

# Probably Approximately Correct Ontology Completion with `pacco` (Extended Abstract)

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Ontology learning, Angluin’s learning framework, PAC learning,

## 1. Introduction

Traditionally, ontology construction is conducted by knowledge engineers whose task is to formalize relevant concepts of the application domain and the relationships among them. This process is tedious and error-prone, and thus various methods to facilitate construction and, to some extent, ensure completeness of the resulting ontology by querying a domain expert have been investigated. For instance, learnability of axioms in lightweight DLs from a given set of interpretations has been studied in [1], and exact learnability of lightweight DL ontologies in Angluin’s framework via queries has been studied in [2]. In [3, 4, 5, 6, 7], learning DL concepts and axioms from given examples has been investigated. Learning a DL ontology that is inseparable from a target ontology w.r.t. a query language and a fixed dataset via querying an oracle has been investigated in [8]. In a recent work [9], PAC learning of DL concepts has been studied. For an overview on learning DL ontologies see [10, 11]. In another line of work, Formal Concept Analysis (FCA) [12] was employed for mining an  $\mathcal{EL}^+$ -basis of all axioms holding in a given model [13, 14]. In [15], a further approach for mining bases with a predefined and fixed role depth of concept expressions was proposed. In a more recent work [16], an approach for efficient mining of axioms from graph dataset was introduced and evaluated on large datasets.

The FCA-based approaches have the disadvantage that, in the worst case, they issue exponentially many queries to the domain expert. In [17], we presented an approach combining the advantages of both lines of research for completing the missing information in the ontology w.r.t. some given set of concept descriptions via issuing only polynomially many (w.r.t. the relevant quantities) queries to an expert. In the present work, we improve and implement the approach from [17] and present a PAC version of the ontology-completion method, which was initially introduced in [18] and implemented in [19]. Our solution is based on an algorithm for PAC learning a Horn envelope of an arbitrary propositional formula [20], which is itself based on an algorithm for exactly learning Horn formulas [21]. Our setting is the following. Given:

- an initial TBox  $\mathcal{T}_0$ ;
- an expert  $\mathcal{E}$  able to answer subsumption queries w.r.t. a TBox  $\mathcal{T}_{\mathcal{E}}$  that is unknown to us;
- a set  $\mathcal{C}$  of concept descriptions built over the signature of  $\mathcal{T}_{\mathcal{E}}$ ;
- a sampling oracle  $\mathcal{U}$  that, when called, returns a subsumption query over  $\mathcal{C}$  according to a probability distribution  $\mathcal{D}_{\mathcal{U}}$ ;
- a probability  $\delta$  and error bound  $\epsilon$  with  $0 < \epsilon, \delta < 1$ ;

compute a TBox  $\mathcal{T}$  such that  $\mathcal{T}_0 \subseteq \mathcal{T}$  and  $\mathcal{T}$  is, with probability at least  $1 - \delta$ , an  $\epsilon$ -approximation of  $\mathcal{T}_{\mathcal{E}}$ . Building on our work [17], we introduce `pacco`<sup>1</sup>, a tool for probably approximately correct (PAC) completion of DL TBoxes.

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<sup>1</sup><https://github.com/sertkaya/pacco>

**Definition 1.** If  $\mathcal{T}$  is a TBox and  $\mathcal{C}$  is a finite set of concept descriptions, we denote by  $\text{Impl}(\mathcal{T}, \mathcal{C}) = \{X \rightarrow Y \mid X, Y \subseteq \mathcal{C} \text{ and } \mathcal{T} \models \Box X \sqsubseteq \Box Y\}$  the set of implications corresponding to GCIs over conjunctions of concepts from  $\mathcal{C}$  entailed by  $\mathcal{T}$ .

Our approach to solving the above problem w.r.t.  $\mathcal{T}_0$ ,  $\mathcal{C}$ , and  $\mathcal{E}$  is to learn a set  $\mathcal{L}$  of implications that approximates  $\text{Impl}(\mathcal{T}^\mathcal{E}, \mathcal{C})$ . For this, we need an oracle capable of answering membership and equivalence queries with respect to  $\text{Impl}(\mathcal{T}^\mathcal{E}, \mathcal{C})$ . A membership query asks whether a certain subset of  $\mathcal{C}$  is a model of  $\text{Impl}(\mathcal{T}^\mathcal{E}, \mathcal{C})$ , and an equivalence query asks for a subset of  $\mathcal{C}$  that is a model of either  $\text{Impl}(\mathcal{T}^\mathcal{E}, \mathcal{C})$  or  $\mathcal{L}$  but not both. We rely on a (*domain*) *expert oracle*  $\mathcal{E}$  answering *subsumption queries* over  $\mathcal{C}$  of the form  $\mathcal{T}^\mathcal{E} \models \Box X \sqsubseteq C$ , where  $X \subseteq \mathcal{C}$  and  $C \in \mathcal{C}$ .

## 2. Probably Approximately Correct Completion

As a measure of approximation for TBoxes, we consider how often the two TBoxes give the same answers to subsumption queries:

**Definition 2.** Let  $\mathcal{T}_1$  and  $\mathcal{T}_2$  be two TBoxes,  $\mathcal{C}$  be a finite set of concept descriptions, and  $\mathcal{D}$  be a probability distribution of subsumption queries over  $\mathcal{C}$ . We define  $\text{dist}_{\mathcal{C}}^{\mathcal{D}}(\mathcal{T}_1, \mathcal{T}_2)$ , the  $\mathcal{C}$ - $\mathcal{D}$ -distance between  $\mathcal{T}_1$  and  $\mathcal{T}_2$ , as the probability of getting different responses to a subsumption query w.r.t.  $\mathcal{T}_1$  and  $\mathcal{T}_2$ :  $\Pr_{\mathcal{D}}(Q \mid (\mathcal{T}_1 \models Q) \Leftrightarrow (\mathcal{T}_2 \not\models Q))$ , where  $Q$  is a subsumption query over  $\mathcal{C}$ . For a given  $0 < \epsilon < 1$ , we call a TBox  $\mathcal{T}$  an  $\epsilon$ - $\mathcal{C}$ - $\mathcal{D}$ -approximation of  $\mathcal{T}^*$  if  $\text{dist}_{\mathcal{C}}^{\mathcal{D}}(\mathcal{T}, \mathcal{T}^*) \leq \epsilon$ . If, in addition,  $\mathcal{T}^* \models \mathcal{T}$ , i.e., the set of models of  $\mathcal{T}^*$  is a subset of the set of models of  $\mathcal{T}$ , then we say that  $\mathcal{T}$  is an *upper  $\epsilon$ - $\mathcal{C}$ - $\mathcal{D}$ -approximation* of  $\mathcal{T}^*$ .

**Proposition 1.** If  $\mathcal{T}$  is an upper  $\epsilon$ - $\mathcal{C}$ - $\mathcal{D}$ -approximation of  $\mathcal{T}^*$ , then the  $\mathcal{C}$ - $\mathcal{D}$ -distance between  $\mathcal{T}$  and  $\mathcal{T}^*$  is equal to  $\Pr_{\mathcal{D}}(Q \mid \mathcal{T} \not\models Q \text{ and } \mathcal{T}^* \models Q)$ .

Our algorithm has access to a *sampling oracle*  $\mathcal{U}$  simulating a user that, when called, returns a subsumption query over  $\mathcal{C}$  according to a probability distribution  $\mathcal{D}_{\mathcal{U}}$ . Given parameters  $0 < \epsilon, \delta < 1$ , it uses oracles  $\mathcal{E}$  and  $\mathcal{U}$  to compute a TBox  $\mathcal{T}$  that, with probability at least  $1 - \delta$ , is an upper  $\epsilon$ - $\mathcal{C}$ - $\mathcal{D}_{\mathcal{U}}$ -approximation of  $\mathcal{T}^\mathcal{E}$ .

The algorithm internally maintains a list of implications over  $\mathcal{C}$  such that, after termination, the corresponding set of GCIs is a desired approximation of  $\mathcal{T}^\mathcal{E}$ . It starts with the empty list  $\mathcal{L}$  of implications and uses counterexamples obtained from (simulated) equivalence queries to update  $\mathcal{L}$ . In the algorithms from [21] and [20], both positive and negative counterexamples are possible. A positive counterexample may be returned if the current hypothesis contains an implication not entailed by the target Horn formula. In our setting, such an implication corresponds to a CGI not entailed by the target TBox  $\mathcal{T}^\mathcal{E}$ . Since our goal is to obtain an upper approximation of  $\mathcal{T}^\mathcal{E}$ , we do not allow such GCIs. Therefore, we modify the algorithm to make sure that  $\mathcal{L}$  always contains only implications corresponding to GCIs entailed by  $\mathcal{T}^\mathcal{E}$ . This ensures that the resulting approximation of  $\mathcal{T}^\mathcal{E}$  is an upper approximation.

We simulate every equivalence query with several calls to the sampling oracle  $\mathcal{U}$ . If not all valid GCIs are entailed by  $\text{GCI}(\mathcal{L}) = \{\Box X \sqsubseteq \Box Y \mid X \rightarrow Y \in \mathcal{L}\}$ , we expect an equivalence query to return a subset  $X$  of  $\mathcal{C}$  closed under  $\mathcal{L}$  but not under  $\text{Impl}(\mathcal{T}^\mathcal{E}, \mathcal{C})$ . Since we aim at an  $\epsilon$ -approximation, we must be able to obtain, with an appropriate probability, such  $X$  whenever  $\text{dist}_{\mathcal{C}}^{\mathcal{D}}(\text{GCI}(\mathcal{L}), \mathcal{T}^\mathcal{E}) > \epsilon$ .

Every iteration of the algorithm starts with a search for a counterexample  $X$  to  $\text{GCI}(\mathcal{L})$ . For the  $i$ th iteration, our algorithm makes  $\lceil \log_{1-\epsilon} \frac{\delta}{i(i+1)} \rceil$  attempts to generate a counterexample [22]. Since  $\mathcal{L}$  contains only valid implications, the counterexample, if found, is always negative and is a model of  $\mathcal{L}$ . The rest of the iteration eliminates some counterexample  $Y \subseteq X$  by making sure that, for every  $C \in \mathcal{C}$ , the responses to the query  $\Box Y \sqsubseteq C$  w.r.t.  $\text{GCI}(\mathcal{L})$  and  $\mathcal{T}^\mathcal{E}$  are identical. To do this, the algorithm searches  $\mathcal{L}$  for the first implication  $U \rightarrow V$  such that  $Y = U \cap X \neq U$  and  $\mathcal{T}^\mathcal{E} \models \Box Y \sqsubseteq C$  for some  $C \in V \setminus Y$ . If such an implication is found, it is refined to  $Y \rightarrow \text{Compl}(Y)$ , where  $\text{Compl}(Y)$  is the *completion* of  $Y$  w.r.t.  $\mathcal{C}$  and  $\mathcal{T}^\mathcal{E}$ , i.e.,  $\text{Compl}(Y) = \{C \in \mathcal{C} \mid \mathcal{T}^\mathcal{E} \models \Box Y \sqsubseteq C\}$ . Otherwise, a new implication  $X \rightarrow \text{Compl}(X)$  is added to the end of  $\mathcal{L}$ . In both cases, the new implication is valid w.r.t.

$\delta$	$\epsilon$	Number of				Execution time (sec.)
		samples	queries	generated axioms	unrecovered axioms	
0.01	0.01	17502	18628	28	33	283
	0.1	3467	6474	23	35	147
	0.2	1690	4142	18	38	67
	0.3	359	1226	6	46	10
0.1	0.01	14456	16172	27	32	254
	0.1	3347	6431	22	34	132
	0.2	1084	2719	10	43	44
	0.2	745	2458	9	44	34
0.2	0.01	14965	16371	26	36	221
	0.1	3338	6383	22	36	128
	0.2	502	1533	6	47	16
	0.3	209	773	3	48	5
0.3	0.01	14332	16039	26	34	226
	0.1	3560	6923	22	34	140
	0.2	1375	3735	15	39	56
	0.3	19	88	1	49	<1

**Table 1**

Evaluation results completing GO-Plant. All values are rounded arithmetic means of 5 different runs.

$\mathcal{T}^{\mathcal{E}}$ . Note that the computation of completion requires  $O(|\mathcal{C}|)$  queries to the oracle  $\mathcal{E}$  and no calls to the sampling oracle  $\mathcal{U}$ . Combining this with the results in [20], we obtain

**Theorem 1.** Given a domain expert oracle  $\mathcal{E}$  and a sampling oracle  $\mathcal{U}$ , our algorithm computes, with probability at least  $1 - \delta$ , an upper  $\epsilon$ - $\mathcal{C}$ - $\mathcal{D}$ -approximation of  $\mathcal{T}^{\mathcal{E}}$ . The number of queries to  $\mathcal{E}$  and  $\mathcal{U}$  posed by the algorithm is polynomial in  $|\mathcal{C}|$ ,  $1/\epsilon$ ,  $1/\delta$ , and the minimal size of an implication set equivalent to  $\text{Impl}(\mathcal{T}^{\mathcal{E}}, \mathcal{C})$ .

### 3. Experimental Results

We evaluated our approach on a subset of the Gene Ontology (GO) [23], namely, GO-Plant<sup>2</sup>, which contains 97 classes and 155 logical axioms, among them 145 subclass axioms between concept names. For our experiments, we randomly deleted 50 of these subclass axioms and completed the resulting ontology with pacco in various test settings using the uniform distribution of subsumption queries in the sampling oracle. As base set  $\mathcal{C}$ , we took all the 97 concept names. As expert, we used the Hermit reasoner in conjunction with GO-Plant.

As expected, increasing  $\epsilon$  results in a smaller number of generated axioms and a larger number of unrecovered axioms, i.e., TBoxes that have larger  $\mathcal{C}$ - $\mathcal{D}$ -distance to the original TBox. The effect of varying  $\delta$  is much smaller, since the number of sampling queries used to simulate an equivalence query depends linearly on  $1/\epsilon$  and only logarithmically on  $1/\delta$ .

We compared the resulting TBoxes with the original GO-Plant using the ontology diff tool ecco<sup>3</sup> [24]. Ecco finds differences between ontologies and reports them in separate categories, one of which contains axioms from one ontology not entailed by the other. These are listed under *unrecovered axioms* in Table 1. The numbers remain relatively large even for small values of  $\epsilon$ . The reason is that  $\epsilon$  corresponds to the target maximal distance between the entire expert and completed TBoxes rather than between what has been removed and what has been recovered.

In future, we plan to explore the empirical behavior of our algorithm by simulating the expert  $\mathcal{E}$  and the sampling oracle  $\mathcal{U}$  from data with different distributions. For example, we may sample frequently occurring subsets of  $\mathcal{C}$  to learn GCIs with high *support* in the sense of association rule mining. It could also be interesting to sample infrequent subsets of  $\mathcal{C}$  and learn GCIs with low support. One could

<sup>2</sup><https://geneontology.org/docs/download-ontology>

<sup>3</sup><https://github.com/rsgoncalves/ecco>

say that GCIs highly supported by data can be accepted without resorting to an expert, whereas, for low-support GCIs, it is important to get a confirmation. It may be worthwhile to develop a modification of the algorithm to approximately complete both a TBox and an ABox w.r.t. a specific interpretation. This may require a slightly different notion of approximation accounting for the information contained in the ABox to be completed.

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