## Constraint Programming and Biology: Introduction

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ACP Summer School in Constraint Programming Wrocław, September 2012

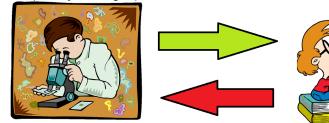
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- Biology is an incredible source of challenging problems for computer science
- Problems are often hidden or confused and emerge only after long discussions with biologist, physics, chemists, physicians, and so on (briefly, biologist)

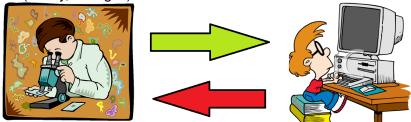


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#### Overview

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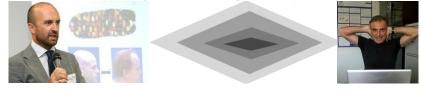


 Solving one of these problems can be of unpredictable importance for life sciences and medicine

 Some problems are of little interest for computer science but of great importance for biologist (eg developing scripts for automatization of sequences of simple tasks).

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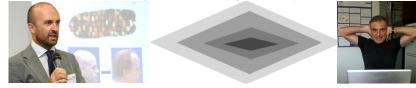
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• Some problems are polynomial time solvable but the input size is huge (e.g., a DNA string). These problems require fast string matching algorithms. They are important and challenging but the constraint programming approach is not the best suited for them.

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- Some problems are polynomial time solvable but the input size is huge (e.g., a DNA string). These problems require fast string matching algorithms. They are important and challenging but the constraint programming approach is not the best suited for them.
- ✓ We will not deal with the two kinds of problems above in these lectures

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- We love studying and solving these simplified models and at the end we win (at least for little inputs, and using some "reasonable" heuristics) but the risk is that with these simplifications our solutions are useless for biologists.
- We will focus on this family. CP techniques are perfect for NP problems. And sometimes our solutions are not useless! 4 3 5 4 3 5 5

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#### **Problems for Bioinformatics**

Bioinformatics can be seen as the area of computer science that deal with modeling and solving problems for Biology.

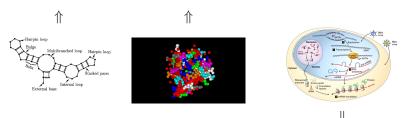
We have several families of problems.

- Those concerning DNA and genes
- Those concerning the transcription DNA → RNA and the structure of RNA
- Those concerning the translation RNA → proteins and the structure of proteins
- Those concerning the interaction between molecules and the behavior/interaction of systems of molecules (e.g. cells), till the modeling of living organisms.

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#### **Areas of Bioinformatics**

- Genomics. Study of the genomes. Huge amount of data, fast algorithms (not always), limited to sequence analysis.
- Structural Bioinformatics. Study of the folding process of bio-molecules. Less structural data than sequence data available.



Systems Biology. Study of complex interactions in biological systems. High level of representation.

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#### Why Constraint Programming?

(At least) two main reasons:

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• Models are rarely stable (and also the problems change quickly). Modifying a CP-modeling is easy and fast.

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- Models are rarely stable (and also the problems change quickly). Modifying a CP-modeling is easy and fast.
- Linear Programming is not enough (in particular for modeling energy models)

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#### What we'll see in mode details

We'll focus on some challenging problems and how modeling them using constraints:

- Genomics:
  - Haplotype Inference
  - Phylogenetic trees
- Systems Biology:
  - Reasoning on Biological Networks
- Structural Bioinformatics:
  - RNA secondary structure prediction
  - protein structure prediction (on/off lattice)
- ⇒ For these problems I have prepared the encodings in CLP(FD) (tested with BProlog—free). Link in my home page.

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#### Some introductory references

- P. Clote and R. Backofen. *Computational Molecular Biology*. An Introduction. Wiley, 2000.
- Nice introductory slides by Sebastian Will (MIT) http: //math.mit.edu/classes/18.417/Slides/intro.pdf
- A movie on DNA replication http://www.youtube.com/watch?v=teV62zrm2P0
- A movie on DNA transcription http://www.youtube.com/watch?v=5MfSYnItYvg
- A movie on Protein synthesis http://www.youtube.com/ watch?v=lpb5s2F1pyM&feature=related
- A movie on Systems Biology http://www.youtube.com/ watch?v=HNP1EAYLhOs&feature=fvwrel

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# Some references on Constraints and Bioinformatics

- P. Barahona, L. Krippahl, and O. Perriquet. *Bioinformatics: A Challenge to Constraint Programming*. In Hybrid Optimization The Ten Years of CPAIOR, Springer, 2011.
- Workshops on Constraint-based methods for Bioinformatics: WCB05 (Sitges), WCB06 (Nantes), WCB07 (Porto), WCB08 (Paris), WCB09 (Lisbon), WCB10 (Edinburgh), WCB11 (Perugia), WCB12 (Budapest).

Formerly: Workshops on Constraints and Bioinformatics/ Biocomputing in CP'97 and CP'98.

- Constraints, Volume 13. Special Issue on Bioinformatics and Constraints, 2008.
- Algorithms for Molecular Biology 7:15–17 (Thematic Series of AMB on Constraints and Bioinformatics), 2012.

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## Acknowledgments

(in advance)

- School organizers: ACP, Krzysztof Apt, Witold Charatonik, Leszek Pacholski, ...
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