

# OWL Model of Clinical Trial Eligibility Criteria Compatible With Partially-known Information

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**Abstract.** Clinical trials are important for patients, for researchers and for companies. One of the major bottlenecks is patient recruitment. This task requires to match a great quantity of information about the patient with numerous eligibility criteria, in a logically-complex combination. Moreover, the patient's information required by some of the eligibility criteria may not be available at the time of pre-screening. In such situations, the classic approach based on negation as failure ignores the distinction between a trial for which patient eligibility should be rejected and trials for which patient eligibility cannot be asserted, which results in underestimating recruitment. We propose an OWL design pattern for modeling eligibility criteria based on the open world assumption to address the missing information problem.

**Keywords:** open world assumption, ontology design pattern, clinical trial eligibility criteria

## 1 Introduction

A major focus in all clinical trials is the recruitment of patients. Adequate enrollment provides a base for projected participant retention, resulting in evaluative patient data. Identification of eligible patients for clinical trials (from the principal investigator perspective) or identification of clinical trials in which the patient can be enrolled (from the patient perspective) is an essential phase of clinical research and an active area of medical informatics research. The National Cancer Institute identified several barriers that health care professionals claim in regards to clinical trial participation [1]. Among those barriers, lack of awareness of appropriate clinical trials is frequently mentioned.

Automated tools that help perform a systematic screening either of the potential clinical trials for a patient, or of the potential patients for a clinical trial could overcome this barrier [2]. Efforts have been dedicated to provide a uniform access to heterogeneous data from different sources. The Biomedical Translational Research Information System (BTRIS) is being developed at NIH

to consolidate clinical research data [3]. It is intended to simplify data access and analysis of data from active clinical trials and to facilitate reuse of existing data to answer new questions. STRIDE [4] is a platform supporting clinical and translational research consisting of a clinical data warehouse, an application development framework for building research data management applications and a biospecimen data management system. The i2b2 framework integrates medical record and clinical research data [5] and SHRINE [6] handles several sources by providing a federated query tool for clinical data repositories. The ObTiMA system relies on OWL and SWRL to perform semantic mediation between heterogeneous data sources [7]. Lezcano et al. propose an architecture based on OWL to represent patients data from archetypes, and on SWRL rules to perform the reasoning [8]. Several other efforts have been dedicated to the formal representation of clinical trials eligibility criteria to support automated reasoning [9]. Weng et al. performed an extensive literature review [10]. Ross et al. conducted a survey of 1,000 criteria randomly selected from ClinicalTrials.gov and found that 80% of them had a significant semantic complexity [11], with 40% involving some temporal reasoning. Tu et al. proposed an approach to convert free text eligibility criteria into the computable ERGO formalism [12]. O'Connor et al. developed a solution based on OWL and SWRL that supports temporal reasoning and bridges the gap between patients specific data and more general eligibility criteria [13]. The ASTEC (Automatic Selection of clinical Trials based on Eligibility Criteria) project aims at automating the search of prostate cancer clinical trials patients could be enrolled to [14]. It features syntactic and semantic interoperability between the oncologic electronic medical records and the recruitment decision system using a set of international standards (HL7 and NCIT), and the inference method is based on ERGO [15]. The EHR4CR project aims at facilitating clinical trial design and patient recruitment by developing tools and services that reuse data from heterogeneous electronic health records. The TRANSFoRm project has similar objectives for primary care.

All these works on data and criteria representation, integration and reasoning are motivated by the requirement to have the necessary information available at the time of processing the patient's data, and assume that somehow, that will be the case. Missing information that is required for deciding whether a criterion is met leads to underestimating recruitment. Solutions for circumventing this difficulty consist either in making assumptions about the undecided criteria, or in having a pre-screening phase considering a subset of the criteria for which patient's data are assumed to be available. Bayesian belief networks have been used to address the former [16] but require a sensible choice of probability values and may lead to the wrong assumption in particular cases. The latter leaves most of the decision task to human expertise, which provides little added value (if an expert has to handle the difficult criteria, taking the simple pre-screening ones into account adds little to the burden) and is still susceptible to the problem of missing information for the pre-screening criteria.

We propose an OWL design pattern for modeling clinical trial eligibility criteria. This design pattern is based on the open world assumption for handling

missing information. It infers whether a patient is eligible or not for a clinical trial, or if no definitive conclusion can be reached.

## 2 Background

### 2.1 Modeling eligibility criteria

A clinical trial can be modeled as a pair  $\langle (I_i)_{i=0}^n, (E_j)_{j=0}^m \rangle$  where  $(I_i)_{i=0}^n$  is the set of the inclusion criteria, and  $(E_j)_{j=0}^m$  is the set of the exclusion criteria. All the eligibility criteria from  $(I_i)_{i=0}^n \cup (E_j)_{j=0}^m$  are supposed to be independent from the others (at least in the weak sense: the value of criterion  $C_k$  cannot be inferred from the combined values of other criteria). Each criterion can be modeled as an unary predicate  $C(p)$ , where the variable  $p$  represents all the information available for the patient.  $C(p)$  is true if and only if the criterion is met.

A patient is deemed eligible for a clinical trial if all the inclusion criteria and none of the exclusion criteria are met.

$$\boxed{\text{patient eligible} \Leftrightarrow \bigwedge_{i=0}^n I_i(p) \wedge \neg \left( \bigvee_{j=0}^m E_j(p) \right)} \quad (1)$$

Before the final decision on the list of clinical trials a patient is eligible for, there are intermediate pre-screening phases where only the main eligibility criteria of each clinical trial are considered. Such pre-screening sessions rely on subsets of  $(I_i)_{i=0}^n$  and  $(E_j)_{j=0}^m$ , but the decision process remains the same.

For the sake of clarity, in addition to the general case, we will consider a simple clinical trial with two inclusion criteria  $I_0$  and  $I_1$ , and two exclusion criteria  $E_0$  and  $E_1$ .

$$\text{patient eligible} \Leftrightarrow I_0(p) \wedge I_1(p) \wedge \neg(E_0(p) \vee E_1(p)) \quad (2)$$

For example, these criteria could be:

- $I_0$ : evidence of a prostate adenocarcinoma;
- $I_1$ : absence of metastasis;
- $E_0$ : patient older than 70 years old;
- $E_1$ : evidence of diabetes.

According to equation 2, a patient would be eligible for the clinical trial if and only if he has a prostate adenocarcinoma and has no metastasis and is neither older than 70 years old nor has diabetes.

Because of De Morgan's laws, equation 1 is equivalent to:

$$\boxed{\text{patient eligible} \Leftrightarrow \left( \bigwedge_{i=0}^n I_i(p) \right) \wedge \left( \bigwedge_{j=0}^m \neg E_j(p) \right)} \quad (3)$$

Even though equation 1 and equation 3 are logically equivalent, the latter is often preferred because it is an uniform conjunction of criteria. Note that the negations in front of the exclusion criteria are purely formal, as both inclusion and exclusion criteria can represent an asserted presence (e.g. prostate adenocarcinoma for  $I_0$  or of diabetes for  $E_1$ ) or an asserted absence (e.g. metastasis for  $I_1$ ).

For our example:

$$\text{patient eligible} \Leftrightarrow I_0(p) \wedge I_1(p) \wedge (\neg E_0(p)) \wedge (\neg E_1(p)) \quad (4)$$

According to equation 3, a patient would be eligible for the clinical trial if and only if he has a prostate adenocarcinoma and has no metastasis and is not older than 70 years old and has not diabetes.

## 2.2 The problem of unknown Information

**Distinction between the patients that we know are not eligible and those that we do not know if they are eligible** When a part of the information necessary for determining if at least one criterion is met is unknown, the conjunction of equation 3 can never be true. This necessarily makes the patient not eligible for the clinical trial, whereas the correct interpretation of the situation is that the patient cannot be proven to be eligible. This is different from proving that the patient is not eligible, and indeed, in reality the patient can sometimes be included by assuming the missing values (cf. next section).

For our fictitious clinical trial, we consider a population of nine patients covering all the combinations of “*True*”, “*False*” or “*Unknown*” for the inclusion criterion  $I_1$  and the exclusion criterion  $E_1$ . Table 1 presents the value of equation 4 and correct inclusion decision for the nine combinations. Among the five patients ( $p_2$ ,  $p_5$ ,  $p_6$ ,  $p_7$  and  $p_8$ ) for which at least a part of the information is unknown, three ( $p_2$ ,  $p_7$  and  $p_8$ ) illustrate a conflict between the value of equation 4 and expected inclusion decision. A strict interpretation of equation 4 leads to the exclusion of the eight patients:

- for three of them ( $p_0$ ,  $p_3$  and  $p_4$ ), all the information is available;
- for two of them ( $p_5$  and  $p_6$ ), some information is unknown, but the available information is sufficient to conclude that the patients are not eligible;
- for the three others ( $p_2$ ,  $p_7$  and  $p_8$ ), however, the cause of rejection is either because one of the inclusion criteria cannot be proven ( $I_1$  for  $p_7$  and  $p_8$ ) or because one of the exclusion criteria cannot be proven to be false ( $E_1$  for  $p_2$  and  $p_8$ ).

Therefore, if we generalize, equation 3 alone is not enough in the case of partially-known information to make the distinction between the patients we know are not eligible (the first two categories, so this also includes patients for whom a part of the information is unknown) and those we do not know if they are eligible (the third category). This is a problem because patients from the first two categories should be excluded from the clinical trial, whereas those from the third category should be considered for inclusion.

Patient	$I_0$	$I_1$	$E_0$	$E_1$	$I_0 \wedge I_1 \wedge \neg E_0 \wedge \neg E_1$	Decision
$p_0$	T	T	F	T	F	Exclude ( $E_1$ )
$p_1$	T	T	F	F	T	Include
$p_2$	T	T	F	?	<b>F</b> cannot assert $\neg E_1$	<b>Propose</b> (assume $\neg E_1$ )
$p_3$	T	F	F	T	F	Exclude (both $\neg I_1$ and $E_1$ )
$p_4$	T	F	F	F	F	Exclude ( $\neg I_1$ )
$p_5$	T	F	F	?	F	Exclude ( $\neg I_1$ )
$p_6$	T	?	F	T	F	Exclude ( $E_1$ )
$p_7$	T	?	F	F	<b>F</b> cannot assert $I_1$	<b>Propose</b> (assume $I_1$ )
$p_8$	T	?	F	?	<b>F</b> cannot assert $I_1$ cannot assert $\neg E_1$	<b>Propose</b> (assume both $I_1$ and $\neg E_1$ )

**Table 1.** Evaluation of equation 4 and correct inclusion decision for all the possible values of  $I_1$  and  $E_1$ , with possibly unknown information

**Assuming values for criteria** Currently, the case of each patient diagnosed with cancer is examined in a multidisciplinary meeting (MDM), gathering (oncologists, pathologists, surgeons,...). The goal is to determine collectively the best therapeutic strategy for the patient, including consideration of potential inclusion into clinical trials. This preliminary stage is called pre-screening because it takes place before obtaining informed consent (i.e., before enrollment). It mainly relies on retrospective data coming from the patient health record. At this point, all the information necessary for determining the status of each inclusion and exclusion criteria may not be available, but the rationale is to focus on the clinical trials the patient may be eligible for. It should be noted that the missing items may differ between patients. One solution could be to assume the values of the unknown criteria in order to go back to a situation where inclusion or exclusion could be computed using equation 3.

In this case:

- inclusion criteria for which the available information is not sufficient to compute the status are considered to be met;
- exclusion criteria for which the available information is not sufficient to compute the status are considered not to be met.

Therefore, in the case where the available information is not sufficient to compute the status of a criterion, a different status is assumed depending on whether the criterion determines inclusion or exclusion.

Referring to our fictitious clinical trial, the lack of information about the absence of metastasis would lead to the assumption that  $I_1$  is true, whereas the lack of information about diabetes would lead to the assumption that  $E_1$  is false.

This situation raises several issues:

- a different status is assumed depending on whether the criterion determines inclusion or exclusion;
- the assumed status depends on the nature of the criterion (i.e. inclusion or exclusion) and not on its probability;
- one has to remember that the value for at least a criterion has been assumed in order to qualify the inferred eligibility (adamant for  $p_0$  or  $p_1$  vs “under the assumption that...” for  $p_2$ ,  $p_7$  and  $p_8$ );
- this qualification can be difficult to compute (the status of  $E_1$  is unknown for both  $p_2$  and  $p_5$ , but  $p_5$  can be confidently excluded whereas  $p_2$  can be included assuming  $E_1$ ).

### 2.3 The extent of the missing information problem

To determine the extent of the missing information problem, we analyzed the 286 prostate cancer cases examined during the weekly urology multidisciplinary meetings at Rennes’ university hospital between October 2008 and March 2009. This involved 252 patients: 25 of them were examined during two different MDM, and 5 were examined during three different MDM. Before the MDM, the patient’s data are collected in a form with 59 fields. The form supports the distinction

between known and unknown values (e.g. for “antecedent of neoplasm”, the possible answer are “yes”, “no”, “not specified”).

Overall, 58.64% of the values were unknown. On average, for each case studied in a MDM, 34.6 fields (among 59) had an unknown value.

All of the 286 cases studied had at least some of the 59 fields with an unknown value. Indeed, the case with the most fields filled still missed 19 of them.

54 fields (91.53% of 59) had a missing value in at least one of the 286 cases. The five fields that were systematically filled were: the patient identifier, the MDM date, the patient’s gender, the tumor anatomic site and the primary histological type.

During this period, 4 clinical trials related to prostate cancer running at Rennes Comprehensive Cancer Center were considered during the MDM. Table 2 presents the composition of the clinical trials fields and their proportion of missing information. It shows that for each clinical trial, all the patients had at least one missing field that prevented formula 3 to be true (regardless of the values of the known fields).

	CT1	CT2	CT3	CT4
Nb inclusion fields	15	19	17	10
Nb exclusion fields	10	9	13	11
Nb common fields	3	0	3	3
Missing values	50.06%	61.72%	52.99%	42.99%
Nb patients with all inclusion fields known	0	0	1	1
Nb patients with all exclusion fields known	4	3	0	1
Nb patients with all fields known	0	0	0	0
Nb eligible patients	30	23	7	2

**Table 2.** Importance of unknown information during pre-screening for the four clinical trials of interest

### 3 Methods

We propose an OWL design pattern for modeling clinical trial eligibility criteria. We then explain how the reasoning unfolds using the fictitious clinical trial from table 1. We validate our approach by verifying if the inferred outcome corresponds to the expected value from table 1. We evaluate our approach on the four clinical trials related to prostate cancer and the 286 cases mentioned at section 2.3. This allows us to quantify the impact of missing information on inclusion rates, as we have seen that in some cases, even partially-known information can lead to certain rejection.

## 4 Results

### 4.1 Eligibility criteria design pattern

- for each criterion, create a class  $C_i$  (at this point, we do not care if it is an inclusion or an exclusion criteria, or both) and possibly add a necessary and sufficient definition representing the criterion itself (or use SWRL);
- for each criterion, create a class  $Not\_C_i$  defined as  $Not\_C_i \equiv Criterion \sqcap \neg C_i$ . This process can be automated;
- for each clinical trial, create a class  $Ct\_k$  (placeholder);
- for each clinical trial, create a class  $Ct\_k\_include$  as a subclass of  $Ct\_k$  with a necessary and sufficient definition representing the conjunction of the inclusion criteria and of the exclusion criteria (cf. equation 3) ( $Ct\_k\_include \equiv \prod_{i=0}^n I_i \sqcap \prod_{j=0}^m Not\_E_j$ );
- for each clinical trial, create a class  $Ct\_k\_exclude$  (placeholder) as a subclass of  $Ct\_k$ ;
- for each clinical trial, create a class  $Ct\_k\_exclude\_at\_least\_one\_exclusion\_criterion$  as a subclass of  $Ct\_k\_exclude$  with a necessary and sufficient definition representing the disjunction of the exclusion criteria ( $Ct\_k\_exclude\_at\_least\_one\_exclusion\_criterion \equiv \bigsqcup_{j=0}^m E_j$ );
- for each clinical trial, create a class  $Ct\_k\_exclude\_at\_least\_one\_failed\_inclusion\_criterion$  as a subclass of  $Ct\_k\_exclude$  with a necessary and sufficient definition representing the disjunction of the negated inclusion criteria ( $Ct\_k\_exclude\_at\_least\_one\_failed\_incl\_criterion \equiv \bigsqcup_{i=0}^n Not\_I_i$ );
- represent the patient's data with instances (Fig. 1 and 2). For the sake of simplicity, we will make the patient an instance of as many  $C_i$  as we know he matches criteria, and as many  $Not\_C_j$  classes as we know he does not match criteria, even if this is ontologically questionable (a patient is not an instance of a criterion). How the patient's data are reconciled with the criteria by making the patient an instance of the criteria is not specified here: it can be manually, or automatically with necessary and sufficient definitions or SWRL rules for the  $C_i$  and  $Not\_C_j$  classes.

### 4.2 Reasoning

If all the required information is available, after classification the patient will be an instance of each  $C_i$  or  $Not\_C_i$ , and therefore will also be instantiated as either  $Ct\_k\_include$  (like  $p_1$  in Fig. 3),  $Ct\_k\_exclude\_at\_least\_one\_exclusion\_criterion$  or  $Ct\_k\_exclude\_at\_least\_one\_failed\_inclusion\_criterion$  (so at least we are doing as well as the other systems).



The screenshot displays a software interface with two main panels. The left panel, titled "Class Hierarchy", shows a tree structure of classes. The root is "owl:Thing", which branches into "ClinicalTrial" and "Criterion". "ClinicalTrial" further branches into "Ct\_DL", "Ct\_DL\_exclude", "Ct\_DL\_include", and "Ct\_DL\_propose". "Ct\_DL\_exclude" has two sub-classes: "At\_least\_one\_exclusion\_criterion" and "At\_least\_one\_failed\_inclusion\_criterion". "Criterion" has a sub-class "Patient (9)". The "Patient (9)" class is highlighted in yellow. The right panel has two tabs: "Asserted" (selected) and "Inferred". Below the tabs is a list of "Asserted Instances" containing nine entries labeled p0 through p8, each with a purple diamond icon. Below this list is a section titled "Asserted Types" containing five entries: "E\_1", "Not\_E\_0", "I\_1", "Patient", and "I\_0", each with a yellow circle icon. At the bottom of the interface, there is a horizontal scroll bar and a small dropdown menu.

Fig. 1. A patient for who all the information is available

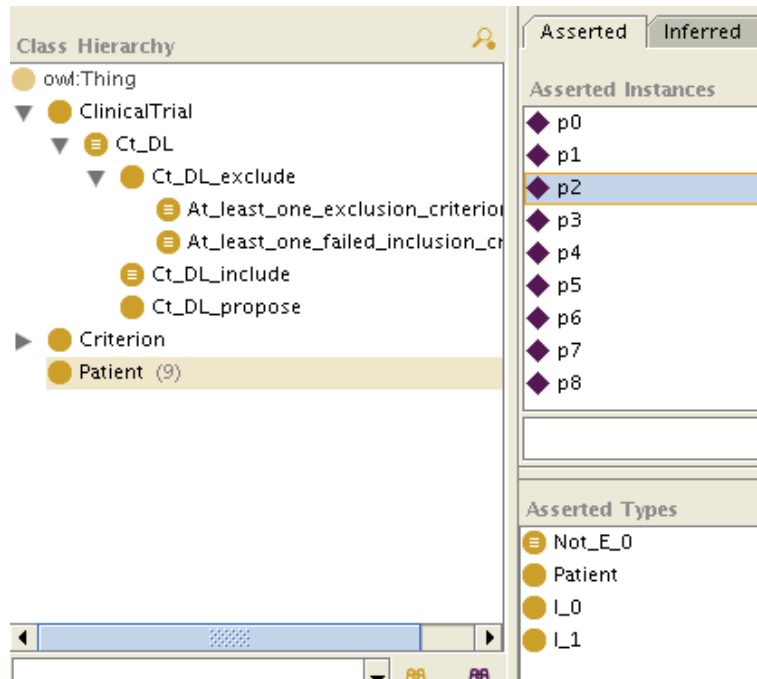


Fig. 2. A patient for who some information is unknown (here about  $E_1$ )

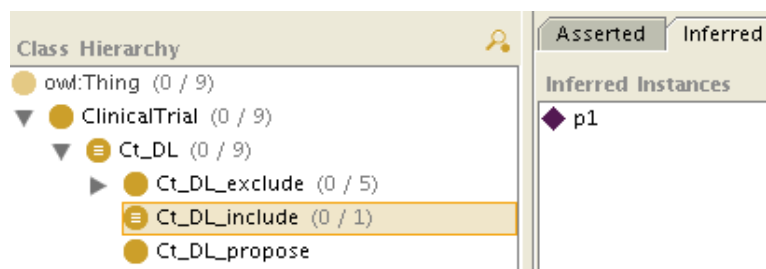
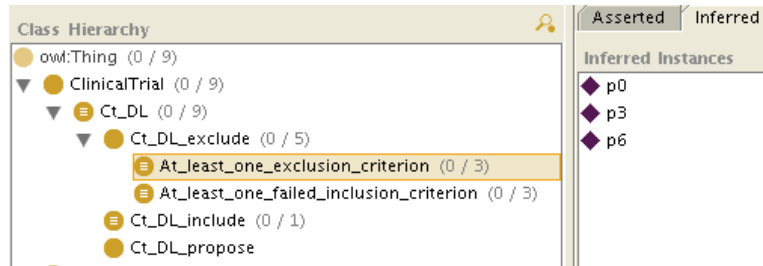
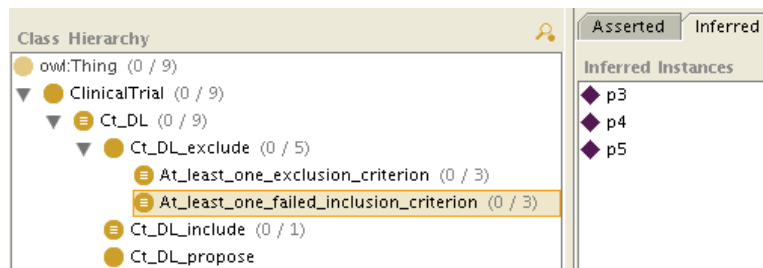


Fig. 3. The class modeling clinical trial inclusion after classification (here patient  $p_1$  can be included)



**Fig. 4.** The class modeling clinical trial exclusion because at least one of the exclusion criteria has been met after classification (here patients  $p_0$ ,  $p_3$  and  $p_6$  match the definition)



**Fig. 5.** The class modeling clinical trial exclusion because at least one of the inclusion criteria failed to be met after classification (here patients  $p_3$ ,  $p_4$  and  $p_5$  match the definition)

If not all the information is available, because of the open world assumption, there will be some criteria for which the patient will neither be classified as an instance of `C.i` nor of `Not_C.i` (Fig. 2), so he will not be classified as an instance of `Ct.k.include` either. However, the patient may be classified as an instance of `Ct.k.exclude_at_least_one_exclusion_criterion` or of `Ct.k.exclude_at_least_one_failed_inclusion_criterion`. As both are subclasses of `Ct.k.exclude`, we will conclude that the patient is not eligible for the clinical trial. We will even know if it is because he matched an exclusion criterion (like  $p_0$ ,  $p_3$  and  $p_6$  in Fig. 4), because he failed to match an inclusion criterion (like  $p_3$ ,  $p_4$  and  $p_5$  in Fig. 5), or both (like  $p_3$ ).

If the patient is neither classified as an instance of `Ct.k.include` nor of `Ct.k.exclude` (or its subclasses), then we will conclude that the patient can be considered for the clinical trial, assuming the missing information will not prevent it (like  $p_2$ ,  $p_7$  and  $p_8$ , who do not appear in Figs. 3, 4 and 5, consistently with Table 1). By retrieving the criteria for which the patient is neither an instance of `C.i` nor of `Not_C.i`, we will know which information is missing.

### 4.3 Validation

We modeled our fictitious clinical trial from section 2.1 as well as the nine combinations of values from section 2.2<sup>4</sup>. All the results were identical to the decision of table 1.

### 4.4 Evaluation

We evaluated our model on the first clinical trial (work is ongoing on the three others)<sup>5</sup>. Among the 286 cases, 0 were formally eligible, 122 were potentially eligible, and 164 were not eligible. The 30 cases that were identified as eligible by the experts during the multidisciplinary meetings were all among the 122 proposed by our system (precision was 0.24; recall was 1.0).

It should be noted that the *a posteriori* analysis of the 92 cases proposed by our model but not by the MDM revealed that several were not proposed even if they formally met the eligibility criteria because their Gleason score was deemed too low. We added an inclusion criterion requiring patients to have a Gleason score superior or equal to 7. This resulted in 54 cases potentially eligible, among which were 25 of the 30 actually eligible (precision was 0.46; recall was 0.83). The five false negative cases had a Gleason score of 6. Among the 29 false positive, at least 15 were rejected during the MDM because of additional information not available at the time of pre-screening: 8 because new results indicated that they did not have cancer, 3 because too much information was missing and 4 because other elements such as a relatively young age resulted in proposing a surgical treatment instead of the clinical trial.

<sup>4</sup> <http://www.u936.univ-rennes1.fr/dameron/clinicalTrial/ct-validation.tgz>

<sup>5</sup> <http://www.u936.univ-rennes1.fr/dameron/clinicalTrial/ct-getug14.tgz>

## 5 Discussion

The analysis of the first clinical trial demonstrates that missing information would have lead to the rejection of all the 30 patients proposed as eligible by the experts during the multidisciplinary meetings. Our approach correctly identified these 30 cases among the 122 it proposed as potentially eligible. This shows that our system confidently rejects non-eligible cases, which leaves more time to examine the others during the multidisciplinary meetings. Moreover, precision can be significantly improved by adding pragmatic criteria that further discriminate the patients who would not be considered as eligible even if they meet the pre-screening criteria. Note that this second step can be kept separated from the formal determination of eligibility but is useful both for the acceptance of the system by the experts and for maintaining the efficiency of the multidisciplinary meetings.

Missing information can partially be handled even with reasoning based on negation as failure using *ad hoc* conversion between inclusion and exclusion criteria. For example, the inclusion criterion “*absence of ischemic heart disease*” can be converted into the exclusion criterion “*presence of ischemic heart disease*”. The former will probably never be met because a patient’s record only mentions ischemic heart disease when they are present, whereas the latter will (correctly) only exclude those patients having evidence of ischemic heart disease. The problem is that if “*absence of ischemic heart disease*” had been an exclusion criterion, it would likewise have been converted into the inclusion criterion “*presence of ischemic heart disease*” and the system would have (incorrectly, at least during pre-screening) rejected patients whose record does not mention the presence nor the absence ischemic heart disease. Moreover, a criterion can be an inclusion criterion for a clinical trial and an exclusion criterion for another trial, so this strategy is not a general solution to the problem of missing information.

Reasoning about the conjunction of the eligibility criteria should be handled by OWL, which supports the open world assumption, rather than by related technologies such as SWRL which do not. It would be possible to write a SWRL rule that represents the conjunction of criteria (cf. formula 3). However, it is impossible to distinguish situations where we know that one criterion is not met from those where we cannot determine if it is met, because in both cases the rule will not fire.

Potential applications of our approach are not limited to clinical trials [16]. They cover all clinical decision situations where some information may be missing. We are currently adapting this approach for the determination of pacemaker alerts severity [17]. Electronic health records and clinical reports have been shown to exhibit large amounts of redundant information [18, 19], but Pakhomov et al. observed a discordance between patient-reported symptoms and their (lack of) documentation in the electronic medical records [20]. They noted that this has important implications for research studies that rely on symptom information for patient identification and may have clinical implications that must be evaluated for potential impact on quality of care, patient safety, and outcomes.

## 6 Conclusion

We showed that ignoring the missing information problem for automatic determination of clinical trial eligibility lead to over-estimate rejection. Systems based on negation as failure infer that the patient is not eligible if it cannot be proved that the is eligible, whereas the situations where it cannot be determined that the patient is eligible nor that he is not eligible should be identified and treated separately. A retrospective analysis of 252 patients with prostate cancer showed that for the four clinical trials of interest, all the patients had at least one missing value that had them rejected whereas 62 of them were actually eligible for at least one of the clinical trials.

We proposed a modeling strategy of eligibility criteria in OWL that leveraged the open world assumption to address the missing information problem. Our approach was able to distinguish a clinical trial for which the patient is eligible, a clinical trial for which we know that the patient is not eligible and a clinical trial for which the patient may be eligible provided that further pieces of information (which we can identify) can be obtained.

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