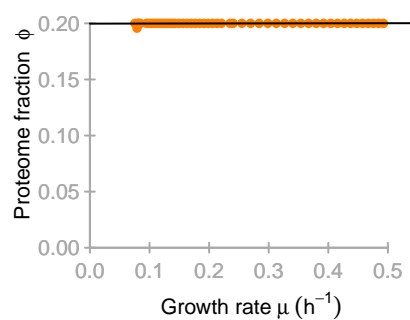
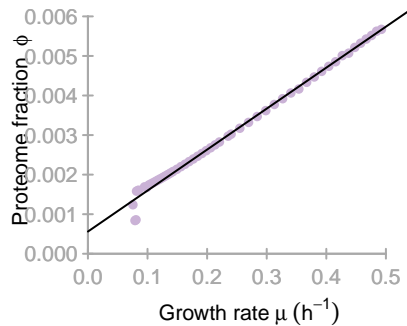
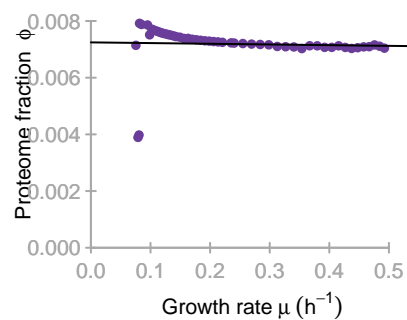
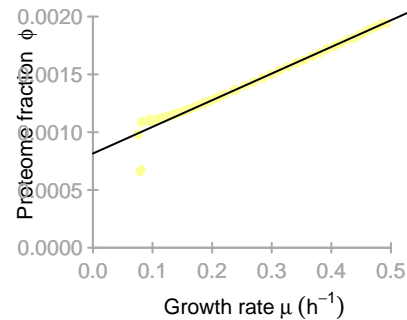
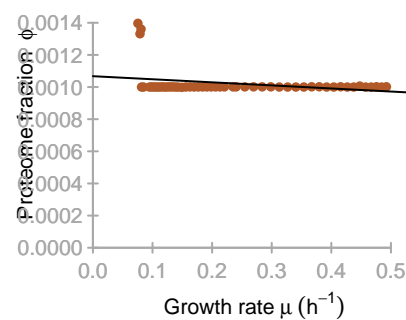
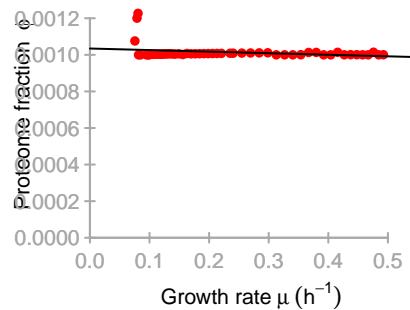
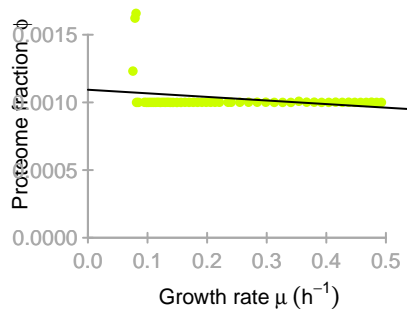
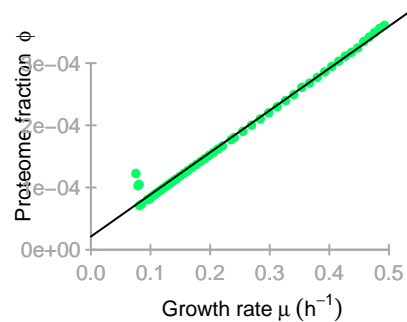
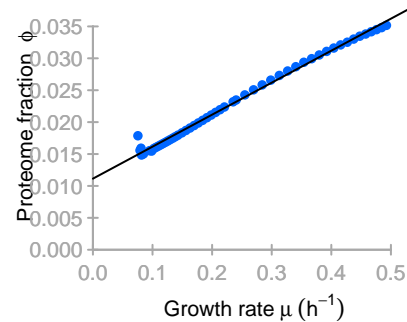
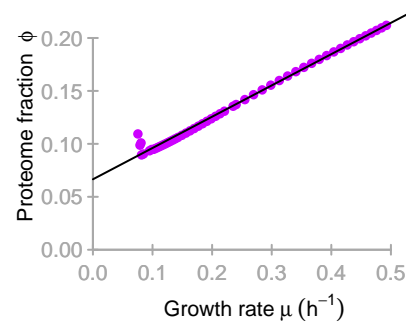
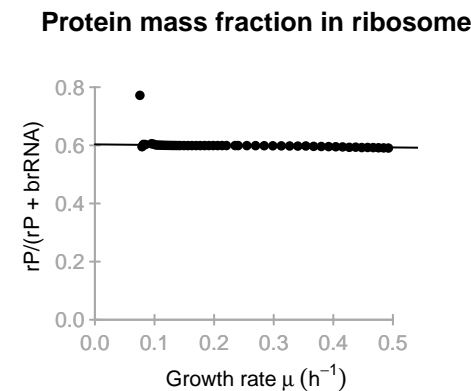
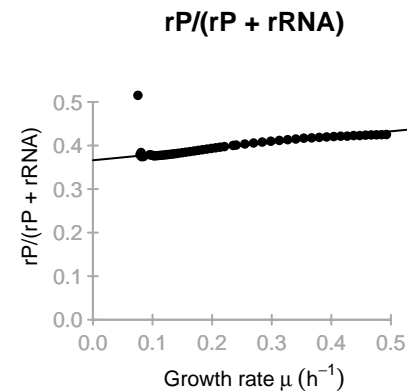
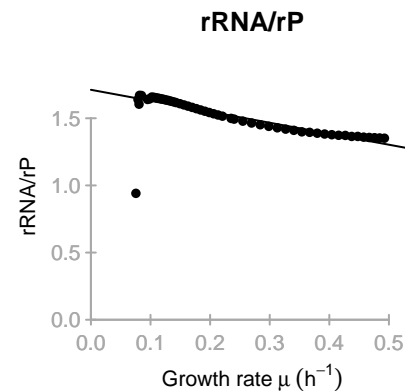
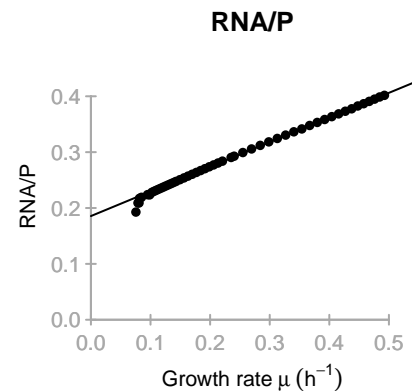
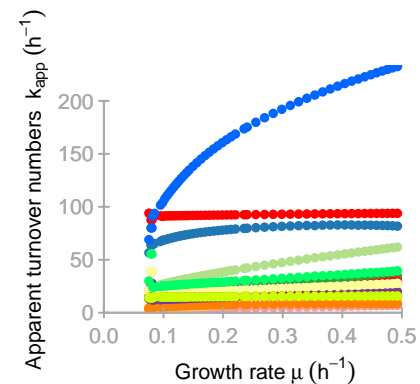
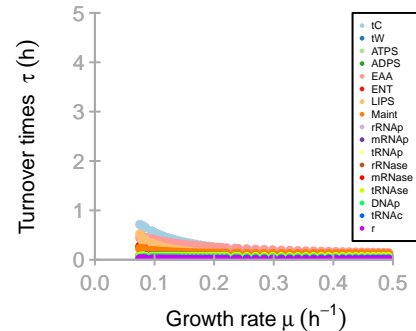
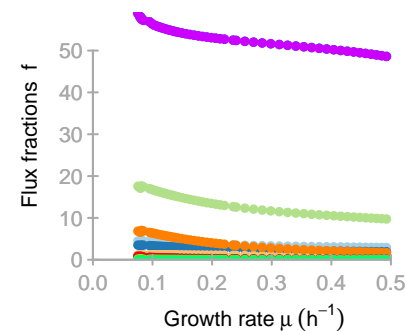
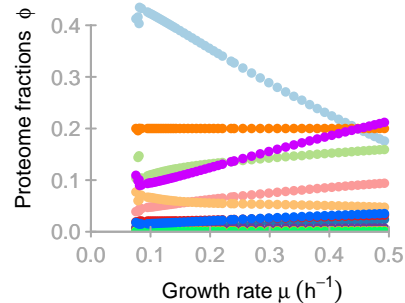
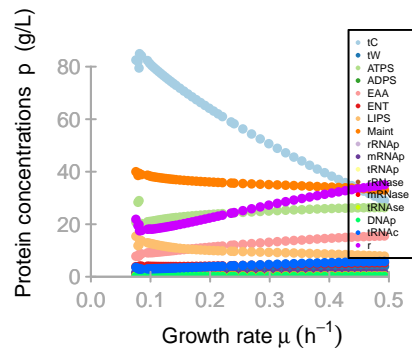
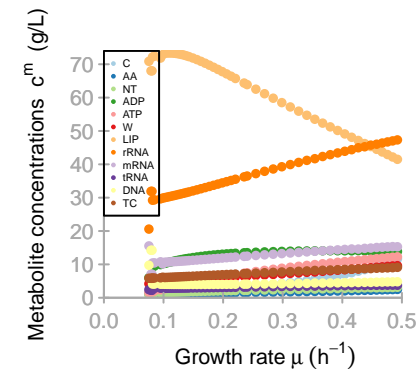
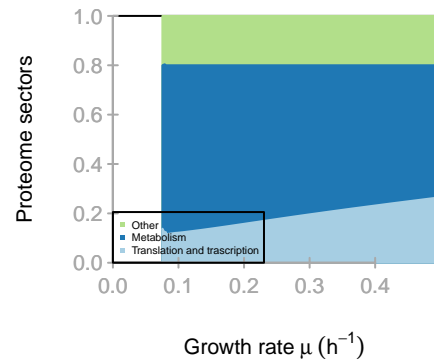
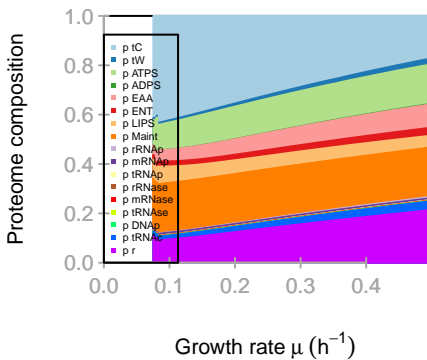
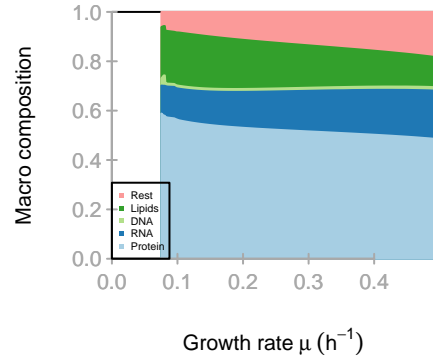
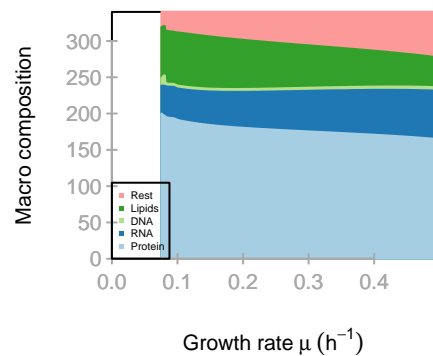
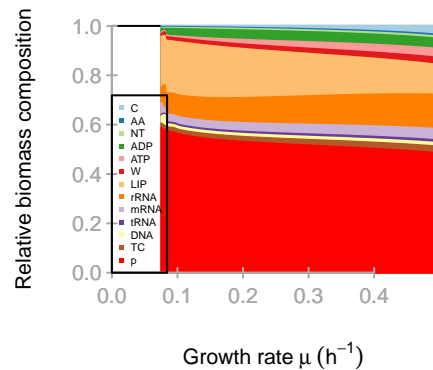
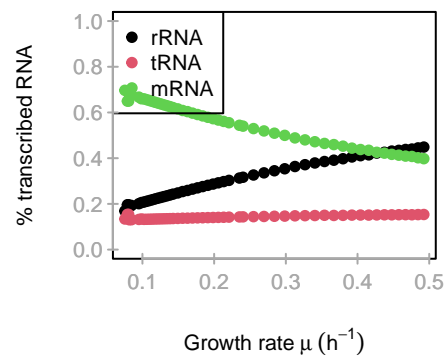
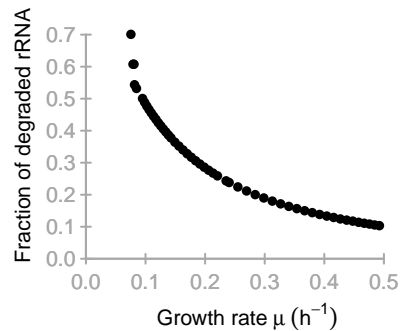
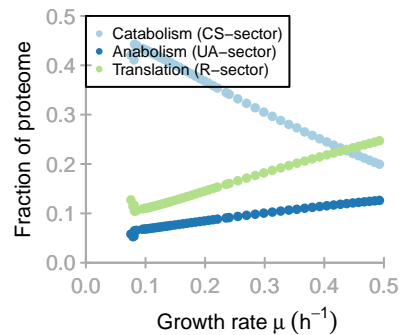


**Maint****rRNAp****mRNAp****tRNAp****rRNase****mRNase****tRNase****DNAp****tRNAc****r**





### Proteome sectors



**M**

[illegible]

## K

[illegible]

**KA**[illegible]

**kcat**[illegible]



# Keq

[1,]	[,1] 200	[,2] 1	[,3] 300	[,4] 5	[,5] 10	[,6] 50	[,7] 320	[,8] Inf	[,9] Inf	[,10] Inf	[,11] Inf	[,12] Inf	[,13] Inf	[,14] Inf	[,15] Inf	[,16] Inf	[,17] Inf
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## minimal phi constraint

[1,]	[,1] 0	[,2] 0	[,3] 0	[,4] 0	[,5] 0	[,6] 0	[,7] 0	[,8] 0.2	[,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]  
         0      [,9]  
         0      [,10]  
         0      [,11]  
         0      [,12]  
         0      [,13]  
         0      [,14]  
         0      [,15]  
         0      [,16]  
         0      [,17]  
         0