

Sharing Parameter by Conjugation for Knowledge Graph Embeddings in Complex Space

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Abstract

A Knowledge Graph (KG) is the directed graphical representation of entities and relations in the real world. KG can be applied in diverse Natural Language Processing (NLP) tasks where knowledge is required. The need to scale up and complete KG automatically yields Knowledge Graph Embedding (KGE), a shallow machine learning model that is suffering from memory and training time consumption issues. To mitigate the computational load, we propose a parameter-sharing method, i.e., using conjugate parameters for complex numbers employed in KGE models. Our method improves memory efficiency by 2x in relation embedding while achieving comparable performance to the state-of-the-art non-conjugate models, with faster, or at least comparable, training time. We demonstrated the generalizability of our method on two best-performing KGE models 5★E (Nayyeri et al., 2021) and ComplEx (Trouillon et al., 2016) on five benchmark datasets.

1 Introduction

A Knowledge Graph (KG) is a representation of confident information in the real world and employed in diverse Natural Language Processing (NLP) applications, e.g., recommender system, question answering, and text generation. A triple in the form of $(head, relation, tail)$ is widely used as the representation of elements in the KG instead of raw text for scalability. Cite $(clinician, synset_domain_topic_of, psychology)$ as an example, *clinician* and *psychology* is the head and tail entity respectively, and *synset_domain_topic_of* is the relation of the head entity pointing to the tail entity.

Knowledge Graph Embedding (KGE) models are designed for automatic link prediction. Relations in KG have multiple categories, e.g., symmetry, antisymmetry, inversion, and hierarchical. Missing links indicate incomplete ties between en-

tities and are a common phenomenon as finding the missed connections is labor-intensive work.

The theoretical space complexity of KGE models are often $\mathcal{O}(n_e d_e + n_r d_r)$, which is proportional to the number of KG elements, i.e. entities n_e and relations n_r , and embedding dimension d_e, d_r respectively. Scaling a KG is problematic as n_e, n_r can go up to millions; also because KGE models are often shallow machine learning models composed of simple operations, e.g., matrix multiplication. Caution that a shallow model needs a large dimension size d to depict the data feature, yielding the issue of the drastic increase of embedding parameters (Dettmers et al., 2018).

KGE models represented using complex numbers have state-of-the-art performance, while they demand high memory costs. E.g., if using one of the best models ComplEx (Trouillon et al., 2016) to create embedding for the benchmark dataset FB15K whose $n_e = 14,951, n_r = 1,345$, and the best-performing dimensionalities $d_e = 4000, d_r = 4000$, will result in the parameter size of 65,184,000. Considering the data type 64-bit integer (signed), who has a size of 8 bytes in PyTorch, the memory cost will be $65,184,000 \times 8 \approx 497$ MB. A KG for real-world application could have a much larger size, e.g., IBM’s KG contains entities > 100 million and relations > 5 billion, which is actively in use and continually growing (Noy et al., 2019), would need > 148 TB memory to do link prediction task.

Inspired by the improved performance of complex number representation and Non-Euclidean models where transformation parameters attempt to interact rather than be independent, we intuited the idea of sharing parameters for memory efficiency.

We demonstrate a parameter-sharing method for complex numbers employed in KGE models. Specifically, our method formulates conjugate parameters in appropriate dimensions of the transformation functions to reduce relation parameters. By

using our method, models can reduce their space complexity to $\mathcal{O}(n_e d_e + n_r d_r / 2)$, which means the relation embedding size is half the original model. In the second place, using conjugate parameters may help save training time, especially on the datasets who have more parameter patterns. Further, our method can be easily applied to various complex number represented models.

We verified our method on two best-performing KGE models, i.e., ComplEx (Trouillon et al., 2016) and 5★E (Nayyeri et al., 2021). The experiments were conducted on five benchmark datasets, i.e., FB15K-237, WN18RR, YAGO3-10, FB15K, and WN18, by which we empirically show that our method reserves the models’ ability to achieve state-of-the-art results. We also see 31% training time saved on average for 5★E in addition to the memory. Our method is implemented in PyTorch¹ and the code with hyperparameter settings² are available online.

2 Related Works

We describe the categorizations of KGE models according to the representation method and the vector space that inspired our idea.

Representation Method Real and complex number representations are used to quantify entities and relations.

Translation approaches including TransE (Bordes et al., 2013) and its variants (Ji et al., 2015; Lin et al., 2015) describe embeddings using real number representation. Although these simple models cost fewer parameters, they can only encode two or three relation patterns, e.g., TransE cannot encode symmetric relations.

ComplEx (Trouillon et al., 2016) creates embedding with complex number representation, which can handle a wider variety of relations than using only real numbers, among them symmetric and antisymmetric relations (Trouillon et al., 2016). 5★E (Nayyeri et al., 2021) utilizes Möbius transformation, a projective geometric function that supports multiple simultaneous transformations in complex number representation and can embed entities in much lower ranks.

Vector Space Euclidean and Non-Euclidean spaces are practiced for the calculation of triple plausibility.

Factorization models such as RESCAL (Nickel et al., 2011) and DistMult (Yang et al., 2014) employ element-wise multiplication in Euclidean space. Correspondingly, the plausibility of a triple is measured according to the angle of transformed head and tail entities.

MuRP (Balazevic et al., 2019) minimizes hyperbolic distances other than Euclidean. It needs fewer parameters than its Euclidean analog. ATTH (Chami et al., 2020) leverages trainable hyperbolic curvatures for each relation to simultaneously capture logical patterns and hierarchies. Compared with Euclidean, the Hyperbolic models can save more structures using variational curvatures in different areas to depict hierarchical relations.

Relational Constrain on Parameters Replacing real number with complex number representation enables the imaginary part to have an effect on the real part parameters, the boosted performance of which indicates the **hidden relation** among parameter. Using hyperbolic space other than Euclidean enables the distances or angles at different positions to vary, the increased accuracy hints us to add **various constraints** on parameters. Learning from the work by Hayashi and Shimbo (2017), the potential of improving representations through conjugate symmetric constraint is revealed. Therefore, we hypothesize the efficiency of relational parameters and propose a parameter-sharing method using conjugate numbers.

3 Method

Complex number employed in current KGE models enforces **multiplicative constraint** on representations; our method further adds **conjugate constraint** within the parameters. Note that we don’t reduce the dimensions of the parameters, instead, we share the dimensions.

We economize 50% of the memory in relation embedding by sharing half of the parameters in the conjugate form. Our approach is at least comparable in accuracy to the baselines. In addition, our method reduces calculation in the regularization process, e.g., for the 5★E model, 31% of training time is saved on average for five benchmark datasets.

3.1 Preliminaries

Link prediction task inquires if a triple (h, r, t) constructed by existing head and tail entities $h, t \in \mathbb{V}^{d_e}$

¹<https://pytorch.org/>

²github.com/xincanfeng/dimension

and relations $r \in \mathbb{V}^{d_r}$ (\mathbb{V}^d is a d -dimensional vector space) is true or not. In KGE models, the relations are often represented as the transformation function ϑ that maps a head entity into a tail entity which are described as vectors in corresponding space, i.e., $\vartheta(h) = t$. Then, the score function $f : \mathbb{V}^{d_e} \times \mathbb{V}^{d_r} \times \mathbb{V}^{d_e} \rightarrow \mathbb{R}$ returns the plausibility p of constructing a true triple: $f(h, r, t) = p(\vartheta(h), t)$.

$a, b, c, d \in \mathbb{C}$ denote the parameters in the relation embedding matrices. $x \in \mathbb{C}$ is the parameter of the entity embedding matrices. a_i, x_i are the parameters of the submatrices of $[a]$ and $[x]$ respectively. $Re(z)$ is the real part of the complex number z , \bar{z} is the complex conjugate of z .

Complex This is the first and one of the best-performing complex models in Euclidean space. Trouillon et al. (2016) demonstrated that complex number multiplication could capture antisymmetric relations while retaining the efficiency of the dot product, i.e., linearity in both space and time complexity. Balancing between model expressiveness and parameter size is also discussed as the keystone of KGE. However, targeting SOTA is still computational-expensive because Trouillon et al. (2016) didn't solve the performance deterioration problem when reducing parameters directly.

Performance deterioration can be severe whenever the KG needs to be expanded because the mispredicted links could lead to further misinformation. Hence we should always endeavour to adopt the best-performing embedding size in doing link prediction task, even though it could be hundreds of TB.

To obtain the best results, Complex needs embedding size of $rank = 2000$ on dataset FB15K-237, WN18RR, FB15K, WN18, and $rank = 1000$ on dataset YAGO3-10. $rank$ denotes the vector dimension of a single-functional parameter. Each entity and relation in this model needs $2 \times rank$ parameters, representing real and imaginary part, respectively.

In this model, relations are represented as the real part of low-rank matrix $[a]$, which act as weights on each entity dimension x , followed by a projection onto the real subspace. The transformation of Complex is

$$x \rightarrow [a]x \rightarrow ax. \quad (1)$$

5★E This is a novel model applying complex numbers in Non-Euclidean space. Nayyeri et al. (2021) tackled the problem of multiple subgraph

structures in the neighborhood, e.g., combinations of path and loop structures. Unlike the Complex model, they replaced the dot product with the Möbius function which has several favorable theoretical properties. This model subsumes Complex in that it embeds entities in much lower ranks, i.e., about 25% or even smaller to achieve the state-of-the-art performance. However, 5★E is inferior to Complex in that it needs almost the same large size of relation parameters to do much more sophisticated calculation.

Following the hyperparameter search range of Nayyeri et al. (2021), the embedding sizes we tested for 5★E to obtain the best result are $rank = 500$ for all datasets. Each entity needs $2 \times rank$ parameters, and each relation needs $8 \times rank$ parameters that function differently.

In this model, relations are represented as $\begin{bmatrix} a & b \\ c & d \end{bmatrix}$. The transformation function ϑ of 5★E is

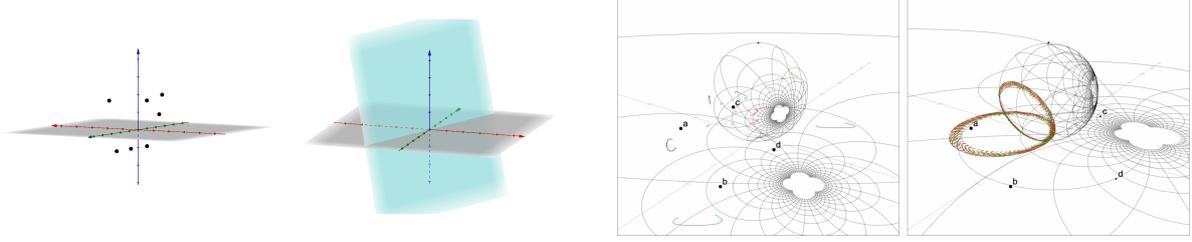
$$x \rightarrow \begin{bmatrix} x \\ 1 \end{bmatrix} \rightarrow \begin{bmatrix} a & b \\ c & d \end{bmatrix} \begin{bmatrix} x \\ 1 \end{bmatrix} \rightarrow \frac{ax + b}{cx + d}. \quad (2)$$

Möbius function ϑ is capable of representing various relations simultaneously because it combines five subsequent transformations: $\vartheta = \vartheta_4 \circ \vartheta_3 \circ \vartheta_2 \circ \vartheta_1$, where $\vartheta_1 = x + \frac{d}{c}$ is describing translation by $\frac{d}{c}$, $\vartheta_2 = \frac{1}{x}$ is describing inversion and reflection w.r.t. real axis, $\vartheta_3 = \frac{bc-ad}{c^2}x$ is describing homothety and rotation, and $\vartheta_4 = x + \frac{a}{c}$ is describing translation by $\frac{a}{c}$.

3.2 Method Formulation

Let $[a_1 \ a_2]$ denotes the relation embedding matrix. Our method constrains half of the parameters a_2 using the complex conjugate of the other half \bar{a}_1 , i.e., $a_2 = \bar{a}_1$; it is model-dependent to specify which parameters are suitable for conjugation. We formulated our method on above two baseline models.

Complex By using our method, the original model Complex is adapted to the parameter-sharing model Complex, where relations are represented as the real part of low-rank matrices with conjugate parameters. Specifically, we set the original square relation embedding matrices $[a_1 \ a_2]$ to be half the normal parameters and the other half their conjugation, i.e., $[a_1 \ \bar{a}_1]$. In this model, since each parameter is functioning equally, the positions of the conjugate parameters can be set randomly.



(a) Transformed entities by ComplEx (left) and Complex (right). In the left graph, a black point describes a transformed entity, and the vector values of a point are unrelated in each dimension. While in the right graph, half of the value $z_i, i \in [1, d/2]$ of a vector z_1, z_2, \dots, z_d that is describing a point are constrained as the other half $z_i, i \in [d/2 + 1, d]$ correspondingly. The linear constrain $z_i = a_i x_i + b_i y_i$ is illustrated in the right graph.

(b) Transformed entities by $5^\star E$ (left) and $5^\star \epsilon_n$ (right). Note that we illustrate the negative conjugated model instead of the positive conjugated one for simplicity in plotting. Blue traces are the original entities and their projections in the Non-Euclidean space. Green traces are the multiple copies of the blue traces under iterations of the Möbius transformation. Red traces are the inverse of green traces. Apparently, the right graph has much neater geometric properties.

Figure 1: Transformed entities illustrated in 3D

$5^\star \epsilon$ Our method transforms the original model $5^\star E$ into the parameter-sharing model $5^\star \epsilon$, where relations are represented as the real part of low-rank matrices using conjugate parameters. Specifically, we set the original square relation embedding matrices $\begin{bmatrix} a & b \\ c & d \end{bmatrix}$ to be half the normal parameters and the other half their conjugate parameters, i.e., $\begin{bmatrix} a & b \\ \bar{b} & \bar{a} \end{bmatrix}$. In this model, parameters play distinct roles at different positions, and the best conjugation positions are the principal and secondary diagonal positions. Note that experiments showed that, the following negative conjugation method, i.e., $\begin{bmatrix} a & b \\ -\bar{b} & \bar{a} \end{bmatrix}$, achieves similar performance as above. Although the negative conjugation on this model is equivalent to restricting the original Möbius function to the unitary Möbius transformation, our approach is much more general to a variety of representations.

3.3 Transformation Analysis

Complex Let $a_2 = \bar{a}_1$, then the transformation of conjugate model Complex is

$$[x_1 \ x_2] \rightarrow [a_1 \ \bar{a}_1] [x_1 \ x_2] \rightarrow [a_1 x_1 \ \bar{a}_1 x_2]. \quad (3)$$

We can see that the resulted relation embedding is constrained to $[a_1 \ \bar{a}_1]$ other than $[a_1 \ a_2]$; the predicted tail entity is constrained to $[a_1 x_1 \ \bar{a}_1 x_2]$ instead of $[a_1 x_1 \ a_2 x_2]$ in original model, which does not narrow the rang of relation or tail embedding since the a_1, x_2 can be any value. Further, since tail entities also act as head entities, we can

say that the range of both the entities and relations are not constrained.

$5^\star \epsilon$ Let $c = \bar{b}, d = \bar{a}$, then the transformation of conjugate model $5^\star \epsilon$ is

$$x \rightarrow \begin{bmatrix} x \\ 1 \end{bmatrix} \rightarrow \begin{bmatrix} a & b \\ \bar{b} & \bar{a} \end{bmatrix} \begin{bmatrix} x \\ 1 \end{bmatrix} \rightarrow \frac{ax + b}{\bar{b}x + \bar{a}}. \quad (4)$$

The five subsequent transformations turn into: $\vartheta_1 = x + \frac{a}{b}$ which depicts translation by $\frac{a}{b}$, $\vartheta_2 = \frac{1}{x}$ which depicts inversion and reflection w.r.t. real axis, $\vartheta_3 = \frac{bb - a\bar{a}}{\bar{b}^2} x$ which depicts homothety and rotation, and $\vartheta_4 = x + \frac{a}{b}$ which depicts translation by $\frac{a}{b}$. We can see that, although the relation parameters are constrained comparing to the original model, the five sub-transformations are reserved in this conjugate model.

Characteristics For this reason, we consider our conjugate models retain expressiveness in function level for various relation patterns compared to their original counterparts. The difference between original models and our conjugate models is that, the latter ones have more linear constrain in its value of each embedding parameter, as illustrated in Figure 1.

3.4 Reduced Calculation

Sharing half of the parameters also reduces the computation for the regularization terms into half, where each parameter of relation is squared to the sum. For example, the original calculation $r_1^2 + r_2^2$ is turned into $r_1^2 \times 2$ in both baseline models, where r_1, r_2 denote the real or imaginary part of a complex number, and in which r_1 represents the shared

Dataset	#Training	#Validation	#Test	Ent	Rel	Exa
FB15K-237	272,115	17,535	20,466	14,541	237	544,230
WN18RR	86,835	3,034	3,134	40,943	11	173,670
YAGO3-10	1,079,040	5,000	5,000	123,188	37	2,158,080
FB15K	483,142	50,000	59,071	14,951	1,345	966,284
WN18	141,442	5,000	5,000	40,943	18	282,884

Table 1: Datasets statistics. #: Split in terms of number of triples; Ent: Entities; Rel: Relations; Exa: Examples.

parameter. However, the final time consumption depends on multiple aspects, such as formulation and coding, thus is not necessarily reduced.

4 Experiments

4.1 Experimental Setup

Metrics We followed the standard evaluation protocol for KGE models. T : the rank set of truth, r_i : the rank position r of the first true entity for the i -th query. We computed two rank-based metrics: (i) Mean Reciprocal Rank (MRR), which computes the arithmetic mean of reciprocal ranks of all true entities from the ranked list of answers to queries T , and (ii) Hits@ N ($N = 1, 3, 10$), which counts the true entities \mathbb{I} and calculate their proportion in the truth T in top N sorted predicted answers list.

$$\text{MRR} = \frac{1}{T} \sum_{i=1}^T \frac{1}{r_i} \quad (5)$$

$$\text{Hits@}N = \frac{1}{T} \sum_{r \in T, r \leq N} \mathbb{I} \quad (6)$$

We also use additional metric Time (seconds/epoch) to measure how many seconds each training epoch costs to demonstrate the time saved by our method. To do this, we conducted all experiments using the same GPUs. GeForce GTX 1080 Ti is used for all datasets except for the largest dataset YAGO3-10 who needs a larger GPU and we used Tesla V100S-PCIE-32GB for it.

Datasets We evaluated our method on five widely used benchmark datasets (See Table 1). FB15K (Bordes et al., 2013) is a subset of Freebase, the contents of which are general facts. WN18 (Bordes et al., 2013) is a subset of Wordnet, a database that features lexical relations between words. YAGO3-10 (Dettmers et al., 2018) is the largest common dataset, which mostly describes attributes of persons, and contains entities associated with at least ten different relations.

As was first noted by Toutanova and Chen (2015), FB15K and WN18 suffer from test leakage through inverse relations, e.g., the test set frequently contains triples such as $(s, \textit{hyponym}, o)$ while the training set contains its inverse $(o, \textit{hypernym}, s)$. To create a dataset without this property, they introduced FB15K-237, a subset of FB15K where inverse relations are removed. WN18RR was created for the same reason by Dettmers et al. (2018).

We adopted all of the five datasets for comprehensive comparison of models.

Hyperparameter Settings We explored the influence of hyperparameter settings to our method. To do this, we used the best hyperparameter settings for the original models (marked as ∇ or no mark), and applied the same settings on our conjugate models and ablation models. We adopted the best hyperparameter settings for ComplEx provided by Nayyeri et al. (2021), and fine-tuned the best hyperparameters ourselves for $5^\star E$ since there was no published best hyperparameter settings for this model at the time we did the experiments. We also fine-tuned the best hyperparameters for one of our conjugate model $5^\star \epsilon$ (noted as \diamond) to explore the upper bound.

We selected the hyperparameters based on the MRR on the validation set. Our grid search range referred to but was larger than Nayyeri et al. (2021). The optional optimizers are {Adagrad, Adam, SGD}. The range of embedding dimensions are {100, 500} with learning rates range in {1E-02, 5E-02, 1E-01}. The batch sizes attempted range in {100, 500, 1000, 2000}. Regularization coefficients are tested among {2.5E-03, 5E-03, 1E-02, 5E-02, 1E-01, 5E-01}.

5 Results

5.1 Main Results and Analysis

The main experimental results are shown in Table 2 and Table 3. The numbers with boldface indicate

Model	Time	MRR	H@1	H@3	H@10
ComplEx	42±8	0.366±4e-4	0.271	0.402	0.558
Complex	46±11	0.363±5e-4	0.268	0.400	0.555
5★E	18±3	0.350±8e-4	0.257	0.386	0.538
5★ ϵ_{∇}	14±4	0.353±7e-4	0.259	0.390	0.541
5★ ϵ_{\diamond}	17±9	0.354±8e-4	0.259	0.391	0.544

(a) FB15K-237

Model	Time	MRR	H@1	H@3	H@10
ComplEx	139±21	0.488±1e-3	0.442	0.503	0.579
Complex	146±45	0.475±9e-4	0.433	0.488	0.558
5★E	16±1	0.490±5e-4	0.444	0.506	0.587
5★ ϵ_{∇}	11±1	0.493±8e-4	0.442	0.512	0.588
5★ ϵ_{\diamond}	-	-	-	-	-

(b) WN18RR

Model	Time	MRR	H@1	H@3	H@10
ComplEx	370±2	0.577±1e-3	0.502	0.622	0.712
Complex	371±2	0.574±2e-3	0.500	0.618	0.707
5★E	415±2	0.574±2e-3	0.502	0.617	0.701
5★ ϵ_{∇}	297±1	0.576±2e-3	0.505	0.619	0.702
5★ ϵ_{\diamond}	-	-	-	-	-

(c) YAGO3-10

Table 2: Link prediction results on FB15K-237, WN18RR, YAGO3-10 datasets. Time, MRR and H@n are presented as mean (\pm standard deviation).

the best results among all the models.

We mainly tested whether the conjugate models perform consistent with their original counterparts, especially whether they can achieve the same state-of-the-art results. We conducted one set of experiments using the best hyperparameters of the original models (marked as ∇ or no mark), and the other set of experiments tuning the hyperparameters for one of our conjugate model (marked as \diamond).

The results show that both Complex and 5★ ϵ consistently achieve results comparable to their original models on the datasets without test set leakage, including the largest dataset, i.e., YAGO3-10; and obtain the same optimal accuracies as the original models on all datasets with possibly-required fine-tuning. From the perspective of training time, we see 5★ ϵ spends 31% less time on average for all datasets; and both conjugate models perform substantially best in training time on datasets FB15K, who have the most relations.

Complex Under the best hyperparameter settings of the original model, the performance of Complex are consistently comparable with ComplEx on all five datasets. We speculate the reason for the consistent but tiny performance drop might come from the computation precision, but we will leave it as our future studies.

Model	Time	MRR	H@1	H@3	H@10
ComplEx	346±124	0.855±1e-3	0.823	0.874	0.910
Complex	293±16	0.855±1e-3	0.827	0.871	0.907
5★E	42±9	0.812±1e-3	0.767	0.840	0.889
5★ ϵ_{∇}	26±0	0.794±2e-3	0.743	0.827	0.882
5★ ϵ_{\diamond}	29±5	0.813±2e-3	0.766	0.844	0.894

(a) FB15K

Model	Time	MRR	H@1	H@3	H@10
ComplEx	57±3	0.951±3e-4	0.944	0.954	0.961
Complex	58±5	0.950±3e-4	0.945	0.953	0.960
5★E	43±6	0.952±5e-4	0.946	0.955	0.962
5★ ϵ_{∇}	29±6	0.949±6e-4	0.944	0.953	0.959
5★ ϵ_{\diamond}	26±2	0.952±3e-4	0.947	0.955	0.962

(b) WN18

Table 3: Link prediction results on FB15K and WN18 datasets. Instructions for this table are the same as those in Table 2.

Although our method reduces the computation, applying the method on this model requires splitting and concatenating matrices to keep the shape of outputs which incurs additional time-consuming operations. Consequently, the total time cost is not reduced much. However, training time on dataset FB15K, who has the most relations, becomes very stable.

Overall results imply our conjugate model Complex is at least comparative with its baseline model ComplEx.

5★ ϵ Under the best hyperparameter settings of the original 5★E, the conjugate 5★ ϵ_{∇} consistently achieve competitive results on the datasets FB15K-237, WN18RR and YAGO3-10. The tiny but consistent accuracy enhancement on these three datasets is probably caused by similar programming artifacts as observed in ComplEx.

We hypothesize that the accuracy fluctuation of 5★ ϵ_{∇} on FB15K and WN18 is caused by the test leakage issue which makes the model sensitive to its hyperparameter setting. Because the only difference of these two datasets comparing to their subsets FB15K-237 and WN18RR is the 81% and 94% inverse relations (Toutanova and Chen, 2015), i.e., (*s*, *hyponym*, *o*) and (*o*, *hypernym*, *s*) in the training set and the test set respectively, which is known as test leakage. Note that the accuracy fluctuation was simply solved by fine-tuning the hyperparameters (See results marked as 5★ ϵ_{\diamond}).

Notice that in Table 2, we didn't report the fine-tuned results of 5★ ϵ_{\diamond} on datasets WN18RR and YAGO3-10, because the results obtained with the original settings ∇ is already the best.

Model	Time	MRR	H@1	H@3	H@10
$5^\star\epsilon_\nabla$	14±4	0.353±7e-4	0.259	0.390	0.541
$5^\star\epsilon_n$	13±2	0.353±8e-4	0.259	0.389	0.541
$5^\star E_r$	16±0	0.326±1e-3	0.238	0.357	0.505
$5^\star\epsilon_v$	13±1	0.264±4e-4	0.192	0.288	0.404
$5^\star\epsilon_h$	12±0	0.301±4e-4	0.221	0.329	0.458

(a) FB15K-237

Model	Time	MRR	H@1	H@3	H@10
$5^\star\epsilon_\nabla$	11±1	0.493±8e-4	0.442	0.512	0.588
$5^\star\epsilon_n$	14±3	0.485±1e-3	0.432	0.506	0.589
$5^\star E_r$	16±0	0.410±3e-3	0.391	0.417	0.447
$5^\star\epsilon_v$	12±5	0.026±2e-4	0.015	0.025	0.045
$5^\star\epsilon_h$	14±3	0.026±3e-4	0.016	0.025	0.046

(b) WN18RR

Model	Time	MRR	H@1	H@3	H@10
$5^\star\epsilon_\nabla$	297±1	0.576±2e-3	0.505	0.619	0.702
$5^\star\epsilon_n$	298±1	0.574±1e-3	0.502	0.618	0.701
$5^\star E_r$	416±2	0.569±2e-3	0.499	0.611	0.695
$5^\star\epsilon_v$	297±1	0.562±8e-4	0.488	0.607	0.695
$5^\star\epsilon_h$	298±1	0.546±1e-3	0.471	0.592	0.680

(c) YAGO3-10

Table 4: Ablation studies on FB15K-237, WN18RR, YAGO3-10 datasets. Instructions for this table are the same as those in Table 2.

Training time in this model was reduced by 22%, 31%, 28%, 38% and 33% on each dataset respectively, and 31% on average. $5^\star E$ has eight parameter matrices in the coding. By using our method, the parameter matrices are directly reduced to four with no additional coding operations, which makes the significant saved training time.

Above results mean our conjugate model $5^\star\epsilon$ exceeds the baseline model $5^\star E$ in all respect of accuracy, memory-efficiency and time footprint.

5.2 Ablation Studies

We did two kinds of ablation studies. The results are shown in Table 4 and Table 5. We know the reduced calculation is mainly in the regularization process because we only use half of the parameters. Thus we experimented where the regularization term is only half of the parameters on the original model (See results for $5^\star E_r$) to explore whether the effect of our method is similar to the reduced parameters regularization.

Then we experimented with conjugations in different positions to explore how the models perform differently. We set negative conjugation $c = -\bar{b}, d = \bar{a}$ in model $5^\star\epsilon_n$, where half of the conjugate parameters are using negative conjugation instead of positive conjugation; we set vertical

Model	Time	MRR	H@1	H@3	H@10
$5^\star\epsilon_\nabla$	26±0	0.794±2e-3	0.743	0.827	0.882
$5^\star\epsilon_n$	31±12	0.799±2e-3	0.750	0.831	0.883
$5^\star E_r$	37±1	0.807±3e-3	0.760	0.838	0.888
$5^\star\epsilon_v$	31±7	0.801±8e-4	0.753	0.833	0.885
$5^\star\epsilon_h$	28±2	0.787±2e-3	0.735	0.822	0.877

(a) FB15K

Model	Time	MRR	H@1	H@3	H@10
$5^\star\epsilon_\nabla$	29±6	0.949±6e-4	0.944	0.953	0.959
$5^\star\epsilon_n$	26±0	0.952±3e-4	0.946	0.955	0.962
$5^\star E_r$	40±0	0.943±9e-4	0.935	0.950	0.954
$5^\star\epsilon_v$	31±11	0.892±2e-3	0.836	0.944	0.958
$5^\star\epsilon_h$	26±0	0.822±2e-3	0.719	0.920	0.949

(b) WN18

Table 5: Ablation studies on FB15K and WN18 datasets. Instructions for this table are the same as those in Table 2.

conjugation $c = \bar{a}, d = \bar{b}$ in model $5^\star\epsilon_v$, where parameters are conjugated in their vertical direction instead of the diagonal direction; and we let $b = \bar{a}, d = \bar{c}$ in model $5^\star\epsilon_h$, the horizontal conjugation, where parameters are conjugated in their horizontal direction.

The studies show that, first, by comparing the accuracy of $5^\star\epsilon$ and $5^\star E_r$, we know that reducing parameters in the regularization process hurts the accuracy significantly, which indicates our conjugation method indeedly reserves model’s ability even when the parameters are reduced. Second, the negative conjugate model $5^\star\epsilon_n$ performs as well as $5^\star\epsilon$. Last but not least, conjugate method should choose suitable positions, e.g., $5^\star\epsilon_v$ and $5^\star\epsilon_h$ do not perform as well.

5.3 Statistical Methods

To clarify the difference between original models and their conjugate models, we took the highest mean as the best result, with the standard deviation as a secondary judgement, and ultimately two-sample t-tests (See Table 6 in Appendix) are conducted to decide whether two similar results can be considered statistically equivalent and which is the best.

The two-sample t-test estimates if two population means are equal. Here we use the t-test to judge if the Time or MRR means of two models are equal. We set significance level $\alpha = 0.05$, and the null hypothesis assumed that the two data samples are from normal distributions with unknown and unequal variances. (h, p) means the result h and p -value of the hypothesis test. $h = 1, 0$. $h = 1$

means rejection to the null hypothesis at the significance level α . $h = 0$ indicates the failure to reject the null hypothesis at the significance level α . $p \in [0, 1]$ is a probability of observing a test statistic as extreme as, or more extreme than, the observed value under the null hypothesis. A small p value suggests suspicion on the validity of the null hypothesis.

To prepare data for the t-tests, experiments on ComplEx, Complex, 5★E, 5★ ϵ_{∇} , 5★ ϵ_{\diamond} and 5★ ϵ_n are conducted 17 times each. Apart from that, the 5★E_r, 5★ ϵ_v , 5★ ϵ_h apparently perform worse than the former six models, thus the t-tests are not needed and their experiments are conducted 5 times each.

Most of our t-tests were done among the original model and its conjugate models as the distribution differs significantly if the base model is different. However, since the accuracies among different models are similar on the YAGO3-10 and WN18 datasets, we did several supplementary t-tests (indicated in *italics*). The supplementary t-tests showed that the distributions are different indeed when based on different original models even though they appear to be similar. On the contrary, there exist similar distributions among the results distribution of the original model and its conjugate model.

5.4 Advantages of Parameter Sharing

Approaching for the best accuracy in link prediction task has the trade off of misinformation effect or inevitable high memory and time costs. Our parameter-sharing method by using half conjugate parameters is very easy to apply and can help control these costs, and potentially no trade off.

The original ComplEx and 5★E each has their own strength in the perspective of accuracy on different datasets; while ComplEx costs much more memory and time than 5★E when compared under similar accuracy.

Our conjugate models consume less memory and time, and not inferior to the original models in accuracy, which shows that our parameter-sharing method makes a complex number represented KGE model superior to itself.

6 Conclusions

We propose using shared conjugate parameters for transformations, which suffices to accurately represent the structures of the KG.

Our method can help scaling up KG with less carbon footprints easily: first, it reduces parameter size and consumes less or at least comparable training time while achieving consistent accuracy as the non-conjugate model, including reaching state-of-the-art results; second, it is easily generalizable across various complex number represented models.

7 Future Work

We would like to deal with the interpretation of the linear constrain of our method. For example, to explore the effect of this method on different relation patterns. Moreover, many KG applications like the work done by [Hongwimol et al. \(2021\)](#) regard visual appeal as important, where appropriate visuals can better convey the points of the data and facilitate user interaction. We can see that the vector representations of transformed entities using this method have more substantial geometric constrains (See transformed entities illustrated in Figure 1). We want to explore if our method can obtain better KG visualization.

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A Two-sample t-test

	Time t-test (h, p)					MRR t-test (h, p)						
	ComplEx	Complex	$5^*\epsilon$	$5^*\epsilon_{\nabla}$	$5^*\epsilon_{\diamond}$	$5^*\epsilon_n$	ComplEx	Complex	$5^*\epsilon$	$5^*\epsilon_{\nabla}$	$5^*\epsilon_{\diamond}$	$5^*\epsilon_n$
ComplEx	-	(0, 2e-1)	-	-	-	-	-	(1, 8e-17)	-	-	-	-
Complex	-	-	-	-	-	-	-	-	-	-	-	-
$5^*\epsilon$	-	-	-	(1, 8e-3)	(0, 6e-1)	(1, 4e-5)	-	-	(1, 1e-12)	(1, 1e-14)	(1, 1e-10)	-
$5^*\epsilon_{\nabla}$	-	-	-	-	(0, 4e-1)	(0, 4e-1)	-	-	-	(1, 6e-3)	(1, 1e-2)	-
$5^*\epsilon_{\diamond}$	-	-	-	-	-	(0, 2e-1)	-	-	-	-	(1, 7e-6)	-
$5^*\epsilon_n$	-	-	-	-	-	-	-	-	-	-	-	-

(a) FB15K-237

	Time t-test (h, p)					MRR t-test (h, p)						
	ComplEx	Complex	$5^*\epsilon$	$5^*\epsilon_{\nabla}$	$5^*\epsilon_{\diamond}$	$5^*\epsilon_n$	ComplEx	Complex	$5^*\epsilon$	$5^*\epsilon_{\nabla}$	$5^*\epsilon_{\diamond}$	$5^*\epsilon_n$
ComplEx	-	(0, 6e-1)	-	-	-	-	-	(1, 3e-28)	-	-	-	-
Complex	-	-	-	-	-	-	-	-	-	-	-	-
$5^*\epsilon$	-	-	-	(1, 4e-11)	-	(1, 6e-3)	-	-	(1, 9e-11)	-	(1, 2e-15)	-
$5^*\epsilon_{\nabla}$	-	-	-	-	-	(1, 2e-2)	-	-	-	-	(1, 4e-21)	-
$5^*\epsilon_{\diamond}$	-	-	-	-	-	-	-	-	-	-	-	-
$5^*\epsilon_n$	-	-	-	-	-	-	-	-	-	-	-	-

(b) WN18RR

	Time t-test (h, p)					MRR t-test (h, p)						
	ComplEx	Complex	$5^*\epsilon$	$5^*\epsilon_{\nabla}$	$5^*\epsilon_{\diamond}$	$5^*\epsilon_n$	ComplEx	Complex	$5^*\epsilon$	$5^*\epsilon_{\nabla}$	$5^*\epsilon_{\diamond}$	$5^*\epsilon_n$
ComplEx	-	(0, 4e-1)	-	-	-	-	-	(1, 3e-6)	(1, 2e-7)	(1, 7e-3)	-	(1, 3e-8)
Complex	-	-	-	-	-	-	-	-	-	-	-	-
$5^*\epsilon$	-	-	-	(1, 5e-47)	-	(1, 1e-46)	-	-	(1, 5e-4)	-	(0, 6e-1)	-
$5^*\epsilon_{\nabla}$	-	-	-	-	-	(0, 7e-1)	-	-	-	-	(1, 3e-4)	-
$5^*\epsilon_{\diamond}$	-	-	-	-	-	-	-	-	-	-	-	-
$5^*\epsilon_n$	-	-	-	-	-	-	-	-	-	-	-	-

(c) YAGO3-10

	Time t-test (h, p)					MRR t-test (h, p)						
	ComplEx	Complex	$5^*\epsilon$	$5^*\epsilon_{\nabla}$	$5^*\epsilon_{\diamond}$	$5^*\epsilon_n$	ComplEx	Complex	$5^*\epsilon$	$5^*\epsilon_{\nabla}$	$5^*\epsilon_{\diamond}$	$5^*\epsilon_n$
ComplEx	-	(0, 1e-1)	-	-	-	-	-	(0, 8e-2)	-	-	-	-
Complex	-	-	-	-	-	-	-	-	-	-	-	-
$5^*\epsilon$	-	-	-	(1, 2e-6)	(1, 3e-5)	(1, 8e-3)	-	-	(1, 2e-21)	(1, 2e-2)	(1, 6e-21)	-
$5^*\epsilon_{\nabla}$	-	-	-	-	(1, 3e-2)	(0, 1e-1)	-	-	-	(1, 1e-23)	(1, 3e-9)	-
$5^*\epsilon_{\diamond}$	-	-	-	-	-	(0, 5e-1)	-	-	-	-	(1, 3e-22)	-
$5^*\epsilon_n$	-	-	-	-	-	-	-	-	-	-	-	-

(d) FB15K

	Time t-test (h, p)					MRR t-test (h, p)						
	ComplEx	Complex	$5^*\epsilon$	$5^*\epsilon_{\nabla}$	$5^*\epsilon_{\diamond}$	$5^*\epsilon_n$	ComplEx	Complex	$5^*\epsilon$	$5^*\epsilon_{\nabla}$	$5^*\epsilon_{\diamond}$	$5^*\epsilon_n$
ComplEx	-	(0, 5e-1)	-	-	-	-	-	(0, 8e-2)	-	-	(1, 1e-17)	(1, 1e-13)
Complex	-	-	-	-	-	-	-	-	-	(1, 5e-21)	(1, 2e-16)	-
$5^*\epsilon$	-	-	-	(1, 1e-7)	(1, 4e-9)	(1, 6e-9)	-	-	(1, 3e-14)	(1, 5e-5)	(0, 2e-1)	-
$5^*\epsilon_{\nabla}$	-	-	-	-	(0, 2e-1)	(0, 8e-2)	-	-	-	(1, 4e-15)	(1, 5e-14)	-
$5^*\epsilon_{\diamond}$	-	-	-	-	-	(0, 1e-1)	-	-	-	-	(1, 3e-5)	-
$5^*\epsilon_n$	-	-	-	-	-	-	-	-	-	-	-	-

(e) WN18

Table 6: t-test (h, p) of Time and MRR on FB15K-237, WN18RR, YAGO3-10, FB15K and WN18 datasets.