

# Efficient Neuro-symbolic Learning of Constraints and Criteria from Sudoku to new functional molecules

(Defresne, Gambardella, et al. 2026)



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de Toulouse

## Inductive and deductive reasoning

- ▶ From observations (solutions) we construct a theory ( $F = m\gamma$ )
- ▶ We then use the theory to make predictions and design objects
- ▶ Until the theory is proven to be incorrect

Sudoku grid with solution

Protein structure with its sequence

The theory is written as a pairwise Graphical Model (a Cost Function Network)

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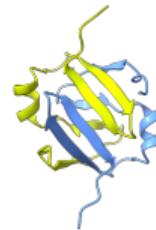
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1	2	6	4	3	7	9	5	8
8	9	5	6	2	1	4	7	3
3	7	4	9	8	5	1	2	6
4	5	7	1	9	3	8	6	2
9	8	3	2	4	6	5	1	7
6	1	2	5	7	8	3	9	4
2	6	9	3	1	4	7	8	5
5	4	8	7	6	9	2	3	1
7	3	1	8	5	2	6	4	9

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## Reminder

- ▶ A set  $X$  of variables  $n$  variables
- ▶ Variable  $x_i$  has domain  $D_i$  max. size  $d$
- ▶ a set of cost functions  $c_{ij} : D_i \times D_j \rightarrow \mathbb{R}^+ \cup \{\infty\}$

## Variables and parameters/costs

- ▶ The cost  $C(t)$  of an assignment  $t$  is the sum of all cost functions on  $t$
- ▶ The cost is linear in the parameters (costs in CF tables)
- ▶ It defines a probability distribution:  $P(t) \propto \exp(-C(t))$  Markov Random Fields

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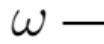
# Structured output prediction (SOP) with a CFN model

2		6		3
7	4	8		
		3		2
8		4	1	
6		5		
		1	7	8
5		9		4

$\omega$

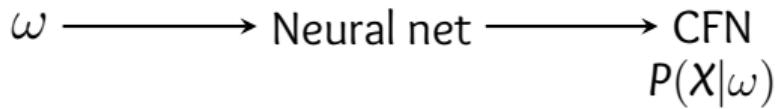
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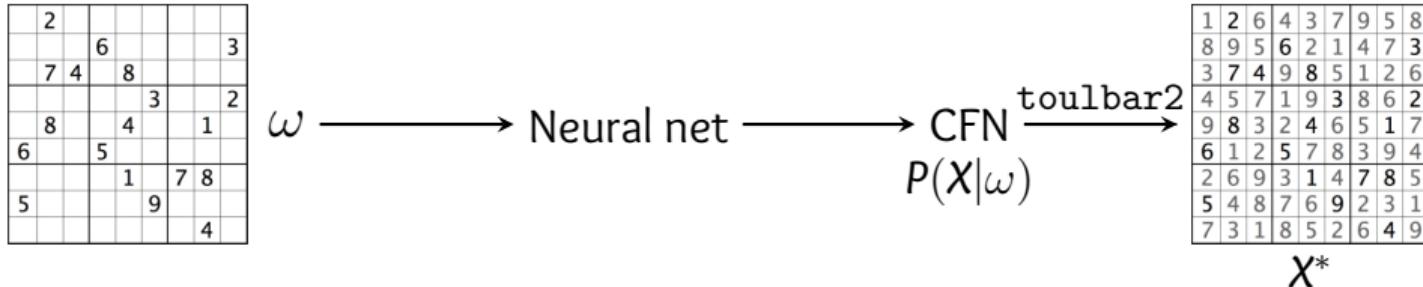
$\omega$   Neural net

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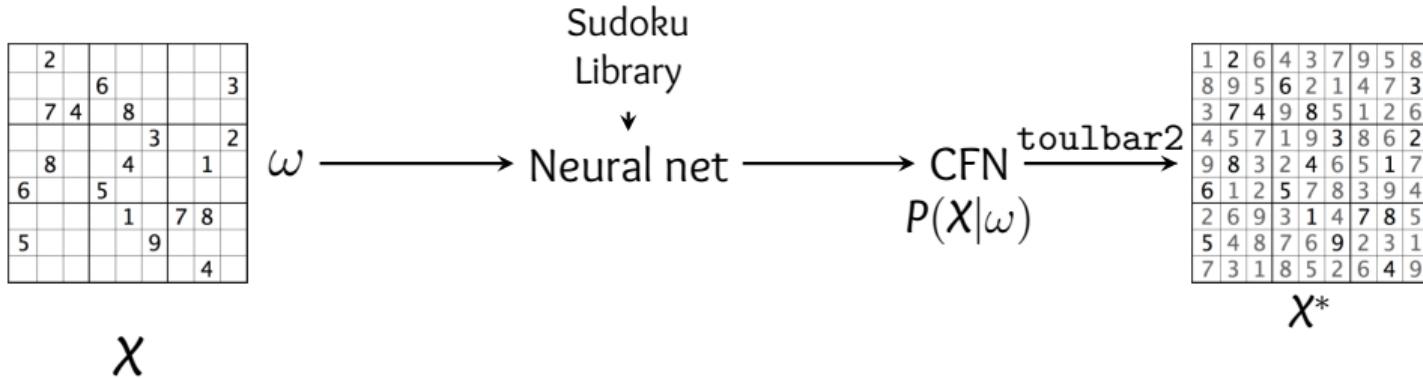
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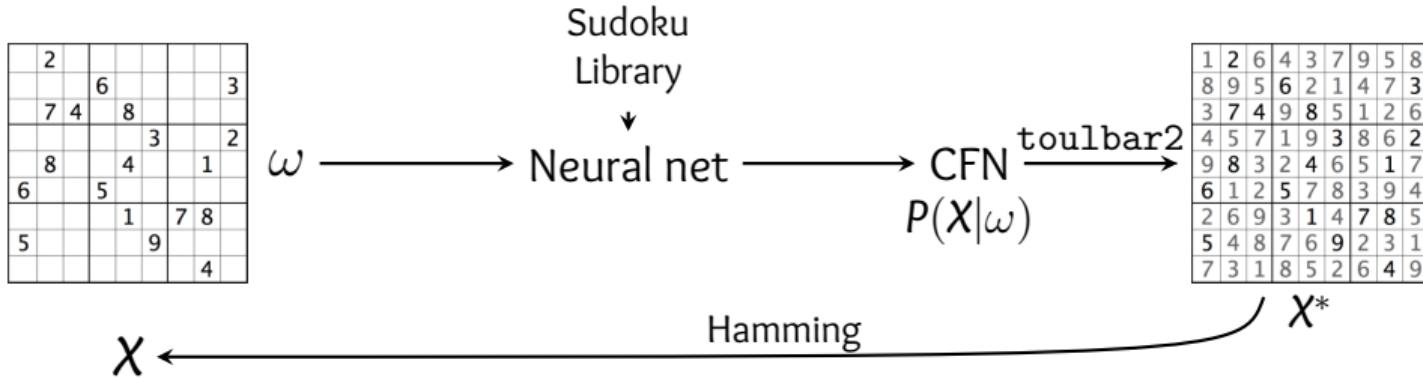
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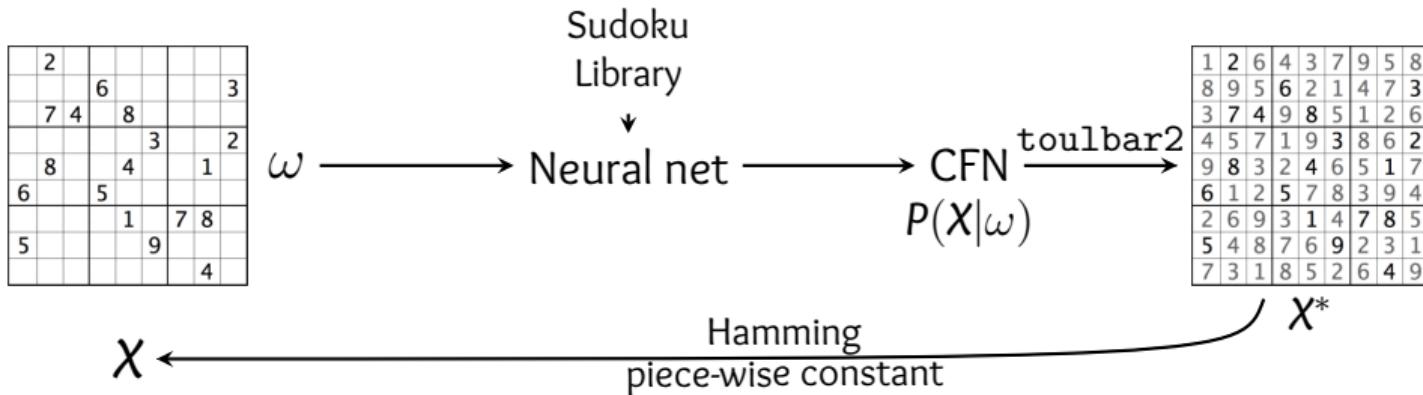
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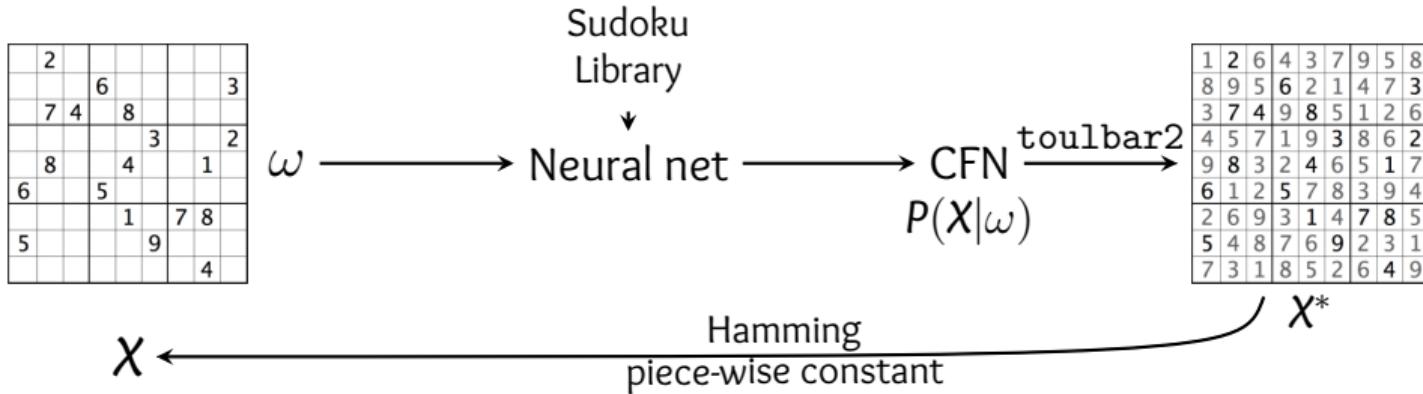
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## Issues

- ▶ Gradients either zero or undefined
- ▶ Requires to repeatedly solve random NP-hard instances

# Structured output prediction (SOP) with a CFN model



Natural choice: the negative loglikelihood

#P-hard

- ▶ Use Besag's pseudo-loglikelihood (1975, efficient)
- ▶ Kicks the solver out of the training loop (scalable training)

(Besag 1975)

- ▶ The Pseudo-LL masks each variable successively in the solution  $X$

(Besag 1975)

$$NLL = - \sum_x \log(P(x))$$

$$NPLL = - \sum_x \sum_{x_i} \log(P(x_i|x_{-i}))$$

- ▶ Nice asymptotic properties
- ▶ The NPLL is a “Fenchel-Young” loss
- ▶ Does not work in practice (high costs)

statistically consistent

(Defresne, Gambardella, et al. 2026; Blondel et al. 2020)

(Montanari et al. 2009)

## Vanishing gradient issue

- ▶ If enough constraints have been learned to force the observed value of  $X_i$  in the context of  $x_{-i}$ , it becomes impossible to learn other constraints.
- ▶ Related to the idempotence of logical information

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## Removing vanishing gradients

We ignore a random fraction of the neighbors/functions when computing  $P(x_i|x_{-i})$

# Easy, hard and extremely hard Sudoku grids

Type	Approach	Acc.	#hints	Train set	Param.	Train time (h)
DL	RRN (Palm et al. 2018)	96.6%	17	180,000	200k	> 50
	Rec. Trans. (Yang et al. 2023)	96.7%	17	180,000	211k	> 50
	Rec. Trans.	76.2–78.2%	17	9,000	-	1.8
	DDPM (Ye et al. 2025)	99.2–100%	33.8	100,000	6M	13.6
	DDPM	0.2%	17	-	-	-
	HRM (G. Wang et al. 2025)	55%	24.8	1,000 × 1,001	27M	>10
Relax+DL	SATNet (P.-W. Wang et al. 2019)	95.1–99.8%	36.2	9,000	600k	2.9
	SATNet	86.1–86.2%	17	-	-	-
CO	(Bessiere et al. 2023)	<b>100%</b>	-	200	-	0.01
CO + ML	(Brouard et al. 2020)	<b>100%</b>	17	9,000	-	1.5
CO+DL	Hinge (Defresne, Barbe, et al. 2023)	<b>100%</b>	17	1,000	180k	>50
	E-PLL (ours)	<b>100%</b>	17	100	22k	0.05
	E-PLL (HRM dataset)	<b>100%</b>	24.8	10 × 100	22k	0.04

Sudokus have only one solution (single target for DL)

- ▶ Existing DL architectures fail on many-solutions Sudokus
- ▶ Corrected using a Reinforcement learning approach
- ▶ Training set with 5 solutions per instance
- ▶ Ability to generate additional solutions

(Nandwani et al. 2021)

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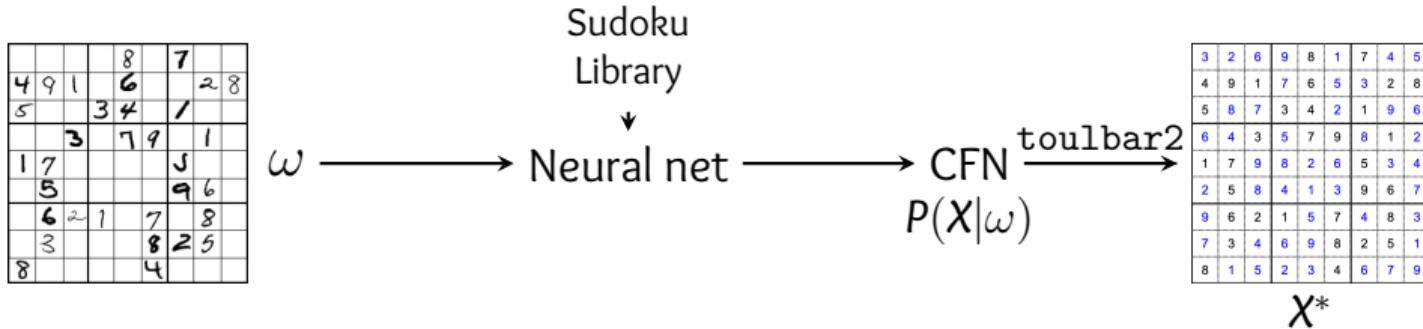
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Sudoku is easy, only one type of constraint

- ▶ Our architecture directly learns how to play Futoshiki
- ▶ Includes both difference and inequality constraints
- ▶ Perfect solving, expected constraints learned

5	>	4	3	>	2	>	1
4		3	1	5	2		
2		1	4	3	5		
3		5	2	1	<	4	
1	<	2	<	5	4	3	

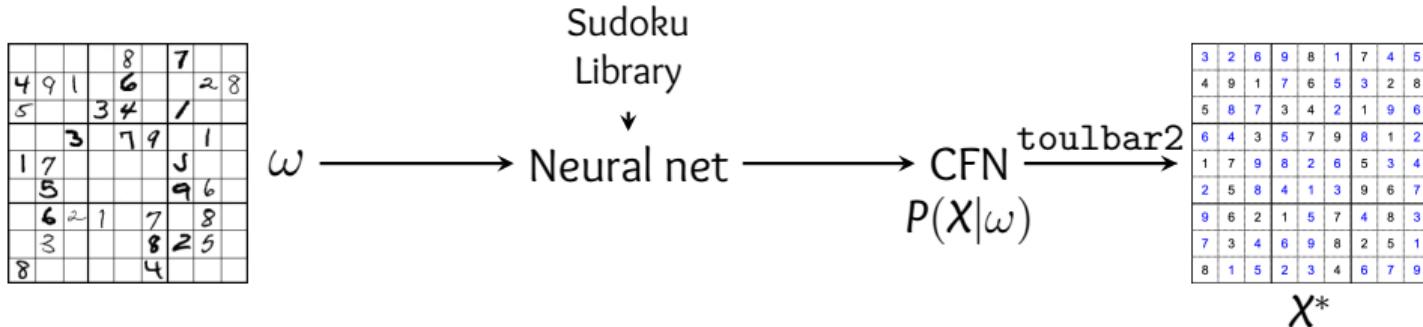
# Learning to play Visual Sudoku



Simultaneously learns to recognize digits and to play the Sudoku

SATNet	Theoretical (no corrections)	Hybrid
63.2 %	74.2%	$94.1 \pm 0.8\%$

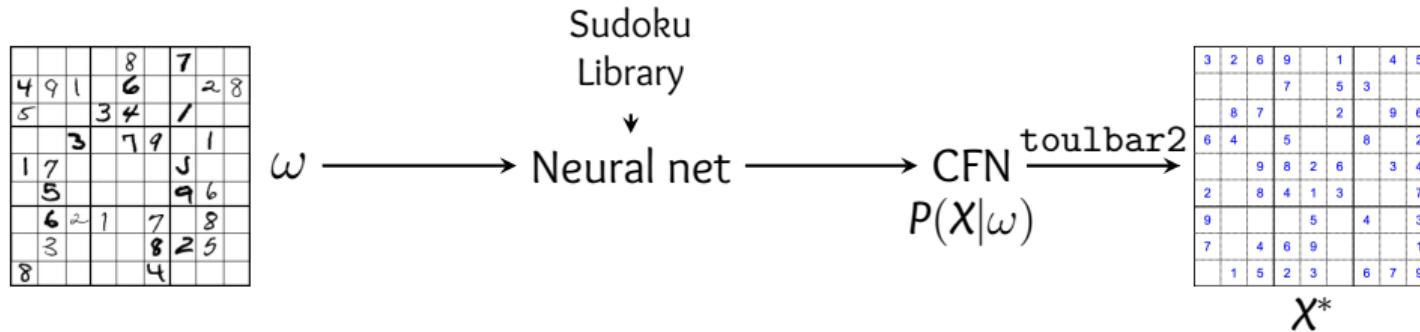
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# Reading numbers without cheating (grounding)



## Grounding issue: a nasty form of data leakage

(Chang et al. 2020)

- ▶ The training set contains images and associated decoded digits (hints).
- ▶ Solved using a complex architecture (InfoGAN+clustering+Distillation) (Topan et al. 2021)
- ▶ E-PLL: missing data, imputation by optimisation
- ▶ Much longer training (a few hours)

# Existing approaches

Approach	MNIST accuracy	Percep.	Solved	Training (h)
SATNet	0.0 %	0.0 %	0.0 %	-
Rec. Trans (Yang et al. 2023)	99.4%	74.8%	75.6%	5.1
NeSy. Prog. (Li et al. 2023)	99.6–99.7%	90.7–93.1%	92.2–94.4%	4.7
<b>E-PLL (Ours)</b>	98.8%	72.9%	93.4%	3.2

## From DFL to Structured Output Prediction

- ▶ Data: pairs  $\langle \omega, c \rangle$  where  $c$  define the criterion parameters
- ▶ Assumes constraints are known
- ▶ we can compute a SOP data-set  $(\omega, X^*)$
- ▶ Aim: minimize regret (difference in real cost of the predicted and optimal solution)

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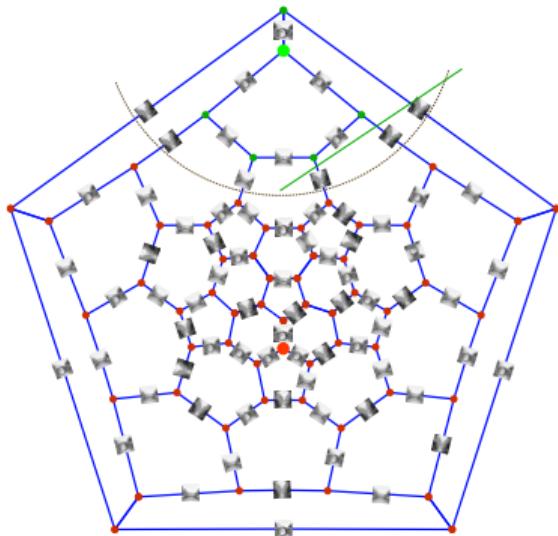
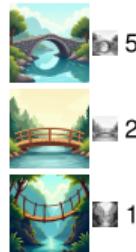
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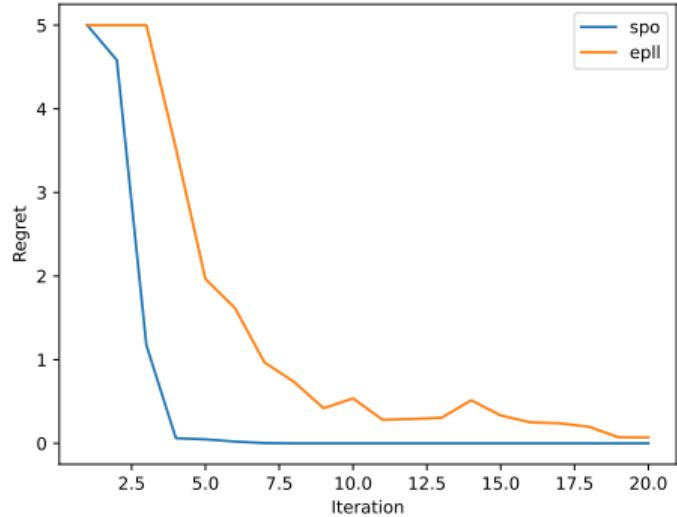
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## The Min/Max-Cut problems

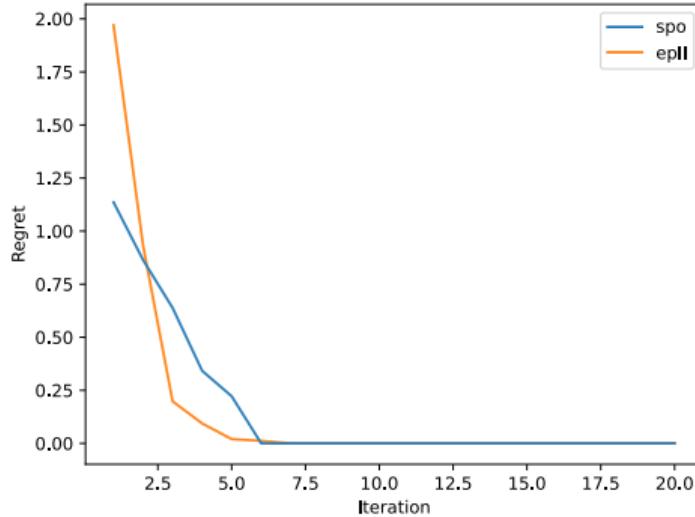
- ▶ one Boolean variable per vertex (cut side)
- ▶ per-edge difference (Min-Cut) or equality (Max-Cut) function scaled by a predicted scalar  $c(\omega)$
- ▶ bridge images with Gaussian noise (std-dev 10)



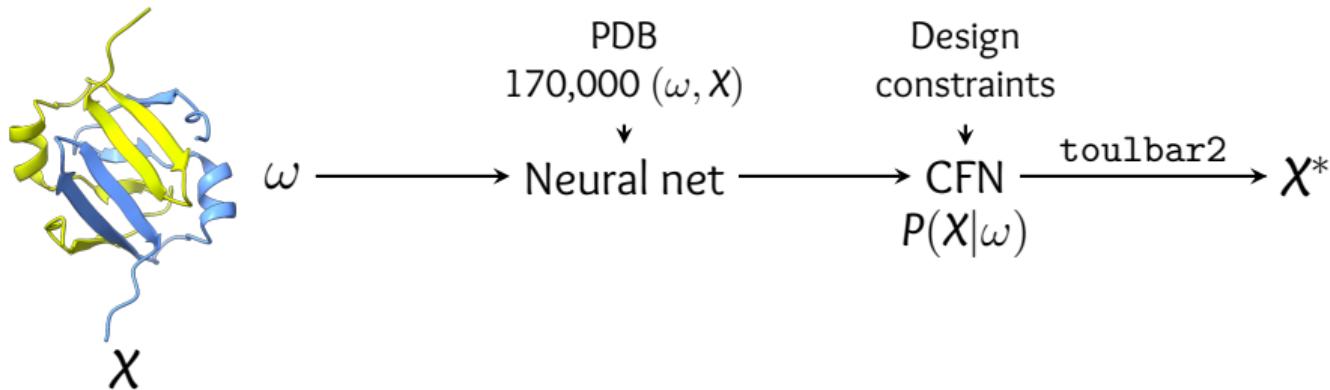
mincut learning task



maxcut learning task



# Learning to design proteins: Effie



## Neural architecture

- ▶ More complex  $SE(3)$ -equivariant neural network
- ▶ Relies on Gated MLPs (post-transformer architecture)

(Liu et al. 2021)

# Optimizing a complete protein sequence

Full redesign of large proteins in the test set

- ▶ Guaranteed `toulbar2` solution expensive
- ▶ Using LR-BCD SDP solver instead (Durante et al. 2022)

Outperforms all-atoms XIX<sup>th</sup>-century physics

- ▶ Metric: Native Sequence Recovery rate (NSR)

Approach	Rosetta	Effie
NSR	17.9%	32.8%

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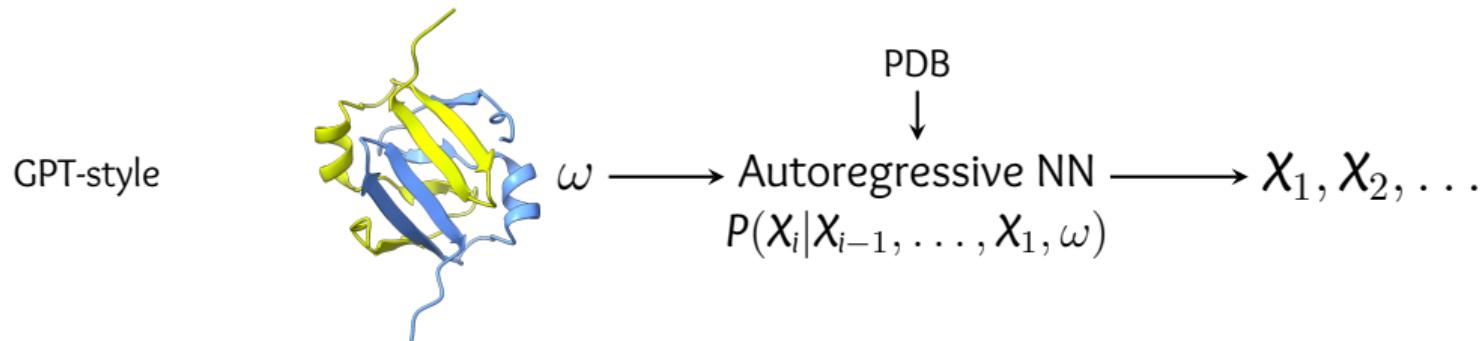
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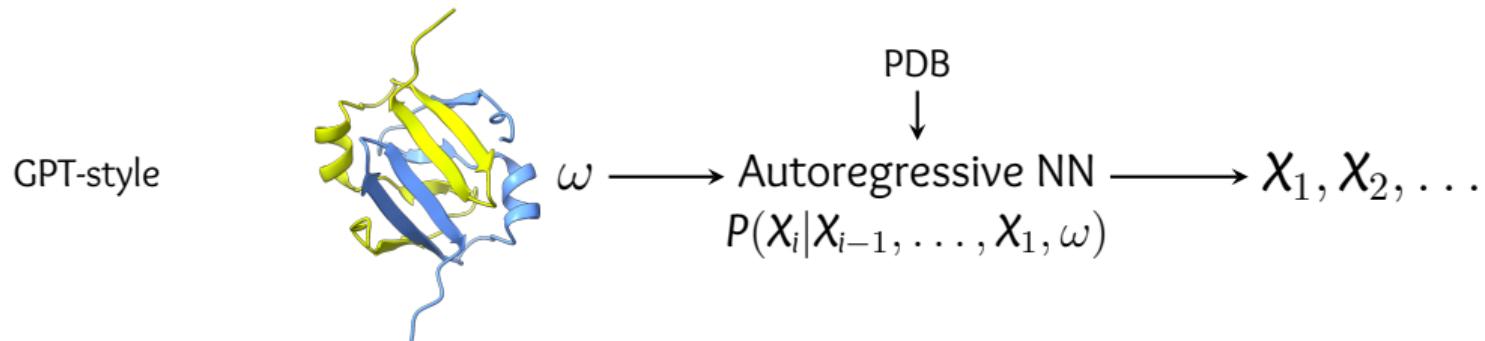
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## Pros and cons

- ▶  $P(X_1, \dots, X_n) = P(X_1)P(X_2|X_1) \cdots P(X_n|X_1, \dots, X_{n-1})$  is a mathematical identity
- ▶ But an easily broken one (e.g., low temperature sampling)
- ▶ Heuristic score instead of NP-hard solving
- ▶ Limited control for design constraints, hard to “reason forward”

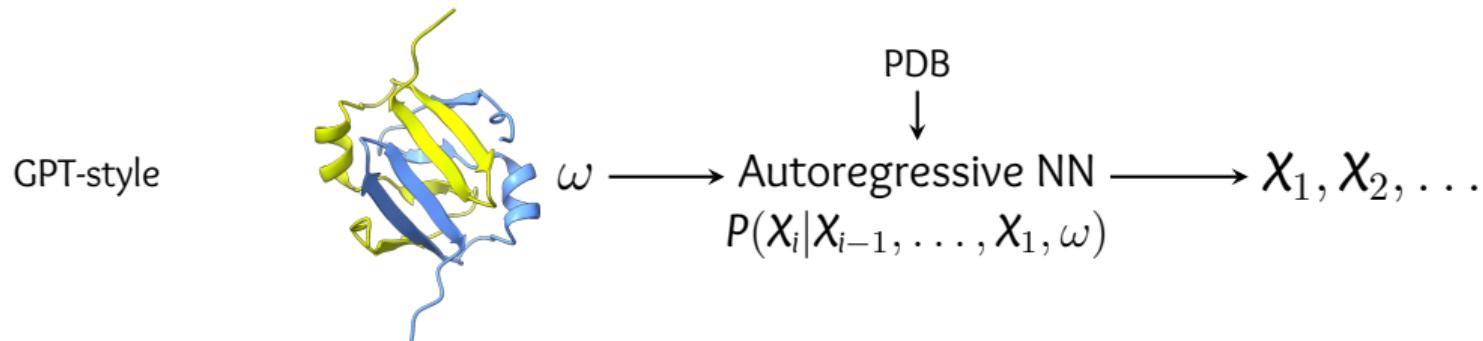
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# Predicting SARS-CoV2 variants



## Enumerate CoViD variants with a bounded number of mutations

- ▶ Uses only the initial March 2020 RBD-ACE2 structure + Effie/toulbar2
- ▶ Relies on a global constraint to bound mutations (Ruffini et al. 2021)
- ▶ Predicts all the first SARS-CoV2 VoCs ( $\alpha, \beta, \gamma, \delta, \kappa, \iota, \lambda$  and  $\mu$ )
- ▶ In a few seconds, on one CPU-thread.

Not achievable by pure autoregressive models (ProteinMPNN).

Previously shown to predict contagious antibody-resistant variants (Colom et al. 2024).

# Design of an enzyme organizing platform

## Design of an heteromeric hexamer

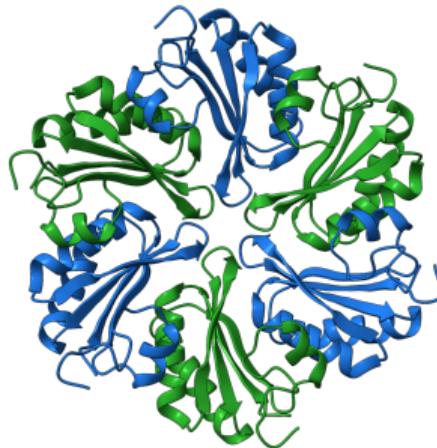
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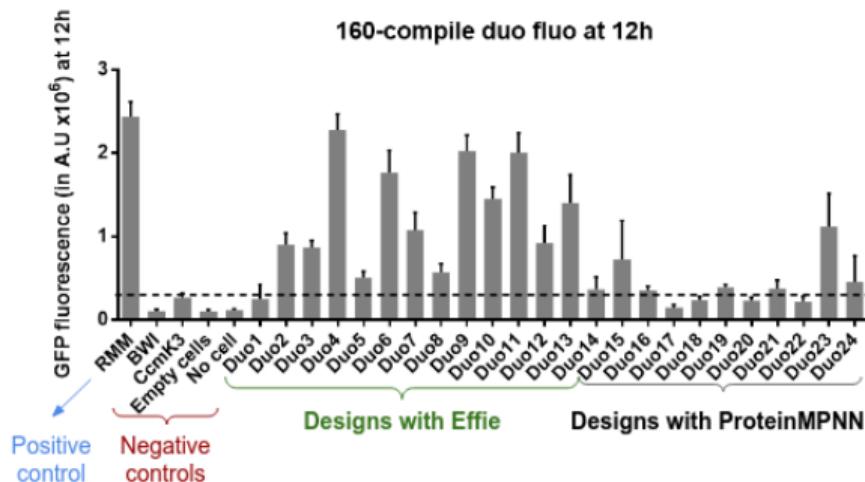
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Scoring →	Effie	PMPNN
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# How often is better than ?

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## A Neural Net, a GM, and a discrete optimizer in a NeSy autoencoder

- ▶ A NeSy Generative AI that benefits from each component
- ▶ Neural Network: ideal to extract a representation of  $P(X|\omega)$  from raw inputs
- ▶ Represented as a GM in a fully explorable and controllable latent layer
- ▶ Using decoding by discrete reasoning (toulbar2) that accepts side constraints
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# Acknowledgments



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S. de Givry (INRA)  
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B. Donald (U. North Carolina)  
K. Roberts (U. North Carolina)  
T. Simonson (Polytechnique)  
J. Cortes (LAAS/CNRS),...



My apologies to those missing in these lists. Even imperfect lists seem better than no list

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