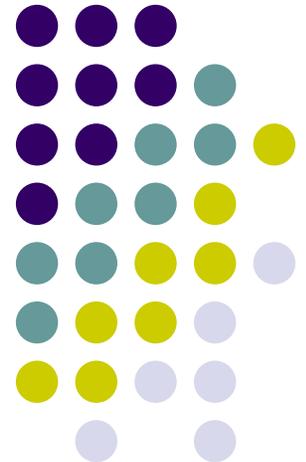


Current characteristics and historical perspective of CS & IT with/for Biology

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Overview



- Introduction
- Historical aspects
- Characteristics
- Brief introduction seminar series programme

Overview



- **Introduction**
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Introduction

- Disclaimer
 - My background
 - High-level overview (tech details in other seminars)
- CS/IT *for* Biology \Rightarrow support for biosciences
- CS/IT *with* Biology \Rightarrow 'new' discipline?
- Biology *for* CS/IT??

Introduction: some terms



Computational Biology

(Meta)Genomics,
Proteomics,
Metabolomics

Theoretical Biology

Ecoinformatics

Computational Chemistry

Environmental engineering

In Silico (molecular) Biology

Mathematics and Biology

Biocomputing

Climate modelling

Agricultural Informatics

Geographical Information Systems

Medical Informatics

Theoretical Ecology

Biomedical Engineering

Bioinformatics

Nanotechnology

Bio-ontologies

Systems Biology

Overview



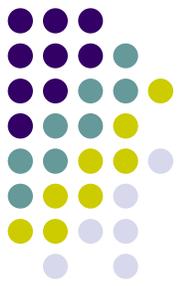
- Introduction
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Historical aspects

- $\pm \geq$ '30s: Systems biology (maths & eco/cellbio)
- $\pm \geq$ '50s:
 - Biomechanics/biomedical engineering
 - Analysis machines (hw/sw)
- $\pm \geq$ '70s:
 - Climate simulations (supercomputers)
 - Agricultural systems (hw/sw)
 - Medical informatics (DSS)
- \pm early '90s: start molbio-DBs





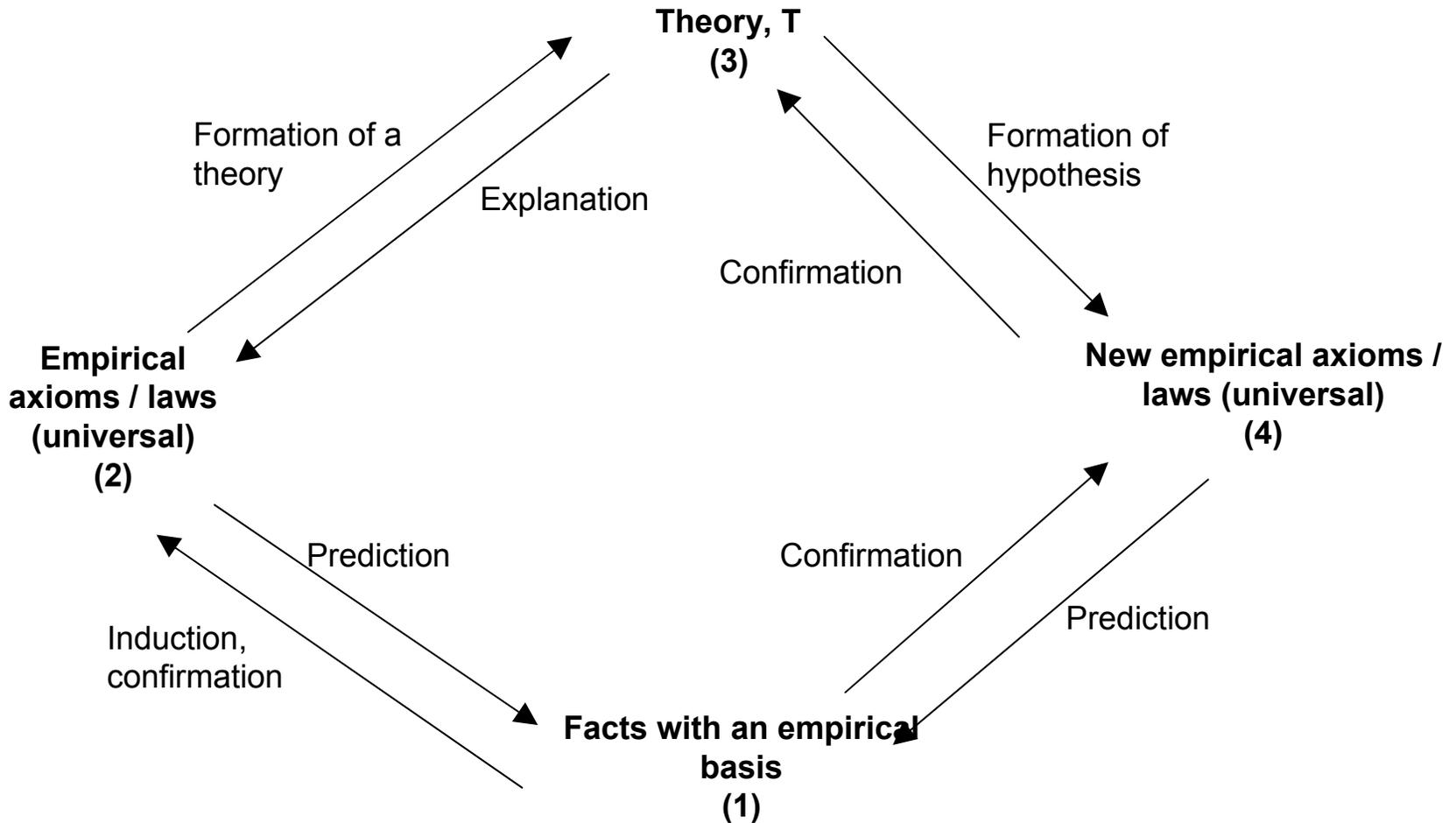
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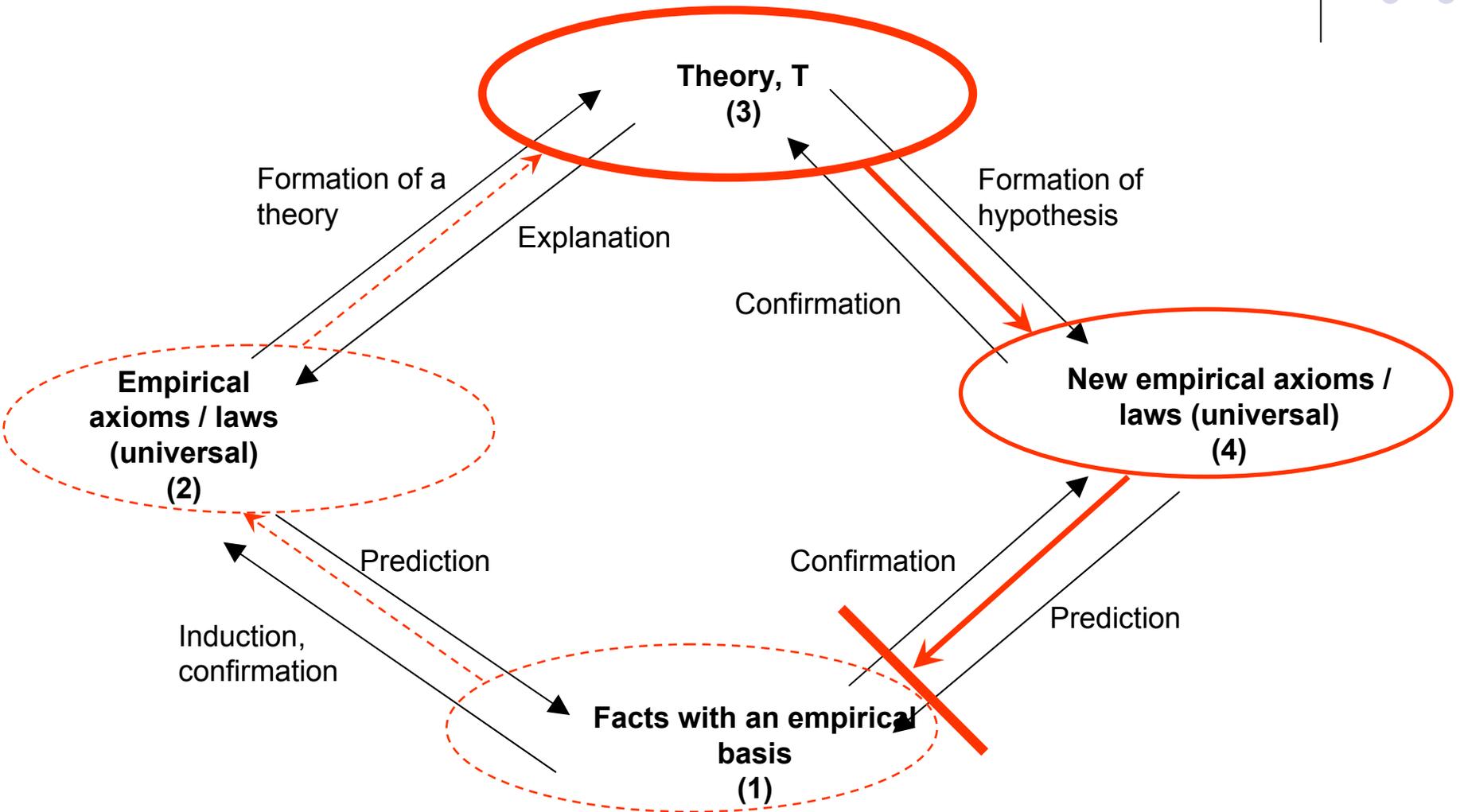


-----MID '90s EXPLOSION-----

Historical aspects



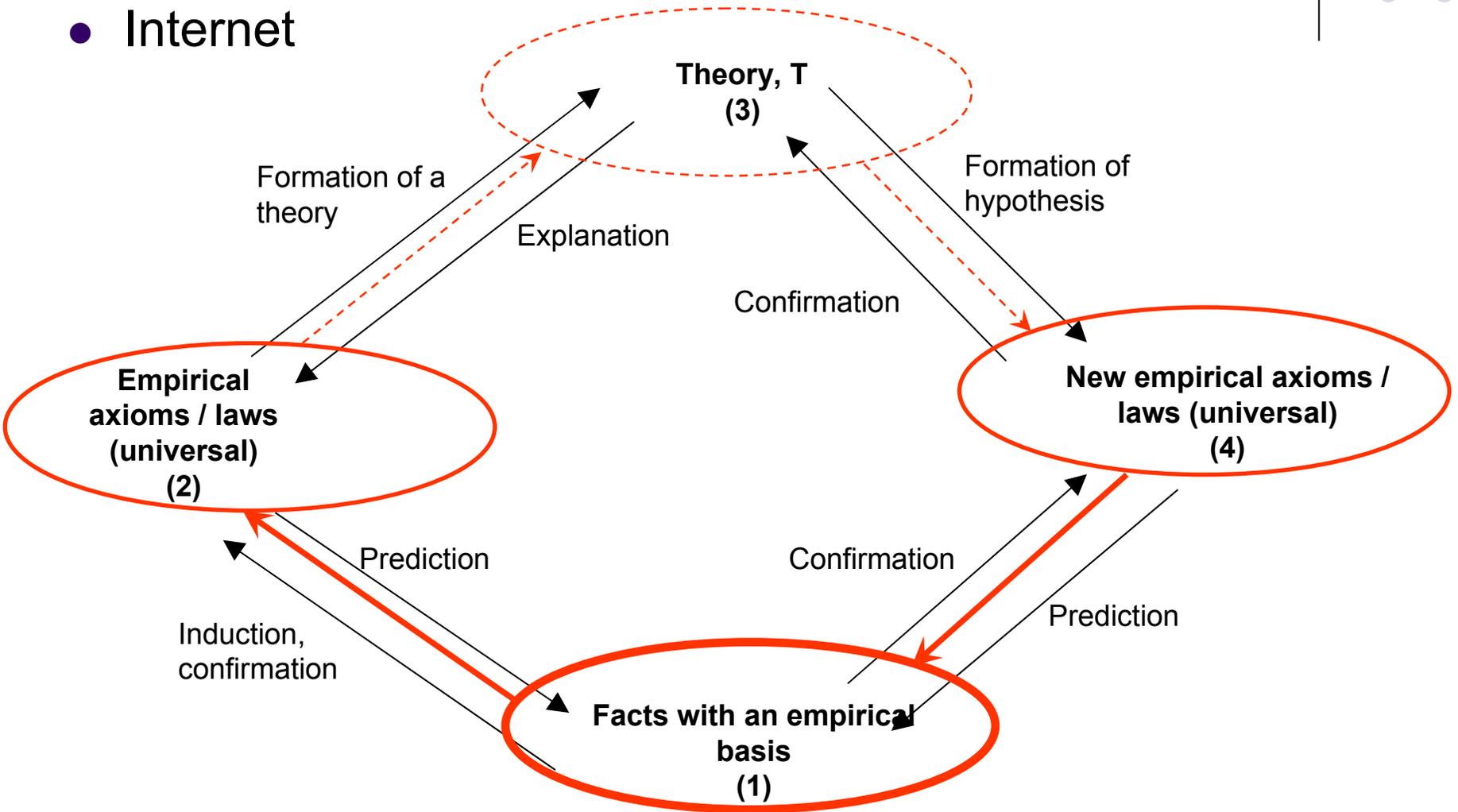
Historical aspects: < mid '90s



Historical aspects: \geq mid '90s



- Cheaper computers
- Internet

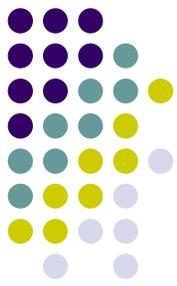


New possibilities for biology, in particular *molecular* biology



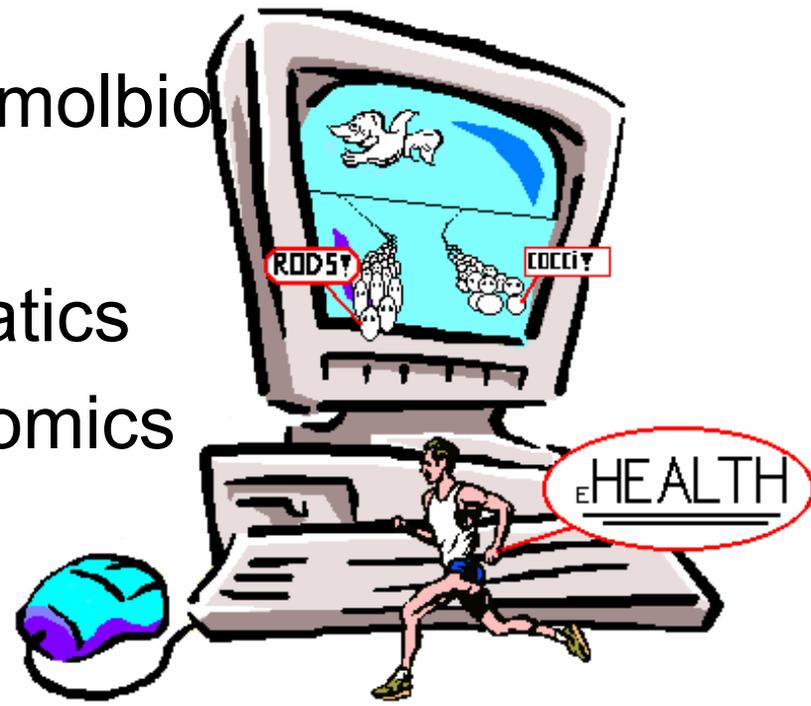
Historical aspects

- $\pm \geq$ mid/late '90s: sequence comparison algorithms, more databases, database integration (DWH, FD), software tools
- 1998: start Gene Ontology Consortium
- $\pm \geq$ '00s: bio-ontologies, computational linguistics (both NLP and NLG), data mining and pattern finding, Grids, workflow, 'hypothesis generation'



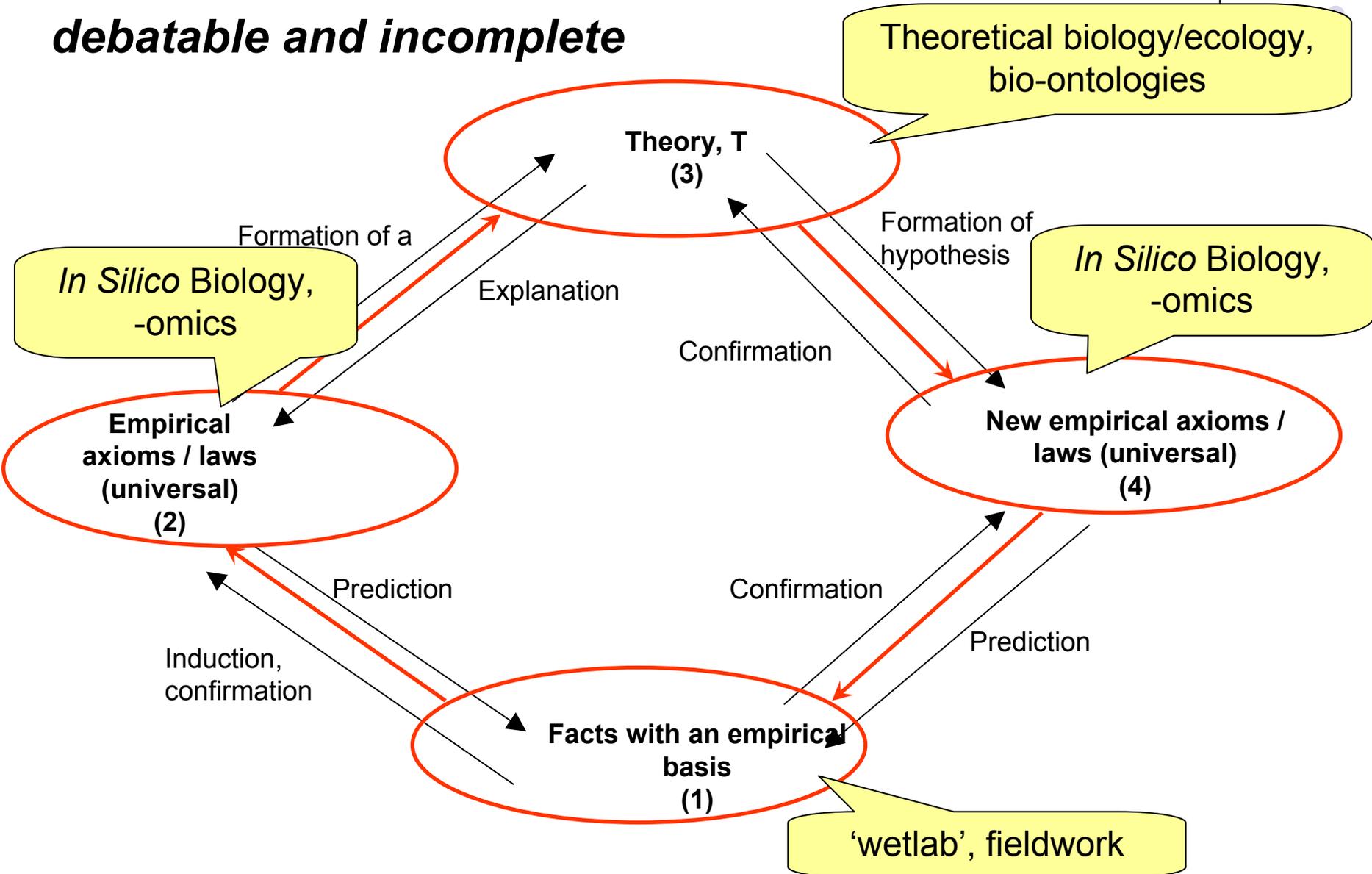
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- bioinformatics: from bio to molbio to biomed (OBO vs. OBO).
Counter-action: ecoinformatics
- systems biology, metagenomics



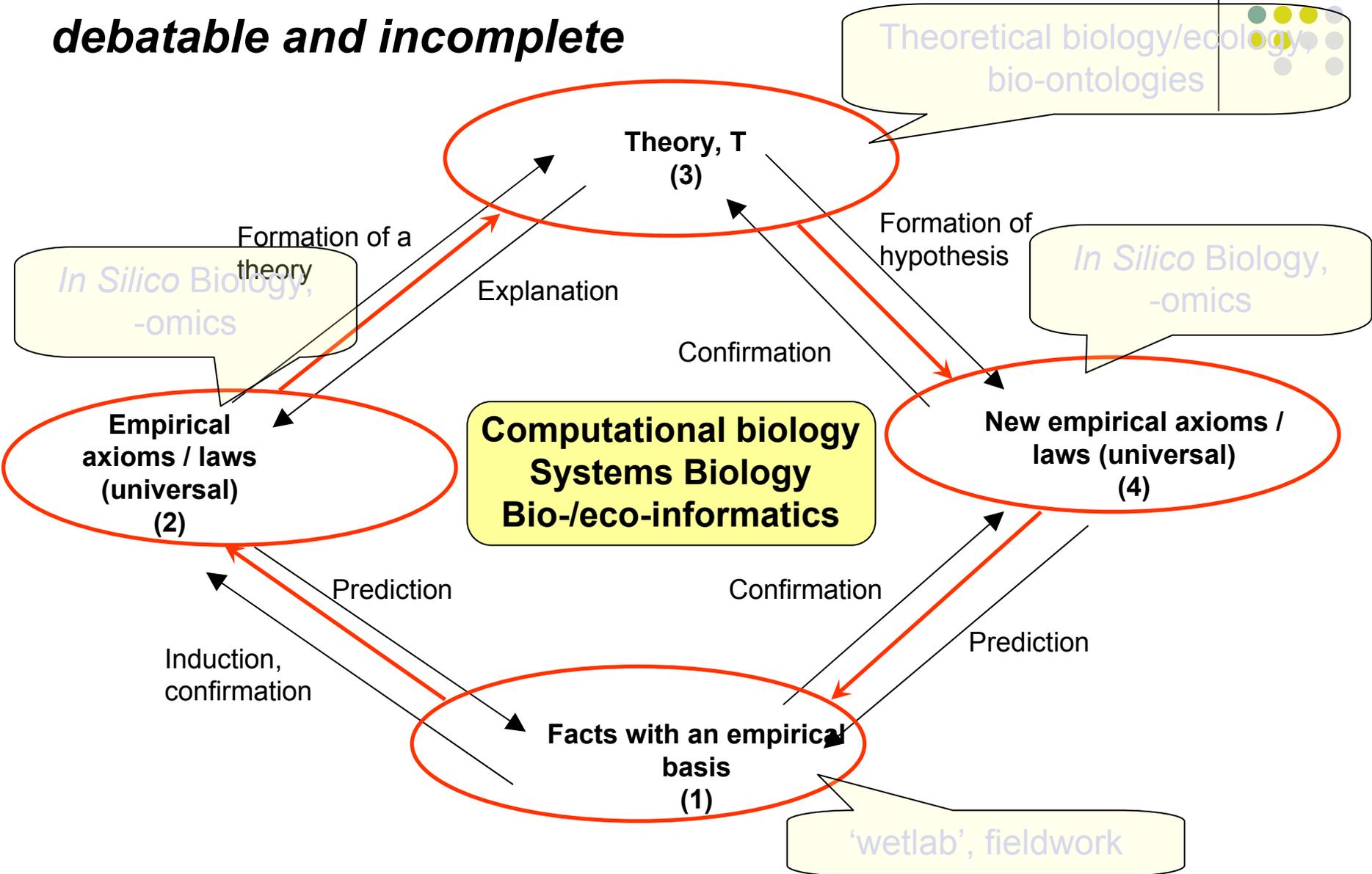
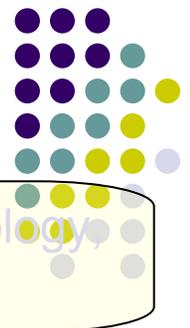
Historical aspects

A few terms
debatable and incomplete



Historical aspects

A few terms
debatable and incomplete



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Characteristics

- (some of the) CS/IT disciplines:

Hardware/Firmware: robotics, grid computing & supercomputing, analyzers

Software: Software engineering (devel., programming), Workflows, Databases (CM, DB devel., integration, temporal DB), distributed processing, Graphics & visualisation, HCI, comp. linguistics, neural networks, KBS, data mining



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- (some of the) Life sciences disciplines:

Core sciences:

Biology: virology, microbiology (bacteriology, fungi), plant sciences, animal sciences (entomology, nematology, ornithology, anatomy, ethology...), taxonomy

Molecular Biology & Chemistry: biochemistry, enzymology, cell physiology, genomics, proteomics, metabolomics, system biology

Ecology & environmental sciences: theoretical and experimental ecology (trophic levels, nutrient cycles, the niche), climatology, system biology

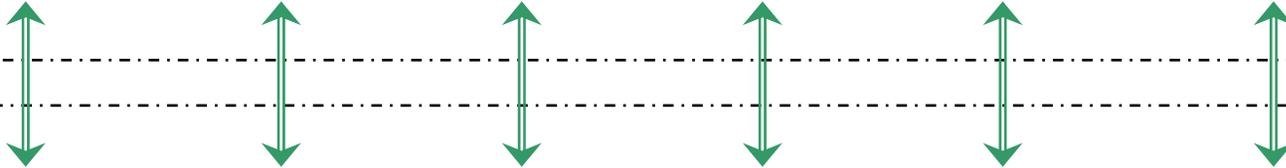
Applied Sciences: biomedicine, agriculture, food science

Characteristics



Combine any you like from the two boxes 😊

robotics, grid computing & supercomputing, analyzers, software engineering (devel., programming), Workflows, Databases (CM, DB devel., integration, temporal DB), distributed processing, Graphics & visualisation, HCI, comp. linguistics, neural networks, KBS, data mining



virology, microbiology (bacteriology, fungi), plant sciences, animal sciences (entomology, nematology, ornithology, anatomy, ethology, ...), taxonomy, biochemistry, enzymology, cell physiology, genomics, proteomics, metabolomics, system biology, theoretical and experimental ecology (trophic levels, nutrient cycles, the niche), climatology, system biology, biomedicine, agriculture, food science



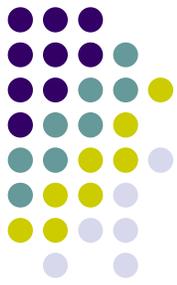
Characteristics

- Is it *inter*disciplinary?

interdisciplinary **teams** vs. interdisciplinary **people**



“An interdisciplinary team is a committee in which members identify themselves as an expert in something else besides the actual scientific problem at hand, and abdicate responsibility for the majority of the work because it's not their field.” [Eddy05]



Characteristics

.... Then, what you get:

- **Many bio-databases:** topical (one or two granularity levels, GOLD, HGVBBase), “species” specific (FlyBase, AceDB), context (Bad Bug Book), primary source vs. boutique DBs (TIGR), ... see also [Galperin05]
- **Many single-issue software tools**, mainly for data analysis of DB content, visualisation, some simulation
- High degree of **autonomy** of development and maintenance of IT tools, most tools poorly maintained, some databases better maintained than others
- **Sub-optimal data management:** data is ill structured, reliability of data becoming an issue, data redundancy, data incompatibility (between DBs and between DB and analysis tools)
- Results reported in many different kinds of journals and conferences: **where & how to find the right info?**
- Education did not keep up (biologist who can program a bit, computer scientist who googles for bio info)
- Differences in paradigms, research approaches, cultures, between science – engineering:
 - CS/IT-based (the motivating/toy example) vs. bio-based (fixes/addressed *this* issue/little problem)
 - Theory - Experiment
 - Mainly technology push
- Difference in use of knowledge (e.g. hierarchical, object-oriented etc vs. associative knowledge) and 'nature of the knowledge' (certainty vs. good-enough, conjectures, change)
- Everybody (claims to) develop(s) the ultimate best solution. There is a lot of stuff, but a) one can't see the forest for the trees on what is there and b) **there is an awakening realisation that the bioinformatics tools at present don't do quite what biologists had in mind the tools would do in helping them to do their research more effectively.** In addition: there are **moving targets**.
- The “jumping on the bandwagon” feeling is very present, there's a lot of money going around in bioinformatics, lots of over-optimistic promises, ethical side largely ignored.

Note: some tools are really useful!

Characteristics



- Is it *interdisciplinary*?

interdisciplinary **teams** vs. interdisciplinary **people**

“An interdisciplinary team is a committee in which members identify themselves as an expert in something else besides the actual scientific problem at hand, and abdicate responsibility for the majority of the work because it's not their field.” [Eddy05]

a) ‘bridge’ function, multilingual

b) apply approach/methodology *a* from discipline *x* to problem *b* from discipline *y*, and v.v. and scale up and merge

Then: “...inventing new ways to look at the world.” [Eddy05]

e.g.: fixing the biologist’s radio [Lazebnik02]

Characteristics



- Is it *interdisciplinary*?
- Or maybe *antedisciplinary*?

Watson & Crick were ornithologist and physicist by training, but did molecular biology, even though the term & discipline didn't exist yet.

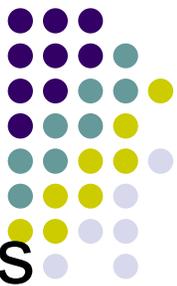
- ⇒ *Will/does/did the combination of CS, IT, and Bio result in one, or more, new disciplines?*
- ⇒ *Does a named set of a few characteristic activities make a (new) (sub)discipline?*

Characteristics

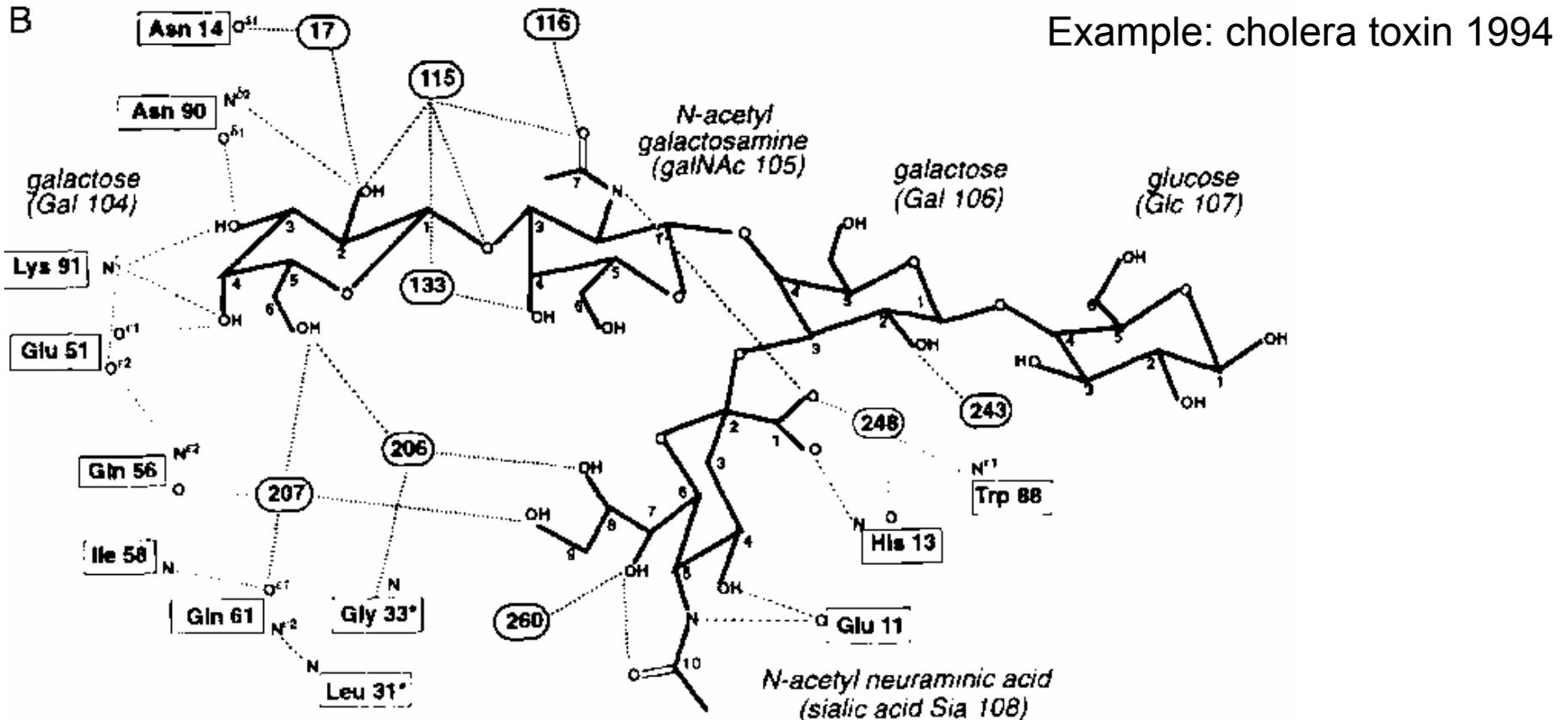


- Some examples of integrative approaches
 - Virtual cell _[ecell]: cell physiologists, (bio)chemists, geneticist, modellers, programmers, visualisation, mathematicians

(Virtual) Cells (& cell components) example



- Basic options:- Schematic & measurements



B: Schematic representation of hydrogen bonding interactions involving the G_{MI} pentasaccharide in subunit B#5 of the pentamer. The peptide residues shown belong to a single monomer, except for the involvement of Gly 33 from an adjacent monomer. Solvent molecules are depicted as ovals. Electron density at the other 4 copies of the pentamer is insufficient to model the full pentasaccharide. In subunit B#5, however, the conformation of the glucose end of the pentasaccharide is stabilized by hydrogen bonds (not shown) to a separate molecule related by the crystallographic symmetry operation $(x + 1/2, y + 1/2, E)$. [meritt94]

(Virtual) Cells

(& cell components)

example

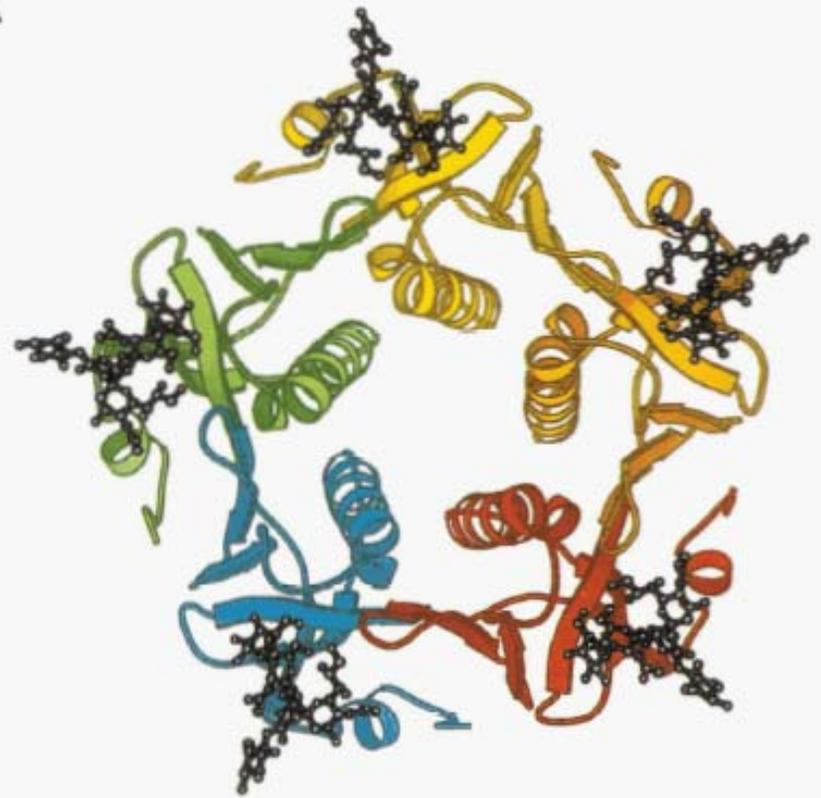
- Basic options

more cholera toxin:

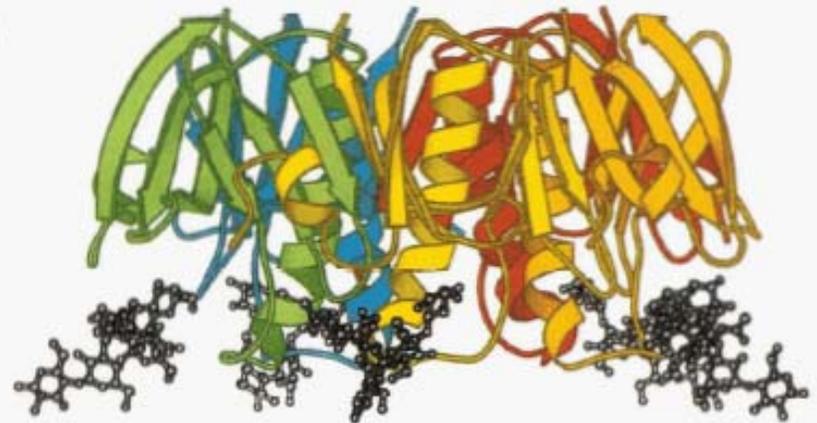
(3D) structure of the protein

[merritt94]

A



B

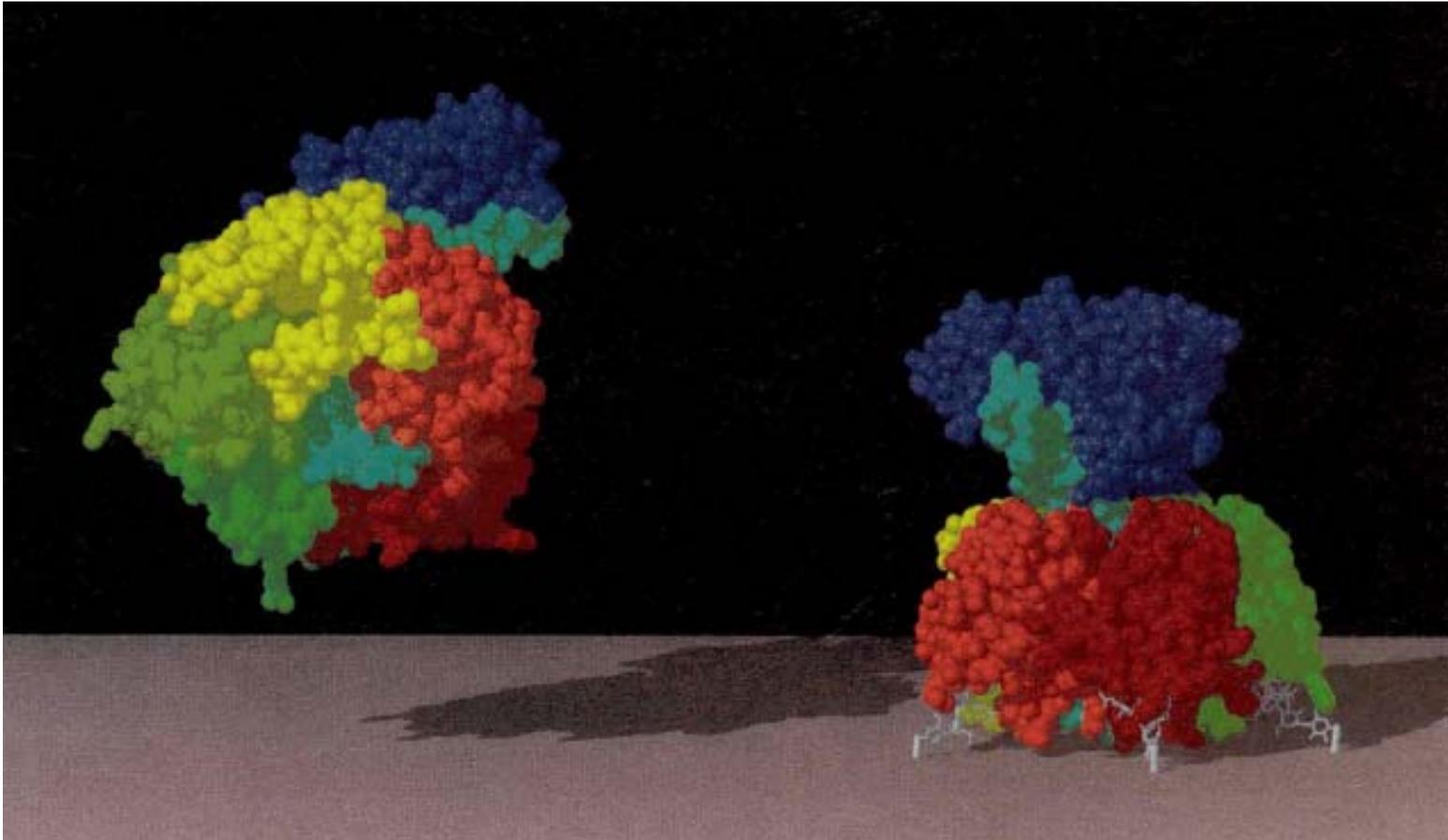


(Virtual) Cells (& cell components) example



- Basic options: even more cholera toxin, now “in action”

the B-pentamer of the toxin binds specifically to the branched pentasaccharide moiety of ganglioside G_{M1} , on the surface of target human intestinal epithelial cells. [meritt94]





(Virtual) Cells example

- Basic options:
 - Schematic e.g. cholera toxin 1994 [meritt94]
- Progress over the past decade, with *new, more, and better*:
 - Simulations for education
 - “fig 9.” http://cellix.imolbio.oeaw.ac.at/Videotour/video_tour_5.html
 - “Translation: the movie” <http://vcell.ndsu.nodak.edu/~christjo/vcell/animationSite/index.htm>
 - Simulations with data
schematic & measurements
 - Videos of real things at (sub)cellular level
see yourself: http://cellix.imolbio.oeaw.ac.at/Videotour/video_tour.html

Characteristics

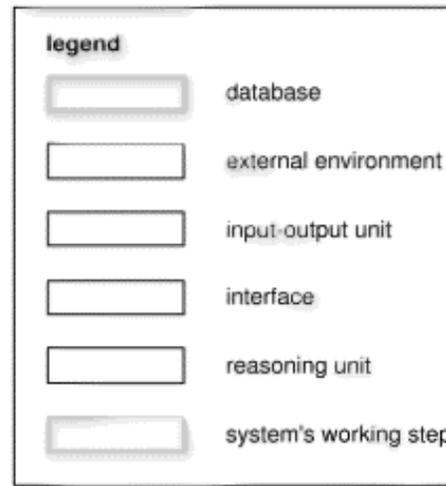
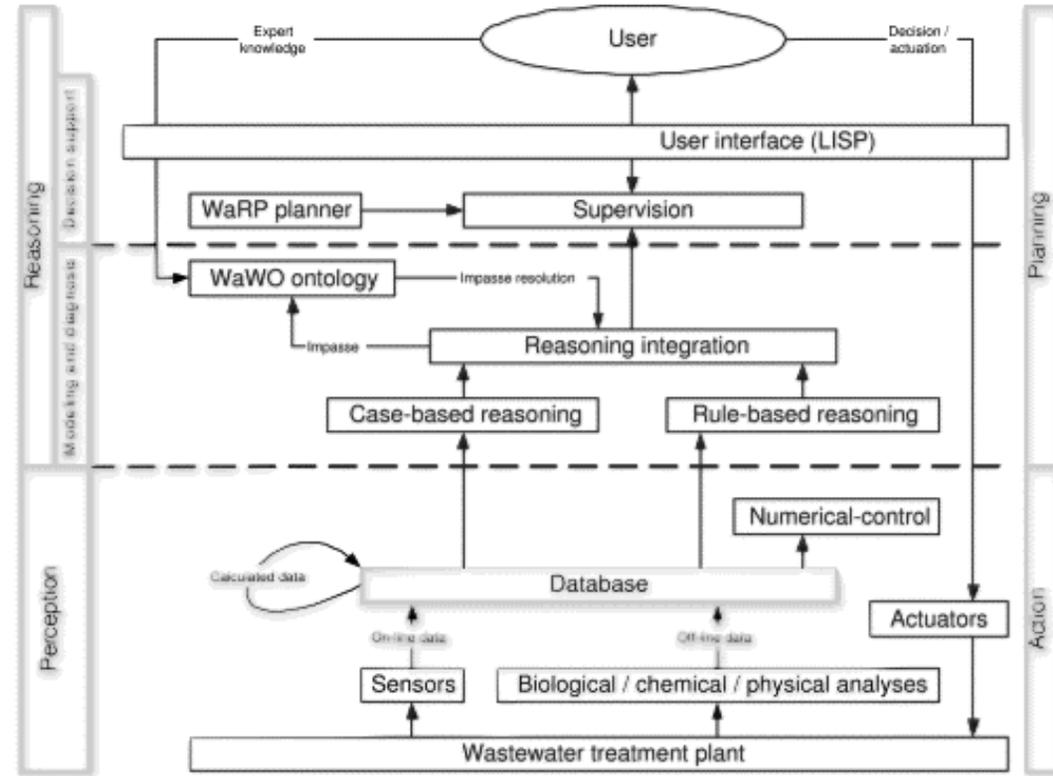


- Some examples of integrative approaches
 - Virtual cell [ecell]: cell physiologists, (bio)chemists, geneticist, modellers, programmers, visualisation, mathematicians
 - Sequence comparisons: geneticists, data miners, mathematicians/statisticians, visualisation, DB, Web-access
 - Applied sciences: e.g. medical / agriculture / environmental engineering. example

Environmental Engineering



- OntoWEDSS: Environmental decision support system for managing active sludge in wastewater treatment plants.





Characteristics

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 - Applied sciences
 - Ecology: example



Ecology - example

- Knowledge Network for Biocomplexity

“...network intended to facilitate ecological and environmental research on biocomplexity. For scientists, the KNB is an efficient way to discover, access, interpret, integrate and analyze complex ecological data from a highly-distributed set of field stations, laboratories, research sites, and individual researchers... focuses on research into informatics and biocomplexity, through the development of software products...” <http://knb.ecoinformatics.org/index.jsp>

⇒ metadata, storage resource broker, distributed data management tool, data integration, quality assurance, hypothesis modelling (Bayesian network), visualisation tools

Characteristics



- Some examples of integrative approaches
 - Virtual cell [ecell]: cell physiologists, (bio)chemists, geneticist, modellers, programmers, visualisation, mathematicians
 - Sequence comparisons: geneticists, data miners, mathematicians/statisticians, ...
 - Applied sciences
 - Ecology
 - Bio-Linux <http://www.biolinux.org/> , 60 software packages
 - BioBrew http://bioinformatics.org/project/?group_id=273, 223 groups
 - Exercise ‘Conservation efforts’

Conservation effort exercise



1. Let X be an endangered species and you want to conserve it.
2. You are in the fortunate position to have enough resources (researchers, technicians, money, etc) to set up a project to fulfill (1).
3. Goal:
how to come up with the “recovery plan”?

Conservation effort exercise



- Known variables:
 - Habitat of species X , the site(s) where the organisms live at present, population census
 - Its diet and predator
 - Dispersal, migration characteristics
- What we need to know:
 - What did change at the sites, and how?
 - Are there any viable other sites to (re)introduce X , if yes, where?
 - Is the gene pool diverse enough?



Characteristics

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 - Medical info system/agriculture
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 - Exercise ‘Conservation efforts’
- Don’t forget ***intradisciplinary* science**



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Titles & Topics

- Ethics
- Computational Systems Biology
- Virtual laboratory for integrative bioinformatics research
- A standardized and dynamic approach for immunogenetics and immunoinformatics
- Querying biological data sources
- Requirements for natural language understanding in referent-tracking based electronic health records
- Ontology design patterns and practical expertise: roles, tasks and techniques in the agricultural domain
- Ecological niche

References & related websites



- Some **Journals & Conferences**:
 - lists: <http://www.lri.fr/~cohen/confSem/confSem.html> , <http://www.conferencealerts.com/bioinform.htm>
- Some **websites & organisations**:
 - Gene Ontology Consortium: <http://www.geneontology.org>
 - Society for Mathematical Biology: <http://www.smb.org/>
 - International Society for Computational Biology: <http://www.iscb.org/>
 - European Molecular Biology Laboratory: <http://www.embl.org/>
 - European Bioinformatics Institute: <http://www.ebi.ac.uk>
 - European Federation for Information Technology in Agriculture, Food and the Environment: <http://www.efita.net/>
 - International Medical Informatics Association: <http://www.imia.org>
- Some **Projects**:
 - Marine genomics: <http://www.marine-genomics-europe.org>
 - Science Environment for Ecological Knowledge: <http://seek.ecoinformatics.org>
 - Semantic Mining in Biomedicine: <http://www.semanticmining.org>
 - Biopattern: <http://www.biopattern.org/index.html>
 - InfoBioMed: <http://www.infobiomed.org>
 - EurEthNet: <http://www.eureth.net>
- Referenced literature:
 - [Ceccaroni04] Ceccaroni, L., Cortés, U. and Sánchez-Marrè, M. OntoWEDSS: augmenting environmental decision-support systems with ontologies. *Environmental Modelling & Software*, 2004, 19 (9): 785-797.
 - [ecell] E-cell project. <http://www.ndsu.nodak.edu/instruct/mcclean/vc/wwwic-vc5.htm> (there are many more projects)
 - [Eddy05] Eddy, S.R. “Antedisciplinary” Science. *PLoS Computational Biology*, 2005, 1(1): e6. <http://compbiol.plosjournals.org/perlserv/?request=get-document&doi=10.1371/journal.pcbi.0010006>
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 - [Lazebnik05] Lazebnik, Y. Can a Biologist Fix a Radio? – or, What I Learned while Studying Apoptosis. *Cancer Cell*, 2002, 2: 179-182. http://www.protein.bio.msu.ru/biokhimiya/contents/v69/pdf/bcm_1403.pdf
 - [Merritt94] Merritt, E.A., Sarfaty, S., Akker, F. van den, L’Hoir, C., Martial, J.A., Hol, W.G.J. Crystal structure of cholera toxin B-pentamer bound to receptor G_{M1} pentasaccharide. *Protein Science*, 1994, 3: 166-175.