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A Mathematical-Physics Approach to Machine Learning

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AI: how (why) does it work?

Al and math, Al and the hard sciences, fantasy or reality?

- National poll: there are about 400 mathematicians, mostly already active in the field and some ready to step in.
- Mainly (but not only) in machine learning.
- ▶ The country (UMI, INDAM, the academic system) has the duty to support them.

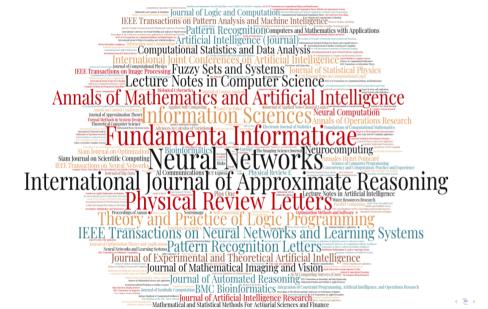
First step: awareness.

Keywords

guantum computation disordered systems dvnamical systems complex systems clustering fuzzy systems segmentation **C**a ta. ence nation algorithms reasoning optimal control ictional VSIS າລເ natural language scalable algorithms statistical high perform KS^{stochastic processes} economy and finance inverse problems wayelet **e** programming spin glassesdeep inference' calculus of variation learning theory diagnostic systems

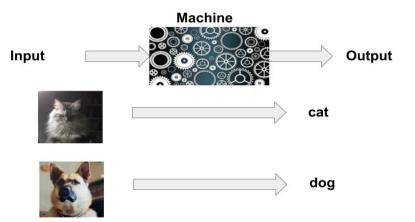
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Journals



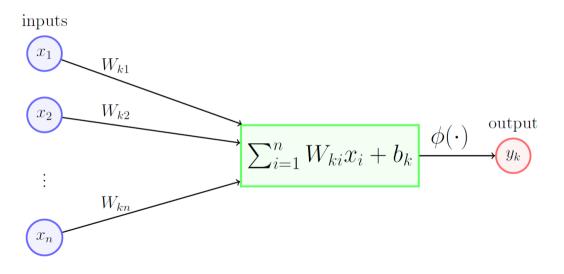
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Machine Learning:



Adjust the parameters, Generalization ability!

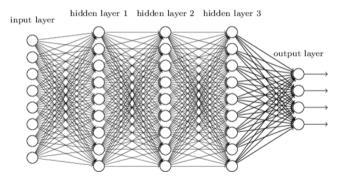
Perceptron algorithm, Rosenblatt 1958



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Bits of history

- Didn't work well, logical and practical objections (but Hinton persisted!)
- Improvements: shallow networks, deep networks Deep neural network



 Large databases, GPU, improved transmission and inversions algorithms, convolutional methods etc.

Efficacy and perspectives

- Nowadays works well in image recognition, language analysis, consumer profiling, and game playing (chess, go, etc).
- Why? Are there principles and laws? Theory and models? Can we make predictions on if it's going to work on different environments? How about efficiency?
- Deep Learning: technology, almost entirely heuristic, like pre-thermodynamic heat-engines.

Open problem (personal) collection

Why is it convenient to grow in depth instead of wideness?

- Why deep networks do not overfit?
- Can we estimate the optimal number of parameters?
- What are the optimal form factors?
- Why good quality minima are easily found? Large entropy.

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Open problem (personal) collection

Three classes of intertwined problems:

Data structure: statistics, signal analysis

Modelisation: physics, mathematics, math-phys

▶ Algorithms: computer science (beyond worse case), optimization

Boltzmann Machines

Math: what is the probabilistic model whose inferential solution is reached with machine learning? Boltzmann Machine!

- Precursors: Sherrington Kirkpatrick 1975, Parisi solution 1979, Hopfield Model 1982
- 1983, Hinton, Sejnowski
- 2009, Hinton, Salakhutdinov

Deep Learning: inverse problem, inference problem, with sampling assigned only in the boundary.

Paramount research: properties of the direct problem under simplifying hypoteses.

Mathematical setting

Let $V_N = \{1, ..., N\}$ be a set of labels for the N particles (or elementary agents, neurons) of a system.

- Spin: to each i ∈ V we attach a binary variable σ_i ∈ {−1, 1} representing the degree of freedom of a single particle.
- Configuration space: Σ_N = {−1,1}^N. A point σ = (σ_i)_{i∈V} ∈ Σ_N represents a (microscopic) configuration of the system.

• Hamiltonian or energy is a (random) function $H_N : \Sigma_N \to \mathbb{R}$

$$H_{N}(\sigma) = -\sum_{\substack{r,s \in \mathcal{S} \\ j \in V_{s}}} \sum_{\substack{i \in V_{r}, \\ j \in V_{s}}} W_{ij}^{(rs)} \sigma_{i} \sigma_{j} - \sum_{r \in \mathcal{S}} \sum_{i \in V_{r}} b_{i}^{(r)} \sigma_{i}$$
(1)

Mathematical Setting

Study the properties of

• Gibbs measure: is a (random) measure on the configuration space Σ_N defined as

$$\mathcal{G}_{N}(\sigma) = \frac{1}{Z_{N}} e^{-\beta H_{N}(\sigma)}$$
⁽²⁾

where $Z_N = \sum_{\sigma \in \Sigma_N} e^{-\beta H_N(\sigma)}$ is the normalization or *partition function*.

- ▶ how does \mathcal{G}_N behaves when $N \to \infty$ (thermodynamic limit)?
- compute the moments generating function in the thermodynamic limit

$$p_N = \frac{1}{N} \log Z_N \tag{3}$$

Multi-species disordered models

- Simplifying assumption: particles can be divided in different species (like in deep learning): let S be a finite set of labels with |S| = K, we assume the vertex set V_N can be written as a disjoint union V_N = ⋃_{s∈S} V_s
- *Relative densities*: we assume that |V_s|/N → α_s ∈ (0, 1) for N → ∞, for each s ∈ S
 for r, s ∈ S
 W^(rs)_{ij} ~ ^{iid} N(^{μ_{rs}}/_{2N}, <sup>Δ_{rs}/_{2N}), (4)
 </sup>

$$b_i^{(s)} \stackrel{\text{iid}}{\sim} \mathcal{N}(\tilde{\mu}_s, \tilde{\Delta}_s)$$
 (5)

invariance under the direct product of the symmetric groups of each specie
 Key result: self averaging Gaussian concentration implies that

$$\lim_{N\to\infty}(p_N-\mathbb{E}p_N)=0, \ W,b-a.s. \tag{6}$$

The convex multi-species model

Consider a finite set (of species) S and the Hamiltonian (1) assuming that:

▶ $\mu = 0$, centered interactions

 \blacktriangleright Δ is a semi-positive definite matrix

 $\blacktriangleright ~\tilde{\mu}$ and $\tilde{\Delta}$ are arbitrary

The property $\Delta \ge 0$ allows to extend the Parisi formula to the multi-species case, namely to express the quenched pressure for $N \to \infty$ as a (infinite dimensional) variational problem.

The core of the problem is the control interacting part of the Hamiltonian

$$H_{N}^{int}(\sigma) = -\sum_{\substack{r,s \in S}} \sum_{i \in V_{r}, \atop j \in V_{s}} J_{ij}^{(rs)} \sigma_{i} \sigma_{j}$$
(7)

This is a centered Gaussian process with covariance

$$\mathbb{E}H_{N}^{int}(\sigma^{1})H_{N}^{int}(\sigma^{2}) = N \sum_{r,s\in\mathcal{S}} \Delta_{rs} \alpha_{r} \alpha_{s} q_{12}^{(r)} q_{12}^{(s)}$$
(8)

where, for each $s \in S$, we define the *relative overlap*

$$q_{12}^{(s)} = \frac{1}{|V_s|} \sum_{i \in V_s} \sigma_i^1 \sigma_i^2$$
(9)

Theorem (Barra-Contucci-Mingione-Tantari '13, Panchenko '13)

If the matrix Δ is positive semi-definite

$$\lim_{N\to\infty} p_N = \lim_{N\to\infty} \mathbb{E}p_N = \inf_x \mathcal{P}(x), \ J - a.s.$$

where $\mathcal{P}(x)$ is K-dimensional functional (generalization of Parisi functional for SK) and x is a cumulative distribution function on $[0,1]^K$ with suitable properties .

Ideas of the proof: The assumption $\Delta \ge 0$ allows to obtain an upper bound for $\mathbb{E}p_N$ dominating the interaction term by a suitable one body system (*replica symmetry breaking interpolation*). The converse bound is obtained exploiting the *synchronization property* of the overlap vector.

Deep Boltzmann machine (DBM)

In the Hamiltonian (1) with K species and centred interactions, consider the species arranged along a linear chain. Only pairs of consecutive species interact, while intra-species interactions as well as long range ones are forbidden. This amounts to the following assumptions on the parameters:

 \blacktriangleright Δ is a non-definite matrix, with zero diagonal and a tridiagonal structure

$$\Delta = egin{pmatrix} 0 & \Delta_{12} & 0 & \cdots & 0 \ \Delta_{12} & 0 & \Delta_{23} & \cdots & 0 \ 0 & \Delta_{23} & 0 & \ddots & 0 \ dots & dots & \ddots & \ddots & \Delta_{K-1,K} \ 0 & 0 & 0 & \Delta_{K-1,K} & 0 \end{pmatrix}$$

- $\mu = 0$, for centered interactions;
- $\tilde{\mu} = \tilde{\Delta} = 0$, in absence of external field.

Theorem (Alberici, Barra, Contucci, Mingione '20)

 $\liminf_{N\to\infty} \mathbb{E} p_N \geq \sup_{a} \mathcal{P}(\theta(a)) ,$

where:

$$\mathcal{P}(heta_1,\ldots, heta_K) \,=\, \sum_{r=1}^K \left(p^{\mathsf{SK}}(heta_r) - p^{\mathsf{Ann-SK}}(heta_r)
ight) + \, p^{\mathsf{Ann.}} \;.$$

 $p^{SK}(\theta_r)$ denotes the limiting quenched pressure of a standard SK model at inverse temperature θ_r , while p^{Ann-SK} denotes its annealed version and p^{Ann} the annealed pressure of the deep Boltzmann machine. Moreover

$$heta_r(a) = \sqrt{lpha_r} \sqrt{rac{1}{a_{r-1}} \Delta_{r-1,r} + a_r \Delta_{r,r+1}}$$

and the supremum is taken over $a = (a_1, \ldots, a_{K-1}) \in (0, \infty)^{K-1}$.

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The annealed pressure is defined as

$$p^{\mathsf{Ann.}} = \lim_{N \to \infty} \frac{1}{N} \log \mathbb{E} Z_N$$

and is easy to compute thanks to the Gaussian nature of the interactions. Our lower bound provides the following result on the annealed regime of the deep Boltzmann machine:

Theorem (Alberici, Barra, Contucci, Mingione - Alberici, Contucci, Mingione '20) If the tridiagonal matrix $M = \Delta \cdot \text{diag}(\alpha_1, \ldots, \alpha_K)$ has spectral radius ≤ 1 , then there exists

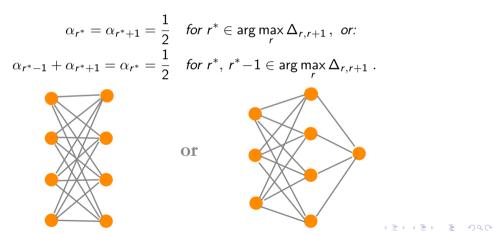
$$\lim_{N\to\infty}\mathbb{E}\,p_N\,=\,p^{\mathrm{Ann.}}\,,$$

namely the deep Boltzmann machine is in the annealed regime.

This result relies on the beautiful algebraic properties of certain Heilmann-Lieb polynomials, which establish a connection between deep Boltzmann machines and monomer-dimer systems.

Minimizing the size of the annealed region for given interaction strengths, one finds the following optimal shapes for the DBM:

Theorem (Alberici, Barra, Contucci, Mingione - Alberici, Contucci, Mingione '20) The maximum of the spectral radius of M over $\alpha_1, \ldots, \alpha_K \ge 0$, $\sum_r \alpha_r = 1$, equals $\max_r \Delta_{r,r+1}$ and is reached if and only if:



M-SK model on the Nishimori line

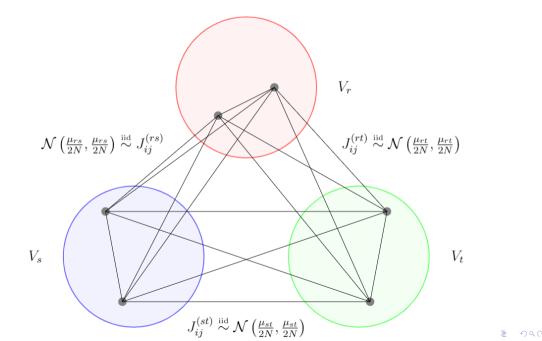
Nishimori line: a subregion of the space of parameters $\mu_{rs}, \Delta_{rs}, \tilde{\mu}_s, \tilde{\Delta}_s$ where $\mu_{rs} = \beta \Delta_{rs}$ and $\tilde{\mu}_s = \beta \tilde{\Delta}_s$. Reabsorbing β it is equivalent to have

$$J_{ij}^{(rs)} \stackrel{\text{\tiny{iid}}}{\sim} \mathcal{N}\left(\frac{\mu_{rs}}{2N}, \frac{\mu_{rs}}{2N}\right) , \quad h_i^{(s)} \stackrel{\text{\tiny{iid}}}{\sim} \mathcal{N}(\tilde{\mu}_s, \tilde{\mu}_s).$$

Identities and correlation inequalities: models in this setting enjoy useful identities $(\langle \cdot \rangle_N \text{ denoting Boltzmann-Gibbs average})$ and inequalities (Contucci, Morita, Nishimori '04,'05):

$$\begin{split} \mathbb{E}[\langle \sigma_i \rangle_N^2] &= \mathbb{E}[\langle \sigma_i \rangle_N], \quad \mathbb{E}[\langle \sigma_i \sigma_j \rangle_N^2] = \mathbb{E}[\langle \sigma_i \sigma_j \rangle_N], \\ \frac{\partial \mathbb{E} p_N}{\partial \mu_{rs}}, \ \frac{\partial \mathbb{E} p_N}{\partial \tilde{\mu}_s}, \ \frac{\partial^2 \mathbb{E} p_N}{\partial \mu_{rs}^2}, \ \frac{\partial^2 \mathbb{E} p_N}{\partial \tilde{\mu}_s^2} \geq 0 \end{split}$$

Replica symmetry: The previous ones imply replica symmetry in the models presented here. The variational principle is indeed finite dimensional.



The convex case on the Nishimori line

The thermodynamic limit is computed by means of the *adaptive interpolation method* (Barbier, Macris '19).

Theorem (Alberici, Camilli, Contucci, Mingione '20)

If μ is positive semidefinite, the thermodynamic limit of the random pressure converges *J*-a.s. and:

$$\lim_{N \to \infty} p_N = \lim_{N \to \infty} \mathbb{E} p_N = \sup_{\mathsf{x} \in [0,1]^K} \bar{p}(\mu, \tilde{\mu}; \mathsf{x})$$

where $\bar{p}(\mu, h; x)$ is a function of K parameters x.

When $\tilde{\mu}_s = 0$ the transition of x towards positive values is controlled by the spectral radius of $M := (\mu_{rs}\alpha_s)_{r,s=1,...,K}$.

- $\rho(M) < 1$: \bar{p} is concave, x = 0 is the unique maximizer;
- $\rho(M) > 1$: x = 0 becomes an unstable saddle point.

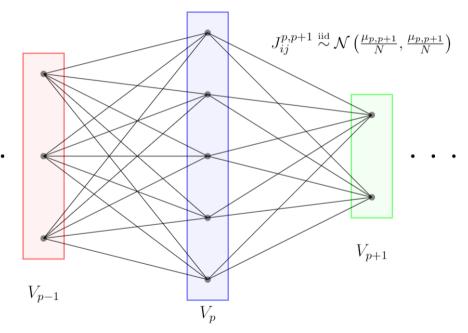
The deep Boltzmann machine on the Nishimori line

The DBM on the Nishimori line corresponds to the choice:

$$\mu = \begin{pmatrix} 0 & \mu_{12} & 0 & \cdots & 0 \\ \mu_{21} & 0 & \mu_{23} & \cdots & 0 \\ 0 & \mu_{32} & 0 & \ddots & 0 \\ \vdots & \vdots & \ddots & \ddots & \mu_{K-1,K} \\ 0 & 0 & 0 & \mu_{K,K-1} & 0 \end{pmatrix},$$

namely, species are arranged in a consecutive way and only adjacent species are allowed to interact. Intra-group interactions are forbidden (0 diagonal elements).

 μ has eigenvalues with alternating sign (symmetric w.r.t. 0). This clearly violates the positivity hypothesis of the convex case.



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The thermodynamic limit of the DBM

Theorem (Alberici, Camilli, Contucci, Mingione '20)

The random pressure of the deep Boltzmann machine on the Nishimori line converges J-a.s. and

$$\lim_{N\to\infty} p_N = \lim_{N\to\infty} \mathbb{E} p_N = \sup_{\mathsf{x}_o} \inf_{\mathsf{x}_e} p_{\mathsf{var}}(\mu, \tilde{\mu}; \mathsf{x}) \;,$$

where x_o and x_e denote the vectors of the odd and even components of $x \in [0, 1)^K$ respectively.

When $\tilde{\mu}_s = 0$ the transition of x towards positive values is controlled by the submatrix (recall $M_{rs} = \mu_{rs}\alpha_s$) $[M^2]^{(oo)} := ((M^2)_{rs})_{r,s \text{ odd } \leq K}$: $\rho([M^2]^{(oo)}) < 1$: x = 0 is the unique optimizer; $\rho([M^2]^{(oo)}) > 1$: the optimizer x = \bar{x} has strictly positive components.

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