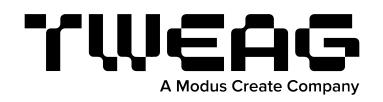
Chainsail: facilitating sampling of multimodal probability distributions

Simeon Carstens







GENERATIVE AI

MEMBERS



NOUR EL MAWASS



GUILLAUME DESFORGES



MARIA Knorps



SIMEON CARSTENS



ALOÏS COCHARD

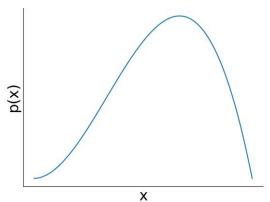


JOE NEEMAN

www.tweag.io/group/genai

Multimodal probability distributions

Unimodal

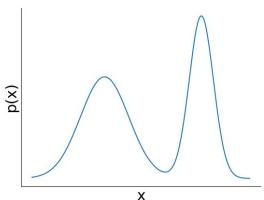


One single mode / bump / region with high probability

Examples:

* Standard distributions (Gaussian, Beta, ...)

Multimodal



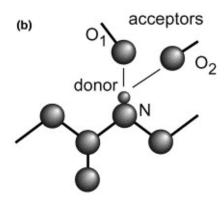
Two or more modes / bumps / regions with high probability

Examples:

- Mixtures (e.g. Gaussian mixtures in soft k-means)
- + Bayesian inference:
 - + Ambiguous data
 - Mismatch between prior distribution and likelihood

Example: ambiguous data in protein structure

Ambiguous distance restraints

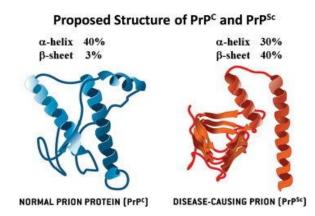


Data says that N is close to O_1 , O_2 or both. Thus,

p(x|D) is multimodal

x: 3D structure of protein, D: data

Protein misfolding

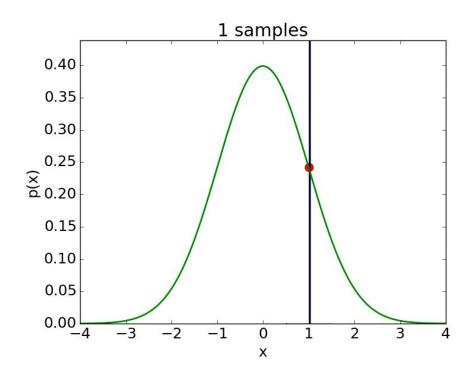


Multiple folded states for the same protein

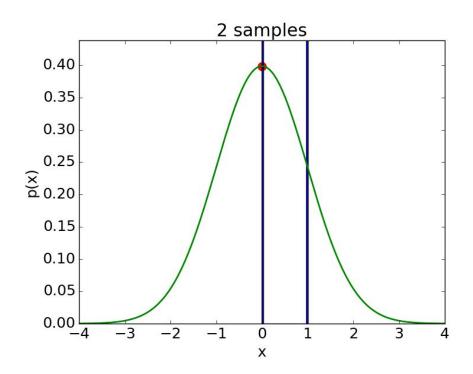
p(x) is multimodal

"Wrong" folded state → neurodegenerative disease

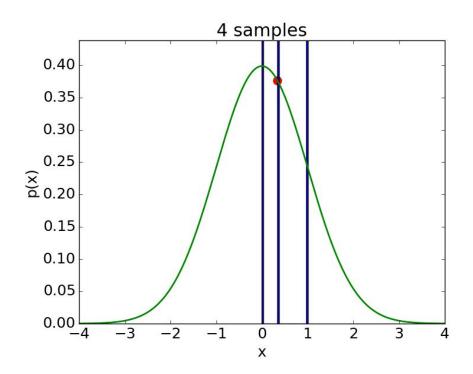
Sampling probability distributions: MCMC



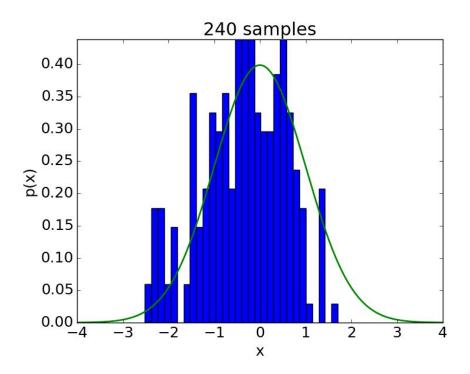
Sampling probability distributions: MCMC



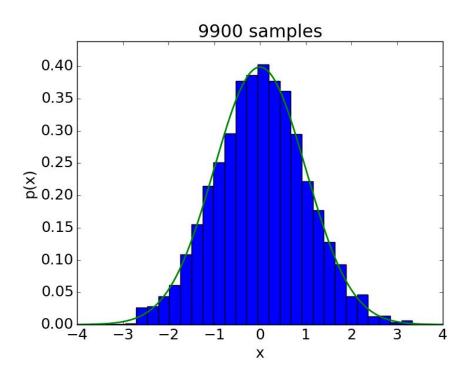
Sampling probability distributions: MCMC



Sampling probability distributions: MCMC

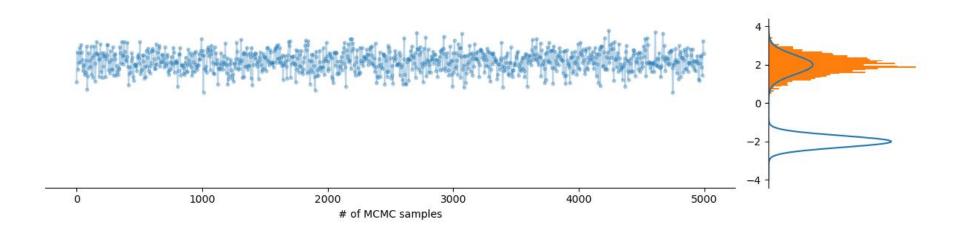


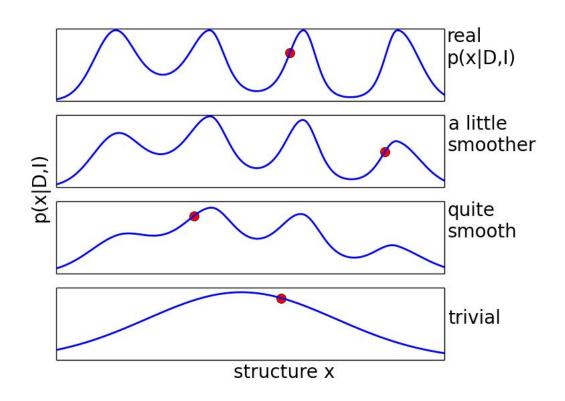
Sampling probability distributions: MCMC

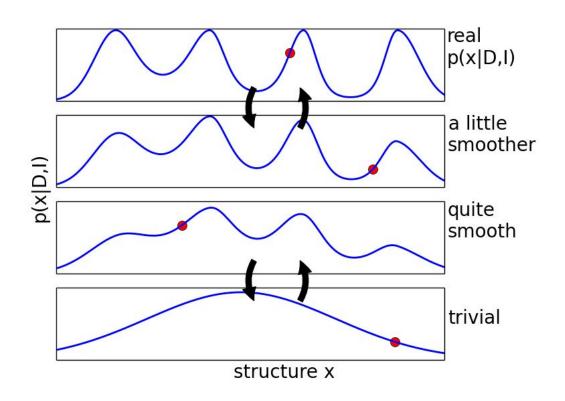


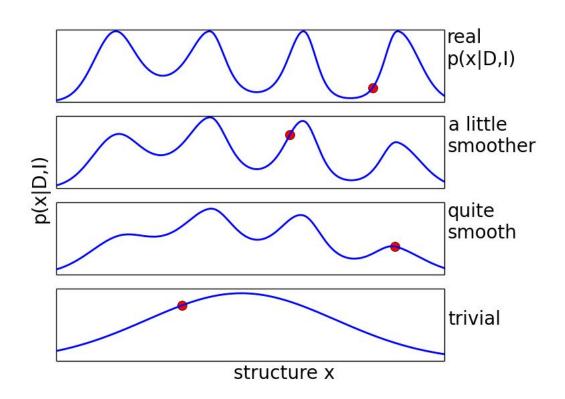
Multimodality: a formidable foe

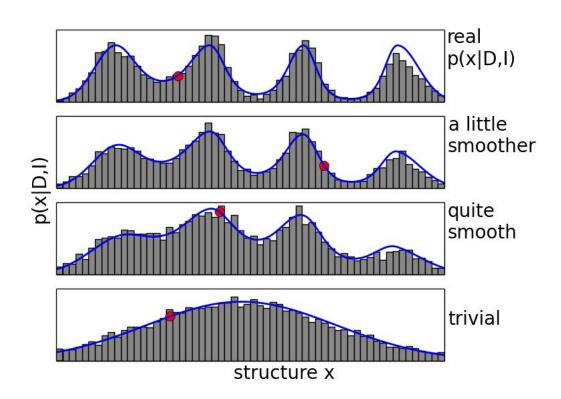
Chain can get trapped in a mode:







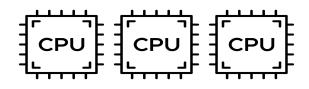




Replica Exchange: problems

Multiple chains in parallel

→ parallel computing required



"Temperatures" difficult to tune

→ experience and trial and error required



Not implemented in probabilistic programming libraries

→ reinvent the wheel for your use case



Chainsail: Replica Exchange on the



A Replica Exchange implementation that

interfaces with existing PPLs (PyMC, Stan)





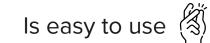
Automatically finds good temperatures $\stackrel{\circ}{=}$ \bigcirc







Scales up and down automatically



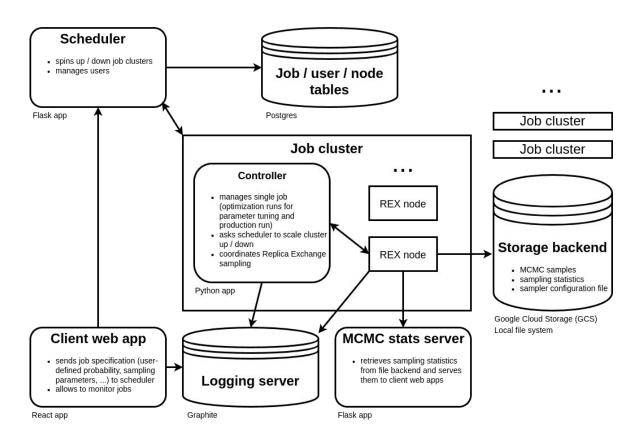


Runs barebone locally, full app locally, and in the cloud





Chainsail: architecture



Chainsail: tech stack

Backend











Frontend



Ops





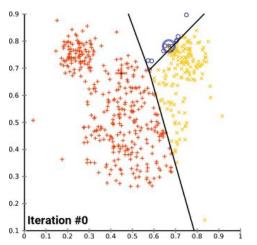






Example: soft k-means

k-means



Iterate over:

- Assign points to cluster with nearest mean
- 2. Recalculate means

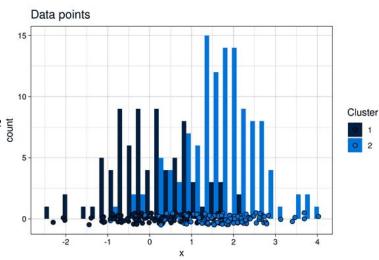
Soft k-means

Probabilistic version of k-means: use Bayesian inference to infer cluster means

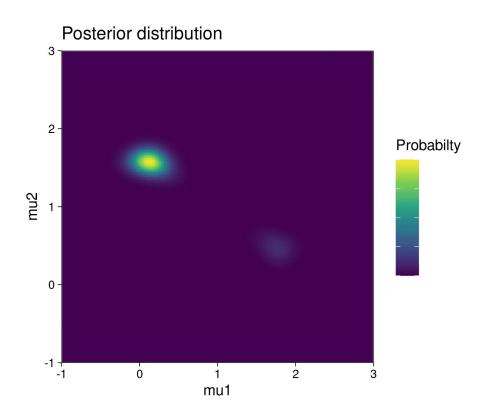
$$p(\mu_1, \mu_2 \mid data) \propto p(data \mid \mu_1, \mu_2) \times p(\mu_1, \mu_2)$$

Likelihood p(data | μ_1 , μ_2):

Assume that data was generated from Gaussian mixture with means μ_1 and μ_2 and unequal weights



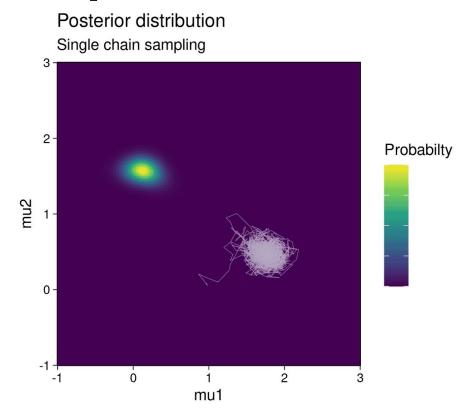
Example: soft k-means



Bimodal posterior distribution for $\mu_1, \, \mu_2$



Example: soft k-means

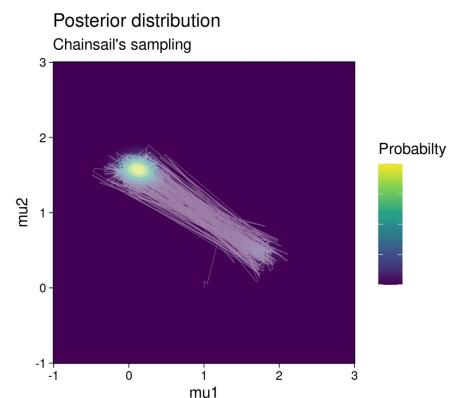


Bimodal posterior distribution for μ_1, μ_2

Hard to sample with one MCMC chain, but...



Example: soft k-means



Bimodal posterior distribution for μ_1, μ_2

Hard to sample with one MCMC chain, but...

... a piece of cake with Chainsail!



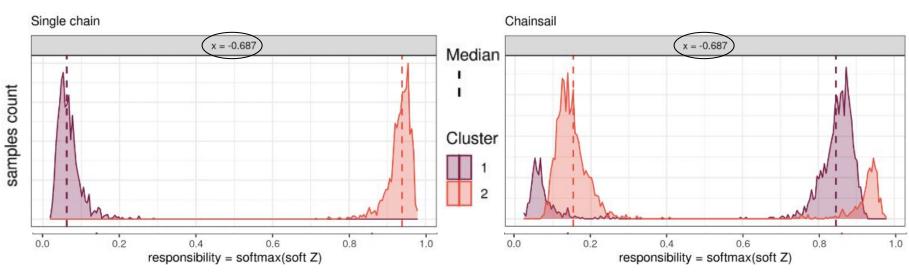
N

Example: soft k-means

Reminder:

- cluster 1 is centered at 0
- cluster 2 is centered at 1.5

Responsibility: measure of how strongly a cluster explains a data point



Wrong cluster assignments:

Low responsibility for cluster 1 High responsibility for cluster 2 Correct cluster assignments:

High responsibility for cluster 1 Low responsibility for cluster 2

... and a small probability for the inverse!

Current issues / things to do

Big picture:

+ Well, they didn't



Real-world use cases needed - reach out to us!

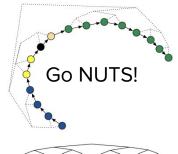
+ Little resources to continue development, but happy to maintain!

Features / functionality:

- Only very basic single-chain sampling implemented (naïve Hamiltonian Monte Carlo)
- * Temperature auto-tuning works well only for "simple" cases

Code / technical debt:

 Code is rotting (Docker images need constant updating, Python dependencies break, ...)





Credits

The Chainsail team:

Dorran Howell
Saeed Hadikhanloo
Etienne Jean
... and many others

Thank you

for your attention!

Attributions:

- + Icons: flaticon.com
- * NUTS figure:

 "The No-U-Turn Sampler [...]",

 Hoffman, M., and Gelman, A.,
 arXiv, 2011
- Ambiguous distance restraints figure:
 Nilges, M: "Ambiguous distance data [...]", Folding and Design, 1997
- Prion figure:
 "Prion Diseases as
 Transmissible Zoonotic
 Diseases", Lee, J. et al., Osong
 Public Health and Research
 Perspectives, 2013
- * Rotten fruit: istockphoto.com
- * "Build it..." meme: imgflip.com

Resources



github.com/tweag/chainsail

(code)

github.com/tweag/chainsail-resources

(examples, documentation)



tweag.io/blog -> <a> "chainsail"

http://chainsail.io

(mostly links...)