

bayesian inference: principles and practice

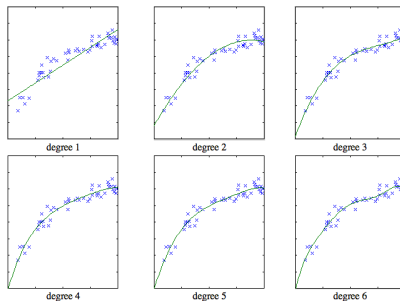
jake hofman

<http://jakehofman.com>

july 9, 2009

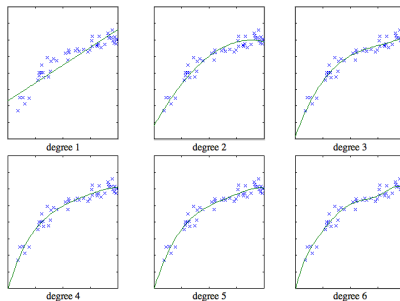
motivation

- would like models that:
 - provide predictive and explanatory power
 - are complex enough to describe observed phenomena
 - are simple enough to generalize to future observations



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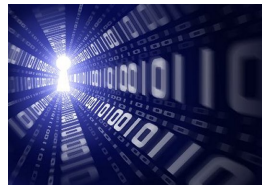
- claim: bayesian inference provides a systematic framework to infer such models from observed data

motivation

- principles behind bayesian interpretation of probability and bayesian inference are well established (bayes, laplace, etc., 18th century)

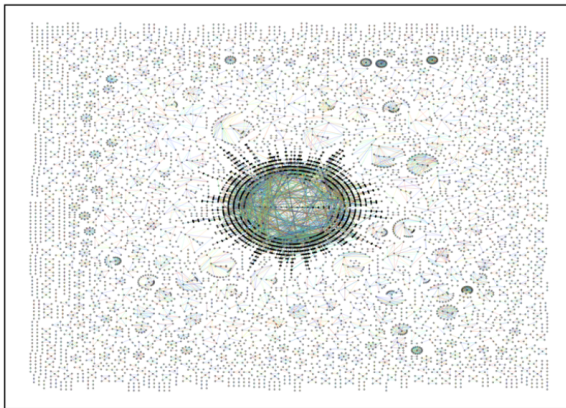


+



- recent advances in mathematical techniques and computational resources have enabled successful applications of these principles to real-world problems

motivation: a bayesian approach to network modularity



outline

- 1 principles (what we'd like to do)
 - background: joint, marginal, and conditional probabilities
 - bayes' theorem: inverting conditional probabilities
 - bayesian probability: unknowns as random variables
 - bayesian inference: bayesian probability + bayes' theorem
- 2 practice (what we're able to do)
 - monte carlo methods: representative samples
 - variational methods: bound optimization
 - references

joint, marginal, and conditional probabilities

joint distribution

$p_{XY}(X = x, Y = y)$: probability $X = x$ *and* $Y = y$

conditional distribution

$p_{X|Y}(X = x|Y = y)$: probability $X = x$ *given* $Y = y$

marginal distribution

$p_X(X)$: probability $X = x$ (*regardless of* Y)

sum and product rules

sum rule

sum out settings of irrelevant variables:

$$p(x) = \sum_{y \in \Omega_Y} p(x, y) \quad (1)$$

product rule

the joint as the product of the conditional and marginal:

$$p(x, y) = p(x|y) p(y) \quad (2)$$

$$= p(y|x) p(x) \quad (3)$$

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inverting conditional probabilities

equate far right- and left-hand sides of product rule

$$p(y|x) p(x) = p(x, y) = p(x|y) p(y) \quad (4)$$

and divide:

bayes' theorem (bayes and price 1763)

the probability of Y given X from the probability of X given Y :

$$p(y|x) = \frac{p(x|y) p(y)}{p(x)} \quad (5)$$

where $p(x) = \sum_{y \in \Omega_Y} p(x|y) p(y)$ is the normalization constant

example: diagnoses a la bayes

- population 10,000
- 1% has (rare) disease
- test is 99% (relatively) effective, i.e.
 - given a patient is sick, 99% test positive
 - given a patient is healthy, 99% test negative

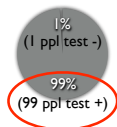
example: diagnoses a la bayes

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- 1% has (rare) disease
- test is 99% (relatively) effective, i.e.
 - given a patient is sick, 99% test positive
 - given a patient is healthy, 99% test negative
- given positive test, what is probability the patient is sick?¹

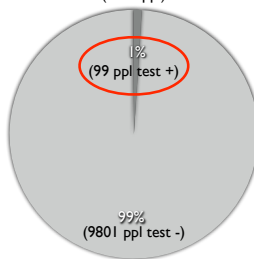
¹follows wiggins (2006)

example: diagnoses a la bayes

sick population
(100 ppl)



healthy population
(9900 ppl)



- 99 sick patients test positive, 99 healthy patients test positive

example: diagnoses a la bayes



- 99 sick patients test positive, 99 healthy patients test positive
- given positive test, 50% probability that patient is sick

example: diagnoses a la bayes

- know probability of testing positive/negative given sick/healthy
- use bayes' theorem to “invert” to probability of sick/healthy given positive/negative test

$$p(sick|test +) = \frac{\overbrace{p(test + | sick)}^{99/100} \overbrace{p(sick)}^{1/100}}{\underbrace{p(test +)}_{198/100^2}} = \frac{99}{198} = \frac{1}{2} \quad (6)$$

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- most “work” in calculating denominator (normalization)

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interpretations of probabilities

(just enough philosophy)

- frequentists: limit of relative frequency of events for large number of trials
- bayesians: measure of a state of knowledge, quantifying degrees of belief (jaynes 2003)

interpretations of probabilities

(just enough philosophy)

- frequentists: limit of relative frequency of events for large number of trials
- bayesians: measure of a state of knowledge, quantifying degrees of belief (jaynes 2003)
- key difference: bayesians permit assignment of probabilities to unknown/unobservable hypotheses (frequentists do not)

interpretations of probabilities

(just enough philosophy)

- e.g., inferring model parameters Θ from observed data \mathcal{D} :
 - frequentist approach: calculate parameter setting that maximizes likelihood of data (point estimate),

$$\hat{\Theta} = \underset{\Theta}{\operatorname{argmax}} p(\mathcal{D}|\Theta) \quad (7)$$

- bayesian approach: calculate distribution over parameter settings given data,

$$p(\Theta|\mathcal{D}) = ? \quad (8)$$

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bayesian probability + bayes' theorem

- bayesian inference:
 - treat unknown quantities as random variables
 - use bayes' theorem to systematically update prior knowledge in the presence of observed data

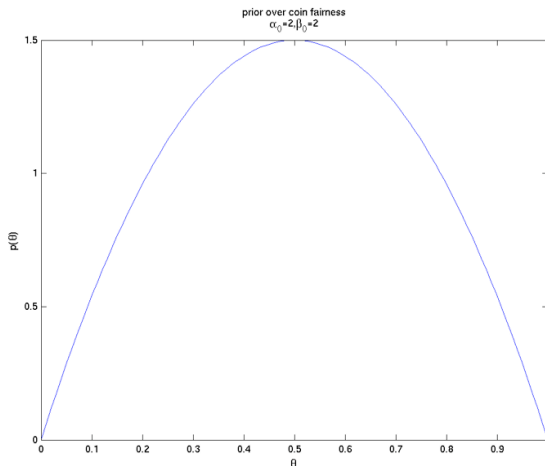
$$\underbrace{p(\Theta|\mathcal{D})}_{\text{posterior}} = \frac{\underbrace{p(\mathcal{D}|\Theta)}_{\text{likelihood}} \underbrace{p(\Theta)}_{\text{prior}}}{\underbrace{p(\mathcal{D})}_{\text{evidence}}} \quad (9)$$

example: coin flipping

- observe independent coin flips (bernoulli trials)
- infer distribution over coin bias

example: coin flipping

prior $p(\theta)$ over coin bias before observing flips

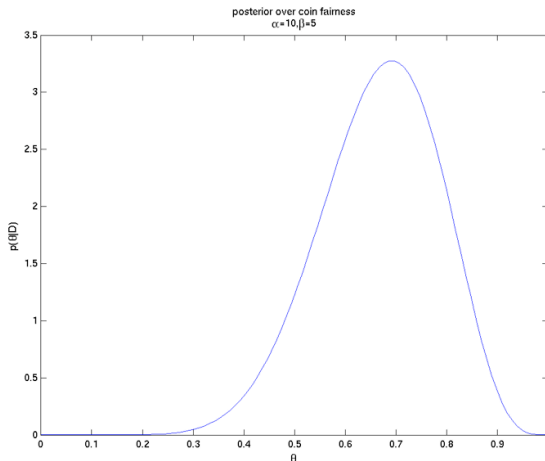


example: coin flipping

observe flips: HTHHHTTTHHHH

example: coin flipping

update posterior $p(\theta|\mathcal{D})$ using bayes' theorem

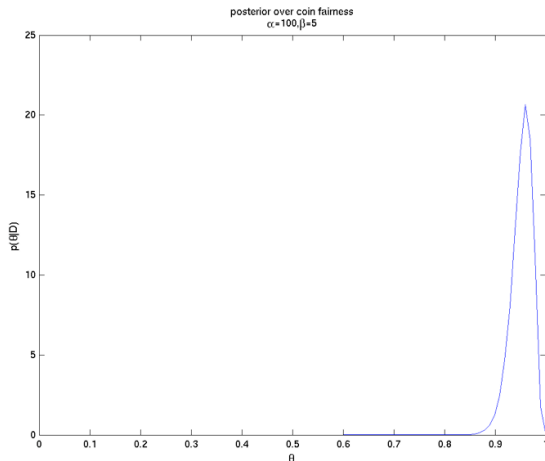


example: coin flipping

observe flips: HHHHHHHHHHHHHHTHHHHHHHHHHHH
HHHHHHHHHHHHHHHHHHHHHHHHHHHHHHHHHHHH
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example: coin flipping

update posterior $p(\theta|\mathcal{D})$ using bayes' theorem



quantities of interest

- bayesian inference maintains full posterior distributions over unknowns
- many quantities of interest require expectations under these posteriors, e.g. posterior mean and predictive distribution:

$$\bar{\Theta} = \mathbb{E}_{p(\Theta|\mathcal{D})} [\Theta] = \int d\Theta \Theta p(\Theta|\mathcal{D}) \quad (10)$$

$$p(x|\mathcal{D}) = \mathbb{E}_{p(\Theta|\mathcal{D})} [p(x|\Theta, \mathcal{D})] = \int d\Theta p(x|\Theta, \mathcal{D}) p(\Theta|\mathcal{D}) \quad (11)$$

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- often can't compute posterior (normalization), let alone expectations with respect to it → approximation methods

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representative samples

- general approach: approximate intractable expectations via sum over representative samples²

$$\Phi = \mathbb{E}_{p(x)} [\phi(x)] = \int dx \underbrace{\phi(x)}_{\text{arbitrary function}} \underbrace{p(x)}_{\text{target density}} \quad (12)$$

²follows mackay (2003), including stolen images

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$$\Downarrow$$
$$\hat{\Phi} = \frac{1}{R} \sum_{r=1}^R \phi(x^{(r)}) \quad (13)$$

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- shifts problem to finding “good” samples

²follows mackay (2003), including stolen images

representative samples

- further complication: in general we can only evaluate the target density to within a multiplicative (normalization) constant, i.e.

$$p(x) = \frac{p^*(x)}{Z} \quad (14)$$

and $p^*(x^{(r)})$ can be evaluated with Z unknown

sampling methods

- monte carlo methods
 - uniform sampling
 - importance sampling
 - rejection sampling
 - ...
- markov chain monte carlo (mcmc) methods
 - metropolis-hastings
 - gibbs sampling
 - ...

uniform sampling

- sample uniformly from state space of all x values
- evaluate non-normalized density $p^*(x^{(r)})$ at each $x^{(r)}$
- approximate normalization constant as

$$Z_R = \sum_{r=1}^R p^*(x^{(r)}) \quad (15)$$

- estimate expectation as

$$\hat{\Phi} = \sum_{r=1}^R \phi(x^{(r)}) \frac{p^*(x^{(r)})}{Z_R} \quad (16)$$

uniform sampling

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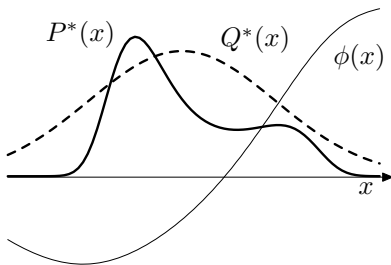
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- requires prohibitively large number of samples in high dimensions with concentrated density

importance sampling

- modify uniform sampling by introducing a sampler density $q(x) = \frac{q^*(x)}{Z_Q}$
- choose $q(x)$ simple enough that $q^*(x)$ can be sampled from, with hope that $q^*(x)$ is a reasonable approximation to $p^*(x)$



importance sampling

- adjust estimator by weighting “importance” of each sample

$$\hat{\Phi} = \frac{\sum_{r=1}^R w_r \phi(x^{(r)})}{\sum_R w_r} \quad (17)$$

where

$$w_r = \frac{p^*(x^{(r)})}{q^*(x^{(r)})} \quad (18)$$

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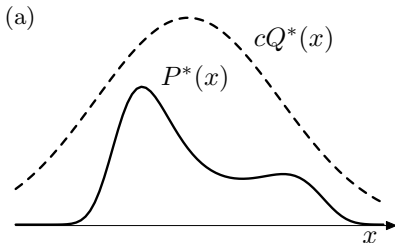
- difficult to choose “good” $q^*(x)$ as well as estimate reliability of estimator

rejection sampling

- similar to importance sampling, but proposal density strictly bounds target density, i.e.

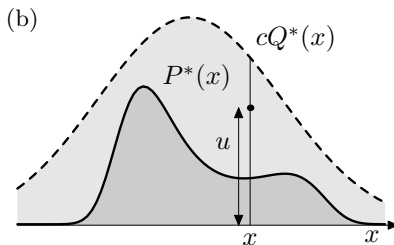
$$cq^*(x) > p^*(x), \quad (19)$$

for some known value c and all x



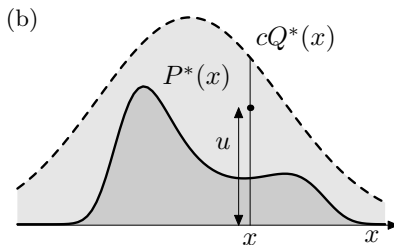
rejection sampling

- generate sample x from $q^*(x)$
- generate uniformly random number u from $[0, cq^*(x)]$
- add x to set $\{x^{(r)}\}$ if $p^*(x) > u$
- estimate expectation as $\hat{\Phi} = \frac{1}{R} \sum_{r=1}^R \phi(x^{(r)})$



rejection sampling

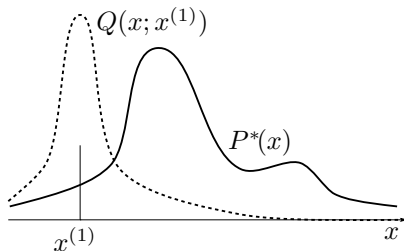
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- c often prohibitively large for poor choice of $q^*(x)$ or high dimensions

metropolis-hastings

- use a local proposal density $q(x'; x^{(t)})$, depending on current state $x^{(t)}$



- construct markov chain through state space, converging to target density
- note: proposal density needn't closely approximate target density

metropolis-hastings

- at time t , generate tentative state x' from $q(x'; x^{(t)})$
- evaluate

$$a = \frac{p^*(x^{(t)})}{p^*(x')} \frac{q(x^{(t)}; x')}{q(x'; x^{(t)})} \quad (20)$$

- if $a \geq 1$, accept the new state; else accept the new state with probability a
- if new state is rejected, set $x^{(t+1)} = x^{(t)}$

metropolis-hastings

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- if $a \geq 1$, accept the new state; else accept the new state with probability a
- if new state is rejected, set $x^{(t+1)} = x^{(t)}$
- effective for high dimensional problems, but difficult to assess “convergence” of markov chain³

³see neal (1993)

gibbs sampling

- metropolis method where proposal density is chosen as conditional distribution, i.e.

$$q(x'_i; x^{(t)}) = p\left(x_i | \{x_j^{(t)}\}_{j \neq i}\right) \quad (21)$$

- useful when joint density factorizes, as in sparse graphical model⁴

⁴see wainwright & jordan 2008

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- useful when joint density factorizes, as in sparse graphical model⁴
- similar difficulties to metropolis, but no concerns about adjustable parameters

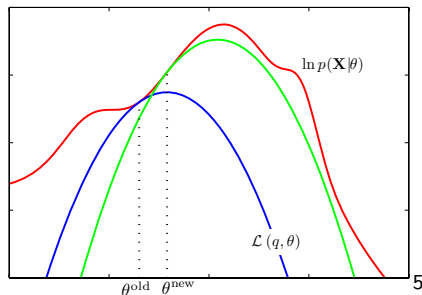
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bound optimization

- general approach: replace integration with optimization
- construct auxiliary function upper-bounded by log-evidence, maximize auxiliary function

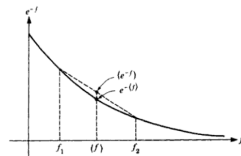


⁵image from bishop (2006)

variational bayes

- bound log of expected value by expected value of log using jensen's inequality⁶:

$$\begin{aligned}-\ln p(\mathcal{D}) &= -\ln \int d\Theta \, p(\mathcal{D}|\Theta)p(\Theta) \\&= -\ln \int d\Theta \, \frac{p(\mathcal{D}|\Theta)p(\Theta)}{q(\Theta)} q(\Theta) \\&\leq -\int d\Theta \, \ln \left[\frac{p(\mathcal{D}|\Theta)p(\Theta)}{q(\Theta)} \right] q(\Theta)\end{aligned}$$



- for sufficiently simple (i.e. factorized) approximating distribution $q(\Theta)$, right-hand side can be easily evaluated and optimized

⁶image from feynman (1972)

variational bayes

- iterative coordinate ascent algorithm provides controlled analytic approximations to posterior and evidence
- approximate posterior $q(\Theta)$ minimizes kullback-leibler distance to true posterior
- resulting deterministic algorithm is often fast and scalable

variational bayes

- iterative coordinate ascent algorithm provides controlled analytic approximations to posterior and evidence
- approximate posterior $q(\Theta)$ minimizes kullback-leibler distance to true posterior
- resulting deterministic algorithm is often fast and scalable
- complexity of approximation often limited (to, e.g., mean-field theory, assuming weak interaction between unknowns)
- iterative algorithm requires restarts, no guarantees on quality of approximation

example: a bayesian approach to network modularity



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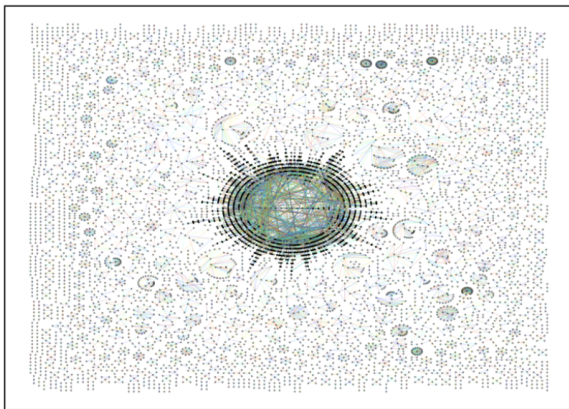
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Session P39: Applications of Complex Networks

Sponsoring Units: GSNP Chair: Narayan Menon, University of Massachusetts, Amherst Morial Convention Center - 231	
Wednesday, March 12, 2008 8:00AM - 8:12AM	P39.00001: Effects of quenched randomness on predator-prey interactions in a stochastic Lotka-Volterra lattice model Uwe C. Tauber , Ulrich Dobramysl
Wednesday, March 12, 2008 8:12AM - 8:24AM	Preview Abstract P39.00002: Dynamical Clustering in Reaction-Dispersion Processes on Complex Networks Vincent David , Marc Timme , Theo Geisel , Dirk Brockmann
Wednesday, March 12, 2008 8:24AM - 8:36AM	Preview Abstract P39.00003: Fluctuations and Food-web Structures in Individual-based Models of Biological Coevolution Per Arne Rikvold , Volkan Sevim
Wednesday, March 12, 2008 8:36AM - 8:48AM	Preview Abstract P39.00004: Metabolic disease network and its implication for disease comorbidity Deok-Sun Lee , Zoltan Oltvai , Nicholas Christakis , Albert-Laszlo Barabasi
Wednesday, March 12, 2008 8:48AM - 9:00AM	Preview Abstract P39.00005: The Human Phenotypic Disease Network Cesar Hidalgo , Nicholas Blumm , Albert-Laszlo Barabasi , Nicholas Christakis

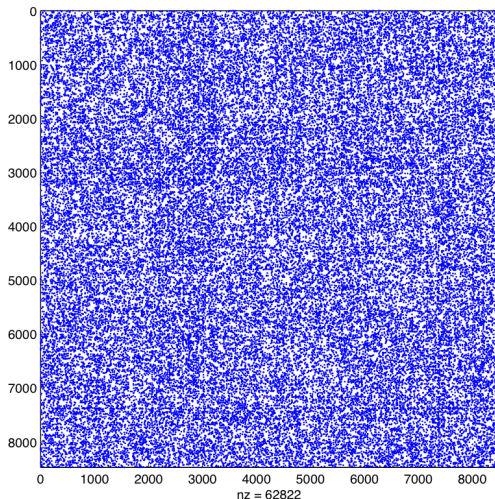
example: a bayesian approach to network modularity

nodes: authors, edges: co-authored papers

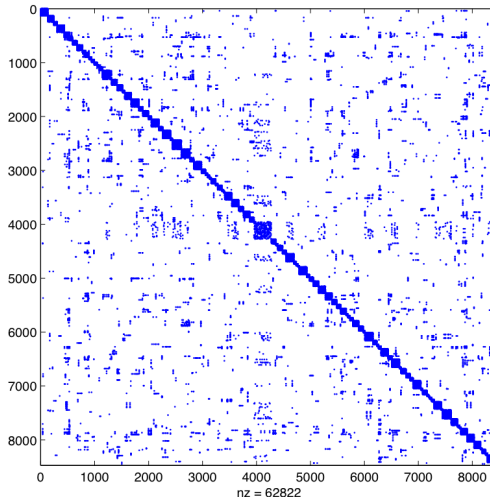


can we infer (community) structure in the giant component?

example: a bayesian approach to network modularity

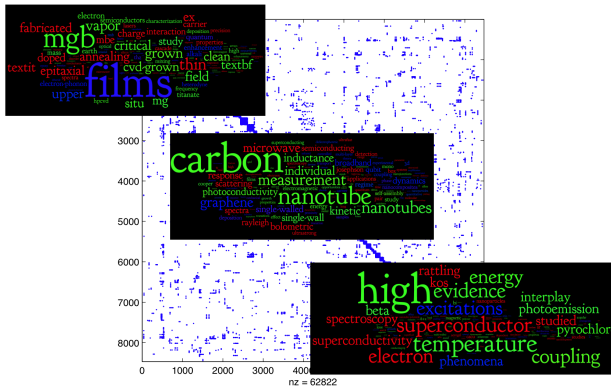


example: a bayesian approach to network modularity



example: a bayesian approach to network modularity

inferred topological communities correspond to sub-disciplines



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