

Adaptive Network Reverse Engineering with L1-Magic

J.D. Stigter and J. Molenaar Biometris, Wageningen University, The Netherlands Email: hans.stigter@wur.nl

Background

Network Reverse Engineering addresses the question of finding the structure of a (unknown) network given input/output data (e.g. metabolic concentrations, transcription data, etc)

➤ We suggest

an innovative design of experiments based on maximization of the information content of the experimental data w.r.t.

Test Case

DREAM6 Challenge – Model 1 (gene regulatory network)



- parameter estimation
- Use of L₁ norm (Compressed Sensing Algorithm) for sparse network reconstruction

Method

- Perform an arbitrary experiment and use the input-output data as a first guess for determination of the parameters in the Jacobi matrix²
- 2. Analyse the resulting linear system, where the primary goal is now to maximize information content w.r.t. the Jacobi matrix parameters, using input or state perturbations

$$\frac{d(\delta x(t))}{dt} = A\delta x + B\delta u$$

3. The results of the second step yield an `optimal' perturbation that can now be used in an *improved* new experiment

Our method yields the interaction matrix (see figure below). The blue dots indicate a *direct* interaction between nodes.



4. If necessary, repeat step (2) with the new parameter estimates as to further improve the condition number of the Fisher Information Matrix



References

1. J.D. Stigter and J. Molenaar, Network Inference via Adaptive



For different noise-levels in the (synthetic) data that were obtained after simulation of the DREAM6 model, we calculated *specificity, sensitivity,* and *accuracy* for our 3 network reconstruction methods: (i) Least-Squares, (ii) L1, and (iii) L1 with optimal experimental design. Below is the so-called ROC plot. **Conclusion**: Optimal design with L1 norm on the

parameter vector clearly has the best performance.

$$1 - Specificity = 1 - \frac{TN}{FP + TN}$$

Sensitivity = $\frac{TP}{TP + FN}$

TP = True Positive FP = False Positive TN = True Negative FN = False Negative





- Optimal Design, to appear in BMC Research Notes, 2012.
 2. H. Schmidt, K. Cho, and E.W. Jacobsen, Identification of small scale biochemical networks based on general type system perturbations, FEBS Journal, 272, p2141—2151, 2005
- Candès E.J., Romberg J.K., Tao, T., Stable signal recovery from incomplete and inaccurate measurements, Communications on Pure and Applied Mathematics, Vol. 59(8), p1207—1223, 2006.
- J.D. Stigter, D. Vries, and K.J. Keesman, On adaptive optimal input design, AIChE, Vol. 52, No. 9, p3290—3296, 2006.