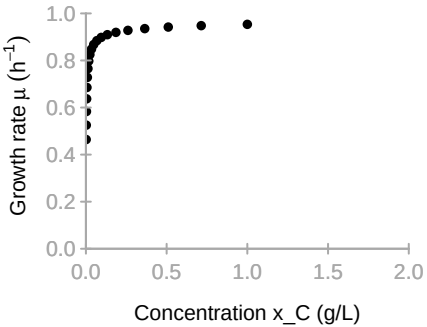
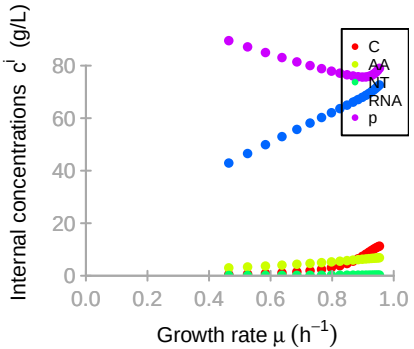
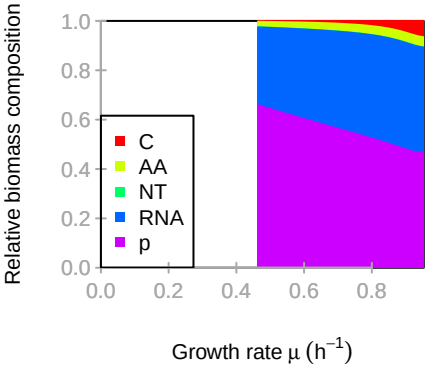
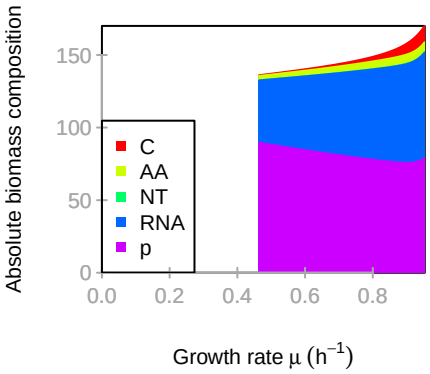
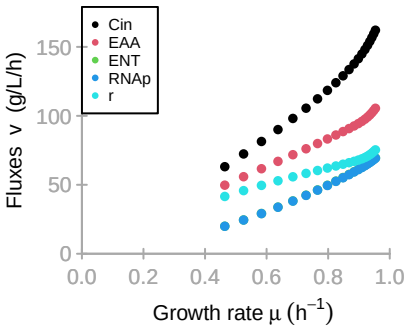
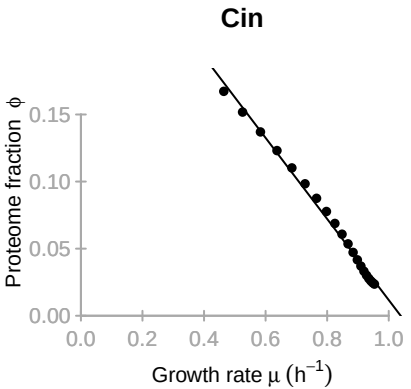

Growth rate μ (h−1)
1.0
0.8
0.6
0.4
0.2
0.0
0.0
0.5
1.0
1.5
2.0
Concentration x_C (g/L)

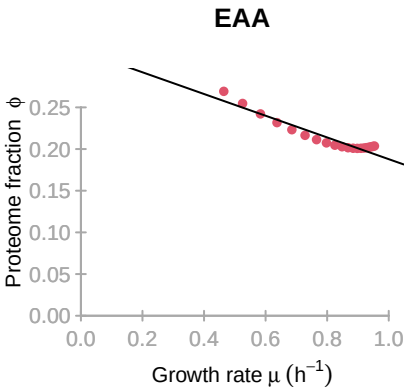

Internal concentrations ci (g/L)
80
60
40
20
0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)
C
AA
NT
RNA
p

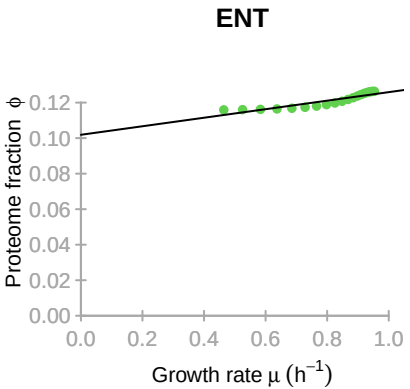

Relative biomass composition
1.0
0.8
0.6
0.4
0.2
0.0
0.0
0.2
0.4
0.6
0.8
Growth rate μ (h−1)
C
AA
NT
RNA
p

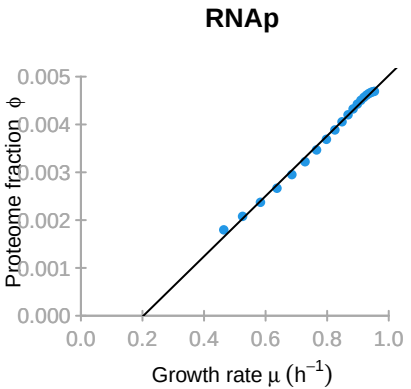

Absolute biomass composition
150
100
50
0
0.0
0.2
0.4
0.6
0.8
Growth rate μ (h−1)
C
AA
NT
RNA
p

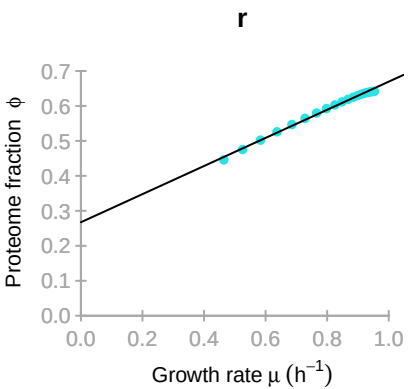

Fluxes v (g/L/h)
150
100
50
0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)
Cin
EAA
ENT
RNAp
r

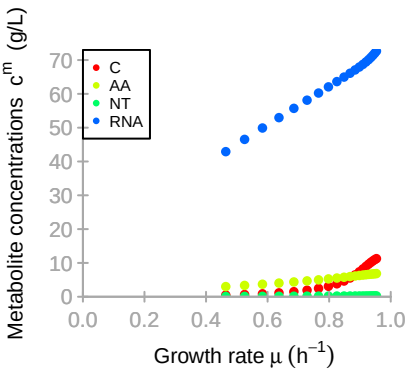

Cin
Proteome fraction ϕ
0.15
0.10
0.05
0.00
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)

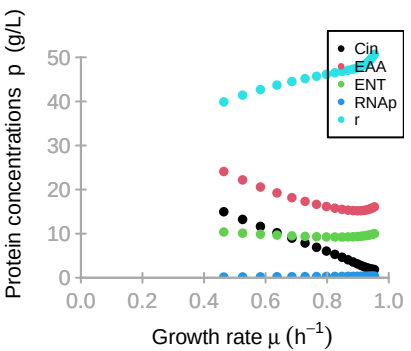

EAA
Proteome fraction ϕ
0.25
0.20
0.15
0.10
0.05
0.00
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)

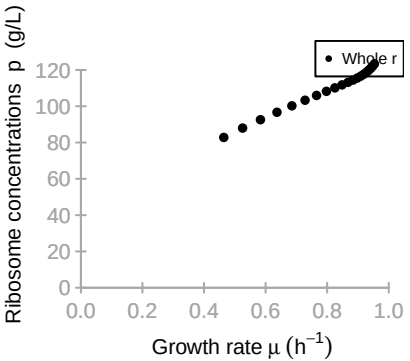

ENT
Proteome fraction ϕ
0.12
0.10
0.08
0.06
0.04
0.02
0.00
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)

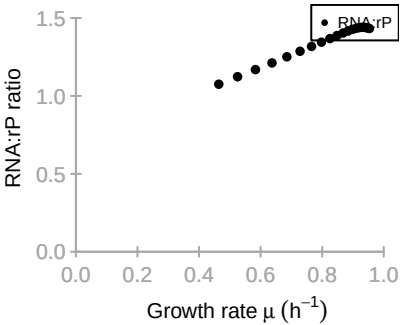

RNAp
Proteome fraction ϕ
0.005
0.004
0.003
0.002
0.001
0.000
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)

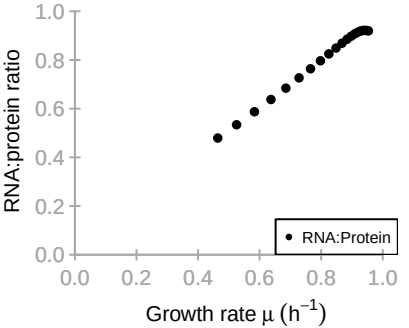

r
Proteome fraction ϕ
0.7
0.6
0.5
0.4
0.3
0.2
0.1
0.0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)

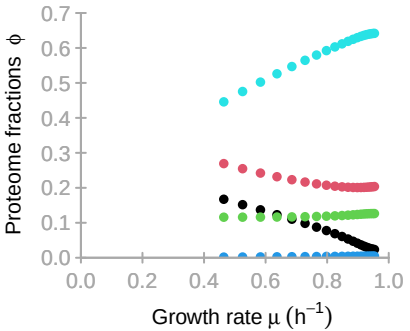

Metabolite concentrations cm (g/L)
C
AA
NT
RNA
70
60
50
40
30
20
10
0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)

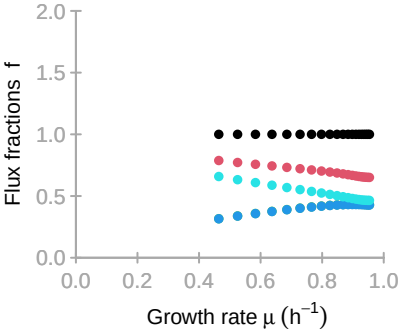

Protein concentrations p (g/L)
Cin
EAA
ENT
RNAp
r
50
40
30
20
10
0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)

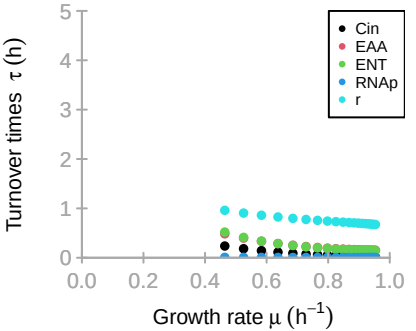

Ribosome concentrations p (g/L)
Whole r
120
100
80
60
40
20
0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)

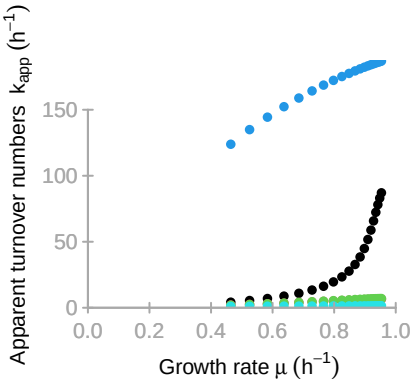

RNA:rP ratio
RNA:rP
1.5
1.0
0.5
0.0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)


RNA:protein ratio
RNA:Protein
1.0
0.8
0.6
0.4
0.2
0.0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)


Proteome fractions ϕ
0.7
0.6
0.5
0.4
0.3
0.2
0.1
0.0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)


Flux fractions f
2.0
1.5
1.0
0.5
0.0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)


Turnover times τ (h)
Cin
EAA
ENT
RNAp
r
5
4
3
2
1
0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h−1)


Apparent turnover numbers k_{app} (h^{-1})
150
100
50
0
0.0
0.2
0.4
0.6
0.8
1.0
Growth rate μ (h^{-1})

M

	Cin	EAA	ENT	RNAp	r
C	1	−1	−0.66	0	0
AA	0	1	−0.34	0	−1
NT	0	0	1	−1	0
RNA	0	0	0	1	0
p	0	0	0	0	1

K

	Cin	EAA	ENT	RNAp	r
[1,]	0.01	0	0	0	0
[2,]	1	1	1	0	0
[3,]	0	20	2	0	0.1
[4,]	0	0	20	0.1	0
[5,]	0	0	0	0	0
[6,]	0	0	0	0	0

KA

	Cin	EAA	ENT	RNAp	r
[1,]	0	0	0	0	0
[2,]	0	0	0	0	0
[3,]	0	0	0	0	0
[4,]	0	0	0	0	0
[5,]	0	0	0	0	100
[6,]	0	0	0	0	0

kcat

	[,1]	[,2]	[,3]	[,4]	[,5]
kcatf	100	7.3846	9.8004	260.2688	3.5828
kcatb	20	1.47692	1.96008	0	0

Keq

	[,1]	[,2]	[,3]	[,4]	[,5]
[1,]	500	100	50	Inf	Inf