



How to Turn Your Knowledge Graph Embeddings into Generative Models

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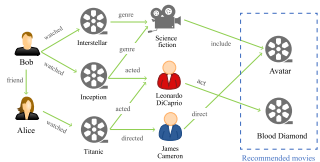
TU Graz, Austria

Antonio Vergari

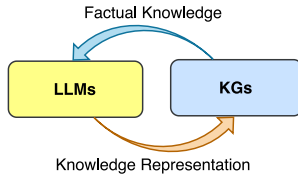
University of Edinburgh, UK

NeurIPS 2023 Oral

Knowledge graphs



Item recommendation



Augment LLMs

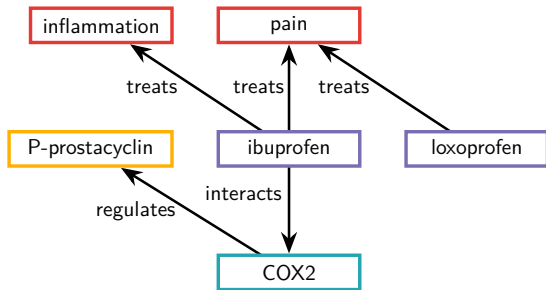


Drug discovery

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

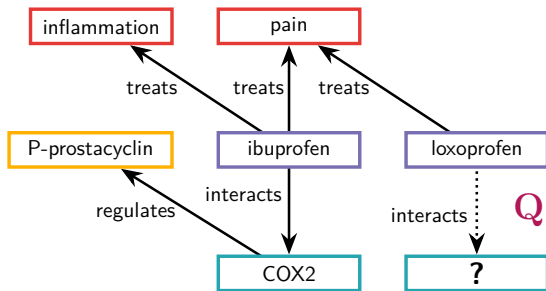
Pan et al., "Unifying Large Language Models and Knowledge Graphs: A Roadmap", 2023

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⟩



- Drugs
- Proteins
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⟨loxoprofen, treats, pain⟩

⟨ibuprofen, treats, pain⟩

⋮

⟨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 ·

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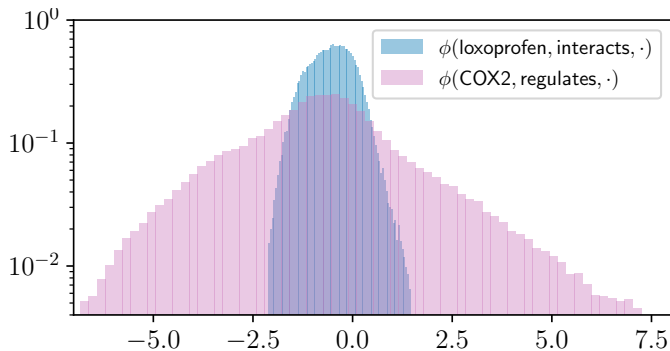
$$\phi_{\text{ComplEx}}(s, r, o) := f(\mathbf{e}_s, \mathbf{w}_r, \mathbf{e}_o) \in \mathbb{R} \quad \mathbf{e}_s, \mathbf{w}_r, \mathbf{e}_o \in \mathbb{C}^d$$

$$1^{\text{st}} \quad \phi_{\text{ComplEx}}(\text{loxoprofen, interacts, **phosp-acid**}) = 2.3 \leftarrow$$

$$2^{\text{nd}} \quad \phi_{\text{ComplEx}}(\text{loxoprofen, interacts, COX2}) = 1.3$$

⋮

Scores ...

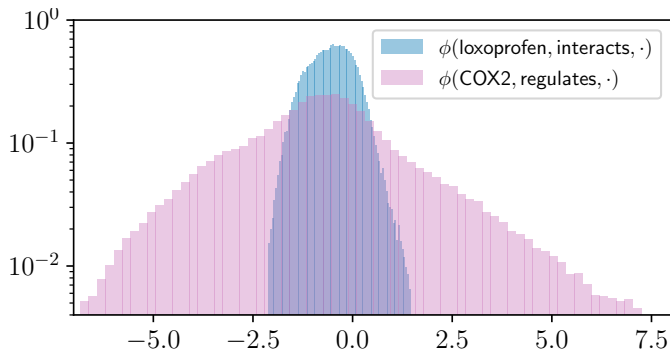


... are difficult to interpret and compare

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

Scores ...



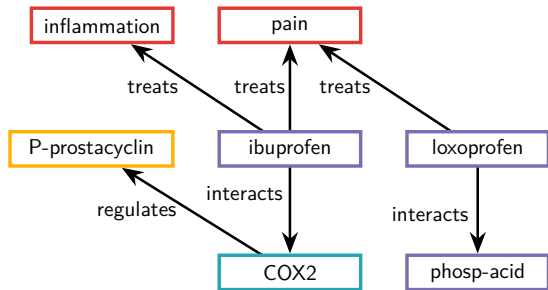
We would like *probabilities* instead !

Issues?

I

How to measure the confidence of predictions ?

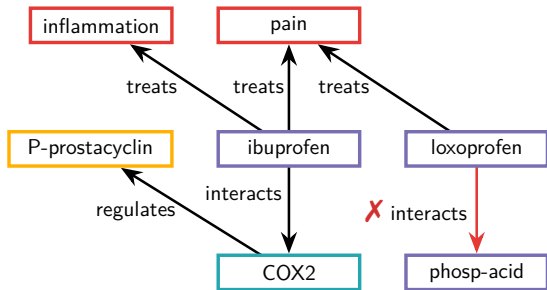
and compare / combine scores



- Drugs
- Proteins
- Symptoms
- Functions

Q: ⟨loxoprofen, interacts, ?⟩

A: ⟨loxoprofen, interacts, **phosp-acid**⟩



- Drugs
- Proteins
- Symptoms
- Functions

Q: ⟨loxoprofen, interacts, ?⟩

A: ⟨loxoprofen, interacts, **phosp-acid**⟩

X

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

Issues?

I

How to measure the confidence of predictions ?

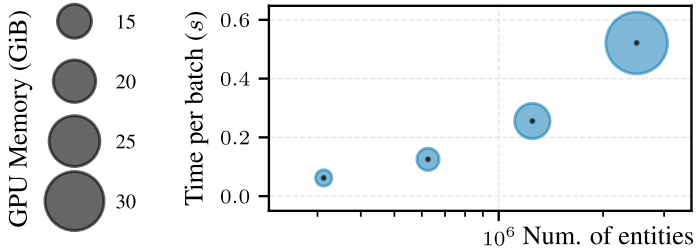
and compare / combine scores

II

How to guarantee the satisfaction of constraints ?

such as domain knowledge

Training is expensive



$107 \cdot 10^6$ entities

Issues?

I

How to measure the confidence of predictions ?

and compare / combine scores

II

How to guarantee the satisfaction of constraints ?

such as domain knowledge

III

How to scale to KGs with millions of entities ?

and be memory efficient

Solutions!

I

Generative models of triples (GeKCs)

calibrated probabilistic predictions, sampling of new triples

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 ...

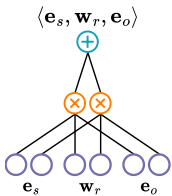
KGE
models

- CP
- RESCAL
- TUCKER
- COMPLEX

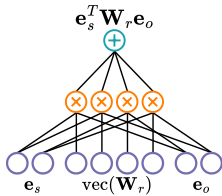
Circuits

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

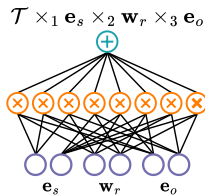
From KGE models to circuits ...



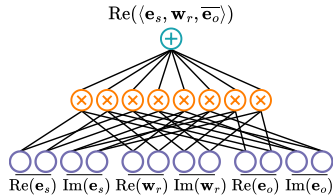
ϕ_{CP}



ϕ_{RESCAL}

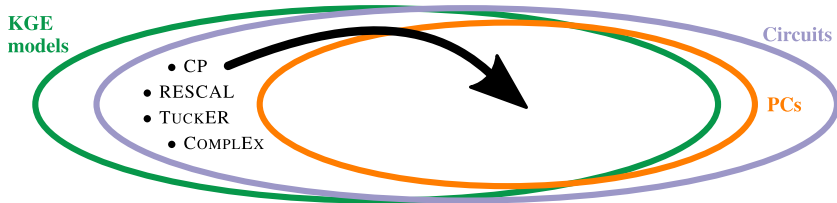


ϕ_{Tucker}



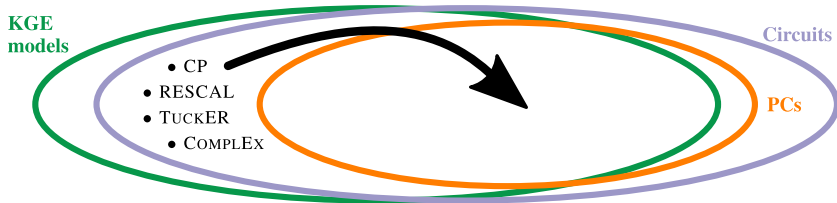
ϕ_{Complex}

... to probabilistic circuits



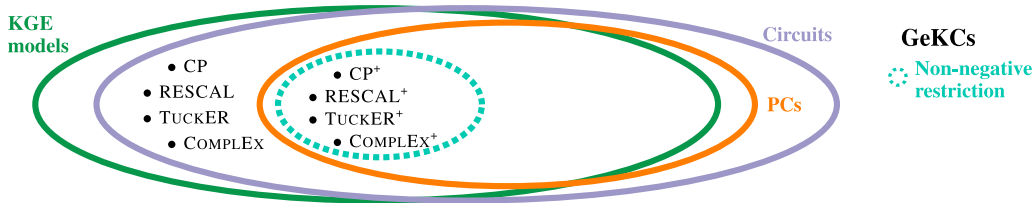
From scores $\phi(s, r, o)$ to **triple probabilities** $p(s, r, o)$

... to probabilistic circuits



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

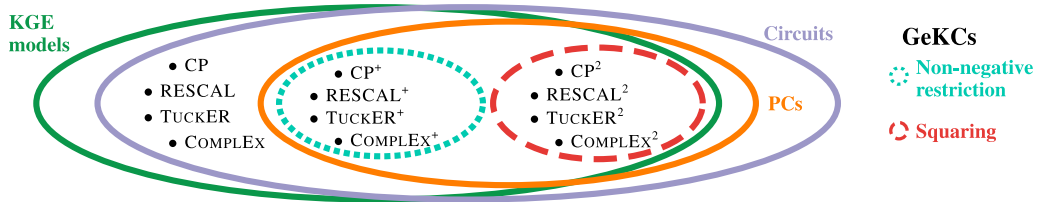
... to probabilistic circuits



Enforce *non-negative embeddings*

⇒ Less accurate on link prediction ...

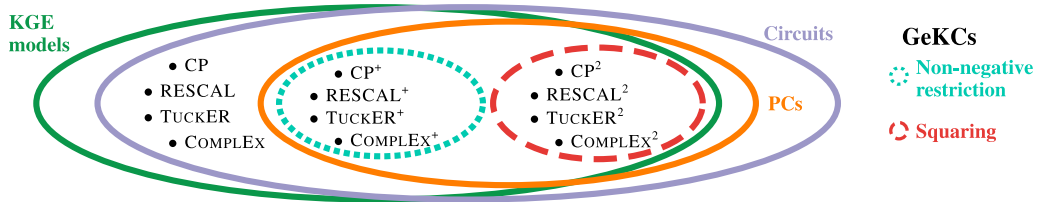
... to probabilistic circuits



Square score functions (unrestricted embeddings)

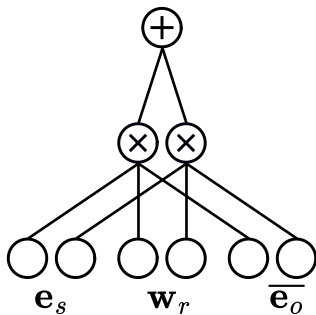
⇒ Competitive on link prediction !

... to probabilistic circuits



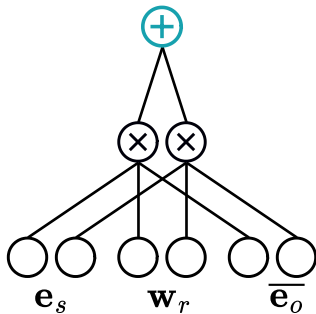
1. Ensure $\phi(s, r, o) \geq 0$, $p(s, r, o) = \frac{1}{Z} \cdot \phi(s, r, o)$

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



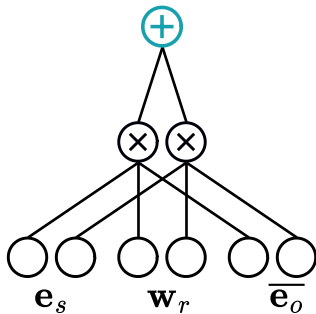
$$Z = \sum_{s \in \mathcal{E}, r \in \mathcal{R}, o \in \mathcal{E}} \phi_{\text{Complex}^+}(s, r, o)$$

The summation over triples computing Z ...



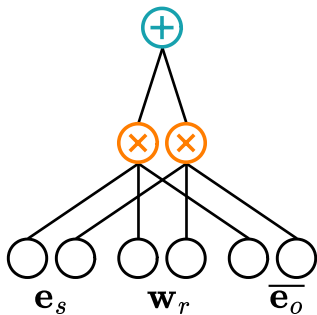
$$Z = \sum_{s \in \mathcal{E}, r \in \mathcal{R}, o \in \mathcal{E}} \sum_{i=1}^d \text{Re} (e_{si} w_{ri} \overline{e_{oi}})$$

The summation over triples computing Z ...



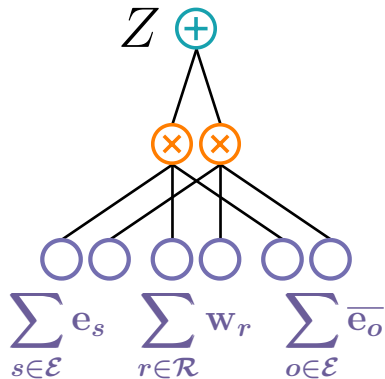
$$Z = \sum_{i=1}^d \sum_{s \in \mathcal{E}, r \in \mathcal{R}, o \in \mathcal{E}} \text{Re}(\mathbf{e}_{si} \mathbf{w}_{ri} \overline{\mathbf{e}_{oi}})$$

... can be pushed ...



$$Z = \sum_{i=1}^d \text{Re} \left[\left(\sum_{s \in \mathcal{E}} \mathbf{e}_{si} \right) \times \left(\sum_{r \in \mathcal{R}} \mathbf{w}_{ri} \right) \times \left(\sum_{o \in \mathcal{E}} \overline{\mathbf{e}_{oi}} \right) \right]$$

... and broken down ...



... thus requiring linear time !

Solutions!

I

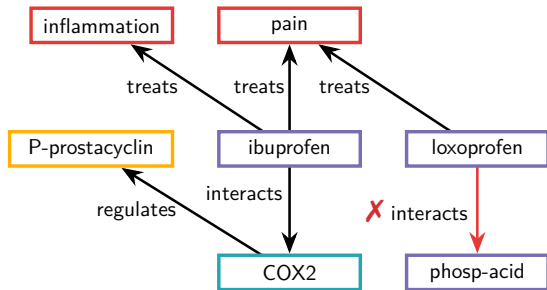
Generative models for KGs (GeKCs)

calibrated probabilistic predictions, sampling of new triples

II

Integrate constraints with guarantees

such as the domain schema



- Drugs
- Proteins
- Symptoms
- Functions

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

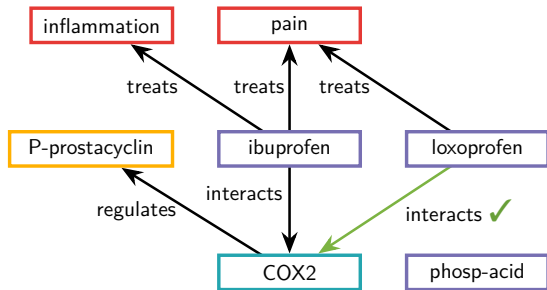
A: $\langle \text{loxoprofen, interacts, phosp-acid} \rangle$

X

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

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





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- Proteins
- Symptoms
- Functions

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

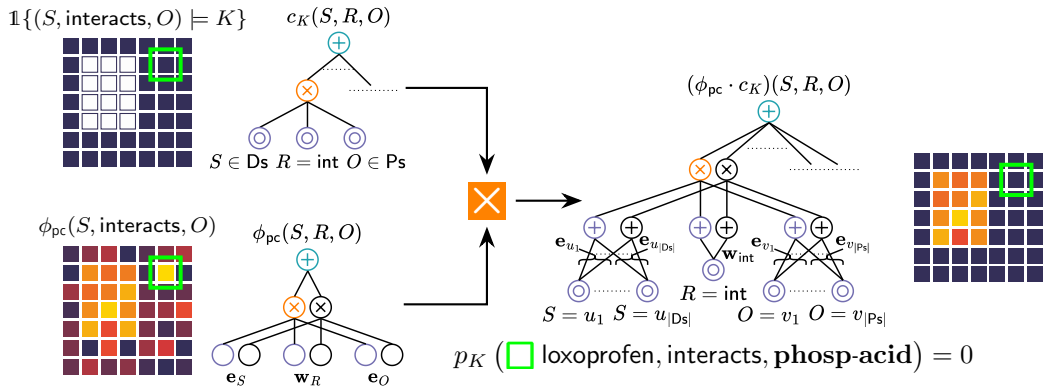
A: $\langle \text{loxoprofen, interacts, COX2} \rangle$



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

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

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



Logical constraints

+

GeKCs

Solutions!

I

Generative models for KGs (GeKCs)

calibrated probabilistic predictions, sampling of new triples

II

Integrate constraints with guarantees

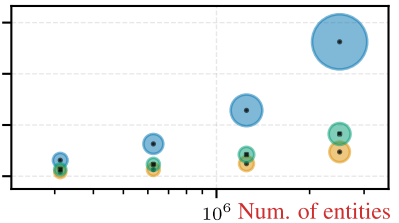
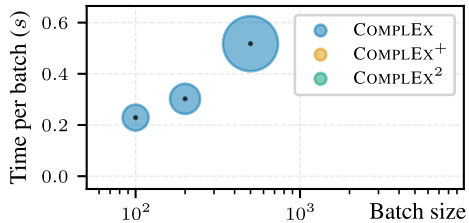
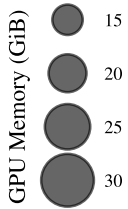
such as the domain schema

III

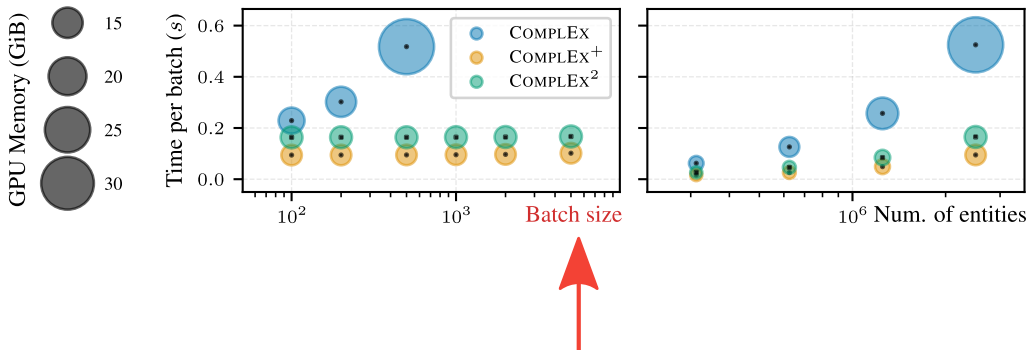
Scale to KGs with millions of entities and triples

speed-up training and save memory

Speed-up training on large KGs



Speed-up training on large KGs



Learning ...

... by discriminative objectives,
generalised as a weighted *pseudo-log-likelihood*

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

Learning ...

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

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

Mean Reciprocal Rank (MRR) ↑

Model	FB15k-237	WN18RR	ogbl-biokg
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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Complex	0.342	0.471	0.829
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GeKCs are competitive with KGE models ...

Mean Reciprocal Rank (MRR) ↑

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Complex ⁺	0.214	0.030	0.503
Complex ²	0.334	0.420	0.858

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

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

I *A generative perspective
of KGE models (GeKCs)*

II *Reliable predictions
with logical constraints*

III *Speed-up training
and reduce costs*

more on circuits

A. Vergari, Y. Choi, and R. Peharz

*Probabilistic Circuits: representations,
inference, learning and applications*

Tutorial @ NeurIPS 2022

Z. Yu, M. Trapp and K. Kersting

Characteristic circuits

Oral @ NeurIPS 2023

Takeaways

I *A generative perspective of KGE models (GeKCs)*

II *Reliable predictions with logical constraints*

III *Speed-up training and reduce costs*

april

april-tools.github.io

about probabilities, reasoning, integrals & logic

Poster Session 1 #1205

Paper



Code



  @lorelloc_

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

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

Semantic consistency scores

Sem@ k scores ↑

Model	k	Embedding size			
		10	50	200	1000
Complex	1	99.68	99.90	99.93	99.94
	20	99.81	99.79	99.85	99.91
	100	99.60	99.44	99.60	99.77
Complex ²	1	82.50	94.22	99.30	99.50
	20	86.50	96.70	99.42	99.64
	100	90.66	97.71	99.23	98.78
D-Complex ²	1	100.00	100.00	100.00	100.00
	20	100.00	100.00	100.00	100.00
	100	100.00	100.00	100.00	100.00

Logical constraints improve small GeKCs

