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Efficient Predicate Invention using Shared NeMuS

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Abstract

Amao is a cognitive agent framework that tackles the invention of predicates with a different strategy as compared to recent advances in Inductive Logic Programming (ILP) approaches like Meta-Intepretive Learning (MIL) technique. It uses a Neural Multi-Space (NeMuS) graph structure to anti-unify atoms from the Herbrand base, which passes in the inductive momentum check. Inductive Clause Learning (ICL), as it is called, is extended here by using the weights of logical components, already present in NeMuS, to support inductive learning by expanding clause candidates with anti-unified atoms. An efficient invention mechanism is achieved, including the learning of recursive hypotheses, while restricting the shape of the hypothesis by adding bias definitions or idiosyncrasies of the language.

1 Introduction

One of the key challenges in Inductive Logic Programming (ILP) is finding good heuristics to search the hypothesis space. In Standard ILP, a good heuristic is one that can arrive quickly at a hypothesis that is both successful and succinct. To achieve this, the efficiency of hypothesis generation depends on the partial or even total order over the Herbrand Base to constrain deduction operations. This work presents a new approach called Inductive Clause Learning (ICL) building on [Mota and Diniz, 2016], which introduces a data structure named Neural Multi-Space (NeMuS) used in Amao, a neural-symbolic reasoning platform that performs symbolic reasoning on structured clauses via Linear Resolution [Robinson, 1965].

Inspired by [Boyer and Moore, 1972], NeMuS is a shared multi-space representation for a portion of first-order logic designed for use with machine learning and neural network methods. Such a structure contains weightings on individual elements (atoms, predicates, functions and constants or variables) to help guide the use of these elements in tasks such as theorem proving, as well as in using them to guide the search in the hypothesis space and improve the efficiency and success of inductive learning. Although it has some similarities to ILP, the hypotheses search mechanism is fundamentally

different. It uses the Herbrand Base (HB) to build up hypothesis candidates using inverse unification (adapted from [Idestam-Almquist, 1993]), and prunes away meaningless hypotheses as a result of inductive momentum between predicates connected to positive and negative examples. In [Mota et al., 2017] inverse or anti-unification [Idestam-Almquist, 1993] was added to allow induction of general rules from ground clauses, which is supported by the idea of regions of concepts. However, the inductive learning algorithm presented did not consider an adequate representation and use of bias, and the invention of predicates called predicate invention. Here we show how this can be achieved without using meta-interpreter level of bias specification or reasoning. It is important to note that weights are still not automatically used, but are taken into account when apparently unconnected literals have a common predicate name.

This paper makes the following contributions: it demonstrates that it is possible to have predicate invention without the use of meta-rules, and consequently, of Meta-Interpretive Learning. It shows that the NeMuS structure can be used for this purpose without generating numerous meaningless hypotheses, as the invention is made during Inductive Clause Learning. For that, we use bias or automated predicate generation. Finally, it demonstrates how invention of recursive rules takes advantage of weights of the logical component representation within NeMuS.

The remainder of this paper is structured as follows: section 2 gives some brief background on inductive logic programming and the Shared NeMuS data structure, sections 3 and 4 describe the implementation of inductive learning in Amao using the Shared NeMuS data structure, then section 5 describes some related work and section 6 discusses the work presented.

2 Background

2.1 Inductive Logic Programming (ILP)

The Inductive Logic Programming (ILP) main challenge, as defined in [Muggleton, 1991], is to search for a logical description (a hypothesis, H) of a target concept, based on set of (positive and negative) examples along with a set called background knowledge (BK). The central idea is that H is a consistent hypothesis, i.e. BK plus the hypothesis H entails the positive examples (e^+) , whilst does not entail the negative

ones (e^-) . Formally, $BK \cup \{H\} \vdash e^+$ and $BK \cup \{H\} \not\vdash e^-$.

Typical ILP systems implement search strategies over the space of all possible hypotheses. To reduce search complexity, such mechanisms rely on partial order of θ -subsumption [Nienhuys-Cheng and De Wolf, 1997], or on a total ordering over the Herbrand Base to constrain deductive, abductive and inductive operations [Muggleton *et al.*, 2015]. As a side-effect, the space of hypotheses grows even more due to quantification over meta-rules by the meta-interpretive process.

The Inductive Clause Learning (ICL) technique is fundamentally different, while its learning results are similar to ILP and MIL. It does not generate hypotheses to then test whether $BK \cup H$ entails positive examples but not the negative ones. Instead, ICL anticipates the elimination of inconsistent hypotheses at each induction step by *colliding* atoms obtained from a search across bindings of constants from e^+ and e^- . This is possible because NeMuS is a network of shared spaces (for constant terms, predicates and clauses), interconnected through weighted bindings pointing to the target space in which an occurrence of an element appears. In what follows this is briefly described.

2.2 Shared NeMuS

NeMuS is an ordered space for components of a first-order language: variables (space 0), atomic constants of the Herbrand Universe (space 1), functions (space 2, suppressed here), predicates with literal instances (space 3), and clauses (space 4), and so on. In what follows vectors are written \boldsymbol{v} , and $\boldsymbol{v}[i]$ or \boldsymbol{v}_i is used to refer to an element of a vector at position i.

Each logical element is described by a vector called T-Node, and in particular each element is uniquely identified by an integer code (an index) within its space. In addition, a T-Node identifies the lexicographic occurrence of the element, and (when appropriate) an attribute position.

Definition 1 (T-Node) Let $c \in \mathbb{Z}$, $a, i \in \mathbb{Z}^+$ and $h, \in \mathbb{N}$. A T-Node (target node) is a quadruple (h, c, i, a) that identifies an object at space h, with code c and occurrence i, at attribute position a (when it applies, otherwise 1). T_N is the set of all T-Nodes. For a vector x, of T-Nodes, with size n, and c is a code occurring in an element of x, then $\iota(c, x) = k$ is the index of c within the T-Node, $0 \le k \le (n-1)$.

As T-Nodes are the building block of our approach, all other elements follows from it and we describe as follows.

NeMuS Binding is an indexed pair $(p, w)_k$ in which $p \in \mathcal{T}_N$, $w \in \mathbb{R}$ and $k \in \mathbb{Z}^+$, such that $n_h(p) = h$, $n_c(p) = c$, $n_a(p) = a$ and $n_i(p) = i$. It represents the importance w of object k over occurrence $n_i(p)$ of object $n_c(p)$ at space $n_h(p)$ in position $n_a(p)$.

Variable Space (0) is a vector $V = [y_1, \ldots, y_n]$, in which each y_i is a vector of bindings. The elements of the variable space represent all of the occurrences of a variables. The logical scope of a variable X is identified by the instances of its bindings.

Constant Space (1) is a vector $C = [x_1, \dots, x_m]$, in which every x_i is a vector of bindings. The function β maps a constant i to the vector of its bindings $\vec{x_i}$, as above.

Compounds (functions, predicates and clauses), are in higher spaces. Their logical components are formed by a vector of T-Nodes (one for each argument), and a vector of Ne-MuS bindings (simply bindings) to represent their instances.

Compound in NeMuS is a vector of T-Nodes, i.e. $\boldsymbol{x}_a^i = [c_1, \dots, c_m]$, so that each $c_j \in \mathcal{T}_N$, and it represents an attribute of a compound logical expression coded as i.

Instance Space (I-Space) of a compound i is the pair (x_a^i, w_i) in which w_i is a vector of bindings. A vector of I-Spaces is a NeMuS *Compound Space (C-Space)*.

A literal (predicate instance), is an element of an I-Space, and so the predicate space is simply a C-Space. Seen as compounds, clauses' attributes are the literals composing such clauses.

Predicate Space (3) is a pair (C_p^+, C_p^-) in which C_p^+ and C_p^- are vectors of C-spaces.

Clause Space (4) is a vector of C-spaces such that every pair in the vector shall be $(x_a^i, [])$.

Note that the order of each space is only defined when they are gathered in the following structure.

Definition 2 (Shared NeMuS) A Shared NeMuS for a set of coded first-order expressions is a 4-tuple (assuming no functions), $\mathcal{N}: \langle \mathcal{V}, \mathcal{S}, \mathcal{P}, \mathcal{C} \rangle$, in which \mathcal{V} is the variable space, \mathcal{S} is the constant space, \mathcal{P} is the predicate space and \mathcal{C} is the clause space.

The next section describes how inductive learning is performed in Amao using NeMuS structure.

3 NeMuS-based Inductive Learning

ICL is based on the concept of Least Herbrand Model (LHM) [Lloyd, 1993] to anticipate the elimination of inconsistent hypotheses, at each induction step, before they are fully generated. This is done by *colliding*, via computing the *inductive momentum*, atoms obtained from bindings of arguments from e^+ (candidates to compose LHM) and e^- . Then, a pattern of linkage across verified literals is identified, anti-unification (adapted from [Idestam-Almquist, 1993]) is applied, and a conjecture is generated. The process repeats until the conjecture becomes a closed and consistent hypothesis.

3.1 Inductive Learning from the Herbrand Base

The problem of inductive learning involves a knowledge base of predicates, called background knowledge (BK), a set E of examples that the logical description H of the target concept (t) should prove (positive examples, e^+) and a set of examples that the target concept should not prove (negative examples, e^-). Figure 1 depicts a possible Herbrand Base that may explain how H entails a positive example for the concept t. In its turn, t can be unary $(p(a_k))$, binary $(p(a_k, a_{k1}))$, etc.

From attribute bindings of t, i.e. $\beta(a_k)$ and possibly $\beta(a_{k1})$, t is connected with attribute mates' bindings. Such connections bridge all concepts they may appear in $(p_1, q_1,$ etc.), like a path $\{c_1, \ldots, c_j\}$ in a graph, until it reaches the binding concept of t's last attribute. The interconnected concepts form a *linkage pattern*. For example, when p_1 , q_1 etc.,

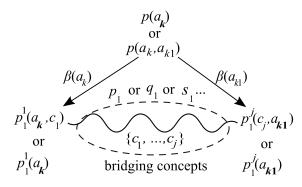


Figure 1: Portion of the HB with bindings of t's attributes.

are all the same concept, then a recursive hypothesis may be generated. Invention and hypotheses generation and will always take place in the *bridging concepts*' region (including the initial and last binding concepts).

The induction method is based on the following aspects: (a) *Inductive momentum* that iteratively "selects" only those atoms not likely to entail e^- , (b) *linkage patterns* among atoms passed (a), based on internal connections (bridging concepts); (c) *anti-unification* substitutes constants in an atom from the Herbrand Base by variables (and optionally (d) *category-first ordering*, [Schramm *et al.*, 2017], useful when BK contains monadic definitions of categories).

3.2 Linkage Patterns and Hypothesis

A special form of intersection ρ identifies the common terms between both literals. For instance, in Figure 1, suppose j=1, i.e. bridging concepts has just two literals of the same concept p_1 : $p_1(a_k,c_1)$ and $p_1(c_1,a_{k1})$. Then $\rho(p_1^1,p_1^2)=c_1$.

Definition 3 (Linkage and Hook-Terms) Let p and q be two predicates of a BK. There is a Linkage between p and q if a same constant, t_h , appears (at least) once in ground instances of p and q. We call t_h a hook-term, computed by $t_h = \rho(p,q)$. The attribute mates t_h w.r.t. an atom p, written $\overline{\rho(p,q)_p}$ is a set of terms occurring in p, but not in q.

With p_1^1, p_1^2 as above, c_1 is their hook term and $\overline{\rho(p_1^1,p_1^2)_{p_1^1}} = \{a_k\}$ and $\overline{\rho(p_1^1,p_1^2)_{p_1^2}} = \{a_{k1}\}$. This would form the definite clause $p(a_k,a_{k1}) \leftarrow p_1(a_k,c_1) \land p_1(c_1,a_{k1})$, which is not generated but it is built using anti-unification to generalize over its hooked ground literals. In the following definition we use the standard notion of a substitution θ as a set of pairs of variables and terms like $\{X_1/t_1,\ldots,X_n/t_n\}$.

Definition 4 (Anti-substitution and Anti-unification) Let G be a first-order expression with no constant term and X_1, \ldots, X_n be free variables of G, e is a ground first-order expression, and t_1, \ldots, t_n are constants terms of e. An anti-substitution is a set $\theta^{-1} = \{t_1/X_1, \ldots, t_n/X_n\}$ such that $G = \theta^{-1}e$, and G is called a simple anti-unification of e. The anti-unification f_{θ}^{-1} maps a ground atom e to its corresponding anti-substitution set that generalizes e, i.e. $f_{\theta}^{-1}(e) = \theta^{-1}$

Note that in the original definition of anti-unification, [Idestam-Almquist, 1993], G should be the generalisation of two ground expressions. Here, as we build a hypothesis by adding literals from a definite clause the definition is

Definition 5 (Anti-unification on linkage terms) Given two literals $p(a_k, a_z)$ and $q(a_z, a_{k1})$. A linkage term, say Z_0 , for their hook term a_z , is a variable that can be placed, by anti-unification, in the hook's position wherever it appears in the ground instance that will produce two non-ground literals.

The definite clause $p(a_k, a_{k1}) \leftarrow p_1(a_k, c_1) \wedge p_1(c_1, a_{k1})$, from Figure 1 (j = 1), $\theta^{-1} = \{a_k/X, a_{k1}/Y, c_1/Z\}$ is incrementally anti-unified, and so the following general clause is found: $p(X, Y) \leftarrow p_1(X, Z) \wedge p_1(Z, Y)$.

This concept is the fundamental operation to generate a hypothesis because it generalizes ground formulas into universally quantified ones. Before describing how negative examples are used we have the following definition.

Definition 6 (Hypothesis) Let H be a formula with no constant term, S be a set of ground atoms formed by concepts and constants from a Herbrand universe H_u , t is a ground atom with k terms (which belongs to the base of constants, H_0) such that $t \notin S$. We say that H is a hypothesis for t with respect to S if and only if there is a set of atoms $E = \{e_1, \ldots, e_n\}$ and a θ^{-1} such that for

- 1. every a of t, there is some $e_i \in E$ and $\rho(t, e_i) = a$
- 2. every e_i and e_{i+1} $\rho(e_i, e_{i+1})$ is not empty.
- 3. when 1 and 2 hold, then $H = \theta^{-1}(\{t\} \cup E)$.

An open hypothesis is one that at least one term of t have not been anti-unified. Thus, 1, 2 and 3 will always generate a closed least hypothesis.

Recall that *learning should involve the generation of a hypothesis and to test it against positive and negative examples*. Such a "test" could be done while the hypothesis is being generated. This is the fundamental role of the following concept. Note that, in the interest of saving space, the following sections shall use logic notation. It is important, however, to recall the definitions given in section 2.2, as the method described runs on top of the Shared NeMuS structure.

3.3 Inductive Momentum

In the following definition, a_k is a constant originating from the path of a positive example, and b_k is another constant originating from a negative example.

Definition 7 (Inductive Momentum) Let (x_a^i, w_i) and (x_a^j, w_j) be two I-spaces of atoms i and j, representing l^+ and l^- atomic formulas (literals) in the Herbrand base. If $\exists k$ and m, from e^+ and e^- , such that $l^+ \in \beta(k)$ and $l^- \in \beta(m)$, i.e. k is an element of x_a^i and m is an element of x_a^j , then the inductive momentum between l^+ and l^- with respect to k and m is

$$I_{\mu}(\boldsymbol{x}_{a}^{i}, \boldsymbol{x}_{a}^{j})_{m}^{k} = egin{cases} inconsistent & if \ i = j, \ and \ & \iota(k, \boldsymbol{x}_{a}^{i}) = \iota(m, \boldsymbol{x}_{a}^{j}) \ consistent & otherwise \end{cases}$$

Note that if i=j, then it is assumed $\|\boldsymbol{x}_a^i\|=\|\boldsymbol{x}_a^j\|$ since they are the same code in the predicate space. When it is clear in the context we shall simply write $I_{\mu}(l^+,l^-)_m^k$ rather than the T-Node vector notation.

Example 1. BK is formed by ground instances of binary and monadic predicates (not limited to them), atoms and Herbrand universe H_u as follows.

- 1. $\{p_1(a,a_1),\ldots,p_k(a_k,a)\},\$
- 2. $\{q_1(a_1,b_1),\ldots,q_j(b_j,a_1)\},\$
- 3. $\{r_1(c_1, a_k), \dots, r_m(a_k, c_m)\},\$
- 4. $\{t_1(b_i),\ldots,s_1(c_1),\ldots,v_1(c_m),\ldots\},\$
- 5. target p(X), with e^+ : p(a) and e^- : $\sim p(b)$.

When the BK is compiled its correspondent NeMuS structure is also built. The induction mechanism, at each step, adds to the premise of a hypothesis the next available atom from the bindings of a constant only if such an atom "resists" the inductive momentum.

Step	partial hypothesis	I_{μ}
1 2	$p(X) \leftarrow p_1(X, Y) p(X) \leftarrow p_1(X, Y)$	n/a $I_{\mu}(q_1(a_1,b_1),r_1(b_1,c_1))_{b_1}^{a_1}$
3	$p(X) \leftarrow p_1(Z, Y)$ $p(X) \leftarrow p_1(X, Y)$ $\land q_2(Z, Y)$	$ \begin{aligned} & = consistent \\ & I_{\mu}(q_1(a_1,b_2),r_2(b_2,c_2))_{b_1}^{a_1} \\ & = consistent \end{aligned} $
n	$p(X) \leftarrow p_1(X, Y) \\ \wedge q_j(Z, Y)$	$I_{\mu}(q_j(b_j, a_1), q_j(b_j, b_1))_{b_1}^{a_1}$ = inconsistent

For a partial hypothesis would be $p(X) \leftarrow p_1(X,Y) \land q_j(Z,Y)$ with $\theta^{-1} = \{a/X, a_1/Y, b_j/Z\}$, but the equivalent path from negative example would reach $q_j(b_j,b_1)$. This would allow p(b) to be also deduced, which is not what it is expected from a sound hypothesis. Thus, this hypothesis is dropped. For this example, a sound hypothesis could be $p(X) \leftarrow p_k(Y,X) \land r_1(Z,Y) \land s_1(Z)$.

Had the target concept be s(X) and positive example $s(c_1)$, then a possible hypothesis generated would be

$$s(X) \leftarrow s_1(X) \wedge r_1(X, Z_0) \wedge p_k(Z_0, Z_1).$$

3.4 Predicate Invention

Predicate invention, according to ILP definition, is a bias defined by the user via a declarative language. It is a way to deal with predicates missing from the BK for the lack of information. Suppose that target concept p of Figure 1 is ancestor(X,Y), BK is the set $B = \{father(jake, alice), mother(matilda, alice), \ldots\}$. There are two different concept relations in which the constant alice participates as a second attribute. There can be many instances of both father and mother, and no constant appear as first argument of both. There seems to be new concept that captures the property that all persons share when appearing as the first attribute of either relation.

On a closer look at Figure 2, it is possible that our general approach to predicate invention generates hypotheses that do not look like what we expect. For example

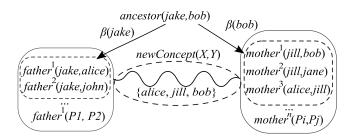


Figure 2: The invented predicate bridges the two regions of concepts.

ancestor(X,Y) given ancestor(jake,bob), might generate $ancestor(X,Y) \leftarrow father(X,Z_0) \land p_1(Z_0,Y)$.

Is is assumed that two concepts, say c_1 and c_2 , are "specialisations" of another concept c whenever there are objects appearing as second argument of both, but can only appear as first in one of them. This is informed to Amao as follows

Consider induction on T knowing E assuming P1 or P2 defines NewP, to mean that the bias we are looking for is NewP \rightarrow P1 or NewP \rightarrow P2.

We say that an *invented predicate bridges two regions of* concepts, and so allowing a more simple generalisation of ground rules into hypothesis. This is illustrated in Figure 2. $target(X,Y) \leftarrow newConcept(X,Z) \wedge target(Z,Y)$

Every time either or both concepts are involved in a hypothesis generation, the new concept is used to intentionally define the target predicate. So, from the figure above the rule base would be

$$newConcept(X,Y) \leftarrow (father(X,Y) \lor mother(X,Y))$$

Of course the new concept is parent and it is not a target concept to consider induction, but shall be used as a bridge or as a base form of a hypothesis, while the target shall be a linear or recursive linkage pattern (in our approach), or tail recursive.

4 Inductive Clause Learning with Invention

The method we are going to present in this section joins all ideas described in section 3. We shall use standard logic program notation for clauses just for readability sake, but recall that Amao language treats $q \leftarrow p$ as $q \lor \neg p$. The general idea of ICL can be summarised in three mains steps.

- to walk across the linkages found in the Herbrand Base in order to select atoms as candidates for composing hypotheses, as well as those to oppose the compositions
- 2. to compute I_{μ} of atoms as candidates for anti-unification that were selected from positive and negative linkages.
- 3. to generalize, via anti-unification, only atomic formulas likely to build consistent hypotheses, i.e. those composed by atoms consistent with respect to I_{μ}

In the following description we shall consider a dyadic theory with no function terms/

4.1 Selecting Candidates to Compose Hypothesis

Given the target t(X,Y), e^+ : $t(a_k,a_{k1})$ and e^- : $t(b_k,b_{k1})$. We access, from the NeMuS of the BK, $\beta(a_k)$ and $\beta(a_{k1})$. The initial view of the space of possible hypotheses that can be formed using atoms from the Herbrand Base and anti-unification is illustrated in Figure 3.

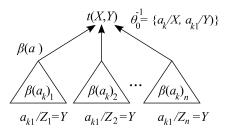


Figure 3: Space of hypotheses formed by Herbrand Base atoms.

Each $\beta(a_k)_i$ in a triangle represents a hypothesis formation branch that can be expanded following the bindings of the attribute in e^+ . Some of them may allow the deduction of e^- , and thus *inductive momentum* is applied to validate fetched atoms. After adding an anti-unified literal from $\beta(a_k)_1$ into the premise of the hypothesis being generated, say H_1 , the next induction step will take a branch from the attribute-mates of a_k to compute I_μ , generalize and so on. This is a depth-first walk across the Herbrand Base. In the breadth-first walk the generation of H_1 is postponed until all triangle branches have been initially exploited. For completeness sake it is implemented breadth-first.

4.2 Computing I_{μ} and Linkages

Accessing the bindings of constants is straightforward, we keep a loop selecting the instances of the literals that appear until the last is verified. Basically it is running while computing I_{μ} and moving across links of the sub-trees (triangles) from Figure 3.

If $I_{\mu}(q_1, \eta)_{b_k}^{a_k} = consistent$ for all $\eta \in \beta(b_k)$, then

If
$$q_1(a_k, a_{k1}) \in \beta(a_k)$$
 and $q_1(a_k, a_{k1}) \in \beta(a_{k1})$, then $H_1: t(X, Y) \leftarrow q_1(X, Y), \theta_1^{-1} = \{a_k/X, a_{k1}/Y\}$

Else if
$$q_1(a_k, a_{k1}) \in \beta(a_k)$$
 and $q_1(a_k, a_{k1}) \notin \beta(a_k)$, then $H_1: t(X, Y) \leftarrow q_1(X, Z_0), \theta_1^{-1} = \{a_k/X, c/Z_0\}$

Otherwise, get another $q_j \in \beta(a_k)$ and repeat the process until there are no more elements to test. In this case there is no hypothesis.

For a consistent H_1 , then there may exist $r_l \in \beta(c)$, and

- a) $r_l \in \beta(a_{k1}): r_l \neq q_1, \, r_l(c,a_{k1})$ is an atom from the Herbrand Base then for $\theta_1^{-1} = \{a_k/X, c/Z_0, a_{k1}/Y\}$ $H_1: t(X,Y) \leftarrow q_1(X,Z_0) \wedge r_l(Z_0,Y)$. (Chain in ILP)
- b) $r_l \notin \beta(a_{k1})$: path can only form a *long linear linkage* pattern. For non dyadic, if $I_{\mu}(r_l, \eta')$ is ok then expand hypotheses : $\beta(c) \{r_l\}$. H_1 's body is added with $r_l(Z_0, Z_1)$ (see expansion illustrated in Figure 4).

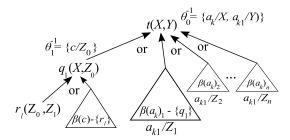


Figure 4: Expanding space of hypotheses following $\beta(a_k)$ and $\beta(c)$.

4.3 "Bias" as Invention of Predicates

Amao performs a similarity training on NeMuS's weights using the vector representation for each constant as well as for literals. Those with similar linkages end up with similar weight values associated to the argument they have and their position within them. Besides, bias may be used to add non targeted new predicates.

Non user bias: "automated" invention

For this, it is necessary "to invent" a predicate, say p_0 , such that H_1 becomes a closed hypothesis. For the sake of space θ^{-1} will be suppressed when anti-substitutions are clear.

 H_1 : $t(X,Y) \leftarrow q_1(X,Z_0) \wedge p_0(Z_0,Y)$, The invented predicate becomes the head of "invented hypothesis", as H_2 : $p_0(X,Y) \leftarrow r_l(X,Z_0)$ with $\theta_1^{-1} = \{c/X,c_k/Z_0\}$, and it becomes the *current open hypothesis*. The search now is guided by $\beta(c_k)$.

User defined bias for invention

When an assumption that r_l defines another concept, say p_b , then H_1 's body would have p_b and H_2 's head would have p_b , rather then p_0 . This would be something like assuming $r_l(X,Y)$ defines $p_b(X,Y)$, then

$$H_1: t(X,Y) \leftarrow q_1(X,Z_0) \wedge p_b(Z_0,Y),$$

 $H_2: p_b(X,Y) \leftarrow r_l(X,Z_0)$

- Assuming $r_l=q_1$, i.e. both are the same predicate (concept region).
 - 1. If $q_1(c, a_{k1}) \in BK$, and no bias given. Simple *linear linkage* pattern (chain)

$$H_1$$
: $t(X,Y) \leftarrow q_1(X,Z_0) \land q_1(Z_0,Y)$, "Shallow" recursive linkage pattern (recursive tail) H_1 : $t(X,Y) \leftarrow q_1(X,Z_0) \land t(Z_0,Y)$ H_2 : $t(X,Y) \leftarrow q_1(X,Y)$

The order they are introduced into the set of clauses is unimportant

- 2. If $q_1(c, a_{k1}) \notin BK$
 - (a) For bias and non dyadic theory: long linear linkage pattern of the same concept would generate $H_1\colon t(X,Y)\leftarrow q_1(X,Z_0)\wedge q_1(Z_0,Z_1)\wedge\ldots\wedge q_1(Z_n,Y).$ Instead, if $q_1(a_k,c)$ and $q_1(c,a_{k1})$ region's weights are similar, then invent of a recursive hypothesis.

$$H_1: t(X,Y) \leftarrow q_1(X,Z_0) \wedge t(Z_0,Y),$$

 $H_2: t(X,Y) \leftarrow q_1(X,Y),$

As there can be many bindings, we close an open hypothesis for each possible combination of bindings. Then, we keep computing the momentum and expanding a new branch for each combination (as explained in sections 4.1 to 4.3).

4.4 A Running Example: the Family Tree

Example 2. Consider the Family Tree, from [Muggleton *et al.*, 2015]. We may request to Amao the following

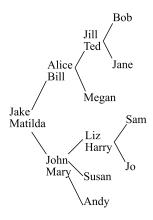


Figure 5: A simple family tree.

consider induction on ancestor(X,Y) **knowing** ancestor(jake,bob) **assuming** father(X,Y) **or** mother(X,Y) **defines** parent(X,Y).

i	atom/hypothesis	θ_i^{-1} / $\beta(a)$
0	ancestor(jake, bob)	$\{jake/X, bob/Y\}$
H_0	$ancestor(X,Y) \leftarrow$	
1	$\beta_1(jake)$	father(jake, alice)
	$\beta_1(bob)$	father(ted, bob)
	$I_{\mu} = consistent$	no hook
bias	-father(X,Y)	match both $\bar{\beta}_1$
for	parent(X, Y)	rename variable
H_0^i	$parent(\bar{X}, \bar{Y}) \leftarrow$	father(X, Y)
	$\{jake/X, bob/Y, alice/Z_0\}$	
H_1	$ancestor(X,Y) \leftarrow$	$parent(X, Z_0)$
	$\beta_1(alice)$	mother(alice, ted)
bias	$-mother(\bar{X},\bar{Y})$	match both $\bar{\beta}_1(\bar{a}\bar{l}i\bar{c}e)$
for	parent(X,Y)	rename variable
H_1^i	$parent(\bar{X}, \bar{Y}) \leftarrow$	$\overline{mother}(\overline{X},\overline{Y})$
H_1^-	reaches maximun body size,	do not add another
	$parent(Z_0, Z_1)$. Check for	region similarity
H_2^i	$ancestor(\bar{X}, \bar{Y}) \leftarrow$	parent(X, Y)
$\bar{H_1}$	$ancestor(X,Y) \leftarrow$	$parent(X, Z_0) \land$
	•	$ancestor(Z_0, Y)$

5 Related Work

Recent advances in Inductive Logic Programming (ILP) ease predicate invention by constraining logical learning operations with higher-order meta-rules, expressions that describe the formats of the rules. These rules have order constraints associated to them (to ensure termination of the proof) and are provided to the meta-interpreter, which attempts to prove the examples. When successful at this task, it then saves the substitutions for existentially quantified variables in the meta-rules [Muggleton, 2017]. This technique has been used to build Metagol [Muggleton et al., 2014; Muggleton et al., 2015], which has been successful in various examples. However, this approach tends to increase the generation of meaningless hypotheses and, consequently, leads to a large hypotheses space. [Cropper and Muggleton, 2016] tackles this challenge by extending Metagol to support abstractions and invention, but it remains a problem. Amao takes a totally different approach by using NeMuS to perform Inductive Clause Learning (ICL). This work extends ICL by using the results of exploring weights of logical components, already present in NeMuS, to support inductive learning by expanding clause candidates with literals which passed in the inductive momentum check. This allows an efficient invention of predicates, including the learning of recursive hypotheses, while restricting the shape of the hypothesis by adding bias definitions or idiosyncrasies of the language.

6 Discussion and Future Work

This paper has shown how the Amao Shared NeMuS data structure can be used in predicate invention without the need to generate meaningless hypotheses. This is achieved via Inductive Clause Learning, with automatic predicate generation that takes advantage of the degree of importance of constant objects. As atomic object, constants of the Herbrand base had never called much attention for logical inference, but only to validate resolution through unification. Here, we showed how they can be used to guide the search for consistent hypothesis in two ways.

First, by walking across their bindings from positive examples which are not rejected by inductive momentum with bindings of negative examples. Second, we use, as a heuristic to a faster generation of potentially recursive hypotheses, the maps or regions of similarities (item 2 of bias for invention), that constant bindings allow us to compute. Such maps as inductive mechanism is demonstrated in [Barreto and Mota, 2019].

Future works will focus on making more efficient use of weighted structures of concepts and their composition to allow learning and reasoning of complex formulae, as well as dealing with noise, uncertainty, and possible worlds. We then aim to incorporate deep learning-like mechanisms by taking advantage of the inherently interconnected *compound spaces* as a sort of layers for convolution when dealing with massive datasets.

Acknowledgement

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