

Role-based representation and inference of biochemical processes

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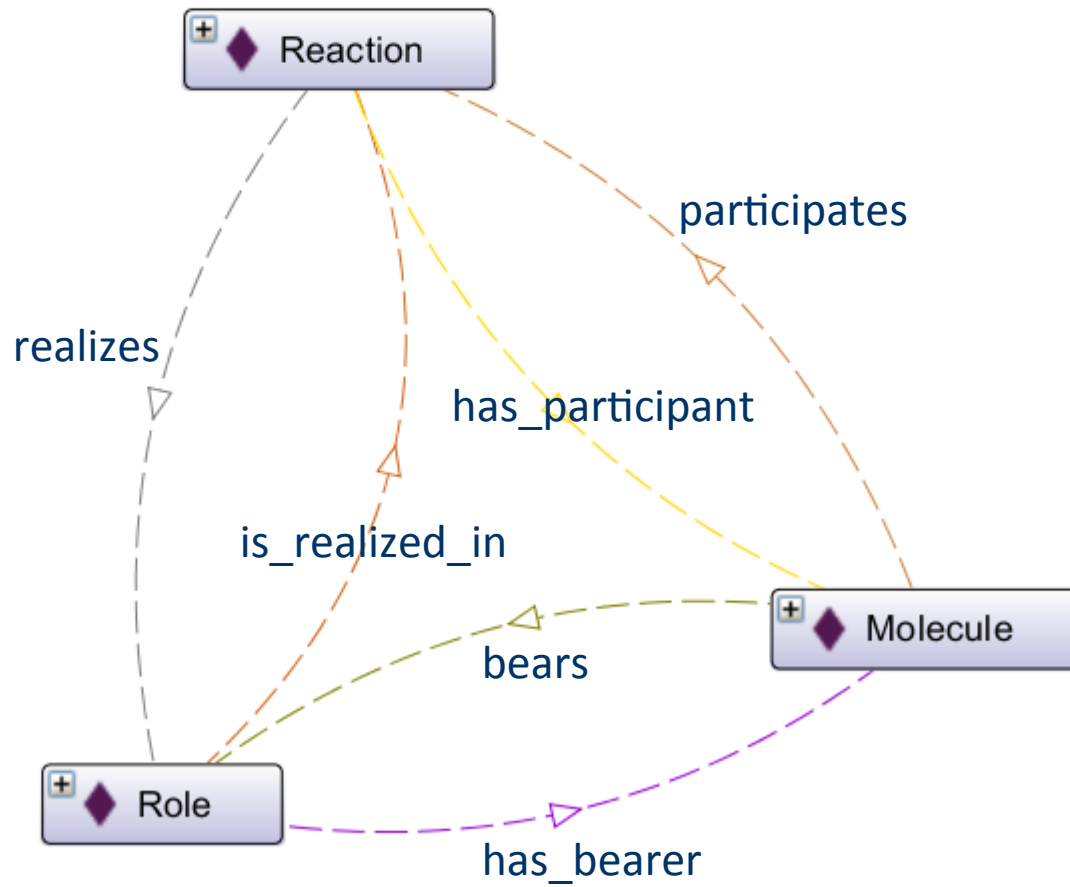
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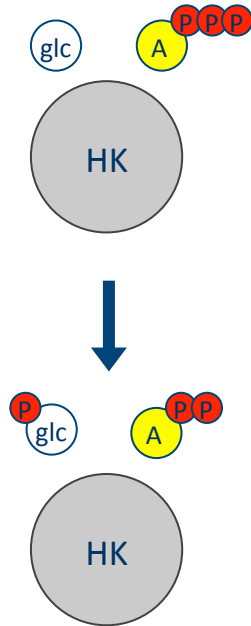
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motivation

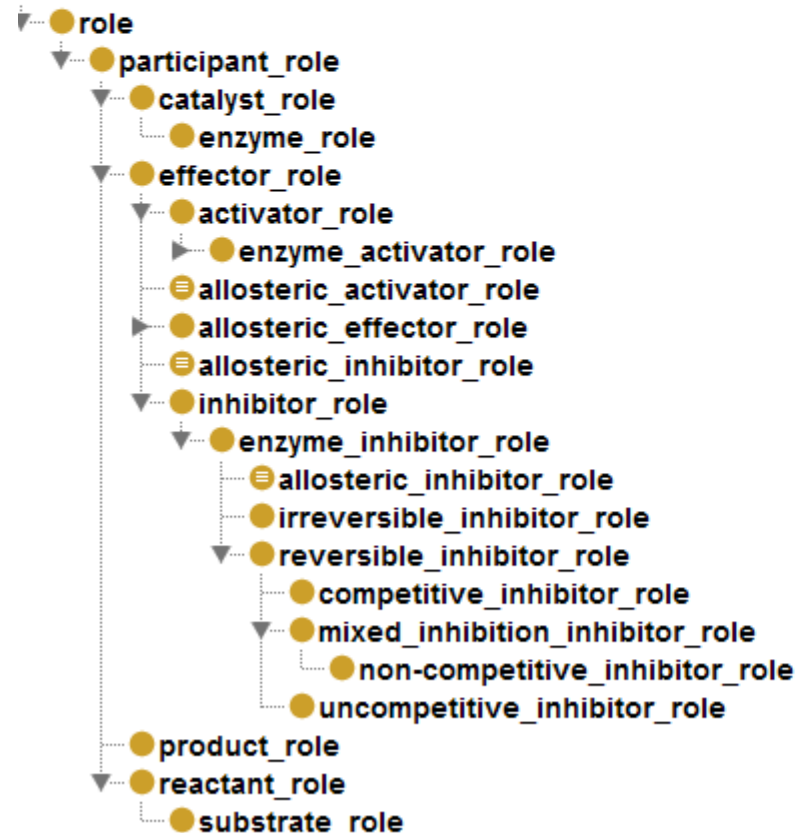
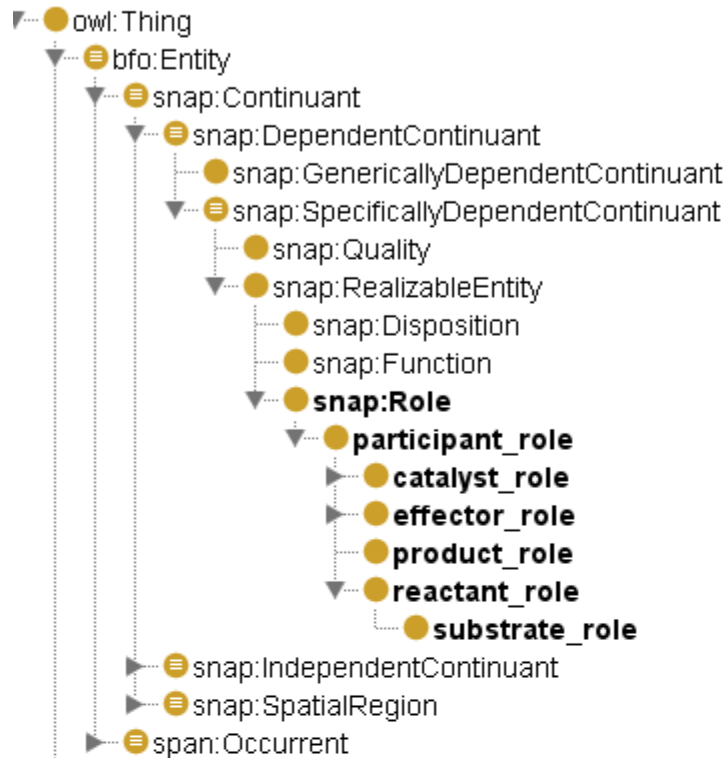
- computational approaches to study biochemistry require machine accessible representations of biochemical knowledge
 - existing representations adopt formal semantics to varying degrees
 - limited expressivity and ability for reasoning
-





- biochemical processes viewed as ***microscopic molecular events***
 - individuals of classes describing chemical species represent ***single molecules***
 - individuals of classes describing biochemical processes represent single molecular events, i.e. ***directed transitions of a chemical system from an initial to a terminal state*** involving individual molecules
-

Roles

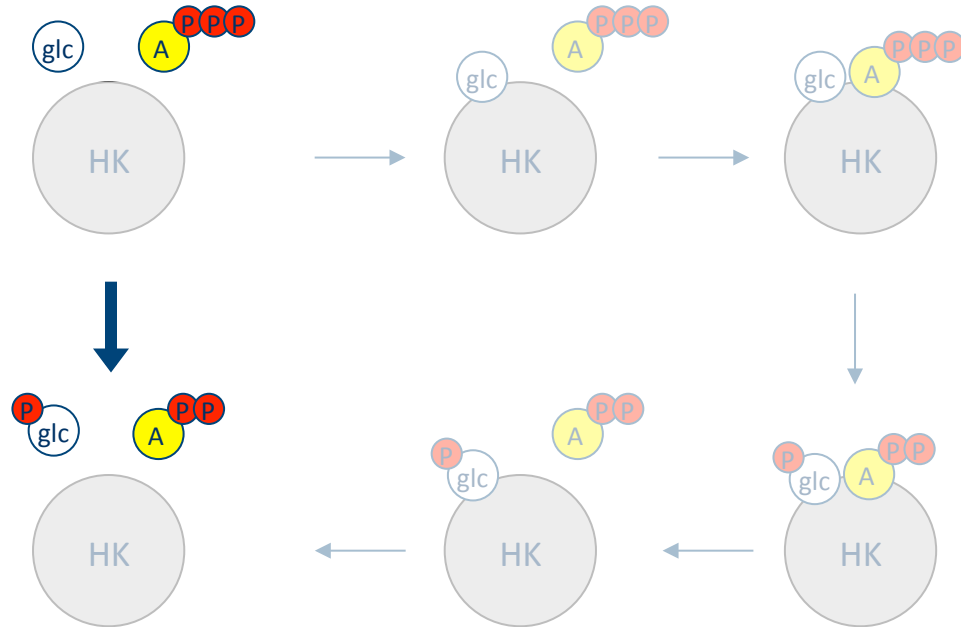


main axis of classification: stoichiometry

$aA + bB \rightarrow xX + yY$ EquivalentTo:
(realizes exactly a (reactant_role and (has_bearer some A))) and
(realizes exactly b (reactant_role and (has_bearer some B))) and
(realizes exactly x (product_role and (has_bearer some X))) and
(realizes exactly y (product_role and (has_bearer some Y))) and
realizes exactly (a+b) reactant_role and
realizes exactly (x+y) product_role

$glc + atp \rightarrow g6p + adp$ EquivalentTo:
(realizes exactly 1 (reactant_role and (has_bearer some glc))) and
(realizes exactly 1 (reactant_role and (has_bearer some atp))) and
(realizes exactly 1 (product_role and (has_bearer some g6p))) and
(realizes exactly 1 (product_role and (has_bearer some adp))) and
realizes exactly 2 reactant_role and
realizes exactly 2 product_role

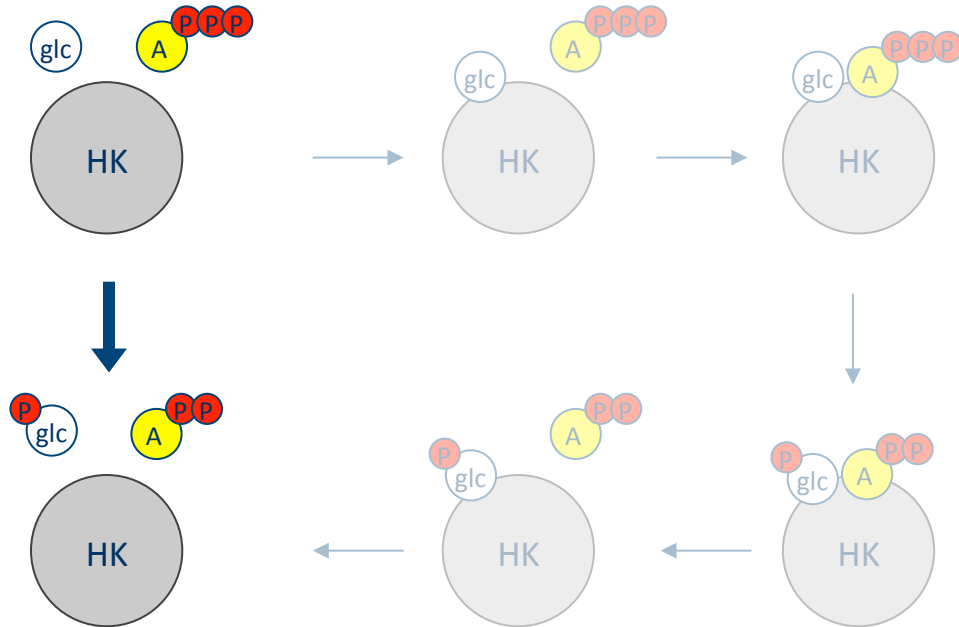
classification by overall stoichiometry



EquivalentTo:

(realizes exactly 1 (reactant_role and (has_bearer some glc))) and (realizes exactly 1 (reactant_role and (has_bearer some atp))) and (realizes exactly 1 (product_role and (has_bearer some g6p))) and (realizes exactly 1 (product_role and (has_bearer some adp))) and realizes exactly 2 reactant_role and realizes exactly 2 product_role

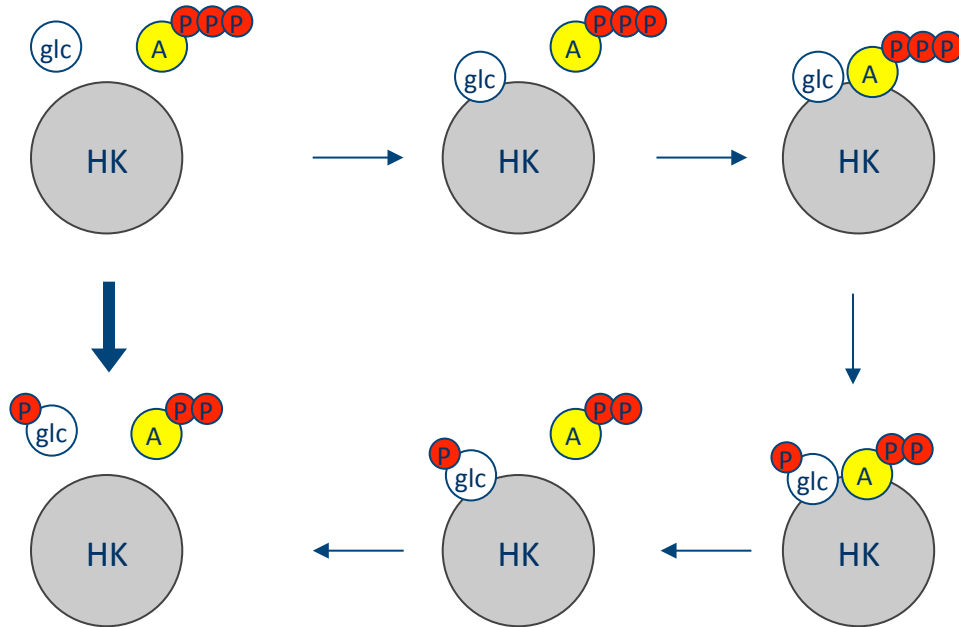
... and catalysis



EquivalentTo:

(realizes exactly 1 (reactant_role and (has_bearer some glc))) and
(realizes exactly 1 (reactant_role and (has_bearer some atp))) and
(realizes exactly 1 (product_role and (has_bearer some g6p))) and
(realizes exactly 1 (product_role and (has_bearer some adp))) and
realizes exactly 2 reactant_role and
realizes exactly 2 product_role and
(realizes exactly 1 (enzyme_role and (has_bearer some hk)))

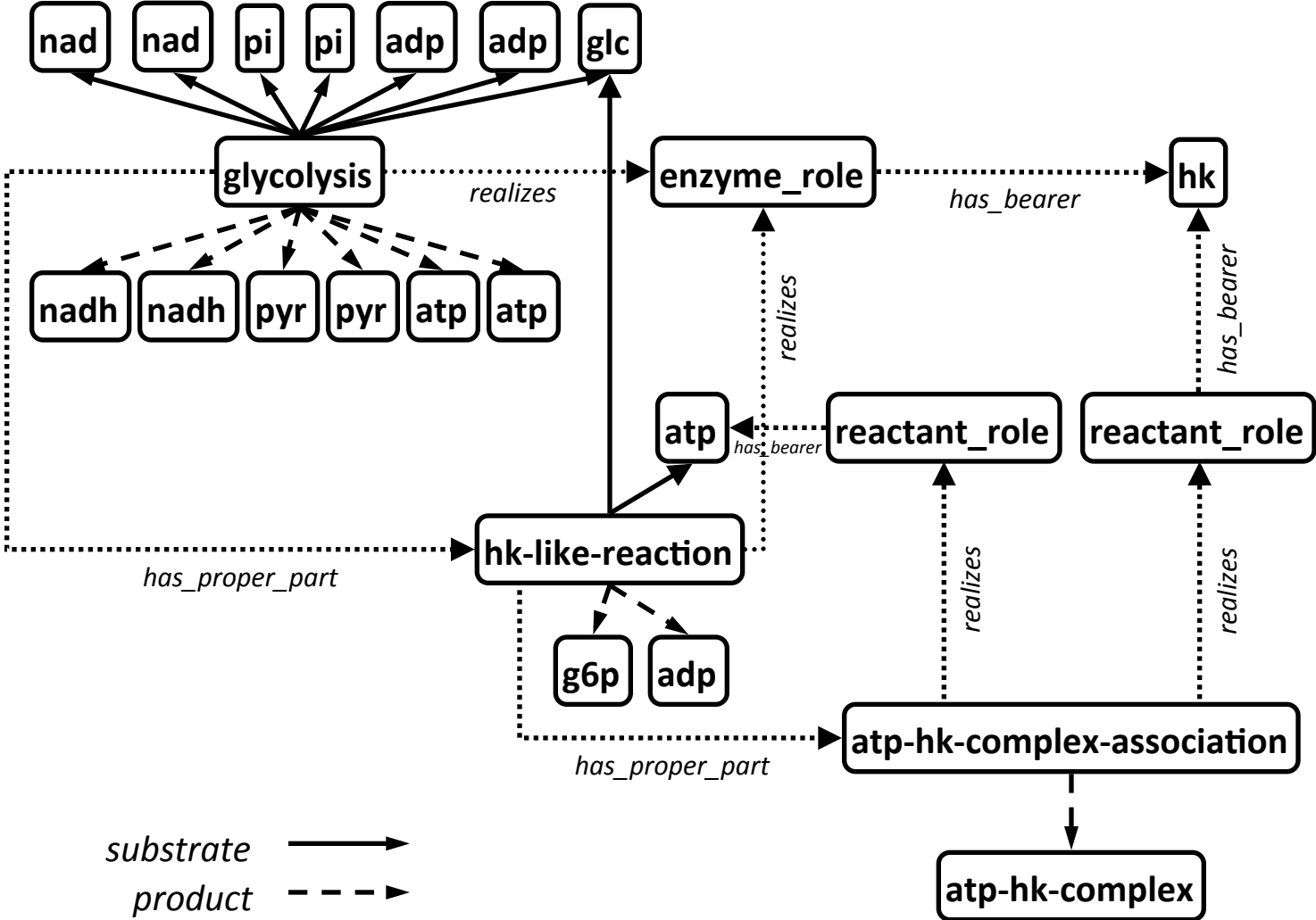
... and mechanism



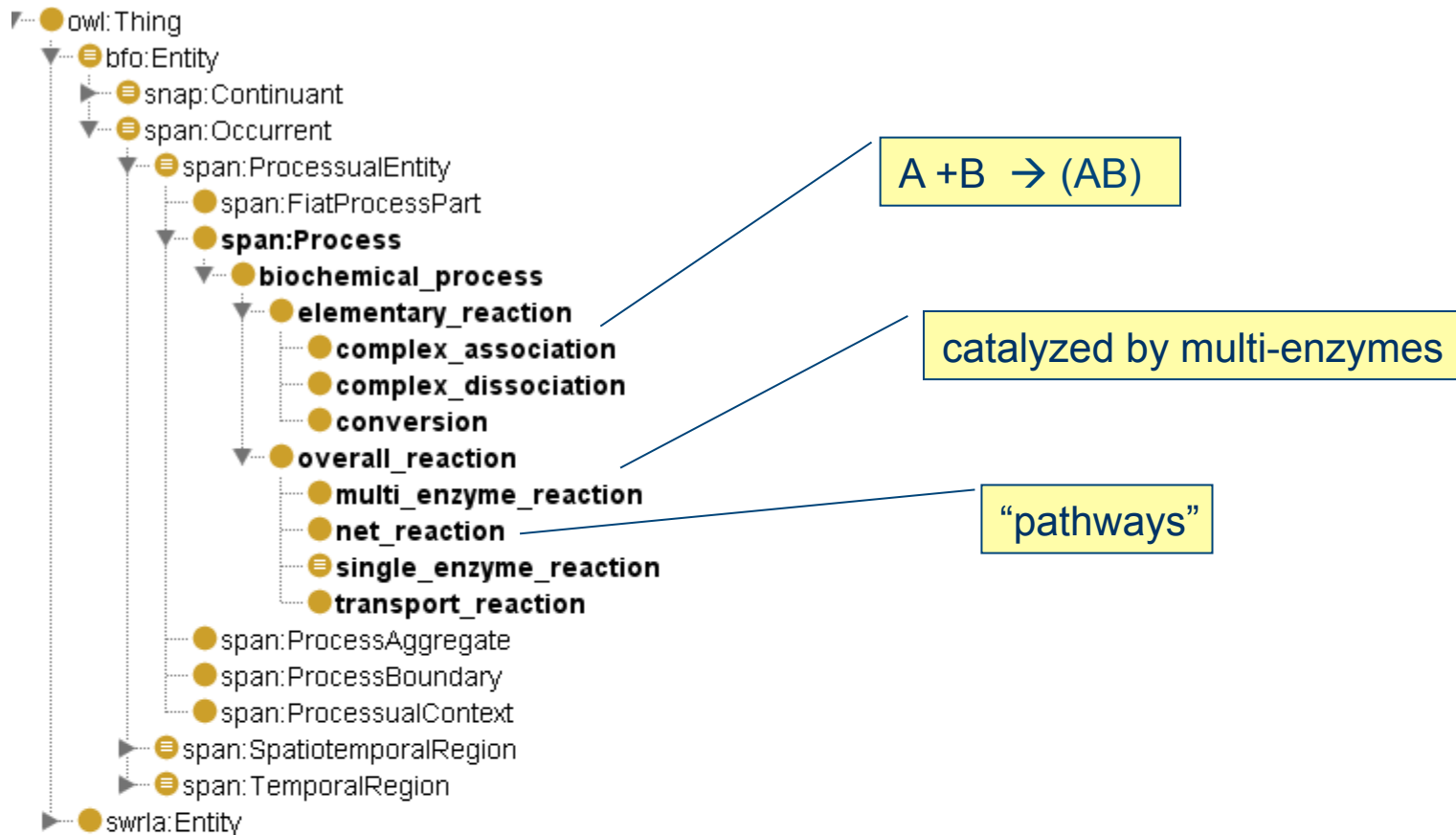
EquivalentTo:

(realizes exactly 1 (reactant_role and (has_bearer some glc))) and (realizes exactly 1 (reactant_role and (has_bearer some atp))) and (realizes exactly 1 (product_role and (has_bearer some g6p))) and (realizes exactly 1 (product_role and (has_bearer some adp))) and realizes exactly 2 reactant_role and realizes exactly 2 product_role and (realizes exactly 1 (enzyme_role and (has_bearer some hk))) and (has_proper_part ...)

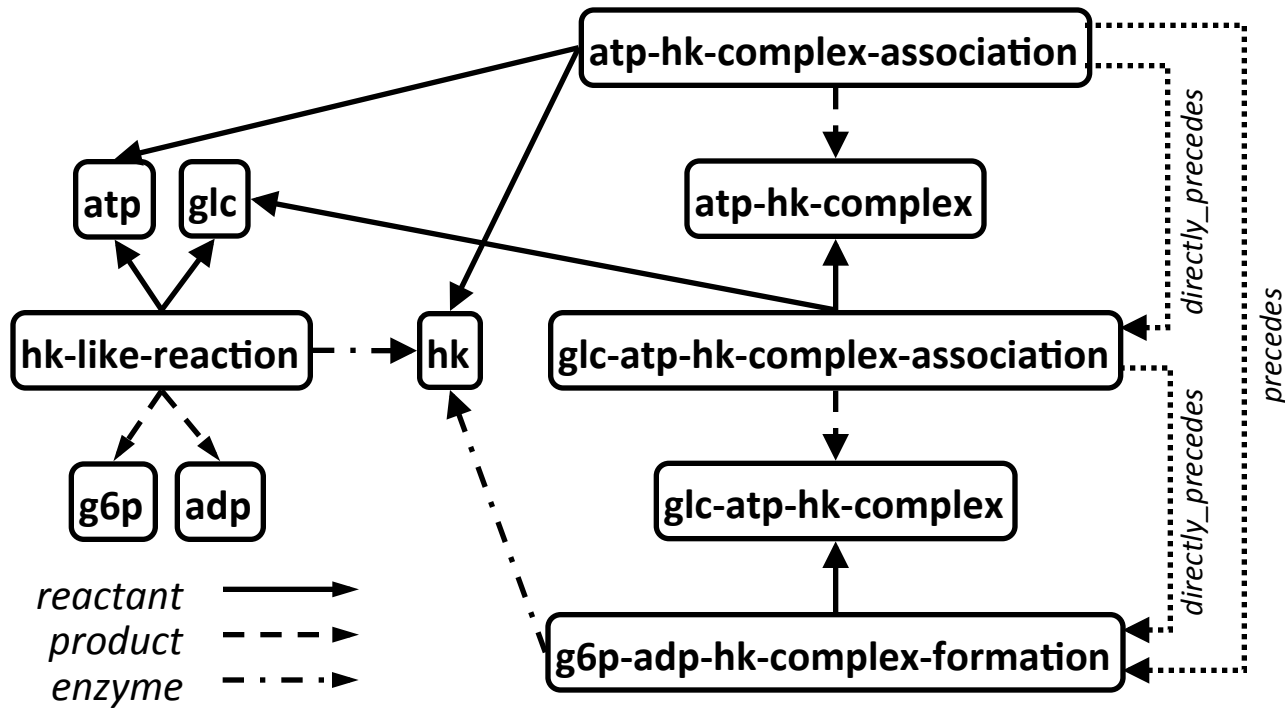
partonomy of processes



process types



shared reactants & products: sequence of processes



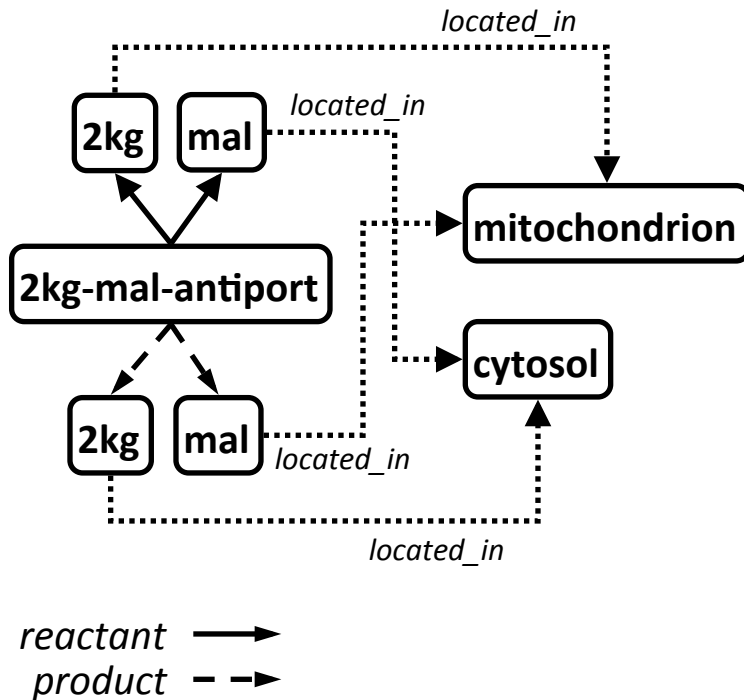
location of processes and transport

glc + atp → g6p + adp in the cytosol of human liver cells

EquivalentTo:

(realizes exactly 1 (reactant_role and (has_bearer some glc))) and
(realizes exactly 1 (reactant_role and (has_bearer some atp))) and
(realizes exactly 1 (product_role and (has_bearer some g6p))) and
(realizes exactly 1 (product_role and (has_bearer some adp))) and
(realizes exactly 2 reactant_role) and
(realizes exactly 2 product_role) and
(occurs_in some (cytosol and (part_of some (cell and (part_of some (liver and (part_of
some Homo_sapiens))))))

location of processes and transport



EquivalentTo:

(realizes exactly 1 (reactant_role and (has_bearer some (2kg and (located_in some mitochondrion)))) and (realizes exactly 1 (reactant_role and (has_bearer some (mal and (located_in some cytosol)))) and (realizes exactly 1 (product_role and (has_bearer some (2kg and (located_in some cytosol)))) and (realizes exactly 1 (product_role and (has_bearer some (mal and (located_in some mitochondrion)))) and realizes exactly 2 reactant_role and realizes exactly 2 product_role

inference of process and entity characteristics

The screenshot displays a class hierarchy viewer with two tabs: 'Class hierarchy' and 'Class hierarchy (inferred)'. The 'Class hierarchy (inferred)' tab is active, showing a tree structure of classes. The root is 'owl: Thing', which branches into 'bfo: Entity', 'snap: Continuant', and 'span: Occurrent'. 'span: Occurrent' further branches into 'span: ProcessualEntity', 'span: FiatProcessPart', and 'span: Process'. 'span: Process' branches into 'biochemical_process', 'elementary_reaction', 'f6p_atp_TO_fbp_adp', 'f6p_atp_TO_fbp_adp_catalyzed', 'g6p_TO_f6p', 'glc_atp_TO_g6p_adp', and 'overall_reaction'. 'f6p_atp_TO_fbp_adp' is highlighted in blue. Below it, 'f6p_atp_TO_fbp_adp_catalyzed' branches into 'f6p_atp_TO_fbp_adp_catalyzed_cytosolic', 'f6p_atp_TO_fbp_adp_catalyzed_cytosolic_human', and 'f6p_atp_TO_fbp_adp_catalyzed_cytosolic_liver_human'. 'span: ProcessAggregate' and 'span: ProcessBoundary' are also visible at the bottom of the tree.

Annotations Usage

Annotations: f6p_atp_TO_fbp_adp

Annotations +

Description: f6p_atp_TO_fbp_adp

Equivalent classes +

- (realizes exactly 1 (product_role and (has_bearer some adp))) and (realizes exactly 1 (product_role and (has_bearer some fbp))) and (realizes exactly 1 (reactant_role and (has_bearer some atp))) and (realizes exactly 1 (reactant_role and (has_bearer some f6p))) and (realizes exactly 2 product_role) and (realizes exactly 2 reactant_role))

Superclasses +

- biochemical_process

inference of process and entity characteristics

- location of reaction participants:
**biochemical_process(?p), occurs_in(?p,?l),
has_participant(?p,?o) -> located_in(?o,?l)**
 - sequence of processes within same location:
**product_role(?r1), reactant_role(?r2), has_bearer(?
r1,?o), has_bearer(?r2,?o), realizes(?p1,?r1),
realizes(?p2,?r2), occurs_in(?p1,?l), occurs_in(?p2,?l)
-> directly_precedes(?p1,?p2)**
-

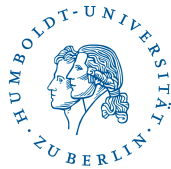
outlook

- integrate with ontological descriptions of chemical entities, cell-types, tissues
 - implement design patterns / rules for classification according to taxonomies of chemicals, locations, reaction mechanisms,...
 - add support for quantitative description of reactions / participants
 - expand on other molecular interactions
 - use for biochemical network reconstruction, integration of pathway datasets, inference of molecular associations...
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Acknowledgments



• Michel Dumontier



• Michael Weidlich



• Matthias König
• Andreas Hoppe
• Hermann-Georg Holzhütter



Federal Ministry
of Education
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