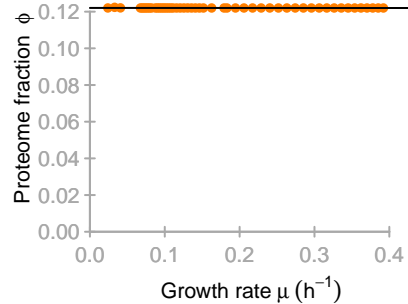
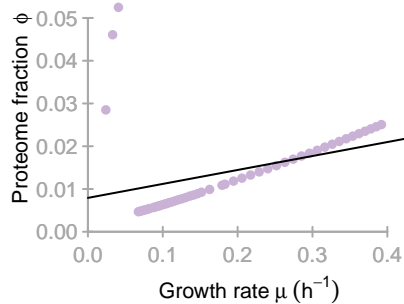
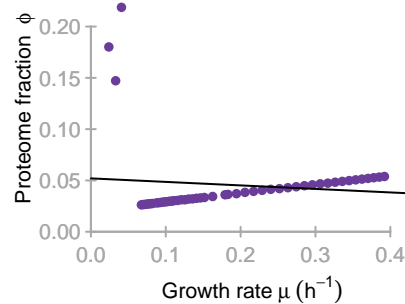
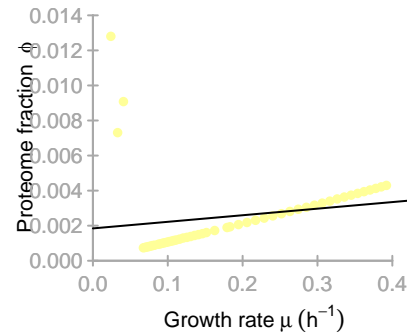
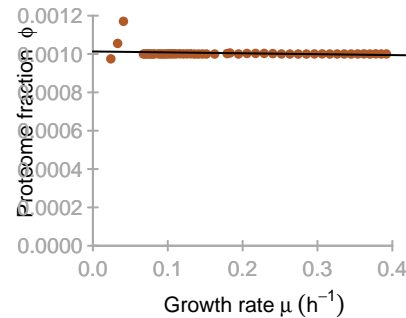
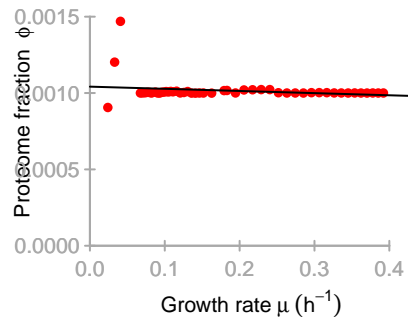
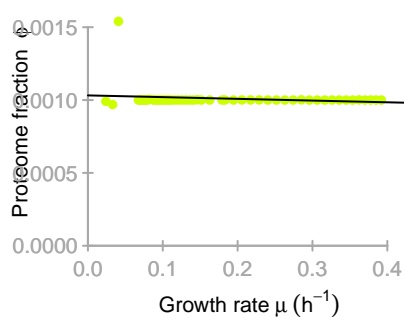
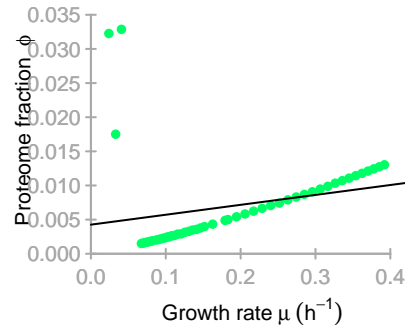
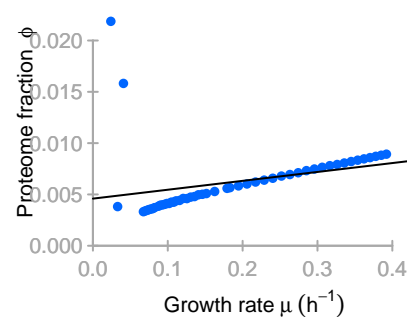
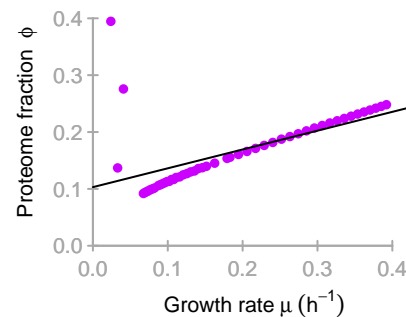
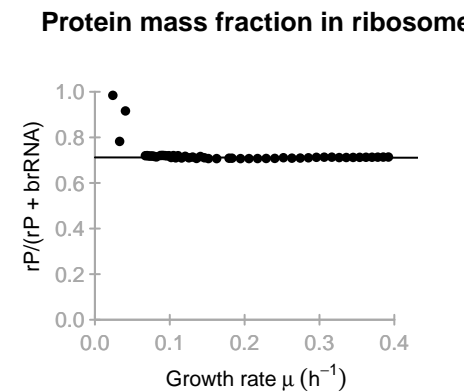
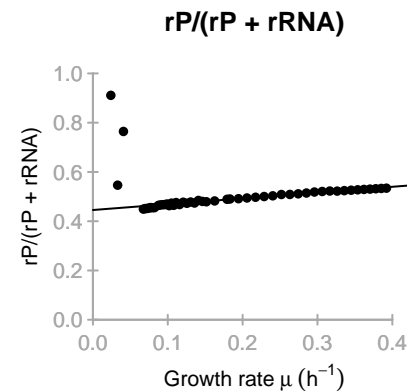
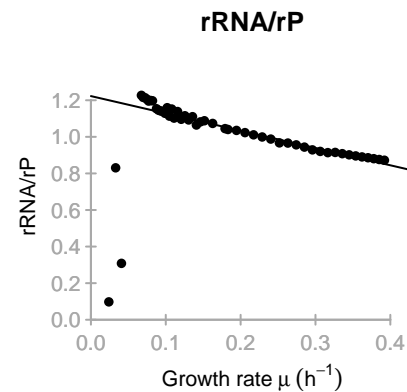
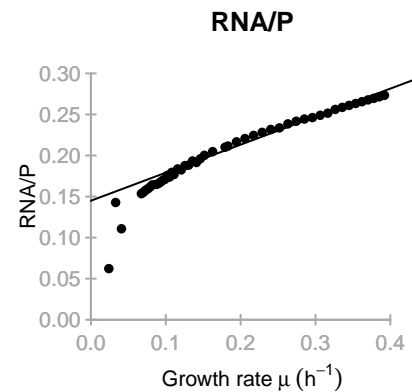
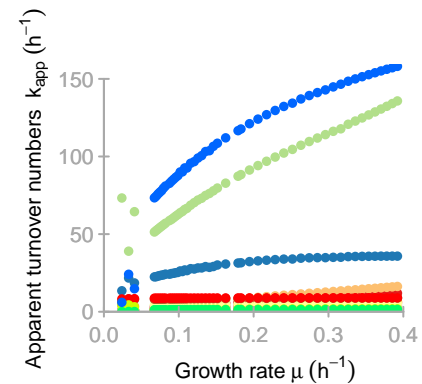
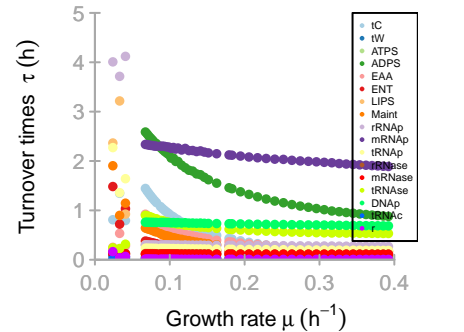
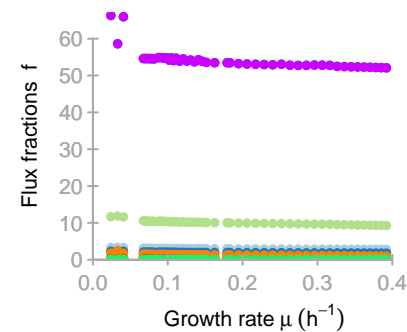
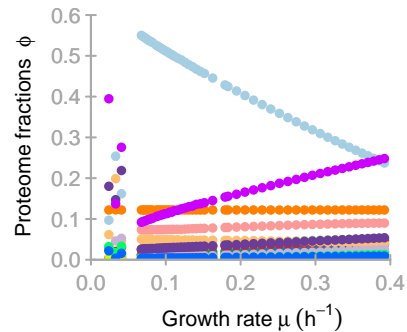
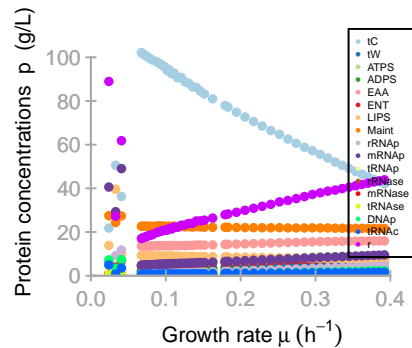
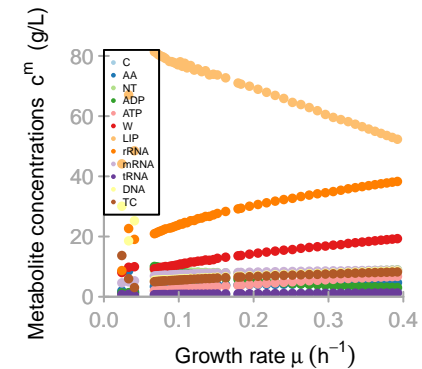
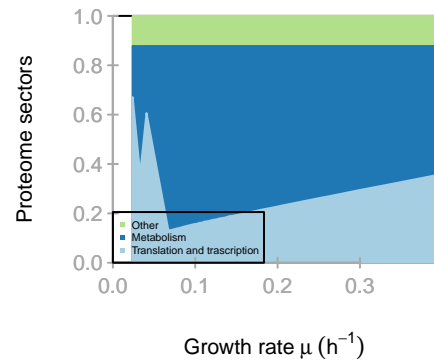
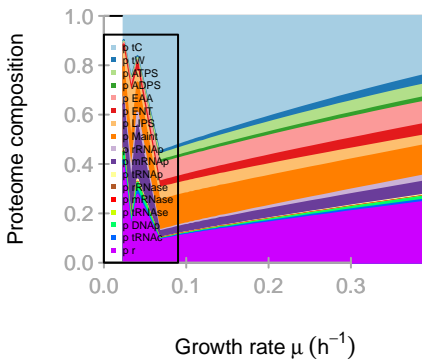
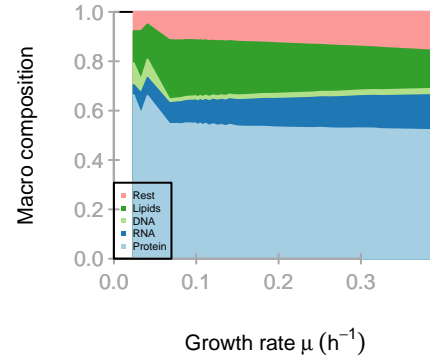
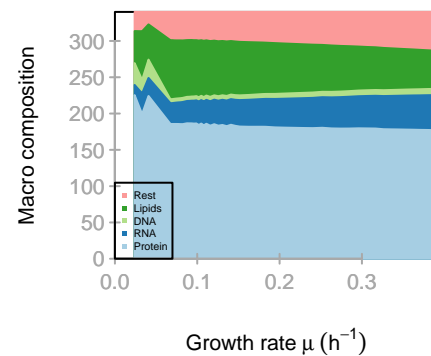
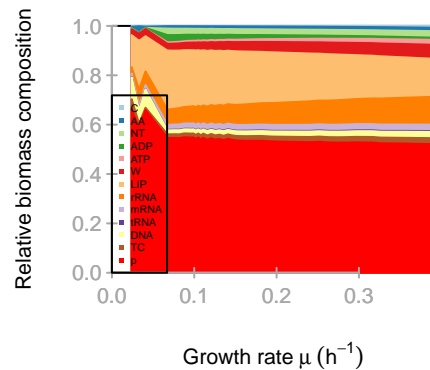
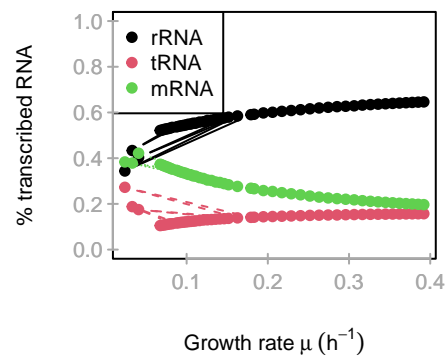
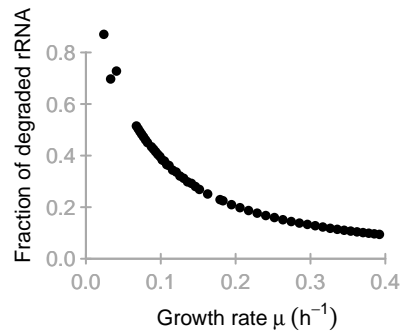
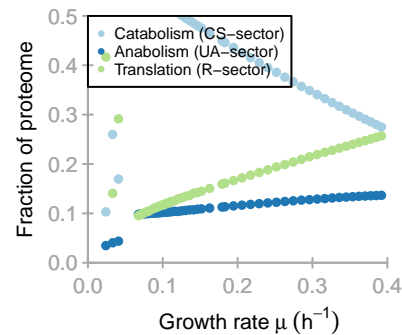


Maint**rRNAp****mRNAp****tRNAp****rRNase****mRNase****tRNase****DNAp****tRNAc****r**





Proteome sectors



M

[illegible]

K

[illegible]

KA[illegible]

Keq

[1,]	[,1] 63	0.357142857142857	[,2] 10.08	[,3] 2	[,4] 8	[,5] 11.5	8.666666666666667	[,6] Inf	[,7] Inf	[,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]
	0.1	0.01	0.1	0.01	0.17	0.04	0.09	0.107	0.01	0.01	0.01	0.001	0.001	0.001	0.02	0.02	0.3

average saturation input

3

minimal phi constraint

[1.]	[1] 0	[2] 0	[3] 0	[4] 0	[5] 0	[6] 0	[7] 0	[8] 0.122	[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]
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 [,2]
 0 0
 [,3]
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 [,4]
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 [,5]
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