

## 1. Simulations in Experiment 4

In Experiment 4, we compare the power of the power-enhanced test with the likelihood-based model selection criterion of [3] and the eigenvalue-based goodness-of-fit test of [2]. We consider networks with  $n$  actors and adjacency matrix  $A \in \{0, 1\}^{K \times K}$ , sampled from a Stochastic Block Model with  $K = 2$  communities, community mixing matrix  $P \in [0, 1]^{K \times K}$  and community assignment multinomial probabilities  $a \in [0, 1]^K$ . We choose a fixed level  $\gamma$  for the PET and the test of [2]. We do not need to define the level for the method of [3], which is not a test but a model selection criterion.

### 1.1. Likelihood-based model selection criterion of [3]

The model selection criterion of [3] leverages the complete data likelihood to consistently estimate the number of communities in a Stochastic Block Model under the regime of growing degrees. For  $K' \in \{1, 2\}$ , we optimize the following mean field approximation of the criterion:

$$\beta(K'; A) = \sup_{(P, a) \in \mathcal{M}_{K'}} \sup_{q \in \mathcal{D}_{K'}} J(q, P, a; A) - \lambda \frac{K'(1 + K')}{2} n \log(n),$$

where  $\mathcal{D}_{K'}$  is the set of product distributions over  $\{0, 1\}^n$ ,  $\mathcal{M}_{K'}$  is the set of proper SBM parameters and  $\lambda$  is a tuning parameter chosen using the procedure described in section 3.2. of [3].  $J(q, P, a; A)$  corresponds to the mean field approximation of the log-likelihood, which can be expressed as

$$\begin{aligned} J(q, P, a; A) &= \sum_{i=1}^n \sum_{k=1}^{K'} \tau_{ik} (-\log(\tau_{ik}) + \log(a_k)) \\ &\quad + \frac{1}{2} \sum_{i \neq j} \sum_{k, l=1}^{K'} \tau_{ik} \tau_{jl} (A_{ij} \log(P_{kl}) + (1 - A_{ij}) \log(1 - P_{kl})), \end{aligned}$$

where  $\{\tau_{ik}\}_{i,k}$  are the multinomial parameters of the mean field distribution  $q$ , with the constraint that  $\sum_k \tau_{ik} = 1$  for all  $i \in \llbracket 1, n \rrbracket$ . [1] showed that for fixed  $(P, a)$ , maximizing  $J(q, P, a; A)$  with respect to  $q$  can be done by solving the fixed point equation

$$\hat{\tau}_{ik} \propto a_k \prod_{\substack{j=1 \\ j \neq i}}^n \prod_{l=1}^{K'} \left( P_{kl}^{A_{ij}} (1 - P_{kl})^{1 - A_{ij}} \right)^{\hat{\tau}_{jl}}.$$

Then, given the variational parameters  $\{\tau_{ik}\}_{i,k}$ , the values of the parameters  $P$  and  $a$  that maximize  $J(q, P, a; A)$  are

$$\hat{a}_k = \frac{1}{n} \sum_{i=1}^n \hat{\tau}_{ik}, \quad \hat{P}_{kl} = \frac{\sum_{i \neq j} \hat{\tau}_{ik} \hat{\tau}_{jl} A_{ij}}{\sum_{i \neq j} \hat{\tau}_{ik} \hat{\tau}_{jl}}.$$

Following the estimation algorithm of [1], we start from an initial guess for  $\{\hat{\tau}_{ik}^{(0)}\}_{i,k}$  (which could be random or using spectral clustering of the regularized Laplacian matrix) and iteratively update  $\{\hat{a}_k\}_k$ ,  $\hat{P}$  and  $\{\hat{\tau}_{ik}\}_{i,k}$  until convergence of  $J(q, P, a; A)$ . For a network  $A$  drawn from our model, we use this procedure to compute  $\beta(1; A)$  and  $\beta(2; A)$ . We call a true positive discovery when  $\beta(2; A) > \beta(1; A)$ , and a false negative discovery when  $\beta(2; A) \leq \beta(1; A)$ .

## 1.2. Eigenvalue-based test of [2]

We also compare the PET to the goodness-of-fit test of [2], which uses the largest singular value of a residual matrix  $\tilde{A}$  obtained by subtracting the estimated block mean effect from the adjacency matrix. We choose an Erdős-Rényi null model. In this setting, we reject the null hypothesis if

$$T_{n,1} := n^{2/3}[\sigma_1(\tilde{A}) - 2] \geq t(\gamma/2),$$

where  $\gamma$  is the level of the test,  $\sigma_1(\tilde{A})$  is the largest singular value of  $\tilde{A}$  and  $t(\gamma/2)$  is the  $\gamma/2$  upper quantile of the Tracy-Widom  $TW_1$  distribution. In the test statistic  $T_{n,1}$ , the residual matrix  $\tilde{A}$  is computed using an estimated community membership vector  $(\hat{z}_i)_{i=1}^n$ . The vector  $(\hat{z}_i)_{i=1}^n$  is obtained from spectral clustering, where we apply  $k$ -means clustering to the rows of the matrix formed by the 2 leading singular vectors of  $A$ . Then,  $\tilde{A}$  can be expressed as

$$\tilde{A}_{ij} = \begin{cases} \frac{A_{ij} - \hat{\Omega}_{ij}}{\sqrt{(n-1)\hat{\Omega}_{ij}(1-\hat{\Omega}_{ij})}}, & i \neq j, \\ 0, & i = j, \end{cases}$$

where  $\hat{\Omega}_{ij} = \hat{P}_{\hat{z}_i \hat{z}_j}$  and

$$\hat{\mathcal{N}}_k = \{i : 1 \leq i \leq n, \hat{z}_i = k\}, \quad \hat{n}_k = |\hat{\mathcal{N}}_k|, \quad \hat{P}_{kl} = \begin{cases} \frac{\sum_{i \in \hat{\mathcal{N}}_k, j \in \hat{\mathcal{N}}_l} A_{ij}}{\hat{n}_k \hat{n}_l}, & k \neq l, \\ \frac{\sum_{i, j \in \hat{\mathcal{N}}_k, i < j} A_{ij}}{\hat{n}_k(\hat{n}_k - 1)/2}, & k = l. \end{cases}$$

## References

- [1] DAUDIN, J.-J., PICARD, F. and ROBIN, S. (2008). A mixture model for random graphs. *Statistics and computing* **18** 173–183.
- [2] LEI, J. (2016). A goodness-of-fit test for stochastic block models. *Ann. Statist.* **44** 401–424.
- [3] WANG, Y. R. and BICKEL, P. J. (2017). Likelihood-based model selection for stochastic block models. *Ann. Statist.* **45** 500–528.