

ASPEN: ASP-Based System for Collective Entity Resolution

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Abstract

In this paper, we present ASPEN, an answer set programming (ASP) implementation of a recently proposed declarative framework for collective entity resolution (ER). While an ASP encoding had been previously suggested, several practical issues had been neglected, most notably, the question of how to efficiently compute the (externally defined) similarity facts that are used in rule bodies. This leads us to propose new variants of the encodings (including Datalog approximations) and show how to employ different functionalities of ASP solvers to compute (maximal) solutions, and (approximations of) the sets of possible and certain merges. A comprehensive experimental evaluation of ASPEN on real-world datasets shows that the approach is promising, achieving high accuracy in real-life ER scenarios. Our experiments also yield useful insights into the relative merits of different types of (approximate) ER solutions, the impact of recursion, and factors influencing performance.

1 Introduction

Entity resolution (ER) is a fundamental problem in data quality which aims at identifying different constants (of the same type) that refer to the same real-world entity (Singla and Domingos 2006; Singla and Domingos 2006). Over time, several variants of ER (also known as record linkage or deduplication) have been investigated, including pairwise matching in a single table and (the more general) *collective* ER, which looks at the joint resolution (match, merge) of entity references across multiple tables (Bhattacharya and Getoor 2007). Given the multi-faceted nature of the ER problem, diverse techniques have been already proposed to tackle it (Christophides et al. 2021), including machine learning (ML), and declarative frameworks based upon logical rules and constraints. Most existing approaches to ER focus on single-pass matching of tuples within a single table or between a pair of tables, and ML methods have obtained remarkable results (Li et al. 2020) for such settings. On the other hand, declarative methods are well suited for complex multi-relational settings as they naturally exploit the relational dependencies to perform collective ER. Moreover, some declarative approaches conduct ER in a *recursive* manner (also called *deep* ER (Deng et al. 2022)), instead of examining entity pairs only once.

LACE is a recently proposed declarative framework (Bienvenu, Cima, and Gutíerrez-Basulto 2022) for collective ER, which employs hard and soft rules to define mandatory and possible merges, and denial constraints (Bertossi 2011) to enforce consistency of the resulting database and constrain the allowed combinations of merges. LACE employs a dynamic semantics in which rule bodies are evaluated over the current induced database, taking into account all previously derived merges. This makes it possible to support recursive scenarios, while ensuring that all merges have a (non-circular) derivation. The semantics of LACE is also global since all occurrences of the matched constants are merged, rather than only those constant occurrences used in deriving the match. Additionally, LACE considers a space of maximal (w.r.t. set inclusion) solutions, which emerges from adopting denial constraints to enforce consistency and restricting which merges can be performed together, effectively creating choices. From this, one can define the notions of possible and certain merges, as those merges that belong to some, respectively all, maximal solutions.

While the theoretical foundations of LACE have been already established, a real-life implementation is to-date absent. Bienvenu, Cima, and Gutíerrez-Basulto (2022) showed that LACE solutions can be faithfully captured by answer set programming (ASP) (Lifschitz 2019; Gebser et al. 2012) stable models. Building upon this, in this paper we present ASPEN, an ASP-based system for collective ER. ASPEN deals with several practical issues, including the question of how to efficiently compute the (externally defined) similarity facts that are used in rule bodies. It also implements new variants of the encodings (including Datalog approximations) and uses different functionalities of ASP solvers to compute (maximal) solutions, and (approximations of) the sets of possible and certain merges. A comprehensive experimental evaluation of ASPEN on real-world datasets shows that the approach is promising, achieving high accuracy in real-life ER scenarios. Our experiments also yield useful insights into the relative merits of different types of (approximate) ER solutions, the impact of recursion, and factors influencing performance, such as the degree of dirtiness and size of datasets. ASPEN also leverages the xclingo (Cabalar and Muñiz 2023) framework for explaining conclusions of

ASP programs to compute the justification of a merge in a solution, making ASPEN a *justifiable* framework for ER.

Related Work We discuss prior work on logic-based approaches to ER; for details on ML and probabilistic methods, see (Christophides et al. 2021). The LACE framework, underlying ASPEN, shares some characteristics with other logic-based ER methods. Similar to approaches based on matching dependencies (MDs) (Bertossi, Kolahi, and Lakshmanan 2013; Deng et al. 2022), ASPEN adopts a dynamic semantics, enabling recursive ER. Like the Datalog-like approaches Dedupalog (Arasu, Ré, and Suciu 2009) and Entity Linking (EL) (Burdick et al. 2016) (and unlike MDs), ASPEN considers a global semantics. Finally, as in the EL framework, ASPEN does not consider only a single solution, but rather a space of maximal solutions, leading to notions of possible and certain merges. While ASP encodings have been proposed for MDs (Bertossi, Kolahi, and Lakshmanan 2013; Bahmani and Bertossi 2017), no implementation nor evaluation are provided. Likewise, there is no implementation of the EL framework. The closest existing implementations of logic-based ER are those of Dedupalog (Arasu, Ré, and Suciu 2009) and MRL (Deng et al. 2022). However, they are not publicly available. The main distinguishing features of ASPEN compared to existing systems are as follows: **i)** Rather than computing a single (possibly non-optimal) solution, ASPEN is not only able to compute, with the guarantee of correctness, a space of maximal solutions but also approximations with different levels of granularity based on different reasoning modes. **ii)** To the best of our knowledge, ASPEN is the first system that is able to explicitly give justifications to merges. We finally note in passing that ASP-based approaches to database repair have been explored (Eiter et al. 2008; Manna, Ricca, and Terracina 2013; Ahmetaj et al. 2022). Additionally, a bespoke ASP system for cleaning healthcare data has also been developed (Terracina, Martello, and Leone 2013).

All programs, code, experiments and data of ASPEN are available at <https://github.com/zi-xiang/Aspen>. Further details on proofs and experiments can be found in the appendix of (Xiang et al. 2024).

2 Preliminaries

We assume infinite sets of *constants* \mathbf{C} and *variables* \mathbf{V} . A (*database*) *schema* \mathcal{S} consist of a finite set of relation symbols, each having an associated arity $k \in \mathbb{N}$. We write $R/k \in \mathcal{S}$ to indicate that R has arity k . A *relational atom* (over schema \mathcal{S}) takes the form $R(t_1, \dots, t_k)$ where $R/k \in \mathcal{S}$ and $t_i \in \mathbf{C} \cup \mathbf{V}$ for $1 \leq i \leq k$, and we call $R(t_1, \dots, t_k)$ a *fact* if $\{t_1, \dots, t_k\} \subseteq \mathbf{C}$. We say that t_i *occurs in position* i of an atom $R(t_1, \dots, t_k)$. A *database* D (over schema \mathcal{S}) is a finite set of facts (over \mathcal{S}). The set of constants occurring in a database D is denoted by $\text{Dom}(D)$. Sometimes it will prove more natural to employ *attributes* rather than (unnamed) positions, writing $R(A_1, \dots, A_k)$ to indicate that (A_1, \dots, A_k) are the attributes of R .

A *conjunctive query* (CQ) over a schema \mathcal{S} takes the form $q(\vec{x}) = \exists \vec{y}. \varphi(\vec{x}, \vec{y})$, where \vec{x} and \vec{y} are disjoint tuples of distinguished and quantified variables, and $\varphi(\vec{x}, \vec{y})$

is a conjunction of relational atoms over \mathcal{S} , with variables drawn from $\vec{x} \cup \vec{y}$. We use $\text{terms}(q)$ for the set of *terms* of q , i.e. the constants and variables appearing in q . The set of *answers* to a CQ $q(\vec{x}) = \exists \vec{y}. \varphi(\vec{x}, \vec{y})$ on a database D (over the same schema), denoted $q(D)$, contains those tuples of constants \vec{c} such that there exists a mapping $h : \text{terms}(q) \rightarrow \text{Dom}(D)$ such that (i) $h(\vec{x}) = \vec{c}$, (ii) $h(t) = t$ for $t \in \text{terms}(q) \cap \mathbf{C}$, and (iii) for every atom $R(t_1, \dots, t_k)$ of q , $R(h(t_1), \dots, h(t_k)) \in D$. We will also consider CQs *with inequalities* (CQ $^\neq$), which may additionally include inequality atoms $t_i \neq t_j$, in which case we require that the mapping h further satisfies (iv) $h(t_i) \neq h(t_j)$ whenever q contains $t_i \neq t_j$. When q has only quantified variables, it is called *Boolean*, and we say that a Boolean CQ $^\neq$ q is *satisfied* in D if $q(D) = \{\emptyset\}$. A *denial constraint* (DC) takes the form $q \rightarrow \perp$, where q is a Boolean CQ $^\neq$. We say that D *satisfies a DC* $q \rightarrow \perp$ just in the case that the CQ $^\neq$ q is not satisfied in D . Functional dependencies (FDs) and primary key constraints are special cases of DCs.

When we speak of complexity, we will always mean *data complexity*, which is measured only in terms of the size of the input database, with all other inputs (e.g. ER specifications and queries) treated as fixed.

3 Entity Resolution Framework

In this section, we recall the syntax and semantics of LACE (Bienvenu, Cima, and Gutiérrez-Basulto 2022). We also introduce new notions that are useful for our system.

3.1 LACE Entity Resolution Specifications

Entity resolution can be formulated as the task of discovering pairs of syntactically distinct database constants that refer to the same entity (we will often use the term *merges* for such pairs). We adopt the LACE framework, which focuses on identifying merges of entity-referencing constants (e.g. paper or author identifiers).

LACE employs hard and soft rules to identify mandatory and possible merges. *Hard rules* and *soft rules*, over schema \mathcal{S} , take respectively the forms:

$$q(x, y) \Rightarrow \text{Eq}(x, y), \quad q(x, y) \dashrightarrow \text{Eq}(x, y) \quad (1)$$

where Eq is a special relation symbol not in \mathcal{S} used to store pairs of merged constants, and $q(x, y)$ is a CQ using relation symbols from \mathcal{S} . Intuitively, a hard (resp. soft) rule states that a pair of constants (c_1, c_2) being an answer to q provides sufficient (resp. reasonable) evidence that c_1 and c_2 refer to the same real-world entity. We use $q(x, y) \rightarrow \text{Eq}(x, y) \in \Gamma$ to denote a generic (hard or soft) rule (note the arrow).

It is natural and useful for rule bodies to use *similarity relations*, i.e. binary relations whose extension is fixed and computed using some external function (e.g. by applying a string similarity measure to a pair of constants and keeping those pairs whose score exceeds a given threshold). We shall thus allow the considered schema to contain such similarity relations and will adopt the more intuitive infix notation $x \approx y$ (possibly with indices) for similarity atoms. Note also that in the present section, we will assume that the similarity facts are provided as part of the input database, leaving the question of how to best compute them to later sections.

Band(<i>bid</i> , <i>name</i> , <i>genre</i> , <i>year</i> , <i>founder</i>)				
<i>bid</i>	<i>name</i>	<i>genre</i>	<i>year</i>	<i>founder</i>
b_1	Pink Floyd	Psy. rock	1965	Barrett
b_2	The Pink Floyd	Prog. rock	1965	Barrett

Song(<i>sid</i> , <i>title</i> , <i>lyricist</i> , <i>bid</i>)			
<i>sid</i>	<i>title</i>	<i>lyricist</i>	<i>bid</i>
s_1	Shine On You Crazy Diamond (I-IV)	Waters	b_1
s_2	Shine On You Crazy Diamond	Waters	b_2
s_3	Shine On You Crazy Diamond (V-IX)	Waters	b_1

$\sigma_{\text{ex}}: \text{Song}(x, t, l, b) \wedge \text{Song}(y, t', l, b) \wedge t \approx t' \dashrightarrow \text{Eq}(x, y)$
 $\rho_{\text{ex}}: \text{Band}(x, n, g, d, f) \wedge \text{Band}(y, n', g', d, f) \wedge n \approx n' \wedge g \approx g' \Rightarrow \text{Eq}(x, y)$
 $\delta_{\text{ex}}: \text{Appear}(s, a, i) \wedge \text{Appear}(s, a, j) \wedge i \neq j \rightarrow \perp$

Appear(<i>sid</i> , <i>album</i> , <i>position</i>)		
<i>sid</i>	<i>album</i>	<i>position</i>
s_1	Wish You Were Here	1
s_2	A Delicate Sound of Thunder	1
s_3	Wish You Were Here	5

We assume that the extension of the similarity relation \approx (restricted to $\text{Dom}(D_{\text{ex}})$) is the reflexive and symmetric closure of $\{(n_1, n_2), (g_1, g_2), (t_1, t_2), (t_2, t_3)\}$, where n_i, g_i , and t_i are the name and genre of band b_i and the title of song s_i , respectively.

Figure 1: A schema \mathcal{S}_{ex} , an \mathcal{S}_{ex} -database D_{ex} , and an ER specification $\Sigma_{\text{ex}} = \langle \Gamma_{\text{ex}}, \Delta_{\text{ex}} \rangle$ for \mathcal{S}_{ex} , where $\Gamma_{\text{ex}} = \{\rho_{\text{ex}}, \sigma_{\text{ex}}\}$ and $\Delta_{\text{ex}} = \{\delta_{\text{ex}}\}$.

Example 1. Figure 1 presents our running example inspired by the MUSIC dataset. Relations *Group*, *Song*, *Appear* of the schema \mathcal{S}_{ex} provide information about music bands, songs, and which songs appear on which albums. The attributes **bid** and **sid** contain identifiers for bands and songs respectively, and the task at hand is to identify which of these identifiers refer to the same band / song. The hard rule ρ_{ex} says that if two band ids have the same founding year, same founder, similar names, and similar genres, they must refer to the same band. The soft rule σ_{ex} states that two song ids likely refer to the same song if they have the same band and lyricist and similar titles. For simplicity, a single similarity relation \approx is used to say which strings count as similar.

To enforce consistency of the inferred merges and to help block false positives, LACE specifications may also include *denial constraints*. For example, the denial constraint δ_{ex} in Figure 1 is an FD for *Appear*, which forbids the occurrence of the same song in different positions within an album.

A LACE *entity resolution (ER) specification* over schema \mathcal{S} takes the form $\Sigma = \langle \Gamma, \Delta \rangle$, where $\Gamma = \Gamma_h \cup \Gamma_s$ is a finite set of hard and soft rules and Δ is a finite set of denial constraints, all over \mathcal{S} . Rulesets are required to satisfy a *sim-safety condition*, whereby the relation positions involved in merges must be distinct from those involved in similarity atoms. The example specification Σ_{ex} is sim-safe as the attributes involved in similarity atoms (**title**, **name**, **genre**) are different from those involved in merges (**bid**, **sid**). Thanks to sim-safety, we may assume w.l.o.g. that the constants appearing in merge positions do not occur in sim positions.

3.2 Semantics of LACE

The semantics of LACE associates with every database D and ER specification Σ a set of solutions, where each solution takes the form of an equivalence relation over $\text{Dom}(D)$, indicating which constants refer to the same entity. Given a set of pairs $P \subseteq S \times S$, we write $\text{EqRel}(P, S)$ to denote the least equivalence relation over S that extends P .

In a nutshell, ER solutions in LACE are obtained by applying the hard and soft rules in such a manner that all hard rules and constraints are satisfied, with the inferred Eq-facts determining the equivalence relation. Importantly, the evaluation

of LACE rules takes into account previously derived merges, making it possible for merges to trigger further merges. Formally, given a database D , and equivalence relation E over $\text{Dom}(D)$, the *database induced by D and E* , denoted D_E , is obtained from D by replacing each constant c by the (uniformly chosen) representative \bar{c} of its equivalence class. The set $q(D, E)$ of *answers to $q(\bar{x})$ w.r.t. D and E* is defined as:

$$q(D, E) = \{(c_1, \dots, c_n) \mid (\bar{c}_1, \dots, \bar{c}_n) \in q(D_E)\}$$

Rule $q(x, y) \rightarrow \text{Eq}(x, y)$ is satisfied in (D, E) if $q(D, E) \subseteq E$, and DC δ is satisfied in (D, E) if δ is satisfied in D_E .

With these notions in hand, we can now define solutions. Given an ER specification Σ and database D , we call E a *candidate solution for (D, Σ)* if one of the following holds:

1. $E = \text{EqRel}(\emptyset, \text{Dom}(D))$
2. $E = \text{EqRel}(E' \cup \{\alpha\}, D)$, where E' is a candidate solution for (D, Σ) and $\alpha \in q(D, E) \setminus E'$ for some $q(x, y) \rightarrow \text{Eq}(x, y) \in \Gamma$

and it is a *solution for (D, Σ)* if additionally (i) $(D, E) \models \Gamma_h$ and (ii) $(D, E) \models \Delta$. We denote by $\text{Sol}(D, \Sigma)$ the set of solutions for (D, Σ) , and let $\text{MaxSol}(D, \Sigma)$ be the set of *maximal solutions*, i.e. solutions E such that there is no solution E' for (D, Σ) with $E \subsetneq E'$.

Example 2. We determine the maximal solutions for $(D_{\text{ex}}, \Sigma_{\text{ex}})$. The initial trivial equivalence relation $E = \text{EqRel}(\emptyset, \text{Dom}(D_{\text{ex}}))$ is not a solution as the hard rule ρ_{ex} requires us to merge b_1 and b_2 . Applying ρ_{ex} , we obtain $E' = \text{EqRel}(\{(b_1, b_2)\}, \text{Dom}(D_{\text{ex}}))$, which is a solution for $(D_{\text{ex}}, \Sigma_{\text{ex}})$ but not a maximal one. Indeed, due to the addition of (b_1, b_2) , both (s_1, s_2) and (s_2, s_3) can now be obtained by applying the soft rule σ_{ex} . Notice, however, that it is not possible to include both of them, as transitivity would force us to include (s_1, s_3) , leading to a violation of δ_{ex} . We thus obtain two maximal solutions for $(D_{\text{ex}}, \Sigma_{\text{ex}})$, namely $E_1 = \text{EqRel}(\{(b_1, b_2), (s_1, s_2)\}, \text{Dom}(D))$ and $E_2 = \text{EqRel}(\{(b_1, b_2), (s_2, s_3)\}, \text{Dom}(D))$.

There can be zero, one, or multiple (maximal) solutions. However, for specifications that do not contain soft rules, there can be at most one solution, and for specifications that do not contain constraints, there is always a single solution.

3.3 Summarizing and Explaining Solutions

When there are multiples solutions, it is useful to be able to summarize them by identifying merges that occur in all or some maximal solutions. Formally, we say that a merge α is *certain* if $\alpha \in E$ for every $E \in \text{MaxSol}(D, \Sigma)$, and it is *possible* if $\alpha \in E$ for some $E \in \text{MaxSol}(D, \Sigma)$ (equivalently, some $E \in \text{Sol}(D, \Sigma)$). We use $\text{CM}(D, \Sigma)$ and $\text{PM}(D, \Sigma)$ for the sets of certain and possible merges.

While certain and possible merges provide natural summarizations, they are unfortunately hard to compute:

Theorem 1. (Bienvenu, Cima, and Gutiérrez-Basulto 2022) *It is NP-complete (resp. Π_2^P -complete) in data complexity to decide if $\alpha \in \text{PM}(D, \Sigma)$ (resp. $\text{CM}(D, \Sigma)$).*

For this reason, it will prove useful to define efficiently computable approximations. To this end, we consider two ways of simplifying a specification $\Sigma = \langle \Gamma_h \cup \Gamma_s, \Delta \rangle$:

- $\Sigma^{\text{lb}} = \langle \Gamma_h, \emptyset \rangle$, i.e. remove soft rules and constraints
- $\Sigma^{\text{ub}} = \langle \Gamma_h \cup \Gamma_{s \rightarrow h}, \emptyset \rangle$, i.e. drop constraints and replace soft rules by the corresponding hard rules ($\Gamma_{s \rightarrow h}$)

As Σ^{lb} and Σ^{ub} do not contain any constraints, they will always yield a unique solution. We can therefore define

- $\text{LB}(D, \Sigma)$ as the unique solution to (D, Σ^{lb})
- $\text{UB}(D, \Sigma)$ as the unique solution to (D, Σ^{ub})

We summarize the properties of the different merge sets:

Theorem 2. *The sets $\text{LB}(D, \Sigma)$ and $\text{UB}(D, \Sigma)$ can be computed in polynomial w.r.t. data complexity. Moreover, if $\text{Sol}(D, \Sigma) \neq \emptyset$, then*

$$\text{LB}(D, \Sigma) \subseteq \text{CM}(D, \Sigma) \subseteq \text{PM}(D, \Sigma) \subseteq \text{UB}(D, \Sigma)$$

and $\text{CM}(D, \Sigma) \subseteq M \subseteq \text{PM}(D, \Sigma)$ for $M \in \text{MaxSol}(D, \Sigma)$.

Example 3. In our toy example, $\text{LB}(D_{\text{ex}}, \Sigma_{\text{ex}})$ and $\text{CM}(D_{\text{ex}}, \Sigma_{\text{ex}})$ coincide, and the only non-trivial pair they contain is (b_1, b_2) . The set $\text{PM}(D_{\text{ex}}, \Sigma_{\text{ex}})$ further contains merges (s_1, s_2) and (s_2, s_3) , whilst $\text{UB}(D_{\text{ex}}, \Sigma_{\text{ex}})$ also contains (s_1, s_3) (not present in any solution).

We will employ *proof trees* to explain why a merge appears in a solution (we provide here some intuitions behind proof trees and refer to App. A for more details). Informally, a proof tree for a merge α in a solution $E \in \text{Sol}(D, \Sigma)$ is a node-labelled tree such that (a) the root node has label α , (b) every leaf node is labelled with a fact from D , and (c) every non-leaf node n is labelled with a pair of constants (d, e) corresponding to a single transitive step or a rule application.

To quantify the number of successive rule applications needed to obtain a merge, we let the *rule-depth of a proof tree* T be the maximum number of rule nodes in any leaf-to-root path in T . The *level of a merge α in a solution E* is 0 if $\alpha = (c, c)$ for some $c \in \mathbf{C}$, and otherwise is the minimum rule-depth of all proof trees of α in E .

4 ASP Encoding and Algorithms

The ASPEN system implements the LACE framework by encoding ER specifications as ASP programs and making calls to ASP solvers to generate and reason about ER solutions.

After recalling some ASP notions, we present the ASP encoding and algorithms employed by ASPEN. We also explore some practical issues that were ignored in the theoretical treatment of LACE, most importantly, the question of how to handle similarity atoms.

We assume familiarity with ASP basics, see (Brewka, Eiter, and Truszczyński 2011; Gebser et al. 2012; Lifschitz 2019) for more details. For our purposes, it is enough to consider *normal rules* and *constraints*. That is, respectively, rules with a single atom head and rules with an empty head. We shall use Π to denote an ASP *program* (a set of rules). Given a program Π , we use $gr(\Pi)$ to denote the set of all ground instantiations rules from Π with constants occurring in Π , and $SM(\Pi)$ to denote the set of all stable models of Π . Determining whether a program has a stable model is the fundamental decision problem in ASP, which is solved using ASP solvers (Gebser et al. 2018). However, more reasoning modes are needed to cover problems encountered in practice. Most modern ASP solvers are also able to *enumerate* (n elements of) $SM(\Pi)$; *project* answers w.r.t. a given set of atoms, and enumerating (n elements of) those projections; computing the intersection (resp. union) of all stable models of Π (*cautious*, resp. *brave* reasoning); and perform *optimizations* by computing some (or enumerating n) elements of $SM(\Pi)$ that minimize a given objective function.

4.1 ASP Encoding of Solutions

Given a LACE specification Σ and a database D , we define an ASP program $\Pi_{(D, \Sigma)}$ containing all the facts in D , and an ASP rule for each (hard or soft) rule in Σ . Consider, for example, the specification Σ_{ex} in Figure 1. Rules ρ_{ex} , σ_{ex} and δ_{ex} are translated as follows:

$$\begin{aligned} \text{eq}(X, Y) \leftarrow & \text{band}(X, N, G, D, F), \text{band}(Y, N', G', D, F), \\ & \text{sim}(G, G', S), S \geq 95, \text{sim}(N, N', S'), S' \geq 95. \end{aligned}$$

$$\begin{aligned} \{\text{eq}(X, Y)\} \leftarrow & \text{song}(X, T, L, B), \text{song}(Y, T', L, B'), \\ & \text{not empty}(L), \text{sim}(T, T', S), S \geq 95, \text{eq}(B, B'). \end{aligned}$$

$$\perp \leftarrow \text{appear}(S, A, I), \text{appear}(S', A, J), \text{eq}(S, S'), I \neq J.$$

Roughly, each relational atom in the body of a (hard or soft rule) is translated into an atom in the body of an ASP rule. The relation eq is used to store mandatory merges (and ultimately solutions to (D, Σ)). Atoms of the form $\text{eq}(X, X')$ in the rule bodies are used to encode that instantiations of X and X' have been determined to denote the same entity. $\Pi_{(D, \Sigma)}$ also includes rules for encoding that eq is an equivalence relation. Soft rules are encoded using a choice rule encoding the possibility of including (X, Y) in eq . We note that choice rules are special syntactic sugar available in ASP.

Differently from the original ASP encoding in (Bienvenu, Cima, and Gutiérrez-Basulto 2022), we encode similarity relations with a relation $\text{sim}_i(X, Y, S)$, where the first two arguments store the pair of constants to be assessed for similarity, while the third argument corresponds to a similarity score. This offers the flexibility of tuning the threshold for the same similarity measure. Facts of the form $\text{sim}_i(X, Y, S)$ are included in D after a preprocessing stage, as discussed in Section 4.3. Notably, the encoding requires a special treatment of null values. Missing values in

databases might be problematic (Fan and Geerts 2012), in particular, when joins need to be performed to evaluate rule bodies. To represent null values in $\Pi_{(D,\Sigma)}$, we use an atom $\text{empty}(nan)$, where nan is a special constant. To encode the fact that merges are not performed on unknown or missing values, we add an atom of the form $\text{not empty}(V)$ in rule bodies, for every joined variable V . This encoding prevents merges that otherwise would result from rule bodies being satisfied when considering two nulls equivalent. With the encoding $\Pi_{(D,\Sigma)}$ in place, solutions of (D, Σ) are then obtained by projecting stable models of $\Pi_{(D,\Sigma)}$ w.r.t. eq .

Theorem 3 ((Bienvenu, Cima, and Gutiérrez-Basulto 2022)). *For every database D and ER specification Σ : $E \in \text{Sol}(D, \Sigma)$ iff $E = \{(a, b) \mid \text{eq}(a, b) \in M\}$ for some stable model $M \in \text{SM}(\Pi_{(D,\Sigma)})$. In particular, $\text{Sol}(D, \Sigma) \neq \emptyset$ iff $\text{SM}(\Pi_{(D,\Sigma)}) \neq \emptyset$ ¹.*

4.2 ASP-based Algorithms

We now explain how to employ the ASP encoding to generate (maximal) solutions and other sets of merges, cf. Sec.3.3.

Solutions. Thanks to Theorem 3, we can obtain a single solution from $\text{Sol}(D, \Sigma)$ by using the ASP solver to generate a stable model of $\Pi_{(D,\Sigma)}$, then projecting onto its eq facts. Likewise, we can enumerate all or a fixed number of solutions by requesting an enumeration of $\text{SM}(\Pi_{(D,\Sigma)})$.

Maximal solutions. The maximal solutions correspond to the stable models of $\Pi_{(D,\Sigma)}$ having a \subseteq -maximal set of eq facts. We rely on `asprin`, a framework for implementing preferences among the stable models of a program (Brewka et al. 2015). In our case, we prefer a model M' over M if its projection to eq is a proper superset of that of M . `asprin` also allows us to compute n optimal stable models of a program.

Lower and upper bound merge sets. To compute $\text{LB}(D, \Sigma)$, we first construct the ASP encoding $\Pi_{(D,\Sigma^{\text{lb}})}$ based on Σ^{lb} and then we use the ASP solver to compute the (unique) answer set of $\Pi_{(D,\Sigma^{\text{lb}})}$, which we project onto the eq relation. We proceed analogously for $\text{UB}(D, \Sigma)$, but using Σ^{ub} .

$$\begin{aligned} \text{eq}(X, Y) \leftarrow & \text{song}(X, T, L, B), \text{song}(Y, T', L, B'), \\ & \text{not empty}(L), \text{sim}(T, T', S), S \geq 95, \text{eq}(B, B'). \end{aligned}$$

Possible merges. To generate $\text{PM}(D, \Sigma)$, it suffices to run the ASP solver in brave reasoning mode, and to project onto the eq relation. To check whether a particular pair (c, c') is a possible merge, we can run the solver on $\Pi_{(D,\Sigma)} \cup \{\perp \leftarrow \text{not eq}(c, c')\}$ observing that $(c, c') \in \text{PM}(D, \Sigma)$ iff this modified program admits a stable model.

Levels. To support our analysis of the impact of recursion, we will need a means of retrieving, for a given solution $E \in \text{Sol}(D, \Sigma)$, all triples (c, c', i) , where $\alpha = (c, c') \in E$ and i is the level of α in E . This can be achieved by considering a variant $\Pi^{\text{lvl}}(i)$ of $\Pi_{(D,\Sigma^{\text{ub}})}$, which is an ASP program that takes an integer i as a parameter and, by taking into consideration also the already derived merges inside E (i.e. those with level $< i$), applies single transitive steps or

rule applications, attaching integer i as the additional third element of the merges obtained in this way.

4.3 Similarity Computation

Our ASP rules contain body atoms $\text{sim}_i(X, X', S)$, but such similarity facts are not present in the data sources and need to be computed via external functions. The question then is how best to compute a sufficient set of sim_i facts to properly evaluate the program, while avoiding making calls to the external functions for every possible pair of values.

A naïve approach would be to compute and store the set $\text{Sim}_{\text{all}}(D)$, containing all facts $\text{sim}_i(c, d, s)$ such that c and d are data values of a form compatible with sim_i and $s = f_i(c, d)$, with f_i the function underlying the similarity relation sim_i . Although it requires only a polynomial number of function calls w.r.t. $|D|$, it is nevertheless extremely costly on even moderately-sized databases. A first improvement would be to only consider those pairs of constants $(c, d) \in R[i] \times R'[j]^2$ such that there is a rule ρ which contains an atom $\text{sim}_i(X, X', S)$ in which X and X' appear respectively in the i th (resp. j th) position of an R -atom (resp. R' -atom). However, as our experiments will show, this improved approach remains memory-consuming and its time consumption grows as the data size grows.

Another idea would be to exploit the structure of the rules so that we only call the similarity functions on pairs of constants that occur in a similarity atom for which the rest of the rule body is satisfied. For example, for rule σ_{ex} , we would remove $\text{sim}(T, T', S)$ and instead store the compared variables (T, T') in a fresh relation `getsim` as follows:

$$\begin{aligned} \text{getsim}(T, T') \leftarrow & \text{song}(X, T, L, B), \text{song}(Y, T', L, B'), \\ & \text{not empty}(L), \text{eq}(B, B'). \end{aligned} \quad (2)$$

Note however that the body still contains $\text{eq}(B, B')$, whose extension is initially unknown. We shall therefore employ a further ingredient: an overapproximation of the eq facts. For this, we could use the program $\Pi_{(D,\Sigma^{\text{ub}})}$, except that we do not know how to evaluate similarity atoms in rule bodies. One option would be to weaken the bodies by dropping all similarity atoms, but this yields a very loose approximation (cf. Appendix B). An alternative is to run the original $\Pi_{(D,\Sigma^{\text{ub}})}$ program, but making *online calls to the similarity functions*³ by replacing literals $\text{sim}(X, Y, S), S \geq \delta$ with $\text{sim}_{\text{ext}}(X, Y) \geq \delta$. We denote by $\Pi_{(D,\Sigma^{\text{ub}}_{\text{ext}})}$ the modified program. This will not only give us an upper bound on the true set of eq facts, but it will also compute a portion of the similarity facts. In fact, a further alternative would be to rely entirely on an online computation of similarity atoms over the original program. However, this is less efficient as results cannot be reused to compute different merge sets and may be time-consuming for the computation of $\text{MaxSol}(D, \Sigma)$ and $\text{PM}(D, \Sigma)$ (we refer to Appendix B for details).

Combining these ideas, we obtain the following approach to similarity computation, on input D, Σ .

² $R[i]$ denotes the constant at position i of R .

³Modern ASP solvers, e.g. `clingo`, support the syntax of external function calls (Kaminski et al. 2023)

¹We note that the proof in (Bienvenu, Cima, and Gutiérrez-Basulto 2022) does not consider nulls, but it can be easily extended.

Phase 1 Compute the unique stable model M_{ub} of $\Pi_{(D, \Sigma_{ext})}$ using an ASP solver with external function calls enabled. Let $U = \{ubeq(c, d) \mid eq(c, d) \in M_{ub}\}$, and let L contain all sim_i facts produced during the computation.

Phase 2 Let Π_{sim} contain, for each rule τ in $\Pi_{(D, \Sigma)}$ that encodes a (soft or hard) rule in Σ with at least one similarity atom in its body, a rule τ_{sim} that is obtained from τ by (i) deleting all similarity atoms and all associated comparison atoms (e.g. $S \geq \delta$), (ii) changing the head atom to $getsim_i(T, T')$, where $sim_i(T, T', S)$ occurs in τ , and (iii) renaming eq as $ubeq$.

Phase 3 Let M_{sim} be the unique stable model of $\Pi_{sim} \cup D \cup U$. For every pair (c, d) such that $getsim_i(c, d) \in M_{sim}$ and there is no $sim_i(c, d, _)$ fact in L , call the associated similarity function f_i on input (c, d) , and create the fact $sim_i(c, d, f_i(c, d))$. Let $Sim_{opt}(D, \Sigma)$ contain L and all of these newly created similarity facts.

Note that in Phase 2, our example rule σ_{ex} would be replaced by (2), but with $ubeq(B, B')$ in place of $eq(B, B')$.

The next result shows that this approach is correct, i.e. we get the same eq facts using Sim_{opt} rather than the full Sim_{all} .

Theorem 4. For every database D and ER specification Σ ,

$$\{M_{eq} \mid M \in SM(\Pi_{(D_{all}, \Sigma)})\} = \{M_{eq} \mid M \in SM(\Pi_{(D_{opt}, \Sigma)})\}$$

where $D_{all} = D \cup Sim_{all}(D)$, $D_{opt} = D \cup Sim_{opt}(D, \Sigma)$, and M_{eq} is the set of eq facts in M .

5 The ASPEN System

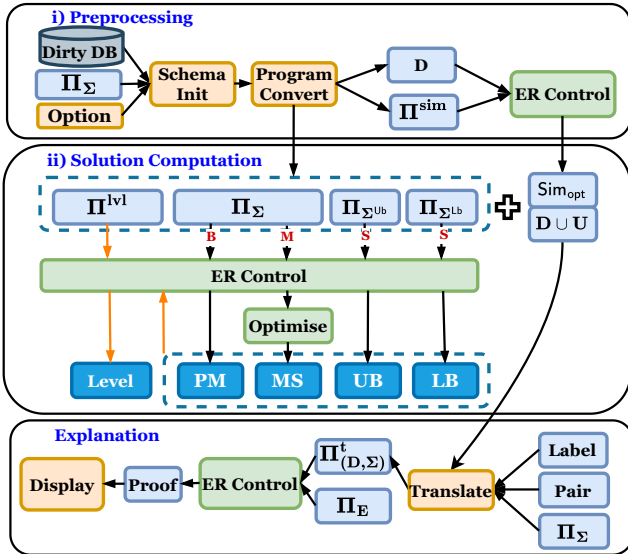


Figure 2: A General Pipeline of ASPEN.

An overview of the pipeline of ASPEN is shown in Figure 2, where boxes coloured light blue, strong blue, orange and green indicate an ASP program, ER solution, Python program and Python-based ASP solver, respectively. Arrows interconnecting these boxes symbolize the

flow of data. The blue dashed boxes highlight a pool of candidates that may be chosen based on the input options.

ASPEN takes as input an ASP encoding of an ER specification (cf. Sec. 4.1), a set of running options (see below) and a dataset in CSV/TSV format containing duplicates with corrupted values, such as missing ones. Depending on the input options, ASPEN outputs either the specified merge sets as shown by the deep blue boxes in the second row of Figure 2, and additionally explanations of all/particular merges (cf. Appendix A). ASPEN comprises two primary phases for generating (approximate) ER solutions:

- i) *Preprocessing Phase:* ASPEN initialises schema information and converts DB tuples into ASP facts, then computes similarity facts.
- ii) *Solution Computation Phase:* ASPEN takes the DB and similarity facts computed before as input, then it computes various types of LACE-based ER solutions as specified in the input options.

The *explanation phase* (third row, Fig. 2) additionally takes as input a set of rule labels and a merge pair to check, and visually displays proof trees of explanations (see App. A).

Preprocessing. Initially, ASPEN uses a Python function called the *Schema Initialiser* to load schema information from the input, including database tuples, schema names, relations, attributes, and foreign keys. This information is stored in a schema instance (Python object), which is used throughout the system workflow. The schema information is then passed to the *Program Converter (PC)*, which generates variants of Π_{Σ} and facts for the computation of different merge sets (cf. Sec. 4.2). During preprocessing, the PC utilises the relation and attributes structure in the schema instance to convert the database tuples into facts, including the reflexive closure of constants on merge positions. Additionally, empty entries in facts are replaced with the special constant *nan*. To precompute the similarity facts, the PC transforms the input ER program into similarity filtering programs Π^{sim} . The resulting programs, along with the generated database facts, are passed to the *ER Controller (ERC)*, a Python object that encapsulates reasoning facilities of the ASP solver *clingo* (Gebser et al. 2012) and algorithms (cf. Sec. 4.2). The ERC then executes the similarity filtering algorithm. The resulting set of triples (compared pair of constants and score) are stored as ternary *sim*-facts, which are then combined with database facts and the set of *ubeqs* to form the fact base, as indicated by the flow arrow from the first row to the second row in Figure 2.

Computing solutions. With the fact base as input, the ERC selects a program variant from the PC based on the selected solution option, as shown in the blue dashed box. Subsequently, the ERC executes grounding and solving with one of the corresponding reasoning modes, as indicated by the red initials in Figure 2. It computes the desired solutions as follows:

1. for the lower and upper bound merge sets (LB, UB) by making a *Standard grounding&solving* call to *clingo*.
2. for a set of maximal solutions *MS- n* with an enumeration limit of n , with *standard grounding* calls on Π_{Σ} but

employing `Maximisation` solving calls supported by the optimiser `asprin`.

- for a set of all possible merges PM using the `Brave` reasoning mode of `clingo`.

In addition, if the merge level is required, the derived merge set will be further combined with the level retrieval program Π^{lvl} to assign levels to each of the merges, cf. orange arrows in the second row of Figure 2.

Explanation. ASPEN uses several `xclingo` functions (Cabalar, Fandinno, and Muñiz 2020; Cabalar and Muñiz 2023) to provide explanations of merges, including: **i)** the `Translate` function, which takes as input a program Π and labels declared upon Π , and outputs a program Π^t , including auxiliary rules to track the original rules that have been triggered, and **ii)** the explanation program Π_E , which constructs explanations by grounding and solving together a stable model M of Π and Π^t .

As illustrated by the third row of Figure 2, ASPEN takes as input a merge pair to be justified and optionally a set of rule labels. A default set of labels is generated automatically capturing rule bodies of the ER program, if no rule labels are provided. Utilising the `Translate` function, ASPEN translates the rules and fact base into a trace program $\Pi_{(D,\Sigma)}^t$ according to the labels. The ERC then combines this with the fact base and the `xclingo` explanation program to derive a proof answer set, which is displayed as a graphical proof tree, explaining the input merge.

6 Experiments

We conducted experiments on real-life ER scenarios, including both pairwise and complex multi-relational datasets. We evaluate the following aspects: (1) effectiveness of our approach to similarity computation, (2) accuracy and efficiency of ASPEN, (3) effect of recursion, (4) factors impacting scalability and (5) use of Datalog engines to compute UB and LB approximations. We also exemplify generated justifications of merges in Appendix A.

6.1 Experimental Setup

Datasets. We consider two pairwise matching datasets from the bibliographic domain: DBLP-ACM (Köpcke, Thor, and Rahm 2010) and CORA, and four multi-relational datasets: **i)** a subset of the IMDB movie dataset (Deng et al. 2022); **ii)** a Music dataset sampled from the Musicbrainz database with synthetic duplicates; **iii)** a dirtier instance of the Music dataset, which contains the same amount of duplicates but a higher percentage of nulls and more syntactical variants on the duplicates; **iv)** a Pokémon dataset, sampled from the Pokémon database with synthetic duplicates and complex inter-table references. For simplicity, we refer to the datasets as *Dblp*, *Cora*, *Imdb*, *Mu*, *MuC* and *Poke*, respectively. Sources and statistics of the datasets can be found in Appendix C, as well as details about duplicates generation, sampling of datasets and the experimental environment.

Similarity measures and metrics. We calculate the syntactic similarities of constants based on their data types (Bahmani, Bertossi, and Vasiloglou 2017). We adopt the commonly

used metrics (Köpcke, Thor, and Rahm 2010) of *Precision* (**P**), *Recall* (**R**) and *F1-Score* (**F**₁). See App. C for details.

6.2 Similarity Filtering

We denote by sim_{opt} the similarity algorithm presented in Section 4.3. We implemented and ran sim_{opt} on all datasets and compared with the baseline procedure sim_{cs} , which computes the similarity of every possible pair of values (Bahmani, Bertossi, and Vasiloglou 2017) occurring in every pair of sim positions appearing in rule bodies.

Data	#At	#Cat	t_{cs}	t_{opt}	M_{cs}	M_{opt}	MRed.
<i>Dblp</i>	5	10.2M	96.3	531.4	512Mb	256Mb	50
<i>Cora</i>	17	0.8M	5.49	932.9	32Mb	4Mb	87.5
<i>Imdb</i>	22	19.6M	89.5	598.9	512Mb	128Mb	75
<i>Mu</i>	72	143.6M	664.03	772.3	4Gb	256Mb	93.8
<i>MuC</i>	72	147.1M	867.5	446.4	4Gb	256Mb	93.8
<i>Poke</i>	104	769.9M	9,419	4,142	32Gb	128Mb	99.6

Table 1: #columns denote the number of attributes, the sum of cross-products of constant pairs found in sim positions of a dataset. **MRed.** stands for the reduction rates of memory usage.

Results. Table 1 presents the results on execution time ($t_{\{\text{opt},\text{cs}\}}$) and memory usage ($M_{\{\text{opt},\text{cs}\}}$). We observe that sim_{opt} requires substantially less space than sim_{cs} , with memory usage reduction rates of 50%, 87.5%, 75% and 99.6% on *Dblp*, *Cora*, *Imdb* and *Poke*, and 93.8% on *Mu*/*MuC*, respectively. Note that in general the reductions become substantially larger as the number of attributes increases from 5 to 104. This is because a key strength of sim_{opt} is to leverage joins on attributes as preconditions for two constants to be compared. Consequently, as the number of attributes within a schema increases, a proportional increase occurs in the preconditions that can be employed to restrict unnecessary comparisons.

As for running time, sim_{cs} tends to be faster than sim_{opt} in datasets of smaller scale, yielding speed advantages of 5.5, 6.5 and 100+ times on *Dblp*, *Imdb* and *Cora*, respectively. Note that as *#Cat* increases this tendency is inverted: on *Mu* both have similar running times, and then on *MuC* and *Poke*, sim_{opt} becomes 2 times faster. This shift might be because time spent on similarity computations outweighs the sim_{opt} program execution time as the number of pairs to be compared grows quadratically for sim_{cs} .

6.3 Main Results

We evaluated performances of the lower LB and upper bound UB approximate solutions, a single maximal solution MS-1⁴, and all possible merges PM. We consider MS-1 as the default output of ASPEN. Additionally, we compared these solutions with two (pairwise) rule-based ER systems: Magellan (Konda et al. 2016) and JedAI (Papadakis et al. 2020). Note that the two closest approaches, Dedupalog (Arasu, Ré, and Suciu 2009) and MRL (Deng et al.

⁴We do not consider all n enumerated maximal solutions as there were only minimal variations among them.

Data	Method	F ₁	(P / R)	t _o	t _g	t _s
Dblp	Magellan	79.97	89.80 72.08	1.71	-	-
	JedAI	95.02	100 90.51	<u>49.16</u>	-	-
	LB	46.09	<u>97.10</u> 30.21	531.63	0.23	0.0039
	MS-1	96.21	95.36 <u>97.07</u>	532.17	0.45	<u>0.32</u>
	PM	<u>95.15</u>	92.85 97.57	538.5	<u>0.35</u>	6.75
	UB	91.11	85.50 97.57	531.41	-	-
Cora	Magellan	79.70	93.09 69.68	<u>131.13</u>	-	-
	JedAI	<u>90.53</u>	<u>95.53</u> 87.72	40.92	-	-
	LB	83.57	99.80 71.87	1,008	7.85	0.29
	MS-1	95.55	94.25 <u>96.79</u>	1,031	20.88	<u>10.50</u>
	PM	95.55	94.25 <u>96.79</u>	1,857.6	20.19	837.58
	UB	87.50	79.66 97.05	999.87	-	-

Table 2: Results on Pairwise Matching. Bold figures indicate the best performing results, and the second-best results are underlined.

2022), are not publicly available. To be comparable, we specified the same preconditions and similarity measures as in ASPEN, and followed the best performing setups for Magellan⁵ and JedAI (Efthymiou et al. 2023). Note that since both systems lack native support for multi-relational table inputs and do not recognise inter-references between tables, for multi-relational datasets, we performed directly pairwise matching for each table (see Appendix D for details).

Accuracy The main results for the pairwise and multi-relational datasets are presented in Tables 2 and 3, respectively. The default output MS-1 consistently achieves the highest F1-score across all datasets, outperforming both Magellan and JedAI by significant margins. On the pairwise benchmarks *Dblp* and *Cora*, MS-1 surpasses Magellan by 16% and 1.1%, and JedAI by 15% and 5% respectively. Substantial performance differences are observed in the multi-relational datasets. When comparing MS-1 with Magellan and JedAI, improvements of 11% and 1.7%, 7.7% and 26%, 28% and 51%, and 81% and 86% are observed on *Imdb*, *Mu*, *MuC* and *Poke* respectively. This shows ASPEN is promising, particularly the multi-relational setup.

We also observed that different (approximate) ASPEN solutions might lead to different performances:

UB vs LB. The comparison between LB and UB reveals extreme results in precision and coverage. LB reached the highest precision (with an average of > 99%) in all but one dataset, but has poor coverage with an average recall of $\approx 43\%$. Given that duplicated tuples often contain different versions of a value, it is not surprising that only few met the strong evidence required by hard rules. UB has the best coverage of the considered (approximate) solutions: on average $\geq 50\%$ higher recall than LB, 0.5%, 0.26%, 0.05%, 0.78% higher than MS-1 on *Dblp*, *Cora*, *Mu*/*MuC*, and slightly higher than PM on *Cora* and *Mu*/*MuC*. However, UB obtains the lowest precision of all solutions in all datasets. This

⁵<https://tinyurl.com/y6hupmrh>

difference can be explained by the presence of additional hard rules (the ones replacing soft rules) for UB, which allow for more merges but also introduce more false positives.

MS-1 & PM vs UB & LB. By contrast, the results for merge sets that make full use of LACE specifications are more balanced. On the one hand, MS-1 and PM outperformed LB in recall by $\approx 44\%$ on average, with slight sacrifices of precision on most datasets, only showing a considerable decrease on *MuC* (11% lower than LB). On the other hand, MS-1 and PM outperformed UB by significant precision margins of 10% and 7.3% on *Dblp* and 14% and 22.4%, on *Cora* and *Poke*, respectively, with only floating-point drops on recall. This highlights the effectiveness of combining soft rules and DCs to encourage the discovery of more merges while enforcing consistency to prevent false merges.

MS-1 vs PM. Observe that PM and MS-1 have identical results in both precision and recall in *Cora*, *Imdb*, *Mu* and *Poke*. A possible explanation is that when DCs are complementary to the soft rules in a specification, the combination behaves like hard rules (Bienvenu, Cima, and Gutiérrez-Basulto 2022). With no room for guessing, the specification derives a unique maximal solution, therefore MS-1 and PM include the same set of merges. Additionally, for *Dblp* and *MuC*, we see that precision of MS-1 respectively is 2.5% and 0.52% higher than that of PM. For these datasets, PM respectively obtained a 0.5% and 0.05% higher recall than MS-1. This underscores the characteristic differences between the two type of solutions: MS-1 prioritises precision while maintaining good coverage, whereas PM is more inclusive but may consequently contain more false positives.

Running Time The overall time, denoted as t_o , consists of preprocessing and ER time. Magellan and ASPEN involve preprocessing steps for blocking or similarity filtering, while JedAI interleaves similarity computation with the ER process, i.e., t_o includes both. For ASPEN, the ER time is further composed of grounding and solving time t_g and t_s . Note that t_o may not always reflect the optimal running times of ASPEN. Indeed, when the simplest LB setup is considered, directly running an ASP encoding with on-line similarity evaluation is much faster (cf. Appendix B). However, to ensure a consistent analysis of the ER times of ASPEN, we present the t_o as the sum of preprocessing and ER time.

As shown by the t_o column of Tables 2 and 3, both Magellan and JedAI significantly outperform MS-1 across all datasets. Magellan achieved speed advantages of 7.8, 8, 13, 16, 150, 311 times on *Cora*, *MuC*, *Mu*, *Poke*, *Imdb* and *Dblp* respectively. Similarly, JedAI is 10, 25, 36, 8.5, 71 and 182 times faster across *Dblp*, *Cora*, *Imdb*, *Mu*, *MuC* and *Poke* respectively. This is largely due to more costly similarity computations in the preprocessing stage, which are necessary to achieve high quality results on the complex multi-relational settings. Indeed, in *Mu*, *MuC* and *Poke*, ASPEN obtains *substantially higher accuracy* than the baselines. We compare t_o of other (approximate) solutions with that of Magellan and JedAI in Appendix D.

When looking at t_g and t_s of the different merge sets, LB is consistently the fastest, with t_g averaging within half

Data	Method	F ₁	(P / R)	t _o	t _g	t _s	Data	Method	F ₁	(P / R)	t _o	t _g	t _s
Imdb	Magellan	88.09	99.80 78.83	3.89	-	-	Mu	Magellan	89.78	98.63 82.38	64.83	-	-
	JedAI	97.49	99.40 95.67	16.65	-	-		JedAI	70.67	87.46 59.30	100.26	-	-
	LB	72.73	100 57.15	600.65	1.73	0.027		LB	64.08	99.79 47.19	798.22	25.85	0.071
	MS-1	99.27	99.39 99.14	609.96	10.27	0.79		MS-1	97.52	99.25 95.58	853.91	79.75	1.86
	PM	99.27	99.39 99.14	643.7	9.94	34.87		PM	97.52	99.25 95.58	1,152.4	78.64	301.51
	UB	99.27	99.39 99.14	598.9	-	-		UB	97.44	99.03 95.90	772.3	-	-
MuC	Magellan	55.54	97.51 38.83	66.87	-	-	Poke	Magellan	7.01	3.97 29.74	260.96	-	-
	JedAI	32.75	73.95 21.02	7.88	-	-		JedAI	2.1	1.08 46.56	23.46	-	-
	LB	53.95	99.79 36.97	474.56	28.01	0.062		LB	28.00	100 16.27	4,144	2.29	0.018
	MS-1	84.10	88.11 80.44	562.37	113.64	2.33		MS-1	88.71	92.88 84.90	4,271.8	127.67	2.17
	PM	83.87	87.59 80.46	893.44	113.26	333.78		PM	88.71	92.88 84.90	4,296	129.04	25.84
	UB	83.55	86.01 81.24	446.4	-	-		UB	77.00	70.43 84.90	4,142	-	-

Table 3: Results on Complex Multi-relational Datasets

a minute and t_s concluding in fractions of a second. This performance is expected, given that LB straightforwardly derives a single set of merges. Regarding MS-1, t_g took the majority of ER time, while solving is consistently much more efficient, terminating in ≤ 10 seconds. For PM, t_g behaved as for MS-1, but it required longer solving time (100 times longer in the worst case). This might be explained by the need to perform brave reasoning. We note that the size of a dataset may not always be the main factor impacting the solving time. For instance, despite containing only 1.9k tuples, t_s of MS-1 and PM on the *Cora* dataset are noticeably longer than on much larger datasets like *Mu* and *Poke*. Similarly, despite *Mu* and *MuC* being of the same size, the t_g of UB, MS-1 and PM on these datasets are very different.

6.4 Effect of Recursion

We executed the levels algorithm described in Section 4.2 on (approximate) solutions derived in our previous experiments on multi-relational datasets. We report accuracy results of UB and MS-1 on the *Poke* dataset and merge increments for various recursion levels of *MuC* and *Poke* in Figures 3 and 4, respectively. Additional results for other datasets can be found in Appendix E. We can observe that achieving convergence of merges requires more than one recursion level. For example, in Figure 3, *Poke* converged at level 2 (red dashed line). Figure 4 shows noticeable increments of merges obtaining 17.8% and 18.8%, 2.4% and 18.8% increment gains in the MS-1 and UB settings on *MuC* and *Poke*, respectively. This shows the efficacy of recursion in discovering new merges utilising merges derived from previous levels.

We present further results on the interplay of recursion and DCs and dataset characteristics in Appendix E.

6.5 Efficiency

Factors impacting scalability. Our efficiency analysis shows that both t_g and t_s increase monotonically with the *size of the data* and the *percentage of duplicates*. The impact on t_s is particularly significant for PM, increasing up to 52 times

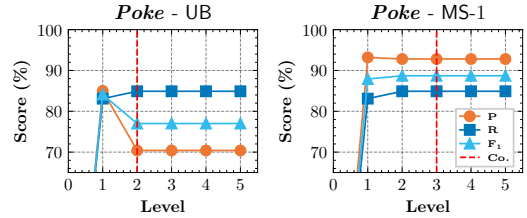


Figure 3: Impact of Recursion on Accuracy

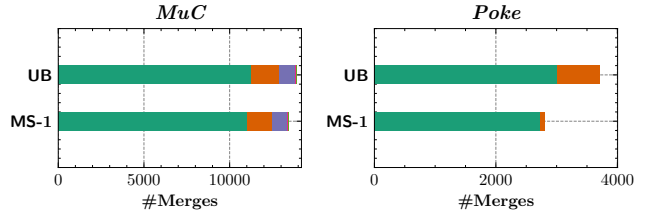


Figure 4: Merge Increments on Levels

when these factors are raised by fivefold. Similarly, lowering *similarity thresholds* consistently increases t_g , while t_s is up to 300 times longer on MS-1 and reaches a time-out (≥ 24 h) on PM. For more details, see Appendix F.

Datalog approximations. We examined the running times of Datalog programs $\Pi_{\Sigma^{ub}}$ and $\Pi_{\Sigma^{ub}}$ on all datasets using ASPEN and the rule engine VLog4j (Carral et al. 2019; Urbani, Jacobs, and Krötzsch 2016). For LB, ASPEN outperforms VLog4j on most datasets, whereas for UB, VLog4j is faster on all multi-relational datasets except *Poke*. These results suggest that the performance of different reasoning engines is impacted by characteristics of the data. See Appendix F for more details.

7 Discussion

We have introduced ASPEN, an ASP-based implementation of the LACE framework for collective, explainable, and re-

cursive ER. Distinguishing features of ASPEN are the consideration of a space of maximal solutions and the ability to supply explanations of derived merges. It is also one of only a handful of systems to natively support collective ER tasks, involving multiple database tables and entity types. A comprehensive experimental evaluation provided insights into how ASPEN performs, in terms of quality metrics and runtime, depending upon which notion of (approximate) solution is employed and how its performance compares to two baseline rule-based ER systems. The overall takeaway is that ASPEN is promising, as it is able to terminate in a reasonable amount of time (ER is typically an occasional offline task) and is competitive and often outperforms the baseline systems w.r.t. quality metrics, in particular, being able to effectively handle more complex multi-relational scenarios.

Our paper showcases entity resolution as an exciting but challenging application for ASP. Indeed, we believe that ER is as an ideal testbed for ASP techniques, as it is an important practical problem that naturally involves many solver functionalities, such as: brave and cautious reasoning (to identify possible and certain merges), preferred answers sets (to generate or reason over maximal solutions), external function calls (for similarity computation), and explanation facilities (to produce justifications of merges). By making ASPEN's code and data publicly available, we hope to facilitate future research on ASP-based approaches to ER.

While our experiments show that ASPEN can successfully handle some real-world ER scenarios, scalability remains an issue, and we expect that both general purpose and dedicated optimizations will be needed to be able to scale up to larger datasets and support even more complex reasoning over ER solutions. In addition to continuing to improve the similarity computation phase, we plan to explore the potential of employing specialized data structures or custom procedures for handling equivalence relations, as has been considered for Datalog reasoners (Nappa et al. 2019; Saheb-lamri et al. 2023). As parallelization has been successfully employed in some rule-based ER systems (Deng et al. 2022), another promising but non-trivial direction would be to see how parallel algorithms can be integrated into ASPEN. For this, we hope to build upon existing work on parallelization of Datalog reasoning (Perri, Ricca, and Sirianni 2013; Ajileye and Motik 2022) and ASP solving (Gebser, Kaufmann, and Schaub 2012).

We also plan to extend ASPEN to handle more expressive ER scenarios, building upon recent extensions to the LACE framework. Our top priority will be support not only global merges of entity-referencing constants, as considered in ASPEN and the original LACE framework, but also local (cell-level) merges of value constants (Bienvenu et al. 2023), so that e.g. some occurrences of 'J. Smith' can be matched to 'Joe Smith' while others are matched to 'Jane Smith'. Another important extension, requiring more significant changes to the ASP encoding, would be to allow for both merges and repair operations, as in the REPLACE framework (Bienvenu, Cima, and Gutiérrez-Basulto 2023), in order to be able to handle constraint violations that cannot be solved solely via merges.

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