

Scalable Coupling of Deep Learning with Logical Reasoning: from Sudoku to new functional molecules

Thomas Schiex

Joint work with S. Barbe, M. Defresne (PhD student)



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 binary Cost Function Network

Human reasoning and scientific discovery

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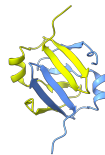
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1	2	6	4	3	7	9	5	8
8	9	5	6	2	1	4	7	3
3	7	4	9	8	5	1	2	6
4	5	7	1	9	3	8	6	2
9	8	3	2	4	6	5	1	7
6	1	2	5	7	8	3	9	4
2	6	9	3	1	4	7	8	5
5	4	8	7	6	9	2	3	1
7	3	1	8	5	2	6	4	9

Sudoku grid with solution



Protein structure with its sequence

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Binary Cost Function Network

- ▶ A set X of variables
- ▶ Variable x_i has domain D_i
- ▶ a set of cost functions

n variables

max. size d

$$c_{ij} : D_i \times D_j \rightarrow \mathbb{R}^+ \cup \{\infty\}$$

Costs and probabilities

- ▶ The cost $C(t)$ of an assignment t is the sum of all cost functions on t
- ▶ Toulbar2 solves the Weighted Constraint Satisfaction Problem
- ▶ A CFN defines a probability distribution: $P(t) \propto \exp(-C(t))$
- ▶ Normalizing constant is #P-hard to compute

$$\min_t C(t)$$

Markov Random Fields

Binary Cost Function Network

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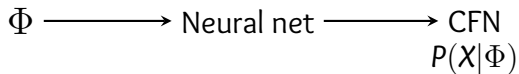
	2						
			6				3
	7	4		8			
					3		2
	8			4			1
6			5				
				1		7	8
5					9		
							4

Φ

	2						
			6				3
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$\Phi \longrightarrow$ Neural net

	2						
			6				3
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6			5				
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	2						
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	7	4		8			
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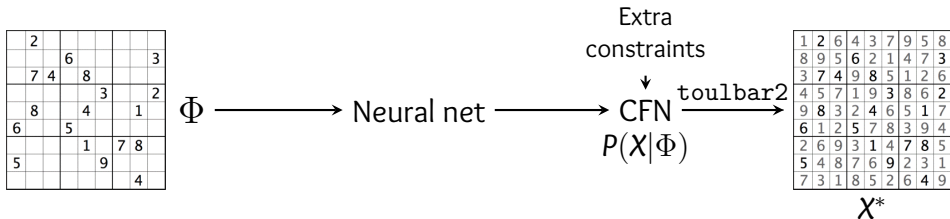
 Φ

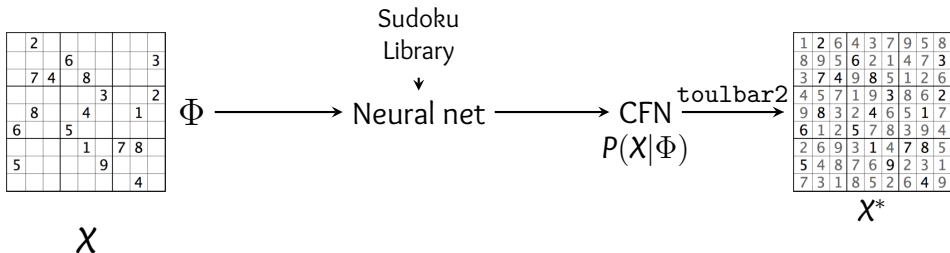
→ Neural net

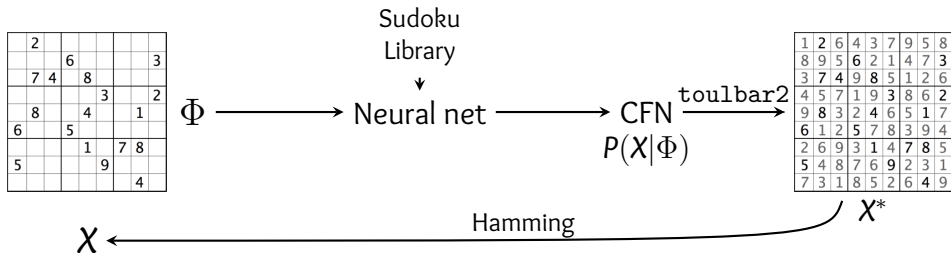
CFN $\xrightarrow{\text{toulbar2}}$
 $P(X|\Phi)$

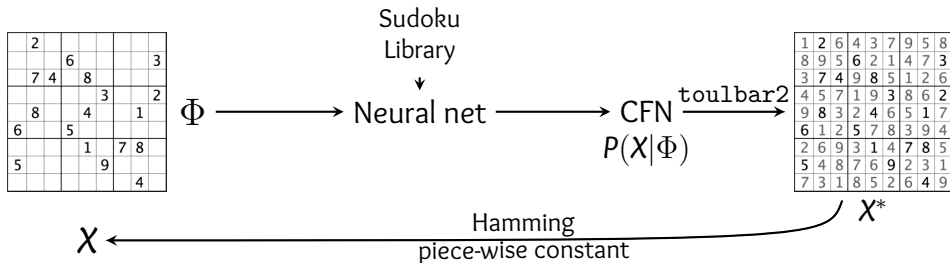
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 X^*



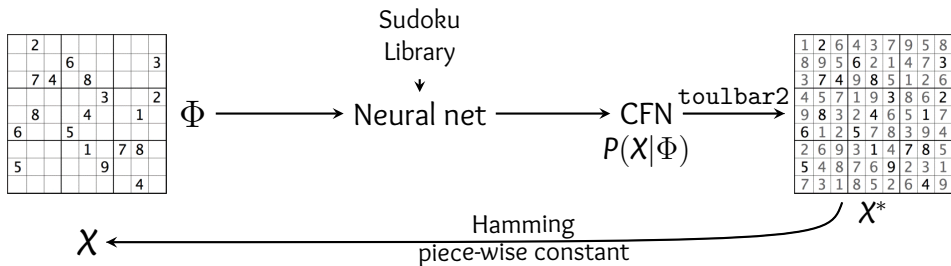






Issues

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

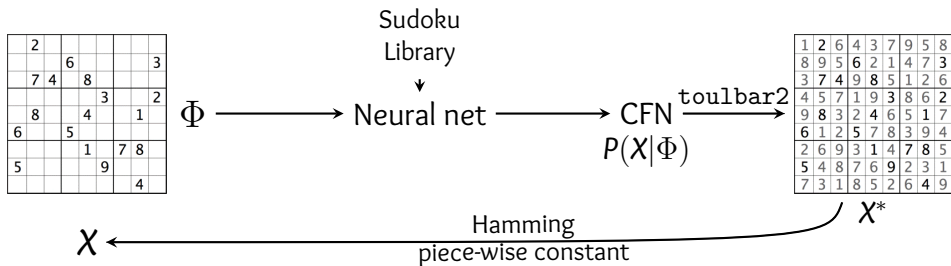


Our solution

- ▶ Introduced a dedicated loss: the E-Pseudo Log Likelihood
- ▶ Kicked the solver out of the training loop (scalable training)

IJCAI'2023

(Defresne et al. 2023)



Approach	Architecture	Acc.	Grids	Training set
RRN NeurIPS18	GNN	96.6%	Hard	180,000
SATNet ICML19	Relaxation	99.8%	Easy	9,000
Hybrid IJCAI23	E-PLL	100%	Hard	200
Symbolic IJCAI23	MaxSAT	100%	Hard	200

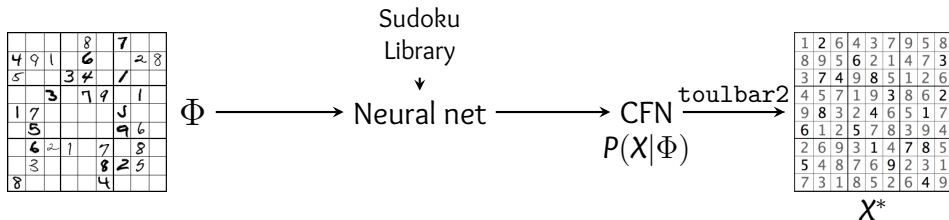
PLL from solution s and current predicted CFN

- ▶ $-\log(P(s))$ would be a nice loss intractable
- ▶ For each variable x_i , we mask x_i in the solution
- ▶ Compute $P(x_i|X_{-i})$ easy
- ▶ Use $-\sum_{x_i} \log(P(x_i|X_{-i}))$ as the pseudo-loss
- ▶ Excellent asymptotic properties statistically consistent
- ▶ Does not work in practice (high costs) Montanari and Pereira 2009

The Emmental Pseudo-loglikelihood

- ▶ When we compute $P(x_i|X_{-i})$
- ▶ We ignore a random fraction of the neighbors
- ▶ Strongly accelerates stochastic gradient optimization

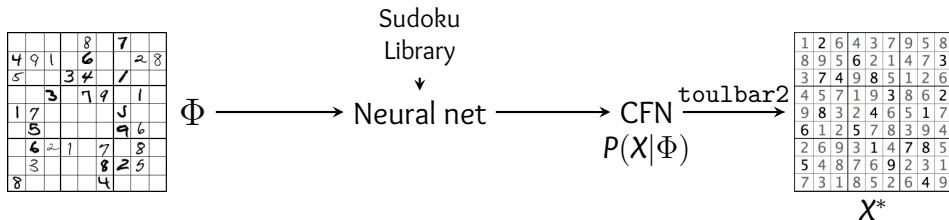
Learning to play Visual Sudoku



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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- ▶ The training set contains images and associated decoded digits (hints).
- ▶ Corrected in (Topan et al. 2021) using a complex architecture (GAN+clustering+Distillation)

Easily solvable in our architecture

- ▶ In the training set, we replace hints with a new “missing value” (0)
- ▶ We connect LeNet output to the CFN generator input
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Sudokus have only one solution (single target for DL)

- ▶ Existing DL architectures fail on many-solutions Sudokus
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- ▶ Training set with 5 solutions per instance
- ▶ Ability to generate additional solutions

(Nandwani et al. 2021)

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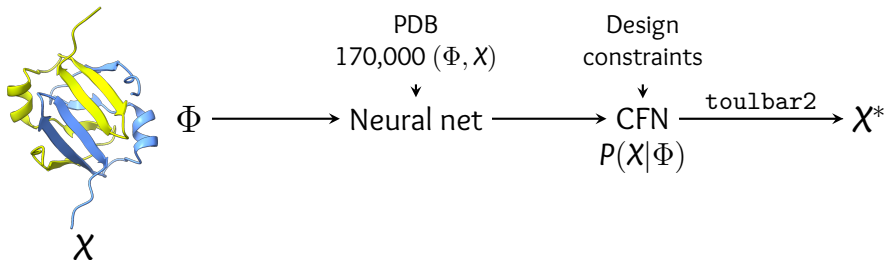
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Sudoku is easy, only one type of constraints

- ▶ 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 design proteins: Effie



Optimizing a complete protein sequence

Full redesign of large proteins in the test set

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

Outperforms all-atoms XIXth-century physics

- ▶ Metric: Native Sequence Recovery rate (NSR)

Approach	Rosetta	Effie
NSR	17.9%	32.8%

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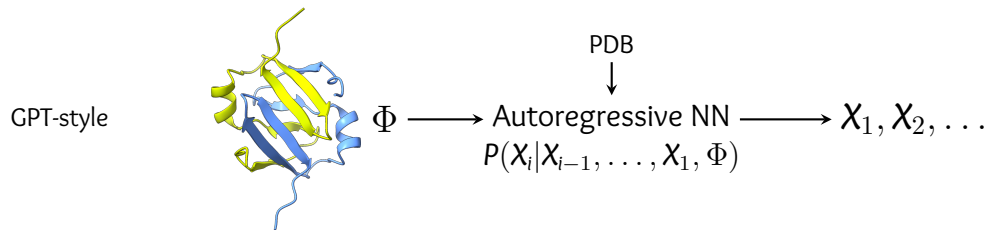
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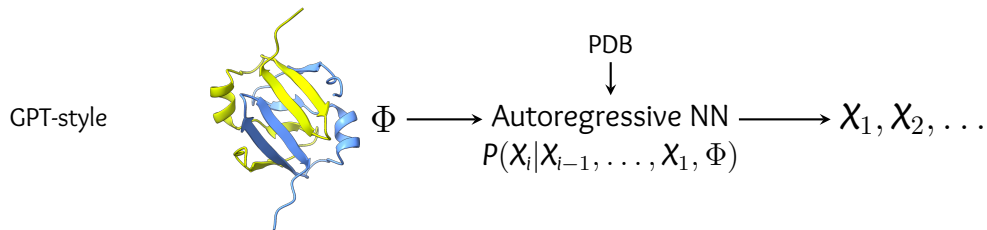
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Pros and cons

- ▶ Heuristic score instead of NP-hard solving
- ▶ Capacity to capture higher-order interactions
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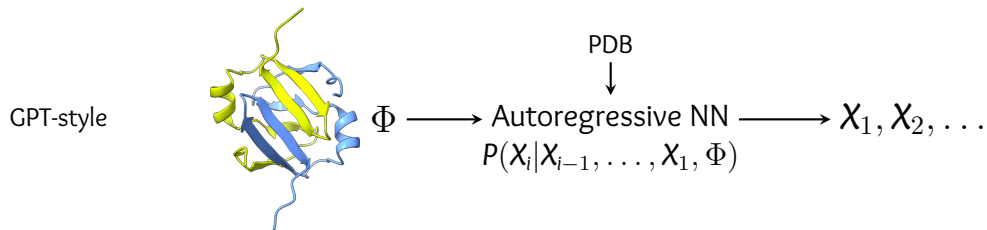
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Enumerate CoViD variants with a bounded number of mutations

- ▶ Uses only the initial March 2020 RBD-ACE2 structure + Effie/toulbar2
- ▶ Relies on (Montalbano et al. 2022) global constraint to bound mutations
- ▶ Predicts all the first SARS-CoV2 VoCs (α , β , γ , δ , κ , ι , λ and μ)
- ▶ In a few seconds, on one CPU-thread.

Not achievable by pure autoregressive models (ProteinMPNN)

Design of an enzyme organizing platform

Design of an heteromeric hexamer

- ▶ Design  and  that self-assemble as  but not as  or 
- ▶ Physics+logic: requires bi-level optimization (NP^{NP} -complete) (Vucinic et al. 2020)
- ▶ Compare Effie+tb2 (NP-complete) with ProteinMPNN, bi-criteria (Buchet et al. 2024)



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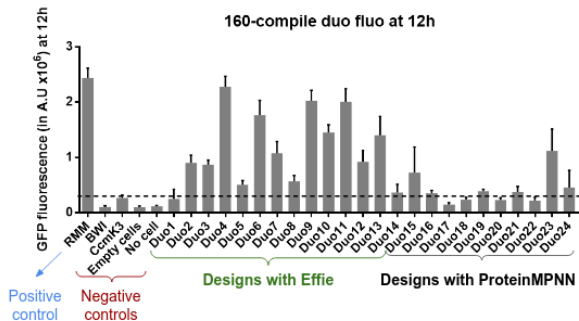


How often is better than ?

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A Neural Net, a CFN and a WCSP solver in a hybrid autoencoder

- ▶ A hybrid generic Generative AI that benefits from each component
- ▶ Neural Network: ideal to extract a representation of $P(X|\Phi)$ from raw inputs
- ▶ Represented as a CFN in a fully explorable and controllable latent layer
- ▶ Using decoding by discrete reasoning (toulbar2)
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What is missing in this architecture to make you happy?

When is Predict-and-optimize useful?

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Acknowledgments



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

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