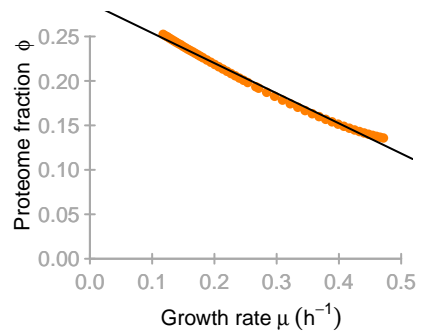
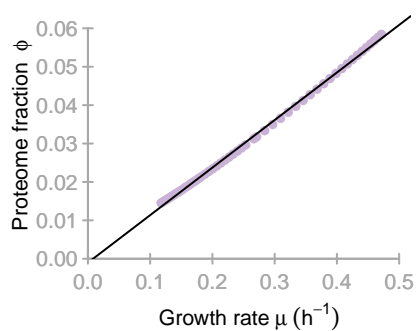
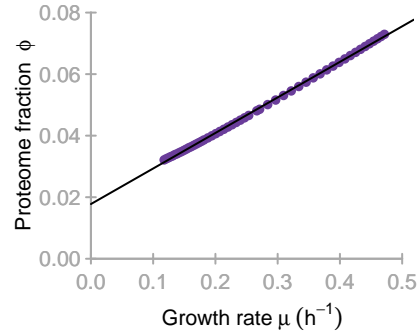
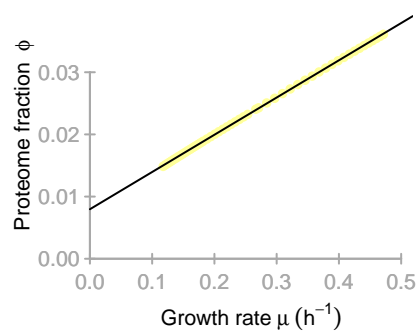
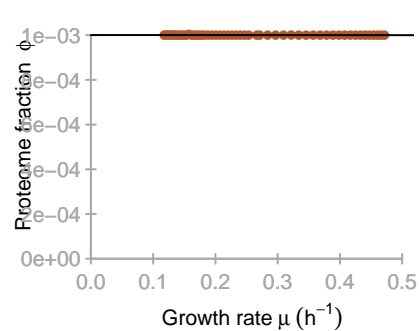
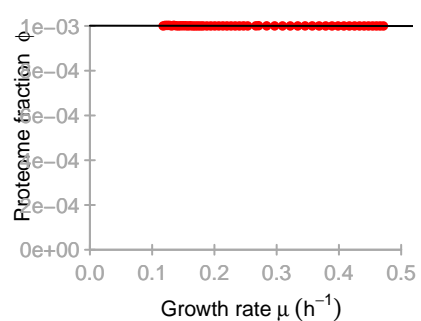
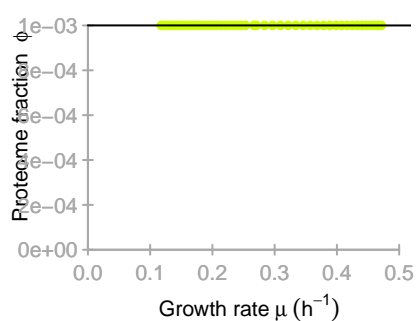
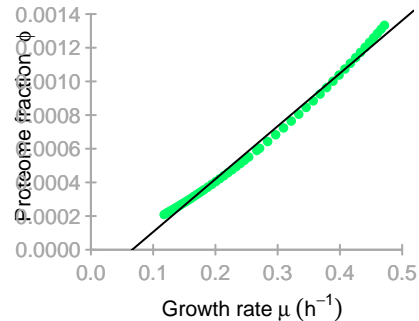
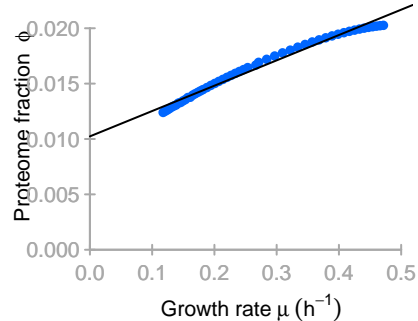
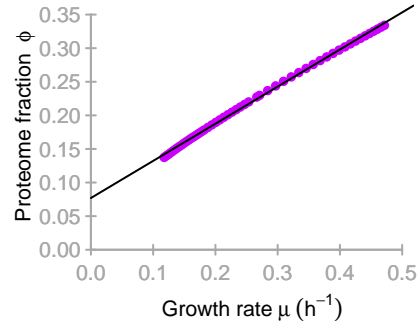
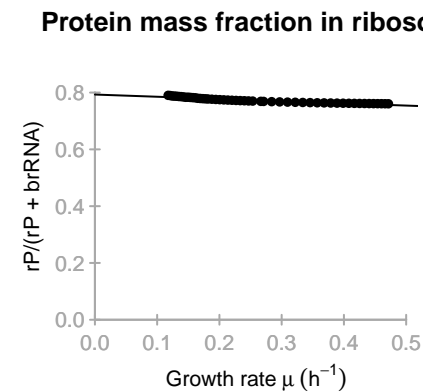
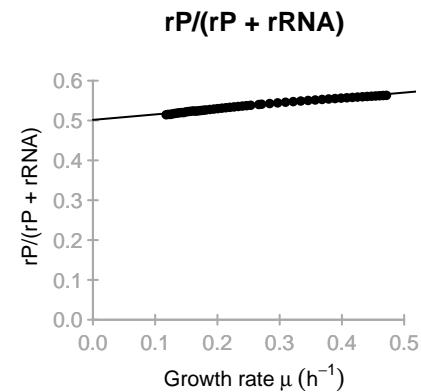
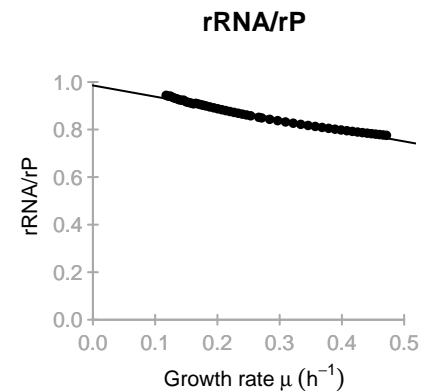
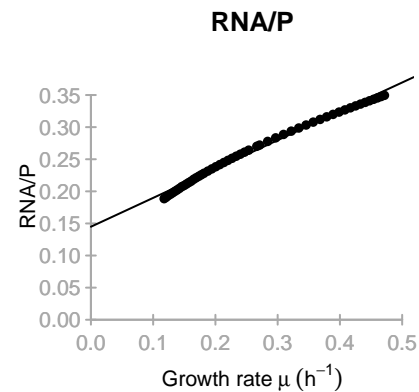
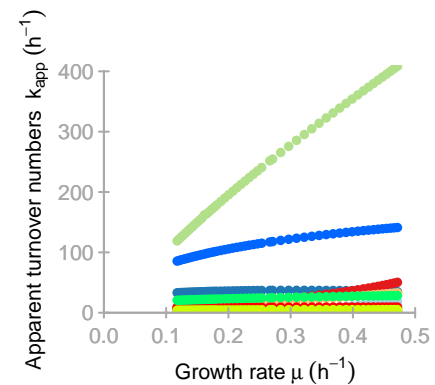
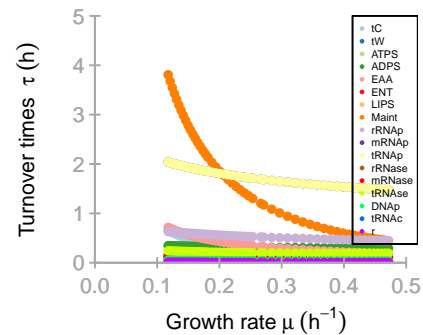
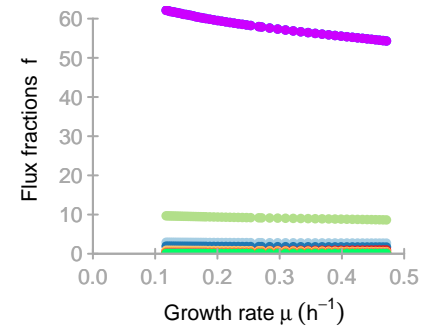
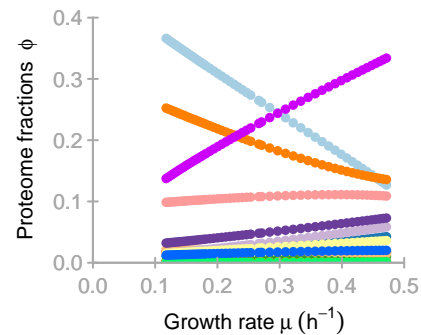
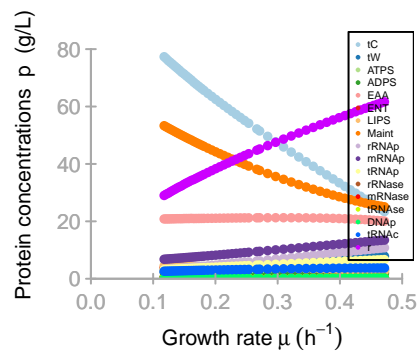
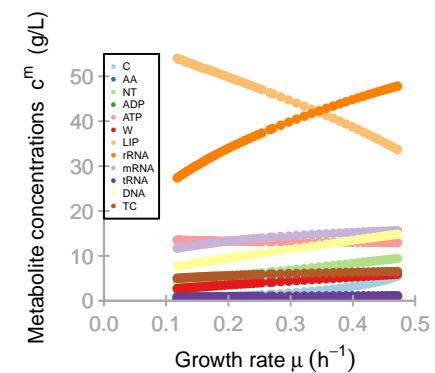
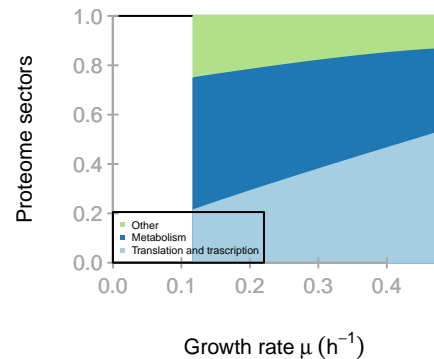
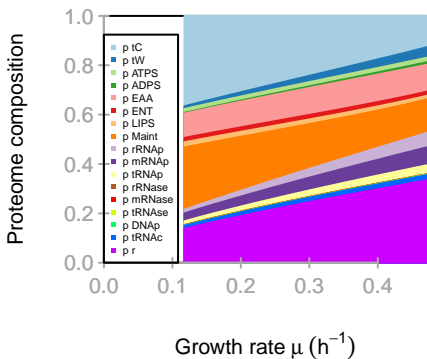
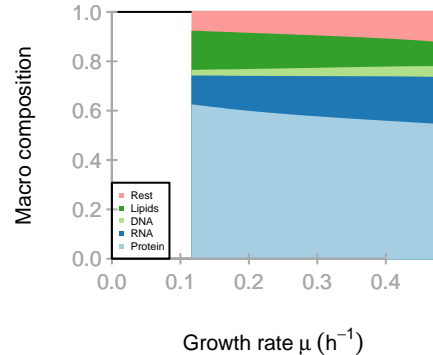
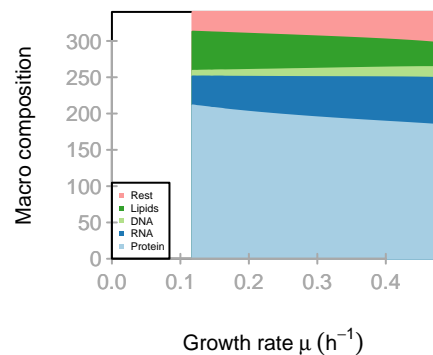
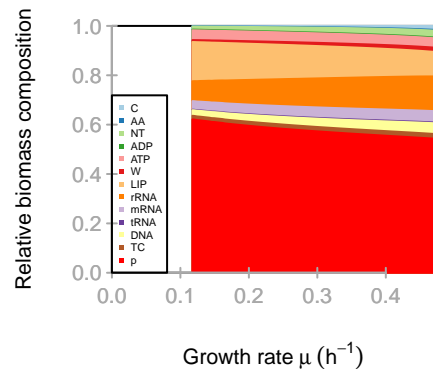
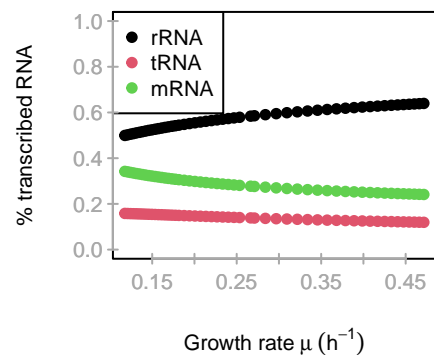
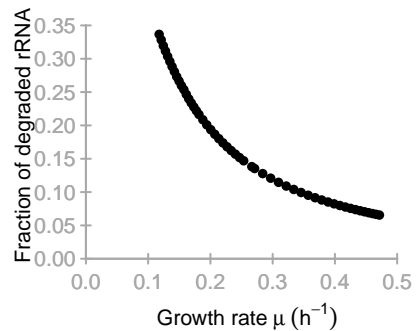
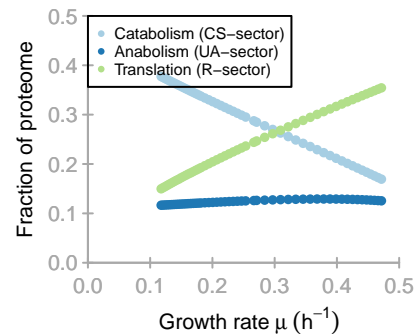


Maint**rRNAp****mRNAp****tRNAp****rRNase****mRNase****tRNase****DNAp****tRNAc****r**





Proteome sectors



M

[illegible]

K

[illegible]

KA[illegible]

Keq

[1.]	[,1] 100	[,2] 5	9.97938144329897	[,3] 5	[,4] 8	[,5] 9	[,6] 50	[,7] 169.6	[,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

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minimal f constraint

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