



How to Turn Your Knowledge Graph Embeddings into Generative Models

Lorenzo Loconte

University of Edinburgh, UK

lorelloc.github.io

Nicola Di Mauro

University of Bari, Italy

Robert Peharz

TU Graz, Austria

Antonio Vergari

University of Edinburgh, UK

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Outline

1

A family of generative models of triples in KGs

based on KGEs and probabilistic circuits

2

Reliable link prediction with logical constraints

ensuring trustworthiness w.r.t. background knowledge

3

Efficiency and experimental results

train for hours rather than days, yet be competitive

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based on KGEs and probabilistic circuits

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Reliable link prediction with logical constraints

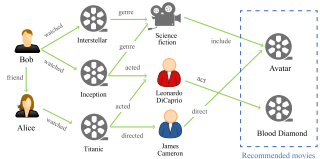
ensuring trustworthiness w.r.t. background knowledge

3

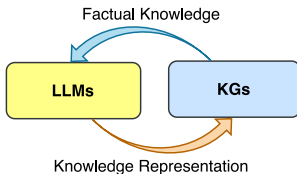
Efficiency and experimental results

train for hours rather than days, yet be competitive

Knowledge graphs



Item recommendation



Augment LLMs

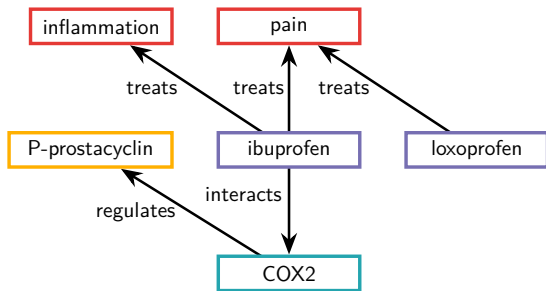


Drug discovery

Guo et al., "A Survey on Knowledge Graph-Based Recommender Systems", 2020

Pan et al., "Unifying Large Language Models and Knowledge Graphs: A Roadmap", 2023

Gogleva et al., "Knowledge Graph-based Recommendation Framework Identifies [...] Resistance in [...] Cell Lung Cancer", 2021



- Drugs
- Proteins
- Symptoms
- Functions

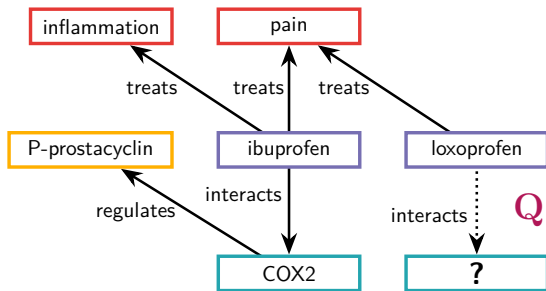
⟨loxoprofen, treats, pain⟩

⟨ibuprofen, treats, pain⟩

⋮

⟨COX2, regulates, P-prostacyclin⟩

⟨ibuprofen, interacts, COX2⟩



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⟨COX2, regulates, P-prostacyclin⟩

⟨ibuprofen, interacts, COX2⟩

Q: ⟨loxoprofen, interacts, ?⟩

KGE models

Knowledge graph embeddings (KGE) models such as ...

Complex Embeddings for Simple Link Prediction

[Théo Trouillon](#), [Johannes Welbl](#), +2 authors [Guillaume Bouchard](#) • Published in International Conference on... 19 June 2016 •

2,142 Citations

Highly Influential Citations 

576

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$$\phi_{\text{Complex}}(s, r, o) := f(\mathbf{e}_s, \mathbf{w}_r, \mathbf{e}_o) \in \mathbb{R} \quad \mathbf{e}_s, \mathbf{w}_r, \mathbf{e}_o \in \mathbb{C}^d$$

KGE models

Knowledge graph embeddings (KGE) models such as ...

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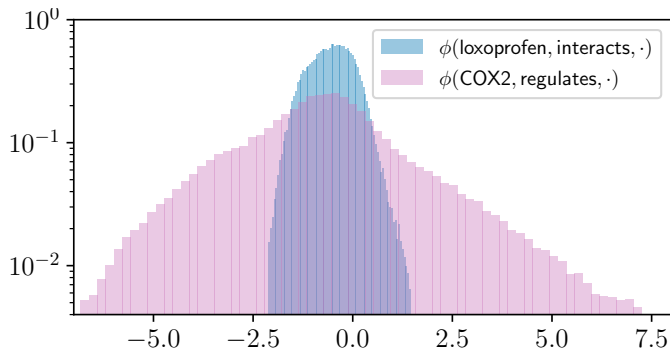
$$\phi_{\text{CompEx}}(s, r, o) := f(\mathbf{e}_s, \mathbf{w}_r, \mathbf{e}_o) \in \mathbb{R} \quad \mathbf{e}_s, \mathbf{w}_r, \mathbf{e}_o \in \mathbb{C}^d$$

$$1^{\text{st}} \quad \phi_{\text{CompEx}}(\text{loxoprofen, interacts, } \mathbf{\text{phosp-acid}}) = 1.17 \Leftarrow$$

$$2^{\text{nd}} \quad \phi_{\text{CompEx}}(\text{loxoprofen, interacts, COX2}) = 0.95$$

⋮

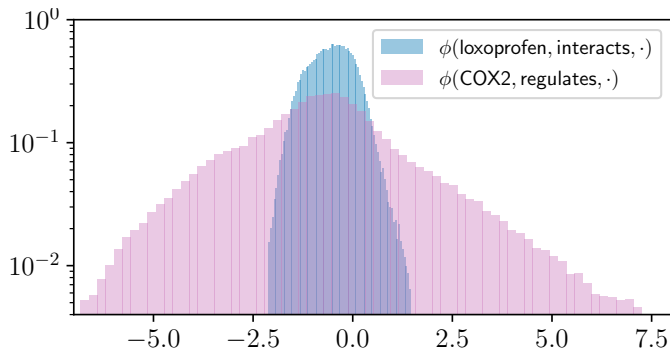
***“Real-valued **scores** are difficult
to **interpret** and **compare**”***



Scores have different orders of magnitude...

Arakelyan, Minervini, and Augenstein, Adapting Neural Link Predictors for Complex Query Answering, 2023

Zhu et al., "A Closer Look at Probability Calibration of Knowledge Graph Embedding", 2023



We would like *triples probabilities* instead !

Arakelyan, Minervini, and Augenstein, Adapting Neural Link Predictors for Complex Query Answering, 2023

Zhu et al., "A Closer Look at Probability Calibration of Knowledge Graph Embedding", 2023

Solutions! (1/3)

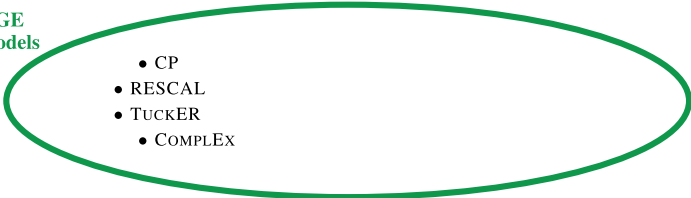
I

Generative models of triples (GeKCs)

calibrated probabilistic predictions by modelling $p(S, R, O)$
sampling of new triples (more later!)

From KGE models ...

**KGE
models**

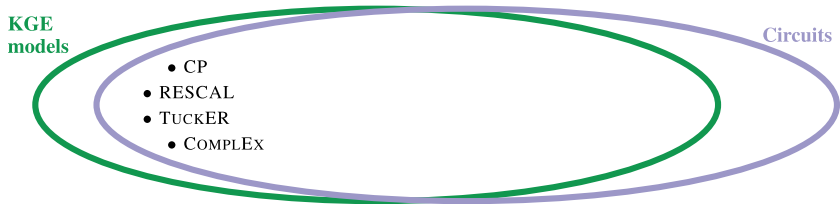
- 
- CP
 - RESCAL
 - TUCKER
 - COMPLEX

Lacroix, Usunier, and Obozinski, "Canonical Tensor Decomposition for Knowledge Base Completion", 2018

Nickel, Tresp, and Kriegel, "A Three-Way Model for Collective Learning on Multi-Relational Data", 2011

Balazevic, Allen, and Hospedales, "TUCKER: Tensor Factorization for Knowledge Graph Completion", 2019

From KGE models to circuits ...



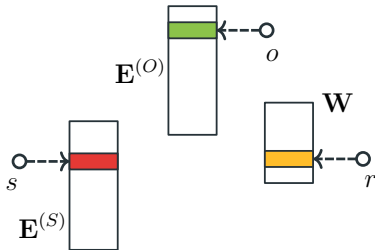
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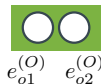
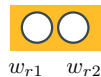
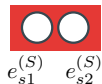
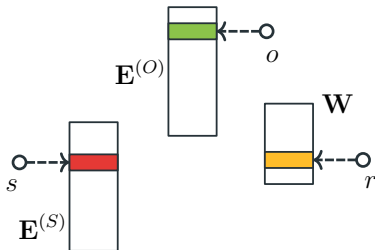
Canonical Polyadic (CP) KGE as a circuit

$$\phi(s, r, o) = \sum_{i=1}^R e_{si}^{(S)} w_{ri} e_{oi}^{(O)}$$



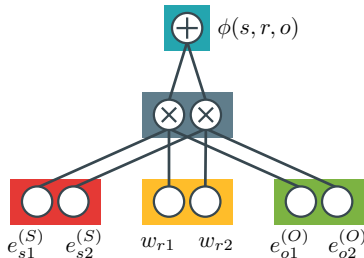
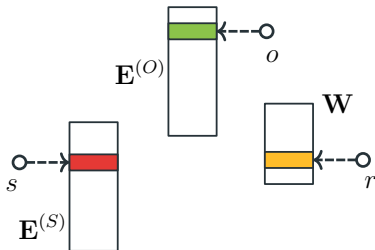
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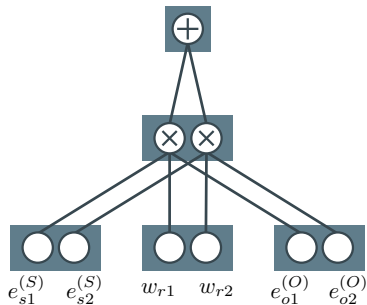


A circuit scoring triples

$$\mathbf{E}^{(S)} = \begin{bmatrix} 0.1 & 1.2 \\ 3.5 & -0.2 \\ -0.1 & 0.2 \end{bmatrix}$$

$$\mathbf{W} = \begin{bmatrix} 2.5 & 0.0 \\ -3.4 & -0.5 \\ -0.1 & 2.2 \end{bmatrix}$$

$$\mathbf{E}^{(O)} = \begin{bmatrix} -2.3 & 1.0 \\ 0.8 & -2.4 \\ 0.7 & 1.5 \end{bmatrix}$$



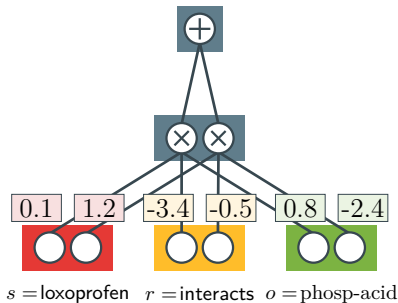
$$\phi_{\text{CP}}(s, r, o) = \sum_{i=1}^R e_{si}^{(S)} w_{ri} e_{oi}^{(O)}$$

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$$\phi_{\text{CP}}(\text{loxoprofen}, \text{interacts}, \text{phosp-acid})$$

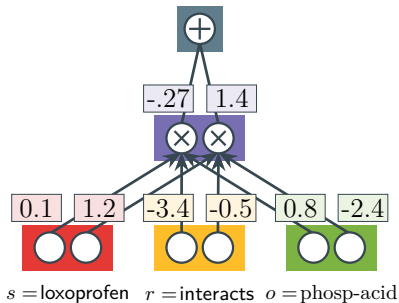
$$= \sum_{i=1}^R e_{\text{loxoprofen},i}^{(S)} w_{\text{interacts},i} e_{\text{phosp-acid},i}^{(O)}$$

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$$\phi_{\text{CP}}(\text{loxoprofen}, \text{interacts}, \text{phospho-acid})$$

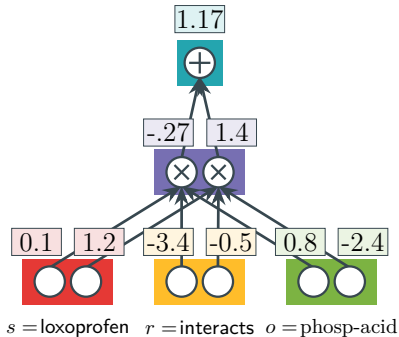
$$= \sum_{i=1}^R e_{\text{loxoprofen},i}^{(S)} w_{\text{interacts},i} e_{\text{phospho-acid},i}^{(O)}$$

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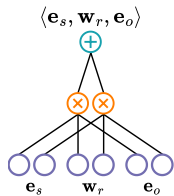
$$\mathbf{E}^{(O)} = \begin{bmatrix} -2.3 & 1.0 \\ 0.8 & -2.4 \\ 0.7 & 1.5 \end{bmatrix}$$



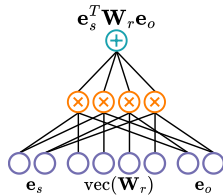
$$\phi_{\text{CP}}(\text{loxoprofen}, \text{interacts}, \text{phospho-acid})$$

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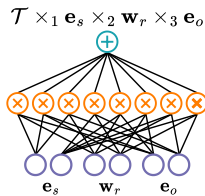
From KGE models to circuits ...



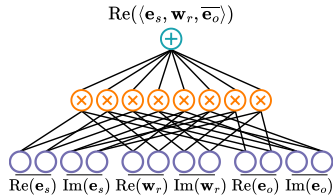
ϕ_{CP}



ϕ_{RESCAL}

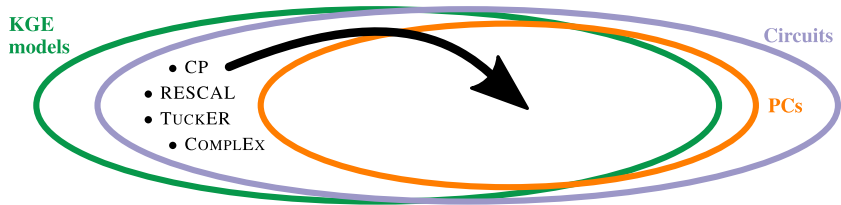


ϕ_{Tucker}



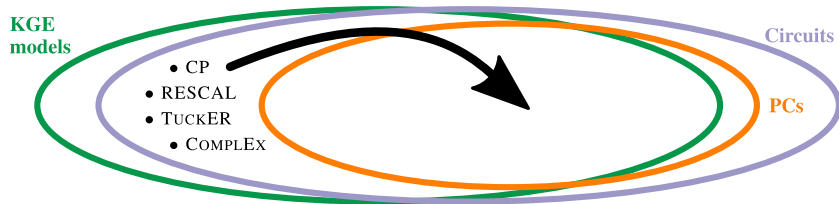
ϕ_{Complex}

... to probabilistic circuits



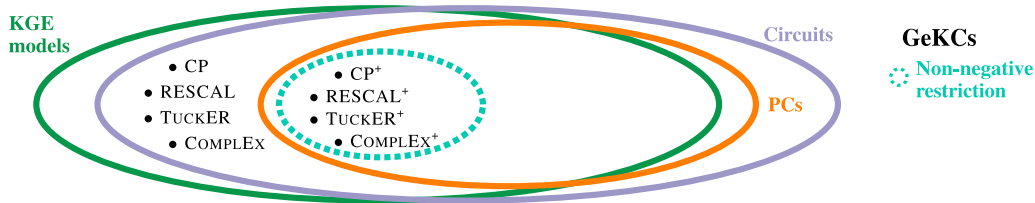
From scores $\phi(s, r, o)$ to **triple probabilities** $p(s, r, o)$

... to probabilistic circuits



1. Ensure $\phi(s, r, o) \geq 0$, $p(s, r, o) = \frac{1}{Z} \cdot \phi(s, r, o)$

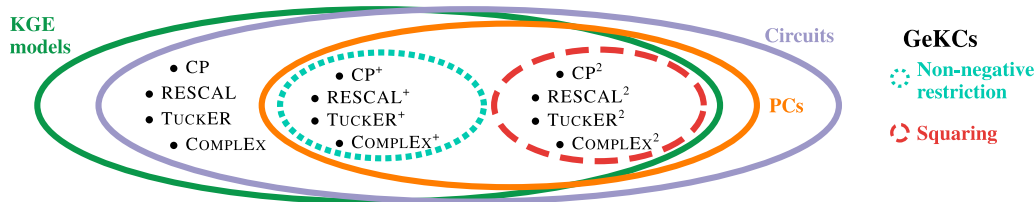
... to probabilistic circuits



Enforce *non-negative embeddings*

⇒ Less accurate on link prediction ...

... to probabilistic circuits



Square score functions (unrestricted embeddings)

⇒ Competitive on link prediction !

Squared circuits

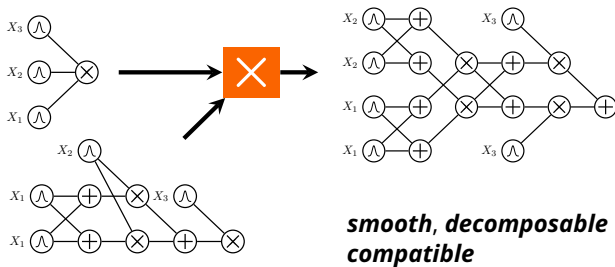
$$p(\mathbf{x}) = \frac{1}{Z} \phi^2(\mathbf{x}) = \frac{1}{Z} \phi(\mathbf{x}) \cdot \phi(\mathbf{x}), \quad \phi(\mathbf{x}) \in \mathbb{R}$$

where parameters and input functions can be **negative**

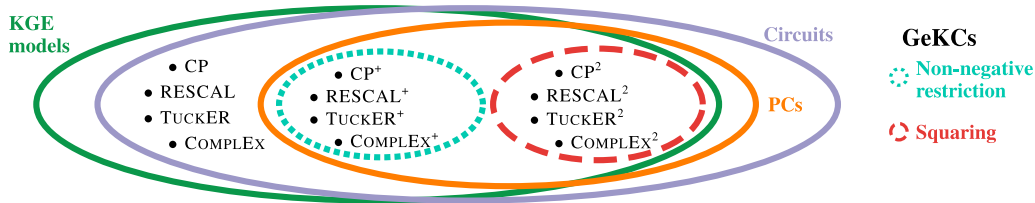
Squared circuits

$$p(\mathbf{x}) = \frac{1}{Z} \phi^2(\mathbf{x}) = \frac{1}{Z} \phi(\mathbf{x}) \cdot \phi(\mathbf{x}), \quad \phi(\mathbf{x}) \in \mathbb{R}$$

Tractable product

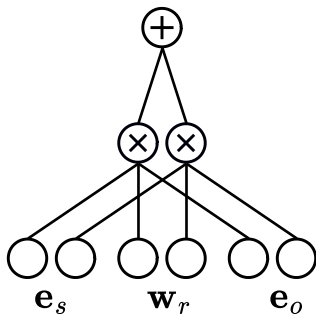


... to probabilistic circuits



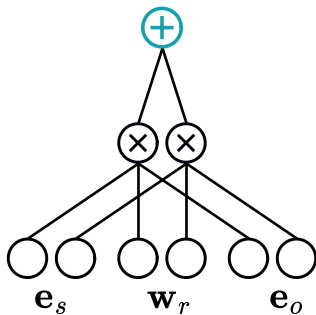
1. Ensure $\phi(s, r, o) \geq 0$, $p(s, r, o) = \frac{1}{Z} \cdot \phi(s, r, o)$

2. Computation of $Z = \sum_{s \in \mathcal{E}, r \in \mathcal{R}, o \in \mathcal{E}} \phi(s, r, o)$



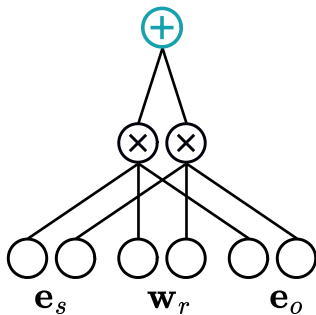
$$Z = \sum_{s \in \mathcal{E}, r \in \mathcal{R}, o \in \mathcal{E}} \phi_{\text{CP}^+}(s, r, o)$$

The summation over triples computing Z ...



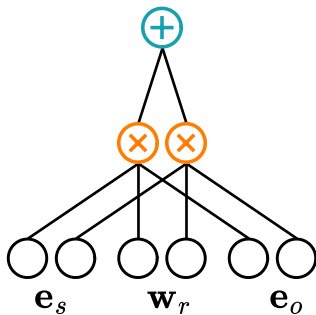
$$Z = \sum_{s \in \mathcal{E}, r \in \mathcal{R}, o \in \mathcal{E}} \sum_{i=1}^d e_{si}^{(S)} w_{ri} e_{oi}^{(O)}$$

The summation over triples computing Z ...



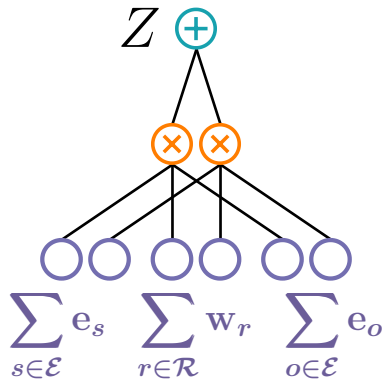
$$Z = \sum_{i=1}^d \sum_{s \in \mathcal{E}, r \in \mathcal{R}, o \in \mathcal{E}} e_{si}^{(S)} w_{ri} e_{oi}^{(O)}$$

... can be pushed (*smoothness*)



$$Z = \sum_{i=1}^d \left(\sum_{s \in \mathcal{E}} e_{si}^{(S)} \right) \times \left(\sum_{r \in \mathcal{R}} w_{ri} \right) \times \left(\sum_{o \in \mathcal{E}} e_{oi}^{(O)} \right)$$

... and broken down (*decomposability*) ...



... thus requiring linear time w.r.t. $|\mathcal{E}|$

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based on KGEs and probabilistic circuits

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Reliable link prediction with logical constraints

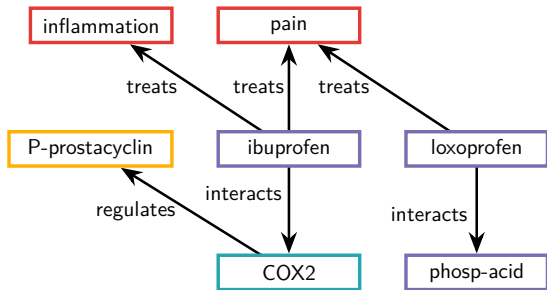
ensuring trustworthiness w.r.t. background knowledge

3

Efficiency and experimental results

train for hours rather than days, yet be competitive

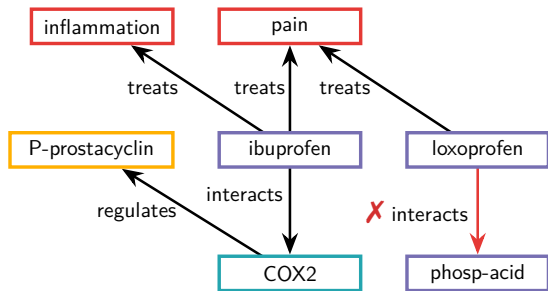
***“KGE models predictions **violate**
even simple **logical constraints**”***



- Drugs
- Symptoms
- Proteins
- Functions

Q: $\langle \text{loxoprofen}, \text{interacts}, ? \rangle$

A: $\langle \text{loxoprofen}, \text{interacts}, \mathbf{\text{phosp-acid}} \rangle$



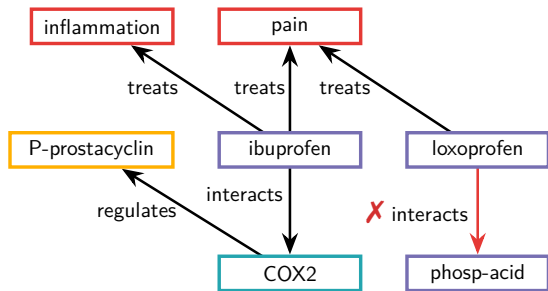
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A: $\langle \text{loxoprofen}, \text{interacts}, \mathbf{\text{phosp-acid}} \rangle$

X

“*interacts*” can only hold
between *drugs* and *proteins*



- Drugs
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- Proteins
- Functions

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A: $\langle \text{loxoprofen}, \text{interacts}, \mathbf{\text{phosp-acid}} \rangle$

X

“*interacts*” can only hold
between *drugs* and *proteins*

Complex predicts a triple violating a constraint !

Solutions! (2/3)

I

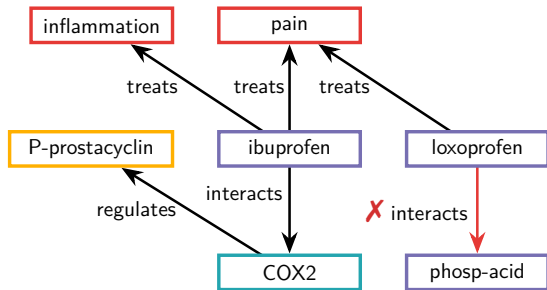
Generative models for KGs (GeKCs)

calibrated probabilistic predictions by modelling $p(S, R, O)$
sampling of new triples (more later!)

II

Integrate constraints with guarantees

such as the domain schema



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- Proteins
- Symptoms
- Functions

Q: $\langle \text{loxoprofen}, \text{interacts}, ? \rangle$

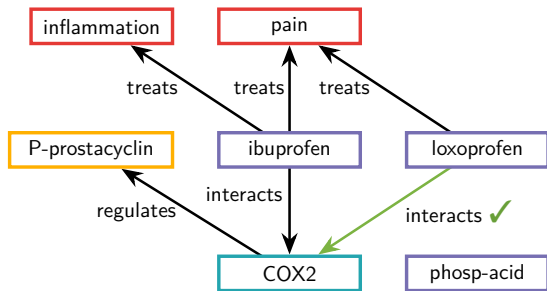
A: $\langle \text{loxoprofen}, \text{interacts}, \text{phosp-acid} \rangle$

X

"*interacts*" can only hold between *drugs* and *proteins*

$p(\text{loxoprofen}, \text{interacts}, \text{phosp-acid}) = 0$





- Drugs
- Proteins
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- Functions

Q: $\langle \text{loxoprofen}, \text{interacts}, ? \rangle$

A: $\langle \text{loxoprofen}, \text{interacts}, \mathbf{COX2} \rangle$



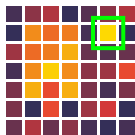
"*interacts*" can only hold between *drugs* and *proteins*

$p(\text{loxoprofen}, \text{interacts}, \text{phosp-acid}) = 0$

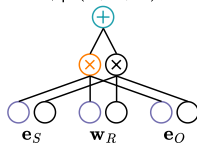
$p(\text{loxoprofen}, \text{interacts}, \mathbf{COX2}) > 0$

GeKCs for logical constraints

$\phi_{pc}(S, \text{interacts}, O)$

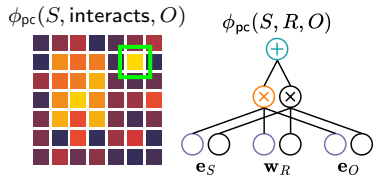
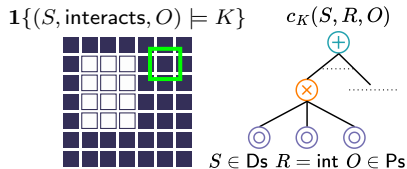


$\phi_{pc}(S, R, O)$



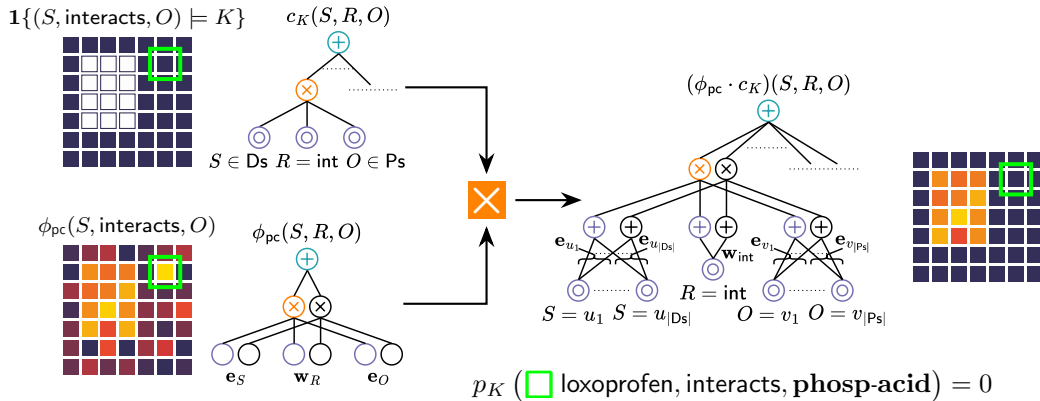
$$p_K(\text{loxoprofen}, \text{interacts}, \text{phosp-acid}) = 0$$

GeKCs for logical constraints



$$p_K(\text{loxoprofen, interacts, phosp-acid}) = 0$$

GeKCs for logical constraints



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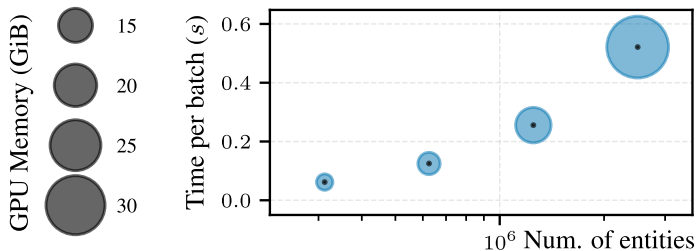
3

Efficiency and experimental results

train for hours rather than days, yet be competitive

***“Training on relatively **large**
knowledge graphs is **expensive**”***

Some benchmarks...



$107 \cdot 10^6$ entities

Solutions! (3/3)

I

Generative models for KGs (GeKCs)

calibrated probabilistic predictions by modelling $p(S, R, O)$
sampling of new triples (more later!)

II

Integrate constraints with guarantees

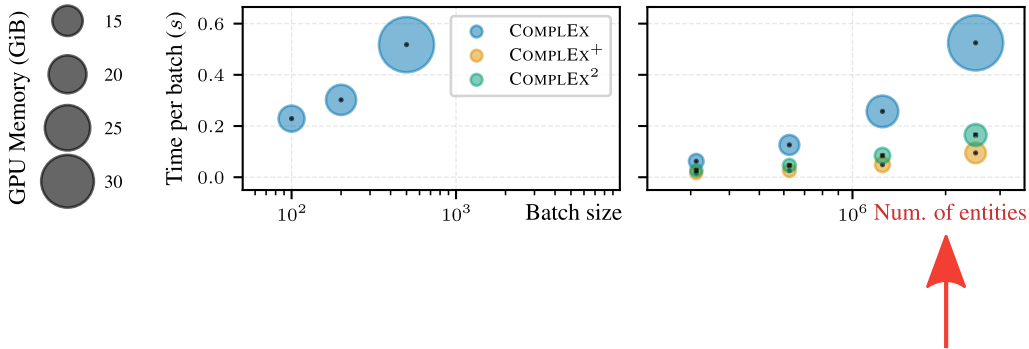
such as the domain schema

III

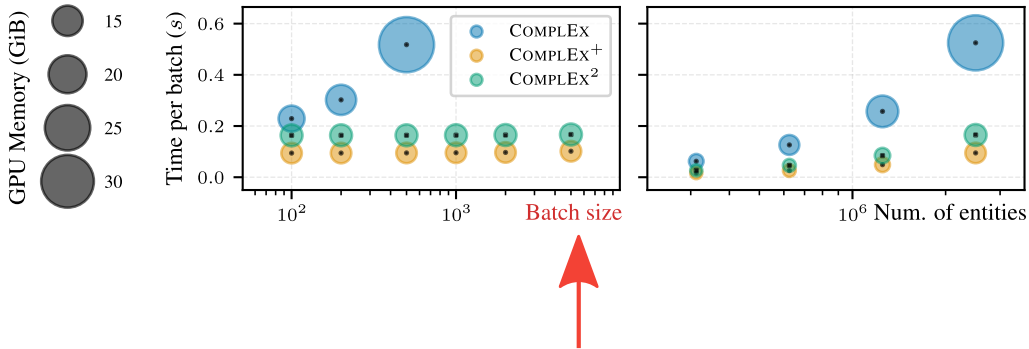
Scale to KGs with millions of entities and triples

using probabilistic training objectives

Speed-up training on large KGs



Speed-up training on large KGs



Learning ...

... by discriminative objectives (*pseudo-log-likelihood*)

$$\mathcal{L}_{\text{PLL}} := \sum_{(s,r,o) \in \mathcal{D}} w_s \log p(s \mid r, o) + w_r \log p(r \mid s, o) + w_o \log p(o \mid s, r)$$

Learning ...

... by discriminative objectives (**pseudo-log-likelihood**)

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... by **maximum-log-likelihood** estimation

$$\mathcal{L}_{\text{MLE}} := \sum_{(s,r,o) \in \mathcal{D}} \log p(s, r, o) = -|\mathcal{D}| \log \mathbf{Z} + \sum_{(s,r,o) \in \mathcal{D}} \log \phi_{\text{pc}}(s, r, o)$$

(faster, as marginalization require a single circuit evaluation)

Mean Reciprocal Rank (MRR) ↑

Model	FB15k-237	WN18RR	ogbl-biogk
CP	0.310	0.105	0.831
CP ⁺	0.237	0.027	0.496
CP ²	0.315	0.104	0.848
ComplEx	0.342	0.471	0.829
ComplEx ⁺	0.214	0.030	0.503
ComplEx ²	0.334	0.420	0.858

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GeKCs are competitive with KGE models ...

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... and achieve the best results on ogbl-biokg

Sampling triples

Kernel triple distance to measure their quality

Sampling triples

Kernel triple distance to measure their quality

Empirical KTD ↓

Model	FB15k-237		WN18RR		ogbl-biokg	
Uniform	0.589		0.766		1.822	
	PLL	MLE	PLL	MLE	PLL	MLE
ComplEx ²	0.326	0.102	0.338	0.278	0.104	0.034

Takeaways

I

A generative perspective of KGE models (GeKCs)

II

Reliable predictions with logical constraints

III

Speed-up training and reduce costs

Takeaways

Questions?

I

A generative perspective of KGE models (GeKCs)

II

Reliable predictions with logical constraints

III

Speed-up training and reduce costs

More link prediction benchmarks

Mean Reciprocal Rank (MRR) ↑

Model	FB15k-237		WN18RR		ogbl-biokg	
	PLL	MLE	PLL	MLE	PLL	MLE
CP	0.310	—	0.105	—	0.831	—
CP ⁺	0.237	0.230	0.027	0.026	0.496	0.501
CP ²	0.315	0.282	0.104	0.091	0.848	0.829
ComplEx	0.342	—	0.471	—	0.829	—
ComplEx ⁺	0.214	0.205	0.030	0.029	0.503	0.516
ComplEx ²	0.334	0.300	0.420	0.391	0.858	0.840

Instantiate GeKCs from KGE models

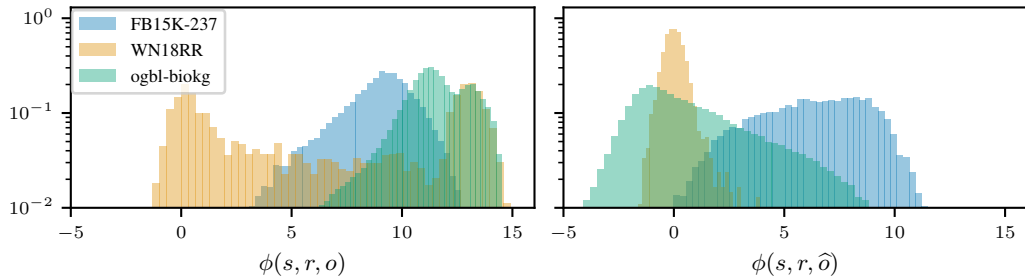
(as a way to initialize the parameters)

Mean Reciprocal Rank (MRR) ↑

Model	FB15k-237		WN18RR		ogbl-biokg	
	PLL	MLE	PLL	MLE	PLL	MLE
ComplEx	0.344	—	0.470	—	0.829	—
ComplEx ²	0.333	0.301	0.416	0.390	0.859	0.839
ComplEx ² ★	0.342	0.340	0.462	0.463	0.859	0.828

Scores are mostly non-negative

(thus squaring them has little effect on the rankings)



Semantic consistency scores

Sem@1 scores ↑

Model	Embedding size			
	10	50	200	1000
ComplEx	99.68	99.90	99.93	99.94
ComplEx ²	82.50	94.22	99.30	99.50
D-ComplEx ²	100.00	100.00	100.00	100.00