

Machine Learning for Probabilistic Modelling

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Hello! These slides were visual aids for a talk, and weren't designed to be read. I've inserted some notes here to summarize what the points were supposed to be, and to give further references.

1. Hierarchical modelling is essential. Many models contain large numbers of 'nuisance variables'. We need to *learn* how these are distributed, because if we make assumptions (including vague or so-called 'uninformative' ones), we'll simply get wrong answers. The model I discussed was referring to: "Inferring the force law in the solar-system from a snapshot", Bovy et al., 2010.

<http://arxiv.org/abs/0903.5308>

More thoughts and references on hierarchical modelling are in my discussion <http://homepages.inf.ed.ac.uk/imurray2/pub/11catchup/catchup.pdf> of <http://dx.doi.org/10.1111/j.1467-9868.2011.01025.x>

Hierarchical models can be hard to infer. Example:

<http://homepages.inf.ed.ac.uk/imurray2/pub/10hypers/>

2. Real-world models have a lot of messy detail:

- 1) complicated instrument-error distributions we may not care about;
- 2) theory encoded in expensive-to-run simulations.

Machine Learning can help.

If we don't want wrong models (point 1.) we need to learn from large amounts of data about our instruments, and from simulation data describing our theories. I've been involved in a series of papers on flexible black-box probabilistic models that could be used here:

<http://homepages.inf.ed.ac.uk/imurray2/pub/11nade/>

<http://homepages.inf.ed.ac.uk/imurray2/pub/13rnade/>

<http://homepages.inf.ed.ac.uk/imurray2/pub/14dnade/>

4. Probabilistic inference methods need extending.

Approximate inference is a heavily-mined and active area. Getting up-to-speed and finding a niche is challenging. However, work in this area is important. In deep and wide graph structures, with billions of observations in some of the plates, it's hard to do fully Bayesian inference.

Some of my work has been on identifying common small inference problems, which are usually only part of an analysis, and developing easier-to-use inference methods for them. E.g.

<http://homepages.inf.ed.ac.uk/imurray2/pub/10ess/>

I'm now also interested in developing easier-to-use methods to summarize and communicate the results of local inferences across large models. I believe the way forward is fitting flexible representations of beliefs, by combining machine learning methods and approximate inference algorithms.

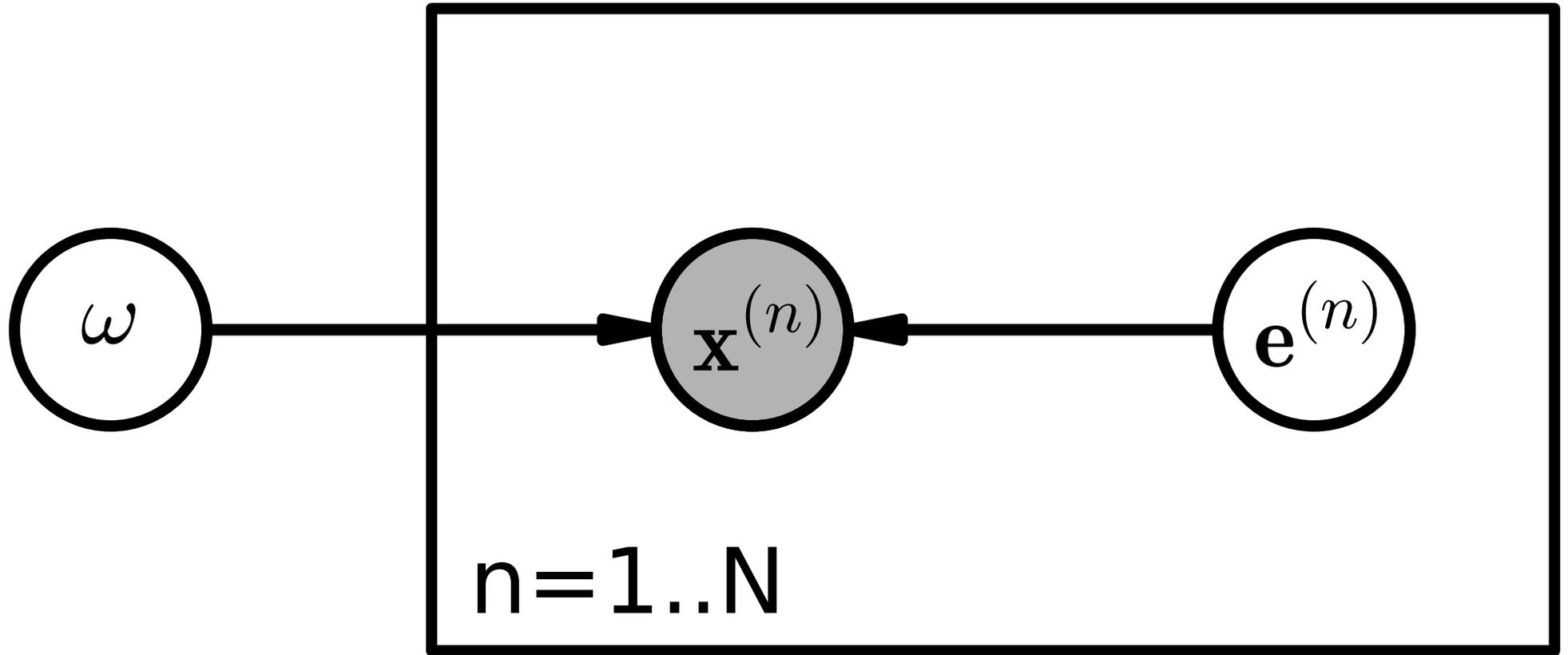
Acceleration law around the sun

$$a(r) = -A \left(\frac{r}{r_0} \right)^{-\alpha}$$

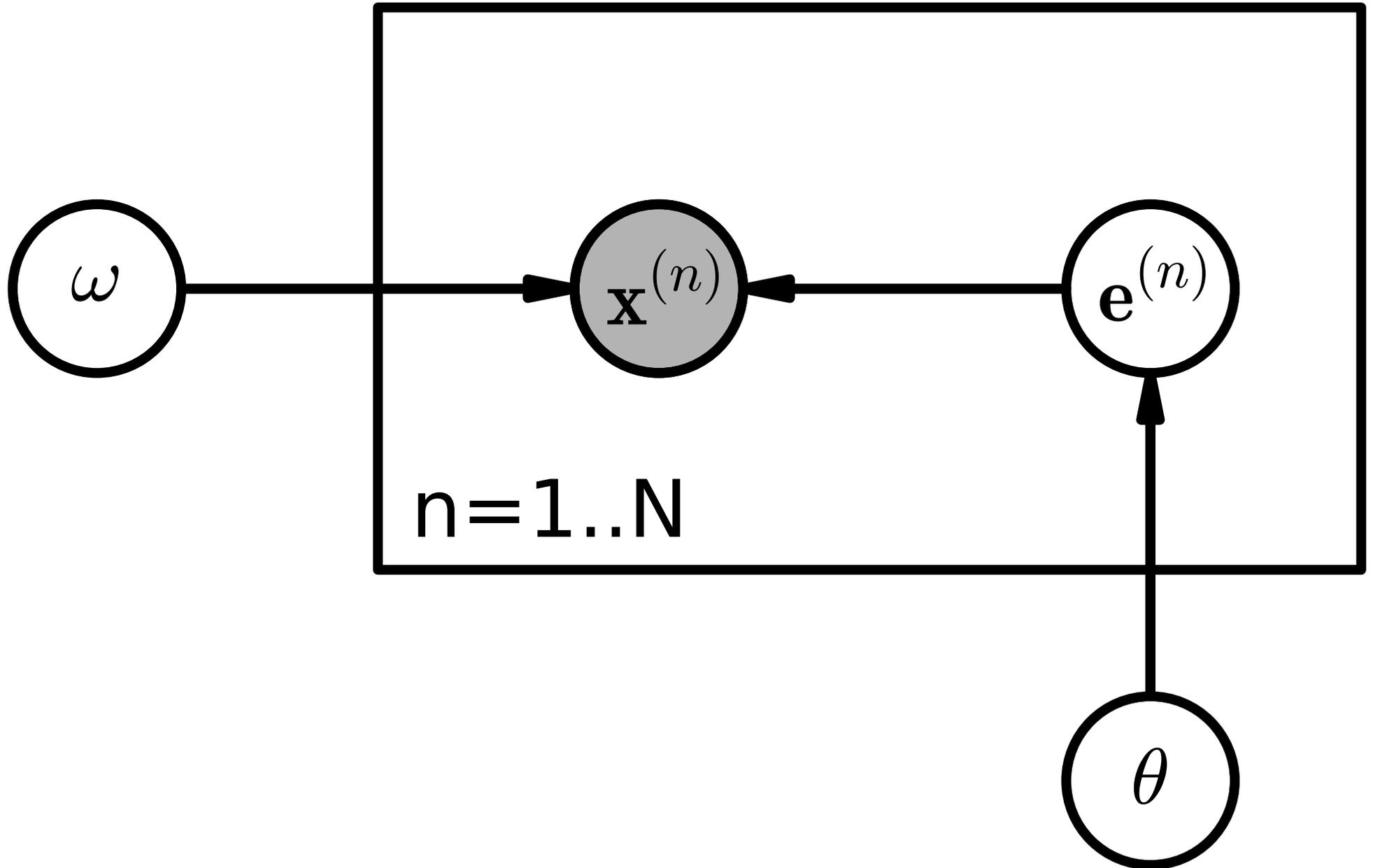
From a snapshot:

8 planet positions and velocities

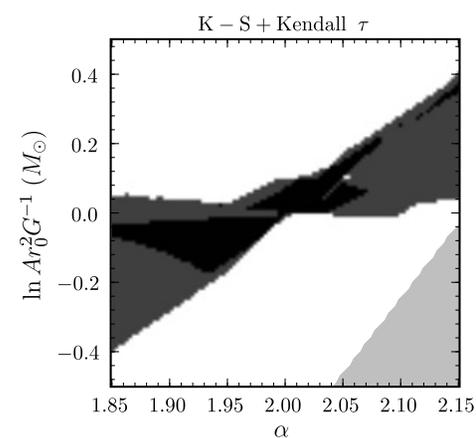
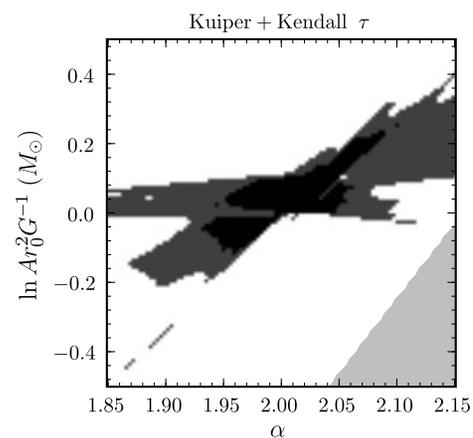
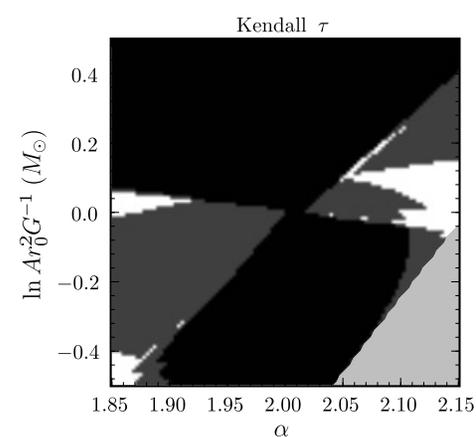
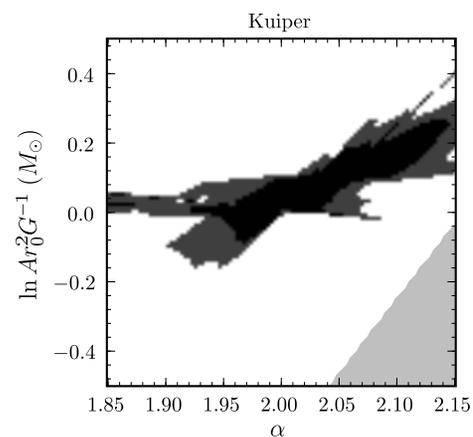
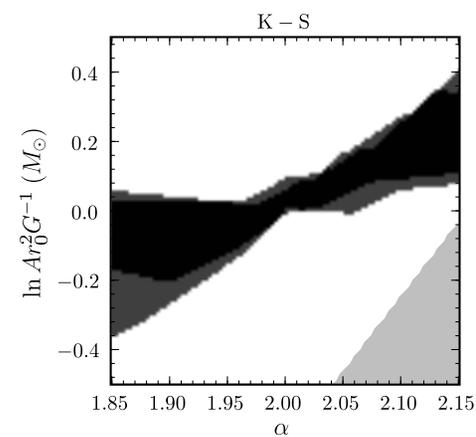
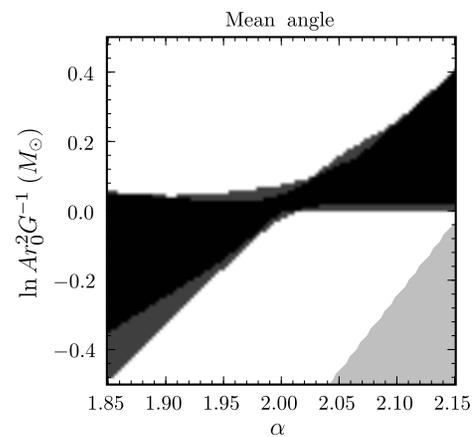
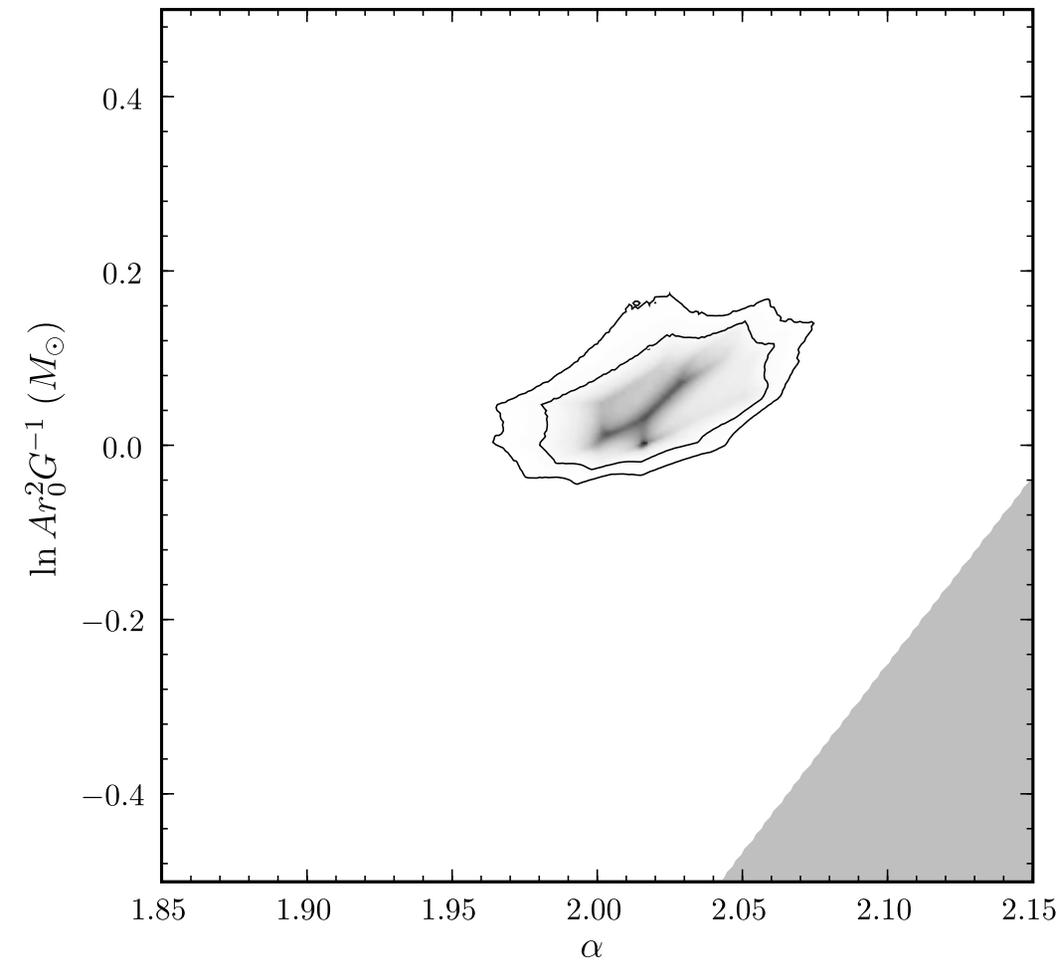
Graphical model



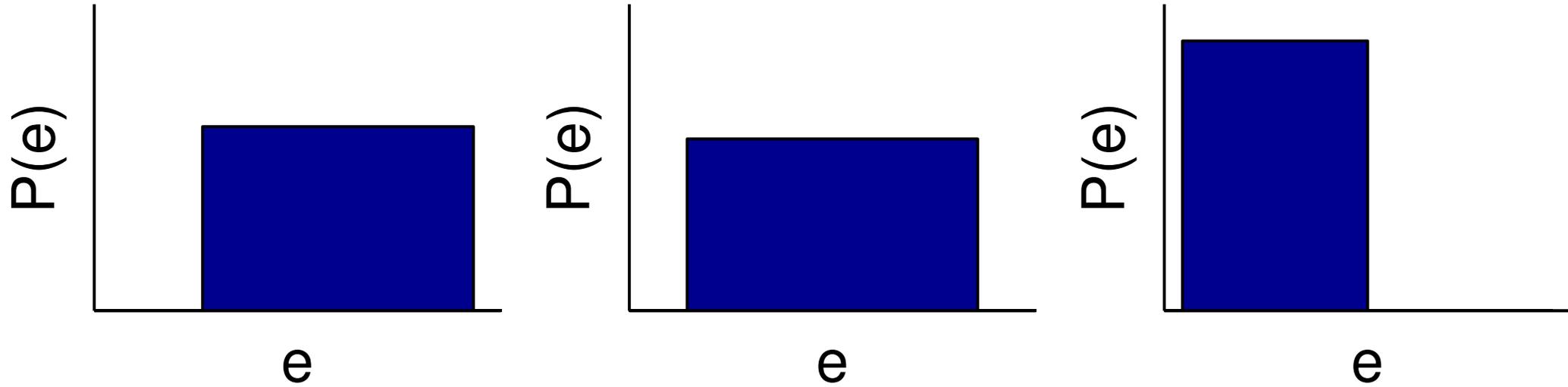
Hierarchical graphical model



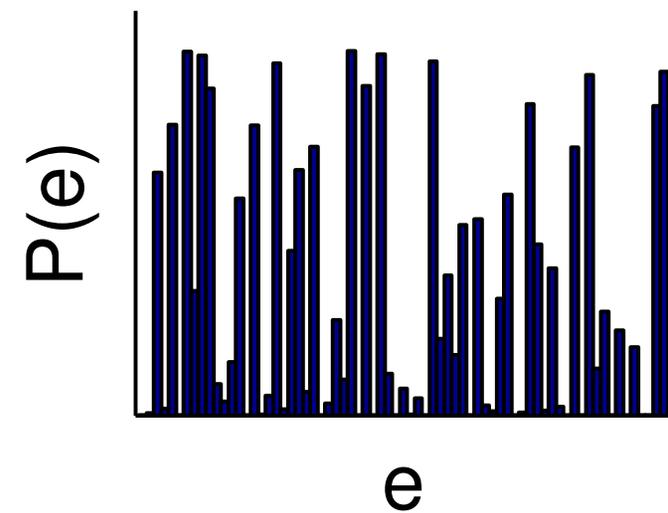
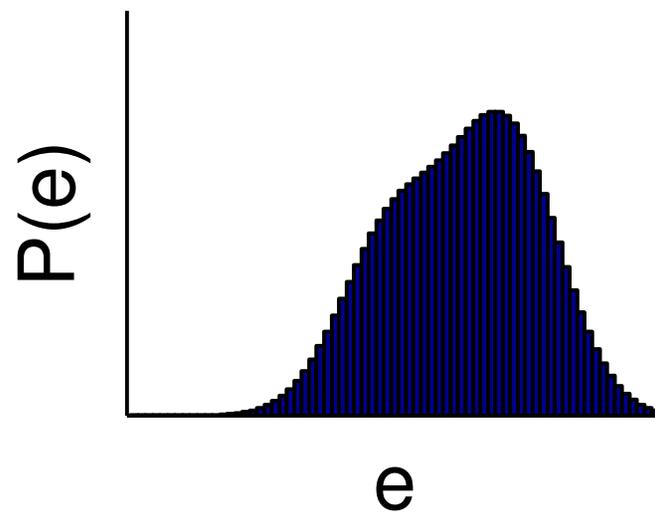
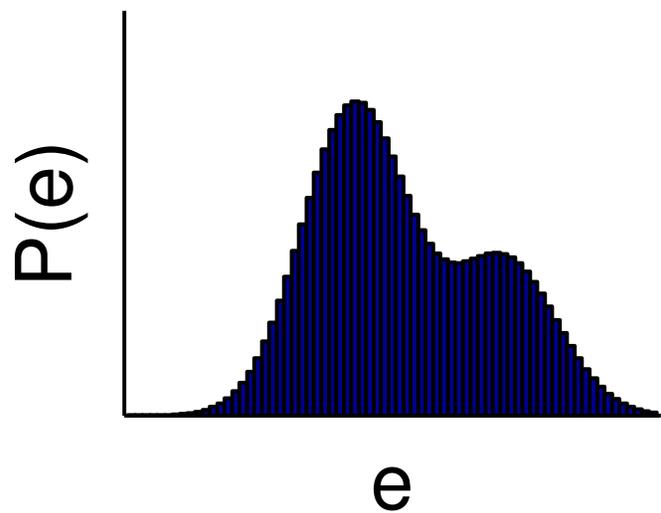
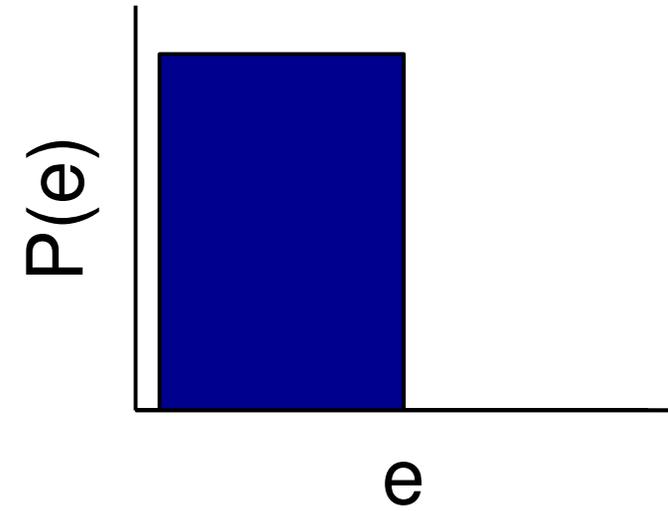
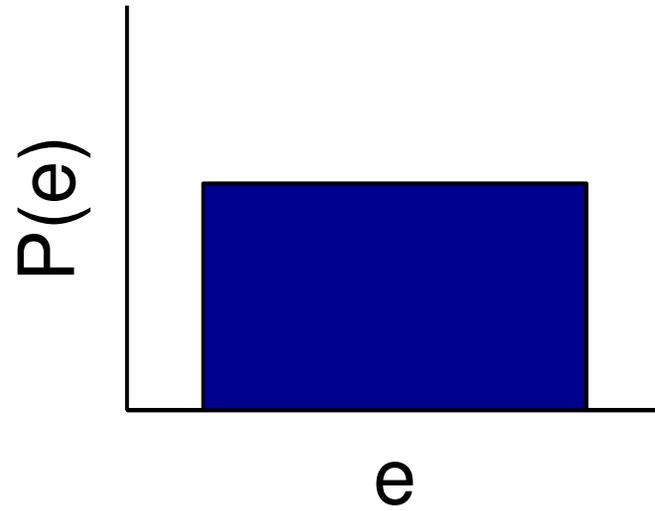
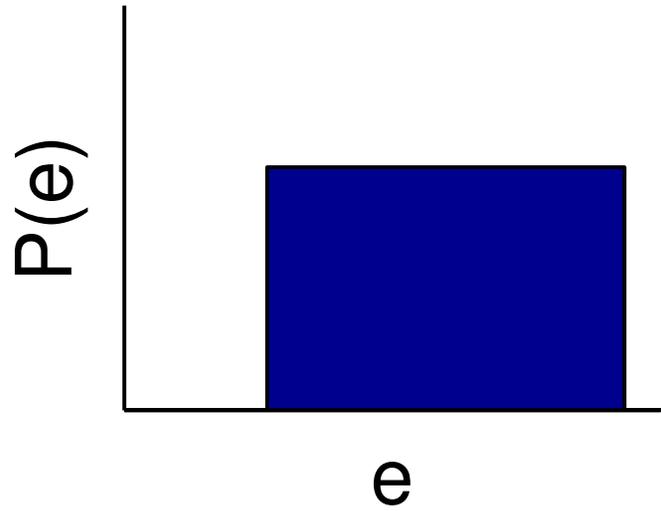
Inferences about the Sun



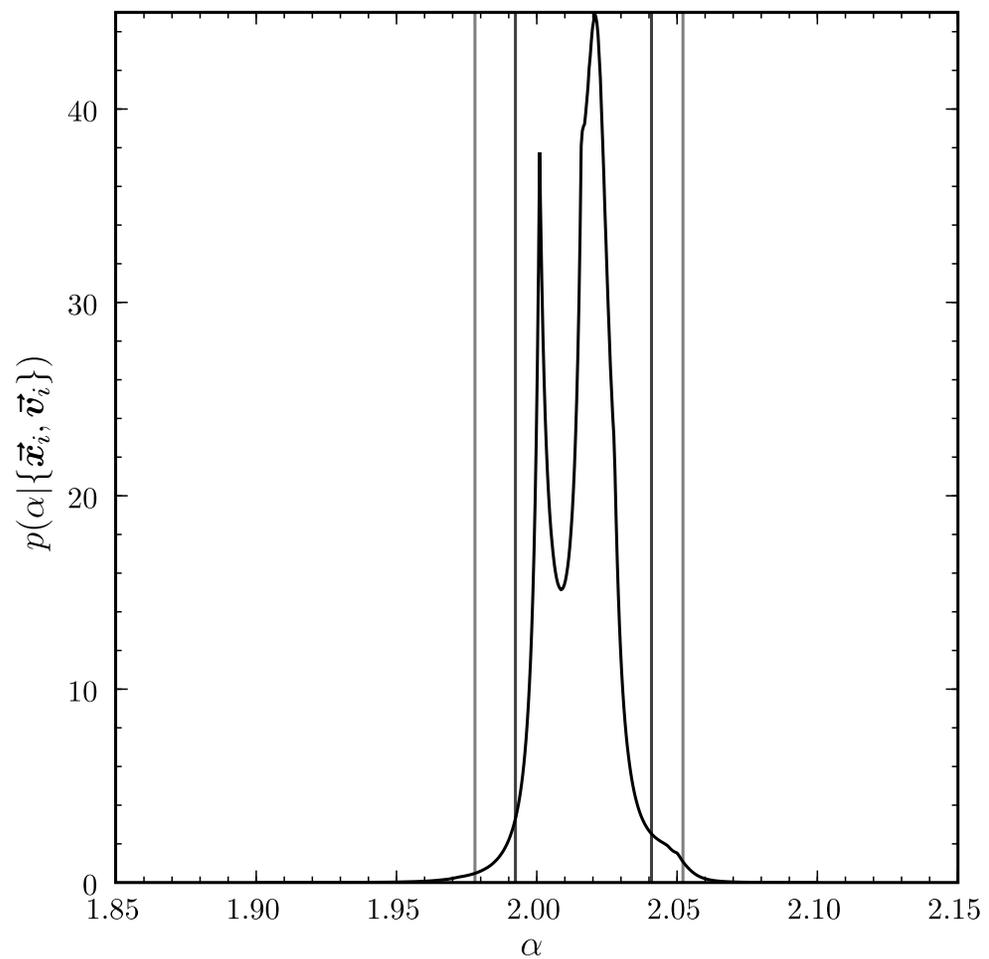
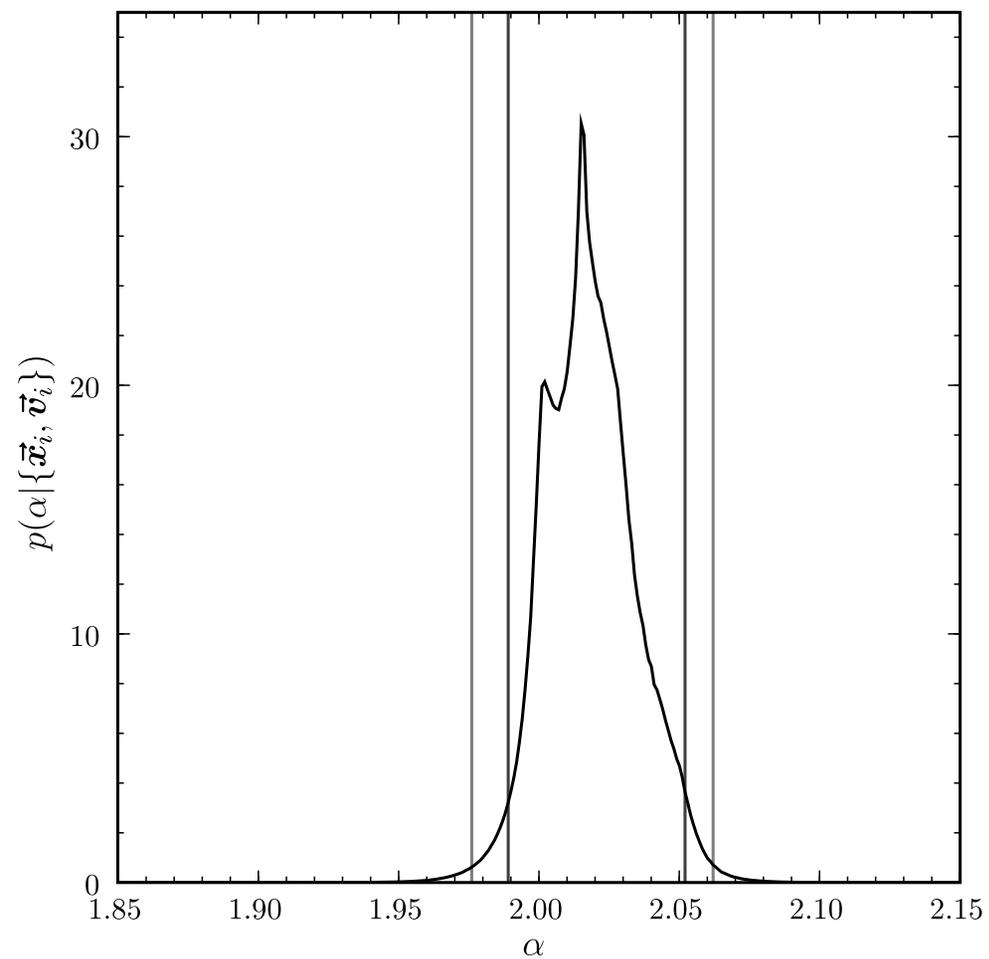
Priors on nuisance distributions

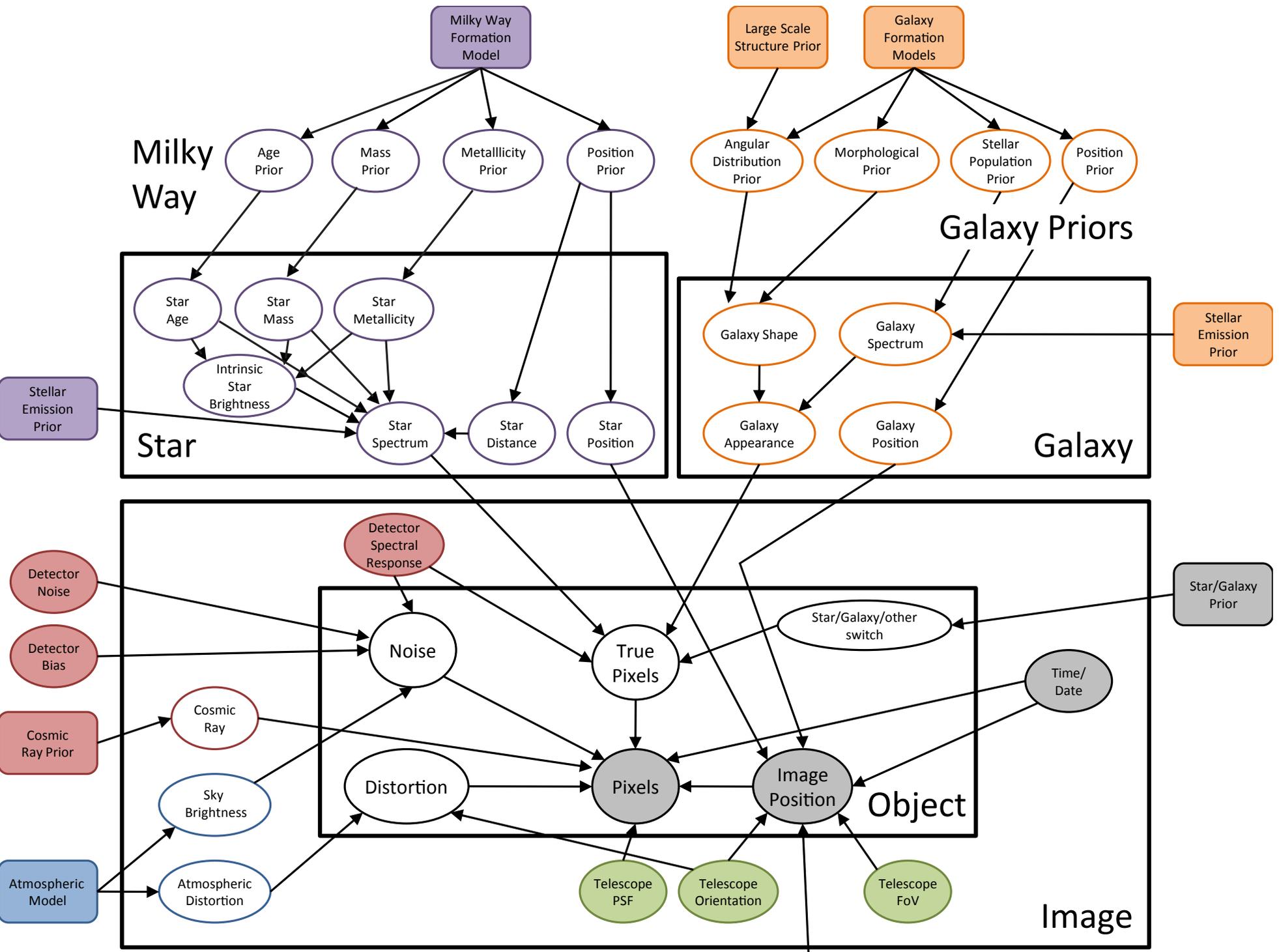


Priors on nuisance distributions



Gravitational exponent





Key: Telescope / Atmosphere / Detector / Star / Galaxy

Machine Learning?

— **Density estimation**

Neural networks and Gaussian processes

— **Inference methods**

Statistical methods: MCMC, etc.

Learning: recognition networks

Representations: communicating results

Appendix Slides

Snapshot of the solar-system

Model for the sun: $\omega = \{\log A, \alpha\}$

Acceleration law, $a(r) = -A [r/r_0]^{-\alpha}$

Model for each planet:

$$\log \epsilon_n \sim p_\epsilon(\cdot | \theta_\epsilon)$$

binding energy

$$e_n \sim p_e(\cdot | \theta_e)$$

radial asymmetry

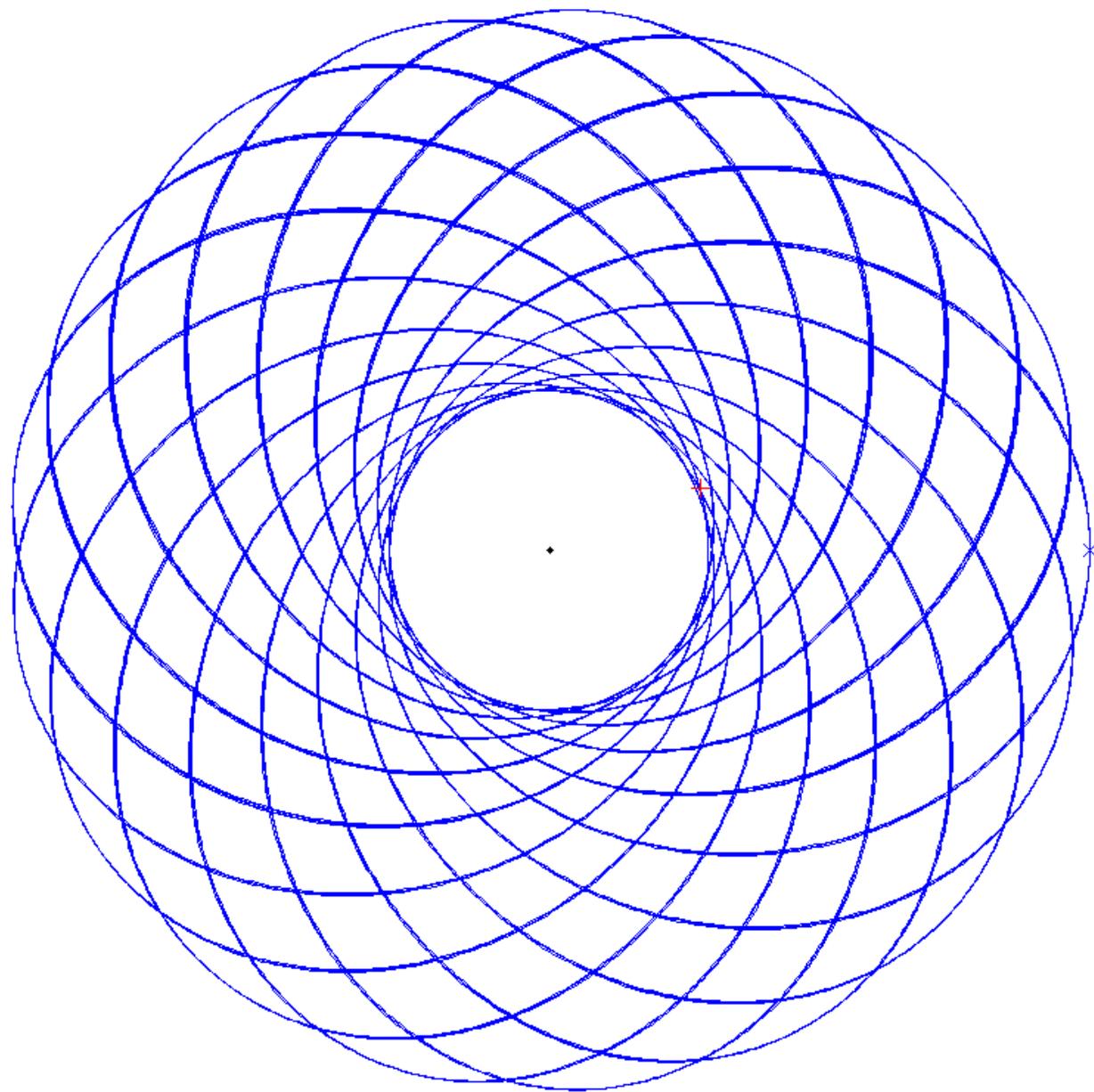
$$f_n \sim \text{Uniform}[0, 1]$$

fraction of time through orbit

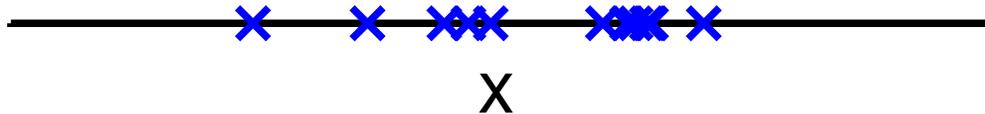
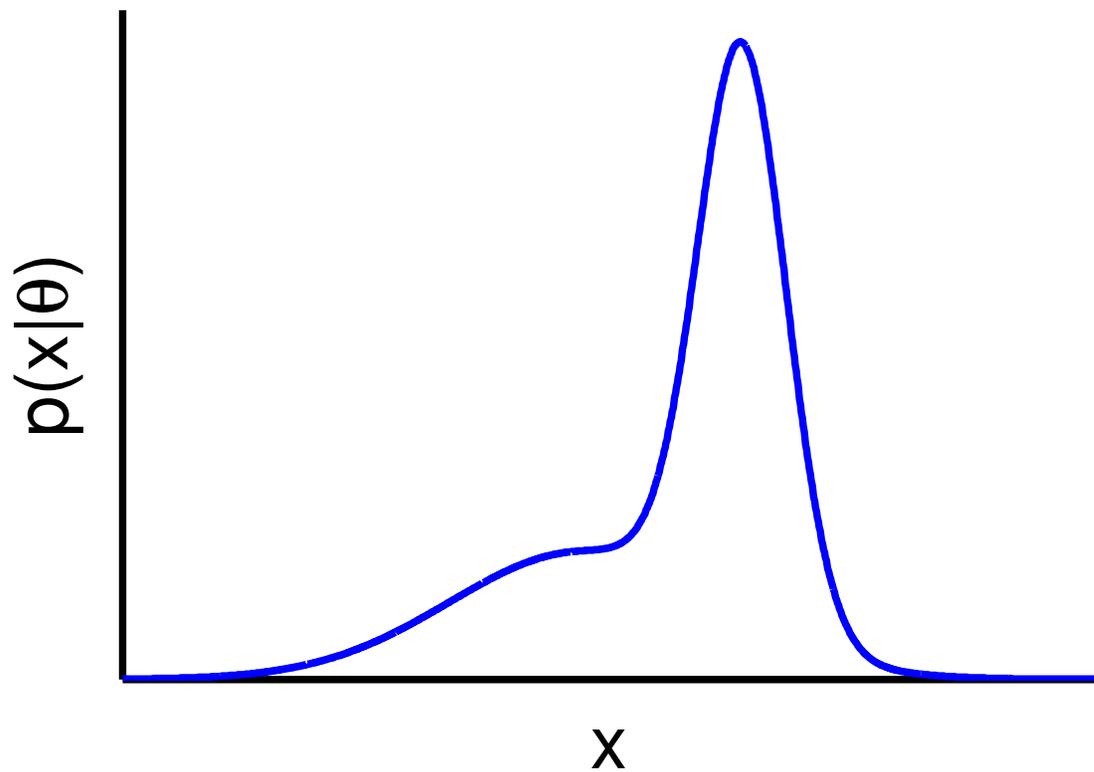
from aphelion (say), t/T_{orbit}

Observations in sky relate to:

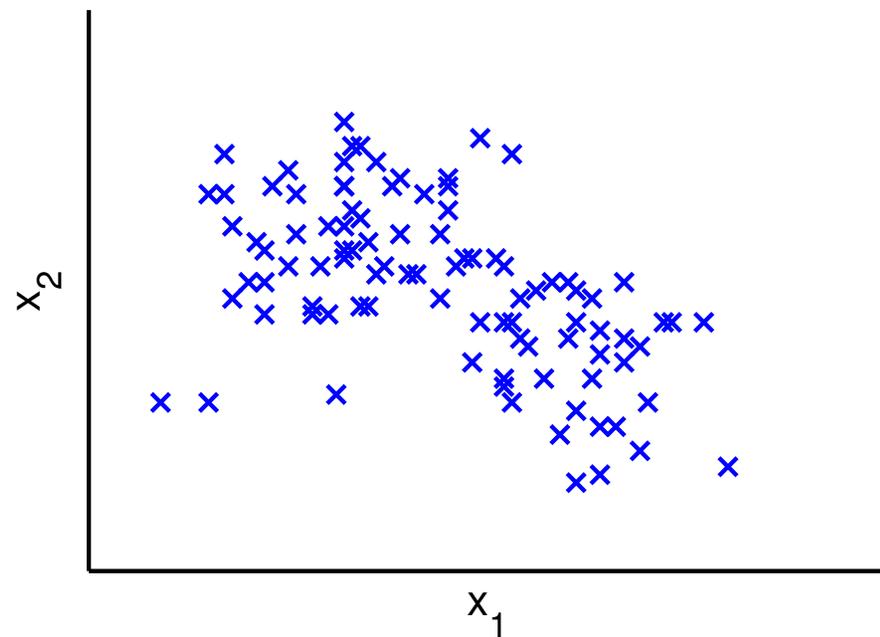
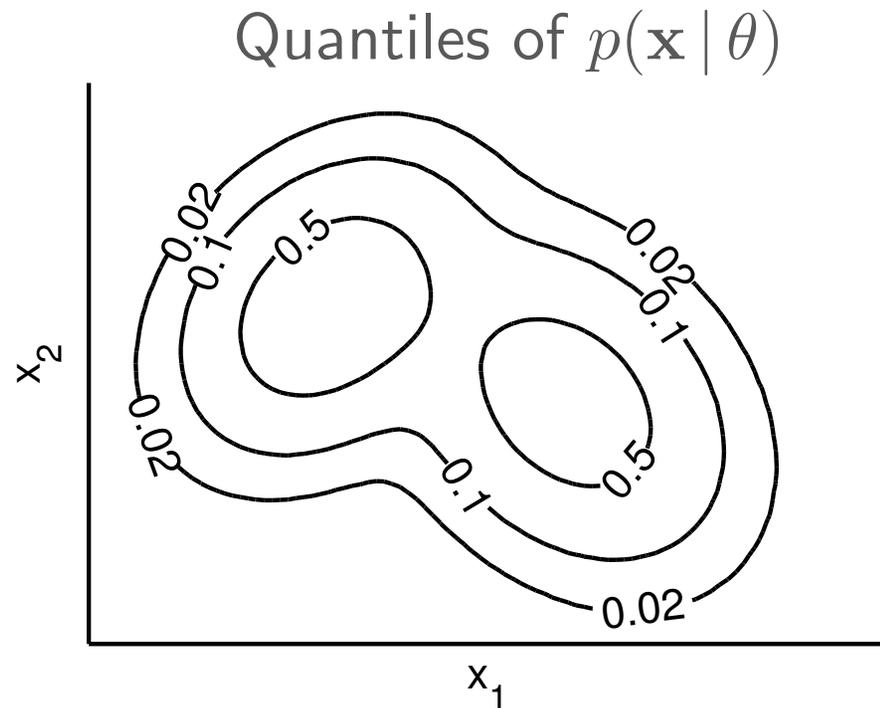
r, v_r, v_t : radial distance, radial velocity, transverse velocity



Density estimation



Task: $\{\mathbf{x}^{(n)}\} \rightarrow \theta$



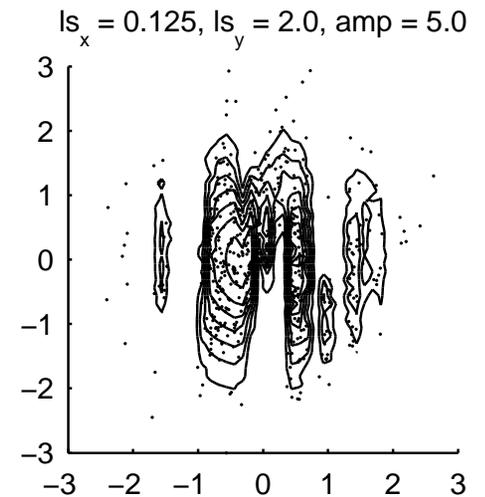
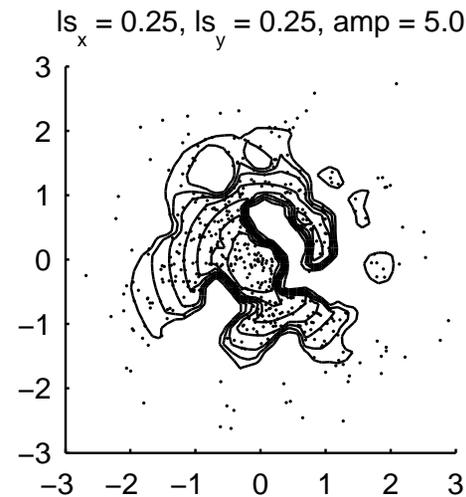
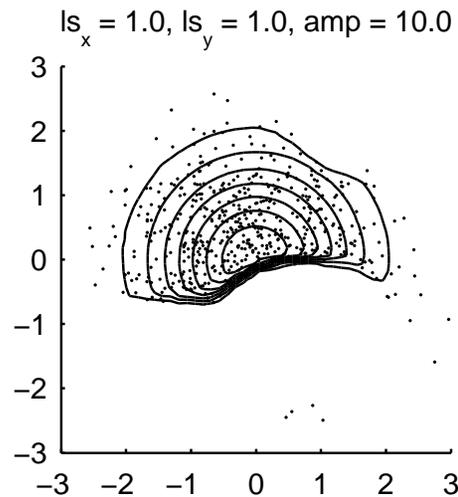
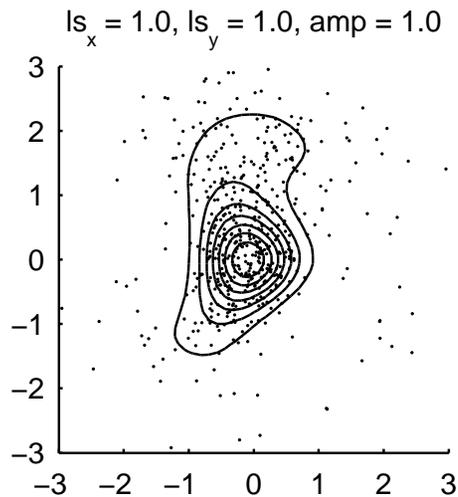
GP Density estimation

$$p(x|\mathbf{f}) = \frac{1}{\mathcal{Z}(\mathbf{f})} \Phi(\mathbf{f}(x)) \pi(x)$$

$$\mathbf{f} \sim \mathcal{GP}$$

Φ = sigmoidal function

π = base measure



Gaussian Process Density Sampler

Adams, Murray and MacKay (2009).

Modelling via the Chain Rule


$$P(x_1)$$

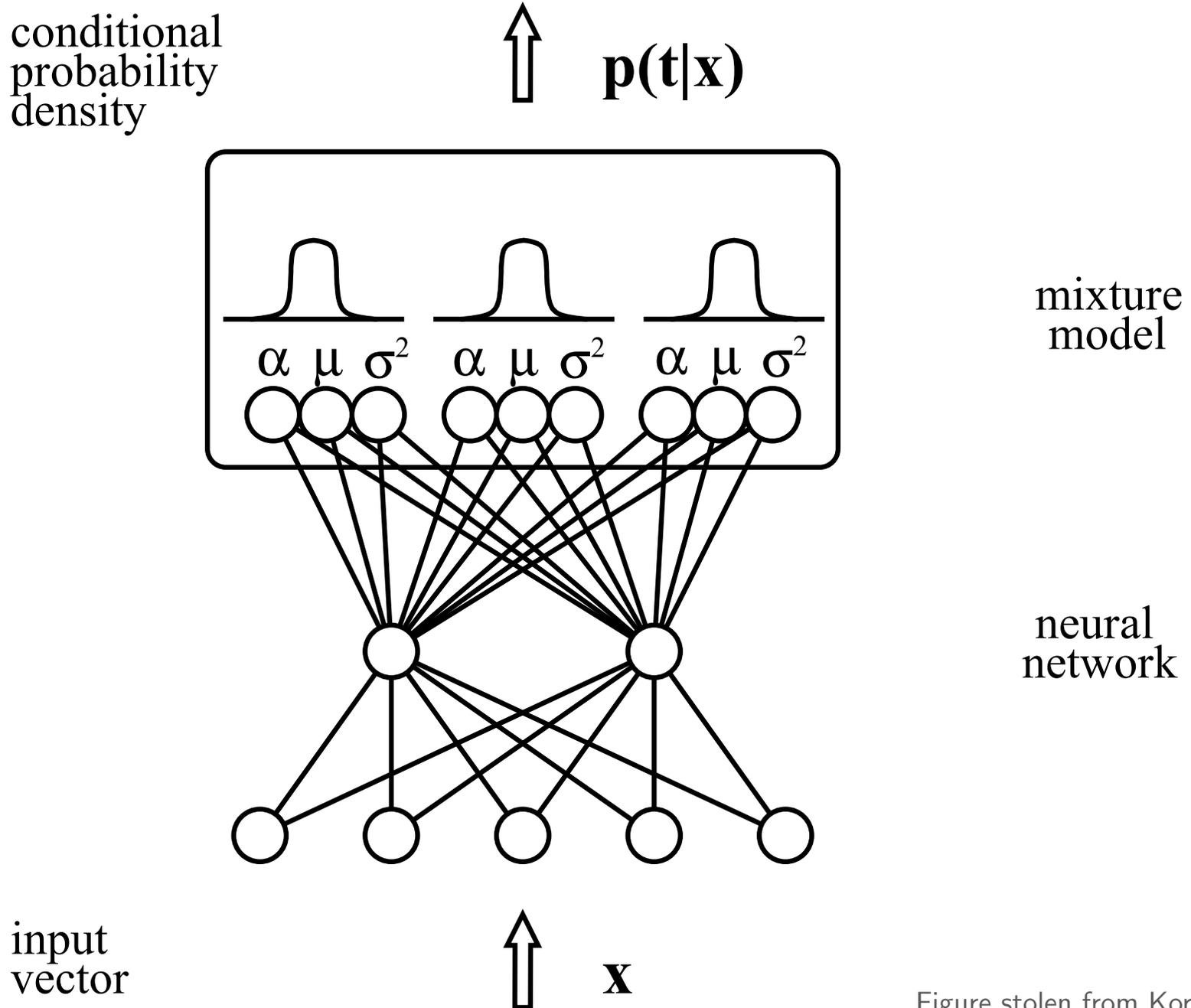

$$P(x_2 | x_1)$$


$$P(x_3 | x_1, x_2)$$

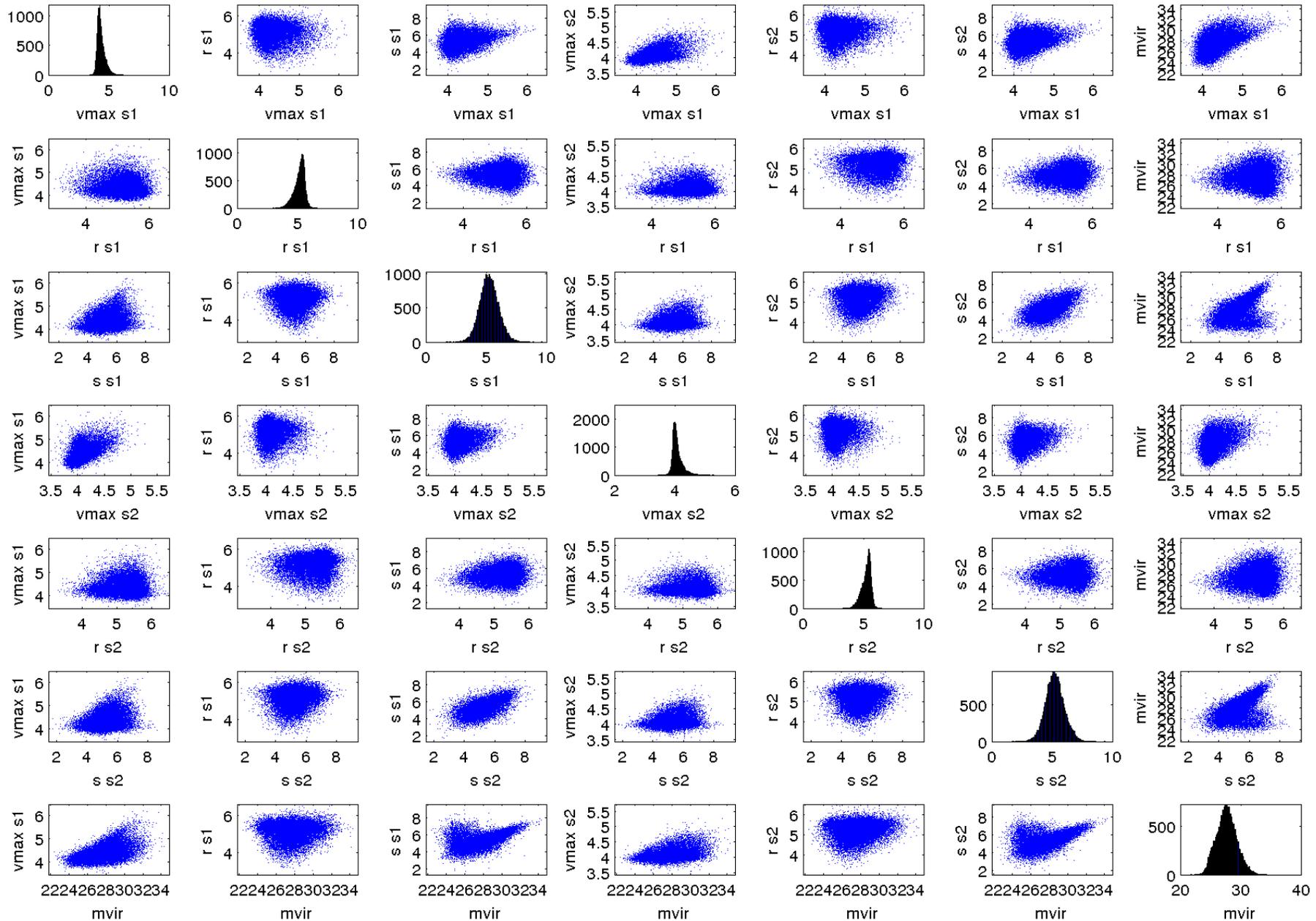

$$P(x_4 | x_1, x_2, x_3)$$

$$P(\mathbf{x}) = P(x_1) \prod_{k=2}^K P(x_k | \mathbf{x}_{<k})$$

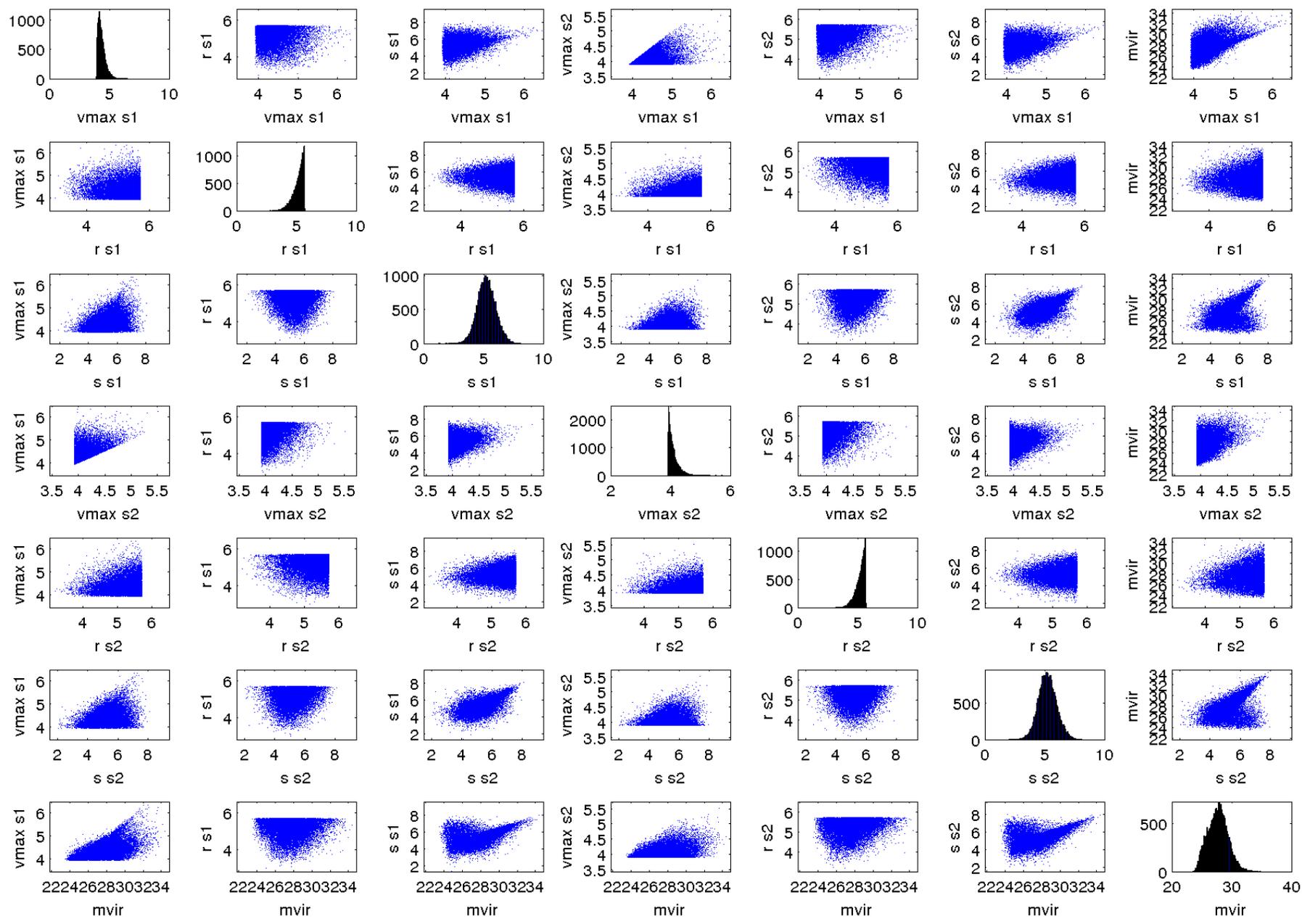
Mixture Density Networks



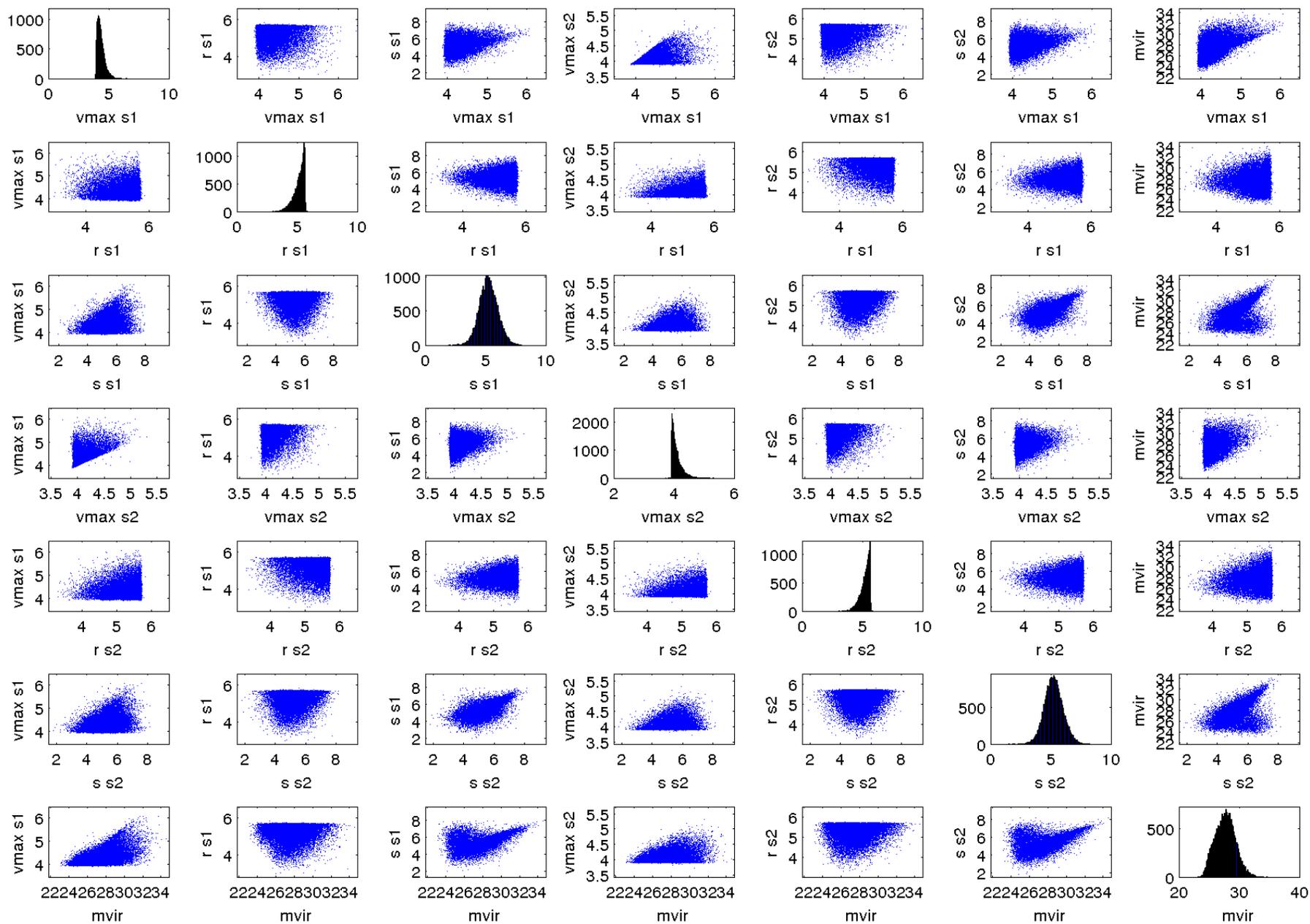
Mixture of Gaussian samples



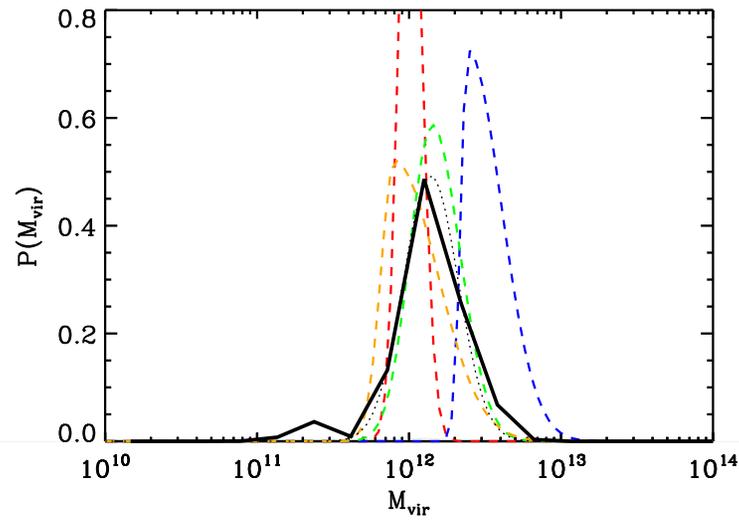
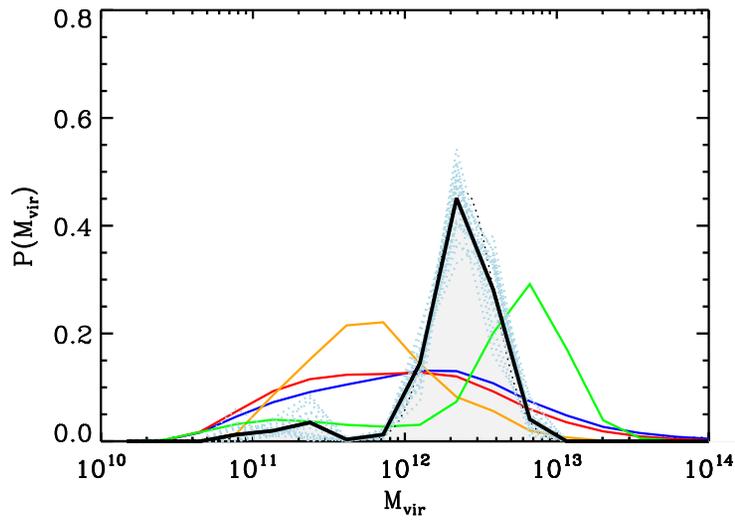
Simulation samples



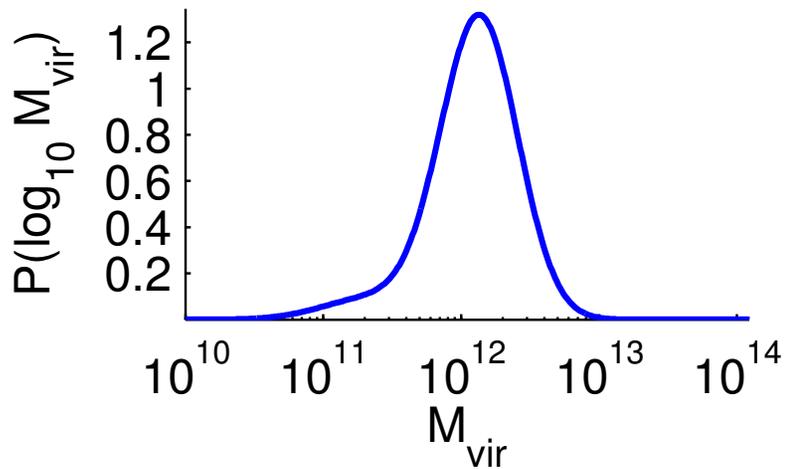
AMDN samples



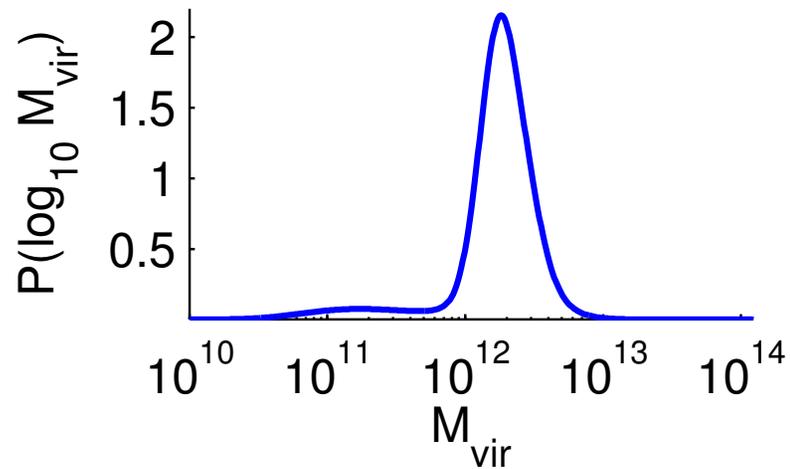
Results of inference



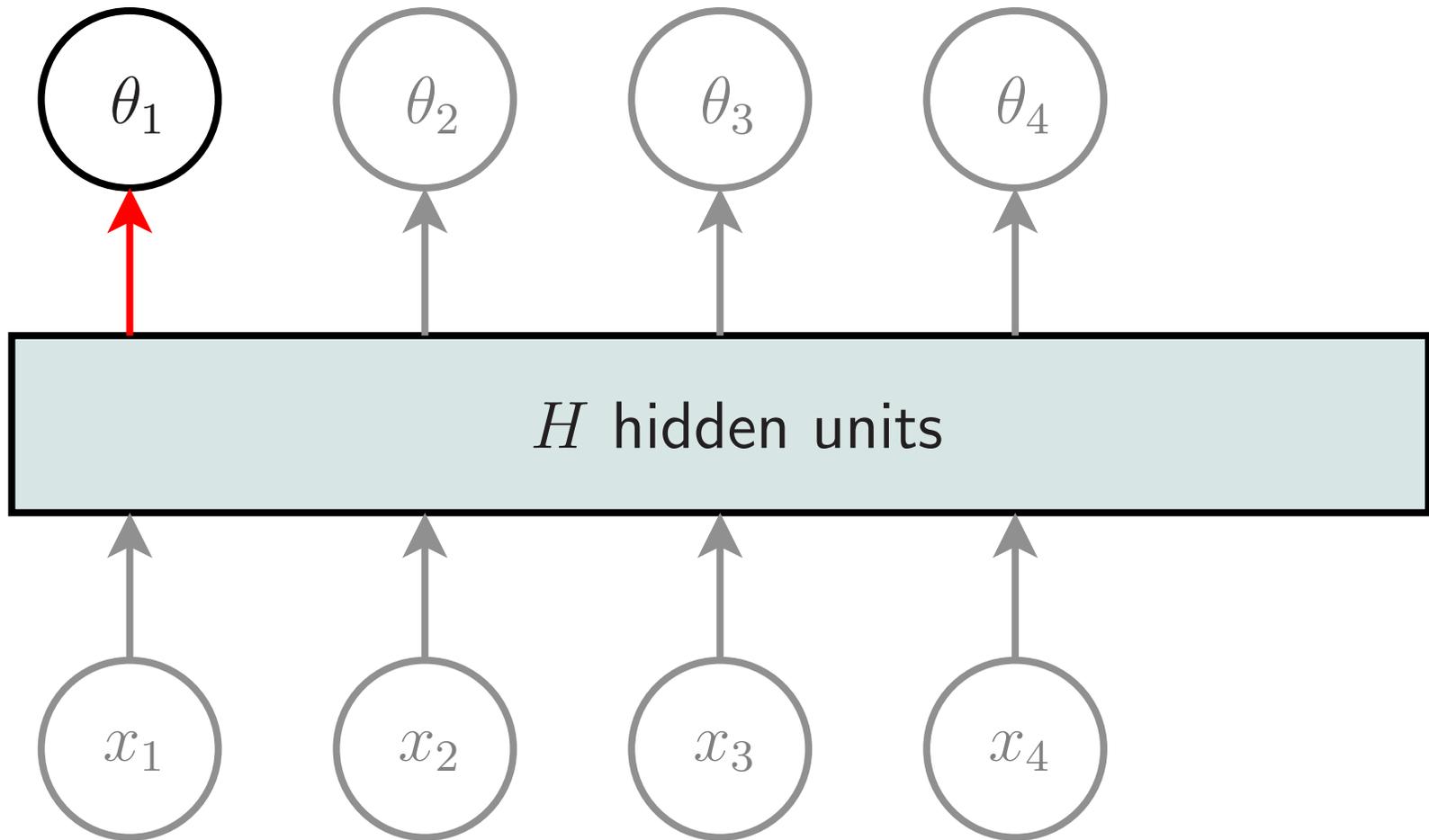
Prior modeled with MoG



Prior modeled with AMDN

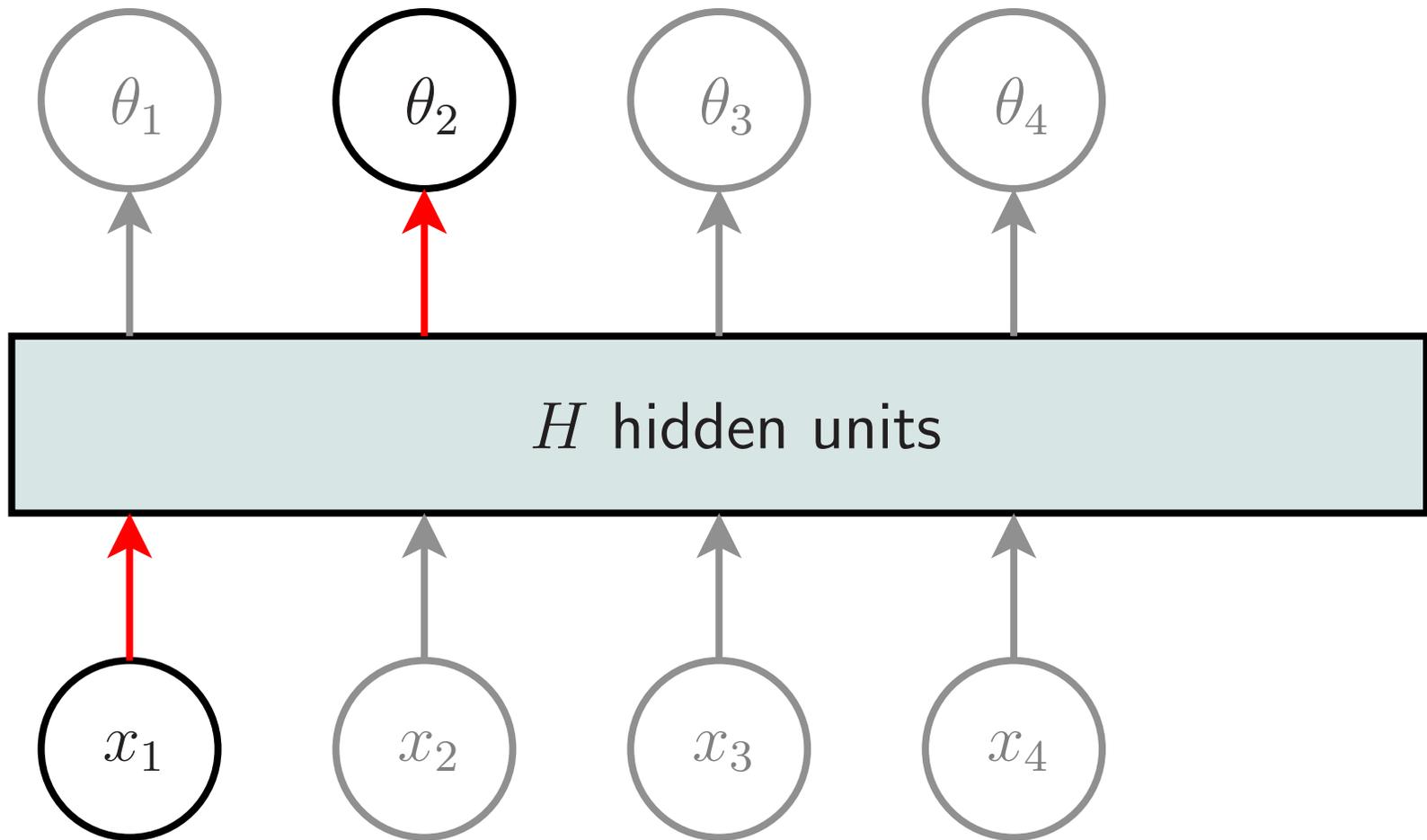


Sequential activation



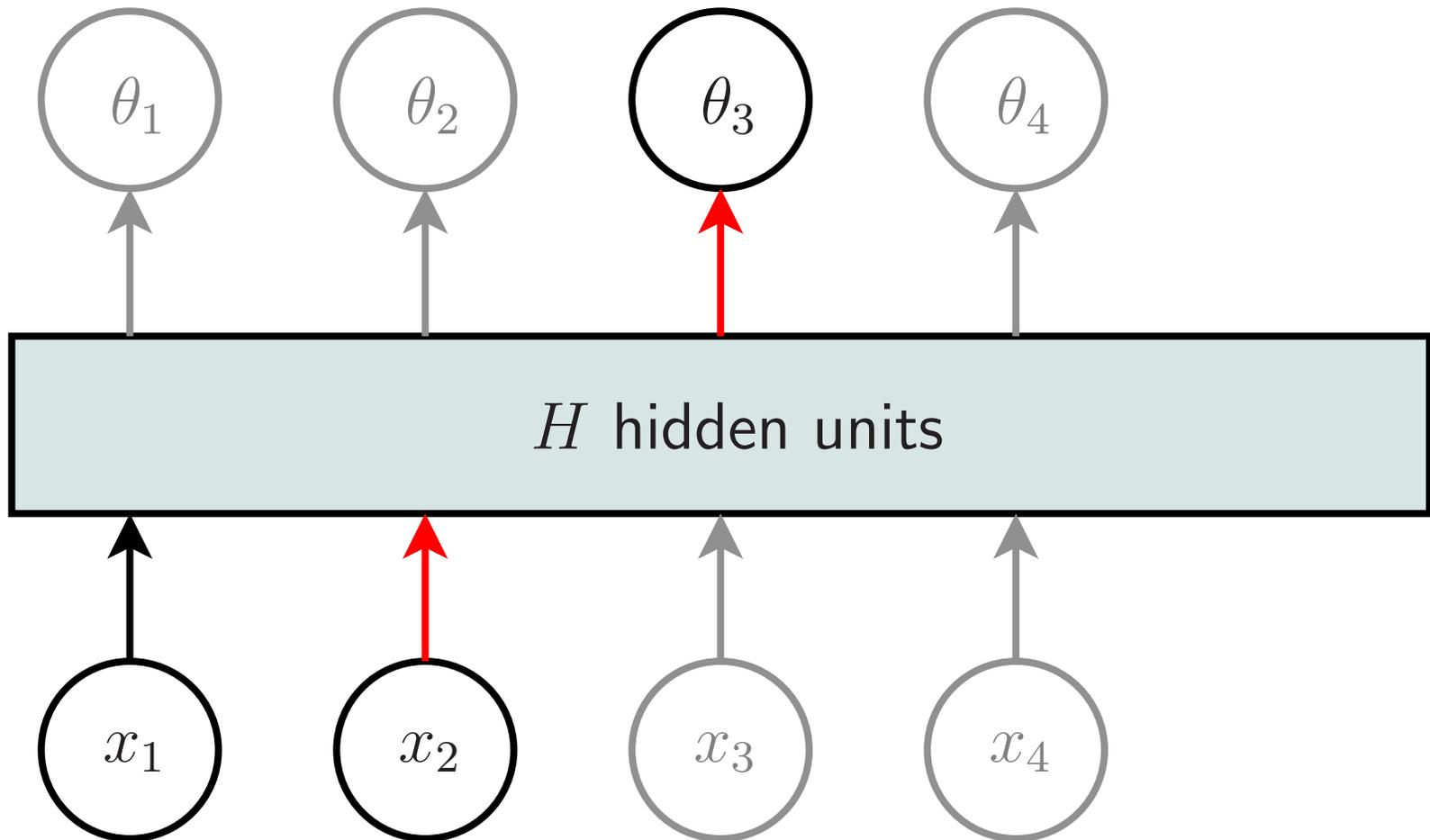
$$P(\mathbf{x}) = P(x_1 | \theta_1) \dots$$

Sequential activation



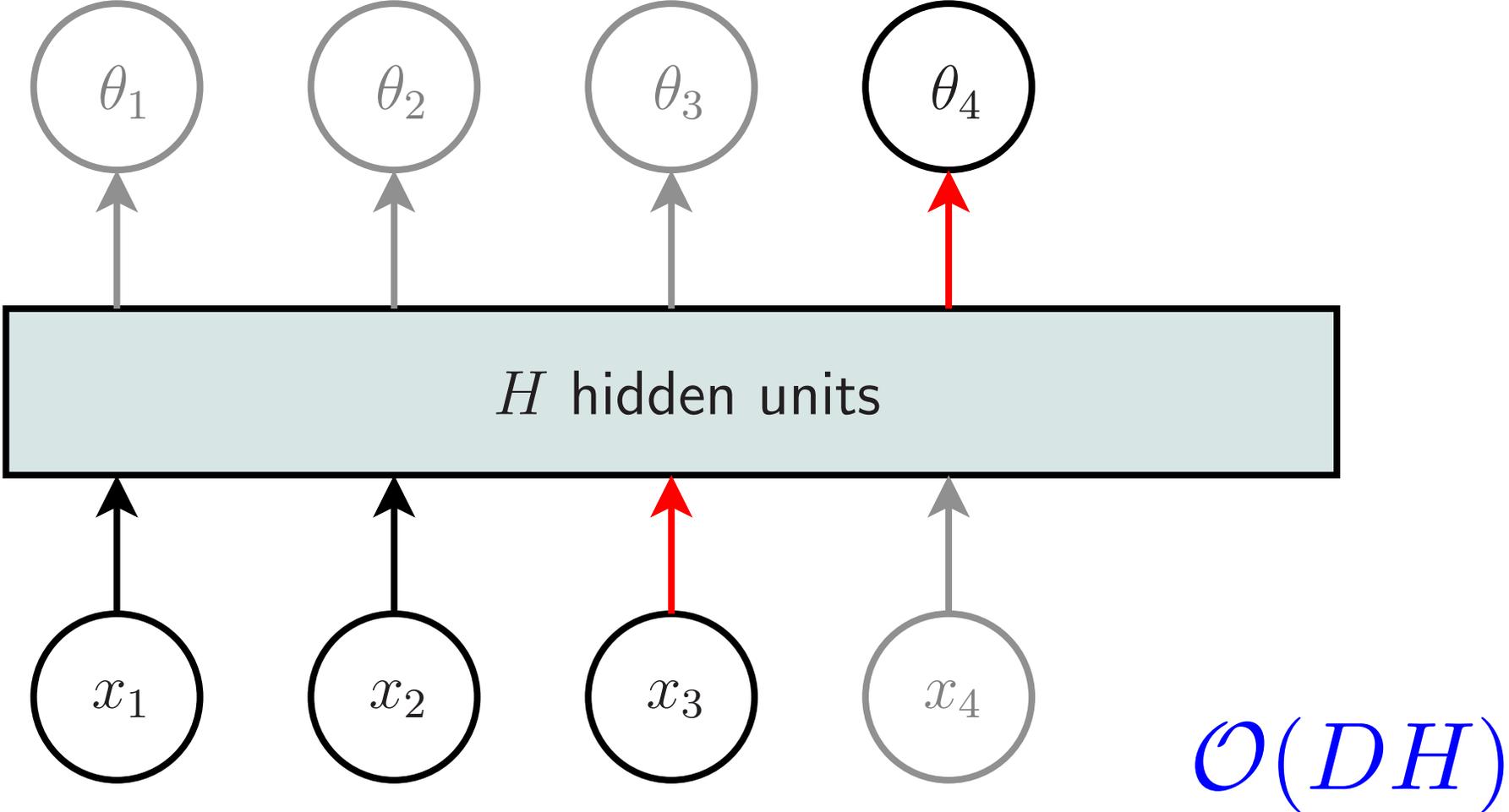
$$P(\mathbf{x}) = P(x_1 | \theta_1) P(x_2 | \theta_2(x_1)) \dots$$

Sequential activation



$$P(\mathbf{x}) = P(x_1 | \theta_1) P(x_2 | \theta_2(x_1)) P(x_3 | \theta_3(x_1, x_2)) \dots$$

Sequential activation



$$P(\mathbf{x}) = P(x_1 | \theta_1) P(x_2 | \theta_2(x_1)) P(x_3 | \theta_3(x_1, x_2)) P(x_4 | \theta_4(x_1, x_2, x_3)) \dots$$

NADE results

Model	ADULT	CONNECT-4	DNA	MUSHROOMS	NIPS-0-12	OCR-LETTERS	RCV1	WEB
MoB	0.00 ± 0.10	0.00 ± 0.04	0.00 ± 0.53	0.00 ± 0.10	0.00 ± 1.12	0.00 ± 0.32	0.00 ± 0.11	0.00 ± 0.23
RBM	4.18 ± 0.06	0.75 ± 0.02	1.29 ± 0.48	-0.69 ± 0.09	12.65 ± 1.07	-2.49 ± 0.30	-1.29 ± 0.11	0.78 ± 0.20
RBM mult.	4.15 ± 0.06	-1.72 ± 0.03	1.45 ± 0.40	-0.69 ± 0.05	11.25 ± 1.06	0.99 ± 0.29	-0.04 ± 0.11	0.02 ± 0.21
RBForest	4.12 ± 0.06	0.59 ± 0.02	1.39 ± 0.49	0.04 ± 0.07	12.61 ± 1.07	3.78 ± 0.28	0.56 ± 0.11	-0.15 ± 0.21
FVSBN	7.27 ± 0.04	11.02 ± 0.01	14.55 ± 0.50	4.19 ± 0.05	13.14 ± 0.98	1.26 ± 0.23	-2.24 ± 0.11	0.81 ± 0.20
NADE	7.25 ± 0.05	11.42 ± 0.01	13.38 ± 0.57	4.65 ± 0.04	16.94 ± 1.11	13.34 ± 0.21	0.93 ± 0.11	1.77 ± 0.20
Normalization	-20.44	-23.41	-98.19	-14.46	-290.02	-40.56	-47.59	-30.16

★ Little variation when changing input ordering:

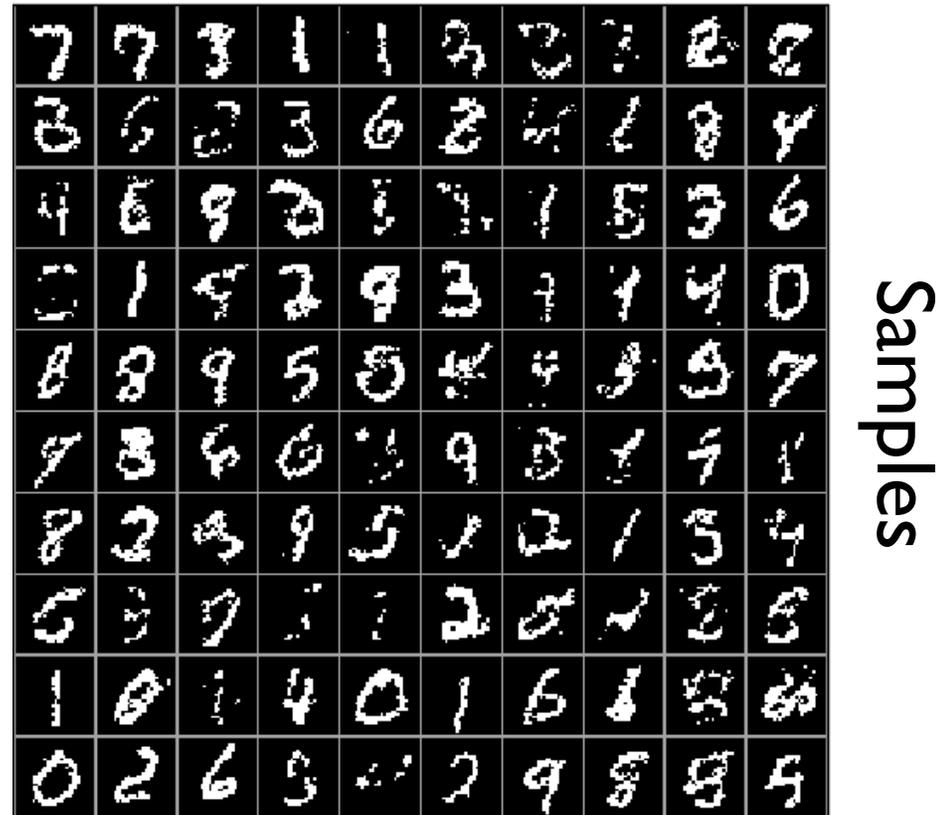
DNA = +/- **0.05**

MUSHROOMS = +/- **0.045**

NIPS-0-12 = +/- **0.15**

NADE results

	Model	Log. Like.
Intractable {	MoB*	-137.64
	RBM (CD1)*	\approx -125.53
	RBM (CD3)*	\approx -105.50
	RBM (CD25)*	\approx -86.34
	FVSBN	-97.45
	NADE	-88.86



RNADE results

Dataset	dim	size	Gaussian	MFA	FVBN	RNADE-MoG	RNADE-MoL
Red wine	11	1599	-13.18	-10.19	-11.03	-9.36	-9.46
White wine	11	4898	-13.20	-10.73	-10.52	-10.23	-10.38
Parkinsons	15	5875	-10.85	-1.99	-0.71	-0.90	-2.63
Ionosphere	32	351	-41.24	-17.55	-26.55	-2.50	-5.87
Boston housing	10	506	-11.37	-4.54	-3.41	-0.64	-4.04

RNADE results

Model	Training LogL	Test LogL
MoG $N = 50$	111.6	110.4
MoG $N = 100$	113.4	112.0
MoG $N = 200$	113.9	112.5
MoG $N = 300$	114.1	112.5
RNADE-MoG $K = 10$	125.9	123.9
RNADE-MoG $K = 20$	126.7	124.5
RNADE-MoL $K = 10$	120.3	118.0
RNADE-MoL $K = 20$	122.2	119.8

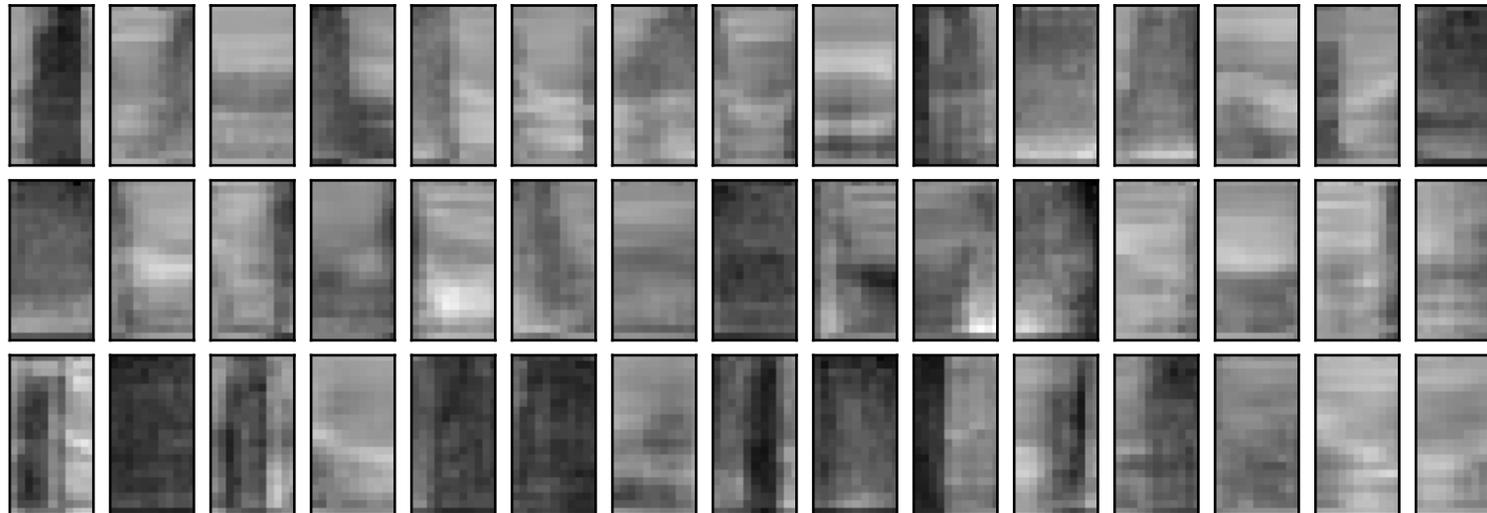


Figure 2: **Top:** 15 datapoints from the TIMIT core-test set. **Center:** 15 samples from a MoG model with 200 components. **Bottom:** 15 samples from an RNADE with 1024 hidden units and output components per dimension. On each plot, time is shown on the horizontal axis, the bottom row displays the energy feature, while the others display the filter bank features (in ascending frequency order from the bottom). All data and samples were drawn randomly.

Deep learning?

One motivation

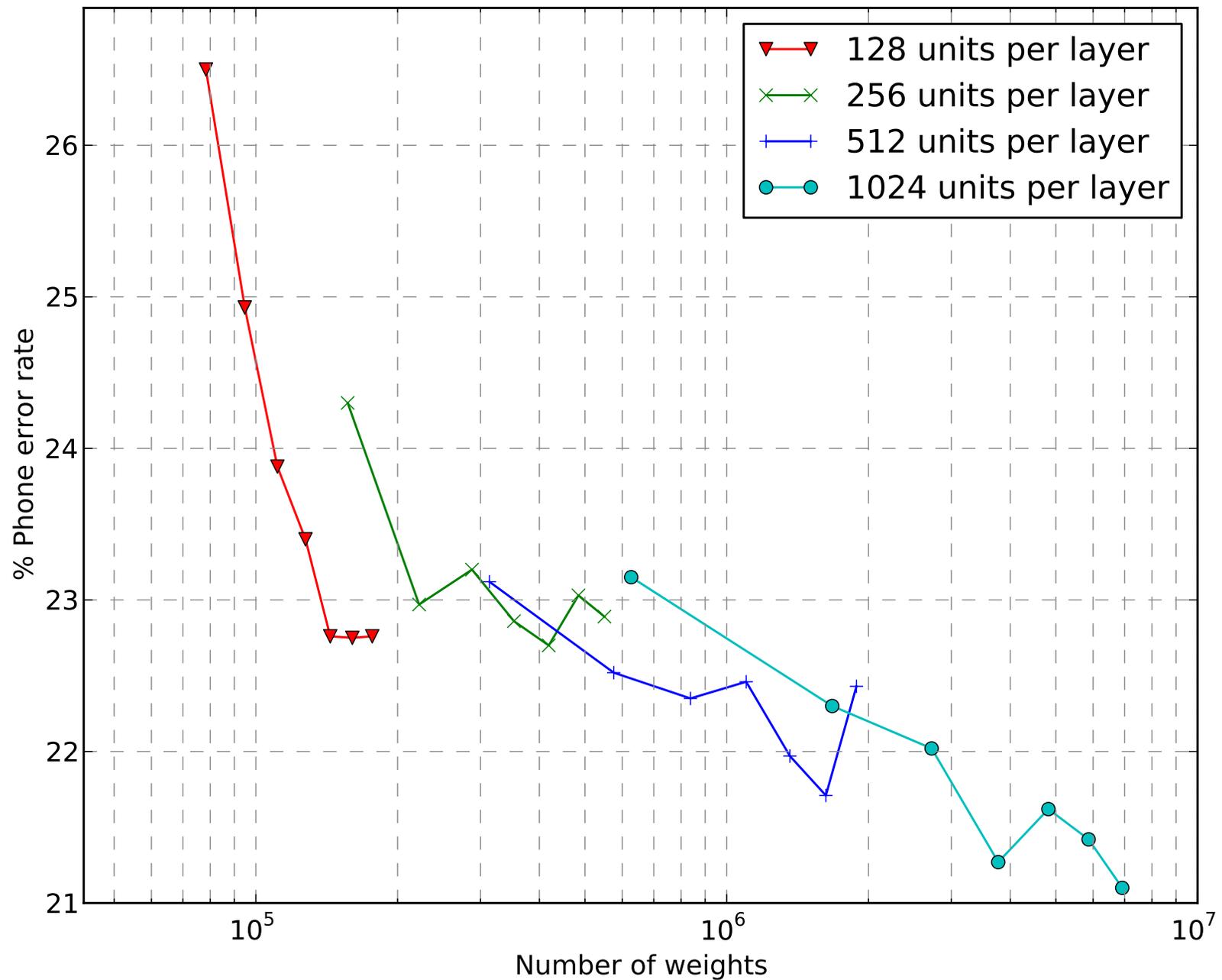
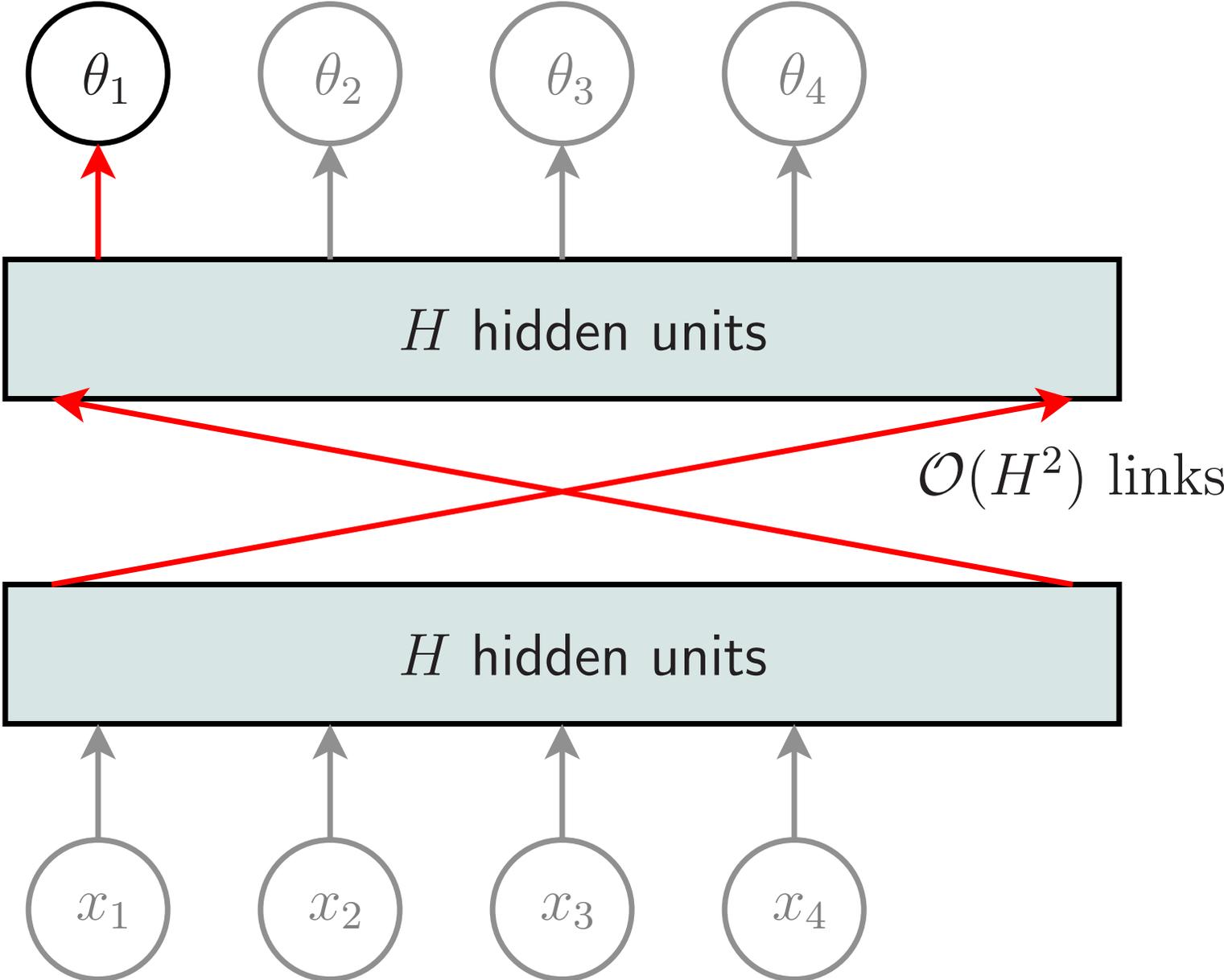
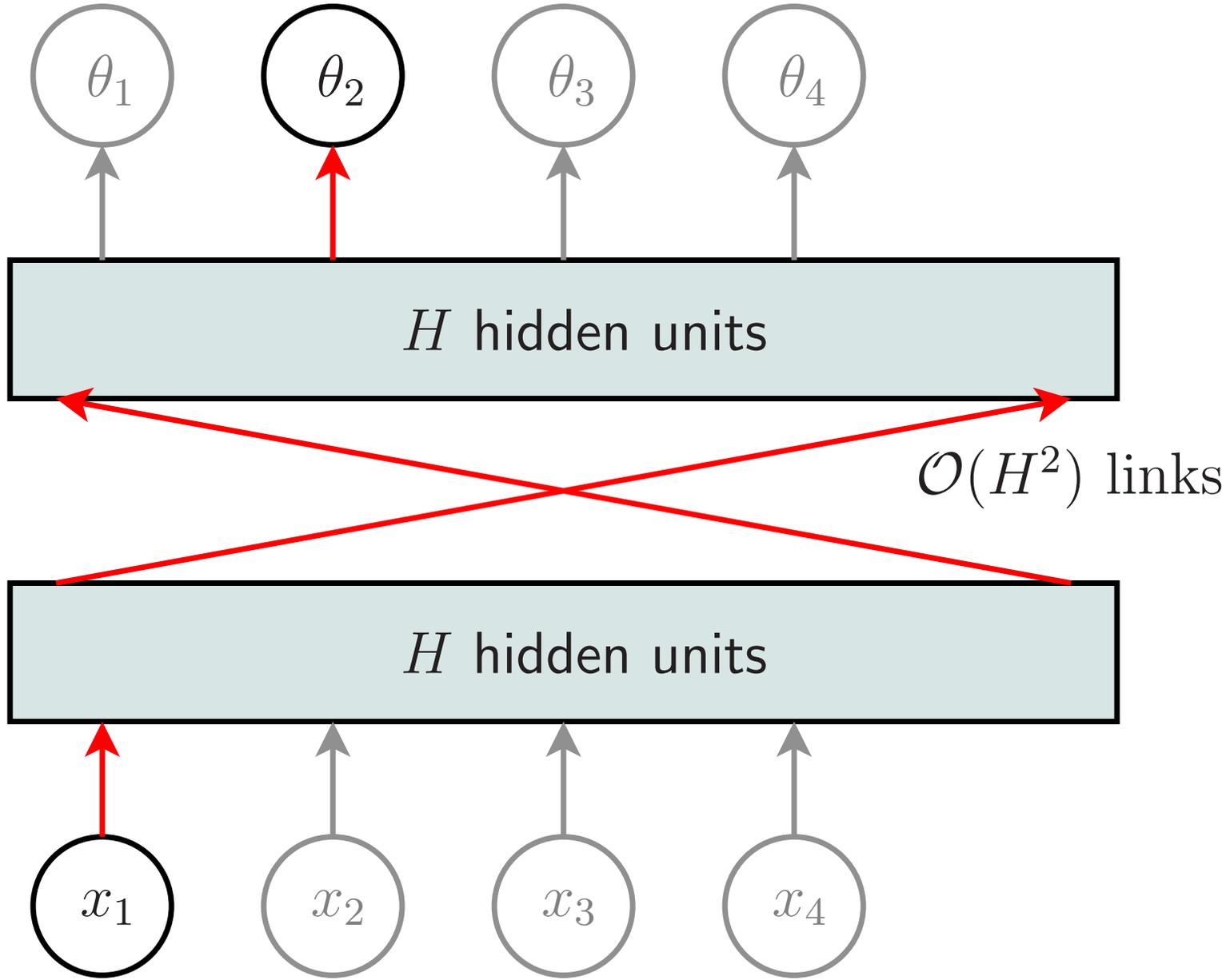


Figure by Benigno Uria

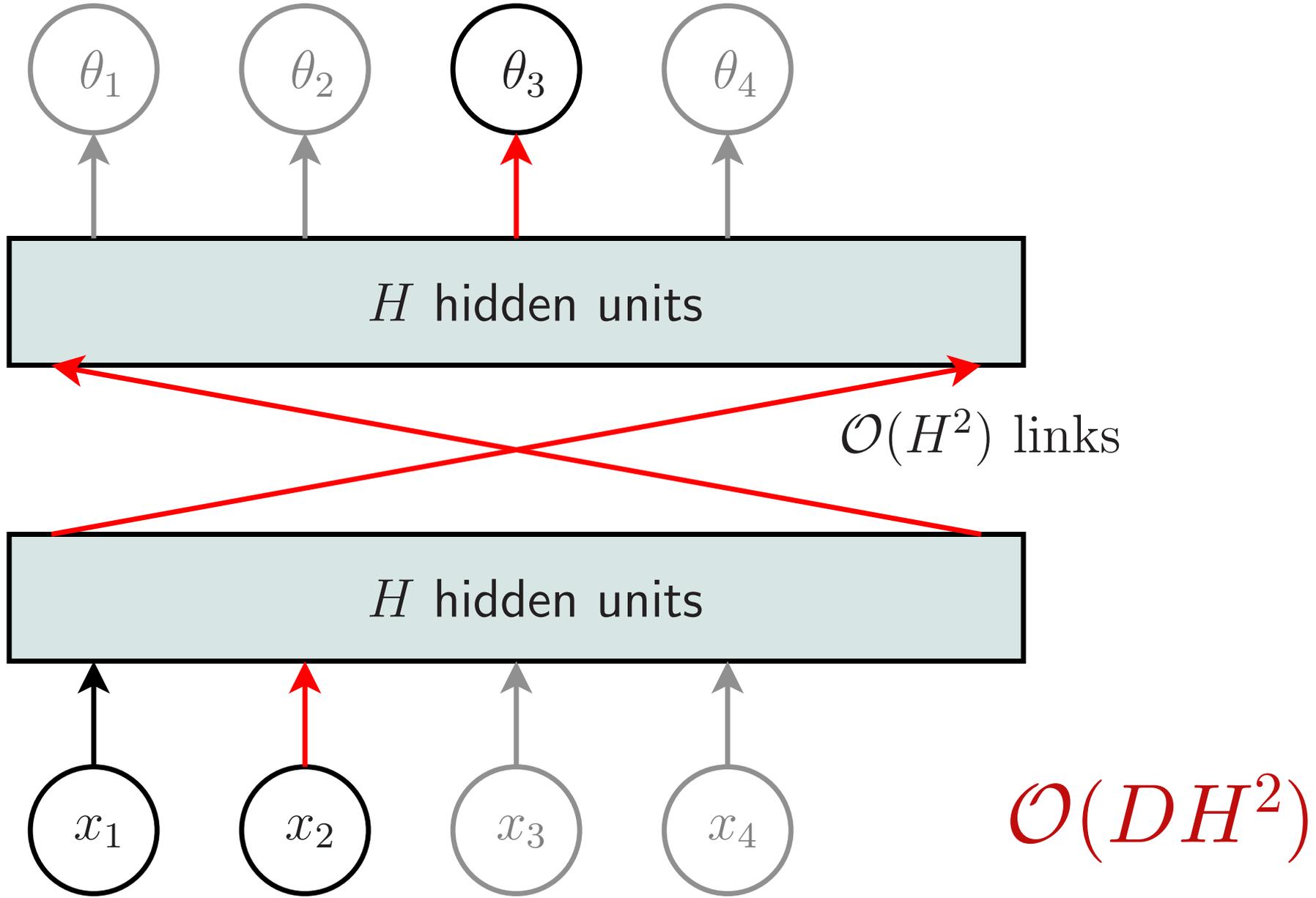
Sequential deep activation



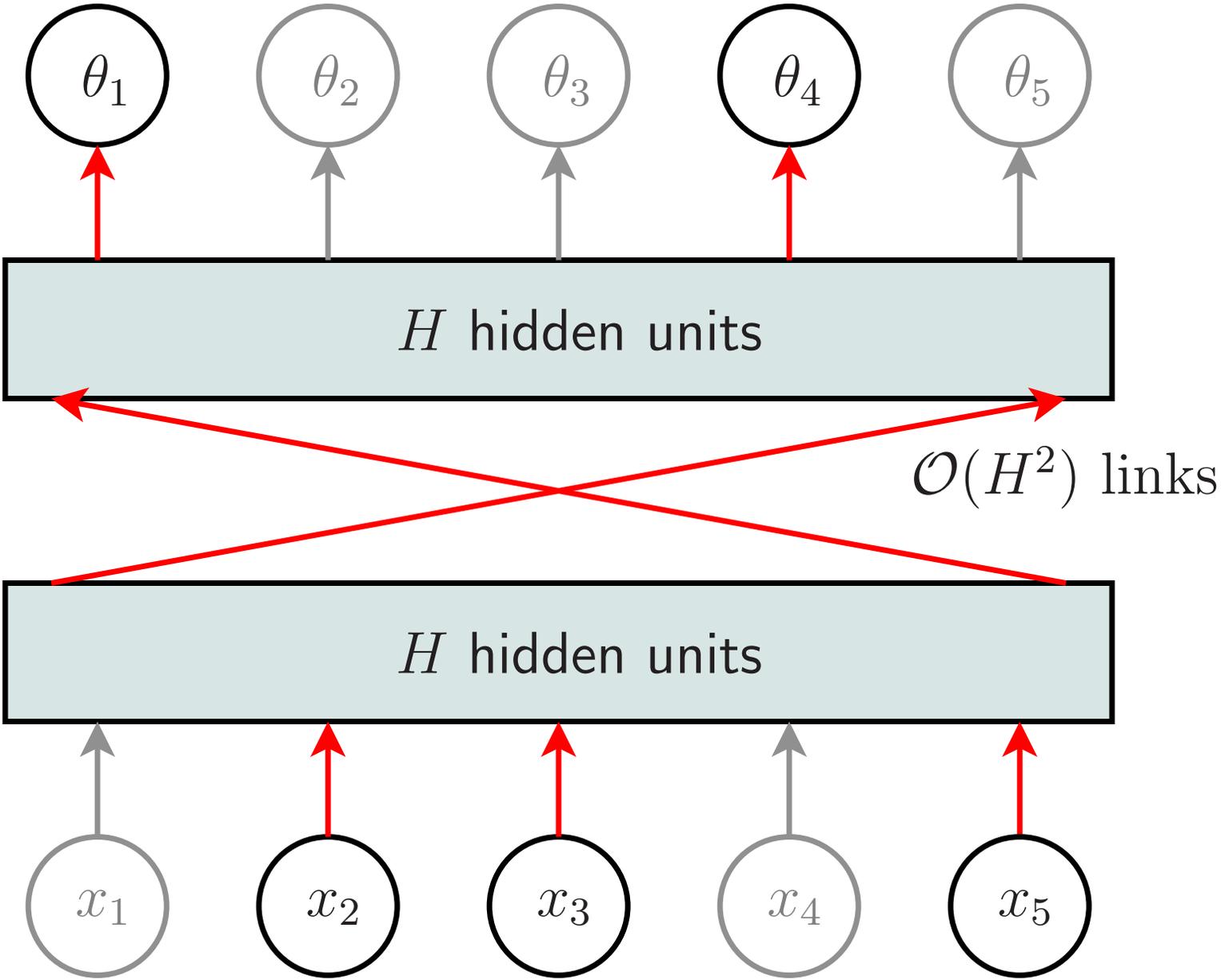
Sequential deep activation



Sequential deep activation



A completing machine



Deep NADE

Train time: $\mathcal{O}(DH + H^2)$ per update

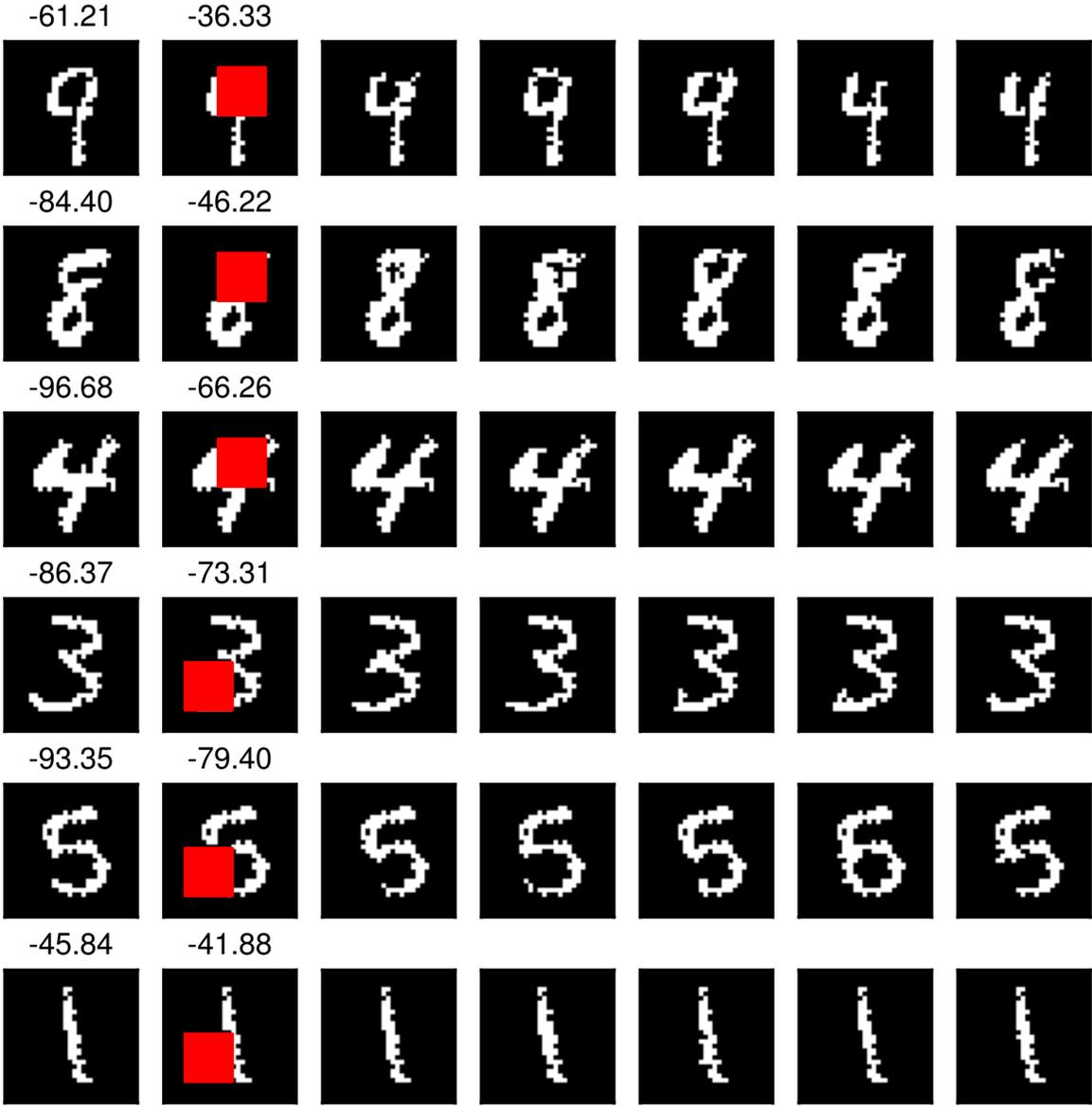
Test time: predict features in any order

(can condition on observations)

Different orderings not consistent:

- Seems bad, but. . .
- have trained large ensemble
- combining different orderings works better

Arbitrary ordering: inpainting



Deep ensembles — results

Small improvements across most UCI datasets

Finally beat MoG on image patches:

Table 4. Average test-set log-likelihood for several models trained on 8 by 8 pixel patches of natural images taken from the BSDS300 dataset. Note that because these are log probability densities they are positive, higher is better.

Model	Test LogL
MoG $K = 200$ (Zoran & Weiss, 2012)	152.8
RNADE 1hl (fixed order)	152.1
RNADE 1hl	143.2
RNADE 2hl	149.2
RNADE 3hl	152.0
RNADE 4hl	153.6
RNADE 5hl	154.7
RNADE 6hl	155.2
EoRNADE 6hl 2 ord.	156.0
EoRNADE 6hl 32 ord.	157.0

ever reported on this task. Ensembles of RNADEs also show

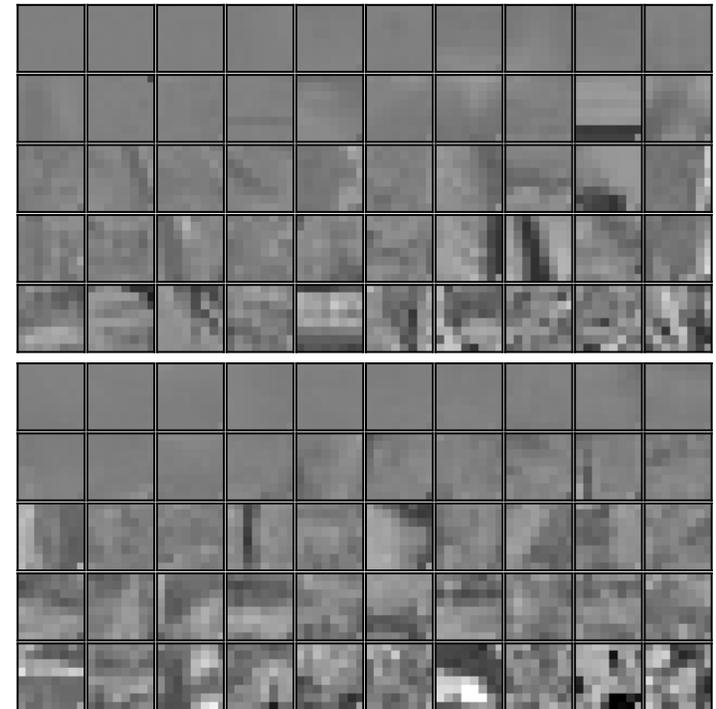


Figure 5. **Top:** 50 examples of 8×8 patches in the BSDS300 dataset ordered by decreasing likelihood under a 6-hidden-layer RNADE. **Bottom:** 50 samples from a 6-hidden-layer RNADE.