

# Tailoring the shape calculus towards quantitative analysis

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# Shape calculus - main features

- Non-deterministic timed calculus representing physical entities moving in 3D
- Processes = 3D shapes + dynamic behaviour
- Processes can move, collide and possibly bind
- Behaviours are specified with a timed CCS-like process algebra with channels aka “type of binders”
- $\langle a, X \rangle$ ,  $X$  is a portion of the surface of the process’s able to bind



# Shape calculus - main features

- Bound processes has behaviour equal to the interleaving of the component processes
- Compound processes can split weakly (no reaction) or split strongly (reaction) dividing in as many pieces as the reaction products
- Without communication (i.e. binding) collisions are considered elastic

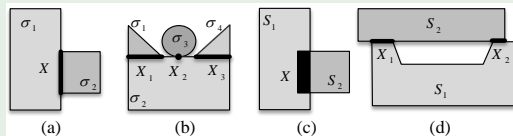


# Shapes in Shape Calculus

## Shape Syntax

$S ::= \sigma \mid S \langle X \rangle S$  where  $\sigma \in \text{Basic}$  and  $X \subseteq \mathbb{R}^3$  is a *non-empty* set of points. The set  $X$  is intended to be the common surface on which the two shapes are attached.

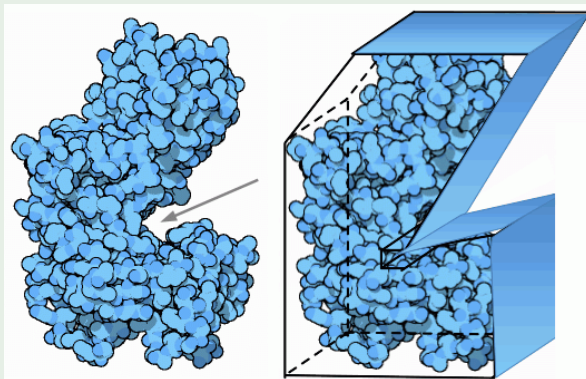
## Examples of compound shapes in 2D



# Shape example

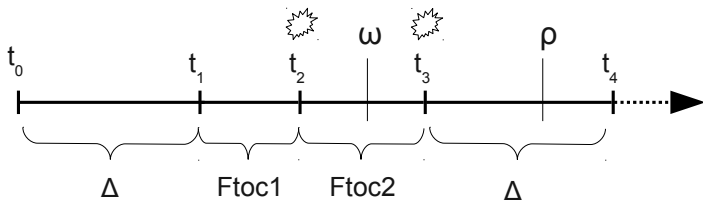
## Representation of enzymatic reaction in Shape Calculus

Shape approximation



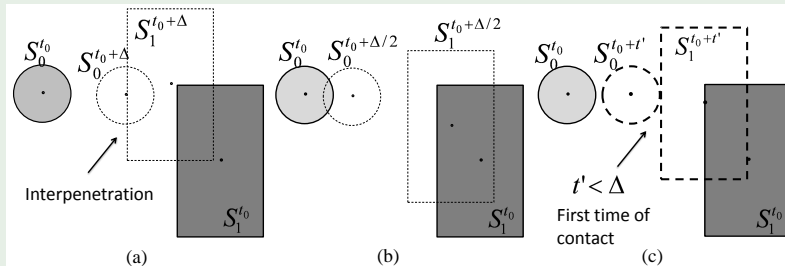
# Time evolution and velocity update

- Time domain  $\mathbb{T} = \mathbb{R}_0^+$  is then divided into an infinite sequence of movement time steps  $t_i$  such that  $t_0 = 0$  and  $t_i = t_{i-1} + \min(\Delta, Ftc(t_{i-1}), Ftr(t_{i-1}))$
- The updating of the velocities is represented by a function  $\text{steer}: \mathbb{T} \rightarrow \text{Shapes} \hookrightarrow \mathbb{V}$  gives the velocity vector  $\text{steer } t$   $S$  to assign to shape  $S$  at time  $t$



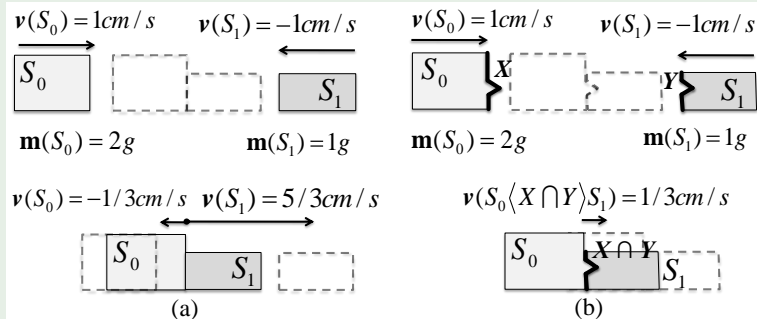
# Collision Detection

## First time of contact



# Collision Response

## Elastic and inelastic collision (one dimensional case)





# Shapes behaviours

## Set $\mathbb{B}$ of *shapes' behaviours* grammar

$$B ::= \text{nil} \mid \langle \alpha, X \rangle . B \mid \omega(\alpha, X) . B \mid \rho(L) . B \mid \epsilon(t) . B \mid B + B \mid K$$

where  $\langle \alpha, X \rangle \in \mathcal{C}$ ,  $L$  is a non-empty subset of  $\mathcal{C}$  whose channels are pairwise incompatible,  $t \in \mathbb{T}$  and  $K$  is a process name in  $\mathcal{K}$ .

## Set 3DP of *3D processes* grammar

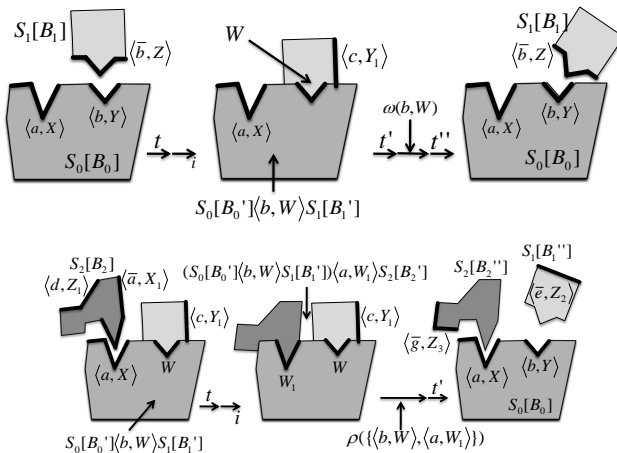
$P ::= S[B] \mid P \langle a, X \rangle P$ , where  $S \in \text{Shapes}$ ,  $B \in \mathbb{B}$ ,  $a \in \Lambda$  and  $X \subseteq \mathbb{R}^3$   
intersection between the surface active sites that are bound

## Modelling Hexokinase in Shape Calculus

$S_h[\text{HEX}]$  where  $\text{HEX} = \langle \text{atp}, X_{ha} \rangle . \text{HA} + \langle \text{glc}, Y_{hg} \rangle . \text{HG}$



# Binding and Splitting



# Modelling HEX, ATP and Glucose behaviours

$$\text{HEX} = \langle \text{atp}, X_{ha} \rangle . \text{HA} + \langle \text{glc}, X_{hg} \rangle . \text{HG},$$

$$\text{HA} = \omega(\text{atp}, X_{ha}) . \text{HEX} + \epsilon(t_h) . \langle \text{glc}, X_{hg} \rangle . \rho(\{\langle \text{atp}, X_{ha} \rangle, \langle \text{glc}, Y_{hg} \rangle\}) . \text{HEX},$$

$$\text{HG} = \omega(\text{glc}, X_{hg}) . \text{HEX} + \epsilon(t_h) . \langle \text{atp}, X_{ha} \rangle . \rho(\{\langle \text{atp}, X_{ha} \rangle, \langle \text{glc}, Y_{hg} \rangle\}) . \text{HEX},$$

where  $X_{ha}, Y_{hg}$  are the surfaces of contact.

$$\text{ATP} = \langle \overline{\text{atp}}, X_{ah} \rangle . (\epsilon(t_a) . \rho(\{\langle \overline{\text{atp}}, X_{ah} \rangle\}) . \text{ADP} + \omega(\overline{\text{atp}}, X_{ah}) . \text{ATP})$$



# Towards quantitative analysis

- Shape Calculus can represent a great variety of scenarios
- Soon it will be integrated with our tool BIOSHAPE for analyse biological phenomena
- Providing formal verification techniques for the Shape Calculus is the next step
- The objective is to find the right abstractions to apply existing quantitative model checking or quantitative equivalence checking techniques



## For instance...

- probabilistic timed automata could be useful to describe in more detail the behaviour of the processes and their interactions
- hybrid automata could be used to specify schemes of motion to be associated to certain classes of processes
- Suitable logic languages for specifying the properties must also be identified



# Logic(s) for...

- verifying that certain 3D configurations are reached
- verifying that a certain molecule concentration is achieved
- verifying occurrence of certain oscillatory behaviours
- verifying that with a certain probability a reaction can occur
- ...



In the end...

*...thanks for your attention! :)*

