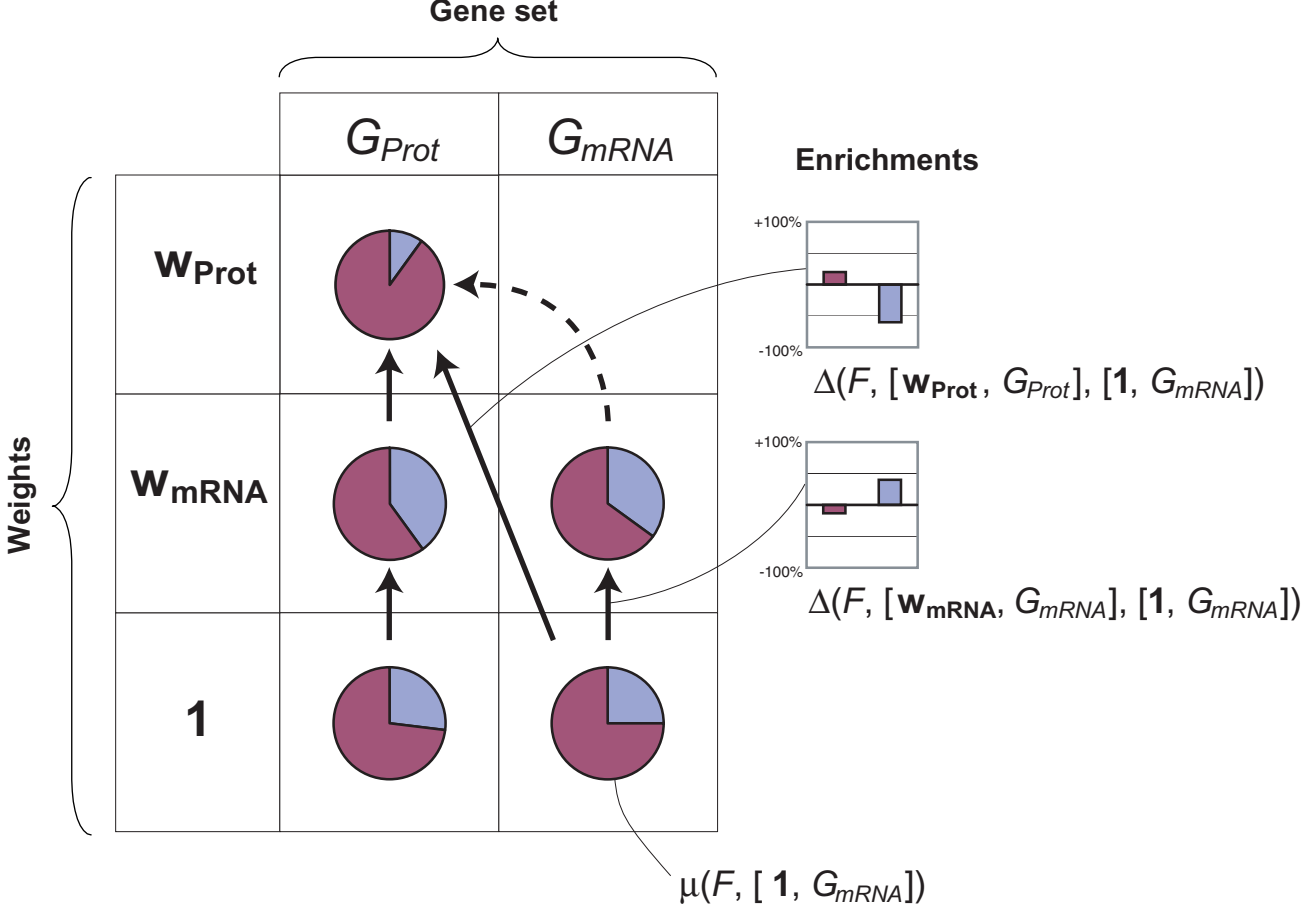
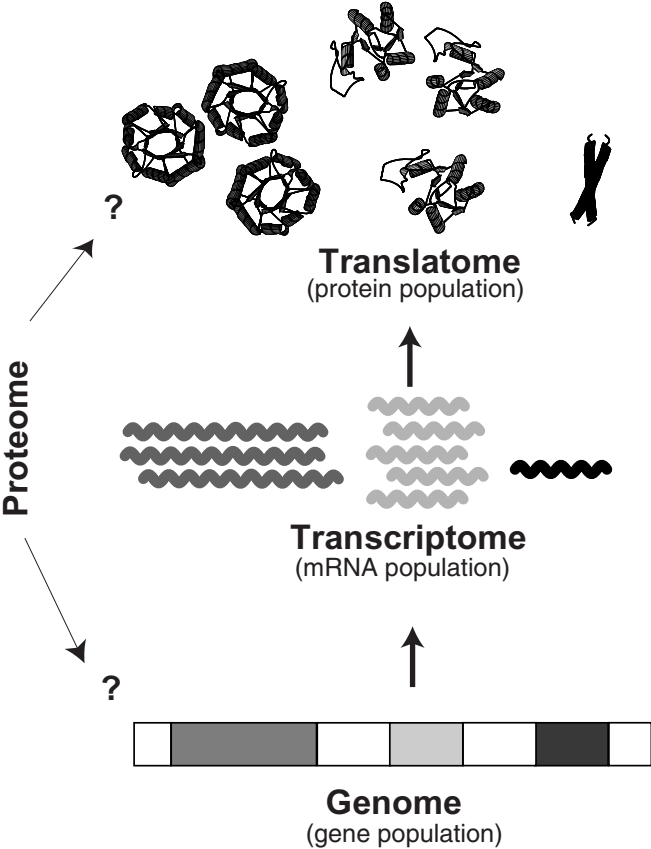
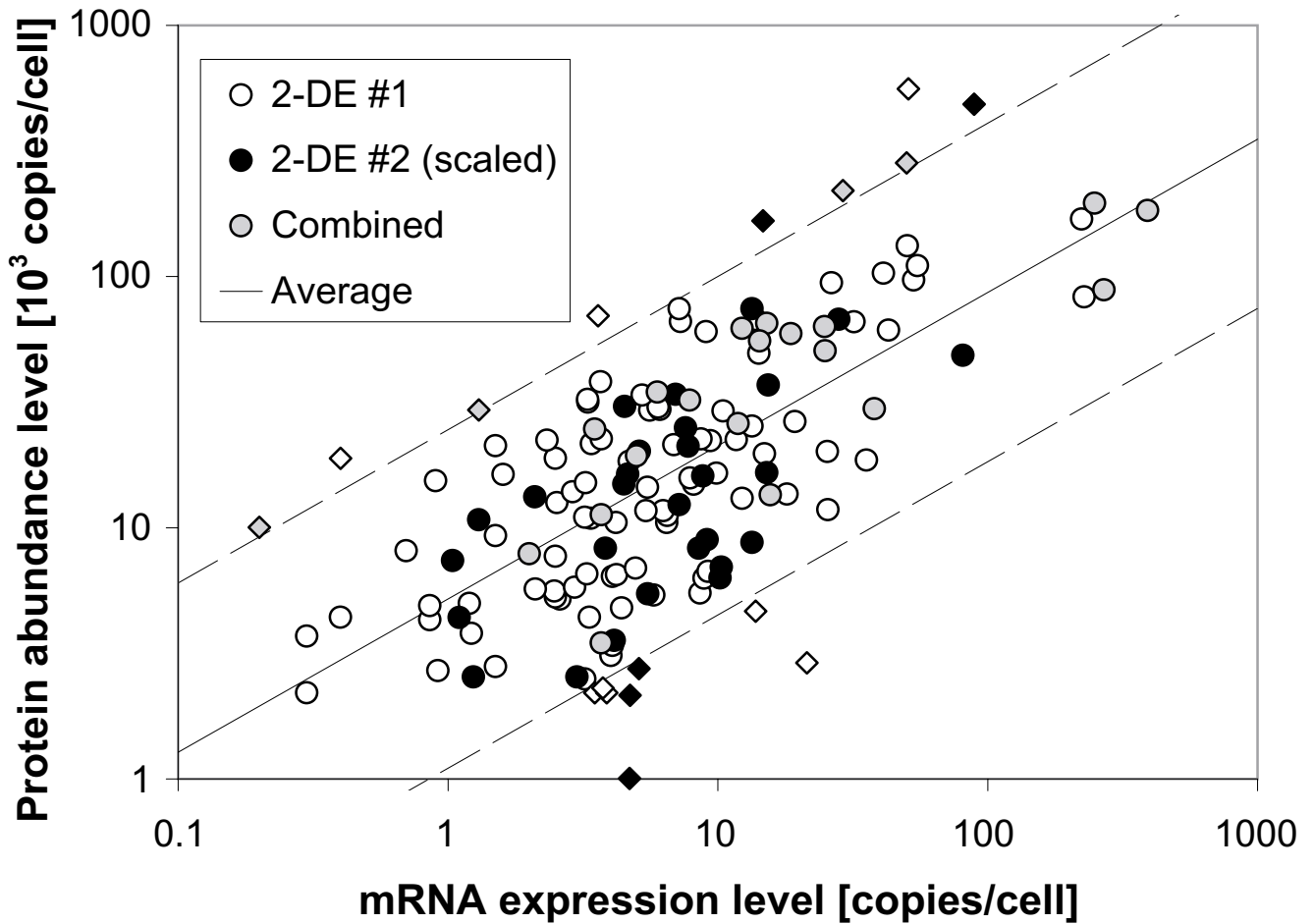


Greenbaum et al - Figure 1



# Greenbaum et al - Figure 2

a

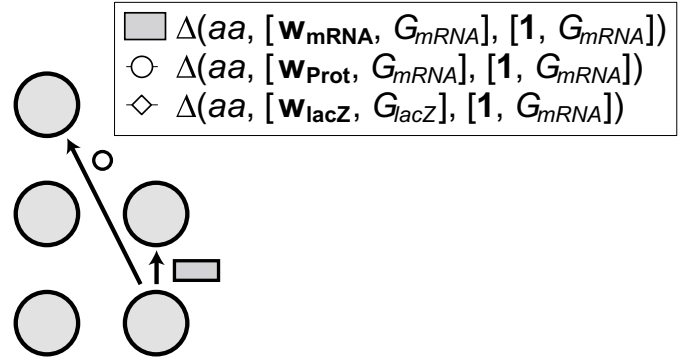
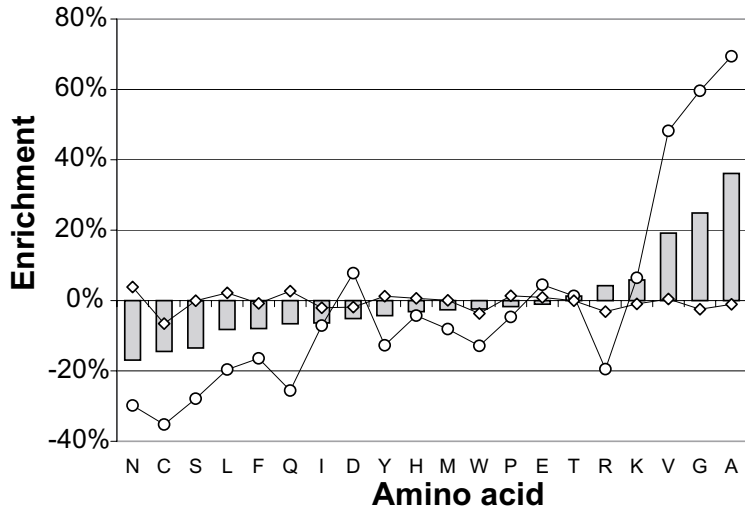


b

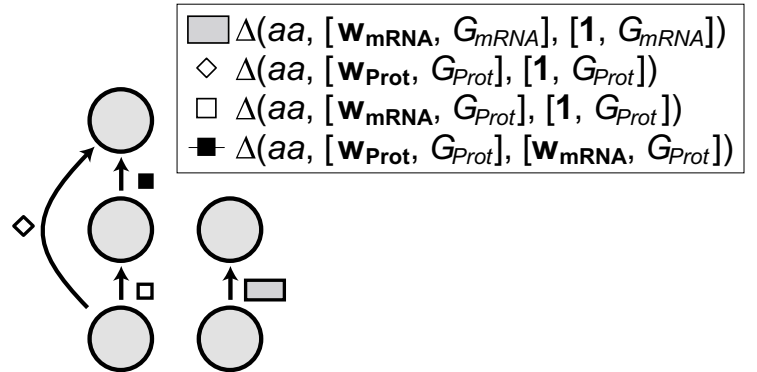
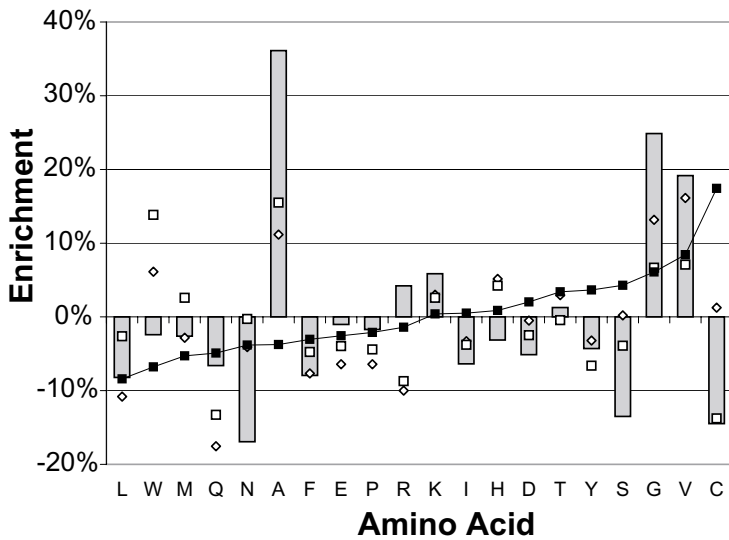
	ORF	FUNCTION	MIPS
above trendline	YBR118W	translation elongation factor eEF1 alpha-A chain	5,30
	YER065C	Isocitrate Lyase	1,2, 30
	YMR303C	Alcohol dehydrogenase II	1, 2, 30
	YOL086C	Alcohol dehydrogenase I	1, 2, 30
	YJR009C	Glyceraldehyde-3-phosphate dehydrogenase 2	1, 2, 30
	YGR192C	Glyceraldehyde-3-phosphate dehydrogenase 3	1, 2, 30
	YJR104C	Copper-zinc superoxide dismutase	11,30
	YML054C	lactate dehydrogenase cytochrome b2	1,2,30
	YJL052W	glyceraldehyde-3-phosphate dehydrogenase 1	1,2,30
below trendline	YKR059W	Translation initiation factor	5,30
	YML008C	S-adenosyl-methionine delta-24-sterol-c-methyltransferase	1,30
	YFL022C	Phenylalanine-- tRNA Ligase beta chain	5,30
	YJL008C	Component of chaperonin-containing T-complex	6,30
	YPL160W	leucine--tRNA ligase	5,30
	YOR361C	translation initiation factor eIF3 subunit	3,5,30
	YCL030C	phosphoribosyl-AMP cyclohydrolase/phosphoribosyl-ATP pyrophosphatase/histidinol dehydrogenase	1
	YNL209W	heat shock protein of HSP70 family	5,30

# Greenbaum et al - Figure 3

**a**

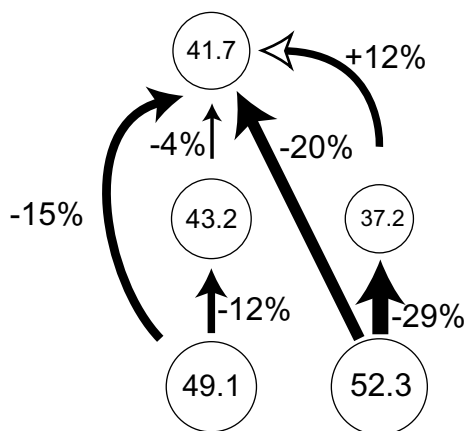


**b**

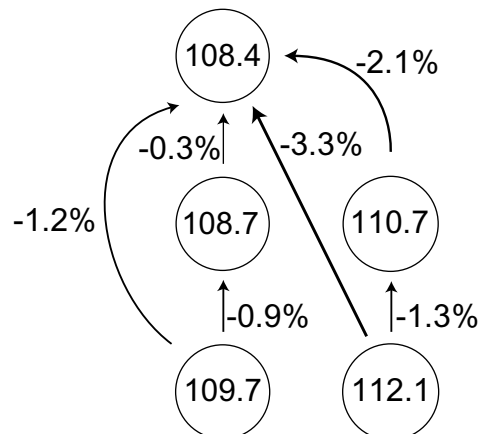


**c**

Molecular weight per ORF [kDa]

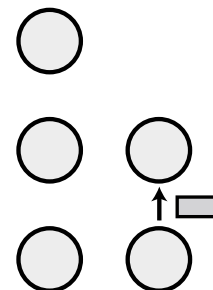
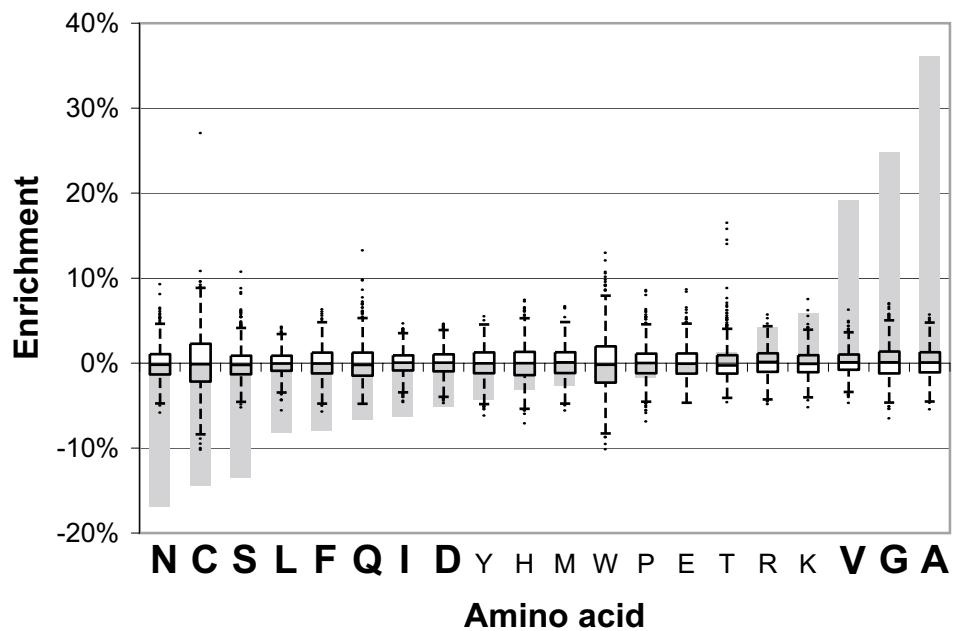


Molecular weight per amino acid [Da]



# Greenbaum et al - Figure 3

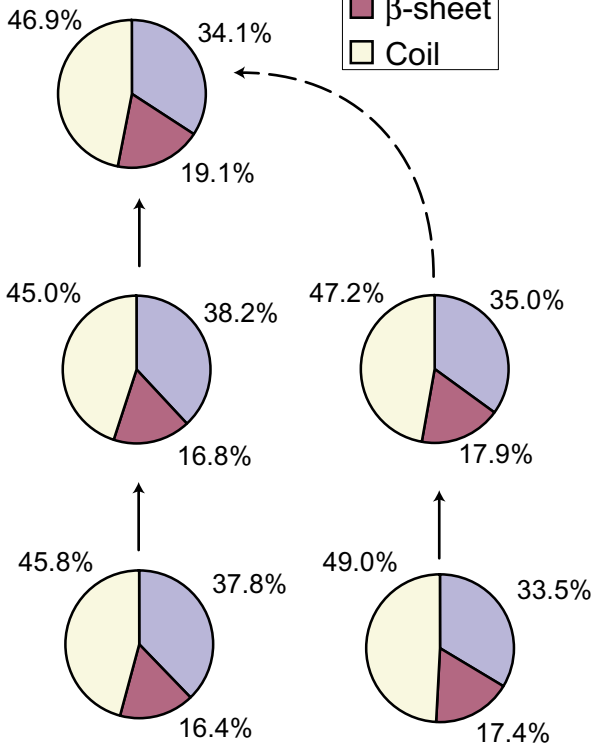
d



# Greenbaum et al - Figure 4

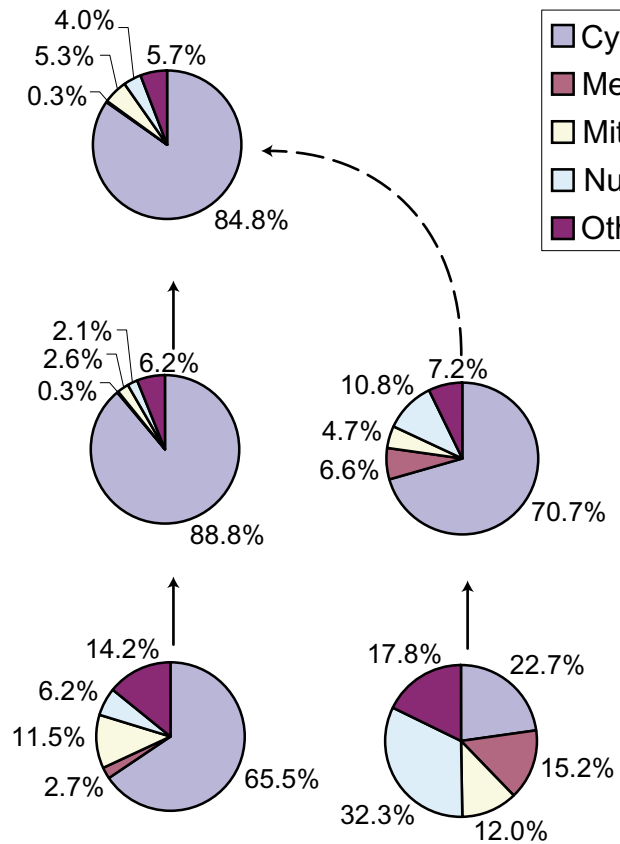
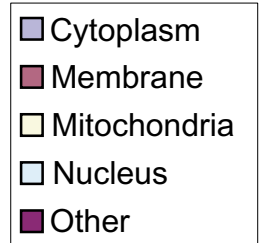
**a**

## Secondary structure



**b**

## Localization



**c**

## Function

