

Ontology development with CASL and HETS

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Overview

- CASL– background and introduction
- Ontology examples in CASL
- Type checking and static analysis

CASL– the Common Algebraic Specification Language

- de facto **standard** for specification of functional requirements
- developed by the “Common Framework Initiative” (an **open** international collaboration)
- approved by **IFIP WG 1.3** “Foundations of Systems Specifications”
- CASL **User Manual** (LNCS 2900) and **Reference Manual** (LNCS 2960) just have appeared

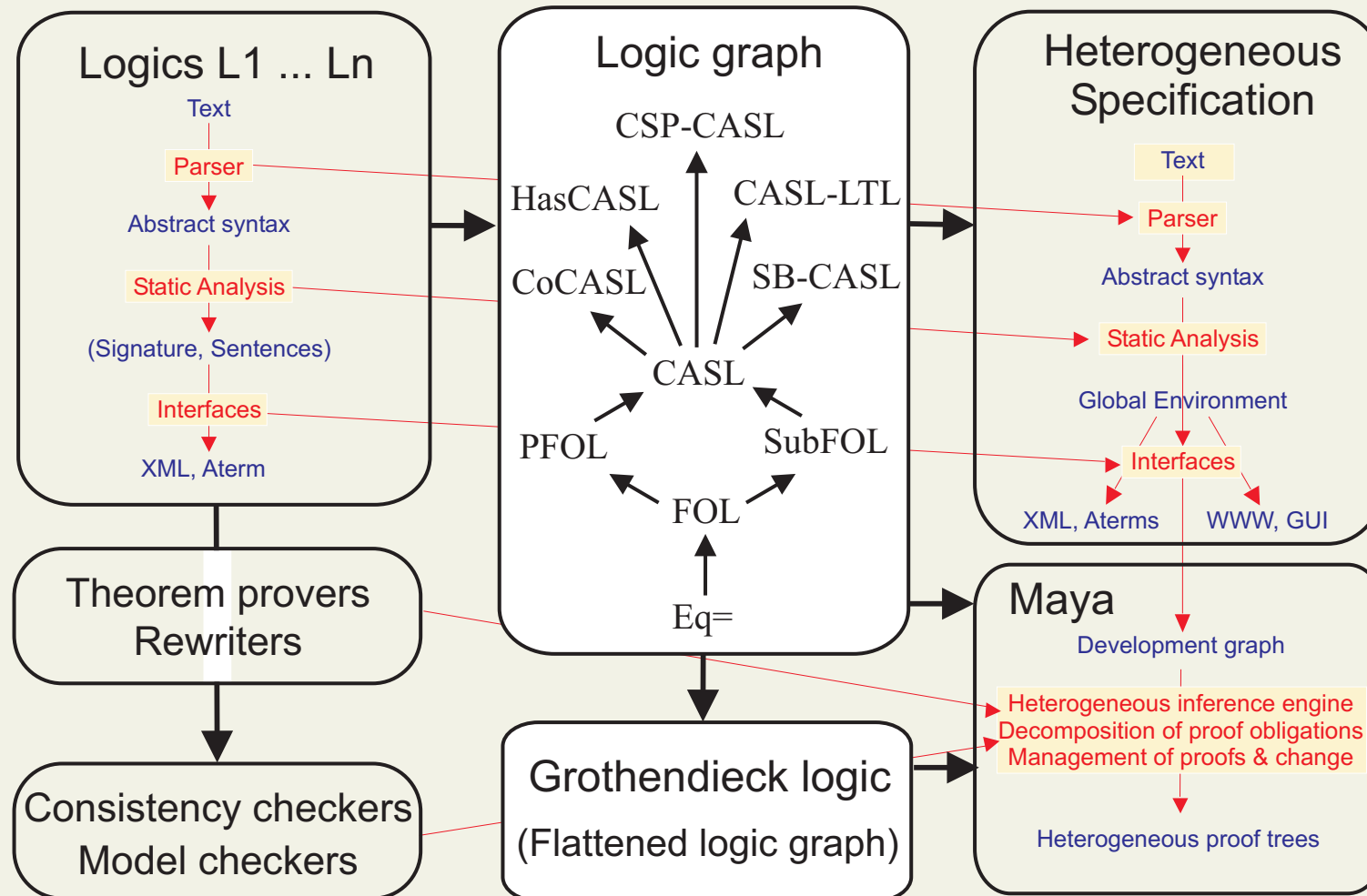
Central features of CASL

- supports both specification of **abstract requirements** (loose specification) as well as description of **specific models** (design specification)
- **In-the-small:**
first-order logic, subsorting, partiality, induction, datatypes
- **In-the-large:**
structuring of specifications and models, libraries distributed over the Internet

CASL in contrast to untyped FOL

CASL	untyped FOL
sorts	unary predicates
sub sorting	implication of unary predicates
just different sorts	disjointness of unary predicates
typed predicates	predicate implies unary predicates
axioms quantified over sorts	disjunction/conjunction of unary predicates implies axiom
typed functions	predicates with unique existential quantification

Architecture of the heterogeneous tool set Hets



Taxonomy with subsorting

%% subsumptions

sorts $A, B < Top$;
 $AA, AB < A$;
 $BA, BB < B$

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sorts $A, B < Top$;
 $AA, AB < A$;
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%% disjointness of concepts

free type $Top ::= \text{sorts } A, B$

%% unary predicates

preds $Un1 : Top$;
 $Un2 : BA$

Show ASCII

Analyse with Hets

(Typed) Roles and Predicates

preds $P1 : AB \times BA;$

$P2 : Top \times B \times BB$

$P3(x, y: B; z: A) \Leftrightarrow P1(z, y) \wedge P2(z, y, x)$

(Typed) Roles and Predicates

preds $P1 : AB \times BA;$

$P2 : Top \times B \times BB$

$P3(x, y: B; z: A) \Leftrightarrow P1(z, y) \wedge P2(z, y, x)$

$P1(z \text{ as } AB, y \text{ as } BA) \wedge P2(z, y, x \text{ as } BB)$

Axioms on sorts (concepts / classes)

Sorts can be defined with a Formula:

sort $C = x: Top \bullet \phi(x)$

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this equivalent to:

$C < Top$ and

$\forall x: Top \bullet x \in C \Leftrightarrow \phi(x)$

Overloading and Subsorting

- embeddings into supersorts work implicitly
- a term that is well typed for a supersort must be projected to the subsort with *as*
- you can have formulas depending on the membership in a sort like this:

$$\forall x: Top \bullet x \in C \Rightarrow \phi(x)$$

If you want a predicate defined on subsorts of a sort you better introduce the predicate like this:

spec TAXO =
 sorts *A*, *B*

```
free type  $Top ::= sorts\ A,\ B$   
end  
spec PARTHOOD =  
    TAXO  
then  
    preds  $P : A \times A;$   
            $P : B \times B$   
     $\forall x: Top$   
    •  $(P(x, x)) \Rightarrow (\phi(x))$  %% is ill typed  
     $\forall x: A$   
    •  $(P(x, x)) \Rightarrow (\phi(x))$  %% is correct  
end
```

Parametrised / Generic Specifications

Rule of Thumb: Keep parameters as small as possible

- a parameter is a basic specification or a named specification
- instantiation of such a specification works by providing all the symbols (and axioms) needed to fit the specified parameter
- an instantiation can add symbols to the signature

Examples of generic specification

library GENERICEXAMPLE2

spec PRIMITIVES =

sorts PD, PED, S, SL, T, TL

free type $PT ::= \text{sorts } PD, PED, S, SL, T, TL$

Examples of generic specification

library GENERICEXAMPLE2

spec PRIMITIVES =

sorts PD, PED, S, SL, T, TL

free type $PT ::= \text{sorts } PD, PED, S, SL, T, TL$

end

spec GENPARTHOOD [**sort** s] =

pred $P : s \times s$

$\forall x, y, z: s$

- $P(x, x)$ %(Ad11)%
- $P(x, y) \wedge P(y, x) \Rightarrow x = y$ %(Ad12)%
- $P(x, y) \wedge P(y, z) \Rightarrow P(x, z)$ %(Ad13)%

end

```
spec GENMEREOLGY [sort  $s$ ] =  
      GENPARTHOOD [sort  $s$ ]  
then  
%% some definitions and axioms  
end
```

```
spec GENMEREOLGY [sort  $s$ ] =  
    GENPARTHOOD [sort  $s$ ]  
then  
%% some definitions and axioms  
end  
spec MEREOLGY =  
    PRIMITIVES  
then GENMEREOLGY [sort  $T$ ]  
then GENMEREOLGY [sort  $S$ ]  
then GENMEREOLGY [sort  $PD$ ]  
end
```

```
spec MEREOLGY_ALTERNATIVE =  
    GENMEREOLGY [PRIMITIVES fit  $s \mapsto T$ ]  
then GENMEREOLGY [PRIMITIVES fit  $s \mapsto S$ ]  
then GENMEREOLGY [PRIMITIVES fit  $s \mapsto PD$ ]  
end
```