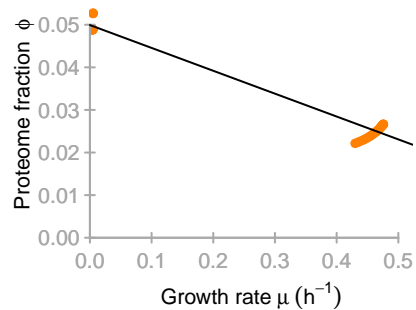
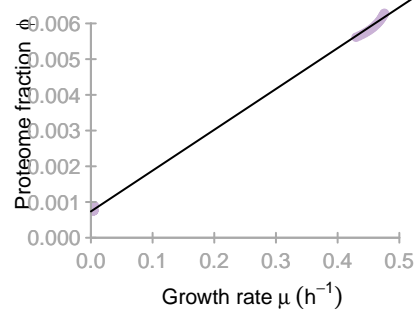
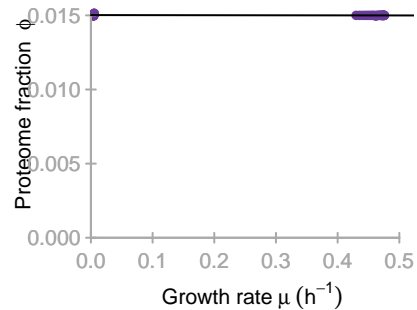
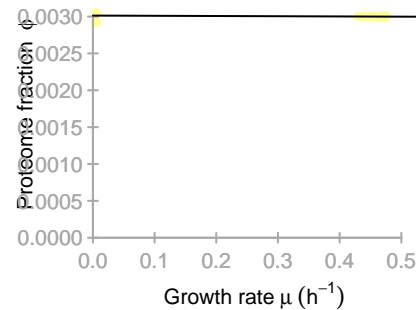
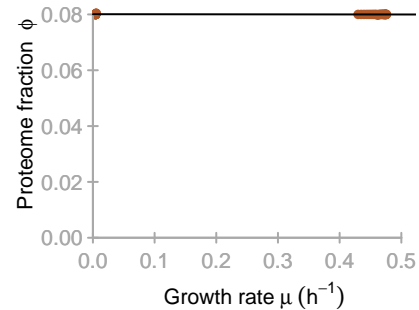
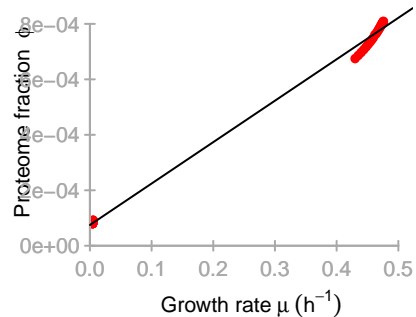
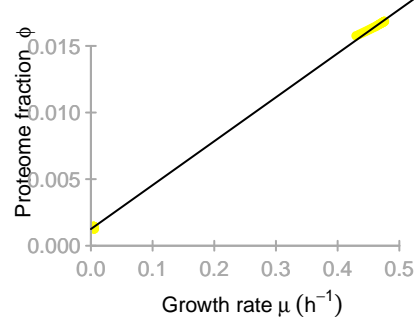
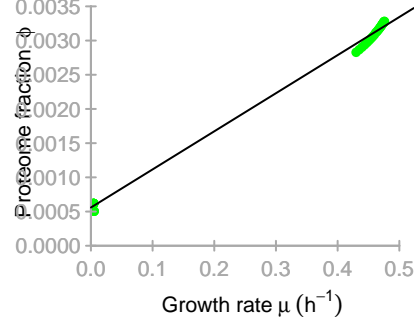
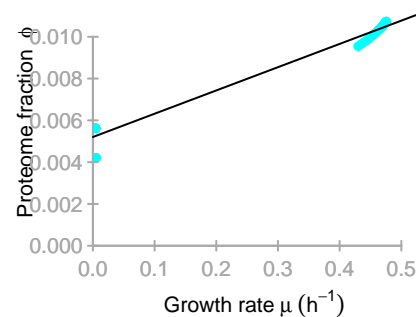
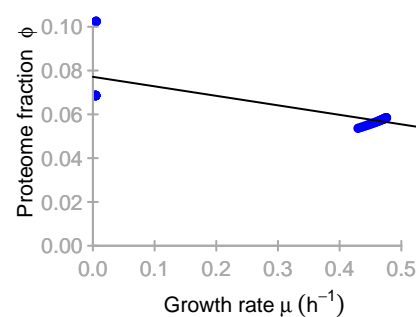
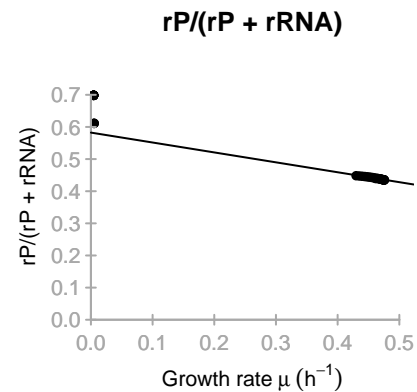
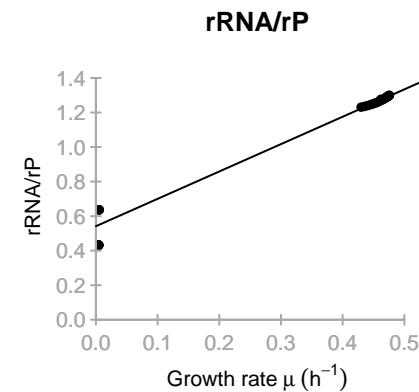
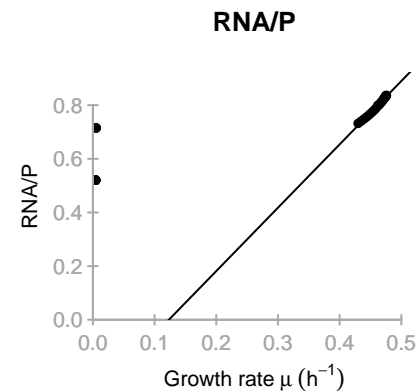
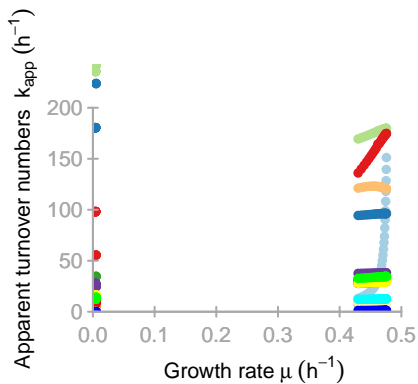
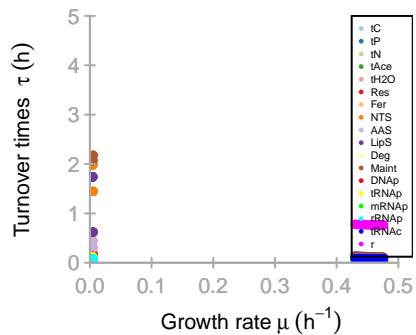
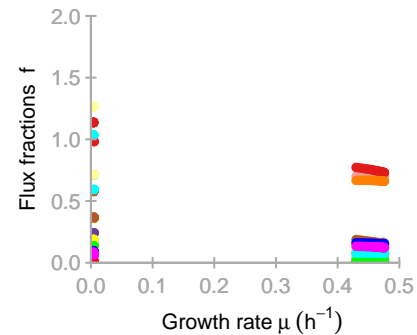
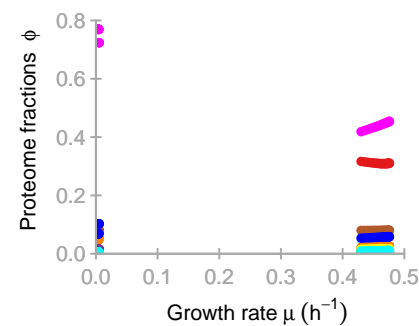
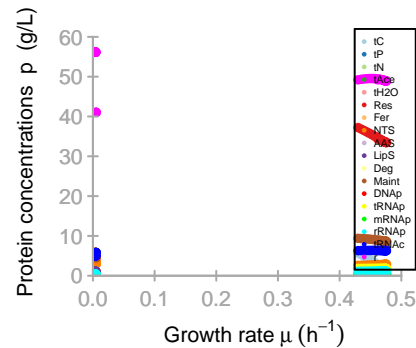
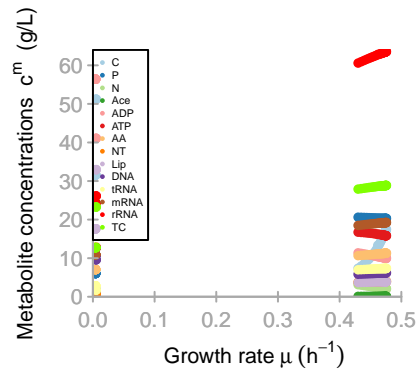
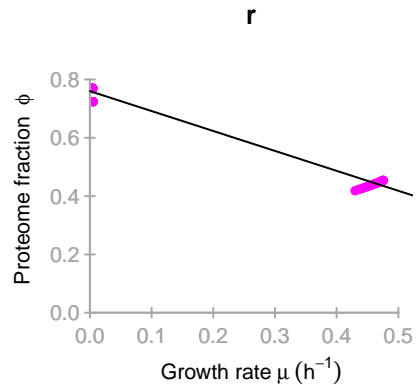
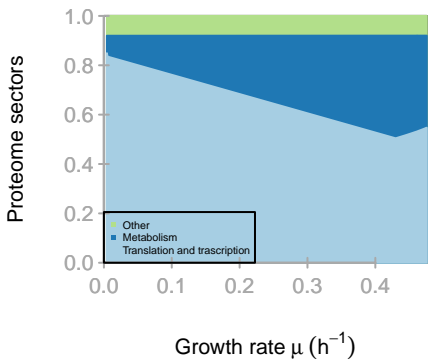
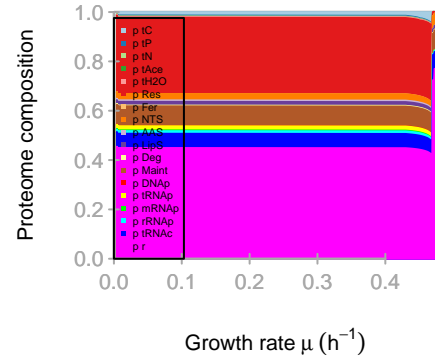
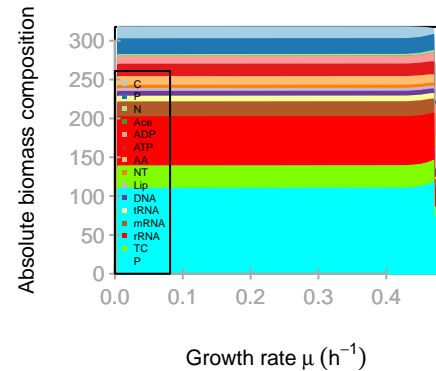
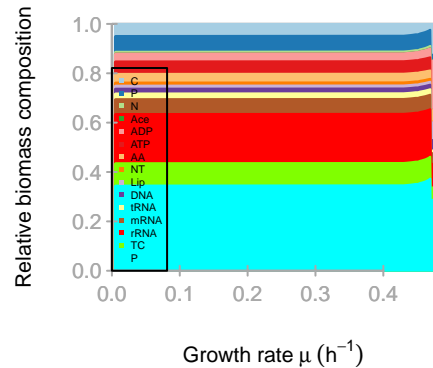
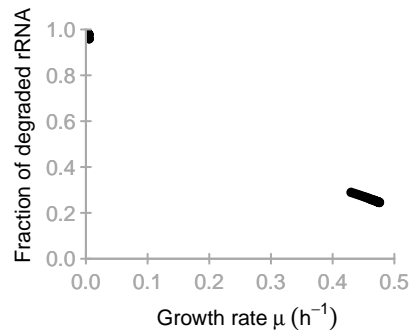
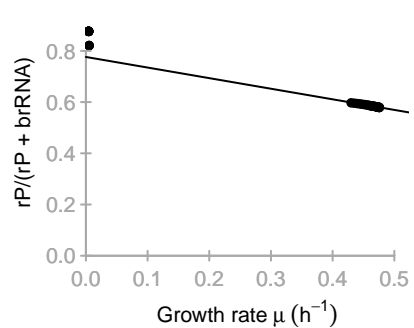


NTS**AAS****LipS****Deg****Maint****DNAP****tRNAp****mRNAp****rRNAp****tRNAc**



Protein mass fraction in ribosome



M

[illegible]

K

[illegible]

KA[illegible]

kcat

	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]	[,8]	[,9]	[,10]	[,11]	[,12]	[,13]	[,14]	[,15]	[,16]	[,17]	[,18]
kcatf	800	500	500	500	1000	20	250	300	80	40	50	10	200	136	96	136	50	4.55
kcatb	50	6	6	6	1	2	2	3	8	4	0	0	0	0	0	0	0	0

Keq

[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]	[,8]	[,9]	[,10]	[,11]	[,12]	[,13]	[,14]	[,15]	[,16]	[,17]
160	833.3333333333333	833.3333333333333	833.3333333333333	10000	0.37037037037037037	41.66666666666667	1666.666666666667	16.66666666666667	100	Inf	Inf	Inf	Inf	Inf	Inf	Inf

minimal phi constraint

[1,]	[,1] 0	[,2] 0	[,3] 0	[,4] 0	[,5] 0	[,6] 0	[,7] 0	[,8] 0	[,9] 0	[,10] 0.015	[,11] 0.003	[,12] 0.08	[,13] 0	[,14] 0	[,15] 0	[,16] 0	[,17] 0	[,18] 0
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minimal f constraint

[1,]	[1] 0	[2] 0	[3] 0	[4] 0	[5] 0.7	[6] 0	[7] 0	[8] 0	[9] 0	[10] 0	[11] 0	[12] 0	[13] 0	[14] 0	[15] 0	[16] 0	[17] 0	[18] 0
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