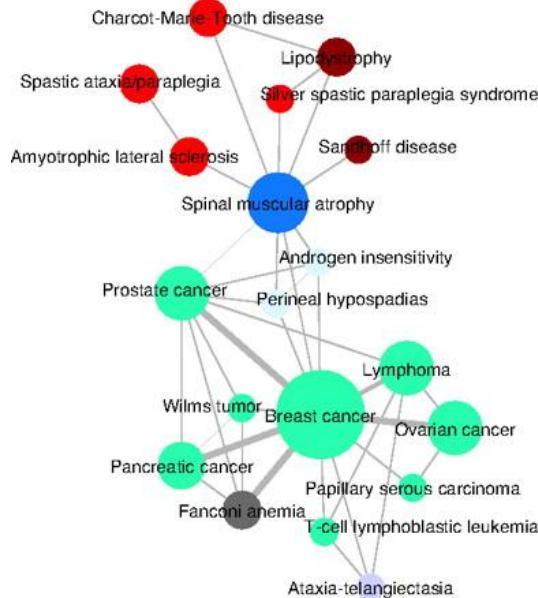


Disease pleiotropy and network modularity

# The human disease network

Construction of the diseaseome bipartite

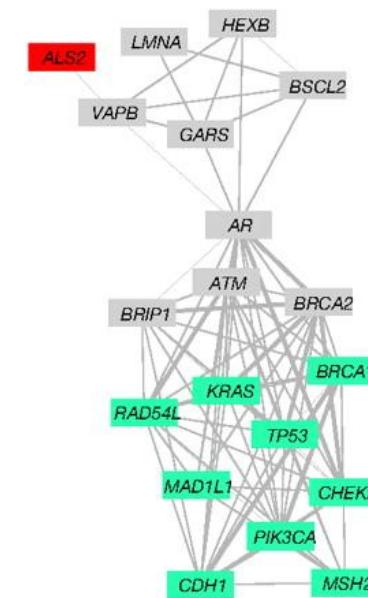
## Human Disease Network (HDN)



**disease phenotype**      **disease genome**

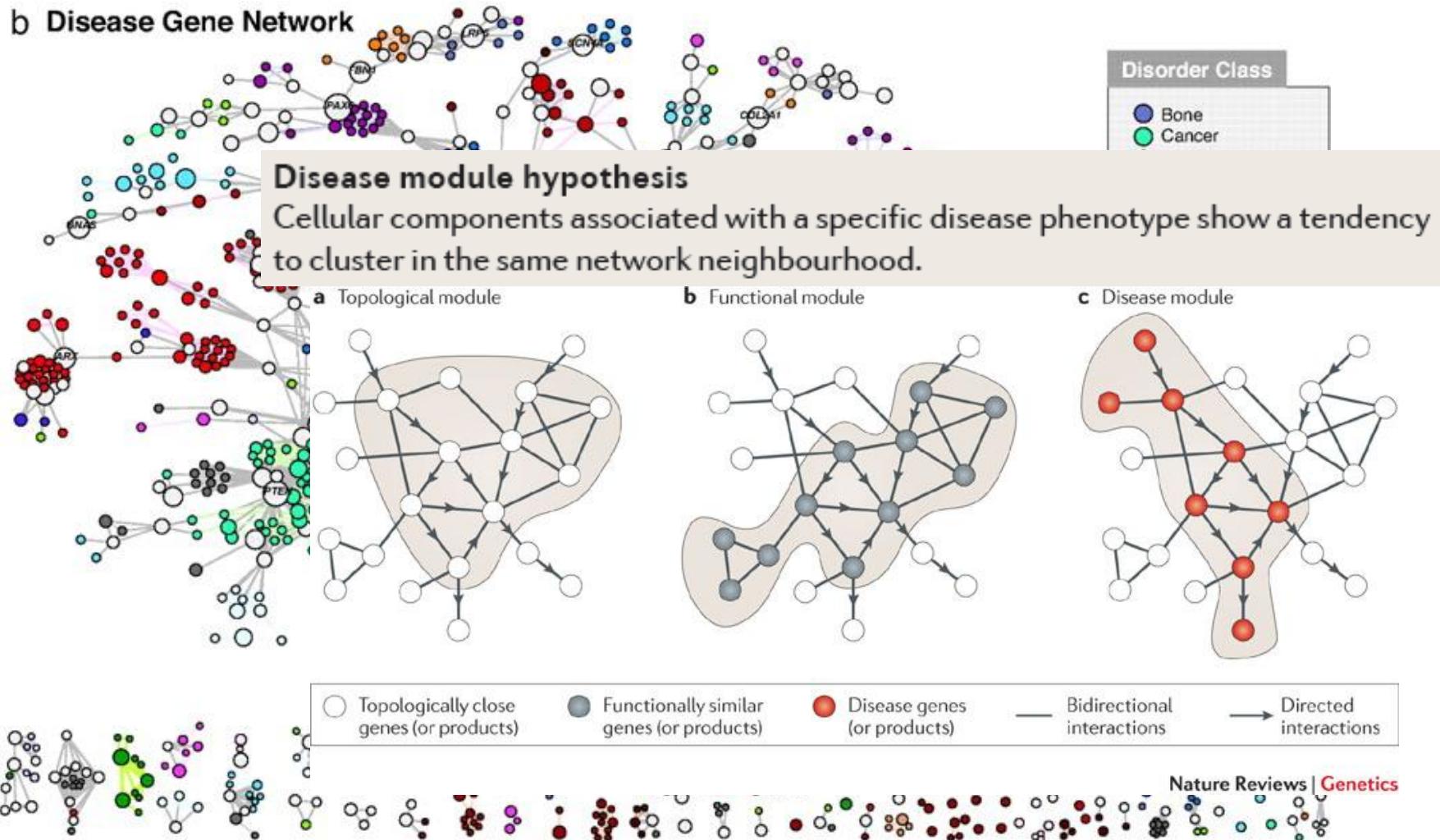
Disease Phenotype	Genes
Ataxia-telangiectasia	ATM
Perineal hypospadias	ATM
Androgen insensitivity	ATM
T-cell lymphoblastic leukemia	BRCA1
Papillary serous carcinoma	BRCA2
Prostate cancer	CDH1
Ovarian cancer	GARS
Lymphoma	HEXB
Breast cancer	KRAS
Pancreatic cancer	LMNA
Wilms tumor	MSH2
Spinal muscular atrophy	PIK3CA
Sandhoff disease	TP53
Lipodystrophy	MAD1L1
Charcot-Marie-Tooth disease	RAD54L
Amyotrophic lateral sclerosis	VAPB
Silver spastic paraplegia syndrome	CHEK2
Spastic ataxia paraplegia	BSCL2
Fanconi anemia	ALS2
	BRI1P1

## Disease Gene Network (DGN)

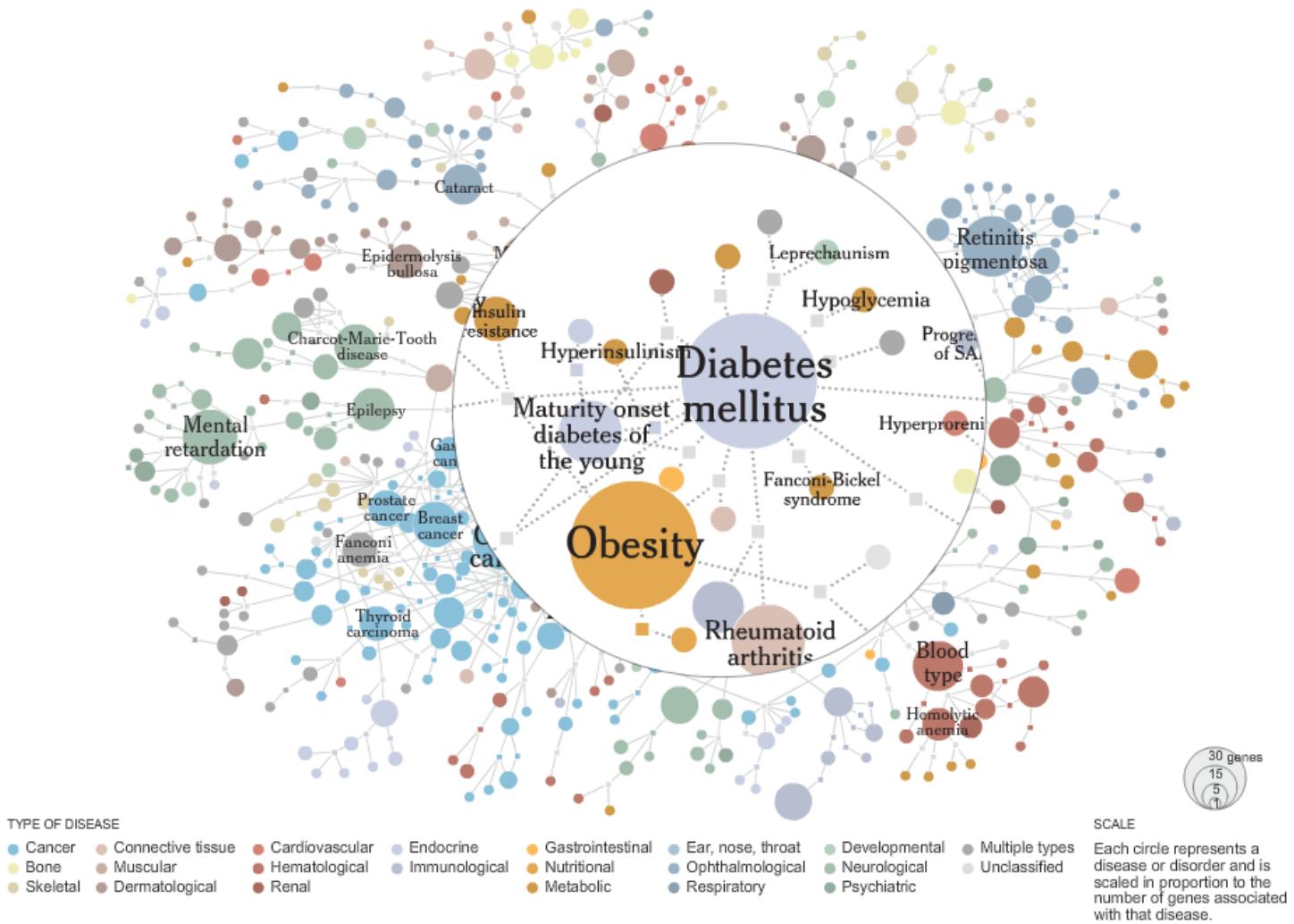


# Disease Gene Network

## b Disease Gene Network



# Example: Diabetes in the Human disease network



# Outline

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*Human disease evolution* *Nature Scientific Reports 2012*  
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2013

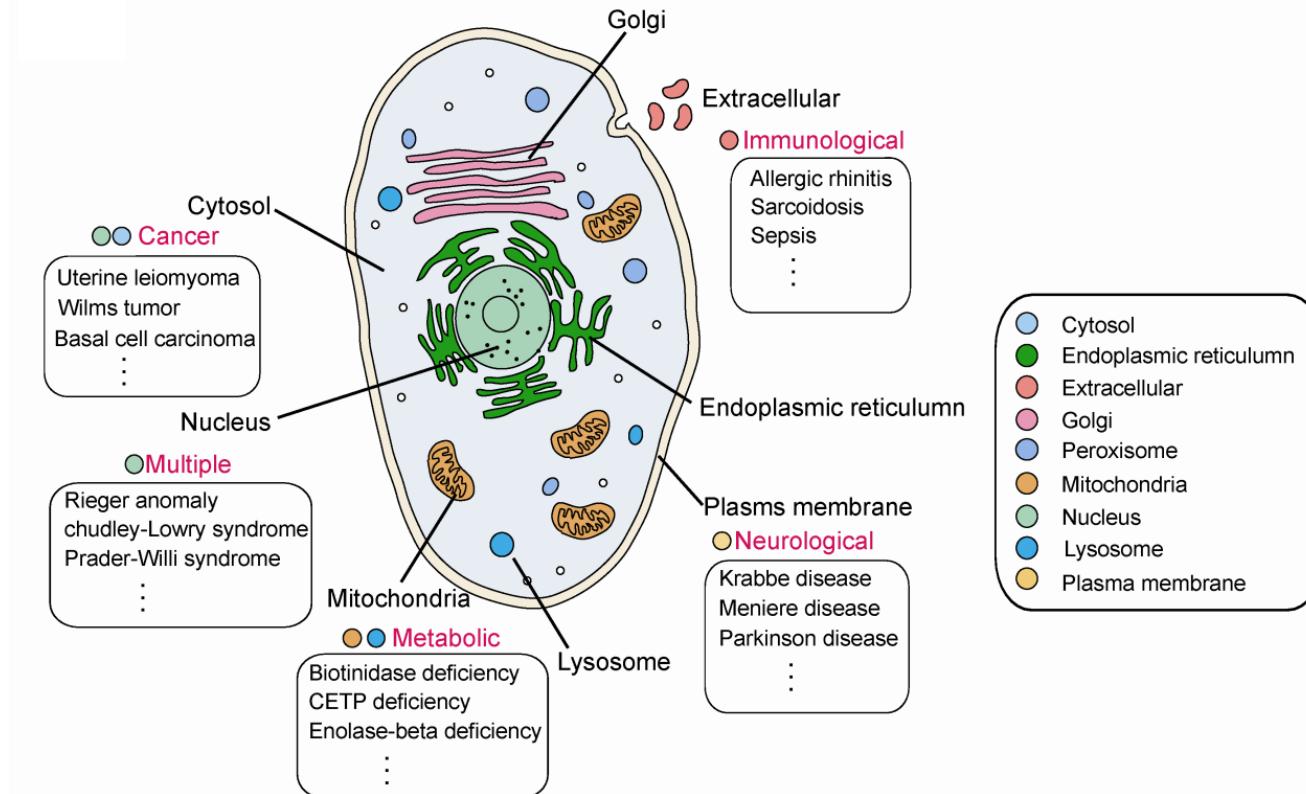
• *Network Clustering → Cancer* *PLoS Comp. Biol. 2011*

• *Network Rewiring and Evolution → Gene essentiality changes* *Nature Scientific Reports 2012*  
*Neuronal Disease* *PLoS Genetics 2012*

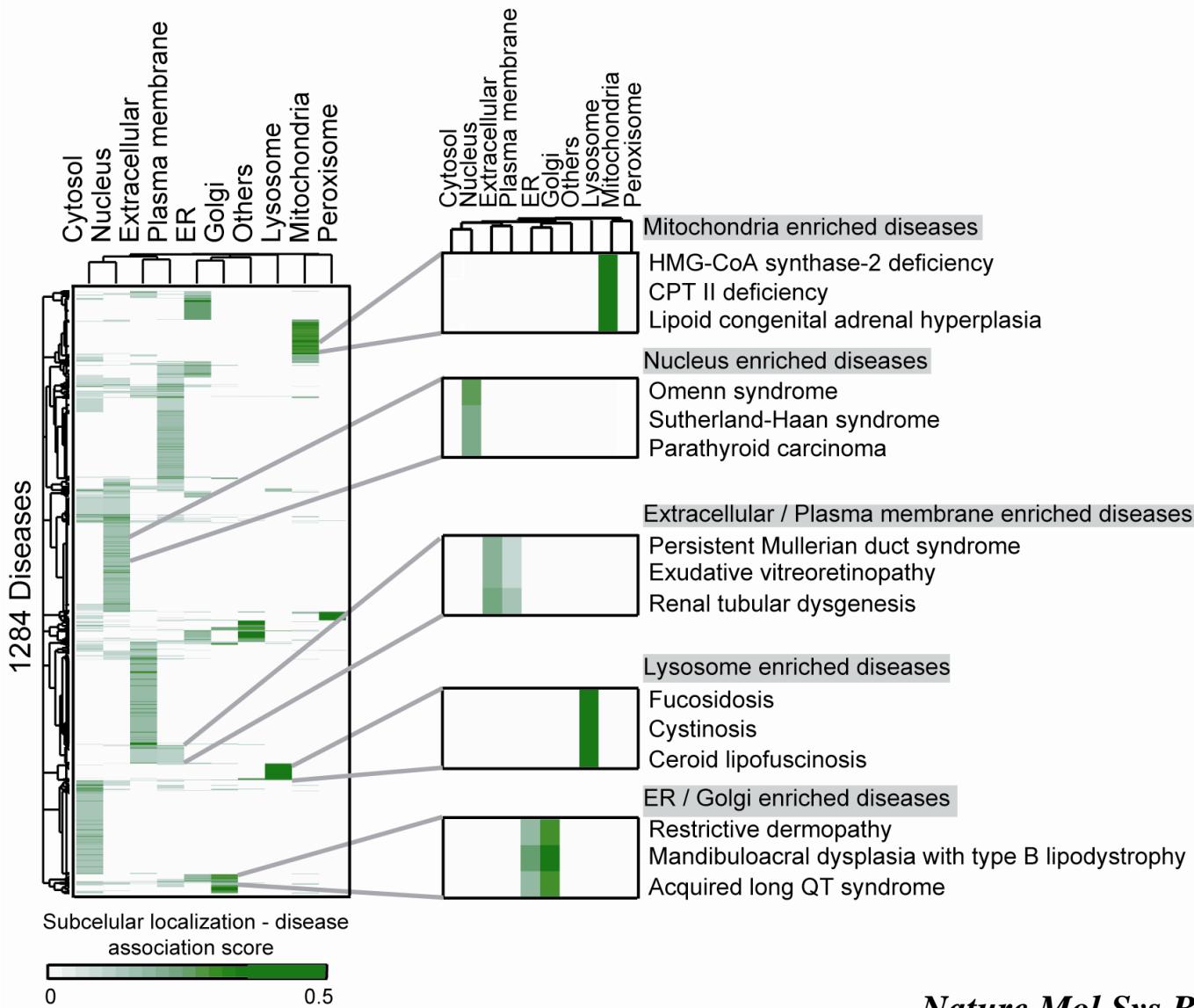
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Protein localization as a principal feature of the etiology and comorbidity of genetic diseases

## Protein subcellular localization and Human diseases

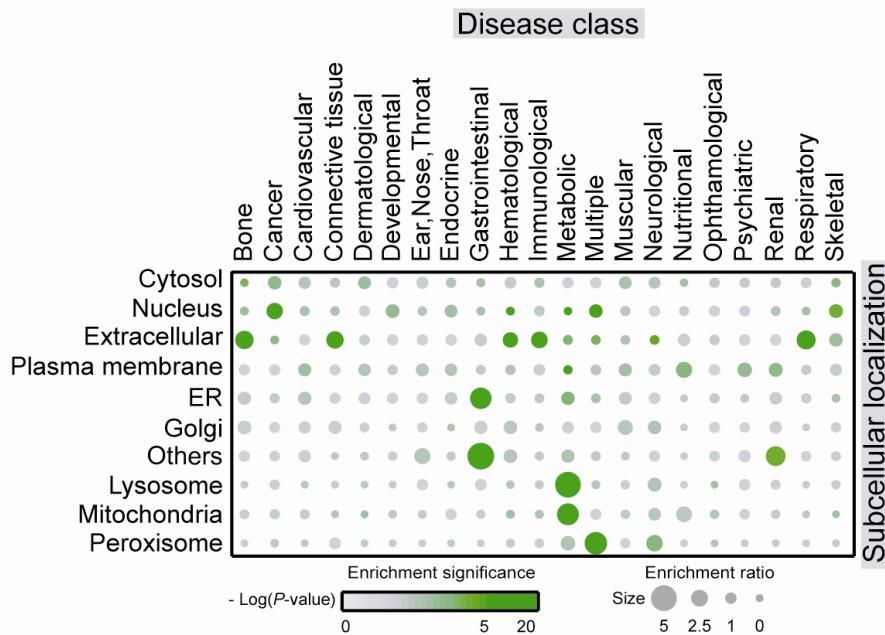


# Relationships between disease-associated proteins and their subcellular localizations

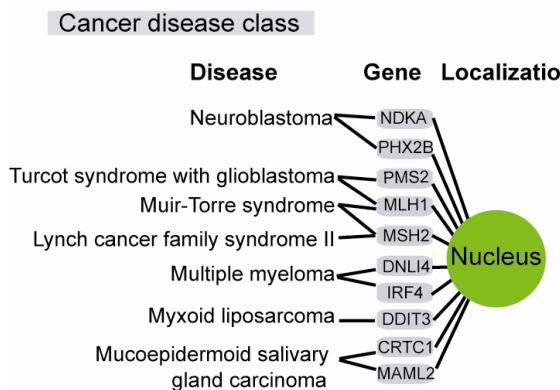


# Correlation between disease classes and subcellular localizations

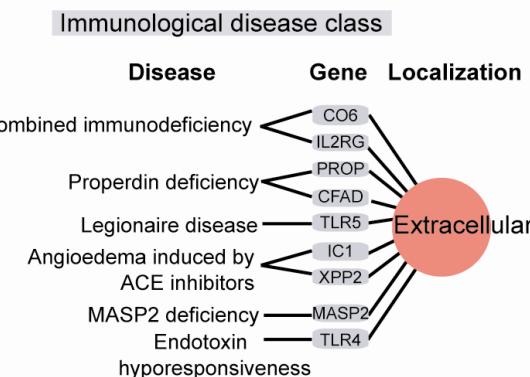
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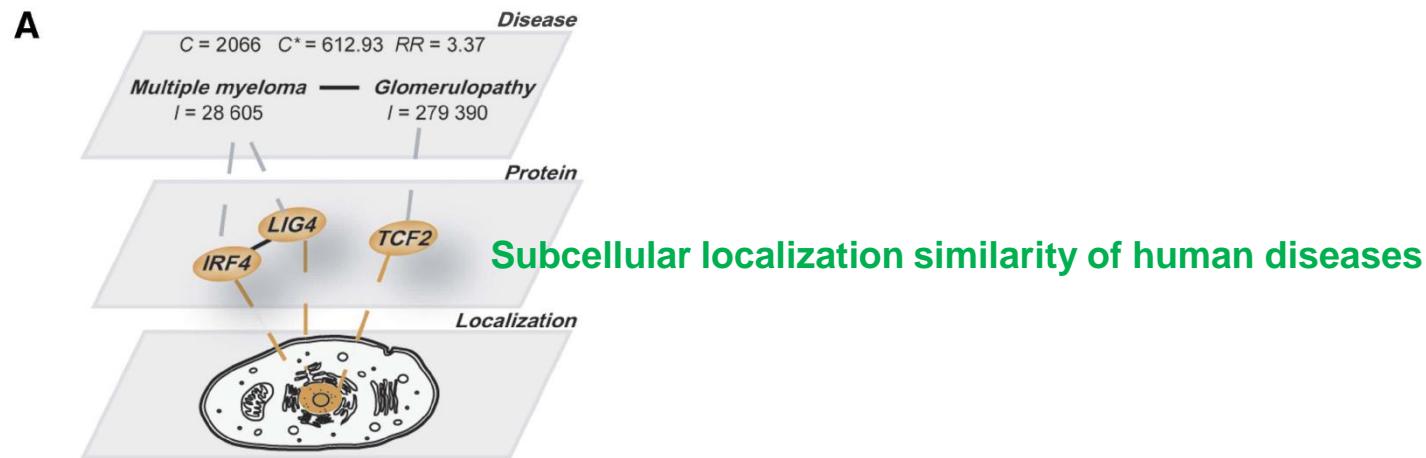
**B**



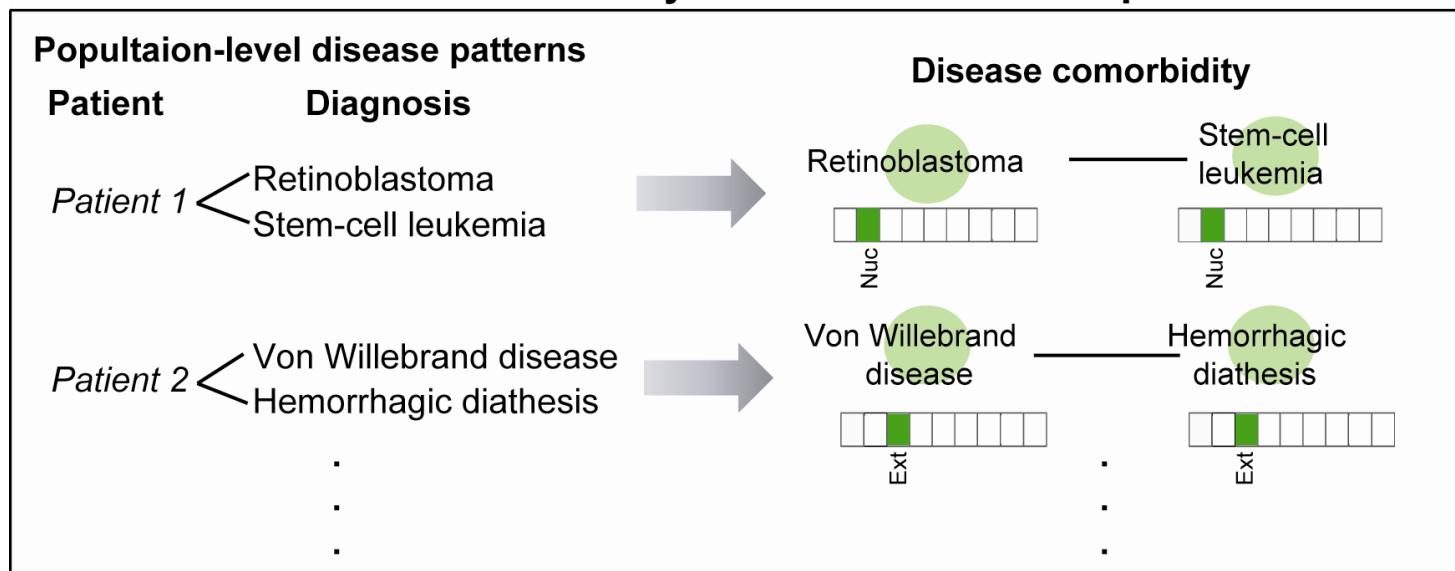
**C**



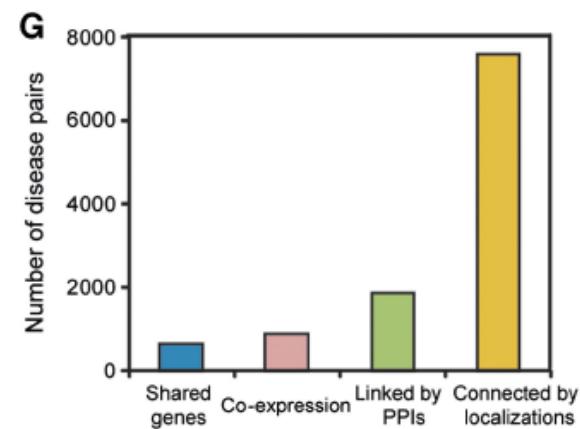
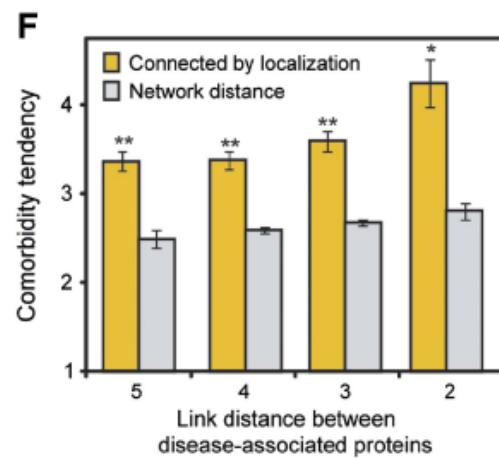
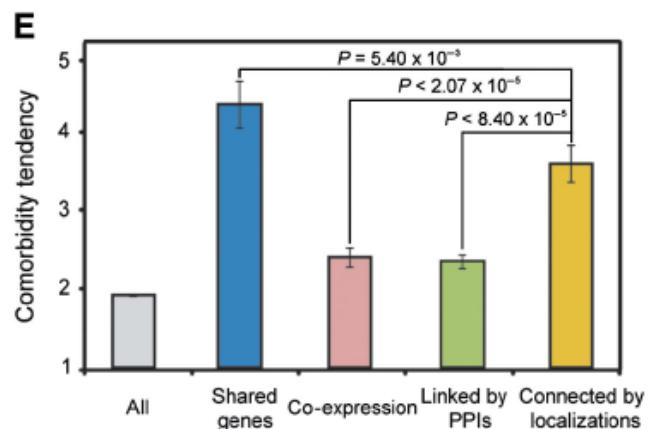
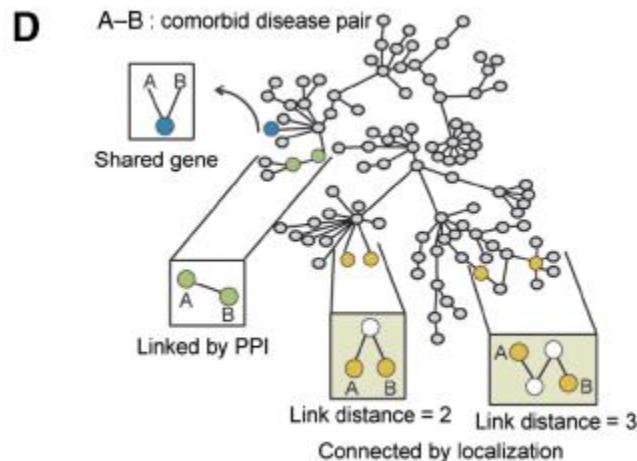
# The implication of subcellular localization for disease comorbidity



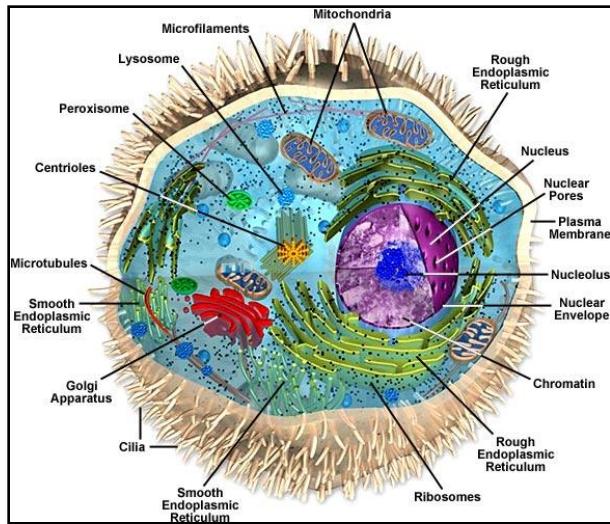
## Subcellular localization similarity of comorbid disease pairs



# The implication of subcellular localization for disease comorbidity



# Disease gene finding through subcellular localization

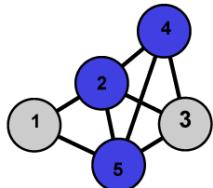


Construction of functional interaction networks through consensus localization predictions of the human proteome.

Park et al. J. Proteome Res., 2009, 8 (7), pp 3367–3376

**A**

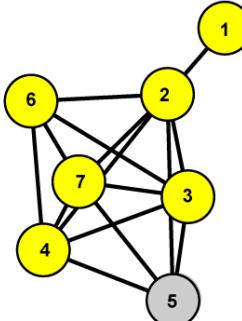
Localization: Plasma membrane  
Disease: Basal cell carcinoma



1: GP107\_Human  
2: PTC1\_Human  
3: HHIP\_Human  
4: SMO\_Human  
5: PTC2\_Human

**B**

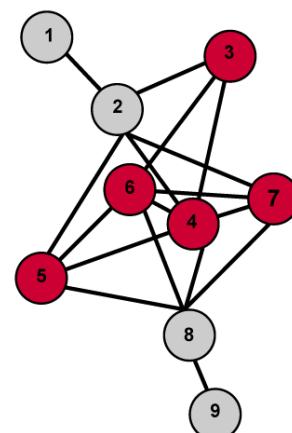
Localization: Cytosol  
Disease: Deafness, autosomal dominant



1: MYO3A\_Human (Con)  
2: WHRN\_Human  
3: MYO15\_Human  
4: OMP\_Human  
5: MYO6\_Human  
6: MYO7A\_Human  
7: USH1C\_Human (Con)

**C**

Localization: Nucleus  
Disease: Mental retardation



1: VEX2\_Human (Con)  
2: ARX\_Human  
3: ZNF81\_Human  
4: JADIC\_Human  
5: AFF2\_Human (Con)  
6: OPHN1\_Human  
7: ZNF41\_Human (Con)  
8: ACSL4\_Human  
9: TMEM9\_Human

> Protein localization information facilitates the identification of disease associated genes

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- Mitochondrial protein network* *Nature Scientific Reports*

2013

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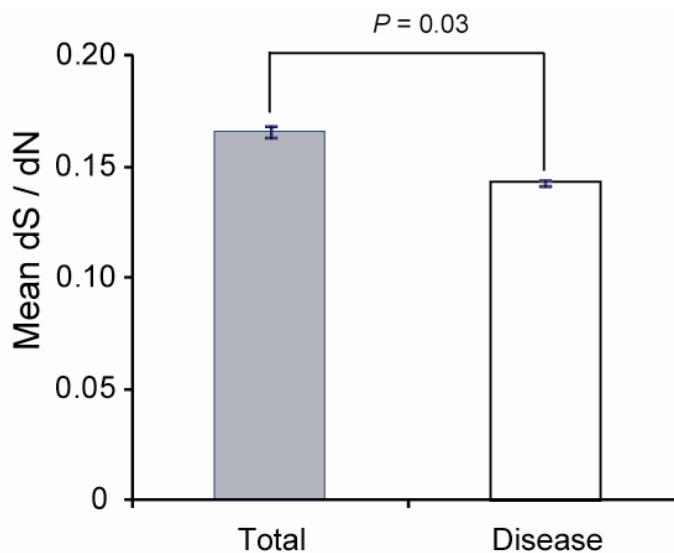
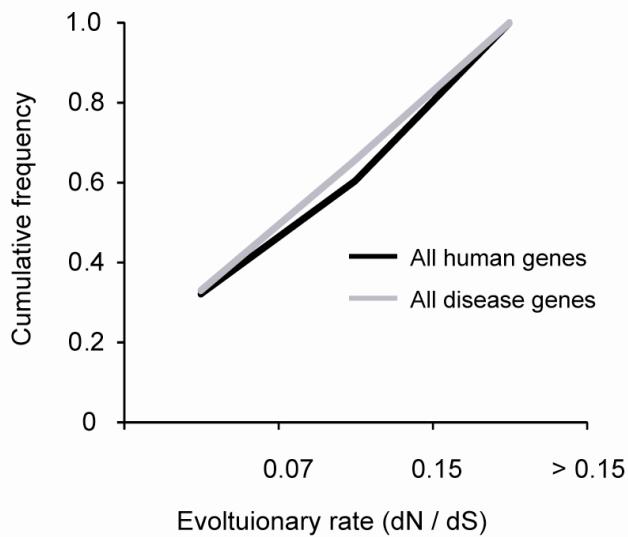
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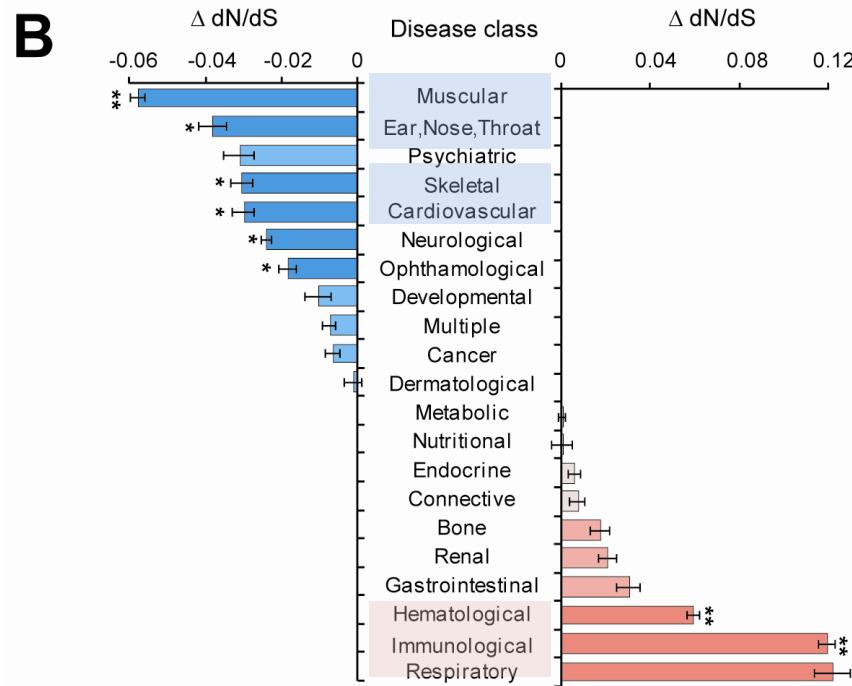
# Human disease genes; fast or slow evolving ?

**A**



Suppl. Figure

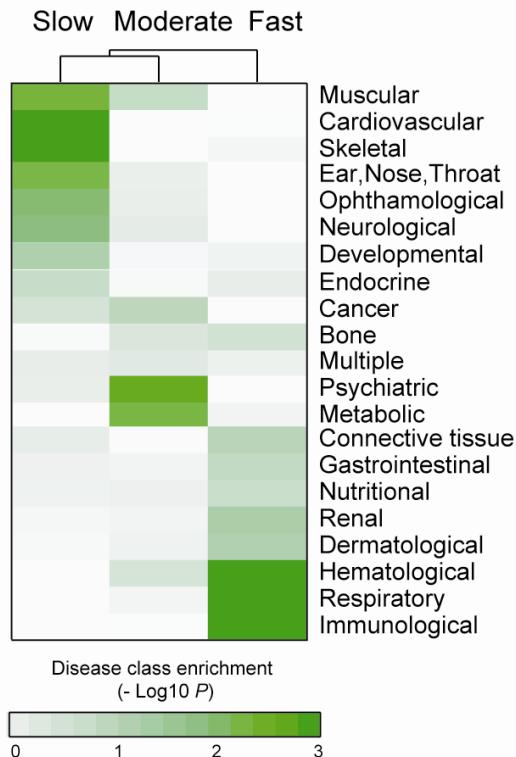
# Human disease genes have diverse evolutionary rates



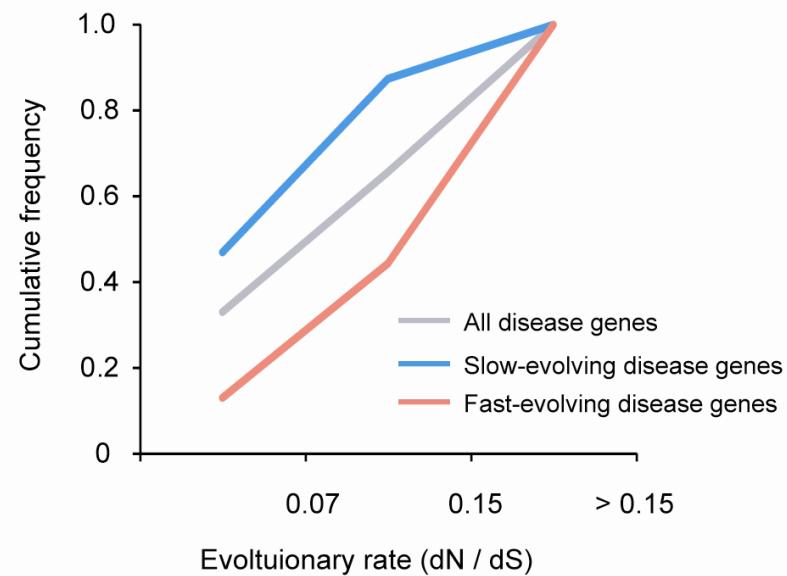
Phenotypically similar disease classes share similar evolutionary history

# Human disease genes have diverse evolutionary rates

C

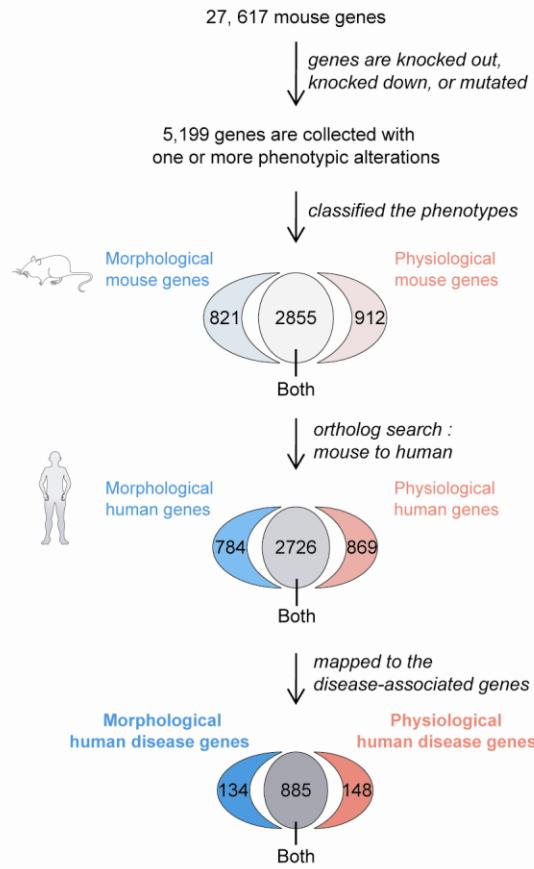


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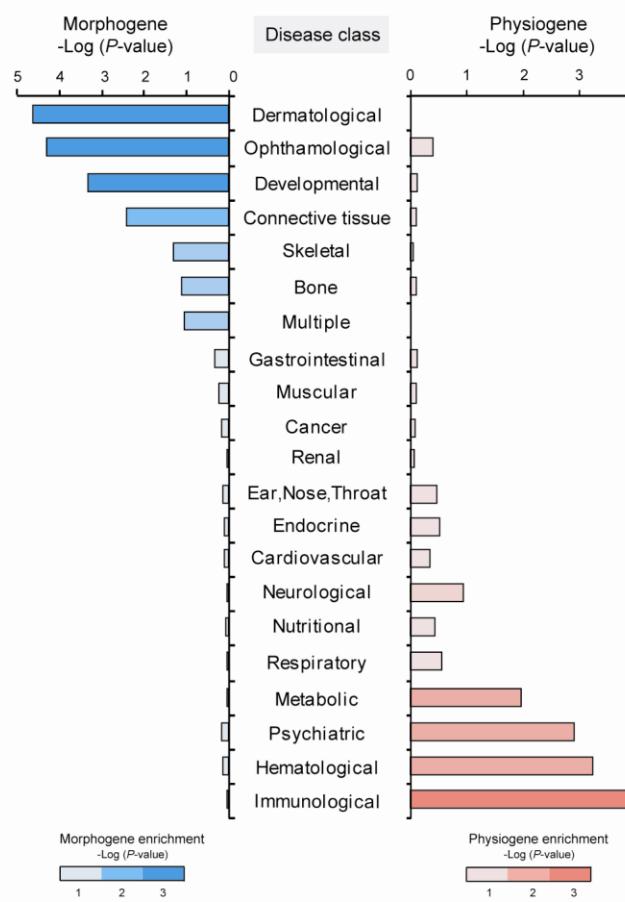


# Morphogenes and physiogenes enriched differently In various disease classes

**A**



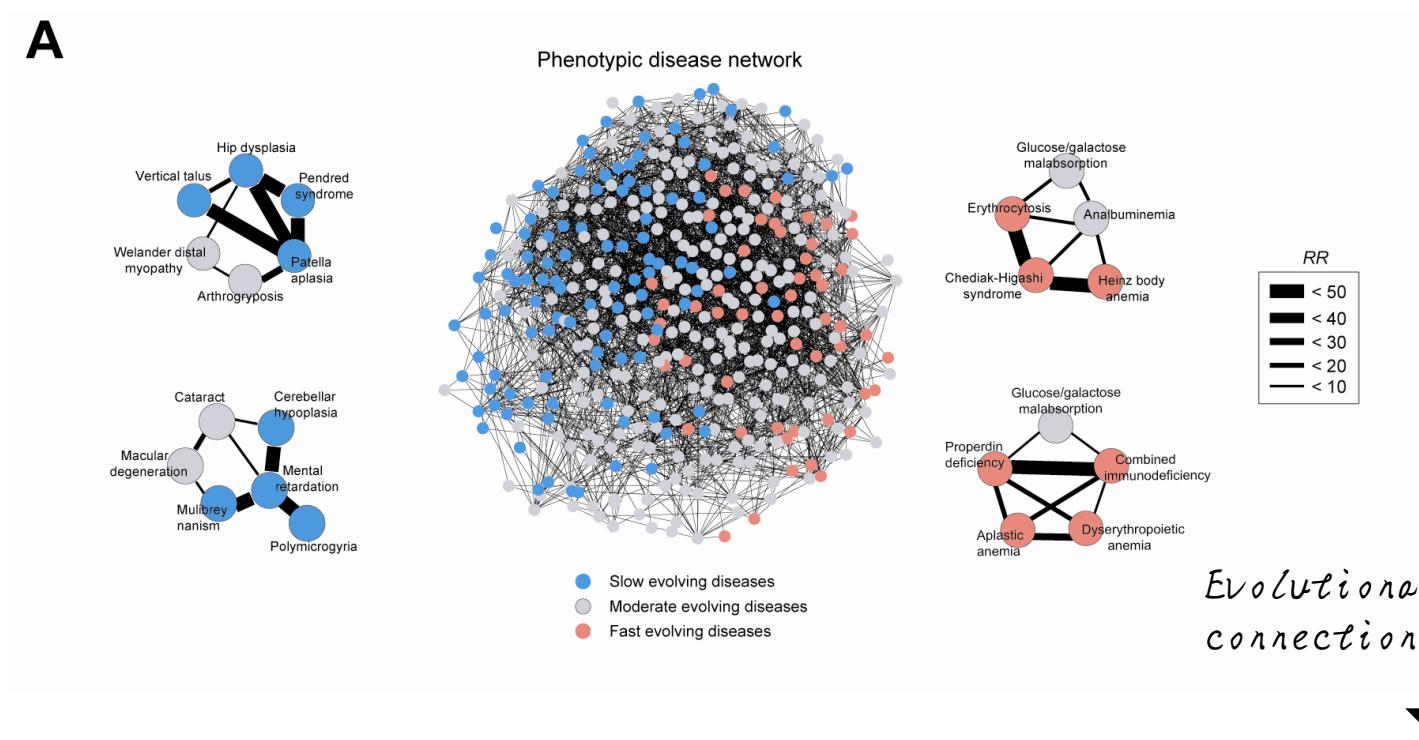
**B**



# Evolution connect genotype to phenotype

## Molecular connections in the comorbid disease pairs

**A**



Evolutionary  
connections

↓

Molecular connections

↓

Phenotypic connections  
: comorbidity

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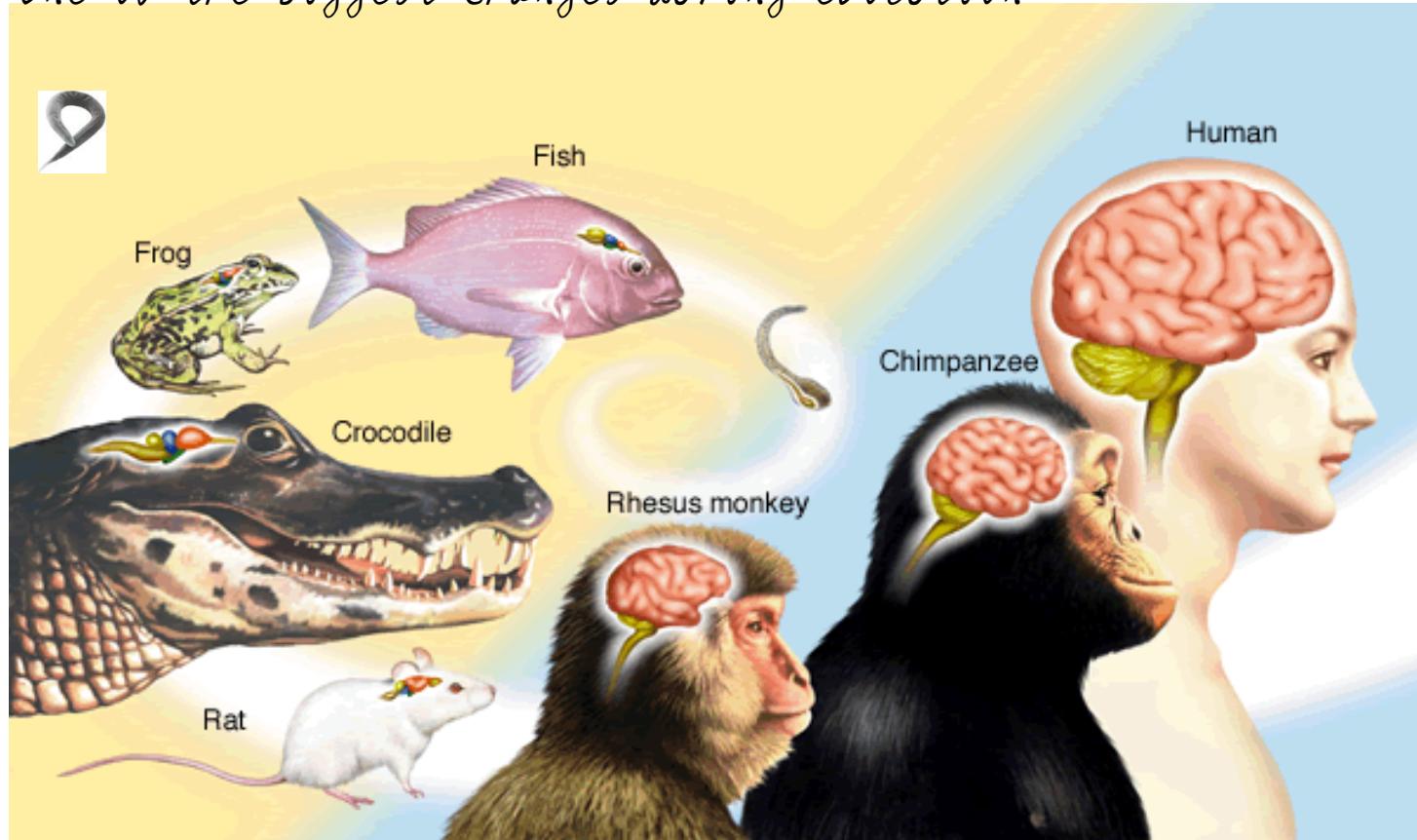
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# Rewiring of PDZ domain-ligand interaction network contributed to eukaryotic evolution

## *Evolution of brain*

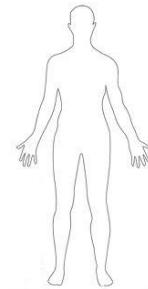
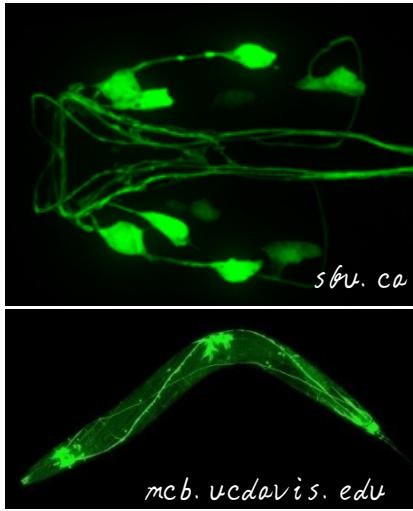
*One of the biggest changes during evolution*



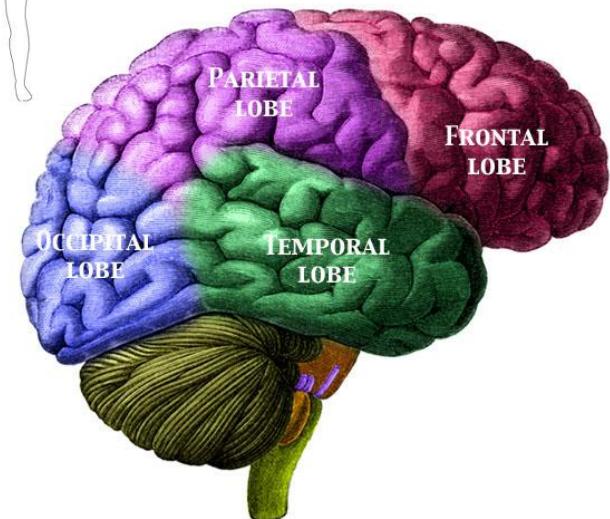
Increase of synaptic diversity (number of brain regions)



*C. elegans* brain



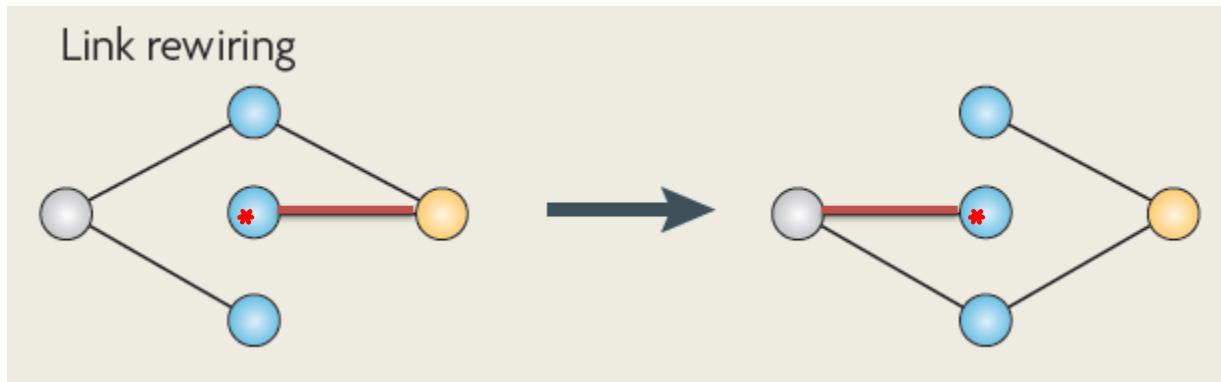
Human brain



[lilachrysikou.wordpress.com](http://lilachrysikou.wordpress.com)

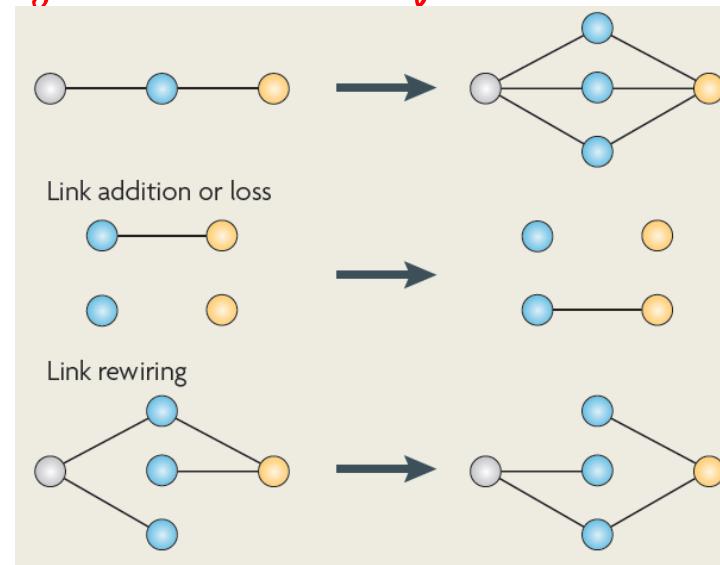
What is the molecular level explanation of neuronal development?

## Rewiring of interactions



T. Yamada and P. Bork, Nat. Rev. Mol. Cell Biol. 2009

Interaction rewiring can reconfigure molecular systems without a gain or loss of gene.

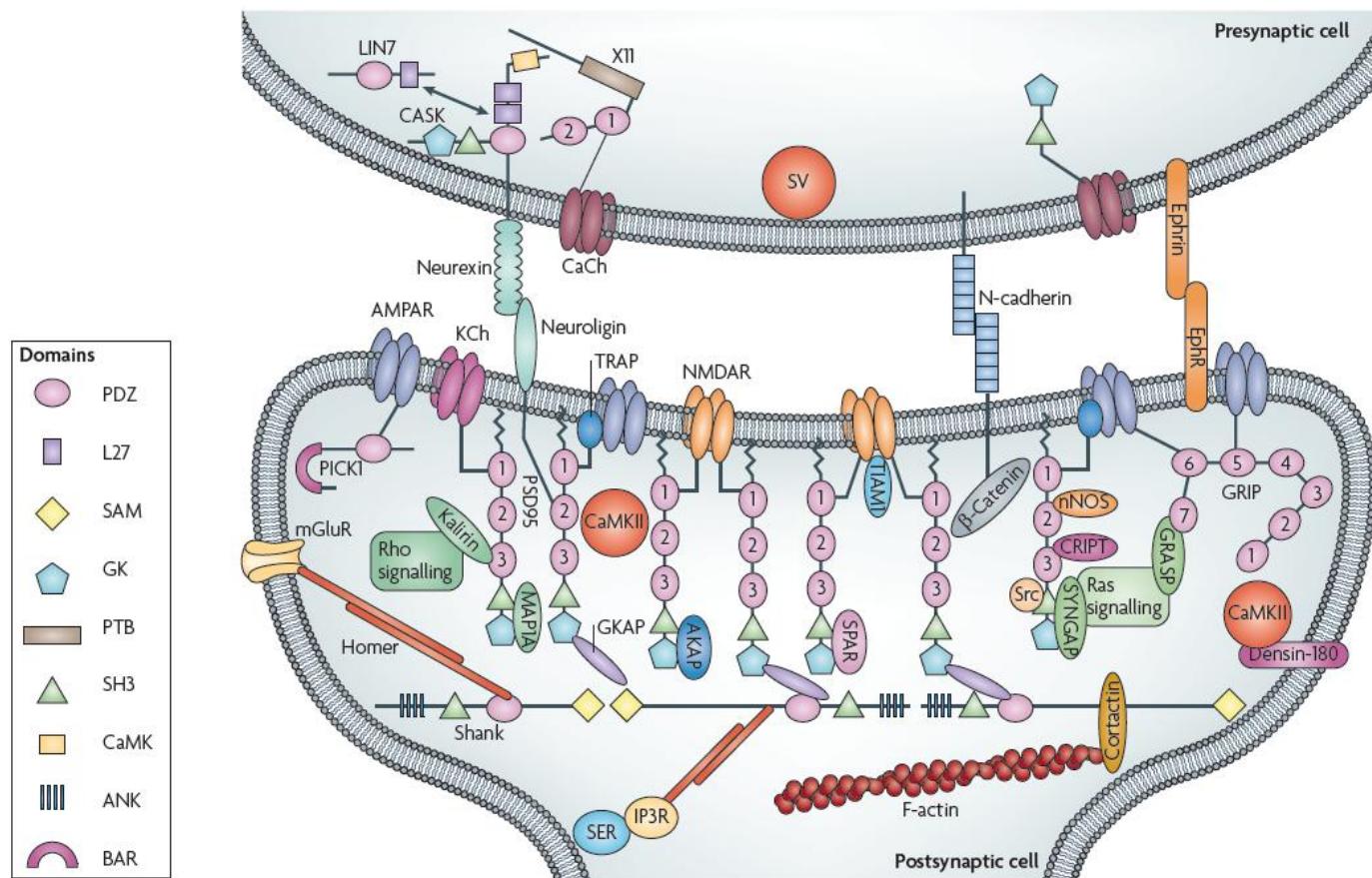


# PDZ protein interactions play important roles in the postsynaptic density (PSD)

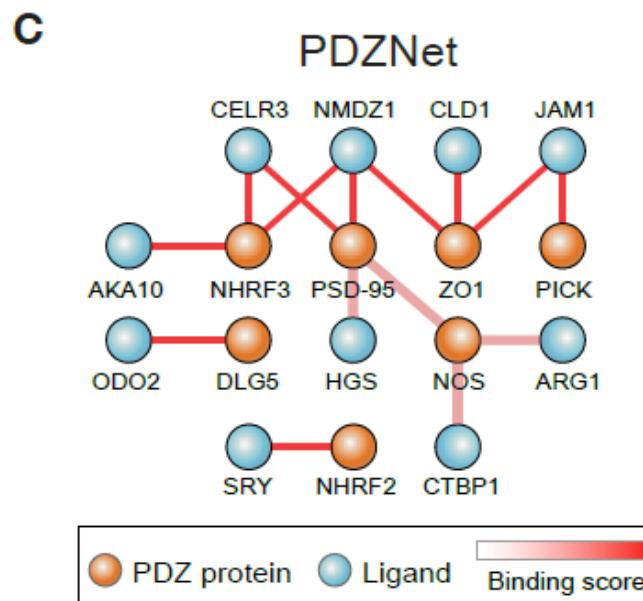
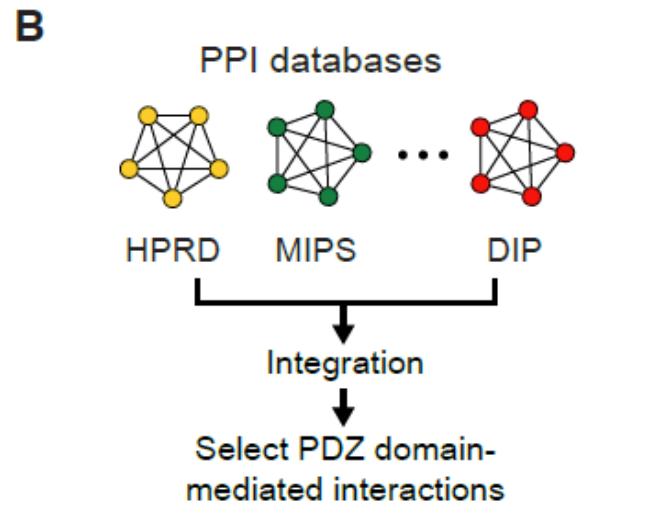
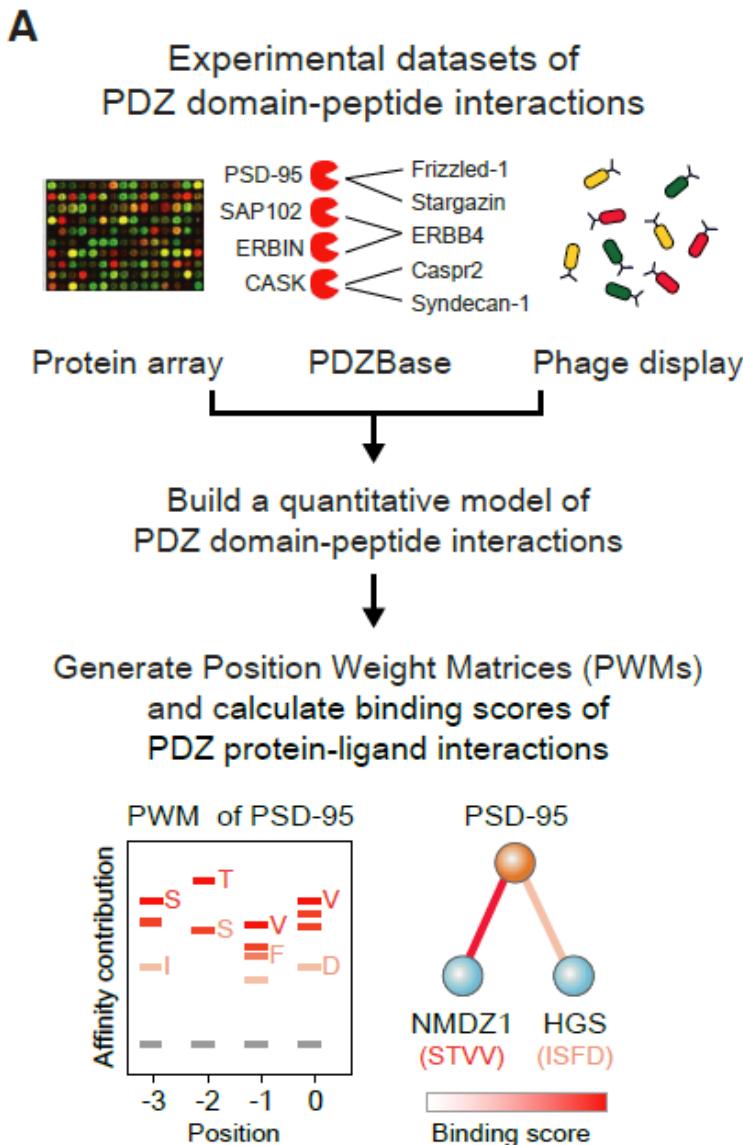
## Organization and dynamics of PDZ-domain-related supramodules in the postsynaptic density

Wei Feng and Mingjie Zhang

NATURE REVIEWS | NEUROSCIENCE | FEBRUARY 2009 |

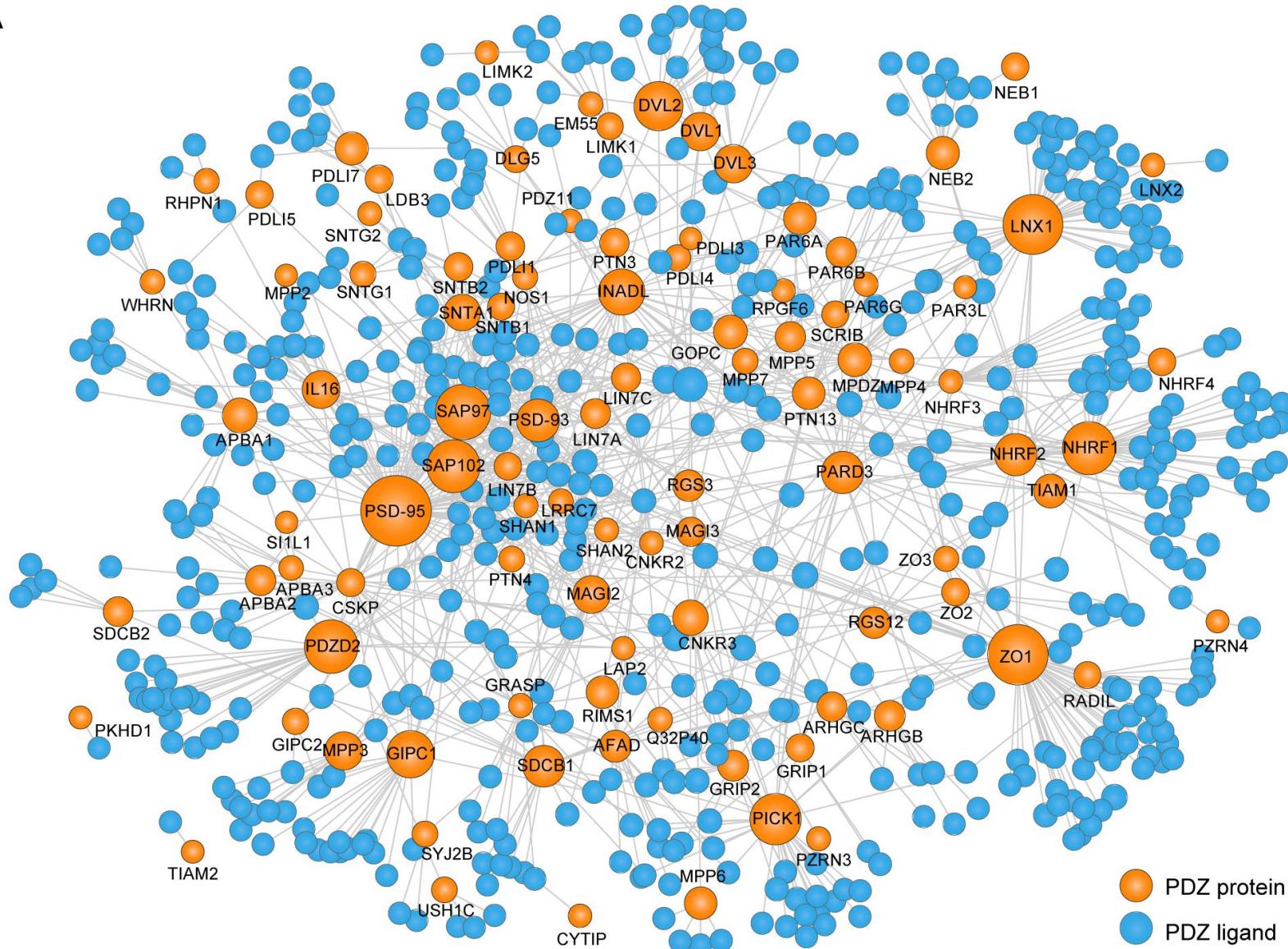


# Construction of human PDZ domain-ligand interaction network (PDZNet)

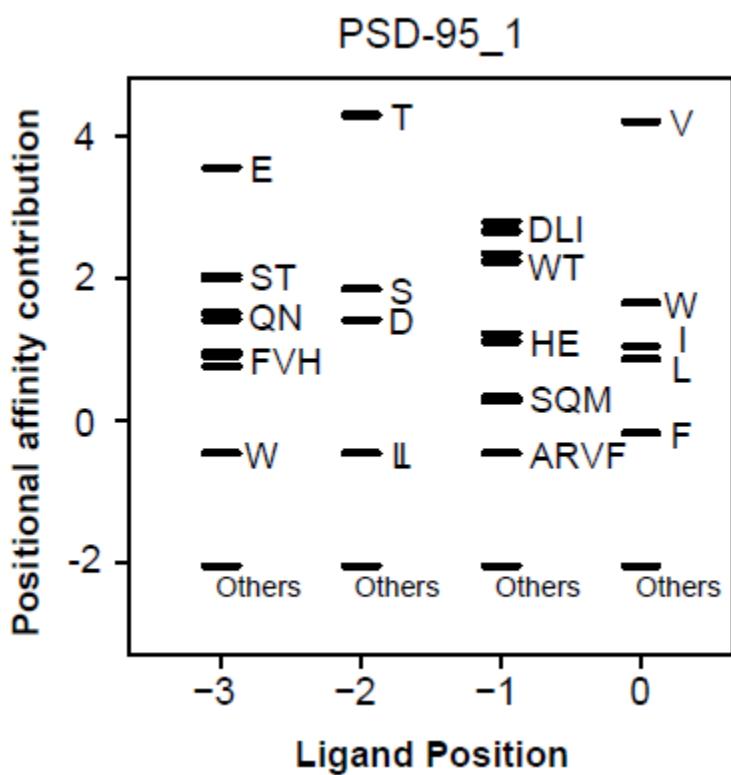


## Human PDZNet

A



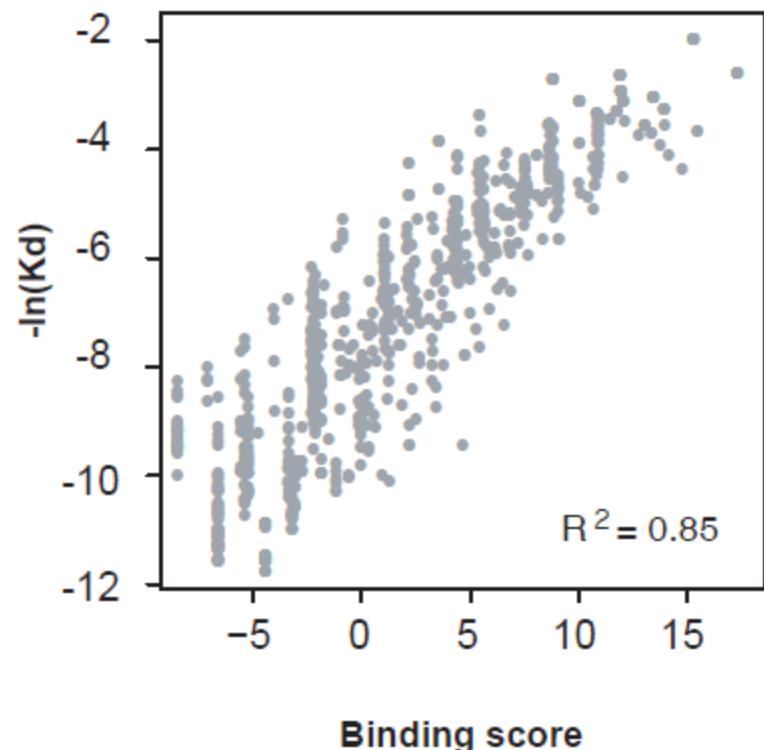
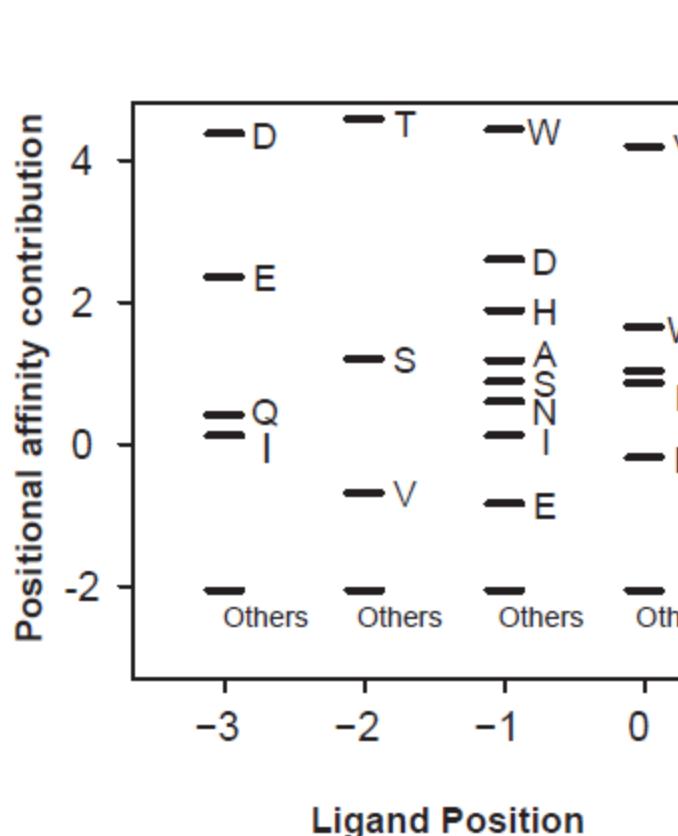
PWM can identify the known binders of PDZ domains



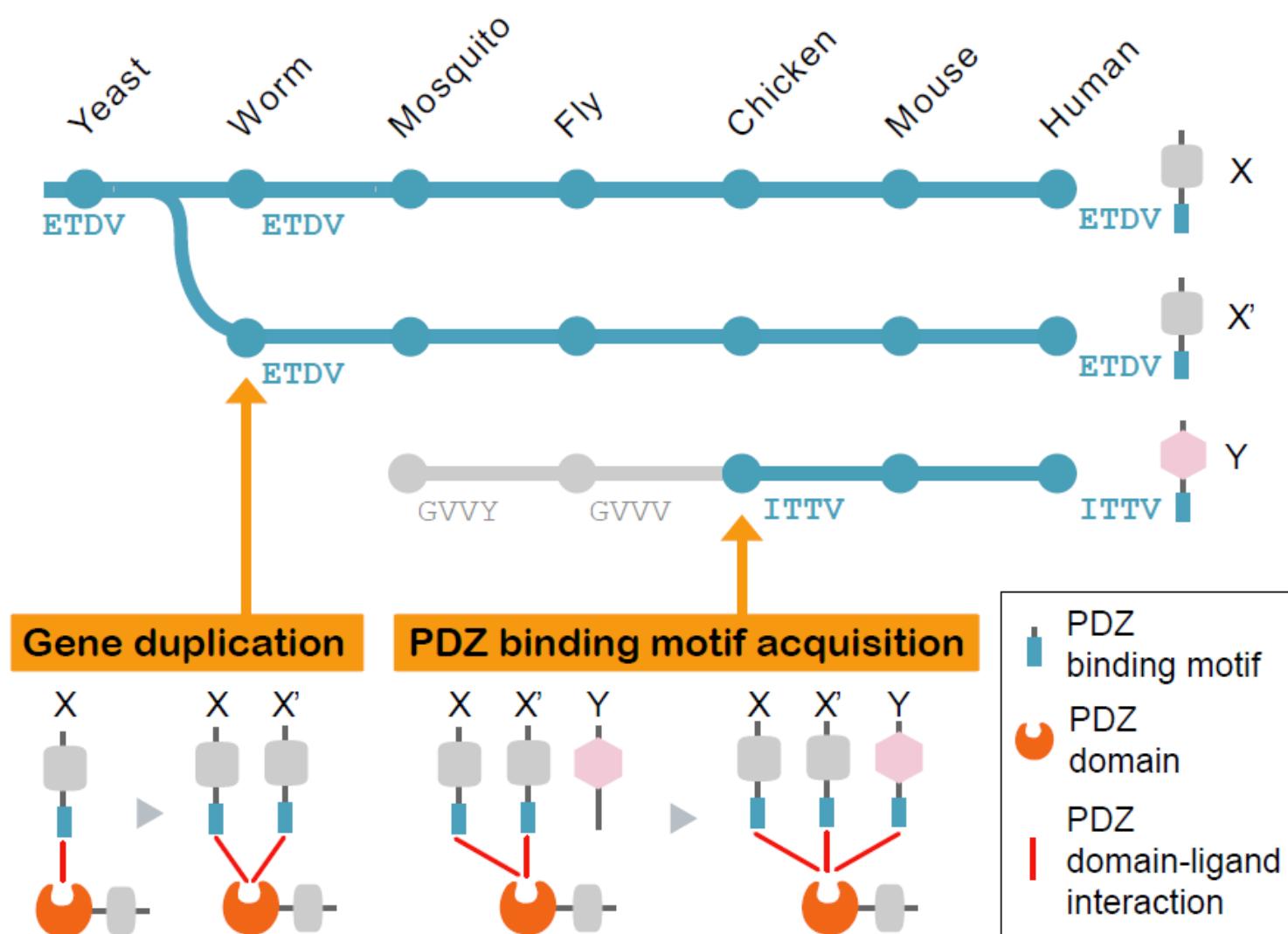
Species	Ligand	C-terminal	Percentile rank (%)	Binding score
Human	PMCA4b	ETSV	99.86	13.05
	PMCA2b	ETSL	99.26	9.49
Mouse	Frizzled-4	ETVV	99.96	13.80
	Frizzled-1	ETTV	99.93	13.49
	Frizzled-2	ETTV	99.93	13.49
	Frizzled-7	ETAV	99.87	13.19
	Sema4c	ESSV	99.59	10.61
	BAI1	QTEV	99.56	10.52
	Stargazin	TTPV	99.36	9.77
Rat	Kv1.4	ETDV	99.98	14.72
	ERBB4	NTVV	99.76	12.67
	NMDAR2A	ESDV	99.72	12.28
	NMDAR2B	ESDV	99.72	12.28
	SynGAP	QTRV	99.19	9.79
	PKC-A	QSAV	98.64	8.99
	Sec8	ITTV	98.12	7.93
	GluR6	ETMA	96.19	6.00

Binding scores correlate well with experimental affinities

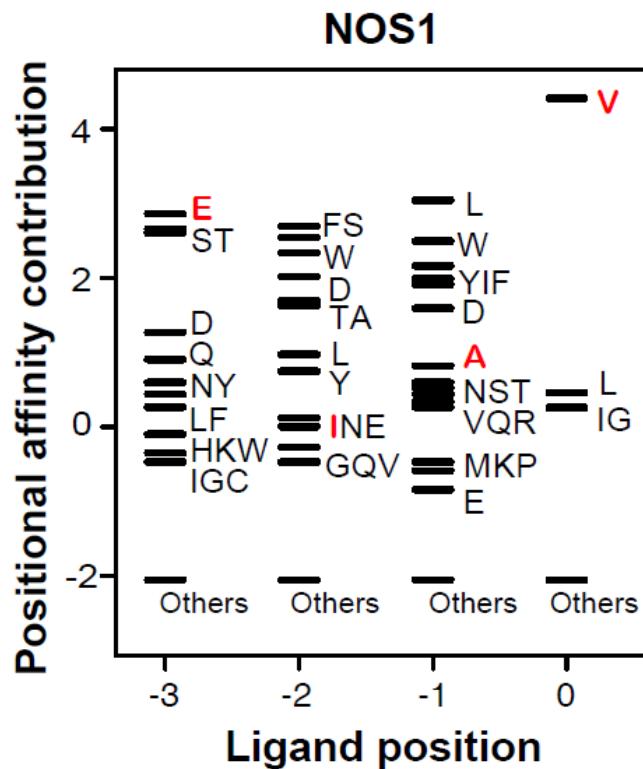
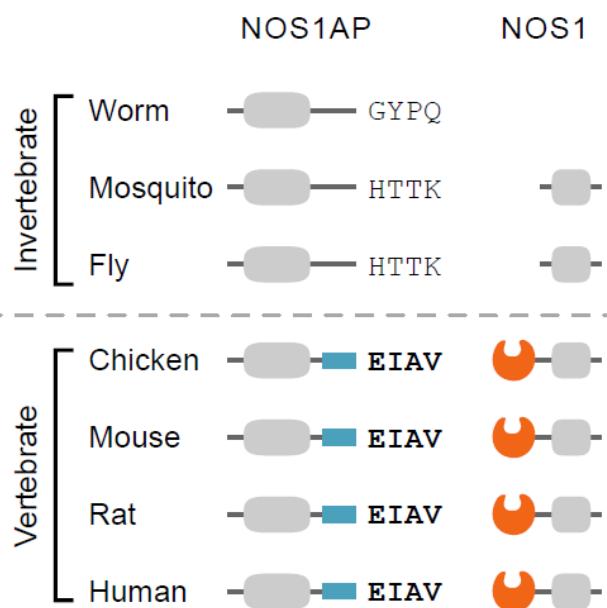
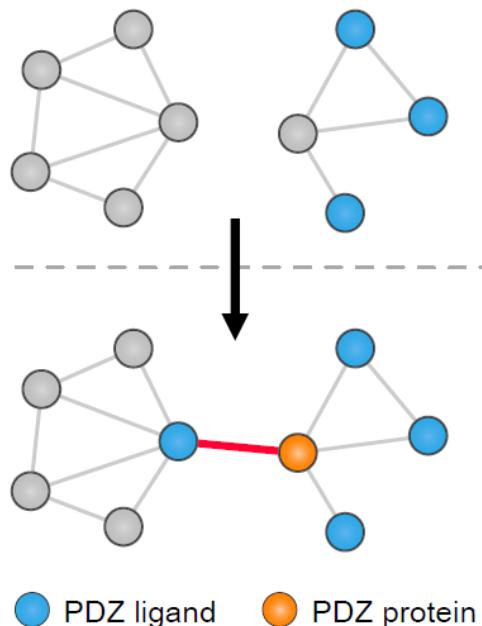
ERBIN human



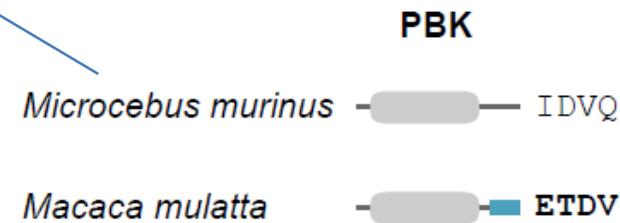
Two evolutionary models describe the expansion of PDZ domain-ligand interactions



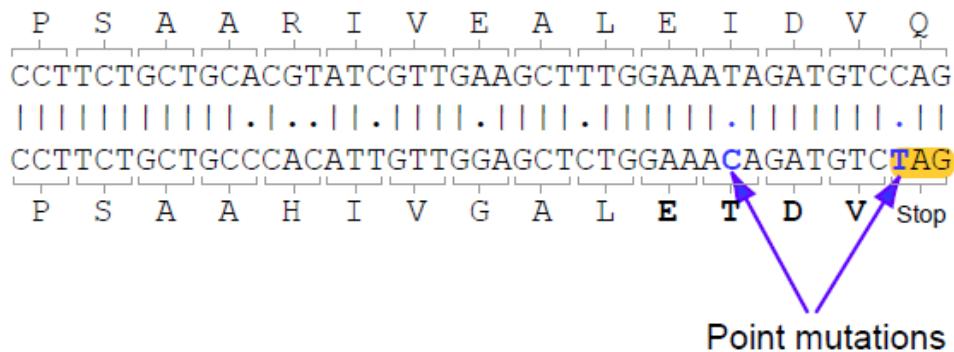
Examples of a PDZ domain-ligand interaction created by sequence mutations



A point mutation generated a PDZ-binding motif in the C-terminal amino acids of the *Macaca mulatta* PBK protein.



*Microcebus murinus*  
*Macaca mulatta*



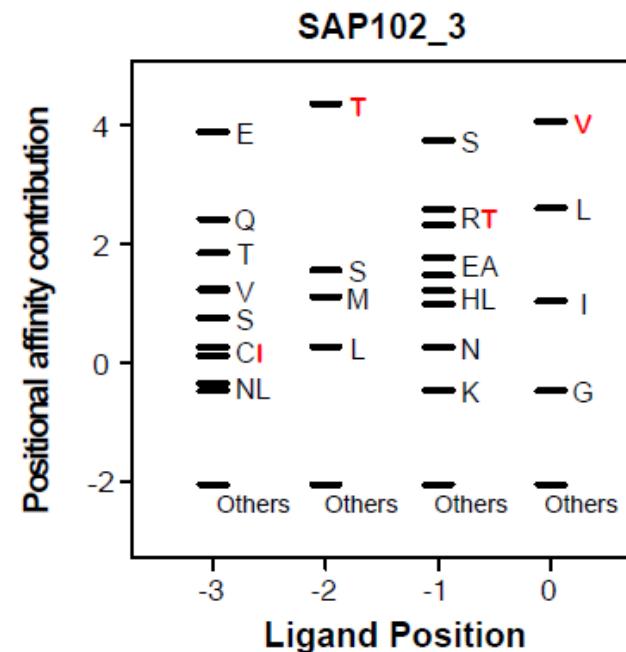
A DNA segment insertion generated a PDZ-binding motif in the C-terminal amino acids of the *Oryzias latipes* EXOC4 protein.



*Anopheles gambiae* EXOC4 GVTV

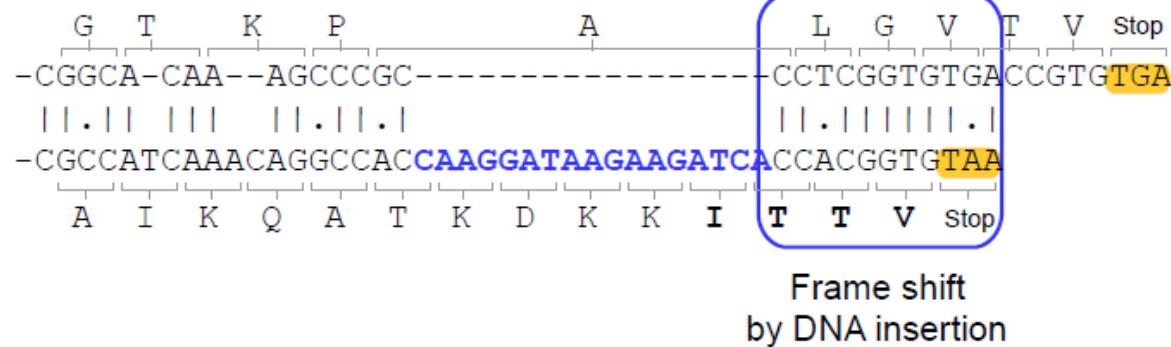


*Oryzias latipes* EXOC4 ITTV

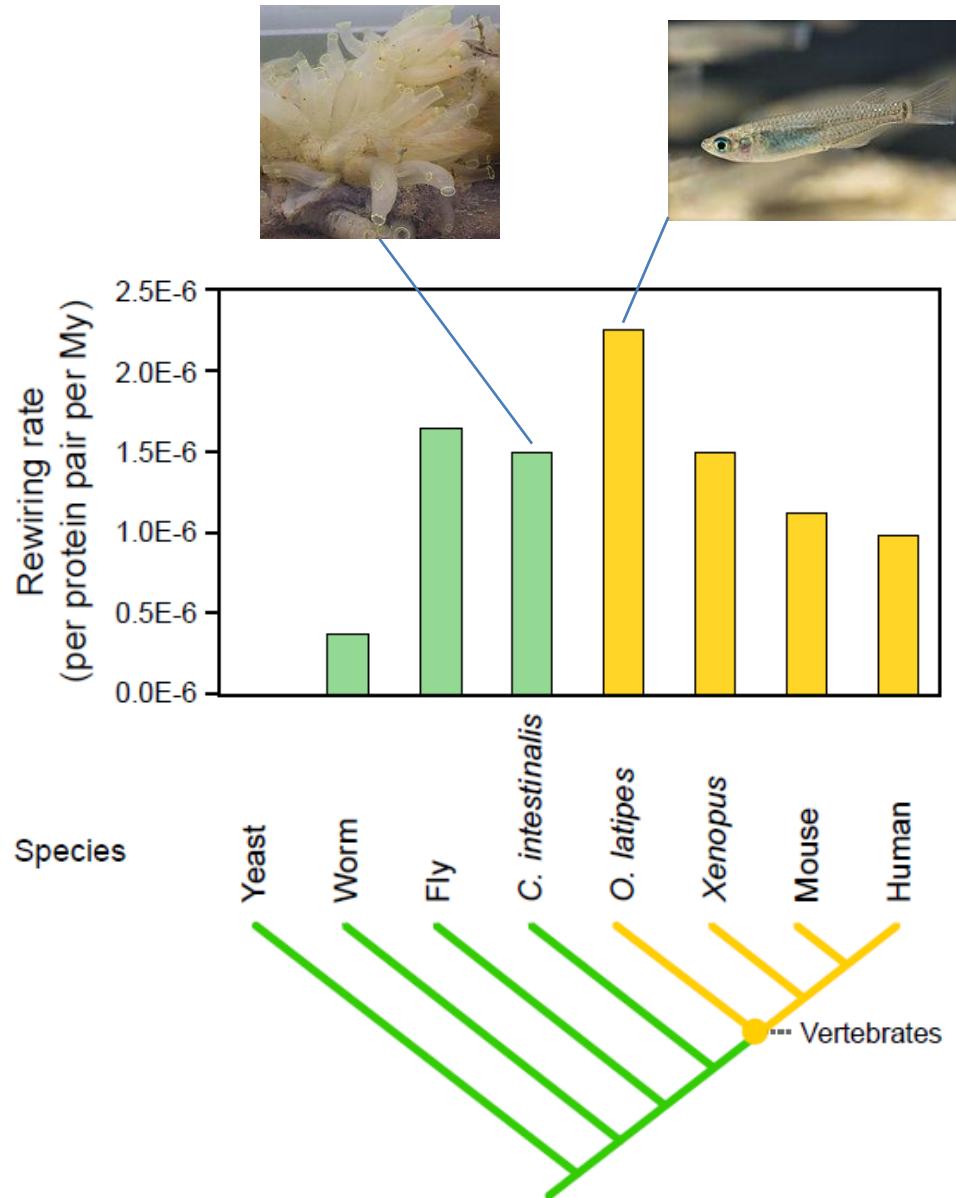


*Anopheles gambiae*

*Oryzias latipes*



Rewiring of PDZ domain-ligand interactions plays an important role in the evolution of nervous systems in vertebrates.



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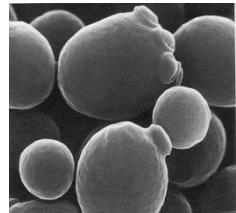
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5. A multifunctional core-shell nanoparticle for dendritic cell-based cancer immunotherapy *Nature Nanotechnology* 2011 6(10):675-682.
6. Network clustering revealed the systemic alterations of mitochondrial protein expression" *PLoS Comp. Biol.* 2011 7(6):e1002093.
7. Protein localization as a principal feature of the etiology and comorbidity of genetic diseases *Nature Mol Sys Biol.* 2011 7:494.

Increase of network complexity has a major impact on gene essentiality changes.



Mechanism of gene essentiality

Complex relationship between genotype and phenotype

Gene essentiality often changes during evolution

*Map2k1* (nonessential in yeast but essential in mouse)

- "Map2k1-/- embryos die at mid-gestation from abnormal development and hypovascularization of the placenta." Vickram Bissonauth, et al. *Development* 2006
- "In the mouse, loss of *Map2k1* function causes embryonic lethality." Valérie Nadeau, et al. *Development* 2009

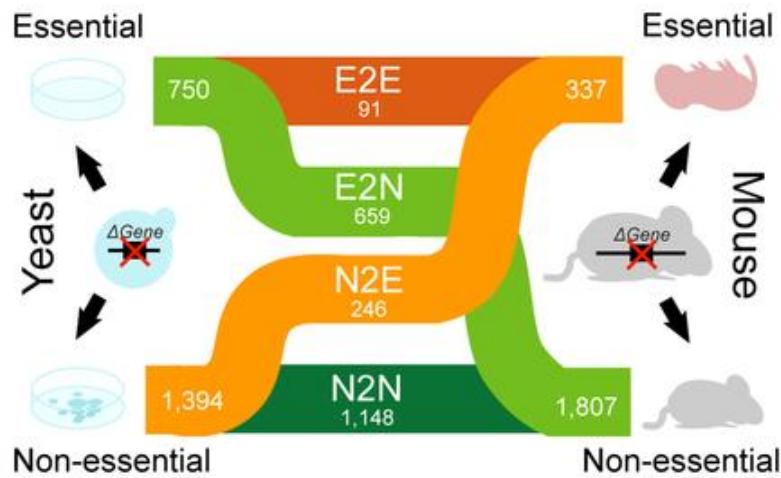
*Ada* (nonessential in yeast but essential in mouse)

- "Adenosine deaminase deficient (*Ada*) mice die perinatally." Alexandra A. J. Migchelsen, et al. *Nat. Genetics* 1995

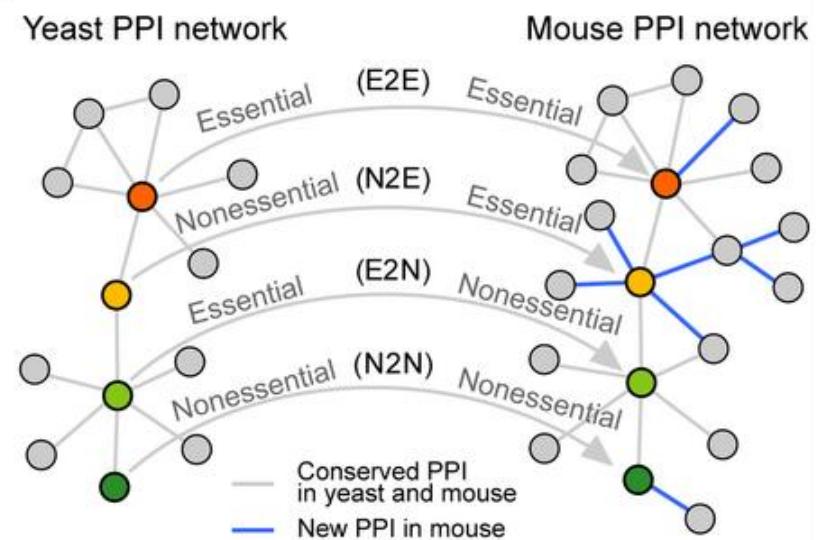
It is unclear how nonessential genes become essential in more complex organisms

# Increase in network connections and gene essentiality changes between yeast and mouse

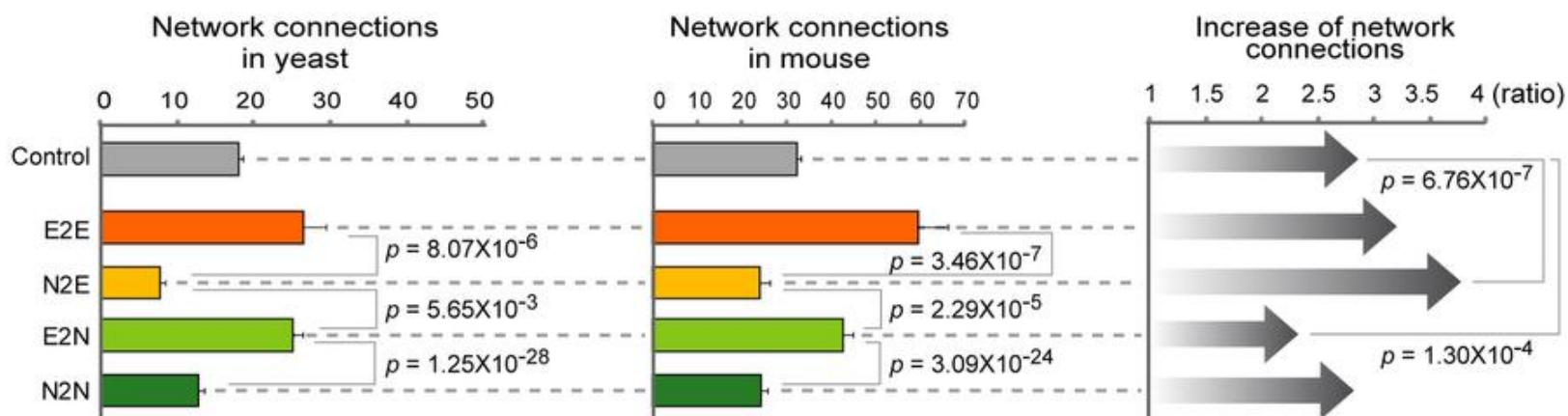
a



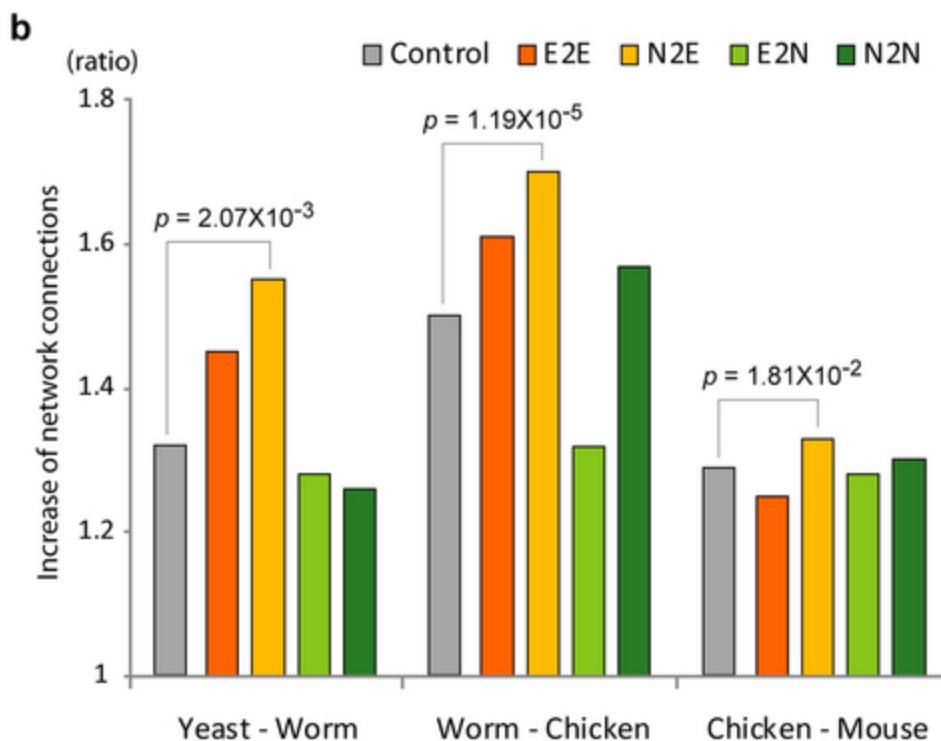
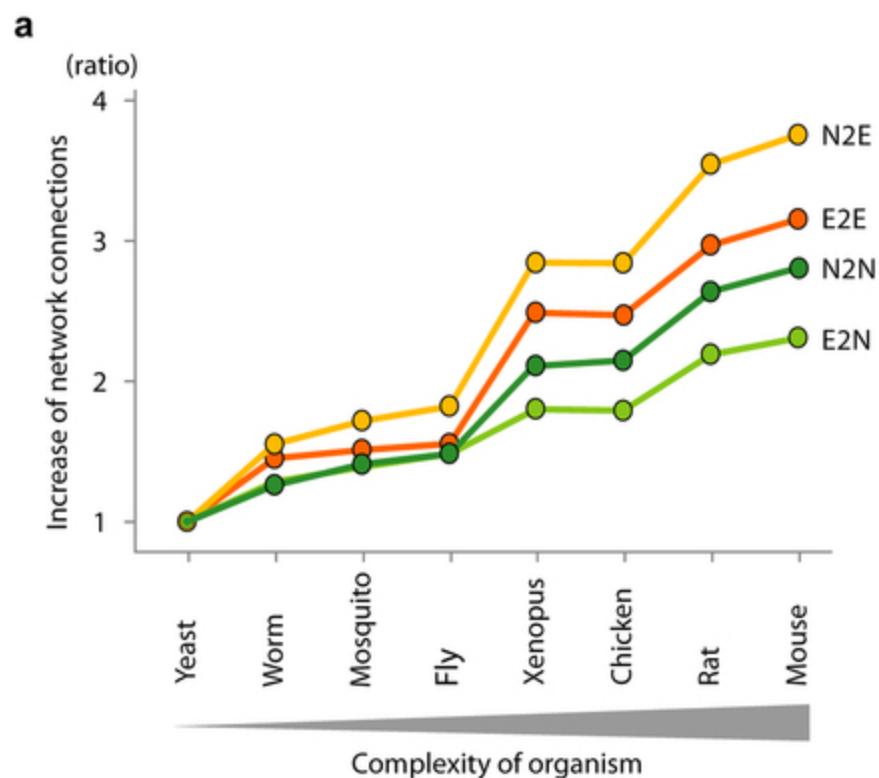
b



c

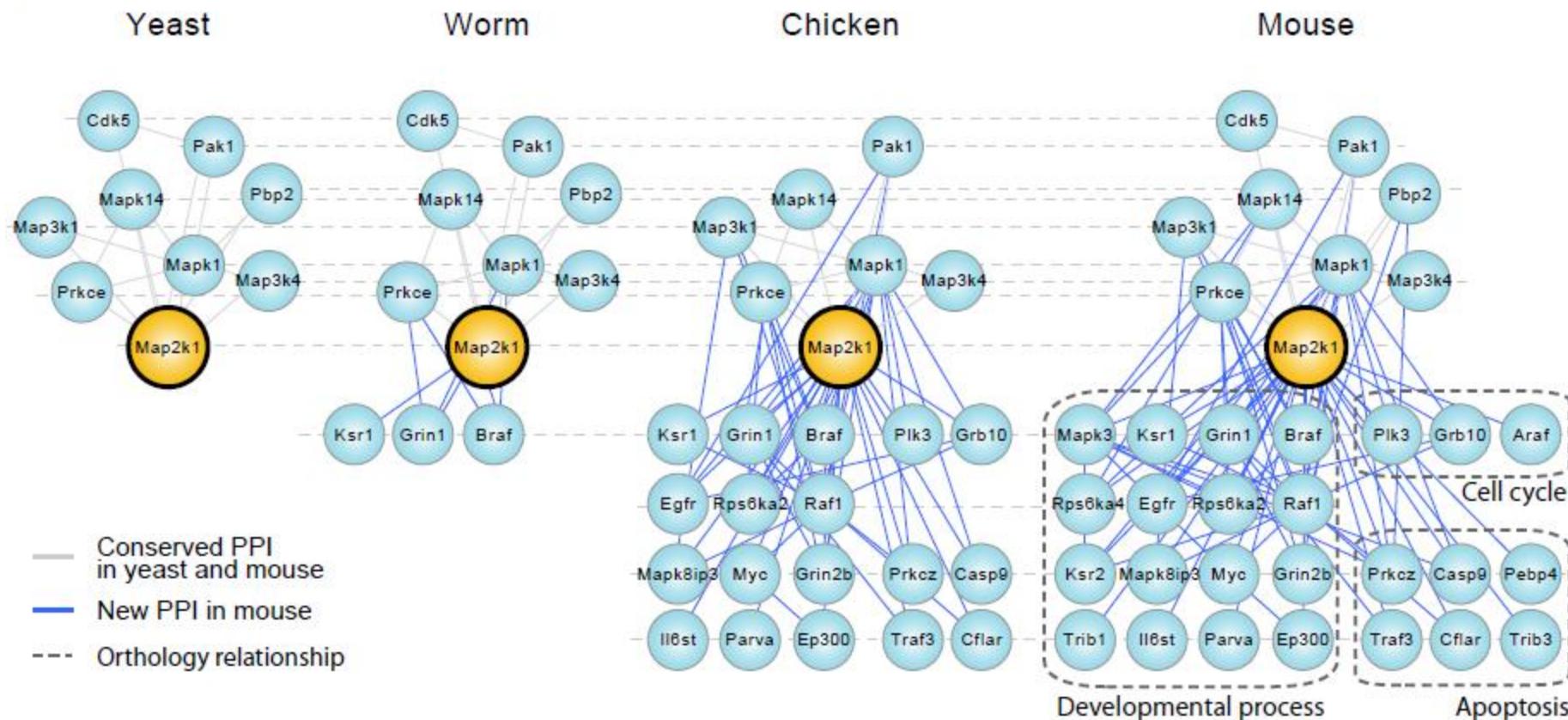


# Comparison of network connections in various species



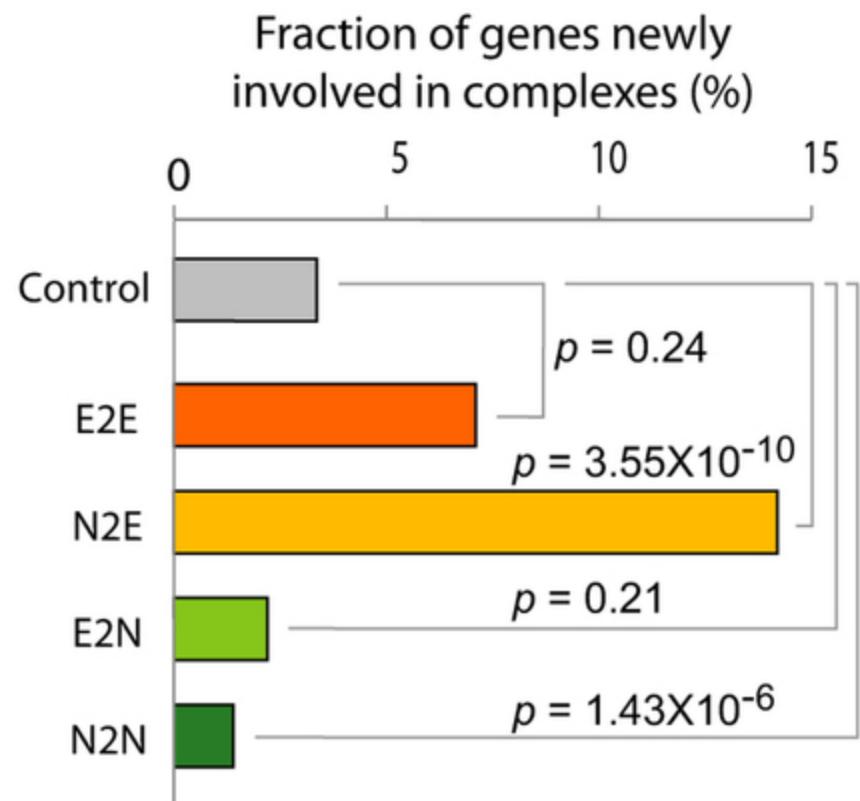
## Network connections of Map2k1 in yeast, worm, chicken, and mouse

C

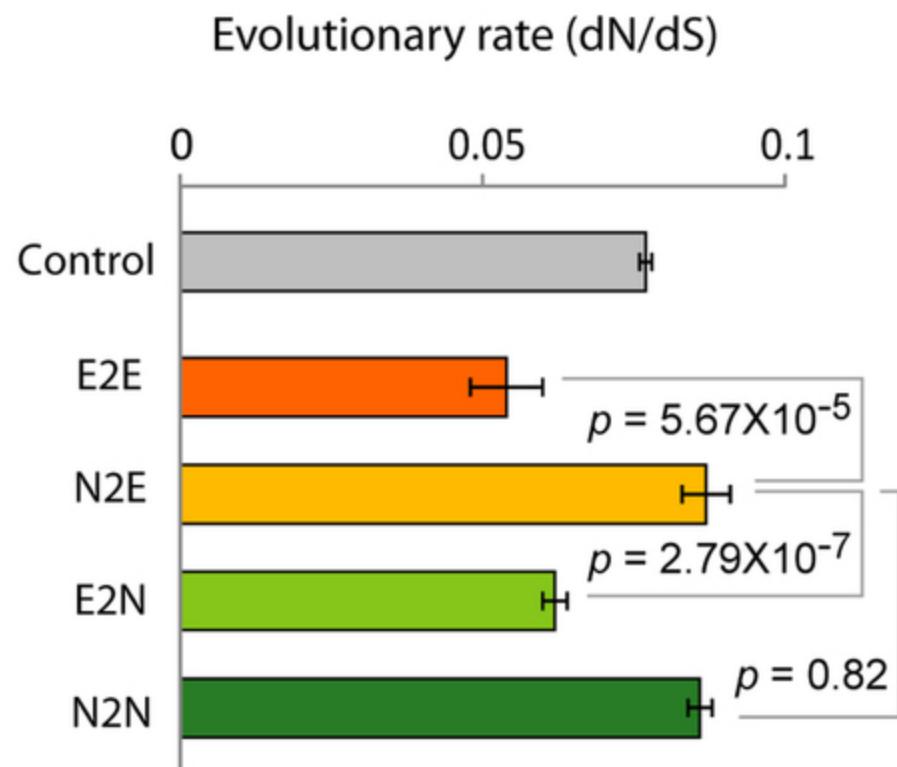


# Protein complex membership and evolution of gene essentiality changes

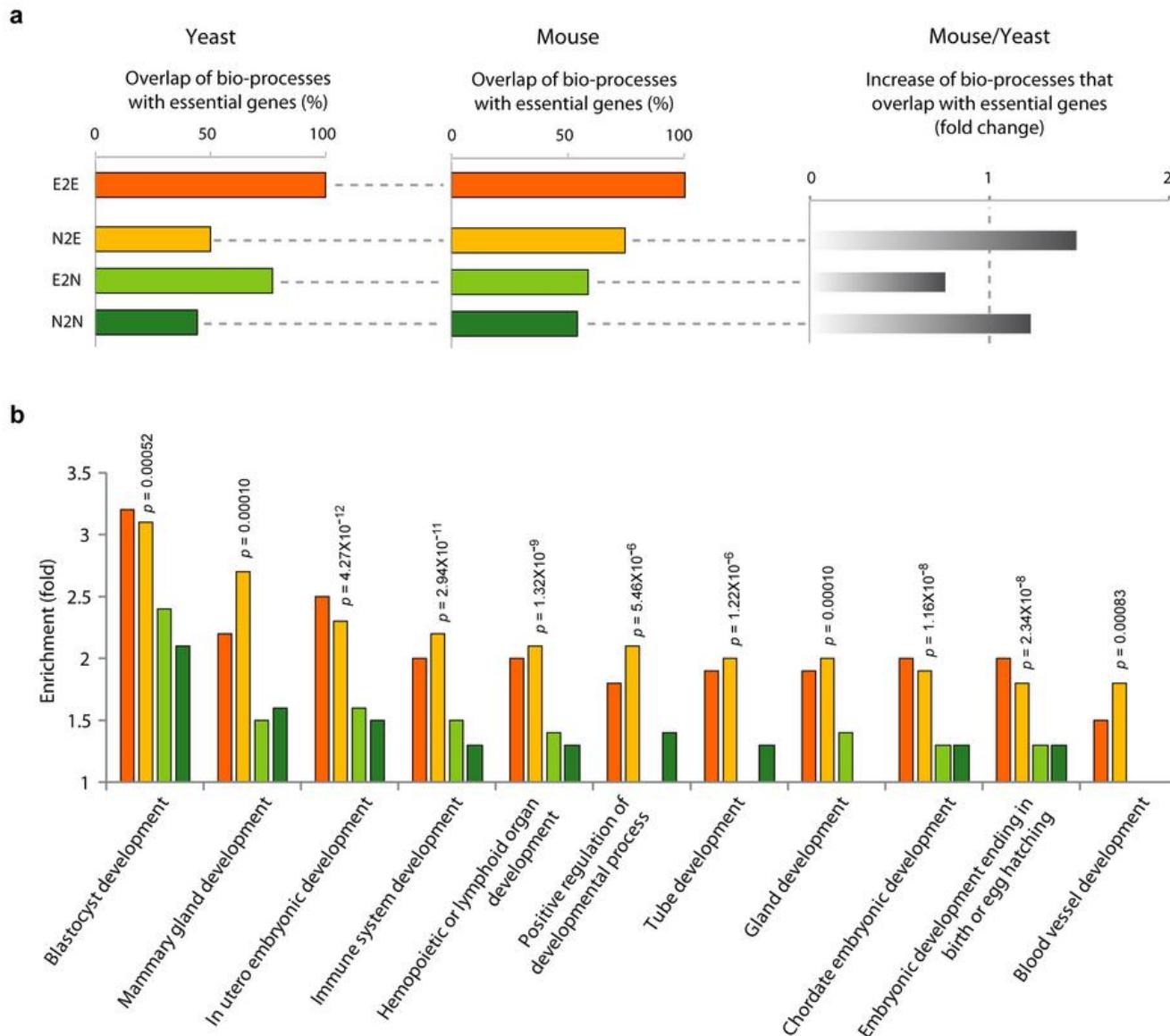
**a**



**b**

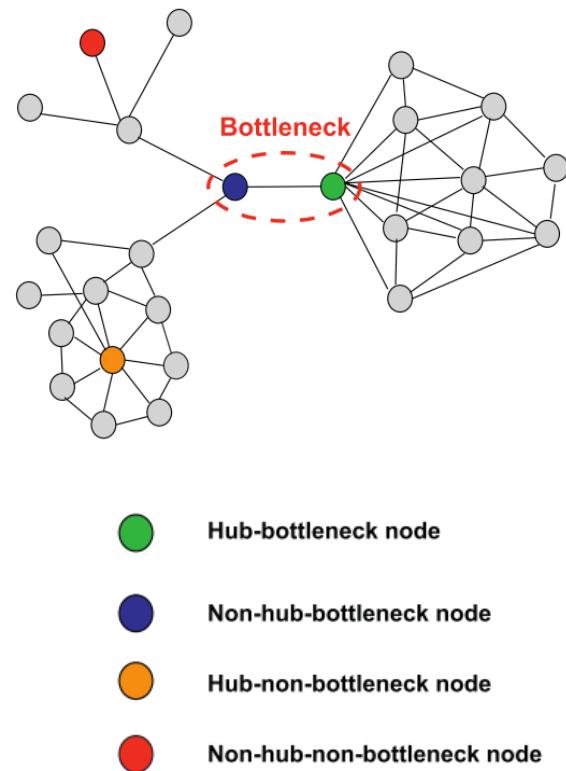
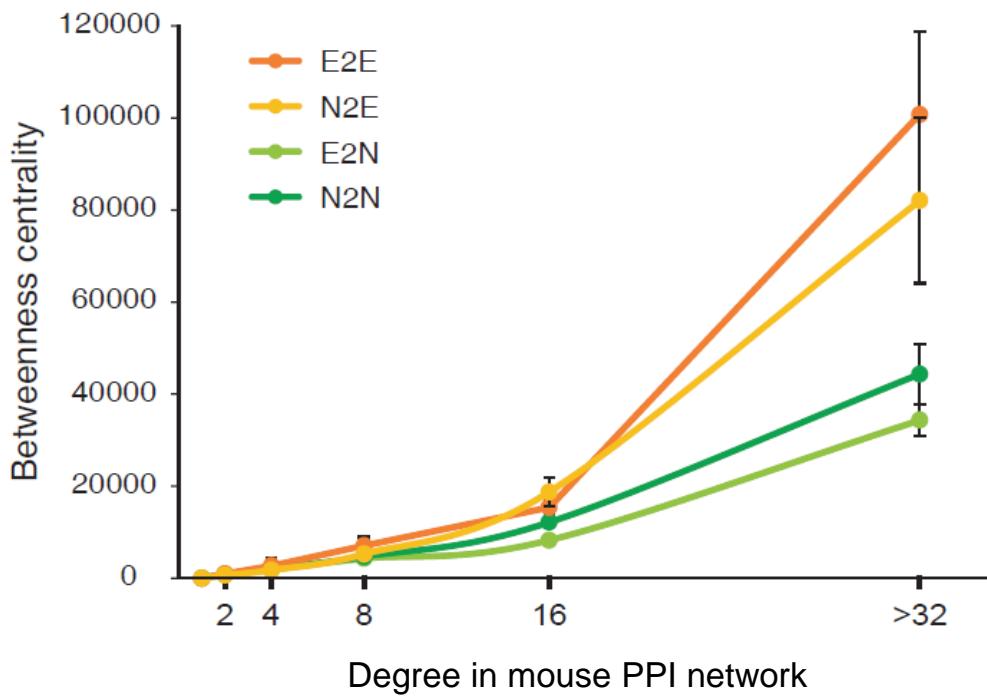


# N2E genes integrated into vital pathways via interaction rewiring



Old genes became essential by participating into vital pathways.

N2E genes often bridge functional modules and control information flow in the PPI network.



Haiyan Yu, Mark Gerstein, Plos CB 2007

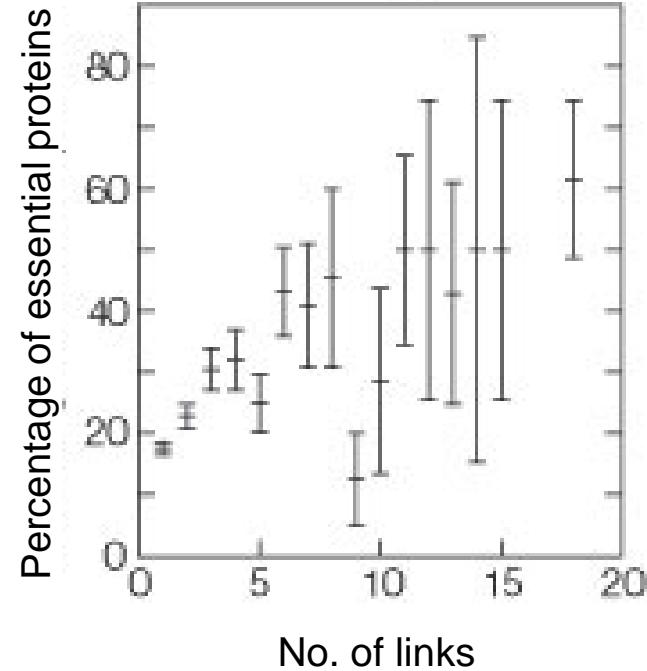
# controversy over the centrality-lethality rule

## Centrality-lethality rule

- The most highly connected proteins in the cell are the most important for its survival.
- But the weak correlation has been a problem.



red, lethal; green, non-lethal;  
orange, slow growth; yellow, unknown



H. Jeong, S. P. Mason, A.-L. Barabási and Z. N. Oltvai  
Nature 2001

The C-L rule dramatically improved for the genes keeping their essentiality both in yeast and mouse

