Inheritance: An overview

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Outline:

Semantic Networks

•Description Logics Definitions vs. Assertions Subsumption Classification

 Inheritance with Exceptions Multiple Paths Conflicted and Preempted Paths Credulous vs. Skeptical Reasoning

What are semantic networks?

Semantic networks can be viewed broadly or narrowly

- Broadly, a semantic network is any graph where
 --- the nodes represent concepts and
 --- the links represent relations between the concepts
- Narrowly, a semantic network is a graph where
 --- the nodes represent some restricted set of concepts
 --- the links represent some very restricted set of relations
 --- a clear semantics is given to nodes and, especially, links

Examples of narrowly defined semantic networks are

- --- terminological or description logics
- ---- inheritance networks (with exceptions)

In the early days of semantic networks (60s, 70s), semantic networks were broadly defined. Anything was okay.

example: Quillian: natural language understanding



Semantic networks have been used for natural language understanding, commonsense knowledge representation



What are we saying here? Is John's finger a part of the orchestra? Does Henny Penny go well with tarragon? Does John fight with himself? Are roast chicken music makers? What's going on? Problem: no semantics for concepts or links

Since the late 1970s, research in semantic networks has focussed on giving a clear semantics for the network (nodes and links) and providing sound and complete inference mechanisms. Networks have thus become much simpler.

Description Logics

- --- equivalent to a subset of first-order logic
- --- has some of the properties of frames: roles, slots, fillers
- --- subsumption and classification algorithms (polynomial)

Inheritance Networks with Exceptions

- --- allows subset, membership, cancels links
- --- polynomial algorithms to determine if there is a path between 2 nodes

Old, unsound networks continue to be used (in medicine, MeSH & UMLS), but there are efforts to be these on a sounder footing

Description Logics

(KL-ONE, Classic, Loom: Brachman, Levesque, Patel-Schneider, etc.)

Aim: provide a language for querying, organization,
 that is considerably more expressive than a database language
 but is much more efficient than first-order logic
 (Note: "pictorial" notion has been discarded)

<concept>::= /* atoms are primitive concepts */ <atom> | (AND <concept1> ... <conceptn>) | (ALL <role> <concept>) | (SOME <role>) <role> ::= <atom> | (RESTR <role> <concept>)

Examples: Can define California-white-wine as (AND WHITE-WINE (ALL region CALIFORNIA-REGION)) Can define Relaxed-meal as (SOME (RESTR course WINE-COURSE))

> Concepts are node in network; all links are subset (is-a) or membership (inst)

The interesting questions here are:

- Given two concepts, does one subsume another? (subsumption) (note that A subsumes B if B is a subset of A)
- Given a concept and a graph, where does the concept fit?

(classification)

The subsumption algorithm: (Brachman & Levesque, AAAI 1984)

To check if a subsumes b:

- Flatten both a and b by removing all nested AND operators So, (AND x (AND y z) w) becomes (AND x y z w)
- Collect all arguments to an ALL for a given role.
 So, (AND (ALL R (AND a b c)) (ALL r (AND X))) becomes (AND (ALL r (AND a b c X)))
- 3. Assuming a is now of the form (AND a1 ... an) b is now of the form (AND b1 ... bm), return true iff

(i) if ai is an atom or a SOME, then one of the bj is ai

(ii) if ai is (ALL r x), then one of the bj is (ALL r y) where x subsumes y.

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Example: (AND person (ALL child doctor)) subsumes (AND (AND (AND person (ALL child rich)) (AND male (ALL (RESTR child rich)) (AND doctor (SOME (RESTR specialty surgery)))))) Complexity of subsumption is $O(n^2)$, where n is length of expression.

Classification is done using subsumption.

Take a concept, find a place in the tree such where it is subsumed by a node, but can't be subsumed by the node's children. Note that one must take into account that several nodes may subsume a particular node.

Worst case would have to examine each node in tree.

Complexity of classification: $m(O(n^2))$, where m is no. of nodes in tree

Note: in practice, power of description logics used for standard inheritance logic + slots (roles) and fillers



When we add slots and fillers, we can say more interesting things. For example, in the drug formulary network, we may be interested in representing facts about price, purpose, control information, etc.

Saying these facts is enabled by the syntax of description logics, which allow: and (all r1 concept1) (all rn conceptn)



Description logics also allow the creation of one concept using conjunction. Thus, we may consider the class of all drugs that are both anticonvulsants and sedatives



What description logics won't let you do: exceptions So there's no way to say that cephalosporins are typically expensive but one particular type is moderately priced.

Well, almost no way. You can "cheat" by adding another role -- a flag -and turning it on if there's an exception



Inheritance with exceptions



Brief Review of Inheritance



But what happens when we need to talk about exceptions?

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We need to represent exceptions within the inheritance network

So we Introduce Inheritance Networks with Exceptions



Note that this potentially changes the semantics of <u>all</u> the links. Obviously it is no longer the case that all mammals are placental! Rather, the Is-a link is a "close-to-subset" link; most mammals are placental. These new links are called defeasible. Some systems mix strict is-a and defeasible is-a; here we stick to all defeasible links

So we Introduce Inheritance Networks with Exceptions



So this is what we are saying here:

All cats are mammals; all kangaroos are marsupials; -all marsupials are mammals; all <u>most</u> mammals have placentas marsupials do <u>not</u> have placentas

Review of Inheritance



Inheritance with Exceptions

- Allows us to specify <u>exceptions</u> to a general class
- <u>Cancels link:</u> nonmonotonicity

We could write this in a nonmonotonic logic: (translational semantics)

mammal(x) : placental(x)

marsupial(x) ==> not placental(x)
kangaroo(x) ==> marsupial(x)

placental(x)

So if Kiri is a kangaroo, we conclude that Kiri is not placental

Note that cancels links also aren't strict; we can have exceptions to exceptions



What we are saying here:

Most animals don't fly; mammals usually don't fly but bats do fly; Birds usually do fly, but penguins don't fly.

Dealing with multiple inheritance:



And there are 3 paths between penguins and non-flying creatures:

penguin isa bird isa animal isa non-flying creature
 penguin isa bird is-not-a non-flying creature
 penguin isa non-flying creature
 which path should we use?

Dealing with multiple inheritance: choosing the right path



It might seem that all we need to do is choose the shortest path, but it turns out that won't always work, especially if we allow redundant links

E.G., consider:



There is a path of length 2 that says baby royal elephants are grey and a path of length 2 that says baby royal elephants are not grey!

Dealing with multiple inheritance: choosing the right path



The shortest-path criterion will not work; instead we use the criterion of <u>specificity</u>

To define specificity, we must give a formal definition of path-based inheritance

(Touretzky, 1986; Horty, Thomason, and Touretzky, 1987, 1990; Stein, 1989, ..., 1992) Formal definition of path-based inheritance (Horty, 1994; Stein, 1992)

- Links may be positive (is-a) or negative (is-not-a; cancels)
- A path is a restricted sequence of positive and/or negative link

positive paths are made up of positive links
negative paths are positive paths, with one negative link at end



• A context: a network and a set of paths (arising out of the network)

In case of multiple paths, how do we choose the "right" path?

A path is <u>inheritable</u> or <u>undefeated</u> in a context if it is:

constructible

i.e. can recursively be built out of paths in the network



Conflicts and Preemptions in more detail:

Conflicts:

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A path of the form \pi(x,\sigma,y) is conflicted with a path of the form \pi(x,\tau,y)
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Penguin

formalizes the notion of <u>specificity;</u> Penguins are more specific than birds What do we do when there are conflicts, preemptions in the network?

•For the Nixon diamond, 2 approaches:

---- conclude that penguins are not fliers

due to specificity)

(the argument that penguins are not fliers

---- accept either that Nixon is a pacifist or that he's not a pacifisit (that is, accept one coherent path: <u>credulous reasoning</u>)

--- decide to conclude nothing about Nixon's pacifism because of the potential conflict (skeptical reasoning)



Should we propagate ambiguity?



Ambiguity Propagation: where it makes sense

(Stein, 1989, 1990, 1992)



seedless grape vine

Whether or not a seedless grape vine is a fruit plant, it's certainly a plant! Classical skeptical inheritance is ambiguity blocking; can't reason beyond conflict If we take "ideally skeptical inheritance" as intersection of credulous extensions, get the desired conclusion: seedless grape vines (and grape vines) are plants **Computing Inheritance**

•Upwards inheritance is efficient (polynomial) downwards inheritance is intractable (NP-hard) (Selman & Levesque, 89)

•Stein gives a polynomial algorithm based on an upward traversal, removal of problematic, preempted edges (AIJ, 1992)