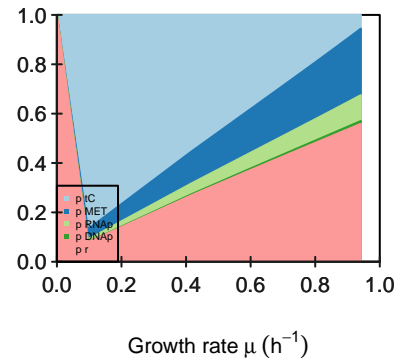
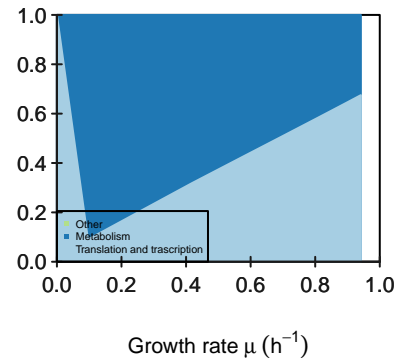


Proteome composition



Proteome sectors



keep_ribosome_kcat FALSE
keep_transport_kcat FALSE
maintenance_fun constant

M

	tC	MET	RNAp	DNAP	r
C	1	-1	0	0	0
AA	0	0.5	0	0	-0.9
NT	0	0.3	-1	-1	0
ATP	0	0.2	0	0	-0.1
rRNA	0	0	1	0	0
DNA	0	0	0	1	0
p	0	0	0	0	1

K

	tC	MET	RNAp	DNAP	r
x_C	0.1	0	0	0	0
C	0	7	0	0	0
AA	0	0	0	0	3
NT	0	0	1	1	0
ATP	0	0	0	0	12
rRNA	0	0	0	0	0
DNA	0	0	0	0	0
p	0	0	0	0	0

KA

	tC	MET	RNAp	DNAp	r
x_C	0	0	0	0	0
C	0	0	0	0	0
AA	0	0	0	0	0
NT	0	0	0	0	0
ATP	0	0	0	0	0
rRNA	0	0	0	0	60
DNA	0	0	4	4	0
p	0	0	0	0	0

kcat

	tC	MET	RNAp	DNAp	r
kcatf	45.3	10	8.4	11.2	5.5
kcatb	5	1	1	1	1

Keq

	[1,]	[,1]	[,2]	[,3]	[,4]	[,5]
[1,]	90.6	1.42857142857143	8.4	11.2	0.152777777777778	

phi input

	[,1]	[,2]	[,3]	[,4]	[,5]
[1,]	0.065	0.352	0.12	0.003	0.46

average saturation input

3

minimal phi constraint

	[1,]	[,1]	[,2]	[,3]	[,4]	[,5]
[1,]		0	0	0	0	0

minimal f constraint

	[1,]	[,1]	[,2]	[,3]	[,4]	[,5]
	0	0	0	0	0	