

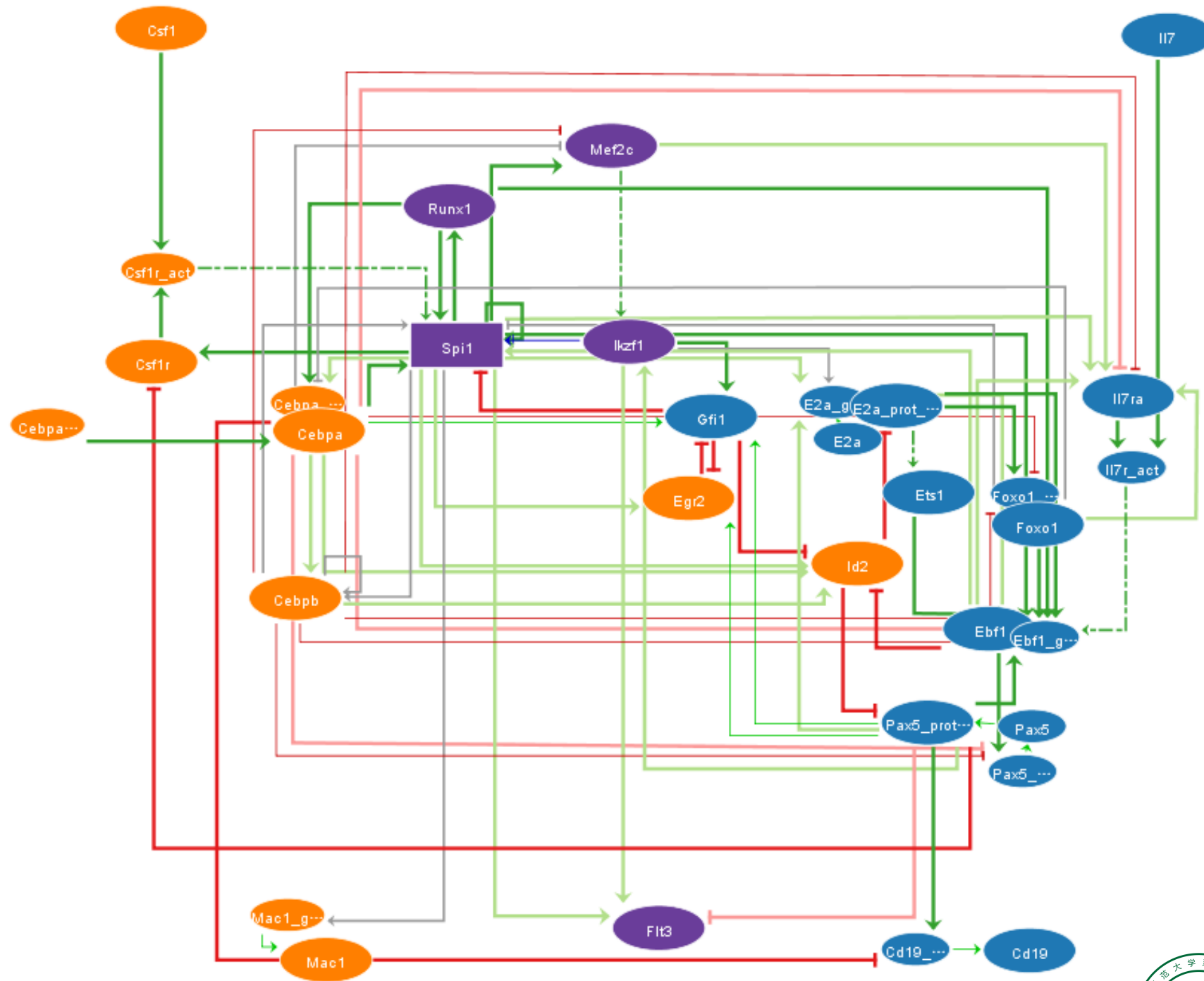
使用深度学习方法重构基因调控网络

报告人：张章

北京师范大学系统科学学院硕士在读



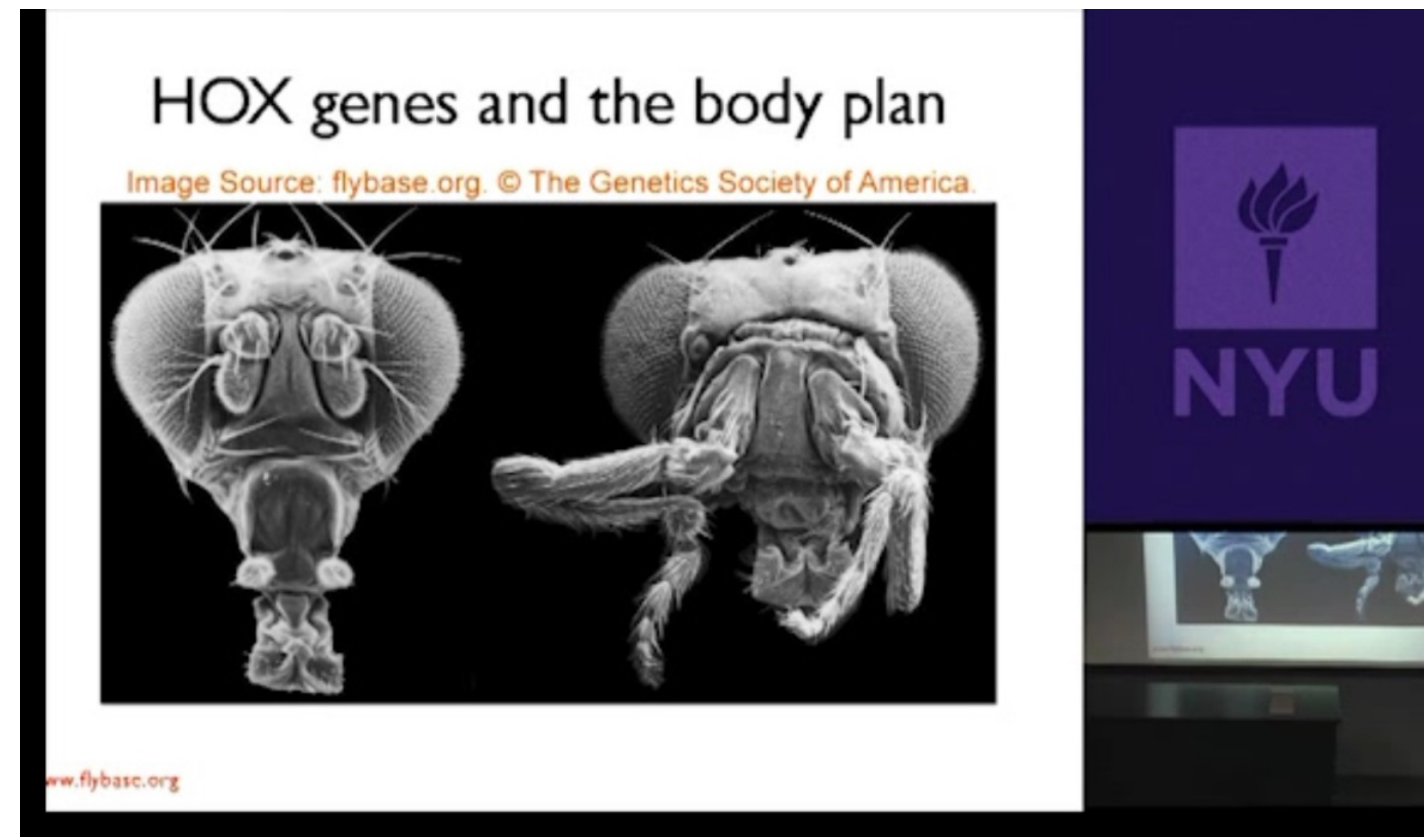
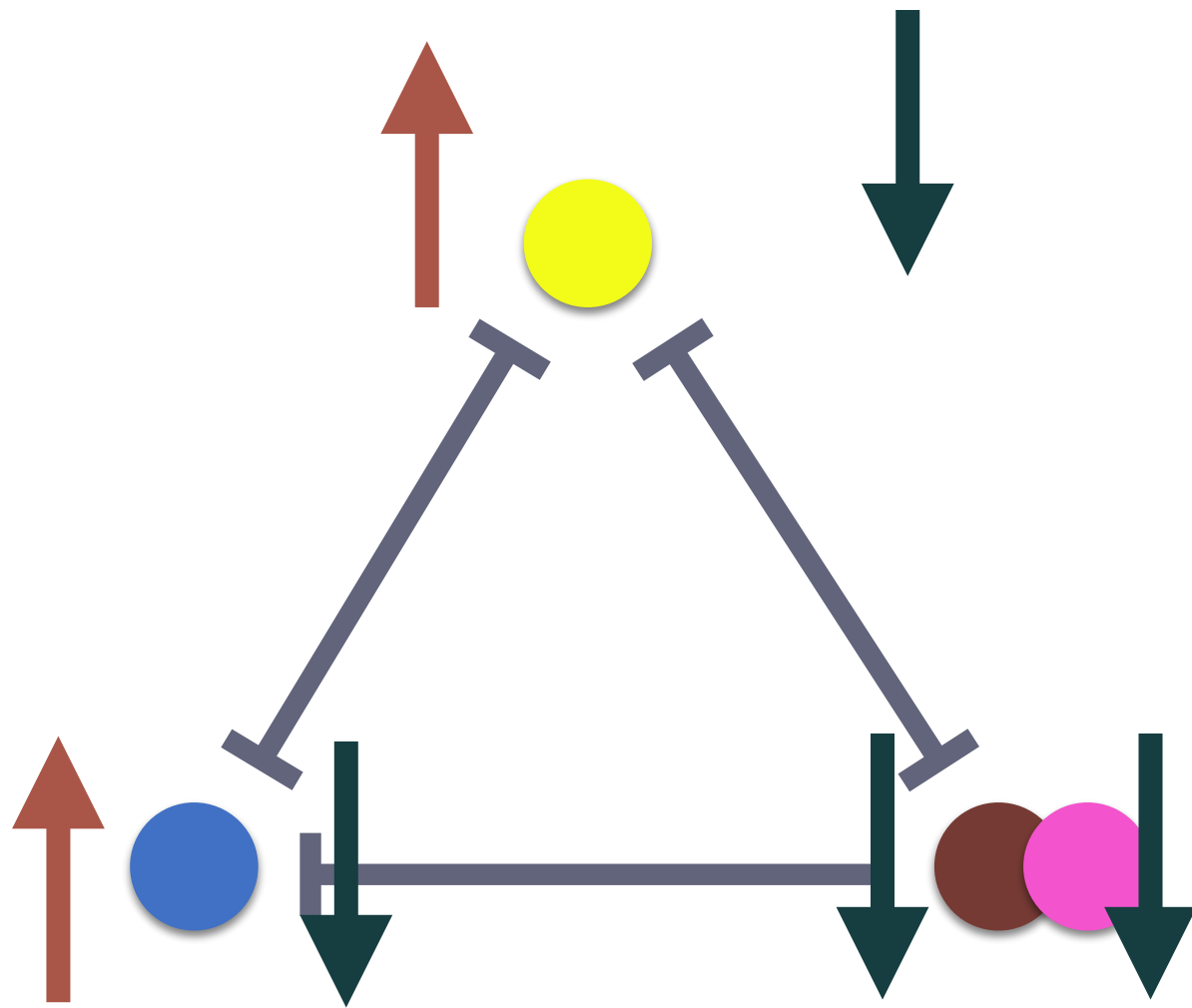
Problem Statement



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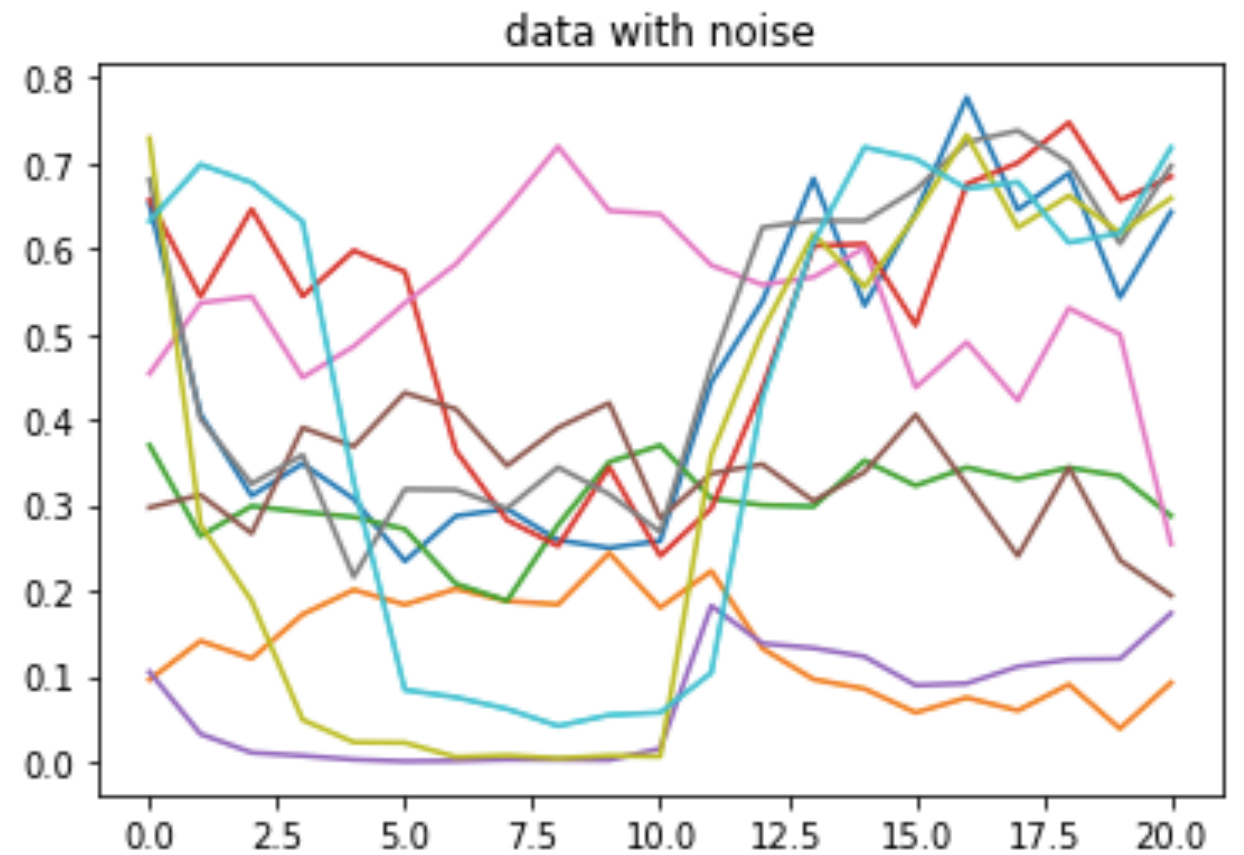
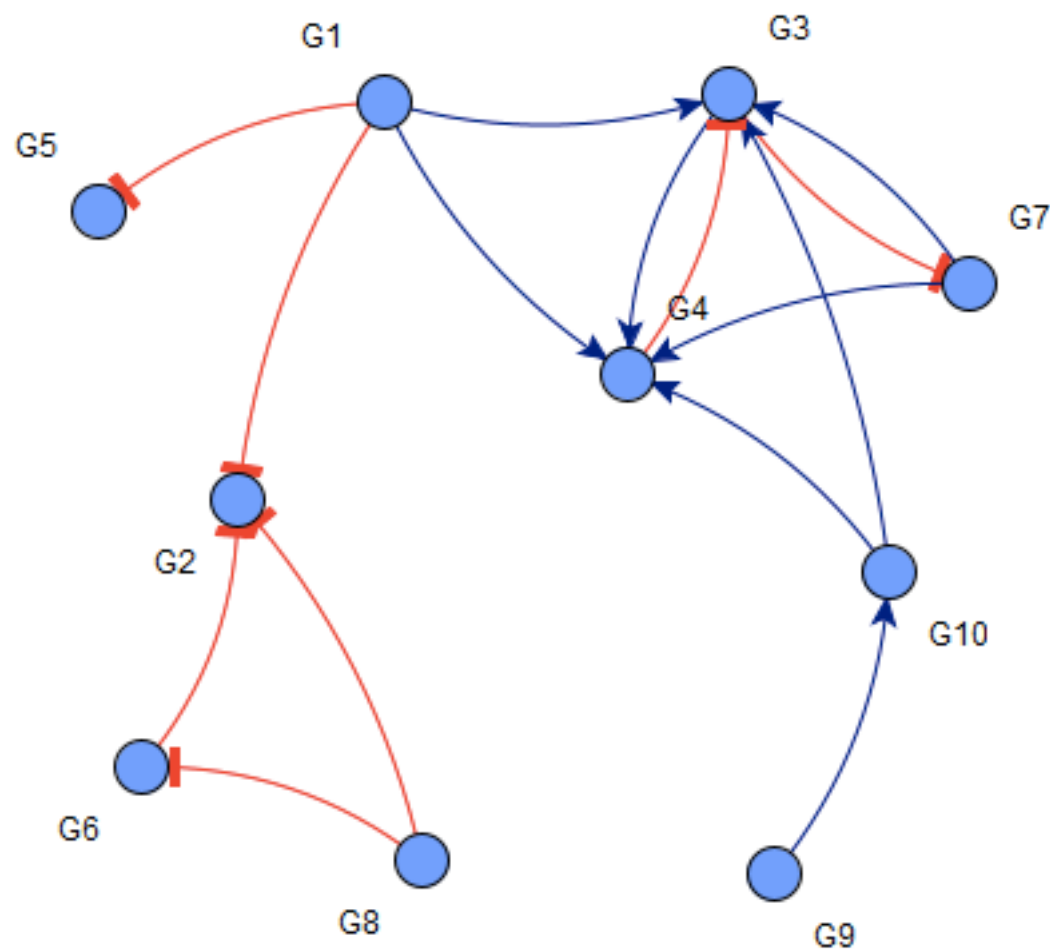


Significance



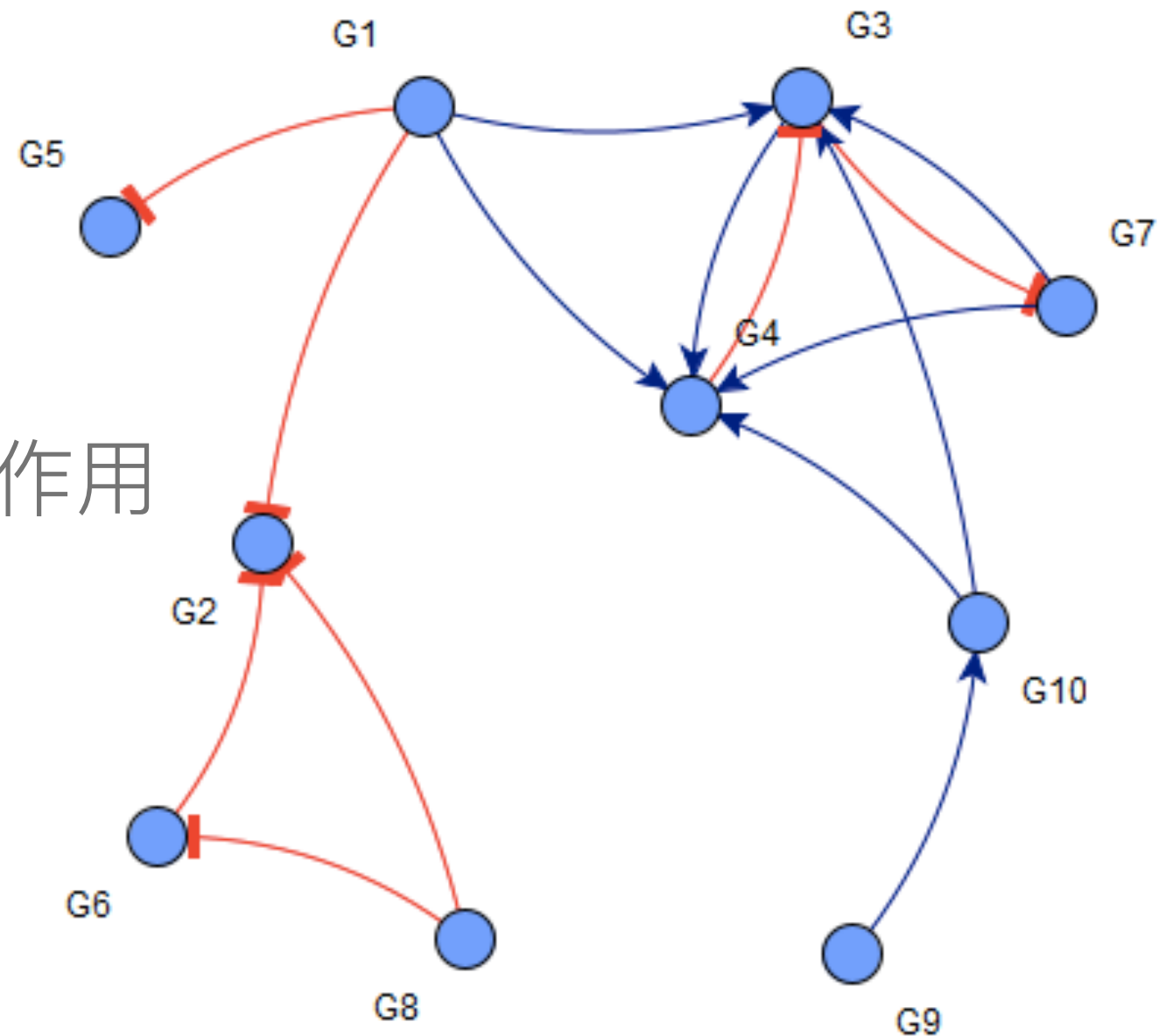
Problem Statement

- 从时序数据序列中重构基因调控网络



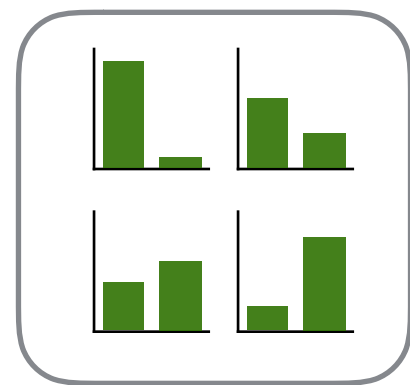
Mathematical model for Simulation 总结, 难度是什么, 用图来表示, 人们的已有认识

- 高度非线性的动力学过程
- 邻居地位不均等的动力学作用

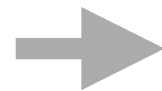


基本思路

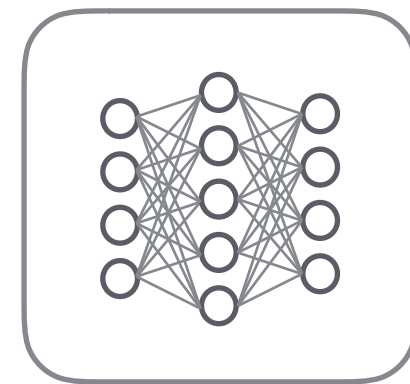
- 用网络生成器采样网络结构
- 用动力学预测器预测未来时序信息
- 共同优化网络生成器和动力学预测器



Network Generator

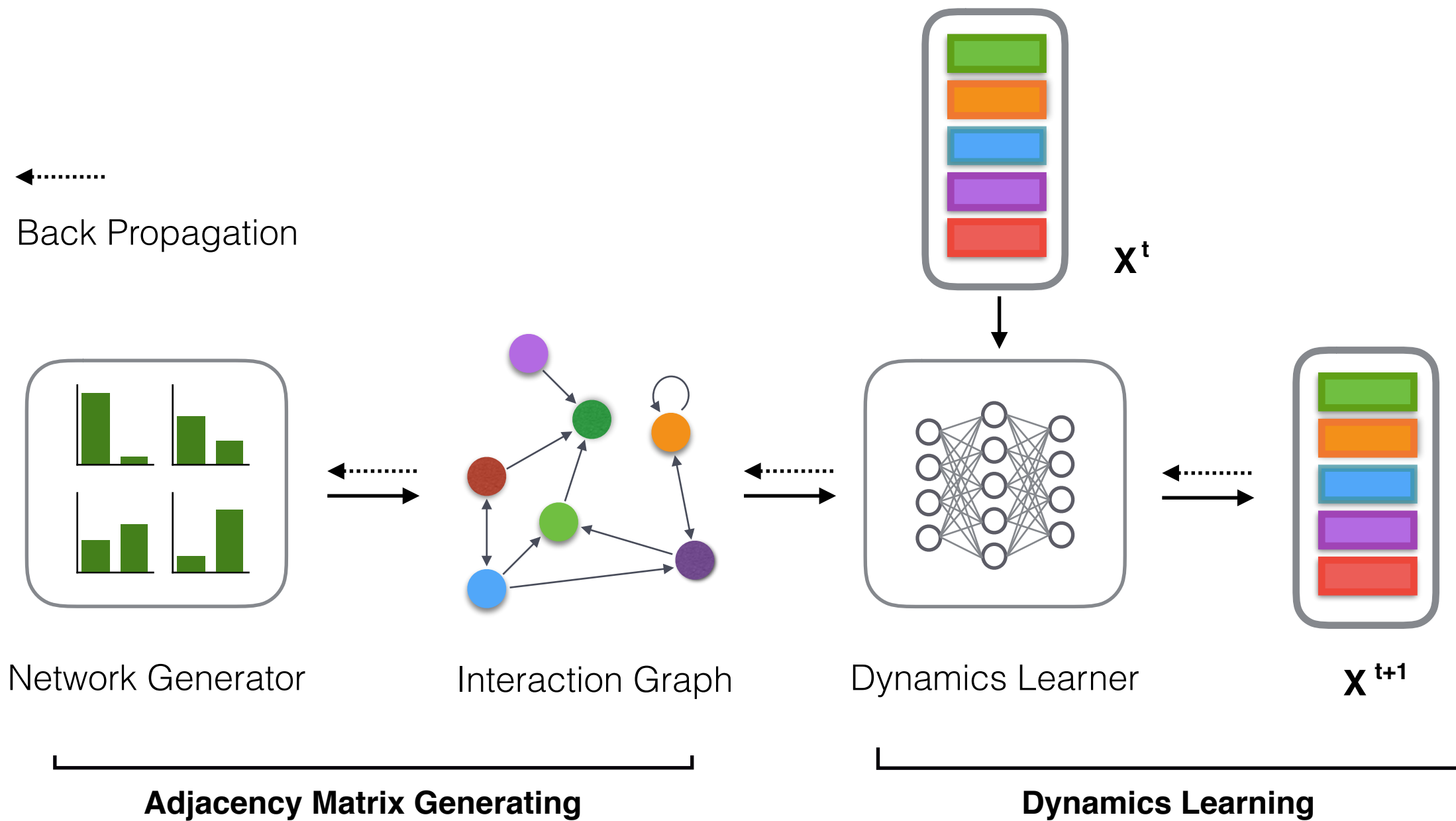


$[[0, 1],$
 $[1, 0]]$

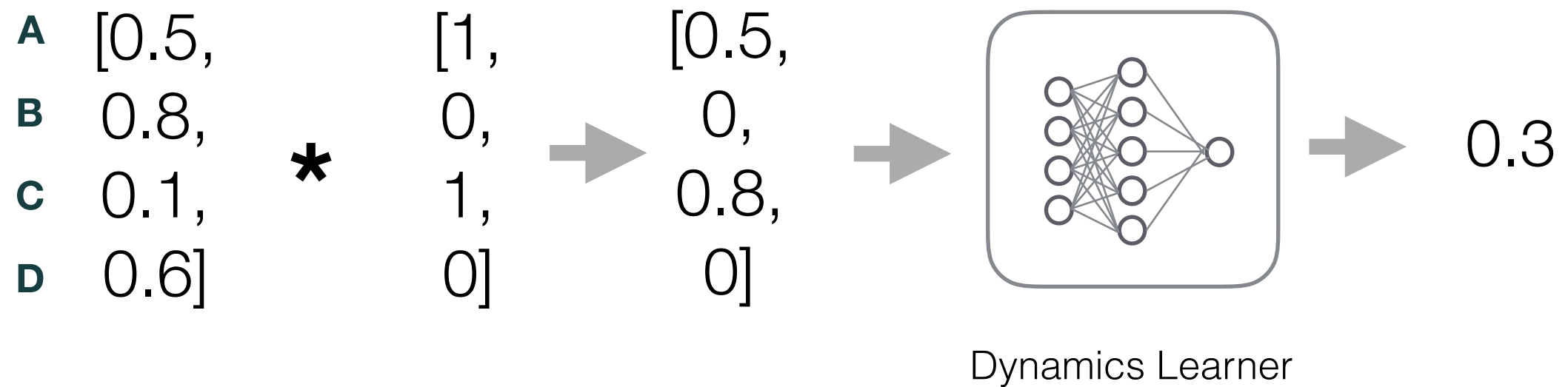
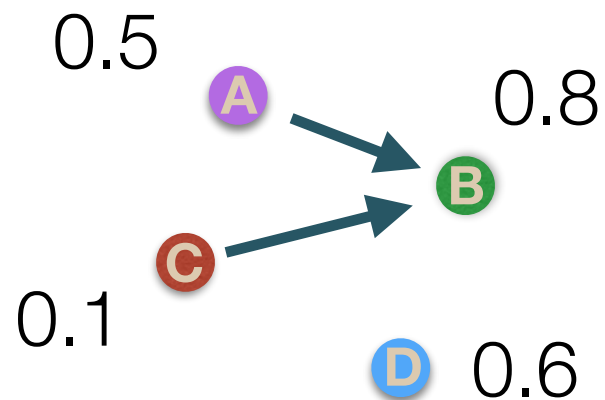


Dynamics Learner

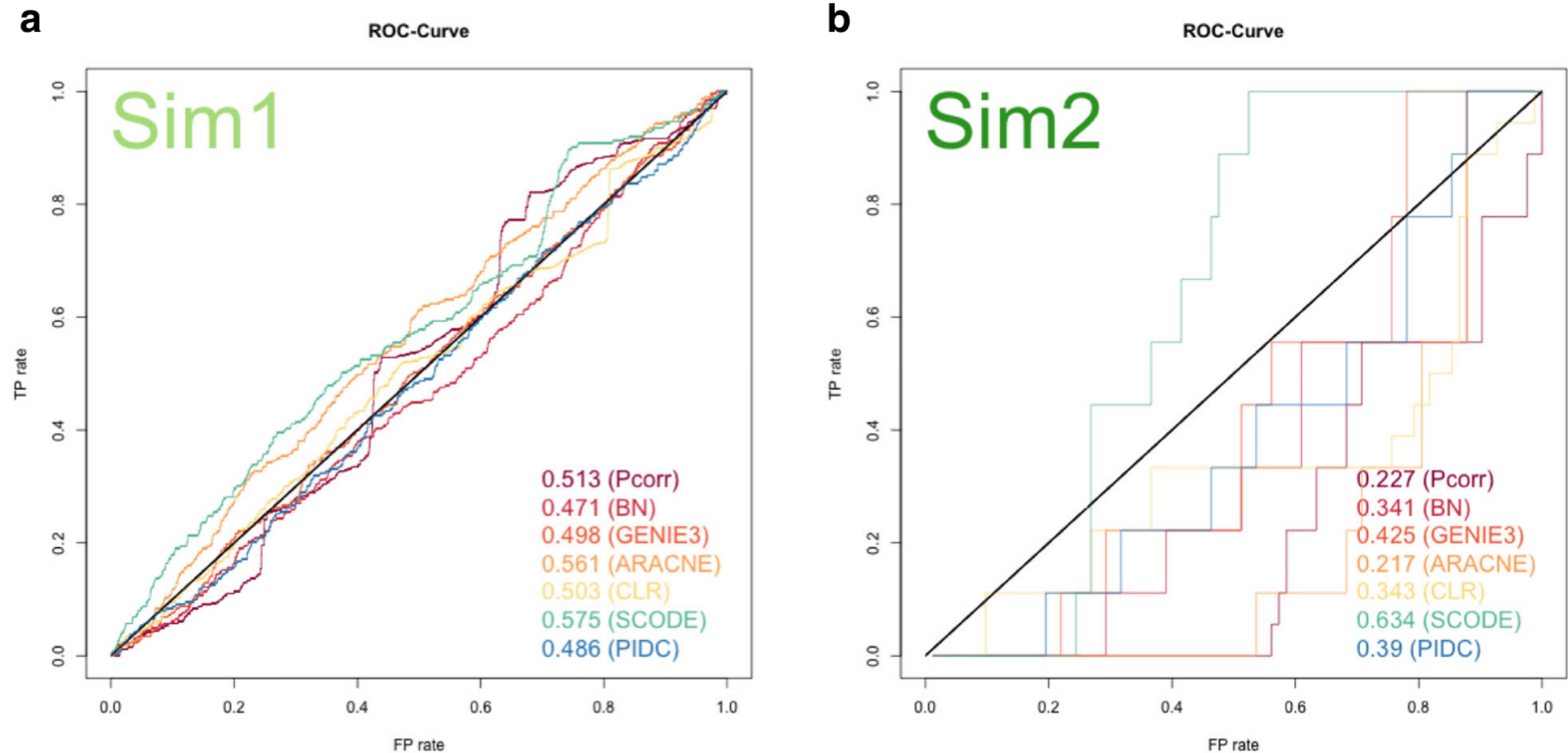
Algorithms Framework



Algorithms Framework: Dyn Learner



Experiments

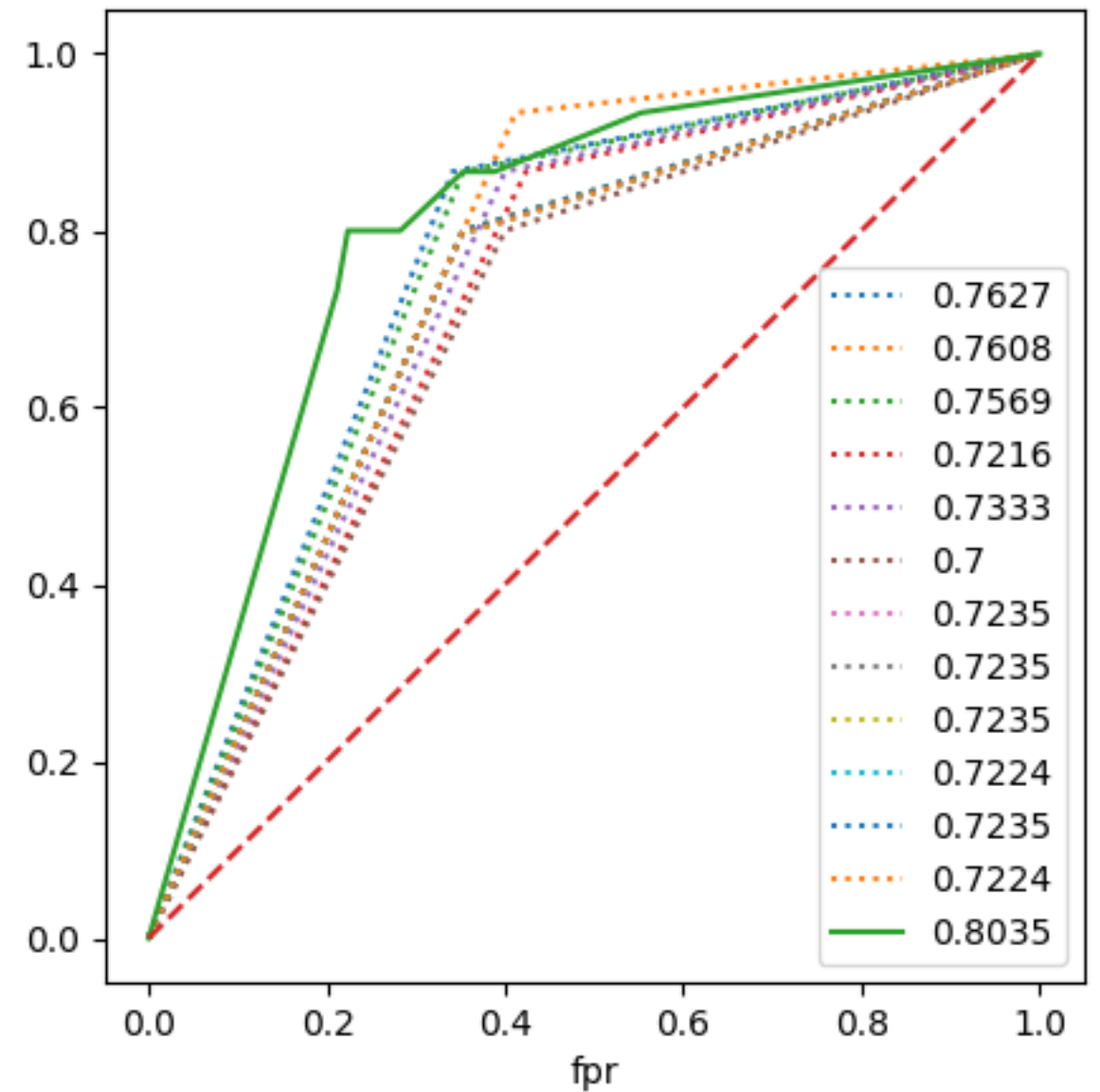
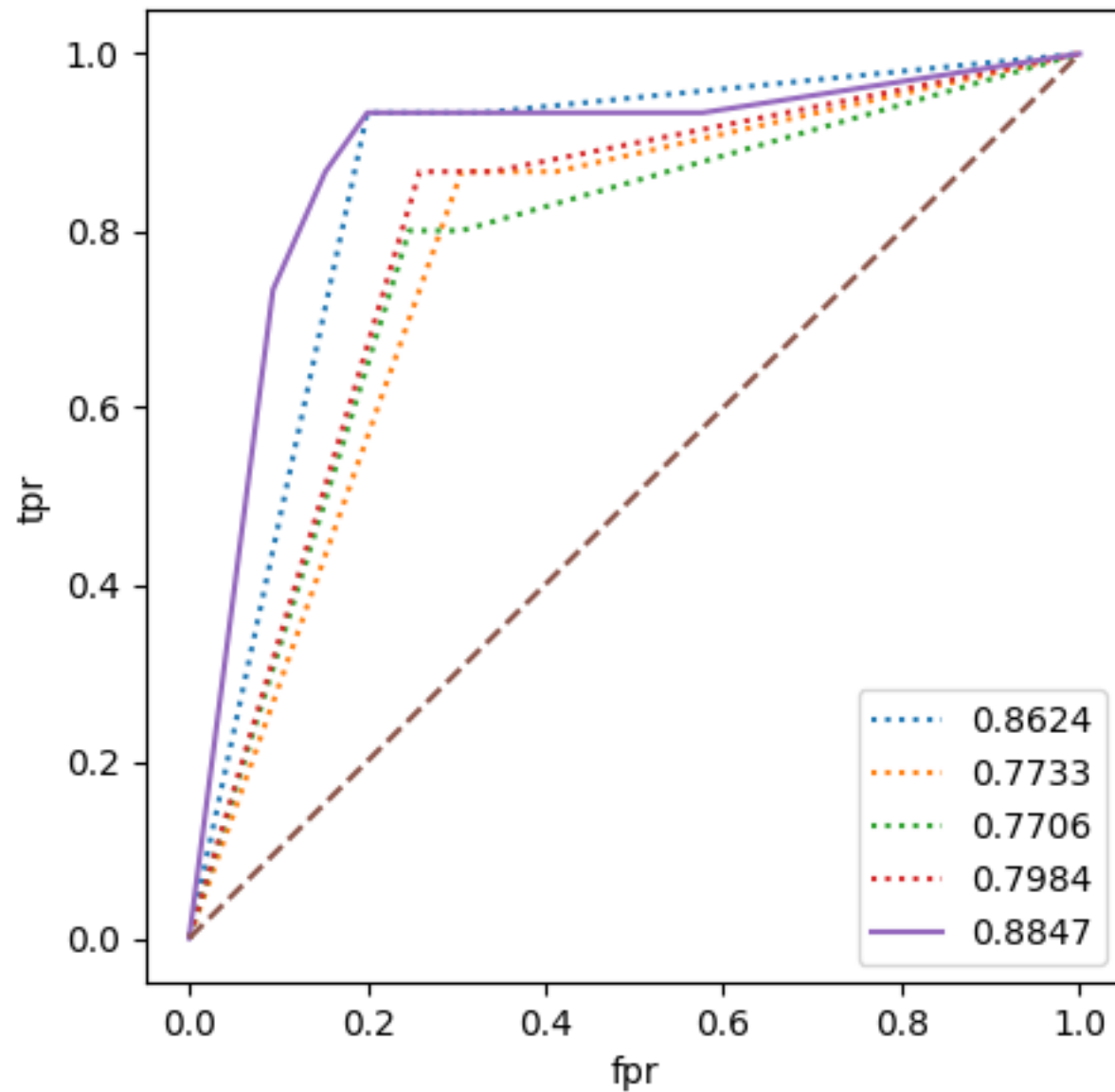


Chen S, Mar J C. Evaluating methods of inferring gene regulatory networks highlights their lack of performance for single cell gene expression data[J]. BMC Bioinformatics, 2018, 19(1).

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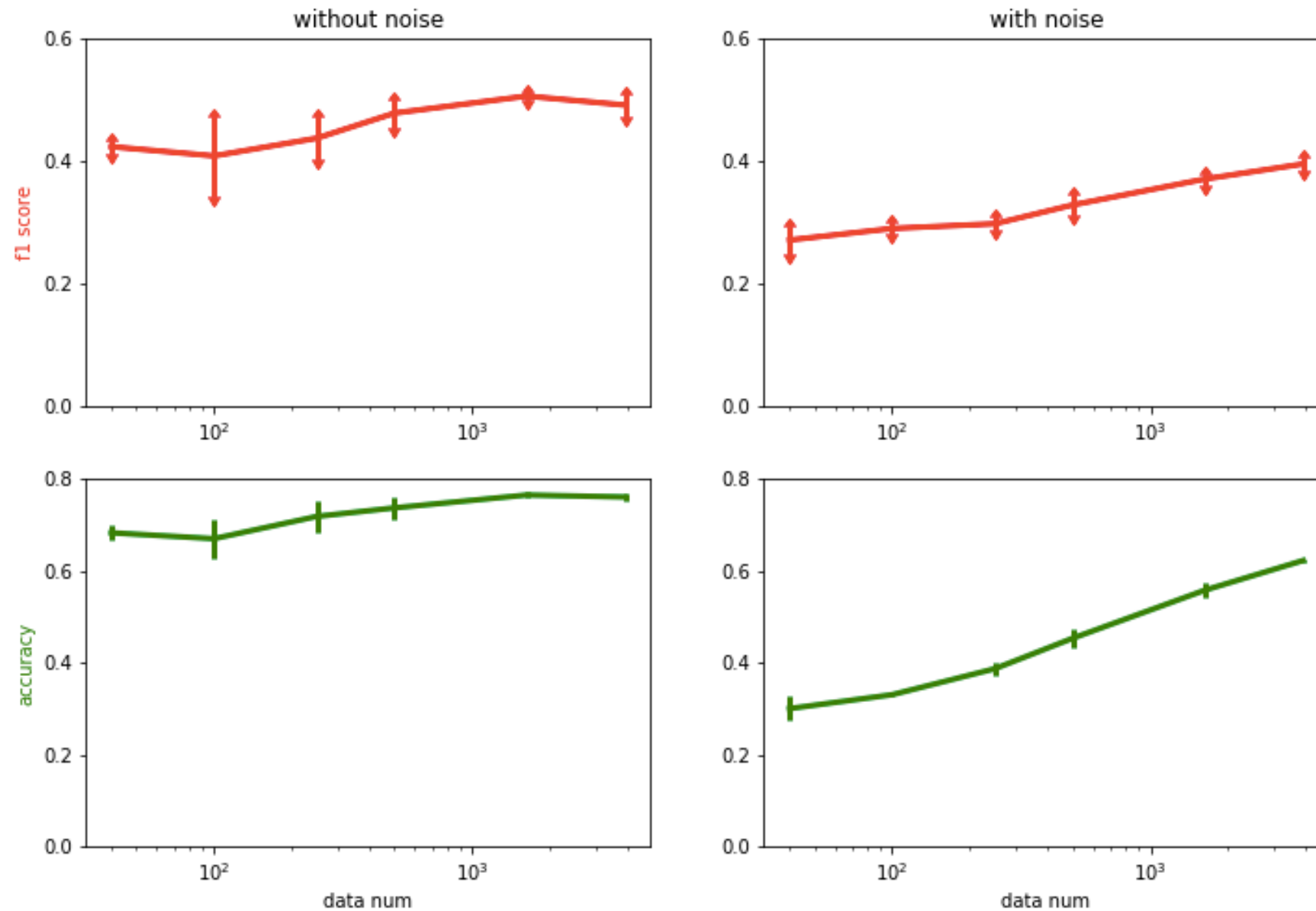
Experiments



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Experiments: data num vs. performance



Future Work

- 复现竞争对手的方法
- 探究时滞效应的影响
- 探究动力学学习器复杂度对重构效果的影响



Conclusion

- 新的，data-driven的重构基因调控网络的方法
- 可以同时完成网络的重构和动力学的预测
- SOTA的效果



Thank you for your listening

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