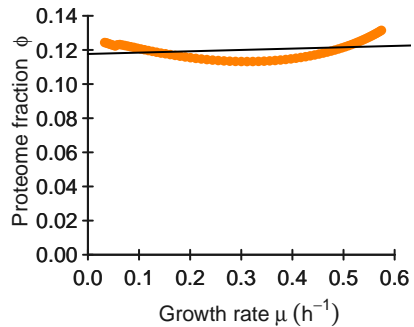
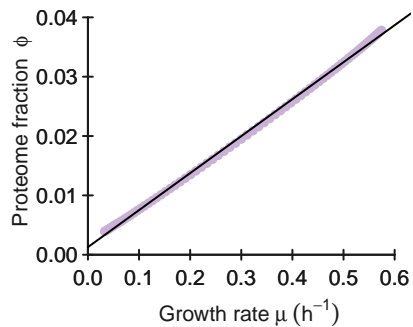
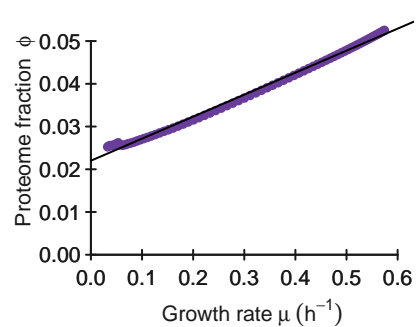
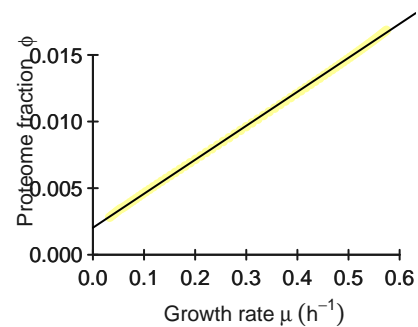
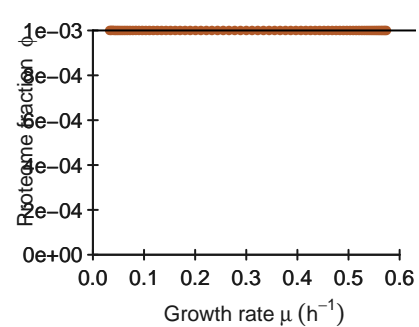
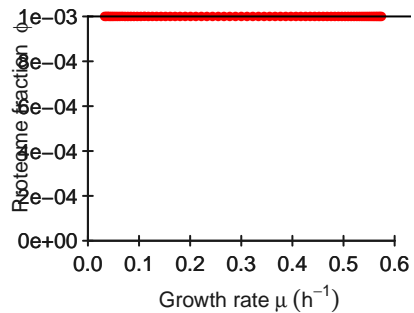
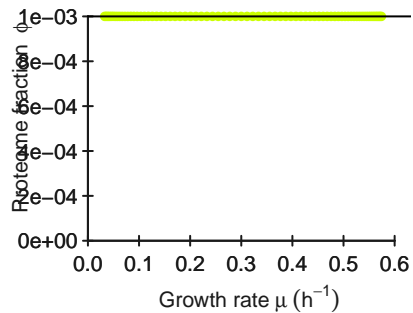
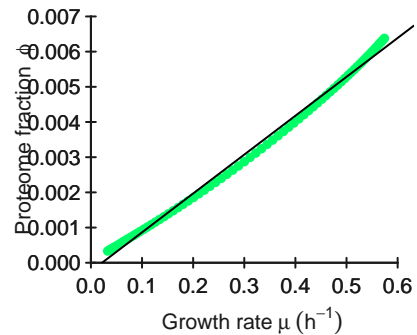
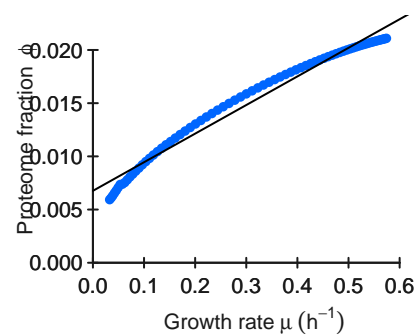
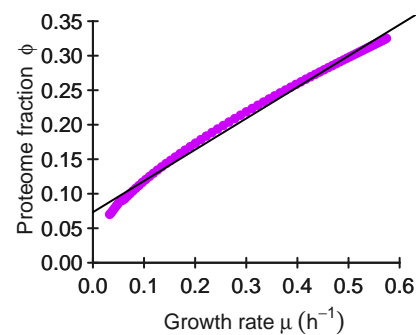
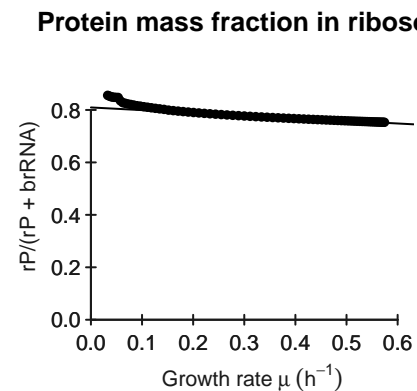
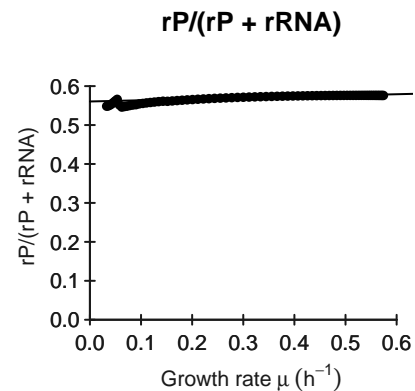
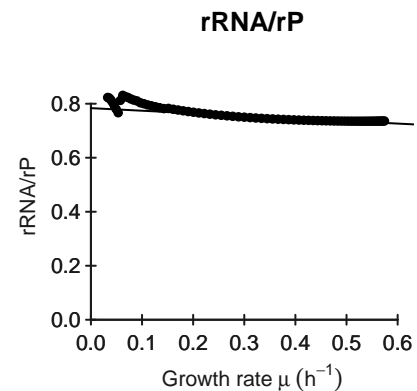
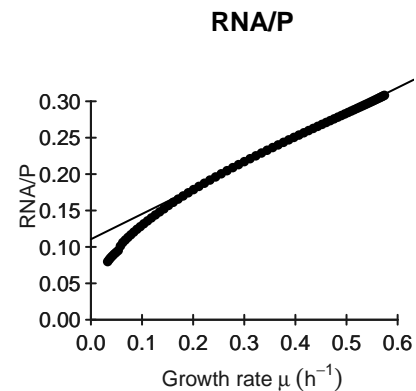
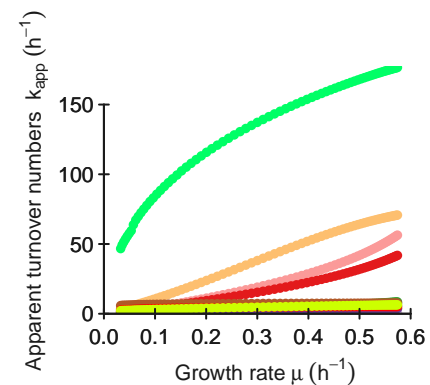
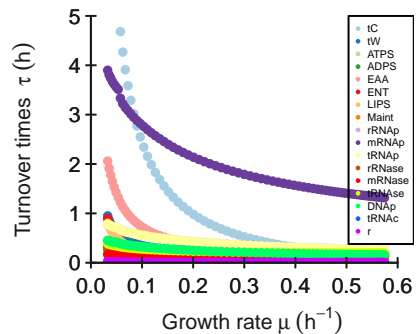
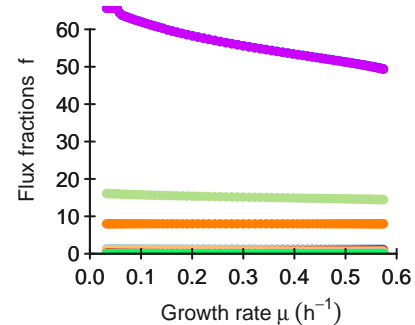
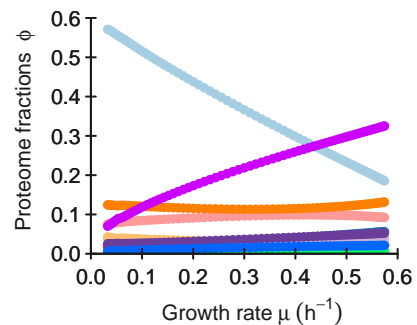
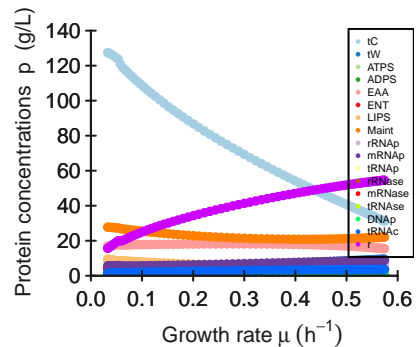
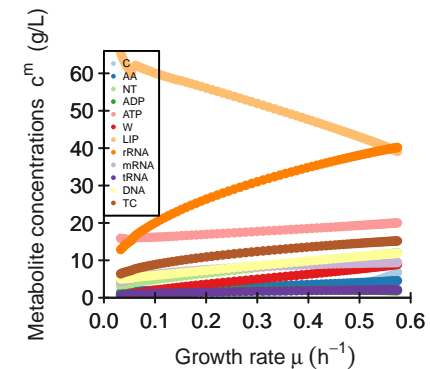
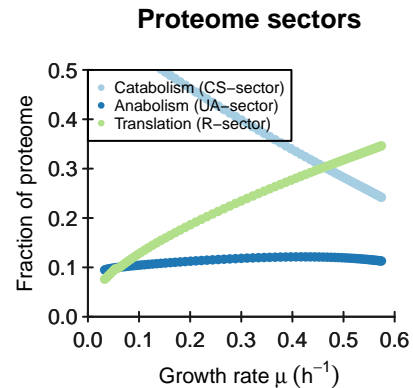
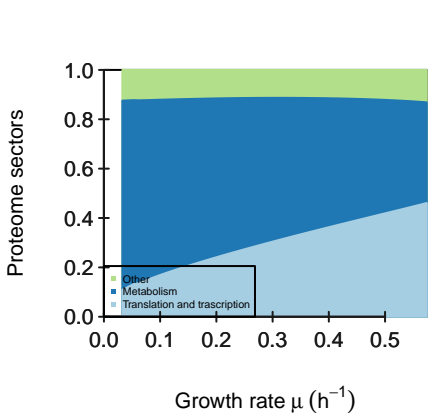
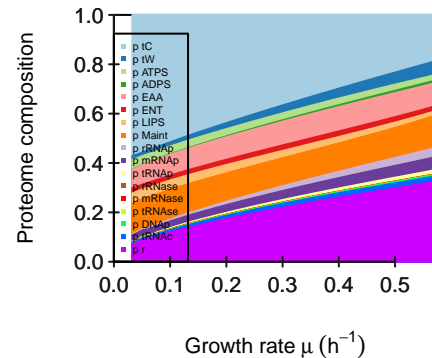
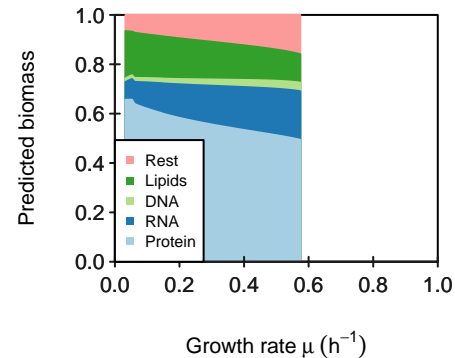
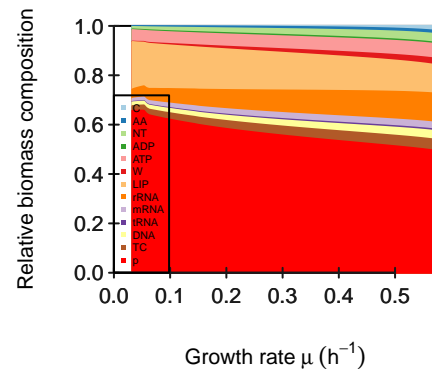
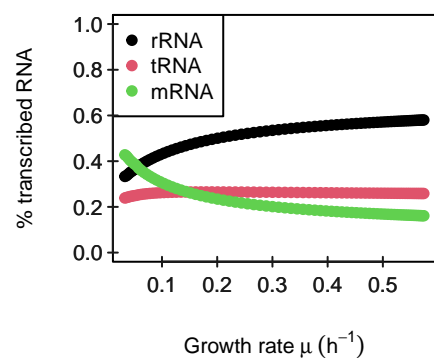
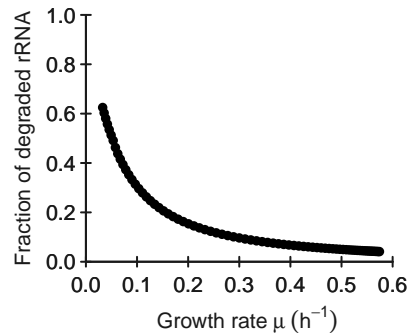


Maint**rRNAp****mRNAp****tRNAp****rRNase****mRNase****tRNase****DNAp****tRNAc****r**





M

[illegible]

K

[illegible]

KA[illegible]

Keq

[1,]	346.6666666666667	[1]	[2]	419.2533333333333	[3]	4.666666666666667	[4]	[5]	72	4.931818181818181	[6]	[7]	91.25	[8]	Inf	[9]	Inf	[10]	Inf	[11]	Inf	[12]	Inf	[13]	Inf	[14]	Inf	[15]	Inf	[16]	Inf	[17]	Inf
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phi input

[1,]	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]	[,8]	[,9]	[,10]	[,11]	[,12]	[,13]	[,14]	[,15]	[,16]	[,17]
	0.089	0.021	0.027	0.005	0.165	0.023	0.031	0.2506	0.0426	0.0213	0.0071	0.002	0.006	4e-04	0.002	0.023	0.284

average saturation input

2

minimal phi constraint

[1,]	[,1] 0	[,2] 0	[,3] 0	[,4] 0	[,5] 0	[,6] 0	[,7] 0	[,8] 0	[,9] 0	[,10] 0	[,11] 0	[,12] 0.001	[,13] 0.001	[,14] 0.001	[,15] 0	[,16] 0	[,17] 0
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minimal f constraint

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