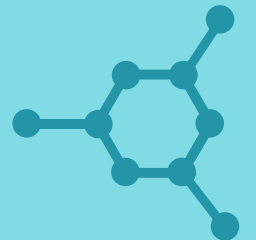
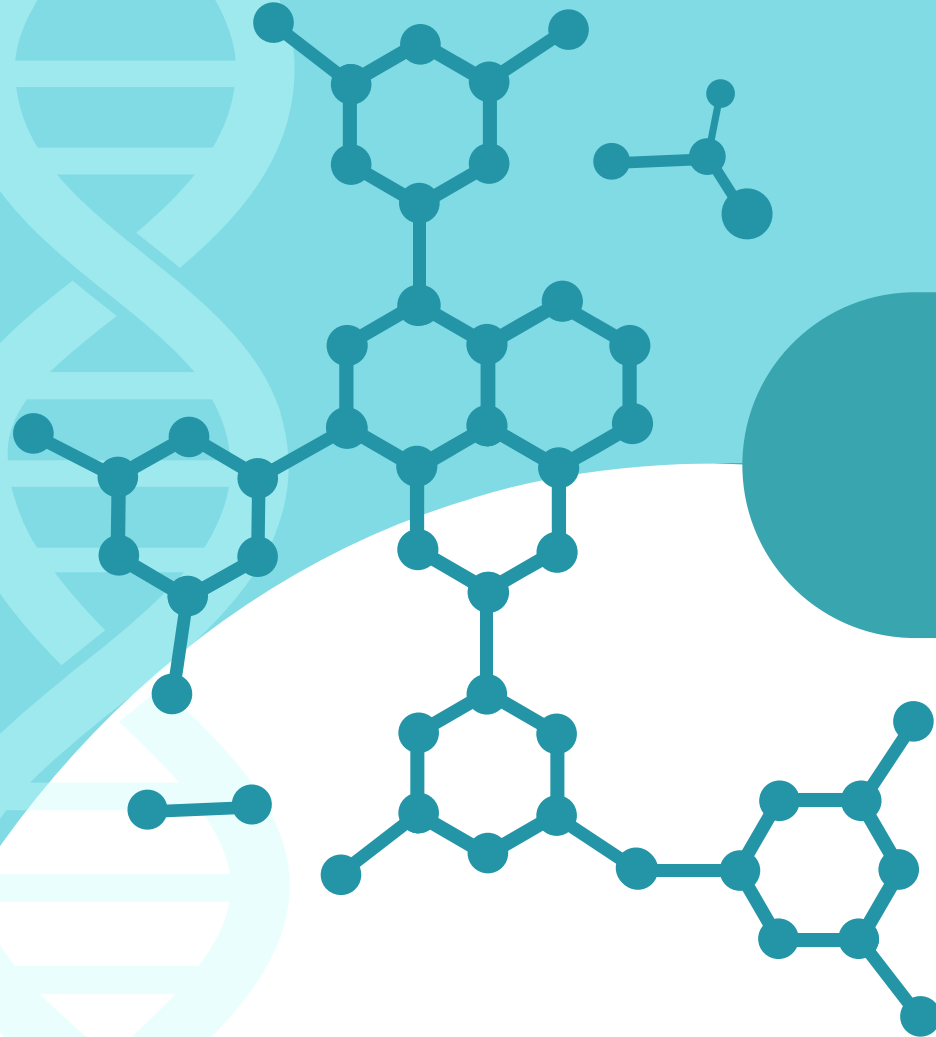




# Interactive models in genetics

Proteins misfolding to detect sickle-cell anemia disease

Stefano Maestri



# Genetic disorders

A disease caused in whole or in part by a change in the DNA sequence away from the normal sequence.

As we unlock the secrets of the human genome (the complete set of human genes), we are learning that nearly all diseases have a genetic component.

Some diseases are caused by mutations that are inherited from the parents and are present in an individual at birth, like *sickle cell disease*.

Other diseases are caused by acquired mutations in a gene or group of genes that occur during a person's life. Such mutations are not inherited from a parent, but occur either randomly or due to some environmental exposure (such as cigarette smoke). These include many cancers, as well as some forms of neurofibromatosis.

## Main causes

**mutations in multiple genes**

multifactorial inheritance disorder

**mutation in one gene**

monogenic disorder

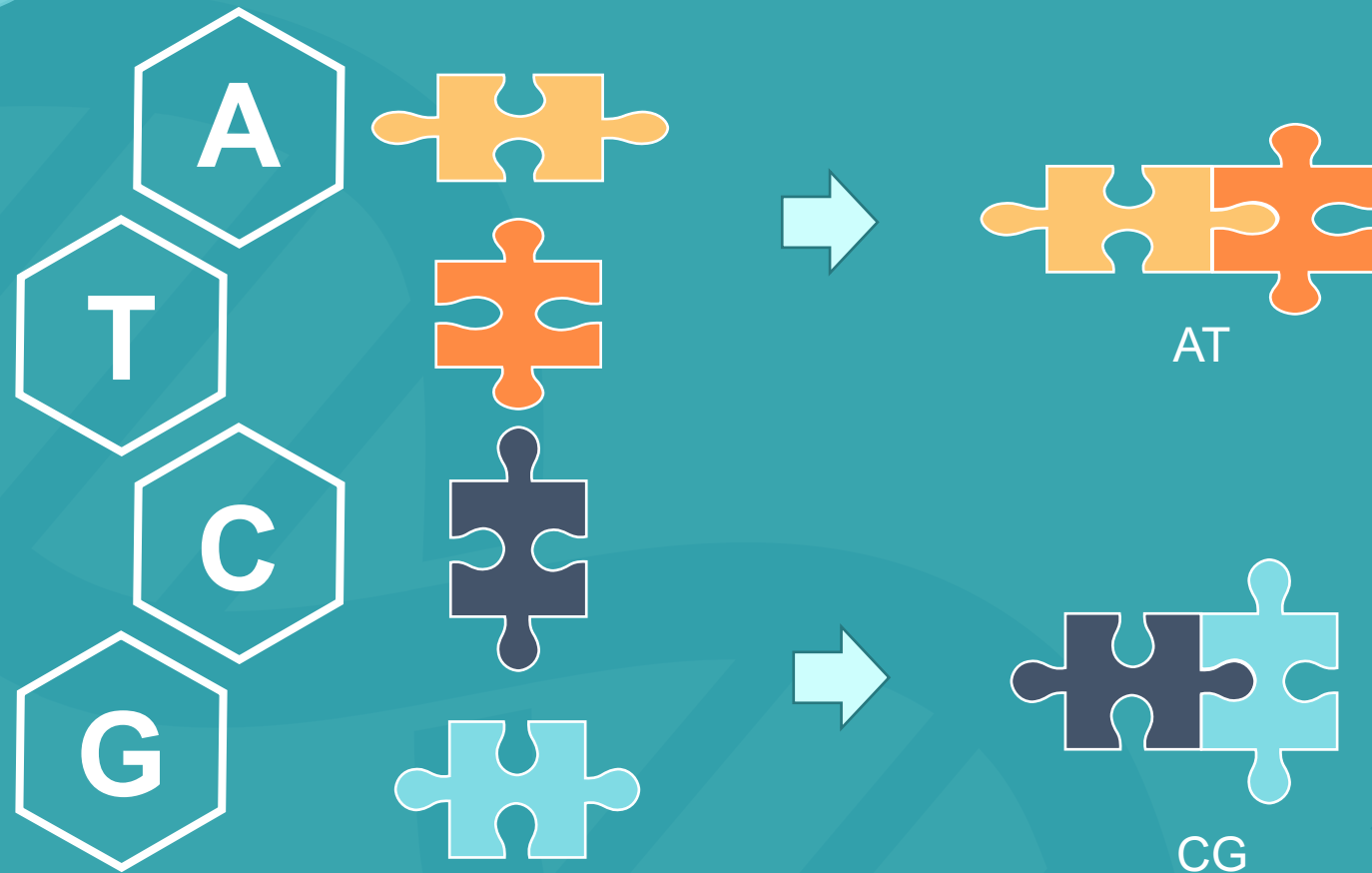
**combination of gene mutations and environmental factors**

**damage to chromosomes**

changes in the number or structure of entire chromosomes, the structures that carry genes

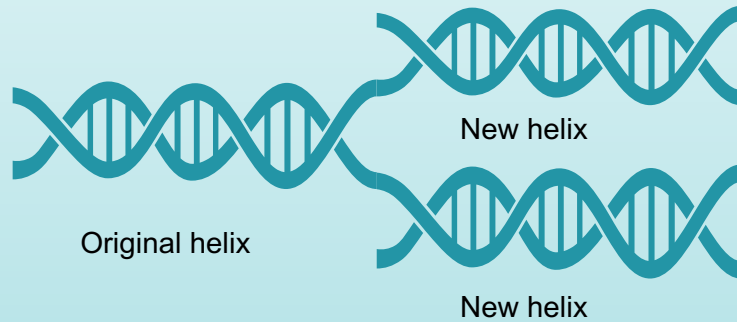
# DNA as a puzzle made of nucleotides

You can think to DNA as a double-helix of puzzle pieces. Each piece represents a nucleotide and can fit only with the complementary one on the other strand.



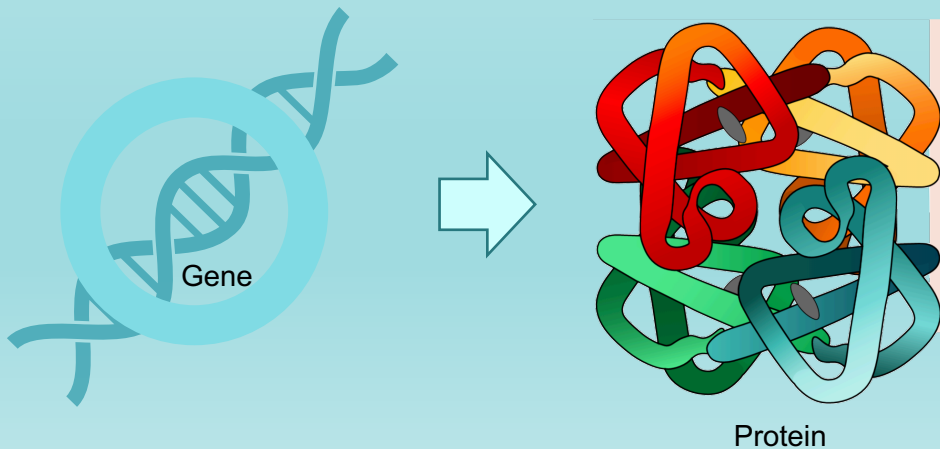
# DNA replication and gene expression

## DNA replication



Basing on nucleotides (puzzle pieces) complementarity, the DNA replication generates two new double helices identical to the original one

## Gene expression



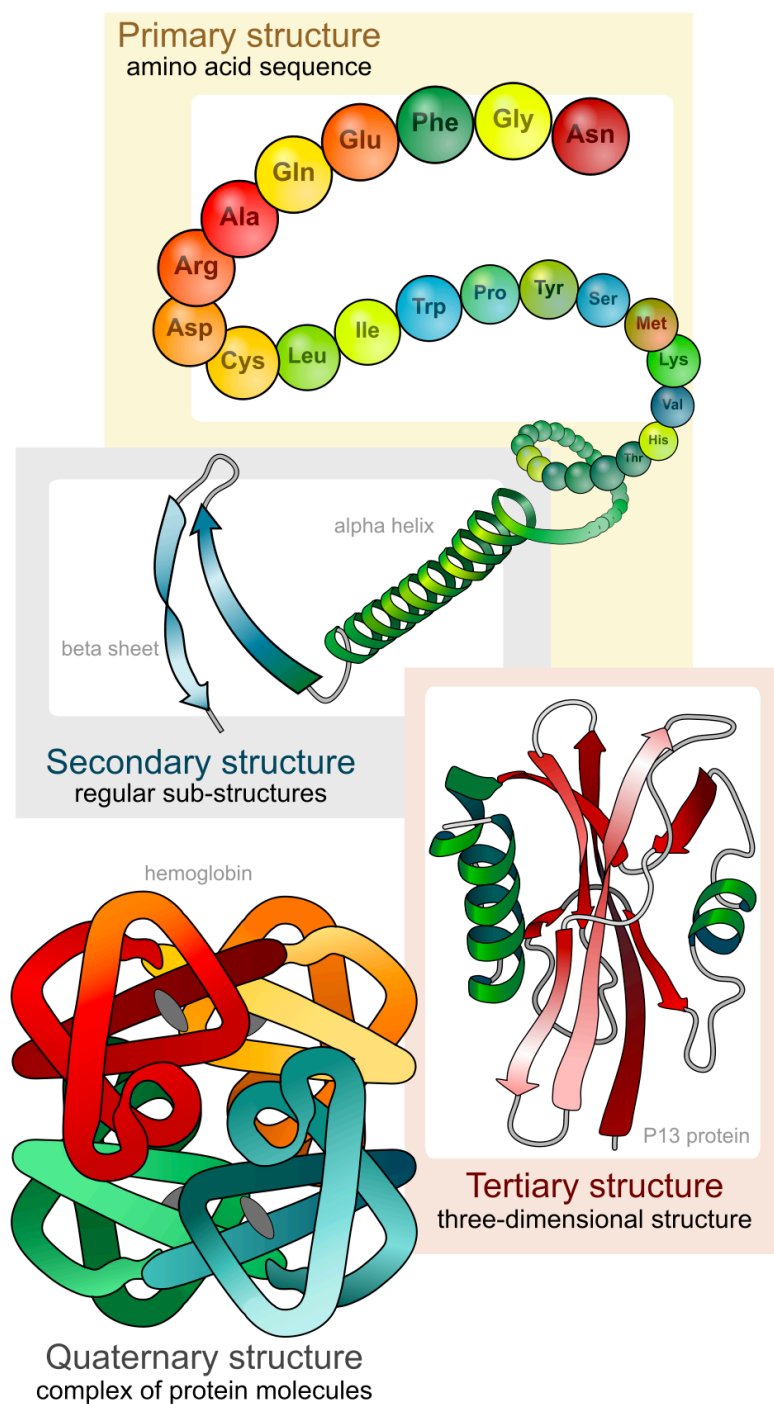
A portion of the DNA sequence (a gene) is used as instruction to produce a new protein

# Protein folding

A protein is made by a sequence of amino acids.

Differently from the pieces that compose DNA, amino acids do not have specific complementarities, but the order in which they are connected is determined during the gene expression, basing on the sequence of nucleotides of the related gene. For that reason we can say that a gene is the instruction for the production of a protein.

Energies of the environment in which the protein is placed and specific properties of each amino acid push the protein to fold into various substructures and eventually reach its final conformation.



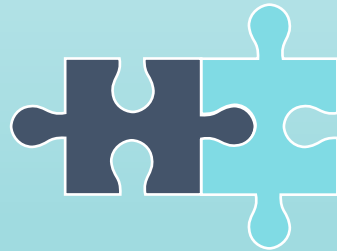
# What is a mutation?



A mutation is a change of the correct nucleotide sequence in a gene. It is often transmitted by inheritance or is caused by an error during the gene replication or expression.

A wrong gene sequence represent an incorrect instruction for the generation of a new protein.

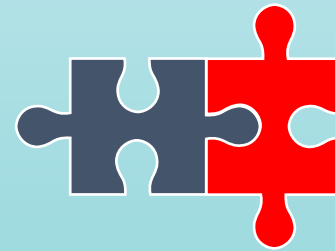
A mutation can affect even a single nucleotide (point mutation).



correct pairing



mutation



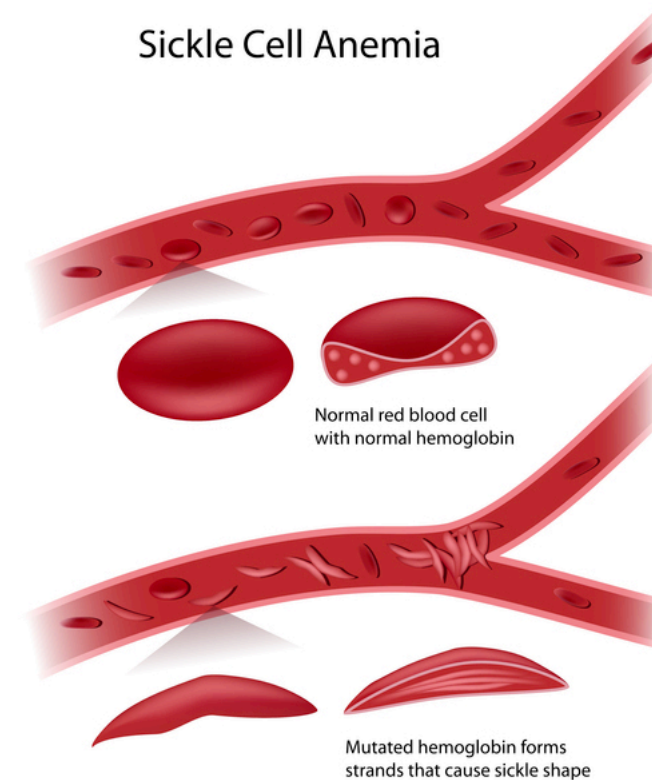
wrong pairing

# Sickle cell disease

Sickle cell disease is a group of disorders that affects hemoglobin, the molecule in red blood cells that delivers oxygen to cells throughout the body. People with this disorder have atypical hemoglobin molecules called hemoglobin S, which can distort red blood cells into a sickle, or crescent, shape.

The hemoglobin S is generated by a point mutation of the related gene, which cause change the production of a wrong amino acid in the protein sequence.

The properties of this amino acid cause the molecule to produce a patch that fits into a pocket of another hemoglobin molecule. This process forms fibrous precipitates, which produce the characteristic sickle shape of the affected red blood cells.

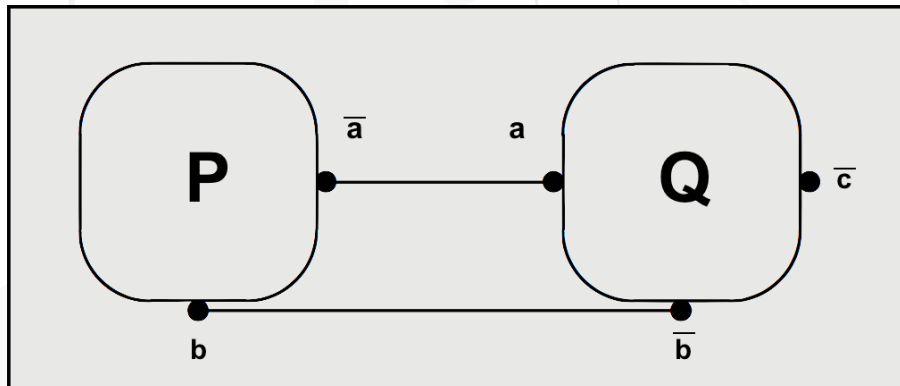


# Modeling the gene expression

## Theoretical tool: Process algebras

Collection of operations for building a new process description from existing ones.

A process can be viewed as a system that exhibit a behavior, a black box with a name and a set of communication channels.



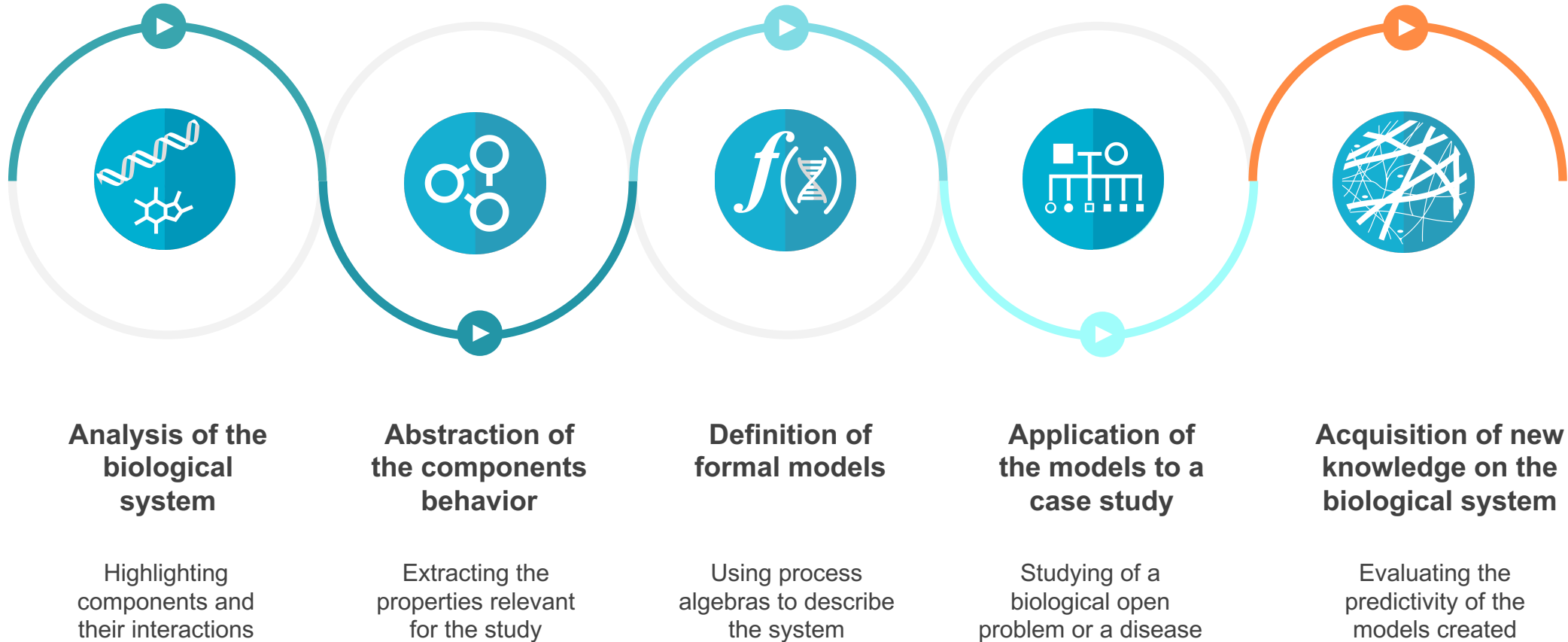
P and Q are processes that communicate with the channels labeled  $a$  and  $b$ ; Q may communicate with a third process via the channel labelled  $c$

## Core idea

Entire molecules, their substructures or even their elementary units as well as their interactions can be modelled as processes. In this way it is also possible to identify similarities among different classes of molecules and in the functions they carry out.



# Modeling Phases



# Results and aims of our approach

