# An overview of embedding models of entities and relationships for knowledge base completion

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#### **Abstract**

Knowledge bases (KBs) of real-world facts about entities and their relationships are useful resources for a variety of natural language processing tasks. However, because knowledge bases are typically incomplete, it is useful to be able to perform *knowledge base completion* or *link prediction*, i.e., predict whether a relationship not in the knowledge base is likely to be true. This paper serves as a comprehensive overview of embedding models of entities and relationships for knowledge base completion, summarizing up-to-date experimental results on standard benchmark datasets.

**Keywords:** Knowledge base completion, link prediction, embedding model, entity prediction.

#### 1 Introduction

Before introducing the KB completion task in detail, let us return to the classic Word2Vec example of a "royal" relationship between "king" and "man", and between "queen" and "woman." As illustrated in this example:  $v_{king} - v_{man} \approx v_{queen} - v_{woman}$ , word vectors learned from a large corpus can model relational similarities or linguistic regularities between pairs of words as translations in the projected vector space (Mikolov et al., 2013; Pennington et al., 2014). Figure 1 shows another example of a relational similarity between word pairs of countries and capital cities:

$$egin{aligned} oldsymbol{v}_{Japan} - oldsymbol{v}_{Tokyo} &pprox & oldsymbol{v}_{Germany} - oldsymbol{v}_{Berlin} \ pprox & oldsymbol{v}_{Portugal} - oldsymbol{v}_{Lisbon} \end{aligned}$$

Let us consider the country and capital pairs in Figure 1 to be pairs of entities rather than word types. That is, we now represent country and capital entities by low-dimensional and dense vectors. The relational similarity between word pairs is presumably to capture a "is\_capital\_of" relationship between country and capital entities. Also, we represent this relationship by a translation vector  $v_{is\_capital\_of}$  in the entity vector space. Thus, we expect:

$$egin{array}{ll} oldsymbol{v}_{Tokyo} + oldsymbol{v}_{is\_capital\_of} - oldsymbol{v}_{Japan} &pprox & oldsymbol{0} \ oldsymbol{v}_{Berlin} + oldsymbol{v}_{is\_capital\_of} - oldsymbol{v}_{Germany} &pprox & oldsymbol{0} \ oldsymbol{v}_{Lisbon} + oldsymbol{v}_{is\_capital\_of} - oldsymbol{v}_{Portugal} &pprox & oldsymbol{0} \end{array}$$

This intuition inspired the TransE model—a well-known embedding model for KB completion or link prediction in KBs (Bordes et al., 2013).

Knowledge bases are collections of real-world triples, where each triple or fact (h,r,t) in KBs represents some relation r between a head entity h and a tail entity t. KBs can thus be formalized as directed multi-relational graphs, where nodes correspond to entities and edges linking the nodes encode various kinds of relationships (García-Durán et al., 2016; Nickel et al., 2016a). Here entities are real-world things or objects such as persons, places, organizations, music tracks or movies. Each relation type defines a certain relationship between entities. For example, as illustrated in Figure 2, the relation type "child\_of" relates person entities with each other, while the relation type "born\_in" relates person entities

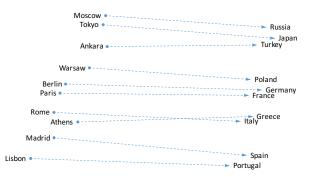


Figure 1: Two-dimensional projection of vectors of countries and their capitals. This figure is drawn based on Mikolov et al. (2013).

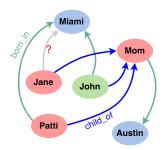


Figure 2: An illustration of (incomplete) knowledge base, with 4 person entities, 2 place entities, 2 relation types and total 6 triple facts. This figure is drawn based on Weston and Bordes (2014).

with place entities. Several KB examples include the domain-specific KB GeneOntology and popular generic KBs of WordNet (Fellbaum, 1998), YAGO (Suchanek et al., 2007), Freebase (Bollacker et al., 2008), NELL (Carlson et al., 2010) and DBpedia (Lehmann et al., 2015) as well as commercial KBs such as Google's Knowledge Graph, Microsoft's Satori and Facebook's Open Graph. Nowadays, KBs are used in a number of commercial applications including search engines such as Google, Microsoft's Bing and Facebook's Graph search. They also are useful resources for many natural language processing tasks such as question answering (Ferrucci, 2012; Fader et al., 2014), word sense disambiguation (Navigli and Velardi, 2005; Agirre et al., 2013), semantic parsing (Krishnamurthy and Mitchell, 2012; Berant et al., 2013) and co-reference resolution (Ponzetto and Strube, 2006; Dutta and Weikum, 2015).

A main issue is that even very large KBs, such as Freebase and DBpedia, which contain billions of fact triples about the world, are still far from complete. In particular, in English DBpedia 2014, 60% of person entities miss a place of birth and 58% of the scientists do not have a fact about what they are known for (Krompaß et al., 2015). In Freebase, 71% of 3 million person entities miss a place of birth, 75% do not have a nationality while 94% have no facts about their parents (West et al., 2014). So, in terms of a specific application, question answering systems based on incomplete KBs would not provide a correct answer given a correctly interpreted question. For example, given the incomplete KB in Figure 2, it would be impossible to answer the question "where was Jane born?", although the question is completely matched with existing entity and relation type information (i.e., "Jane" and "born\_in") in KB. Consequently, much work has been devoted towards knowledge base completion to perform link prediction in KBs, which attempts to predict whether a relationship/triple not in the KB is likely to be true, i.e., to add new triples by leveraging existing triples in the KB (Lao and Cohen, 2010; Bordes et al., 2012; Gardner et al., 2014; García-Durán et al., 2016). For example, we would like to predict the missing tail entity in the incomplete triple (Jane, born\_in,?) or predict whether the triple (Jane, born\_in, Miami) is correct or not.

Embedding models for KB completion have been proven to give state-of-the-art link prediction performances, in which entities are represented by latent feature vectors while relation types are represented by latent feature vectors and/or matrices and/or third-order tensors (Bordes et al., 2013; Socher et al., 2013). This paper overviews the embedding models for KB completion, and then summarizes up-to-date experimental results on the standard evaluation task of entity prediction—which is also referred to as the link prediction task (Bordes et al., 2013).

### 2 A general approach of embedding models for KB completion

Let  $\mathcal{E}$  denote the set of entities and  $\mathcal{R}$  the set of relation types. Denote by  $\mathcal{G}$  the knowledge base consisting of a set of correct triples (h, r, t), such that  $h, t \in \mathcal{E}$  and  $r \in \mathcal{R}$ . For each triple (h, r, t), the embedding models define a *score function* f(h, r, t) of its plausibility. Their goal here is to:

Choose f such that the score f(h, r, t) of a correct triple (h, r, t) is higher than the score f(h', r', t') of an incorrect triple (h', r', t').

For example, TransE defines a score function of  $f_{\text{TransE}}(h,r,t) = -\|\boldsymbol{v}_h + \boldsymbol{v}_r - \boldsymbol{v}_t\|$ , where h,r and t are represented by low dimensional vectors  $\boldsymbol{v}_h, \boldsymbol{v}_r$  and  $\boldsymbol{v}_t$ , respectively. As (Tokyo, is\_capital\_of, Japan) is a correct triple, while (Tokyo, is\_capital\_of, Portugal) and (Lisbon, is\_capital\_of, Japan) are incorrect ones, we would have:  $-\|\boldsymbol{v}_{Tokyo} + \boldsymbol{v}_{is\_capital\_of} - \boldsymbol{v}_{Japan}\| > -\|\boldsymbol{v}_{Tokyo} + \boldsymbol{v}_{is\_capital\_of} - \boldsymbol{v}_{Portugal}\|$ , and  $-\|\boldsymbol{v}_{Tokyo} + \boldsymbol{v}_{is\_capital\_of} - \boldsymbol{v}_{Japan}\| > -\|\boldsymbol{v}_{Lisbon} + \boldsymbol{v}_{is\_capital\_of} - \boldsymbol{v}_{Japan}\|$ . Table 1 in Section 3 summarizes different score functions f(h,r,t).

To learn model parameters (i.e., entity vectors, relation vectors or matrices), the embedding models minimize an objective loss  $\mathcal{L}$ . A conventional objective loss is the margin-based pairwise ranking loss (Bordes et al., 2013):

$$\mathcal{L}_{\text{Margin}} = \sum_{\substack{(h,r,t) \in \mathcal{G} \\ (h',r,t') \in \mathcal{G}'_{(h,r,t)}}} [\gamma - f(h,r,t) + f(h',r,t')]_{+}$$

where  $[x]_+ = \max(0, x)$ ;  $\gamma$  is the margin hyper-parameter; and  $\mathcal{G}'_{(h,r,t)}$  is the set of incorrect triples generated by corrupting the correct triple  $(h, r, t) \in \mathcal{G}$ .

Furthermore, the negative log-likelihood (NLL) of softmax regression (SoftReg) (Toutanova and Chen, 2015) and the NLL of logistic regression (LogReg) (Trouillon et al., 2016) are commonly used in recent work:

$$\mathcal{L}_{\text{SoftReg}} \quad = \quad -\sum_{(h,r,t) \in \mathcal{G}} \left( \frac{\exp\left(f\left(h,r,t\right)\right)}{\sum\limits_{t' \in \mathcal{E} \setminus \{t\}} \exp\left(f\left(h,r,t'\right)\right)} \right. \\ \left. + \frac{\exp\left(f\left(h,r,t\right)\right)}{\sum\limits_{h' \in \mathcal{E} \setminus \{h\}} \exp\left(f\left(h',r,t\right)\right)} \right) \\ \mathcal{L}_{\text{LogReg}} \quad = \quad \sum_{(h,r,t) \in \{\mathcal{G} \cup \mathcal{G}'\}} \log\left(1 + \exp\left(-\mathrm{I}_{(h,r,t)} \cdot f\left(h,r,t\right)\right)\right) \\ \text{with: } \mathrm{I}_{(h,r,t)} = \left\{ \begin{array}{l} 1 \text{ for } (h,r,t) \in \mathcal{G} \\ -1 \text{ for } (h,r,t) \in \mathcal{G}' \end{array} \right.$$

To corrupt the head or tail entities, a common strategy is to uniformly replace the entities when sampling incorrect triples (Bordes et al., 2013), however it results in many false negative labels (Wang et al., 2014). Domain sampling (Krompaß et al., 2015; Xie et al., 2017) generates corrupted triples by sampling entities from the same domain or from the set of relation-dependent entities. The "Bernoulli" trick (Wang et al., 2014) is widely used to set different probabilities for generating head or tail entities: For each relation type r, we calculate the averaged number  $a_{r,1}$  of heads h for a pair (r,t) and the averaged number  $a_{r,2}$  of tails t for a pair (h,r). We then define a Bernoulli distribution with success probability  $\lambda_r = \frac{a_{r,1}}{a_{r,1} + a_{r,2}}$  for sampling: given a correct triple (h,r,t), we corrupt this triple by replacing head entity with probability  $\lambda_r$  while replacing the tail entity with probability  $(1 - \lambda_r)$ .

Recently, Cai and Wang (2018) and Sun et al. (2019) proposed adversarial learning-based strategies for sampling incorrect triples. However, they did not provide a comparison between the adversarial learning-based strategies and the "Bernoulli" trick.

### 3 Specific models

## 3.1 Triple-based embedding models

**Translation-based models:** The Unstructured model (Bordes et al., 2012) assumes that the head and tail entity vectors are similar. As the Unstructured model does not take the relationship into account, it cannot distinguish different relation types. The Structured Embedding (SE) model (Bordes et al., 2011) assumes that the head and tail entities are similar only in a relation-dependent subspace, where each relation is represented by two different matrices. TransE (Bordes et al., 2013) is inspired by models

<sup>&</sup>lt;sup>1</sup>All the losses can also include an L2 regularization on the model parameters, which is not shown for simplification.

Model		Score function $f(h, r, t)$						
Translation	Unstructured	$-\ v_h-v_t\ _{\ell_{1/2}}$						
	SE	$-\ \mathbf{W}_{r,1} v_h - \mathbf{W}_{r,2} v_t\ _{\ell_{1/2}}$ where $\mathbf{W}_{r,1}, \mathbf{W}_{r,2} \in \mathbb{R}^{k  imes k}$						
	TransE	$-\ oldsymbol{v}_h + oldsymbol{v}_r - oldsymbol{v}_t\ _{\ell_{1/2}}$ where $oldsymbol{v}_r \in \mathbb{R}^k$						
	TransH	$-\ (\mathbf{I}-r_pr_p^{ op})oldsymbol{v}_h+oldsymbol{v}_r-(\mathbf{I}-r_pr_p^{ op})oldsymbol{v}_t\ _{\ell_{1/2}} \ \  ext{where}\ oldsymbol{r}_p,oldsymbol{v}_r\in\mathbb{R}^k$ , $\mathbf{I}$ denotes an identity matrix size $k imes k$						
	TransR	$-\ \mathbf{W}_r m{v}_h + m{v}_r - \mathbf{W}_r m{v}_t\ _{\ell_{1/2}} \;  ext{where} \; \mathbf{W}_r \in \mathbb{R}^{n  imes k} \; ,  m{v}_r \in \mathbb{R}^n$						
	STransE	$\ -\ \mathbf{W}_{r,1}m{v}_h+m{v}_r-\mathbf{W}_{r,2}m{v}_t\ _{\ell_{1/2}}$ where $\mathbf{W}_{r,1},\mathbf{W}_{r,2}\in\mathbb{R}^{k imes k}$ , $m{v}_r\in\mathbb{R}^k$						
	TranSparse	$-\ \mathbf{W}_{r,1}(\theta_{r,1})\boldsymbol{v}_h + \boldsymbol{v}_r - \mathbf{W}_{r,2}(\theta_{r,2})\boldsymbol{v}_t\ _{\ell_{1/2}} \text{ where } \mathbf{W}_{r,1}, \mathbf{W}_{r,2} \in \mathbb{R}^{n \times k} \ ; \ \theta_{r,1}, \theta_{r,2} \in \mathbb{R} \ ; \ \boldsymbol{v}_r \in \mathbb{R}^n$						
	TransD	$oxed{-\ (\mathbf{I}+oldsymbol{r}_poldsymbol{h}_p^ op)oldsymbol{v}_h+oldsymbol{v}_r-(\mathbf{I}+oldsymbol{r}_poldsymbol{t}_p^ op)oldsymbol{v}_t\ _{\ell_{1/2}}}  ext{ where } oldsymbol{r}_p,oldsymbol{v}_r,oldsymbol{h}_p,oldsymbol{t}_p\in\mathbb{R}^k$						
	lppTransD	$igg -\ (\mathbf{I}+oldsymbol{r}_{p,1}oldsymbol{h}_p^ op)oldsymbol{v}_h+oldsymbol{v}_r-(\mathbf{I}+oldsymbol{r}_{p,2}oldsymbol{t}_p^ op)oldsymbol{v}_t\ _{\ell_{1/2}} \  ext{where} \ oldsymbol{r}_{p,1},oldsymbol{r}_{p,2},oldsymbol{v}_r,oldsymbol{h}_p,oldsymbol{t}_p\in\mathbb{R}^k$						
	Bilinear	$oldsymbol{v}_h^{ op} \mathbf{W}_r oldsymbol{v}_t \;  ext{ where } \mathbf{W}_r \in \mathbb{R}^{k  imes k}$						
Tensor	DISTMULT	$oldsymbol{v}_h^{ op} \mathbf{W}_r v_t$ where $\mathbf{W}_r$ is a diagonal matrix $\in \mathbb{R}^{k  imes k}$						
& Te	SimplE	$\left  \frac{1}{2} \left( \boldsymbol{v}_{h,1}^{\top} \boldsymbol{W}_r \boldsymbol{v}_{t,2} + \boldsymbol{v}_{t,1}^{\top} \boldsymbol{W}_{r^{-1}} \boldsymbol{v}_{h,2} \right) \right  \text{ where } \boldsymbol{v}_{h,1}, \boldsymbol{v}_{h,2}, \boldsymbol{v}_{t,1}, \boldsymbol{v}_{t,2} \in \mathbb{R}^k \; \; ; \; \boldsymbol{W}_r \text{ and } \boldsymbol{W}_{r^{-1}} \text{ are diagonal matrices} \in \mathbb{R}^{k \times k}$						
ear	SME(bilinear)	$oldsymbol{v}_h^{ op}(\mathbf{M}_1 imes_3oldsymbol{v}_r)^{ op}(\mathbf{M}_2 imes_3oldsymbol{v}_r)oldsymbol{v}_t \  ext{ where } oldsymbol{v}_r\in\mathbb{R}^k \ ; \ \mathbf{M}_1,\mathbf{M}_2\in\mathbb{R}^{n imes k imes k}$						
Bilinear &	TuckER	$\mathbf{M} \times_1 v_h \times_2 v_r \times_3 v_t$ where $v_r \in \mathbb{R}^n$ , $\mathbf{M} \in \mathbb{R}^{k \times n \times k}$ ; $\times_d$ denotes the tensor product along the $d$ -th mode						
	HolE	$sigmoid(oldsymbol{v}_t^ op(oldsymbol{v}_h\staroldsymbol{v}_r))$ where $\star$ denotes circular correlation						
'ork	NTN	$\boxed{ \boldsymbol{v}_r^\top tanh(\boldsymbol{v}_h^\top \mathbf{M}_r \boldsymbol{v}_t + \mathbf{W}_{r,1} \boldsymbol{v}_h + \mathbf{W}_{r,2} \boldsymbol{v}_t + \mathbf{b}_r) \ \text{ where } \boldsymbol{v}_r, \mathbf{b}_r \in \mathbb{R}^n \ ; \ \mathbf{M}_r \in \mathbb{R}^{k \times k \times n} \ ; \ \mathbf{W}_{r,1}, \mathbf{W}_{r,2} \in \mathbb{R}^{n \times k} }$						
netw	ER-MLP	$sigmoid(\mathbf{w}^{\top}tanh(\boldsymbol{W}concat(\boldsymbol{v}_h,\boldsymbol{v}_r,\boldsymbol{v}_t)))$						
ıral 1	ConvE	$\boxed{ \boldsymbol{v}_t^\top sigmoid \left( \boldsymbol{W} vec \left( sigmoid \left( concat(\overline{\boldsymbol{v}}_h, \overline{\boldsymbol{v}}_r) * \boldsymbol{\Omega} \right) \right) \right)}$						
Net	ConvKB	$\mathbf{w}^{ op}$ concat (sigmoid $([oldsymbol{v}_h, oldsymbol{v}_r, oldsymbol{v}_t] * oldsymbol{\Omega}))$						
ctor	ComplEx	$Re\left(oldsymbol{c}_{h}^{T}\mathbf{C}_{r}\hat{oldsymbol{c}}_{t}\right)$ where $Re(c)$ denotes the real part of the complex value $c\in\mathbb{C}$						
Complex vector Neural network		$m{c}_h, m{c}_t \in \mathbb{C}^k \; \; ; \; m{C}_r \in \mathbb{C}^{k  imes k} \;  ext{is a diagonal matrix} \; \; ; \; \hat{m{c}}_t \;  ext{is the conjugate of} \; m{c}_t$						
nple	RotatE	$-\ m{c}_h\circm{c}_r-m{c}_t\ _{\ell_{1/2}}$ ; $m{c}_h,m{c}_r,m{c}_t\in\mathbb{C}^k$ where $\circ$ denotes the element-wise product						
Con	QuatE	$q_h\otimes rac{q_r}{ q_r }ullet q_t\ ;\ q_h,q_r,q_t\in \mathbb{H}^k\  ext{ where}\otimes  ext{and}ullet  ext{denote Hamilton and quaternion inner products}$						
Path	TransE-COMP	$\ -\ oldsymbol{v}_h+oldsymbol{v}_{r_1}+oldsymbol{v}_{r_2}++oldsymbol{v}_{r_m}-oldsymbol{v}_t\ _{\ell_{1/2}}$ where $oldsymbol{v}_{r_1},oldsymbol{v}_{r_2},,oldsymbol{v}_{r_m}\in\mathbb{R}^k$						
Pē	Bilinear-COMP	$oldsymbol{v}_h^{ op} \mathbf{W}_{r_1} \mathbf{W}_{r_2} \mathbf{W}_{r_m} v_t \;  ext{ where } \mathbf{W}_{r_1}, \mathbf{W}_{r_2},, \mathbf{W}_{r_m} \in \mathbb{R}^{k  imes k}$						

Table 1: The score functions f(h,r,t) of several prominent embedding models for KB completion. In these models, the entities h and t are represented by vectors  $v_h$  and  $v_t \in \mathbb{R}^k$ , respectively.  $\ell_{1/2}$  denotes either the L<sub>1</sub>-norm or the squared L<sub>2</sub>-norm. In ConvE,  $\overline{v}_h$  and  $\overline{v}_r$  denote a 2D reshaping of  $v_h$  and  $v_r$ , respectively. In both ConvE and ConvKB models, \* and  $\Omega$  denote a convolution operator and a set of filters, respectively.

such as the Word2Vec Skip-gram model (Mikolov et al., 2013) where relationships between words often correspond to translations in latent feature space. In particular, TransE learns low-dimensional and dense vectors for every entity and relation type, so that each relation type corresponds to a translation vector operating on the vectors representing the entities, i.e.,  $v_h + v_r \approx v_t$  for each fact triple (h, r, t). TransE thus is suitable for 1-to-1 relationships, such as "is\_capital\_of", where a head entity is linked to at most one tail entity given a relation type. Because of using only one translation vector to represent each relation type, TransE is not well-suited for Many-to-1, 1-to-Many and Many-to-Many relationships, such as for relation types "born\_in", "place\_of\_birth" and "research\_fields." For example in Figure 2, using one vector representing the relation type "born\_in" cannot capture both the translating direction from "Patti" to "Miami" and its inverse direction from "Mom" to "Austin."

To overcome those issues of TransE, TransH (Wang et al., 2014) associates each relation with a relation-specific hyperplane and uses a projection vector to project entity vectors onto that hyperplane. TransD (Ji et al., 2015) and TransR/CTransR (Lin et al., 2015b) extend TransH by using two projection vectors and a matrix to project entity vectors into a relation-specific space, respectively. Similar to TransR, TransR-FT (Feng et al., 2016a) also uses a matrix to project head and tail entity vectors. TEKE\_H (Wang and Li, 2016) extends TransH to incorporate rich context information in an external text corpus. lppTransD (Yoon et al., 2016) extends TransD to additionally use two projection vectors for representing each relation. STransE (Nguyen et al., 2016a) and TranSparse (Ji et al., 2016) can be

 $<sup>^2</sup>$ A relation type r is classified Many-to-1 if multiple head entities can be connected by r to at most one tail entity. A relation type r is classified 1-to-Many if multiple tail entities can be linked by r from at most one head entity. A relation type r is classified Many-to-Many if multiple head entities can be connected by r to a tail entity and vice versa.

viewed as direct extensions of TransR, where head and tail entities are associated with their own projection matrices. Unlike STransE, TranSparse uses adaptive sparse matrices, whose sparse degrees are defined based on the number of entities linked by relations. TranSparse-DT (Chang et al., 2017) is an extension of TranSparse with a dynamic translation. ITransF (Xie et al., 2017) can be considered as a generalization of STransE, which allows the sharing of statistic regularities between relation projection matrices and alleviates data sparsity issue. Furthermore, TorusE (Ebisu and Ichise, 2018) embeds entities and relations on a torus to handle TransE's regularization problem which forces entity embeddings to be on a sphere in the embedding vector space.

**Bilinear- & Tensor-based models:** DISTMULT (Yang et al., 2015) is based on the Bilinear model (Nickel et al., 2011; Jenatton et al., 2012) where each relation is represented by a diagonal matrix rather than a full matrix. SimplE (Kazemi and Poole, 2018) extends DISTMULT to allow two embeddings of each entity to be learned dependently. Such quadratic forms are also used to model entities and relations in KG2E (He et al., 2015), TATEC (García-Durán et al., 2016), TransG (Xiao et al., 2016), RSTE (Tay et al., 2017), ANALOGY (Liu et al., 2017) and Dihedral (Xu and Li, 2019). SME(bilinear) (Bordes et al., 2012) is proposed to first separately combine entity-relation pairs (h, r) and (r, t) and then semantically match these combinations, using tensor product. HolE (Nickel et al., 2016b) uses circular correlation—a compositional operator—which can be interpreted as a compression of the tensor product. In addition, TuckER (Balazevic et al., 2019) is a linear model based on the Tucker tensor decomposition of the binary tensor representation of KB triples.

**Neural network-based models:** The neural tensor network (NTN) model (Socher et al., 2013) also uses a bilinear tensor operator to represent each relation while ProjE (Shi and Weninger, 2017) can be viewed as simplified versions of NTN. ER-MLP (Dong et al., 2014) represents each triple by a vector obtained from concatenating head, relation and tail embeddings, then feeds this vector into a single-layer MLP with one-node output layer. ConvE (Dettmers et al., 2018) and ConvKB (Nguyen et al., 2018) are based on convolutional neural networks. ConvE uses a convolution layer directly over 2D reshaping of head-entity and relation embeddings, while ConvKB applies a convolution layer over the embedding triples (here each triple (h, r, t) is represented as a 3-column matrix where each column vector represents a triple element). HypER (Balažević et al., 2019) simplifies ConvE by using a hypernetwork to produce 1D convolutional filters for each relation, then extracts relation-specific features from head entity embeddings. Conv-TransE (Shang et al., 2019) extends ConvE to keep the translational characteristic between entities and relations. InteractE (Vashishth et al., 2020) uses a circular convolution operator and a checkered reshaping function instead of the standard convolution operator and 2D stack reshaping function in ConvE. CapsE (Nguyen et al., 2019) extends ConvKB by stacking a capsule network layer (Sabour et al., 2017) on top of the convolution layer.

Complex vector-based models: Instead of embedding entities and relations in the real-valued vector space, ComplEx (Trouillon et al., 2016) is an extension of DISTMULT in the complex vector space. ComplEx-N3 (Lacroix et al., 2018) extends ComplEx with weighted nuclear 3-norm. Also in the complex vector space, RotatE (Sun et al., 2019) defines each relation as a rotation from the head entity to the tail entity. QuatE (Zhang et al., 2019) represents entities by quaternion embeddings (i.e. hypercomplex-valued embeddings) and models relations as rotations in the quaternion space by employing the Hamilton and quaternion-inner products.

## 3.2 Relation path-based embedding models

All embedding models mentioned above in Section 3.1 only take triples into account. Thus, these models ignore potentially useful information implicitly presented by the structure of the KB. For example, the relation path  $h \xrightarrow{\text{born.in.city}} e \xrightarrow{\text{city.in.country}} t$  should indicate a relationship "nationality" between the h and t entities. Also, neighborhood information of entities could be useful for predicting the relationship between two entities as well. For example, in the KB NELL (Carlson et al., 2010), we have information such as if a person works for an organization and this person also leads that organization, then it is likely that this person is the CEO of that organization.

Many research has shown that relation paths between entities in KBs provide richer context information and improve the performance of embedding models for KB completion (Luo et al., 2015; Liang and Forbus, 2015; García-Durán et al., 2015; Guu et al., 2015; Toutanova et al., 2016; Durán and Niepert, 2018; Takahashi et al., 2018; Chen et al., 2018). In particular, Luo et al. (2015) constructed relation paths between entities and, viewing entities and relations in the path as pseudo-words, then applied Word2Vec algorithms (Mikolov et al., 2013) to produce pre-trained vectors for these pseudo-words. Luo et al. (2015) showed that using these pre-trained vectors for initialization helps to improve the performance of models TransE (Bordes et al., 2013), SME (Bordes et al., 2012) and SE (Bordes et al., 2011). Liang and Forbus (2015) used the plausibility score produced by SME to compute the weights of relation paths.

PTransE-RNN (Lin et al., 2015a) models relation paths by using a recurrent neural network (RNN). Das et al. (2017)'s model and ROPs (Yin et al., 2018) also apply RNN to model the path between an entity pair, however, in contrast to PTransE-RNN, they additionally take the intermediate entities present in the path into account. IRN (Shen et al., 2017) uses a shared memory and RNN-based controller to implicitly model multi-step structured relationships. RTransE (García-Durán et al., 2015), PTransE-ADD (Lin et al., 2015a) and TransE-COMP (Guu et al., 2015) extend TransE to represent a relation path by a vector which is the sum of the vectors of all relations in the path. In Bilinear-COMP (Guu et al., 2015) and PRUNED-PATHS (Toutanova et al., 2016), each relation is a matrix and so it represents the relation path by matrix multiplication. Durán and Niepert (2018) proposed the KB<sub>LRN</sub> framework to combine relational paths with latent and numerical features.

The neighborhood mixture model TransE-NMM (Nguyen et al., 2016b) can be also viewed as a three-relation path model as it takes into account the neighborhood entity and relation information of both head and tail entities in each triple. Neighborhood information is also exploited in R-GCN (Schlichtkrull et al., 2018), SACN (Shang et al., 2019) and KBAT (Nathani et al., 2019), which generalize graph convolutional networks (Kipf and Welling, 2017) and graph attention networks (Veličković et al., 2018) for dealing with highly multi-relational data such as knowledge bases. For computing the final representation of an entity, they make use of layer-wise propagation to accumulate linearly-transformed embeddings of its neighboring entities through a normalized sum with different relational weights. For link prediction, R-GCN, SACN and KBAT apply DISTMULT, Conv-TransE and ConvKB to compute triple scores, respectively.

# 3.3 Other KB completion models

The Path Ranking Algorithm (PRA) (Lao and Cohen, 2010) is a random walk inference technique which was proposed to predict a new relationship between two entities in KBs. Lao et al. (2011) used PRA to estimate the probability of an unseen triple as a combination of weighted random walks that follow different paths linking the head entity and tail entity in the KB. Gardner et al. (2014) made use of an external text corpus to increase the connectivity of the KB used as the input to PRA. Gardner and Mitchell (2015) improved PRA by proposing a subgraph feature extraction technique to make the generation of random walks in KBs more efficient and expressive, while Wang et al. (2016) extended PRA to couple the path ranking of multiple relations. PRA can also be used in conjunction with first-order logic in the discriminative Gaifman model (Niepert, 2016). In addition, Neelakantan et al. (2015) used a RNN to learn vector representations of PRA-style relation paths between entities in the KB. Other random-walk based learning algorithms for KB completion can be also found in Feng et al. (2016b), Liu et al. (2016), Wei et al. (2016), Mazumder and Liu (2017) and Das et al. (2018).

Yang et al. (2017) proposed a Neural Logic Programming (LP) framework to learning probabilistic first-order logical rules for KB reasoning, producing competitive link prediction performances. Feldman et al. (2019) presented an approach to generate sentences from triples via hand-craft templates, and then use the likelihoods produced by the pre-trained BERT (Devlin et al., 2019) for these generated sentences to score the plausibility of the corresponding triples. See other methods for learning from KBs and multi-relational data in Nickel et al. (2016a) and Wang et al. (2017).

Dataset	E	R	#Triples in train/valid/te		
FB15k	14,951	1,345	483,142	50,000	59,071
WN18	40,943	18	141,442	5,000	5,000
FB15k-237	14,541	237	272,115	17,535	20,466
WN18RR	40,943	11	86,835	3,034	3,134

Table 2: Statistics of benchmark experimental datasets.

#### 4 Evaluation task

The standard evaluation task of entity prediction, i.e. link prediction (Bordes et al., 2013), is proposed to evaluate embedding models for KB completion.<sup>3</sup>

**Datasets:** Information about benchmark datasets for KB completion evaluation is given in Table 2. FB15k and WN18 are derived from the large real-world KB FreeBase (Bollacker et al., 2008) and the large lexical KB WordNet (Miller, 1995), respectively. Toutanova and Chen (2015) noted that FB15k and WN18 are not challenging datasets because they contain many reversible triples. Dettmers et al. (2018) showed a concrete example: A test triple (feline, hyponym, cat) can be mapped to a training triple (cat, hypernym, feline), thus knowing that "hyponym" and "hypernym" are reversible allows us to easily predict the majority of test triples. So, datasets FB15k-237 (Toutanova and Chen, 2015) and WN18RR (Dettmers et al., 2018) are created to serve as realistic KB completion datasets which represent a more challenging learning setting. FB15k-237 and WN18RR are subsets of FB15k and WN18, respectively.

# 4.1 Task description

The entity prediction task, i.e. link prediction (Bordes et al., 2013), predicts the head or the tail entity given the relation type and the other entity, i.e. predicting h given (?, r, t) or predicting t given (h, r, ?) where ? denotes the missing element. The results are evaluated using a ranking induced by the function f(h, r, t) on test triples.

Each correct test triple (h, r, t) is corrupted by replacing either its head or tail entity by each of the possible entities in turn, and then these candidates are ranked in descending order of their plausibility score. The "Filtered" setting protocol, described in Bordes et al. (2013), filters out before ranking any corrupted triples that appear in the KB. Ranking a corrupted triple appearing in the KB (i.e. a correct triple) higher than the original test triple is also correct, thus this "Filtered" setting provides a clear view on the ranking performance.

In addition to the mean rank and the Hits@10 (i.e., the proportion of test triples for which the target entity is ranked in the top 10 predictions), which were originally used in the entity prediction task (Bordes et al., 2013), recent work also reports the mean reciprocal rank (MRR).<sup>4</sup> Mean rank is always greater or equal to 1 and the lower mean rank indicates better entity prediction performance, while MRR and Hits@10 scores always range from 0.0 to 1.0, and higher score reflects better prediction result.

# 4.2 Main results

Tables 3 and 4 list recent entity prediction results of KB completion models on FB15k and WN18 and on FB15k-237 and WN18RR, respectively. In Table 3, the first 27 rows report the performance of triple-based models that directly optimize a score function for the triples in a KB, i.e. they do not exploit information about alternative paths between head and tail entities. The next 9 rows report results of models that exploit information about relation paths or neighborhood information. The last 2 rows present results for models which make use of textual mentions derived from a large external corpus. In Table 4, the last 5 rows report results of models that exploit the path or neighborhood information.

In general, Tables 3 and 4 show that the models using external corpus information or employing path information achieve better scores than the triple-based models that do not use such information. In terms

<sup>&</sup>lt;sup>3</sup>Another evaluation task for KB completion is triple classification (Socher et al., 2013), however, it is not as widely used as the link prediction task.

<sup>&</sup>lt;sup>4</sup>See Baeza-Yates and Ribeiro-Neto (2011) for definitions of the mean rank, Hits@10 and MRR. Some recent work additionally reported Hits@1. However, formulas of MRR and Hits@1 show a strong correlation between these two scores. So using Hits@1 might not really reveal any additional insight.

Method         FBIS         W         WIND         WIND         MR         @10         MRR         WIND         MR         MR         PRIA         PRIA <th colspan="2" rowspan="2">Method</th> <th colspan="6">Filtered</th>	Method		Filtered					
TransR (Lin et al., 2015b)         77         68.7         -         225         92.0         -           CTransR (Lin et al., 2015b)         75         70.2         -         218         92.3         -           KG2E (He et al., 2015)         59         74.0         -         331         92.8         -           TransD (Ji et al., 2015)         77.3         -         212         92.2         -           IppTransD (Yoon et al., 2016)         78         78.7         -         270         94.3         -           TransG (Xiao et al., 2016)         82         79.5         -         211         93.2         -           TransParse-DT (Chang et al., 2017)         65         81.0         -         205         94.2         -           NTN (Socher et al., 2013)         -         41.4         0.25         -         66.1         0.53           TransE (Bordes et al., 2013)         -         74.9         0.463         -         94.9         0.938           ComplEx (Trouillon et al., 2016b)         -         73.9         0.524         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.8         0.727         -         94.7         0			FB15k			WN18		
CTransR (Lin et al., 2015b)         75         70.2         -         218         92.3         -           KG2E (He et al., 2015)         59         74.0         -         331         92.8         -           TransD (Ji et al., 2015)         91         77.3         -         212         92.2         -           IppTransD (Yoon et al., 2016)         78         78.7         -         270         94.3         -           TransG (Xiao et al., 2016)         78         79.8         -         470         93.3         -           TransParse (Ji et al., 2016)         82         79.5         -         211         93.2         -           TransParse-DT (Chang et al., 2017)         79         80.2         -         221         94.3         -           NTN (Socher et al., 2013)         -         41.4         0.25         -         66.1         0.53           TransE (Bordes et al., 2016b)         -         73.9         0.524         -         94.9         0.43         0.495           HolE (Nickel et al., 2016b)         -         84.0         0.25         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.4         0.725         - <th></th> <th>MR</th> <th></th> <th>MRR</th> <th>MR</th> <th>@10</th> <th>MRR</th>		MR		MRR	MR	@10	MRR	
KG2E (He et al., 2015)         59         74.0         -         331         92.8         -           TransD (Ji et al., 2015)         91         77.3         -         212         92.2         -           IppTransD (Yoon et al., 2016)         78         78.7         -         270         94.3         -           TransG (Xiao et al., 2016)         98         79.8         -         470         93.3         -           TranSparse (Ji et al., 2016)         82         79.5         -         211         93.2         -           ITransF (Xie et al., 2017)         65         81.0         -         205         94.2         -           NTN (Socher et al., 2013)         -         41.4         0.25         -         66.1         0.53           TransE (Bordes et al., 2016b)         -         74.9         0.463         -         94.3         0.495           Holle (Nickel et al., 2016b)         -         74.9         0.463         -         94.7         0.941           ANAL OGY (Liu et al., 2016)         -         84.0         0.692         -         94.7         0.942           Simple (Kazemi and Poole, 2018)         -         85.4         0.725         -         94.7 <td>TransR (Lin et al., 2015b)</td> <td>77</td> <td>68.7</td> <td>-</td> <td>225</td> <td>92.0</td> <td>-</td>	TransR (Lin et al., 2015b)	77	68.7	-	225	92.0	-	
TransD (Ji et al., 2015)         91         77.3         -         212         92.2         -           IppTransD (Yoon et al., 2016)         78         78.7         -         270         94.3         -           TransG (Xiao et al., 2016)         98         79.8         -         470         93.3         -           Transparse (Ji et al., 2016)         82         79.5         -         211         93.2         -           Transparse-DT (Chang et al., 2017)         79         80.2         -         221         94.3         -           NTN (Socher et al., 2013)         -         41.4         0.25         -         66.1         0.53           TransE (Bordes et al., 2013) [♥]         -         74.9         0.463         -         94.3         0.495           HolE (Nickel et al., 2016b)         -         73.9         0.524         -         94.9         0.938           ComplEx (Trouillon et al., 2016b)         -         84.0         0.692         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.2         0.733         -         94.7         0.942           SimplE (Kazemi and Ichise, 2018)         -         83.2         0.733			70.2	-	218	92.3	-	
IppTransD (Yoon et al., 2016)	KG2E (He et al., 2015)	59	74.0	-	331	92.8	-	
TransG (Xiao et al., 2016)         98         79.8         -         470         93.3         -           TranSparse (Ji et al., 2016)         82         79.5         -         211         93.2         -           TranSparse-DT (Chang et al., 2017)         79         80.2         -         221         94.3         -           NTN (Socher et al., 2013)         -         41.4         0.25         -         66.1         0.53           TransE (Bordes et al., 2016b)         -         74.9         0.463         -         94.3         0.495           HolE (Nickel et al., 2016b)         -         73.9         0.524         -         94.7         0.941           ANALOGY (Liu et al., 2017)         -         85.4         0.725         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.8         0.727         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.8         0.727         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.8         0.727         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.2         0.733	TransD (Ji et al., 2015)	91	77.3	-	212	92.2	-	
TranSparse (Ji et al., 2016)         82         79.5         -         211         93.2         -           TranSparse-DT (Chang et al., 2017)         79         80.2         -         221         94.3         -           NTN (Socher et al., 2013)         -         41.4         0.25         -         66.1         0.53           TransE (Bordes et al., 2013) [♥]         -         74.9         0.463         -         94.3         0.495           HolE (Nickel et al., 2016b)         -         73.9         0.524         -         94.9         0.938           ComplEx (Trouillon et al., 2016)         -         84.0         0.692         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.8         0.727         -         94.7         0.942           TorusE (Ebisu and Ichise, 2018)         -         83.2         0.733         -         95.4         0.947           STransE (Nguyen et al., 2016a)         69         79.7         0.543         206         93.4         0.657           ER-MLP (Dong et al., 2014) [♠]         81         80.1         0.570         299         94.2         0.895           DISTMULT (Yang et al., 2015) [♠]         42         89.3 <td>lppTransD (Yoon et al., 2016)</td> <td>78</td> <td>78.7</td> <td>-</td> <td>270</td> <td>94.3</td> <td>-</td>	lppTransD (Yoon et al., 2016)	78	78.7	-	270	94.3	-	
TransSparse-DT (Chang et al., 2017)         79         80.2         -         221         94.3         -           ITransF (Xie et al., 2017)         65         81.0         -         205         94.2         -           NTN (Socher et al., 2013)         -         41.4         0.25         -         66.1         0.53           TransE (Bordes et al., 2016b)         -         74.9         0.463         -         94.3         0.495           HolE (Nickel et al., 2016b)         -         73.9         0.524         -         94.9         0.938           ComplEx (Trouillon et al., 2016)         -         84.0         0.692         -         94.7         0.941           ANALOGY (Liu et al., 2017)         -         85.4         0.725         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.8         0.727         -         94.7         0.942           Toruse (Ebisu and Ichise, 2018)         -         83.8         0.727         94.7         0.942           Toruse (Bultimer et al., 2016a)         69         79.7         0.543         206         93.4         0.657           ER-MLP (Dong et al., 2014) [♠]         81         80.1         0.798	TransG (Xiao et al., 2016)	98	79.8	-	470	93.3	-	
ITransF (Xie et al., 2017)         65         81.0         -         205         94.2         -           NTN (Socher et al., 2013)         -         41.4         0.25         -         66.1         0.53           TransE (Bordes et al., 2013) [♥]         -         74.9         0.463         -         94.3         0.495           HolE (Nickel et al., 2016b)         -         73.9         0.524         -         94.9         0.938           ComplEx (Trouillon et al., 2016)         -         84.0         0.692         -         94.7         0.941           ANALOGY (Liu et al., 2017)         -         85.4         0.725         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.8         0.727         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.8         0.727         -         94.7         0.942           SimplE (Kazemi and Poole, 2018)         -         83.8         0.727         -         94.7         0.942           STransE (Busiu and Ichise, 2018)         68         83.2         0.733         -         95.4         0.957           DISTMULT (Yang et al., 2014) [♠]         42         87.3	TranSparse (Ji et al., 2016)	82	79.5	-	211	93.2	-	
NTN (Socher et al., 2013)	TranSparse-DT (Chang et al., 2017)	79	80.2	-	221	94.3	-	
TransE (Bordes et al., 2013) [♥]       -       74.9       0.463       -       94.3       0.493         HolE (Nickel et al., 2016b)       -       73.9       0.524       -       94.9       0.938         ComplEx (Trouillon et al., 2016)       -       84.0       0.692       -       94.7       0.941         ANALOGY (Liu et al., 2017)       -       85.4       0.725       -       94.7       0.942         SimplE (Kazemi and Poole, 2018)       -       83.8       0.727       -       94.7       0.942         TorusE (Ebisu and Ichise, 2018)       -       83.2       0.733       -       95.4       0.947         STransE (Nguyen et al., 2016a)       69       79.7       0.543       206       93.4       0.657         ER-MLP (Dong et al., 2014) [♠]       81       80.1       0.570       299       94.2       0.895         DISTMULT (Yang et al., 2015) [♠]       42       89.3       0.798       655       94.6       0.797         ConvE (Dettmers et al., 2018)       44       88.5       0.790       431       95.8       0.951         RypER (Balažević et al., 2018)       40       88.4       0.797       309       95.8       0.951         Rud (Ey	ITransF (Xie et al., 2017)	65	81.0	-	<u>205</u>	94.2	-	
HolE (Nickel et al., 2016b)-73.90.524-94.90.938ComplEx (Trouillon et al., 2016)-84.00.692-94.70.941ANALOGY (Liu et al., 2017)-85.40.725-94.70.942SimplE (Kazemi and Poole, 2018)-83.80.727-94.70.942TorusE (Ebisu and Ichise, 2018)-83.20.733-95.40.947STransE (Nguyen et al., 2014) [ ]8180.10.57029994.20.855ER-MLP (Dong et al., 2015) [ ]4289.30.79865594.60.797ConvE (Dettmers et al., 2018)6487.30.74550495.50.942HypER (Balažević et al., 2019)4488.50.79043195.80.951RotatE (Sun et al., 2019)4088.40.79730995.90.949QuatE (Zhang et al., 2019)4088.40.79730995.90.950ComplEx-N3 (Lacroix et al., 2018)-910.86-960.95TuckER (Balazevic et al., 2019)-89.20.795-95.80.953IRN (Shen et al., 2017)3892.7-24995.3-Prje (Shi and Weninger, 2017)3488.4PTransE-ADD (Lin et al., 2015a)5884.6GAKE (Feng et al., 2016b)11964.8 <tr< td=""><td>NTN (Socher et al., 2013)</td><td>-</td><td>41.4</td><td>0.25</td><td>-</td><td>66.1</td><td>0.53</td></tr<>	NTN (Socher et al., 2013)	-	41.4	0.25	-	66.1	0.53	
ComplEx (Trouillon et al., 2016)       -       84.0 $0.692$ -       94.7 $0.942$ ANALOGY (Liu et al., 2017)       -       85.4 $0.725$ -       94.7 $0.942$ SimplE (Kazemi and Poole, 2018)       -       83.8 $0.727$ - $94.7$ $0.942$ TorusE (Ebisu and Ichise, 2018)       -       83.2 $0.733$ - $95.4$ $0.947$ STransE (Nguyen et al., 2016a)       69 $79.7$ $0.543$ $206$ $93.4$ $0.657$ ER-MLP (Dong et al., 2014) [ ]       81 $80.1$ $0.570$ $299$ $94.2$ $0.895$ DISTMULT (Yang et al., 2015) [ ]       42 $89.3$ $0.798$ $655$ $94.6$ $0.797$ ConvE (Dettmers et al., 2018)       44 $88.5$ $0.790$ $431$ $95.8$ $0.951$ RypER (Balažević et al., 2019)       40 $88.4$ $0.797$ $309$ $95.9$ $0.950$ Quate (Zhang et al., 2019)       17 $90.0$ $0.782$ $162$ $95.9$ $0.950$ TuckER (Balazevic et al., 2019)       38 $92.7$ $0.795$ <td< td=""><td>TransE (Bordes et al., 2013) [♥]</td><td>-</td><td>74.9</td><td>0.463</td><td>-</td><td>94.3</td><td>0.495</td></td<>	TransE (Bordes et al., 2013) [♥]	-	74.9	0.463	-	94.3	0.495	
ANALOGY (Liu et al., 2017)	HolE (Nickel et al., 2016b)	-	73.9	0.524	-	94.9	0.938	
SimplE (Kazemi and Poole, 2018)       -       83.8       0.727       -       94.7       0.942         TorusE (Ebisu and Ichise, 2018)       -       83.2       0.733       -       95.4       0.947         STransE (Nguyen et al., 2016a)       69       79.7       0.543       206       93.4       0.657         ER-MLP (Dong et al., 2014) [♠]       81       80.1       0.570       299       94.2       0.895         DISTMULT (Yang et al., 2015) [♣]       42       89.3       0.798       655       94.6       0.797         ConvE (Dettmers et al., 2018)       64       87.3       0.745       504       95.5       0.942         HypER (Balažević et al., 2019)       44       88.5       0.790       431       95.8       0.951         Rotate (Sun et al., 2019)       40       88.4       0.797       309       95.9       0.949         Quate (Zhang et al., 2019)       17       90.0       0.782       162       95.9       0.950         ComplEx-N3 (Lacroix et al., 2018)       -       91       0.86       -       96       0.95         IRN (Shen et al., 2017)       38       92.7       -       249       95.3       -         PriansE-ADD (Lin et al.	ComplEx (Trouillon et al., 2016)	-	84.0	0.692	-	94.7	0.941	
TorusE (Ebisu and Ichise, 2018)         -         83.2         0.733         -         95.4         0.947           STransE (Nguyen et al., 2016a)         69         79.7         0.543         206         93.4         0.657           ER-MLP (Dong et al., 2014) [ ]         81         80.1         0.570         299         94.2         0.895           DISTMULT (Yang et al., 2015) [ ]         42         89.3         0.798         655         94.6         0.797           ConvE (Dettmers et al., 2018)         64         87.3         0.745         504         95.5         0.942           HypER (Balažević et al., 2019)         44         88.5         0.790         431         95.8         0.951           RotatE (Sun et al., 2019)         40         88.4         0.797         309         95.9         0.949           QuatE (Zhang et al., 2019)         17         90.0         0.782         162         95.9         0.950           ComplEx-N3 (Lacroix et al., 2018)         -         91         0.86         -         96         0.95           TuckER (Balazevic et al., 2017)         38         92.7         -         249         95.3         -           Proje (Shi and Weninger, 2017)         34         <	ANALOGY (Liu et al., 2017)		85.4	0.725	-	94.7	0.942	
STransE (Nguyen et al., 2016a)       69       79.7       0.543       206       93.4       0.657         ER-MLP (Dong et al., 2014) [ ]       81       80.1       0.570       299       94.2       0.895         DISTMULT (Yang et al., 2015) [ ]       42       89.3 $0.798$ 655       94.6       0.797         ConvE (Dettmers et al., 2018)       64       87.3       0.745       504       95.5       0.942         HypER (Balažević et al., 2019)       44       88.5       0.790       431       95.8       0.951         RotatE (Sun et al., 2019)       40       88.4       0.797       309       95.9       0.949         QuatE (Zhang et al., 2019)       17       90.0       0.782       162       95.9       0.950         ComplEx-N3 (Lacroix et al., 2018)       -       91       0.86       -       96       0.95         TuckER (Balazevic et al., 2019)       -       89.2       0.795       -       95.8       0.953         IRN (Shen et al., 2017)       38       92.7       -       249       95.3       -         PrizansE (García-Durán et al., 2015a)       50       76.2       -       -       -       -         PTransE-ADD (Lin et al.,	SimplE (Kazemi and Poole, 2018)	-	83.8	0.727	-	94.7	0.942	
ER-MLP (Dong et al., 2014) [ ]       81       80.1       0.570       299       94.2       0.895         DISTMULT (Yang et al., 2015) [ ]       42       89.3 $0.798$ 655       94.6       0.797         ConvE (Dettmers et al., 2018)       64       87.3 $0.745$ 504       95.5       0.942         HypER (Balažević et al., 2019)       44       88.5       0.790       431       95.8       0.951         RotatE (Sun et al., 2019)       40       88.4       0.797       309       95.9       0.949         QuatE (Zhang et al., 2019)       17       90.0       0.782       162       95.9       0.950         ComplEx-N3 (Lacroix et al., 2018)       -       91       0.86       -       96       0.95         TuckER (Balazevic et al., 2019)       -       89.2       0.795       -       95.8       0.953         IRN (Shen et al., 2017)       38       92.7       -       249       95.3       -         ProjE (Shi and Weninger, 2017)       34       88.4       -       -       -       -         RTransE (García-Durán et al., 2015a)       58       84.6       -       -       -       -         PTransE-RNN (Lin et al., 2015b)	TorusE (Ebisu and Ichise, 2018)	-	83.2	0.733	-	95.4	0.947	
DISTMULT (Yang et al., 2015) [♣] 42 89.3 0.798 655 94.6 0.797 ConvE (Dettmers et al., 2018) 64 87.3 0.745 504 95.5 0.942 HypER (Balažević et al., 2019) 44 88.5 0.790 431 95.8 0.951 RotatE (Sun et al., 2019) 40 88.4 0.797 309 95.9 0.949 QuatE (Zhang et al., 2019) 17 90.0 0.782 162 95.9 0.950 ComplEx-N3 (Lacroix et al., 2018) - 91 0.86 - 96 0.95 TuckER (Balazevic et al., 2019) - 89.2 0.795 - 95.8 0.953 IRN (Shen et al., 2017) 38 92.7 - 249 95.3 - 249 ProjE (Shi and Weninger, 2017) 34 88.4 249 95.3 - 249 PransE-ADD (Lin et al., 2015a) 58 84.6 2 - 2 - 2 - 2 - 2 - 2 - 2 - 2 - 2	STransE (Nguyen et al., 2016a)	69	79.7	0.543	206	93.4	0.657	
ConvE (Dettmers et al., 2018)         64         87.3         0.745         504         95.5         0.942           HypER (Balažević et al., 2019)         44         88.5         0.790         431         95.8         0.951           RotatE (Sun et al., 2019)         40         88.4         0.797         309         95.9         0.949           QuatE (Zhang et al., 2019)         17         90.0         0.782         162         95.9         0.950           ComplEx-N3 (Lacroix et al., 2018)         -         91         0.86         -         96         0.95           TuckER (Balazevic et al., 2019)         -         89.2         0.795         -         95.8         0.953           IRN (Shen et al., 2017)         38         92.7         -         249         95.3         -           ProjE (Shi and Weninger, 2017)         34         88.4         -         -         -         -         -           RTransE (García-Durán et al., 2015)         50         76.2         -         -         -         -           PTransE-ADD (Lin et al., 2015a)         92         82.2         -         -         -         -           GAKE (Feng et al., 2016b)         119         64.8         - <td>ER-MLP (Dong et al., 2014) [♠]</td> <td>81</td> <td>80.1</td> <td>0.570</td> <td>299</td> <td>94.2</td> <td>0.895</td>	ER-MLP (Dong et al., 2014) [♠]	81	80.1	0.570	299	94.2	0.895	
HypER (Balažević et al., 2019)       44       88.5       0.790       431       95.8       0.951         RotatE (Sun et al., 2019)       40       88.4       0.797       309       95.9       0.949         QuatE (Zhang et al., 2019)       17       90.0       0.782       162       95.9       0.950         ComplEx-N3 (Lacroix et al., 2018)       -       91       0.86       -       96       0.95         TuckER (Balazevic et al., 2019)       -       89.2       0.795       -       95.8       0.953         IRN (Shen et al., 2017)       38       92.7       -       249       95.3       -         ProjE (Shi and Weninger, 2017)       34       88.4       -       -       -       -         RTransE (García-Durán et al., 2015)       50       76.2       -       -       -       -         PTransE-ADD (Lin et al., 2015a)       58       84.6       -       -       -       -         GAKE (Feng et al., 2016b)       119       64.8       -       -       -       -         Gaifman (Niepert, 2016)       75       84.2       -       352       93.9       -         Hiri (Liu et al., 2016)       -       70.3       0.603	DISTMULT (Yang et al., 2015) [♣]	42	89.3	0.798	655	94.6	0.797	
RotatE (Sun et al., 2019)         40         88.4         0.797         309         95.9         0.949           QuatE (Zhang et al., 2019)         17         90.0         0.782         162         95.9         0.950           ComplEx-N3 (Lacroix et al., 2018)         -         91         0.86         -         96         0.95           TuckER (Balazevic et al., 2019)         -         89.2         0.795         -         95.8         0.953           IRN (Shen et al., 2017)         38         92.7         -         249         95.3         -           ProjE (Shi and Weninger, 2017)         34         88.4         -         -         -         -           RTransE (García-Durán et al., 2015)         50         76.2         -         -         -         -           PTransE-ADD (Lin et al., 2015a)         58         84.6         -         -         -         -           PTransE-RNN (Lin et al., 2015a)         92         82.2         -         -         -         -           GAKE (Feng et al., 2016b)         119         64.8         -         -         -         -           Hiri (Liu et al., 2016)         -         70.3         0.603         -         90.8	ConvE (Dettmers et al., 2018)	64	87.3	0.745	504	95.5	0.942	
QuatE (Zhang et al., 2019)       17       90.0       0.782       162       95.9       0.950         ComplEx-N3 (Lacroix et al., 2018)       -       91       0.86       -       96       0.95         TuckER (Balazevic et al., 2019)       -       89.2       0.795       -       95.8       0.953         IRN (Shen et al., 2017)       38       92.7       -       249       95.3       -         ProjE (Shi and Weninger, 2017)       34       88.4       -       -       -       -         RTransE (García-Durán et al., 2015)       50       76.2       -       -       -       -         PTransE-ADD (Lin et al., 2015a)       58       84.6       -       -       -       -         PTransE-RNN (Lin et al., 2015a)       92       82.2       -       -       -       -         GAKE (Feng et al., 2016b)       119       64.8       -       -       -       -         Gaifman (Niepert, 2016)       75       84.2       -       352       93.9       -         Hiri (Liu et al., 2016)       -       70.3       0.603       -       90.8       0.691         Neural LP (Yang et al., 2017)       -       83.7       0.76       - </td <td>HypER (Balažević et al., 2019)</td> <td>44</td> <td>88.5</td> <td></td> <td>431</td> <td>95.8</td> <td>0.951</td>	HypER (Balažević et al., 2019)	44	88.5		431	95.8	0.951	
ComplEx-N3 (Lacroix et al., 2018)         -         91         0.86         -         96         0.95           TuckER (Balazevic et al., 2019)         -         89.2         0.795         -         95.8         0.953           IRN (Shen et al., 2017)         38         92.7         -         249         95.3         -           ProjE (Shi and Weninger, 2017)         34         88.4         -         -         -         -           RTransE (García-Durán et al., 2015)         50         76.2         -         -         -         -           PTransE-ADD (Lin et al., 2015a)         58         84.6         -         -         -         -           PTransE-RNN (Lin et al., 2015a)         92         82.2         -         -         -         -           GAKE (Feng et al., 2016b)         119         64.8         -         -         -         -           Gaifman (Niepert, 2016)         75         84.2         -         352         93.9         -           Hiri (Liu et al., 2016)         -         70.3         0.603         -         90.8         0.691           Neural LP (Yang et al., 2017)         -         83.7         0.76         -         94.5         0.	RotatE (Sun et al., 2019)	40	88.4	0.797	309	<u>95.9</u>	0.949	
TuckER (Balazevic et al., 2019)       -       89.2       0.795       -       95.8       0.953         IRN (Shen et al., 2017)       38       92.7       -       249       95.3       -         ProjE (Shi and Weninger, 2017)       34       88.4       -       -       -       -         RTransE (García-Durán et al., 2015)       50       76.2       -       -       -       -         PTransE-ADD (Lin et al., 2015a)       58       84.6       -       -       -       -         PTransE-RNN (Lin et al., 2015a)       92       82.2       -       -       -       -         GAKE (Feng et al., 2016b)       119       64.8       -       -       -       -         Gaifman (Niepert, 2016)       75       84.2       -       352       93.9       -         Hiri (Liu et al., 2016)       -       70.3       0.603       -       90.8       0.691         Neural LP (Yang et al., 2017)       -       83.7       0.76       -       94.5       0.94         R-GCN+ (Schlichtkrull et al., 2018)       -       84.2       0.696       -       96.4       0.819         KB <sub>LRN</sub> (Durán and Niepert, 2018)       44       87.5       0.794	QuatE (Zhang et al., 2019)	17	90.0	0.782	162	<u>95.9</u>	0.950	
IRN (Shen et al., 2017)       38       92.7       -       249       95.3       -         ProjE (Shi and Weninger, 2017)       34       88.4       -       -       -       -         RTransE (García-Durán et al., 2015)       50       76.2       -       -       -       -         PTransE-ADD (Lin et al., 2015a)       58       84.6       -       -       -       -         PTransE-RNN (Lin et al., 2015a)       92       82.2       -       -       -       -         GAKE (Feng et al., 2016b)       119       64.8       -       -       -       -         Gaifman (Niepert, 2016)       75       84.2       -       352       93.9       -         Hiri (Liu et al., 2016)       -       70.3       0.603       -       90.8       0.691         Neural LP (Yang et al., 2017)       -       83.7       0.76       -       94.5       0.94         R-GCN+ (Schlichtkrull et al., 2018)       -       84.2       0.696       -       96.4       0.819         KB <sub>LRN</sub> (Durán and Niepert, 2018)       44       87.5       0.794       -       -       -         TEKE-H (Wang and Li, 2016)       108       73.0       -       114 </td <td colspan="2">ComplEx-N3 (Lacroix et al., 2018)</td> <td><u>91</u></td> <td>0.86</td> <td>-</td> <td>96</td> <td>0.95</td>	ComplEx-N3 (Lacroix et al., 2018)		<u>91</u>	0.86	-	96	0.95	
ProjE (Shi and Weninger, 2017)         34         88.4         -         -         -         -           RTransE (García-Durán et al., 2015)         50         76.2         -         -         -         -           PTransE-ADD (Lin et al., 2015a)         58         84.6         -         -         -         -           PTransE-RNN (Lin et al., 2015a)         92         82.2         -         -         -         -           GAKE (Feng et al., 2016b)         119         64.8         -         -         -         -           Gaifman (Niepert, 2016)         75         84.2         -         352         93.9         -           Hiri (Liu et al., 2016)         -         70.3         0.603         -         90.8         0.691           Neural LP (Yang et al., 2017)         -         83.7         0.76         -         94.5         0.94           R-GCN+ (Schlichtkrull et al., 2018)         -         84.2         0.696         -         96.4         0.819           KB <sub>LRN</sub> (Durán and Niepert, 2018)         44         87.5         0.794         -         -         -           TEKE H (Wang and Li, 2016)         108         73.0         -         114         92.9	TuckER (Balazevic et al., 2019)	-	89.2	0.795	-	95.8	0.953	
RTransE (García-Durán et al., 2015) 50 76.2	IRN (Shen et al., 2017)	38	92.7	-	249	95.3	-	
PTransE-ADD (Lin et al., 2015a)       58       84.6       -       -       -       -         PTransE-RNN (Lin et al., 2015a)       92       82.2       -       -       -       -         GAKE (Feng et al., 2016b)       119       64.8       -       -       -       -         Gaifman (Niepert, 2016)       75       84.2       -       352       93.9       -         Hiri (Liu et al., 2016)       -       70.3       0.603       -       90.8       0.691         Neural LP (Yang et al., 2017)       -       83.7       0.76       -       94.5       0.94         R-GCN+ (Schlichtkrull et al., 2018)       -       84.2       0.696       -       96.4       0.819         KB <sub>LRN</sub> (Durán and Niepert, 2018)       44       87.5       0.794       -       -       -         TEKE-H (Wang and Li, 2016)       108       73.0       -       114       92.9       -	ProjE (Shi and Weninger, 2017)	<u>34</u>	88.4	-	-	-	-	
PTransE-RNN (Lin et al., 2015a)       92       82.2       -       -       -       -         GAKE (Feng et al., 2016b)       119       64.8       -       -       -       -         Gaifman (Niepert, 2016)       75       84.2       -       352       93.9       -         Hiri (Liu et al., 2016)       -       70.3       0.603       -       90.8       0.691         Neural LP (Yang et al., 2017)       -       83.7       0.76       -       94.5 <b>0.94</b> R-GCN+ (Schlichtkrull et al., 2018)       -       84.2       0.696       - <b>96.4</b> 0.819         KB <sub>LRN</sub> (Durán and Niepert, 2018) <b>44 87.5 0.794</b> -       -       -         TEKE-H (Wang and Li, 2016)       108       73.0       - <b>114</b> 92.9       -	RTransE (García-Durán et al., 2015)	50	76.2	-	-	-	-	
GAKE (Feng et al., 2016b)       119       64.8       -       -       -       -         Gaifman (Niepert, 2016)       75       84.2       -       352       93.9       -         Hiri (Liu et al., 2016)       -       70.3       0.603       -       90.8       0.691         Neural LP (Yang et al., 2017)       -       83.7       0.76       -       94.5 <b>0.94</b> R-GCN+ (Schlichtkrull et al., 2018)       -       84.2       0.696       - <b>96.4</b> 0.819         KB <sub>LRN</sub> (Durán and Niepert, 2018) <b>44 87.5 0.794</b> -       -       -         TEKE H (Wang and Li, 2016)       108       73.0       - <b>114</b> 92.9       -	PTransE-ADD (Lin et al., 2015a)	58	84.6	-	-	-	-	
Gaifman (Niepert, 2016)       75       84.2       -       352       93.9       -         Hiri (Liu et al., 2016)       -       70.3       0.603       -       90.8       0.691         Neural LP (Yang et al., 2017)       -       83.7       0.76       -       94.5 <b>0.94</b> R-GCN+ (Schlichtkrull et al., 2018)       -       84.2       0.696       - <b>96.4</b> 0.819         KB <sub>LRN</sub> (Durán and Niepert, 2018) <b>44 87.5 0.794</b> -       -       -         TEKE H (Wang and Li, 2016)       108       73.0       - <b>114</b> 92.9       -	PTransE-RNN (Lin et al., 2015a)	92	82.2	-	-	-	-	
Hiri (Liu et al., 2016) - 70.3 0.603 - 90.8 0.691 Neural LP (Yang et al., 2017) - 83.7 0.76 - 94.5 <b>0.94</b> R-GCN+ (Schlichtkrull et al., 2018) - 84.2 0.696 - <b>96.4</b> 0.819 KB <sub>LRN</sub> (Durán and Niepert, 2018) <b>44 87.5 0.794</b> TEKE_H (Wang and Li, 2016) 108 73.0 - <b>114</b> 92.9 -	GAKE (Feng et al., 2016b)	119	64.8	-	-	-	-	
Neural LP (Yang et al., 2017)       -       83.7       0.76       -       94.5 <b>0.94</b> R-GCN+ (Schlichtkrull et al., 2018)       -       84.2       0.696       - <b>96.4</b> 0.819         KB <sub>LRN</sub> (Durán and Niepert, 2018) <b>44 87.5 0.794</b> -       -       -         TEKE-H (Wang and Li, 2016)       108       73.0       - <b>114</b> 92.9       -	Gaifman (Niepert, 2016)		84.2	-	352	93.9	-	
Neural LP (Yang et al., 2017)       -       83.7       0.76       -       94.5 <b>0.94</b> R-GCN+ (Schlichtkrull et al., 2018)       -       84.2       0.696       - <b>96.4</b> 0.819         KB <sub>LRN</sub> (Durán and Niepert, 2018) <b>44 87.5 0.794</b> -       -       -         TEKE-H (Wang and Li, 2016)       108       73.0       - <b>114</b> 92.9       -	Hiri (Liu et al., 2016)		70.3	0.603	-	90.8	0.691	
R-GCN+ (Schlichtkrull et al., 2018)       -       84.2       0.696       -       96.4       0.819         KB <sub>LRN</sub> (Durán and Niepert, 2018)       44       87.5       0.794       -       -       -         TEKE_H (Wang and Li, 2016)       108       73.0       -       114       92.9       -	Neural LP (Yang et al., 2017)		83.7	0.76	-	94.5	0.94	
TEKE_H (Wang and Li, 2016) 108 73.0 - <b>114</b> 92.9 -	R-GCN+ (Schlichtkrull et al., 2018)		84.2	0.696	-	96.4	0.819	
	KB <sub>LRN</sub> (Durán and Niepert, 2018)		87.5	0.794	-	-	-	
	TEKE_H (Wang and Li, 2016)	108	73.0	-	114	92.9	-	
		82	79.0	-	156	93.2	-	

Table 3: Entity prediction results on WN18 and FB15k. **MR** and @10 denote evaluation metrics of mean rank and Hits@10 (in %), respectively. TransG's results are taken from its latest ArXiv version (https://arxiv.org/abs/1509.05488v7). NTN's results are taken from Yang *et al.* (2015).  $[\heartsuit]$ , [\$\infty]\$ and [\$\infty]\$ denote results taking from Nickel *et al.* (2016b), Ravishankar *et al.* (2017) and Kadlec *et al.* (2017), respectively.

of models not exploiting path or external information, the complex vector-based models (e.g. QuatE, CompleEx-N3 and RotatE) produce the strongest evaluation scores, followed by the neural network-based models (e.g. CapsE, InteractE and HypER).<sup>5</sup> Tables 3 and 4 also show that TransE and DIST-MULT, despite of theirs simplicity, can produce very competitive results (by performing a careful grid search of hyper-parameters).

<sup>&</sup>lt;sup>5</sup>CapsE uses the pre-trained word embeddings for entity vector initialization on WN18RR. It is not surprising that CapsE produces the best MR on WN18RR as many entity names in WordNet are lexically meaningful. It is possible for all other embedding models to utilize the pre-trained word vectors as well. However, averaging the pre-trained word embeddings for initializing entity vectors is an open problem, and it is not always useful since entity names in many domain-specific KBs are not lexically meaningful (Guu et al., 2015).

	Filtered						
Method		FB15k-237			WN18RR		
	MR	@10	MRR	MR	@10	MRR	
IRN (Shen et al., 2017)	211	46.4	-	-	-	-	
KBGAN (Cai and Wang, 2018)	-	45.8	0.278	-	48.1	0.213	
DISTMULT (Yang et al., 2015) [ $\Diamond$ ]	254	41.9	0.241	5110	49	0.43	
ComplEx (Trouillon et al., 2016) [♦]	339	42.8	0.247	5261	51	0.44	
ConvE (Dettmers et al., 2018)	246	49.1	0.316	5277	48	0.46	
ER-MLP (Dong et al., 2014) [♠]		54.0	0.342	4798	41.9	0.366	
HypER (Balažević et al., 2019)		52.0	0.341	5798	52.2	0.465	
TransE (Bordes et al., 2013) [♥]		46.5	0.294	<u>743</u>	56.0	0.245	
ConvKB (Nguyen et al., 2018) [♥]		53.2	0.418	763	56.7	0.253	
CapsE (Nguyen et al., 2019)		59.3	0.523	719	56.0	0.415	
InteractE (Vashishth et al., 2020)		53.5	0.354	5202	52.8	0.463	
RotatE (Sun et al., 2019)		53.3	0.338	3340	<u>57.1</u>	0.476	
QuatE (Zhang et al., 2019)		55.0	0.348	2314	58.2	0.488	
ComplEx-N3 (Lacroix et al., 2018)	-	<u>56</u>	0.37	-	57	0.48	
Conv-TransE (Shang et al., 2019)		51	0.33	-	52	0.46	
TuckER (Balazevic et al., 2019)	-	54.4	0.358	-	52.6	0.470	
Neural LP (Yang et al., 2017)	-	36.2	0.24	-	-	_	
R-GCN+ (Schlichtkrull et al., 2018)		41.7	0.249	-	-	-	
KB <sub>LRN</sub> (Durán and Niepert, 2018)		49.3	0.309	-	-	-	
KBAT (Nathani et al., 2019)		62.6	0.518	1940	58.1	0.440	
SACN (Shang et al., 2019)		54	0.35	-	54	0.47	

Table 4: Entity prediction results on WN18RR and FB15k-237.  $[\diamondsuit]$ ,  $[\diamondsuit]$ , and  $[\heartsuit]$  denote results taking from Dettmers *et al.* (2018), Ravishankar *et al.* (2017) and Nguyen *et al.* (2019), respectively.

#### 5 Discussion and Conclusion

The reasons why much work has been devoted towards developing triple-based models are: (1) additional information sources might not be available, e.g., for KBs for specialized domains, (2) models that do not exploit path information or external resources are simpler and thus typically much faster to train than the more complex models using path or external information, and (3) the more complex models that exploit path or external information are typically extensions of these simpler models, and are often initialized with parameters estimated by such simpler models, so improvements to the simpler models should yield corresponding improvements to the more complex models as well (Nguyen et al., 2016a).

It is worth to further explore those KB completion embedding models for a new application where we could formulate its corresponding data into triples. For example, in Web search engines, we observe useroriented relationships between submitted queries and documents returned by the search engines. That is, we have triple representations (query, user, document) in which for each user-oriented relationship, we would have many queries and documents, resulting in a lot of Many-to-Many relationships. Inspired by this observation, Vu et al. (2017) applied STransE (Nguyen et al., 2016a) for a search personalization task to re-rank the search documents returned by a search engine for users' submitted queries. Other application examples can be also found for recommender systems (Zhang et al., 2016; He et al., 2017; Cao et al., 2019), social relation extraction (Tu et al., 2017) and visual relation detection (Zhang et al., 2017). Future directions might also include: (i) Combining logical rules which contain rich background information and KB triples in a unified KB completion framework, e.g. jointly embedding KBs and logical rules (Guo et al., 2016; Yang et al., 2017). (ii) Recent embedding models for KB completion hold a closed-world assumption where the KBs are fixed (i.e. new entities might not be added easily), thus it would be worth exploring open-world KB completion models to connect unseen entities to the KBs (Shi and Weninger, 2018). (iii) Investigate efficient methods which can be applied to large-scale KBs of millions of entities and relations (Zhang et al., 2020).

In this paper, we have presented a comprehensive overview of embedding models of entity and relationships for knowledge base completion. The paper also provides update-to-date experimental results of the embedding models on the entity prediction (i.e. link prediction) task on benchmark datasets FB15k, WN18, FB15k-237 and WN18RR.

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