EVOLVING ASTEROID STARSHIPS: A BIO-INSPIRED APPROACH FOR INTERSTELLAR SPACE SYSTEMS

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- ² SEAD (Space Ecologies Art and Design)
- ³ Technical University of Berlin, Berlin, Germany
- ⁴ SmartCrops BV, The Hague, the Netherlands





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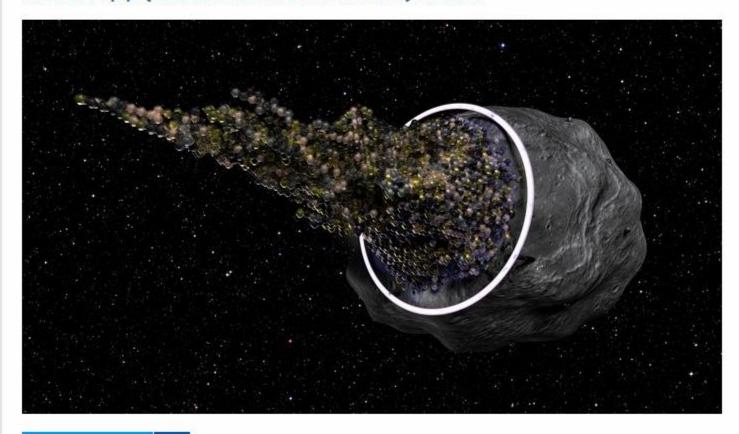
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DETAILS

Title TU Delft E|A|S (Evolving Asteroid Starships) project

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A group of students and researchers at Delft University of Technology are designing a starship capable of keeping generations of crew alive as they cross the gulf between stars – and they've turned to ESA for the starship's life

DSTART the TII Delft Starship Team, is bringing together a wide variety of disciplines to perform advanced

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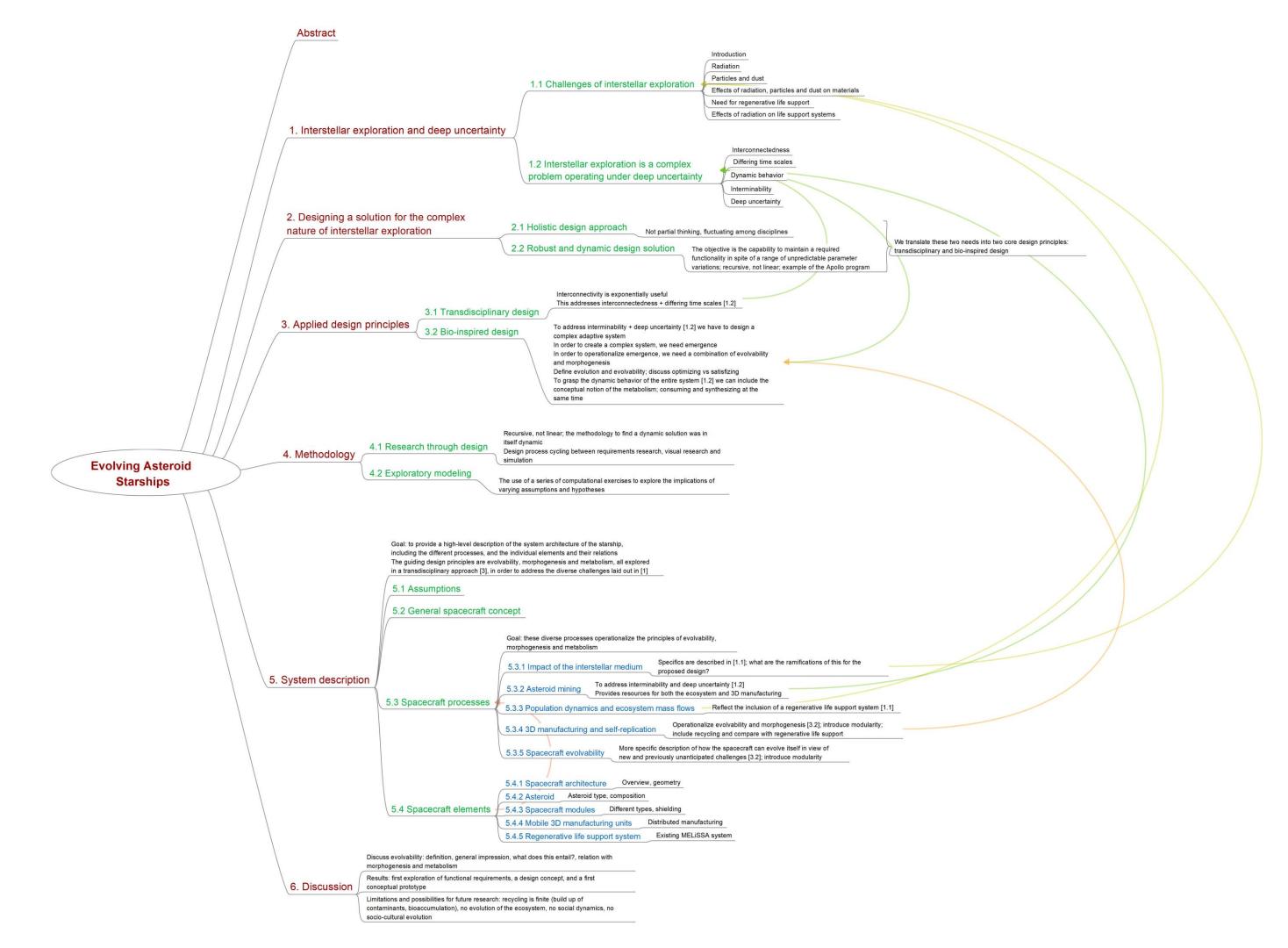
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CHALLENGES OF HUMAN INTERSTELLAR EXPLORATION

Radiation
Particles and dust
Regenerative life support
Not part of this study:

- propulsion & power supply
- social & cultural aspects

HUMAN INTERSTELLAR EXPLORATION IS A COMPLEX PROBLEM OPERATING UNDER DEEP UNCERTAINTY

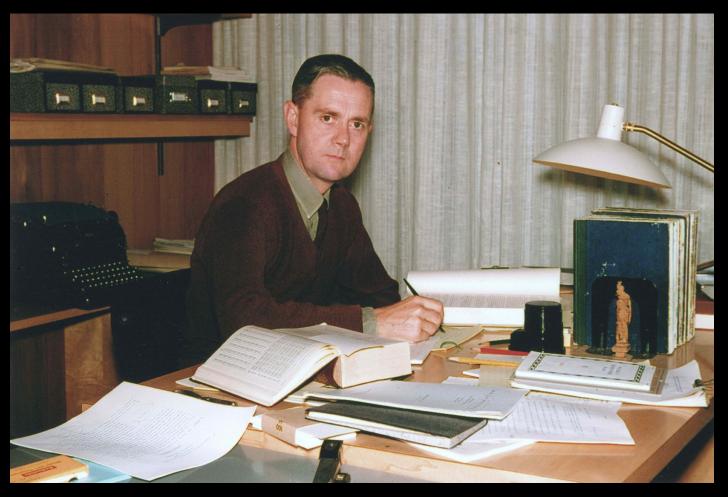
Interconnectedness
Differing time scales
Dynamic behavior
Interminability
Deep uncertainty

BIO-INSPIRED DESIGN FOR HUMAN INTERSTELLAR EXPLORATION

- To address interminability and deep uncertainty we have to design a <u>complex adaptive system</u> (CAS)
- To create a complex system, <u>emergence</u> is needed
- Emergence can be operationalized through a combination of evolution and morphogenesis

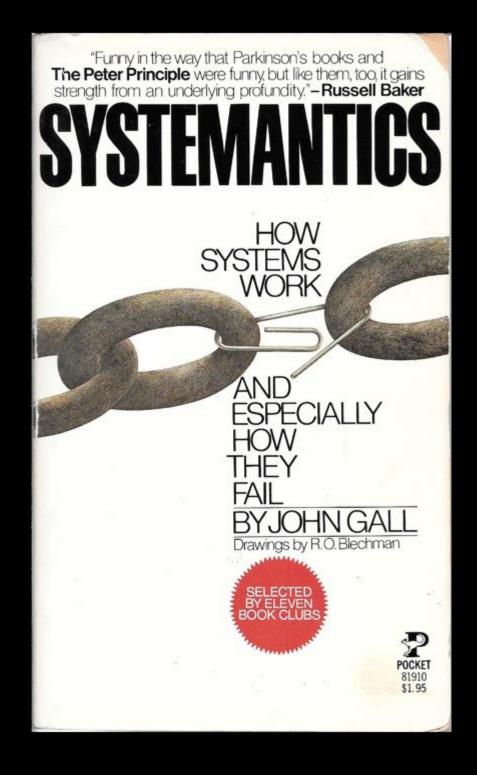
ASHBY'S LAW OF REQUISITE VARIETY

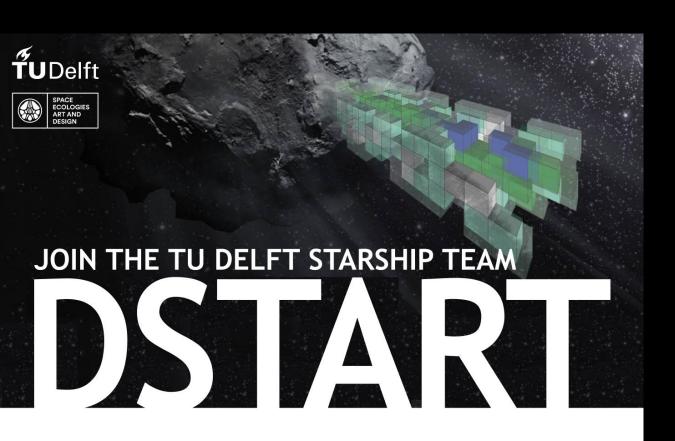
The larger the variety of actions available to a control system, the larger the variety of perturbations it is able to compensate.



GALL'S LAW

A complex system that works is invariably found to have evolved from a simple system that worked. A complex system designed from scratch never works and cannot be patched up to make it work. You have to start over with a working simple system.





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More info: Angelo Vermeulen, a.c.j.vermeulen@tudelft.nl



Mikhail Sirenko **Statistics** Modeling



Arise Wan Architecture **Biomimicry**



Anton Dobrevski Architecture Web design



Alvaro Papic Biological life support Agent-based modeling



Joris Putteneers Architecture Animation



Jimmy Verkooijen Background research Outreach



Farshad Goldoust Astrophysics Modeling



Nils Faber 3D modeling Visualization



Sharon van Rijthoven Aerospace engineering Community management



Daniela Hallak Astrophysics Requirements analysis



Jasper Wennekendonk 3D modeling Visualization



Angelo Vermeulen Model development Biological life support



Brennan Lutkewitte Asteroid mining Requirements analysis

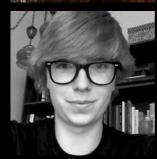
Jason Kiem

Mass flow analysis

Requirements analysis



Amelie Kim 3D modeling *Architecture*

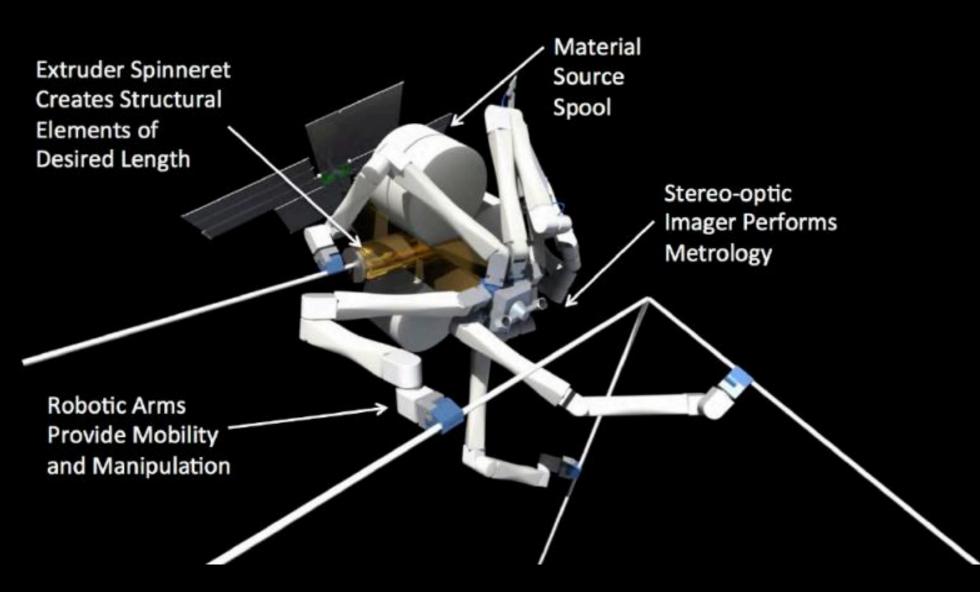


Andreas Theys Aerospace engineering **Programming**

ASTEROID MINING

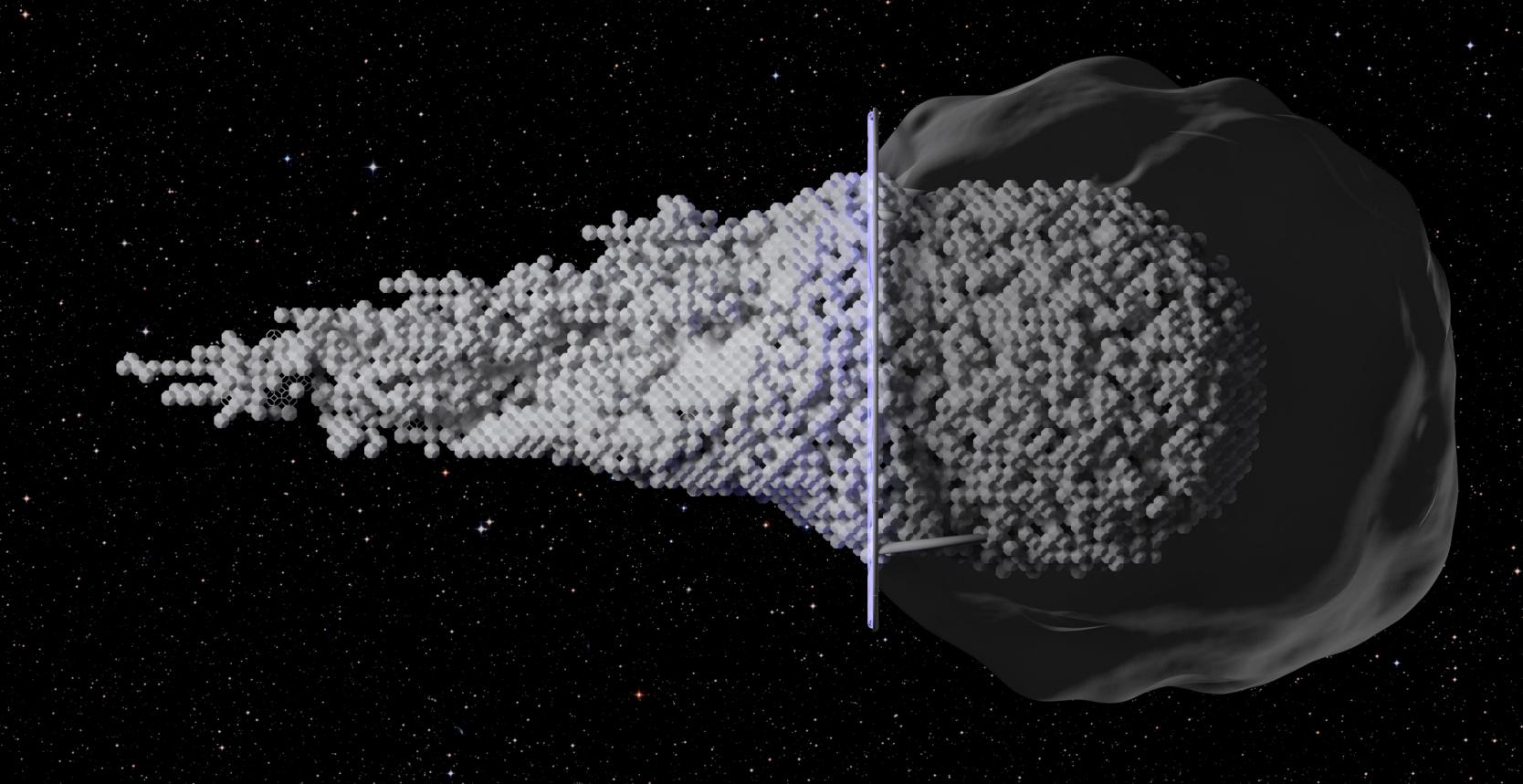
3D MANUFACTURING

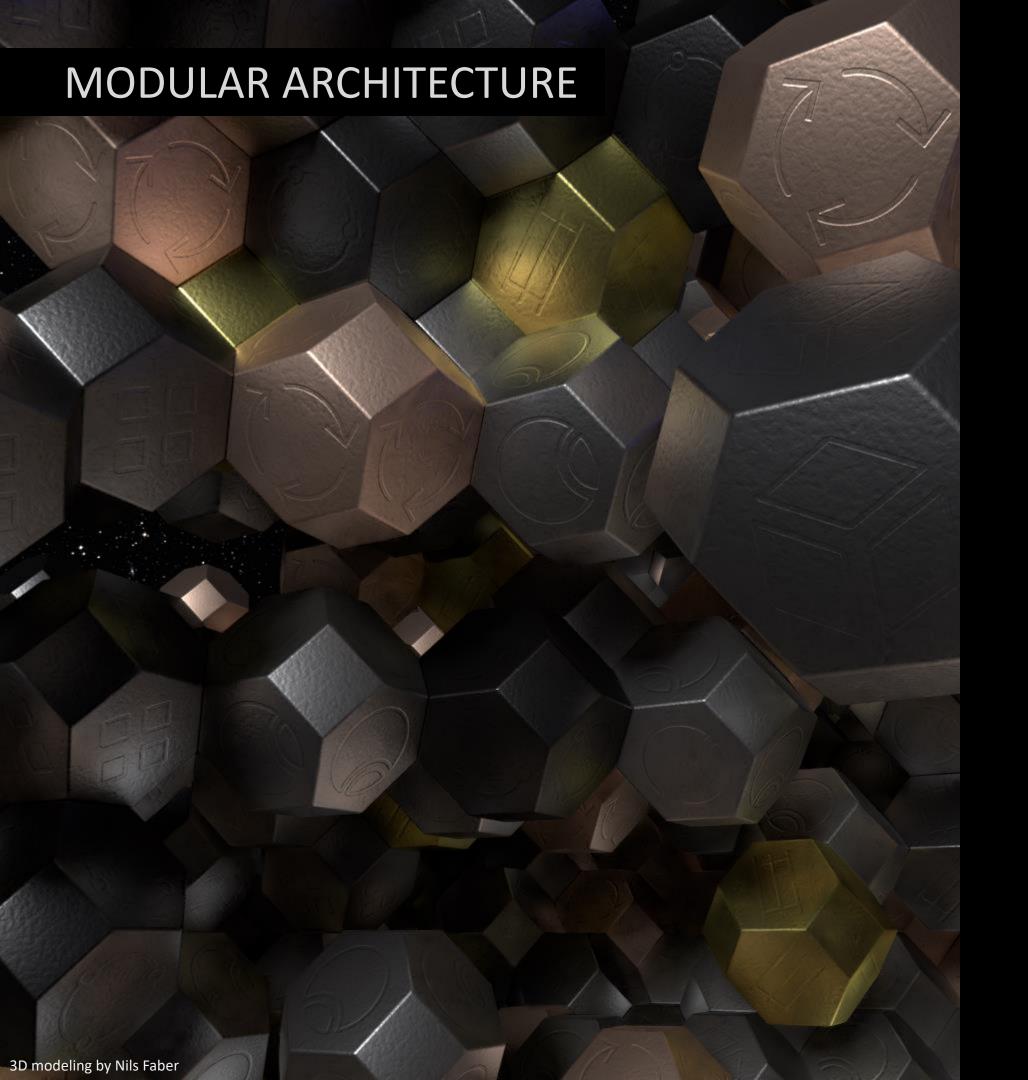




SpiderFab, Tethers Unlimited & NASA, 2013

MODULAR ARCHITECTURE



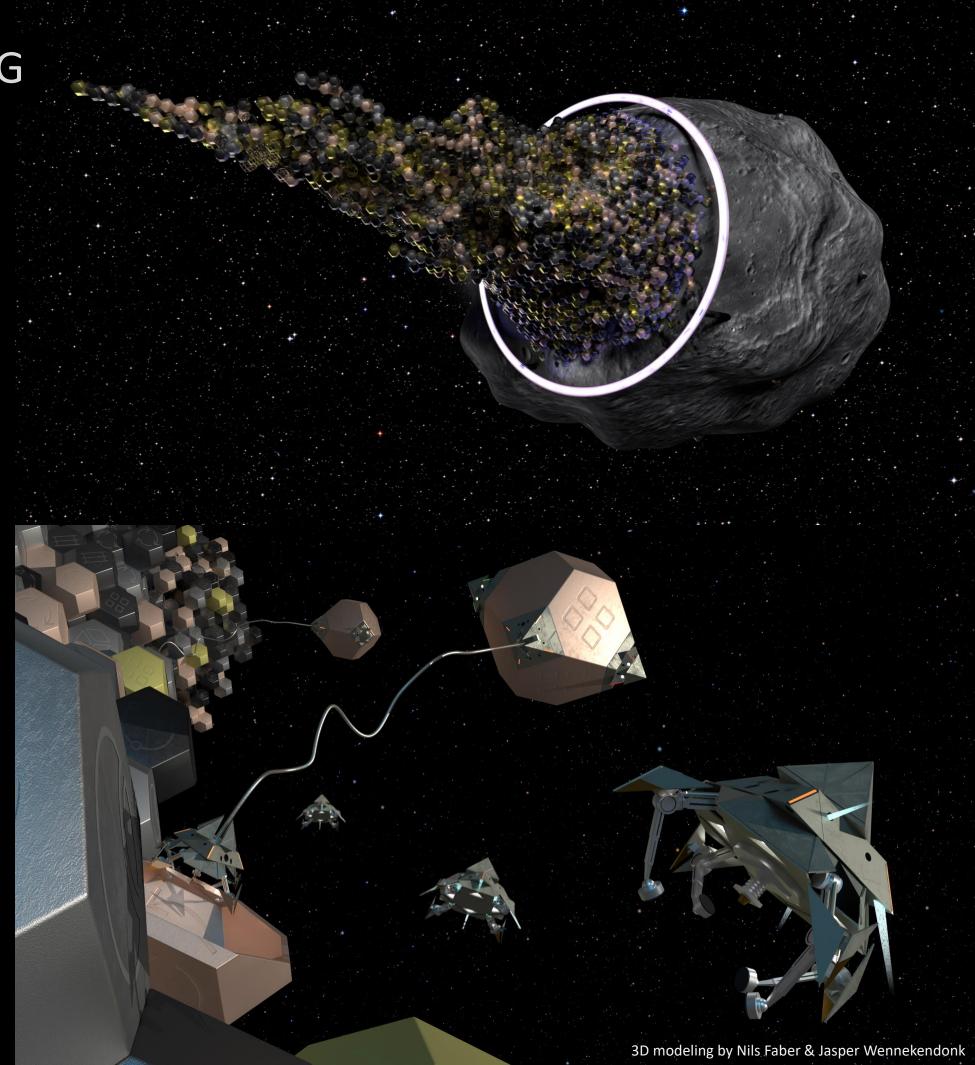


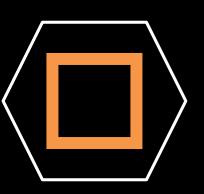
MODULE TYPES

- 1. Regenerative life support
- 2. Habitation
- 3. Radiation shielding
- 4. Collision shielding
- 5. Mining
- 6. Processing
- 7. Manufacturing
- 8. Ore storage
- 9. Refined materials storage

MORPHOGENETIC/EMERGENCE ENGINEERING

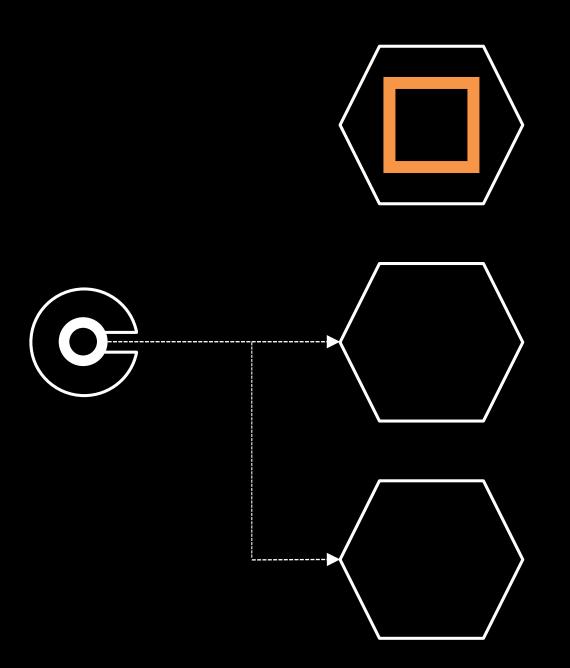




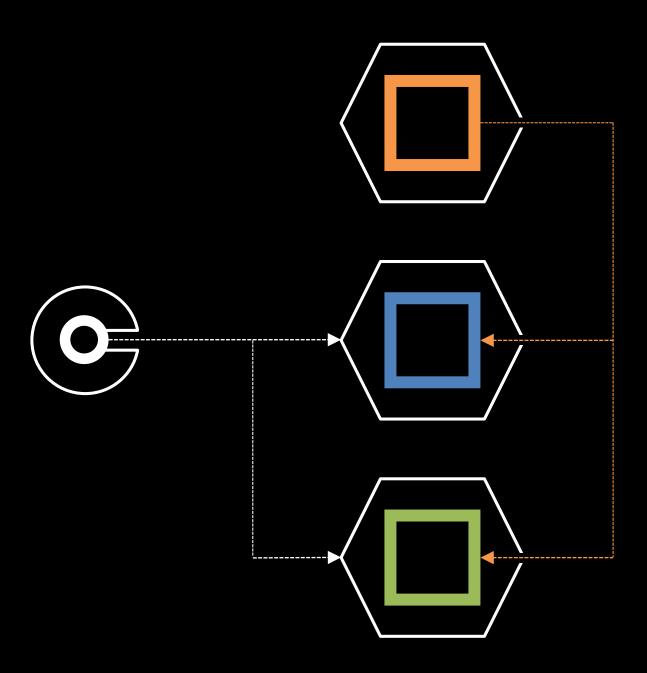




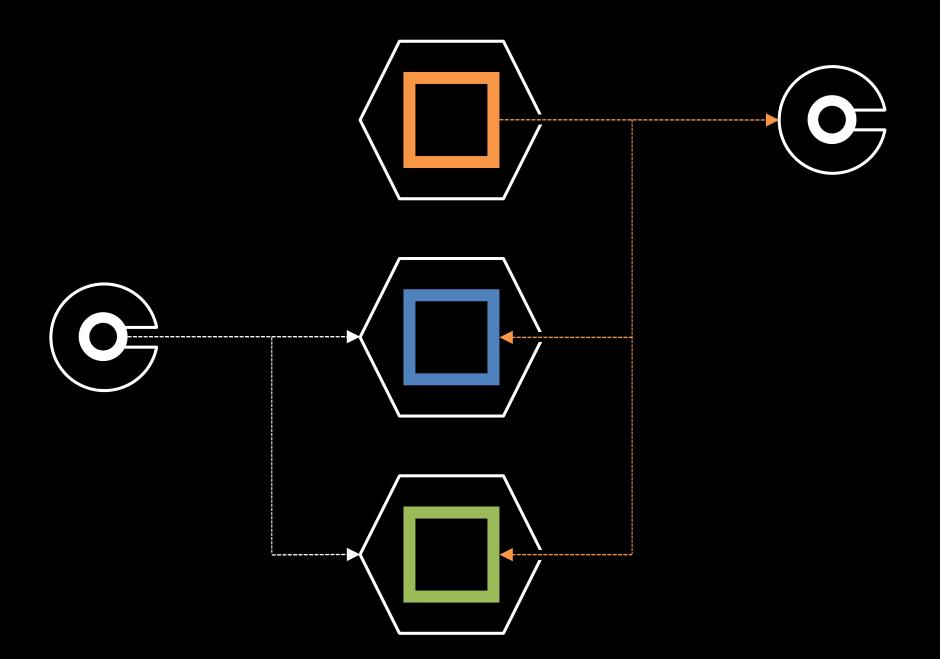
3D manufacturing robotManufacturing module



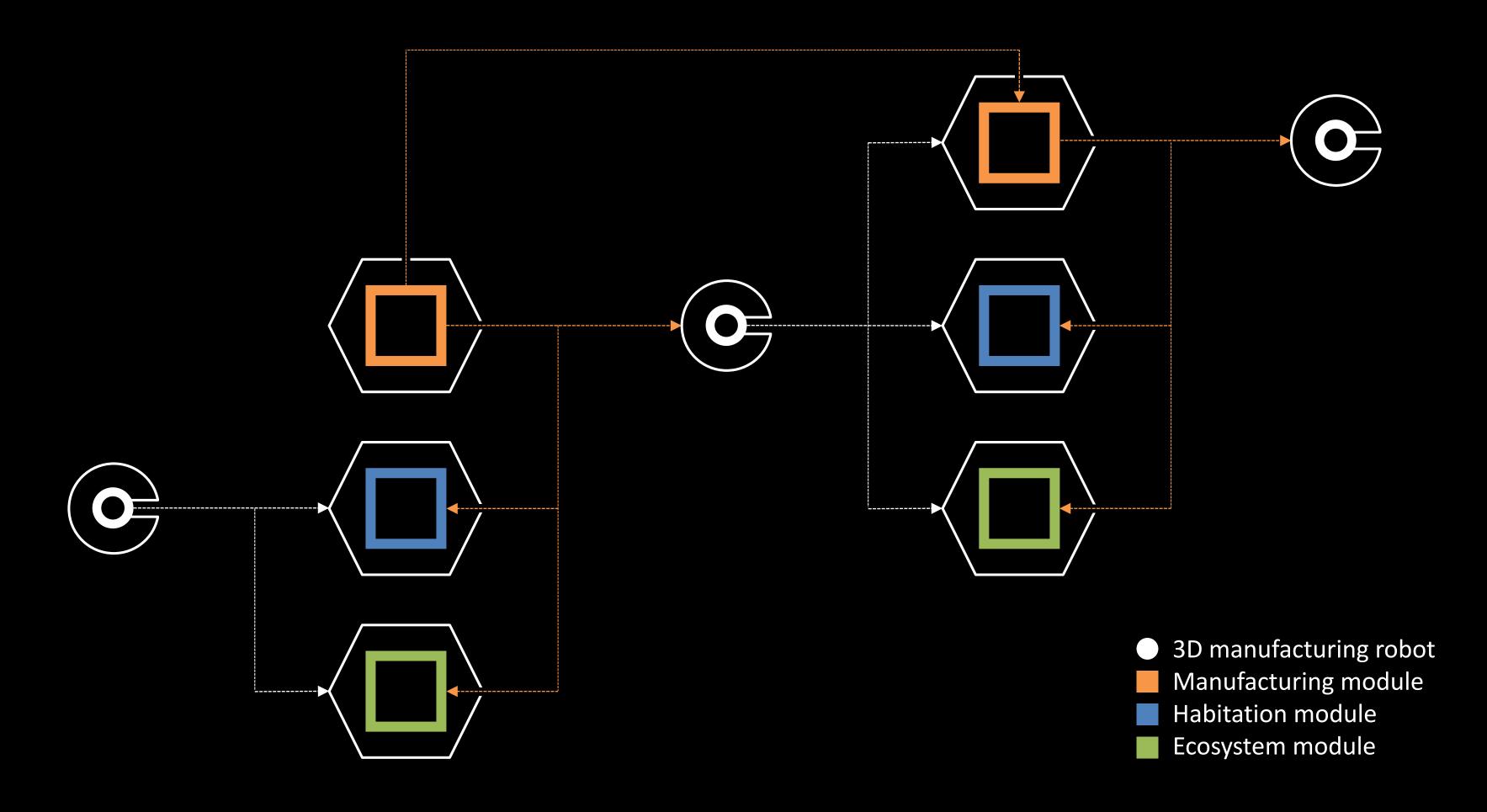
- 3D manufacturing robot
- Manufacturing module



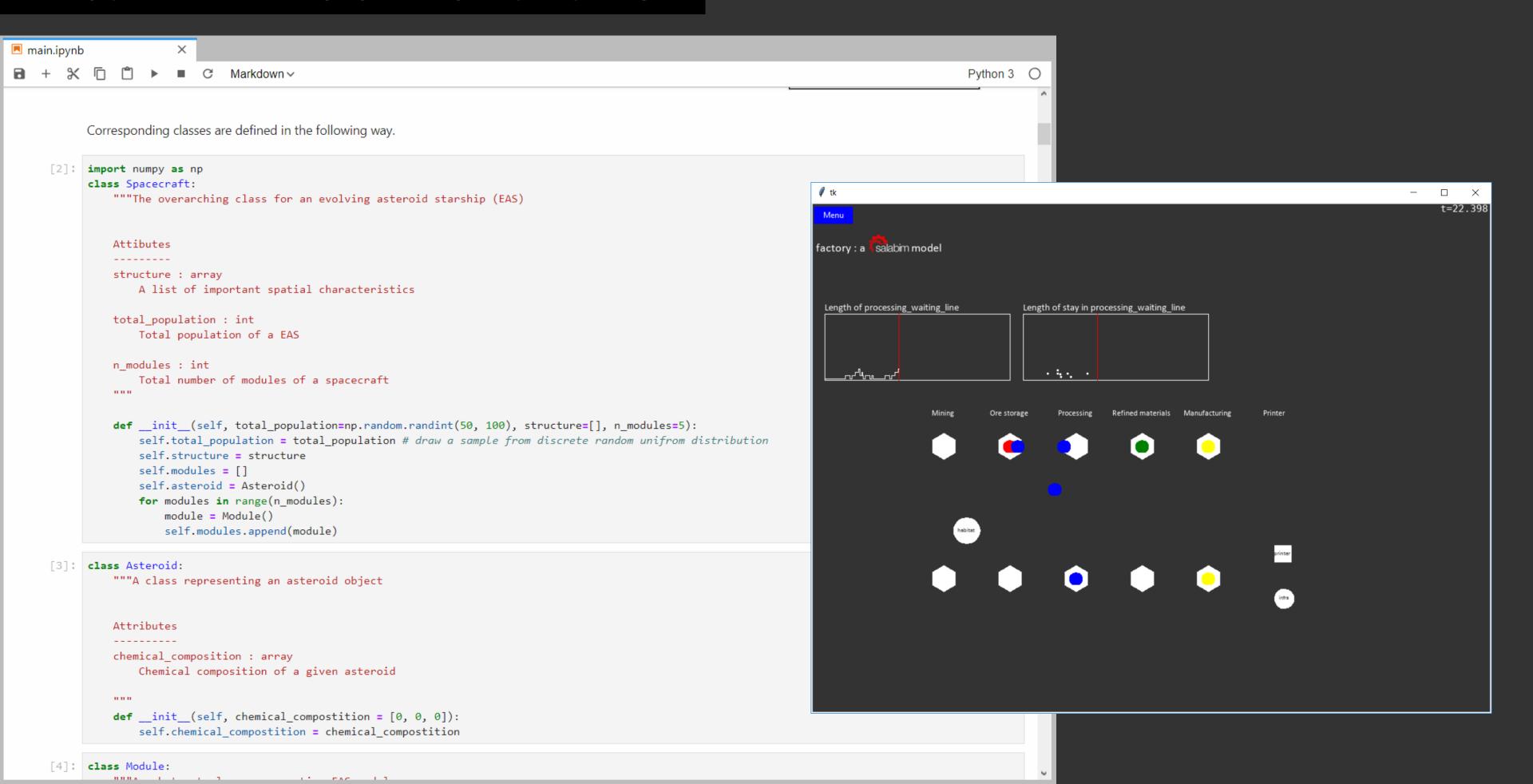
- 3D manufacturing robot
 - Manufacturing module
- Habitation module
- Ecosystem module



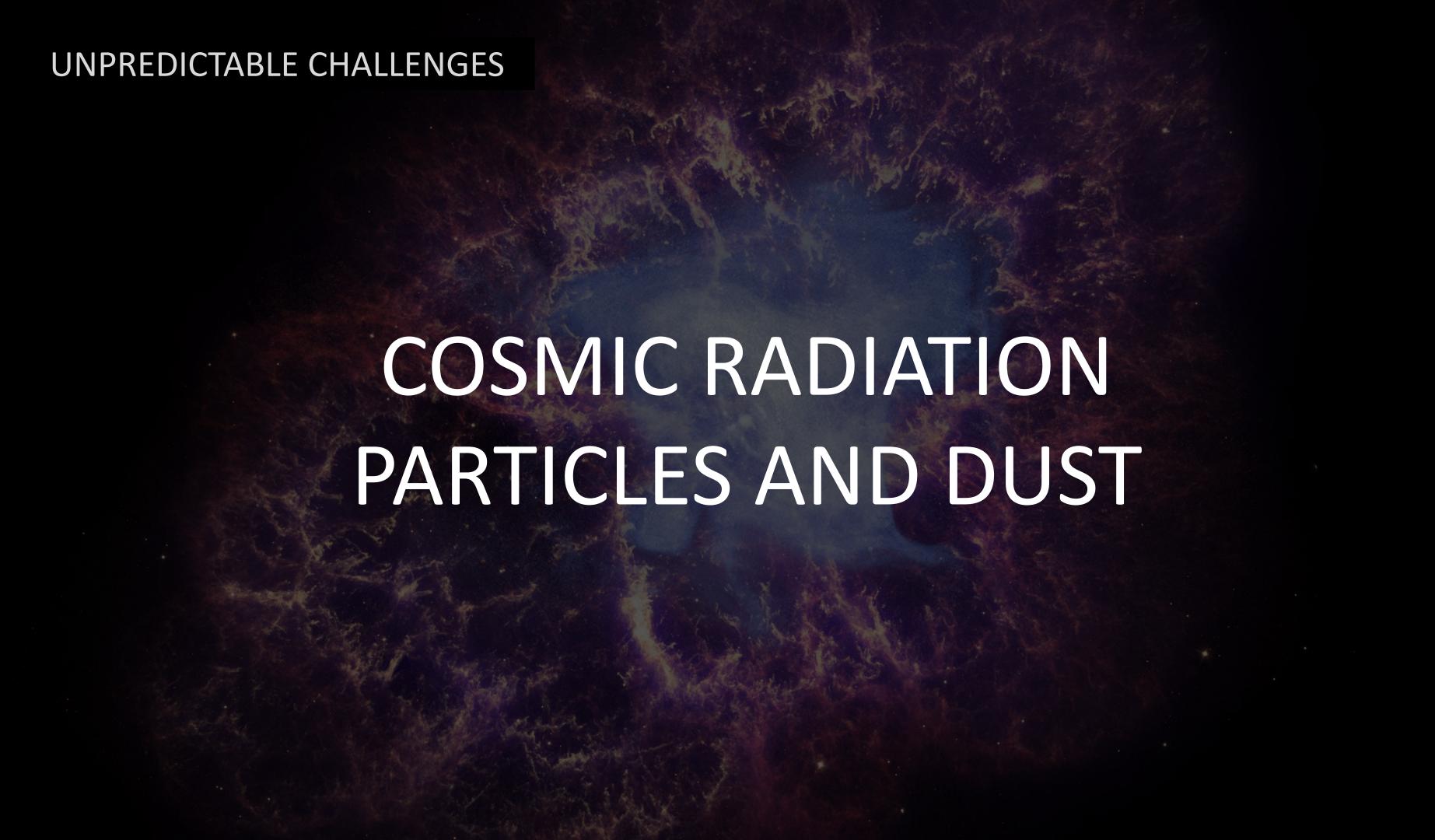
- 3D manufacturing robot
- Manufacturing module
- Habitation module
- Ecosystem module



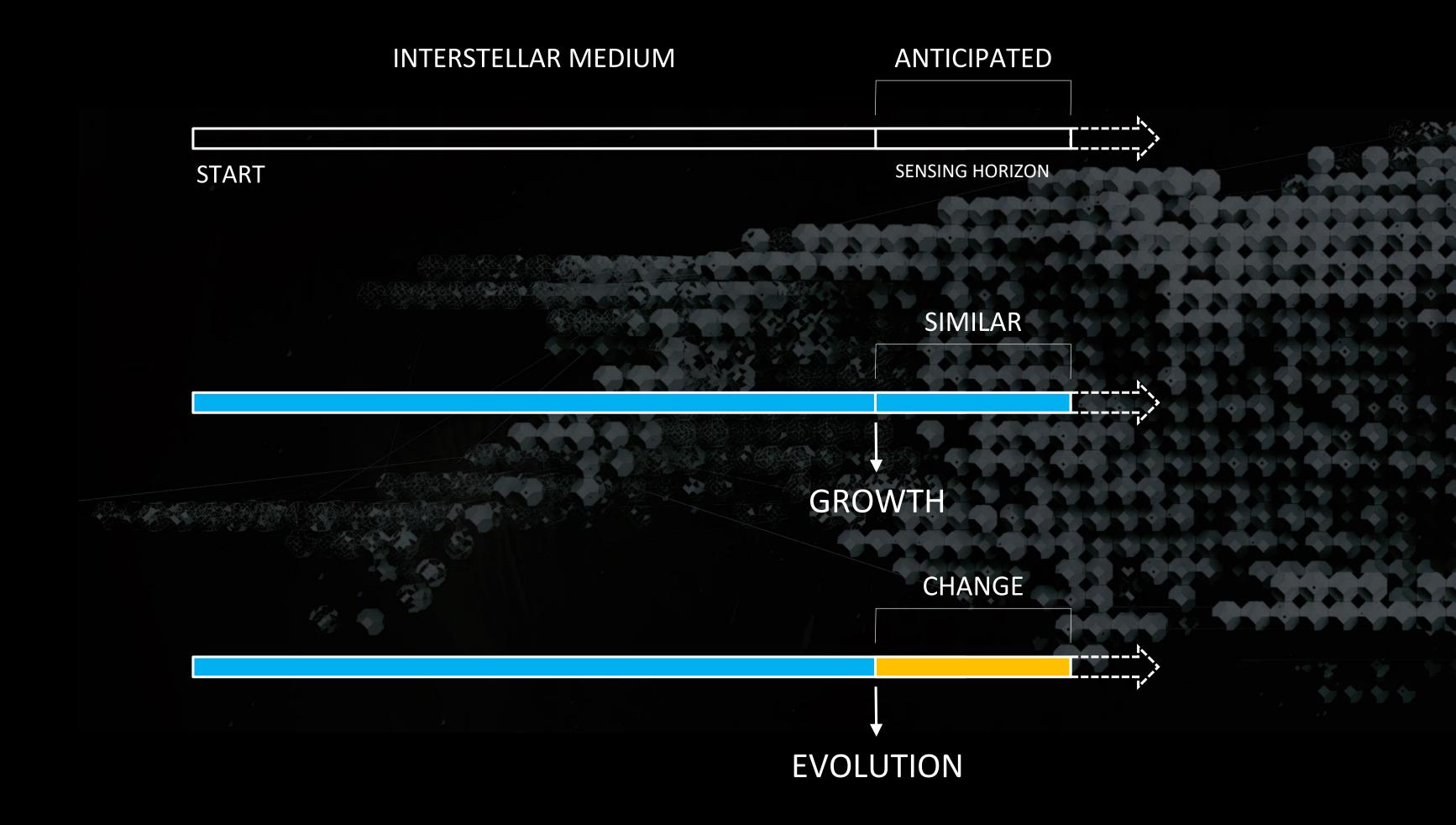
DISCRETE EVENT SYSTEM SPECIFICATION



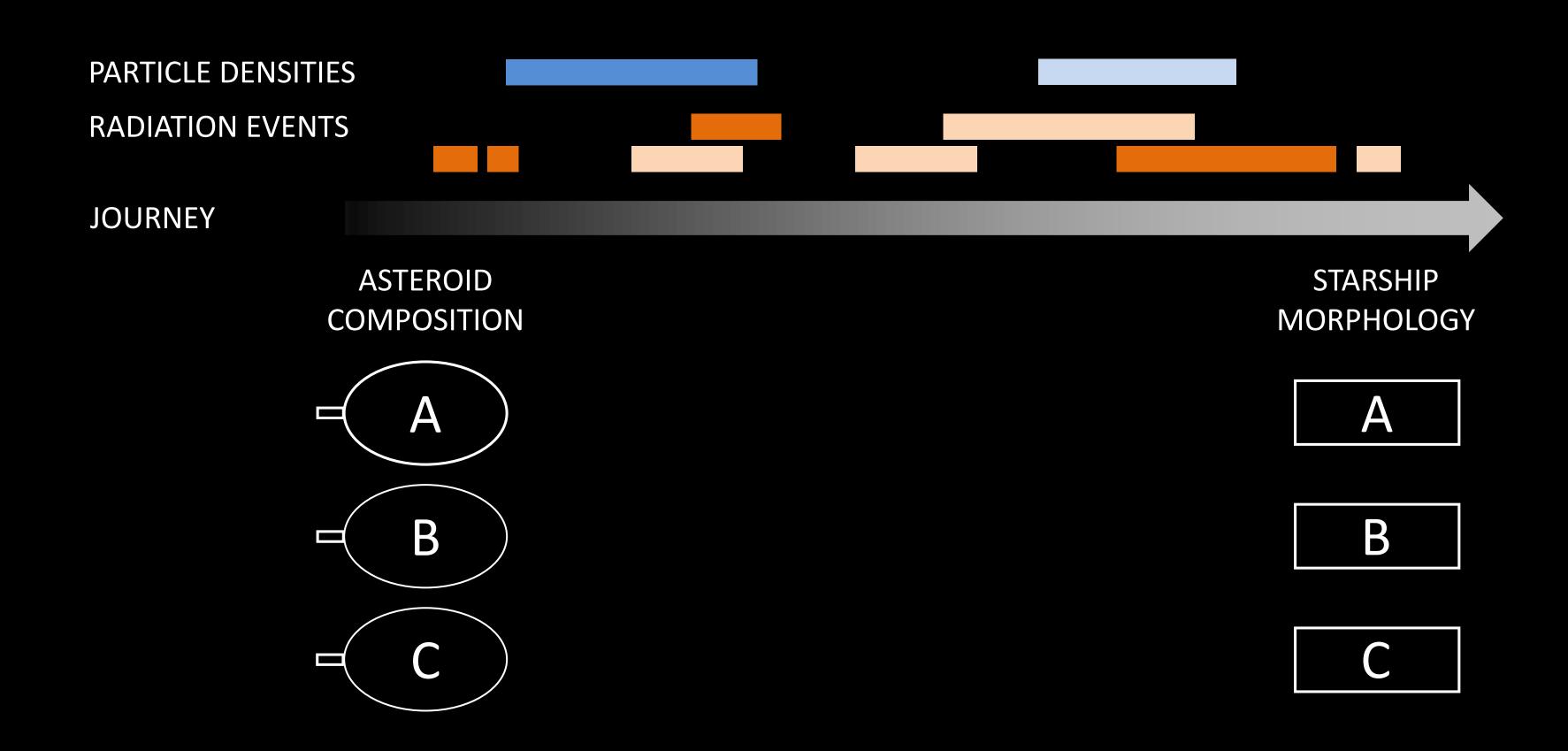




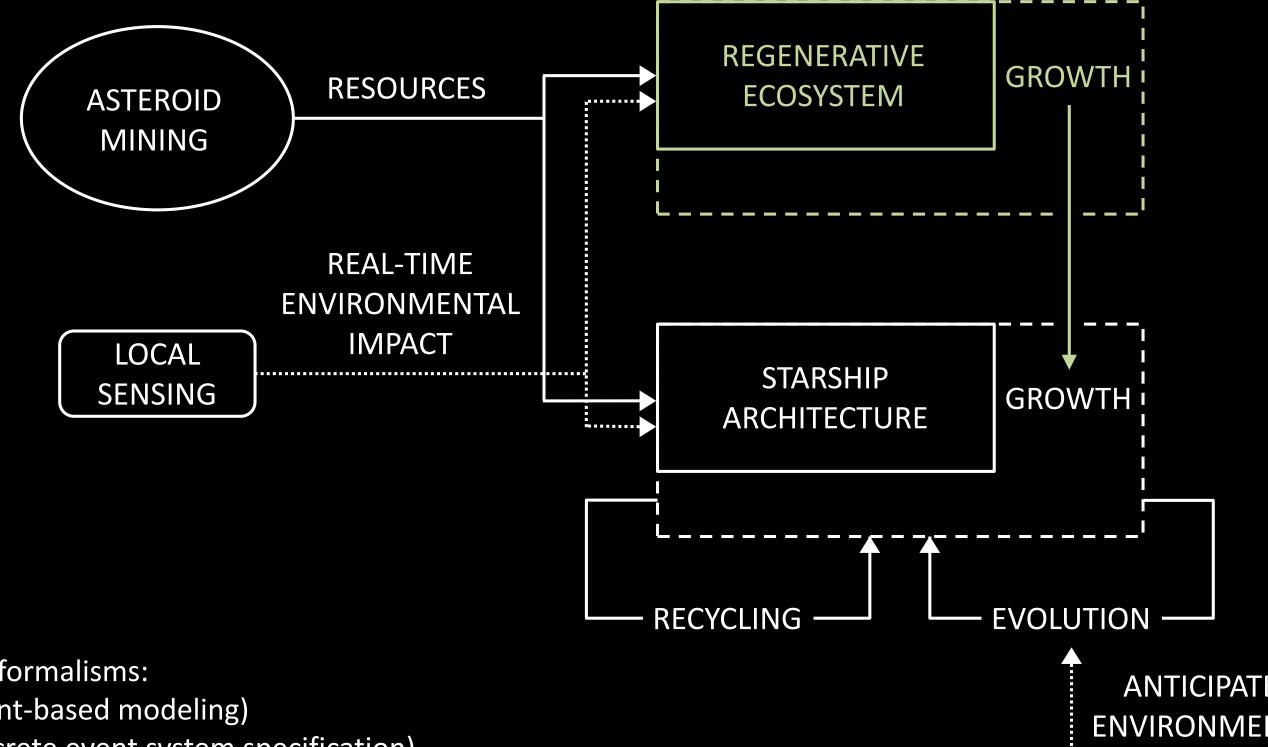
EVOLVABILITY



SIMULATION EXPERIMENTS

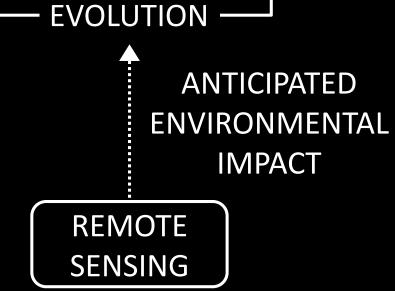


ECOSYSTEM INTEGRATION

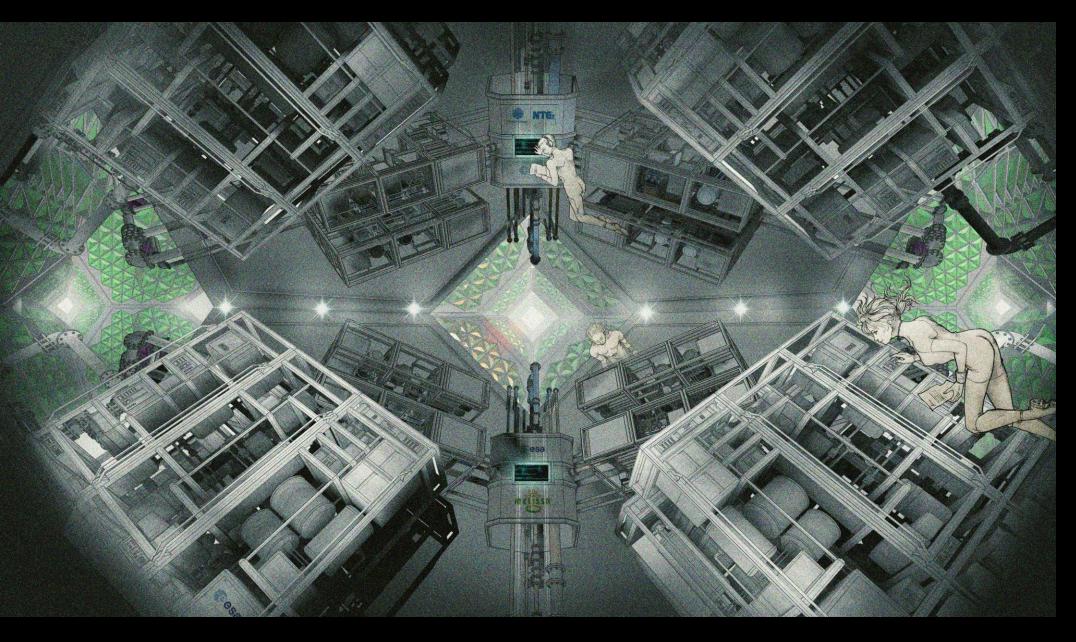


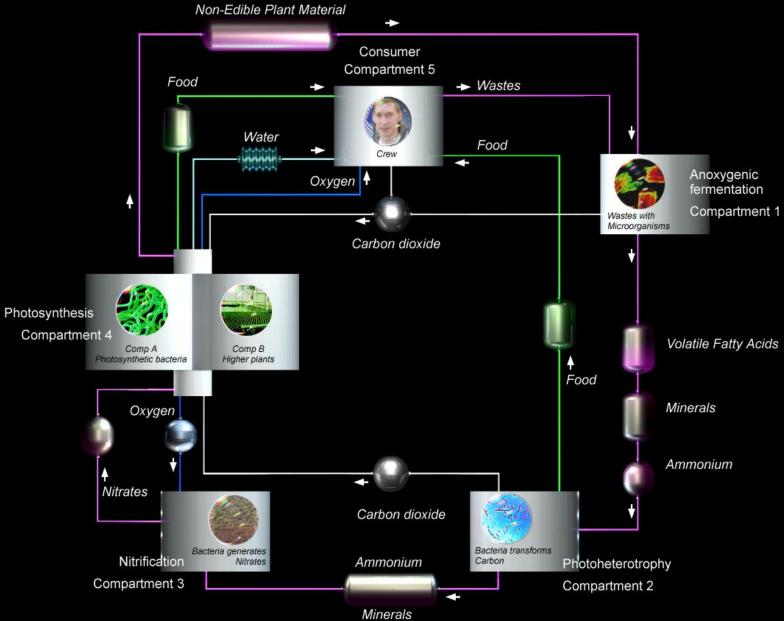
Simulation formalisms:

- ABM (agent-based modeling)
- DEVS (discrete event system specification)
- SD (system dynamics)
- EA (evolutionary algorithms)



ECOSYSTEM INTEGRATION







AGENT-BASED MODELING

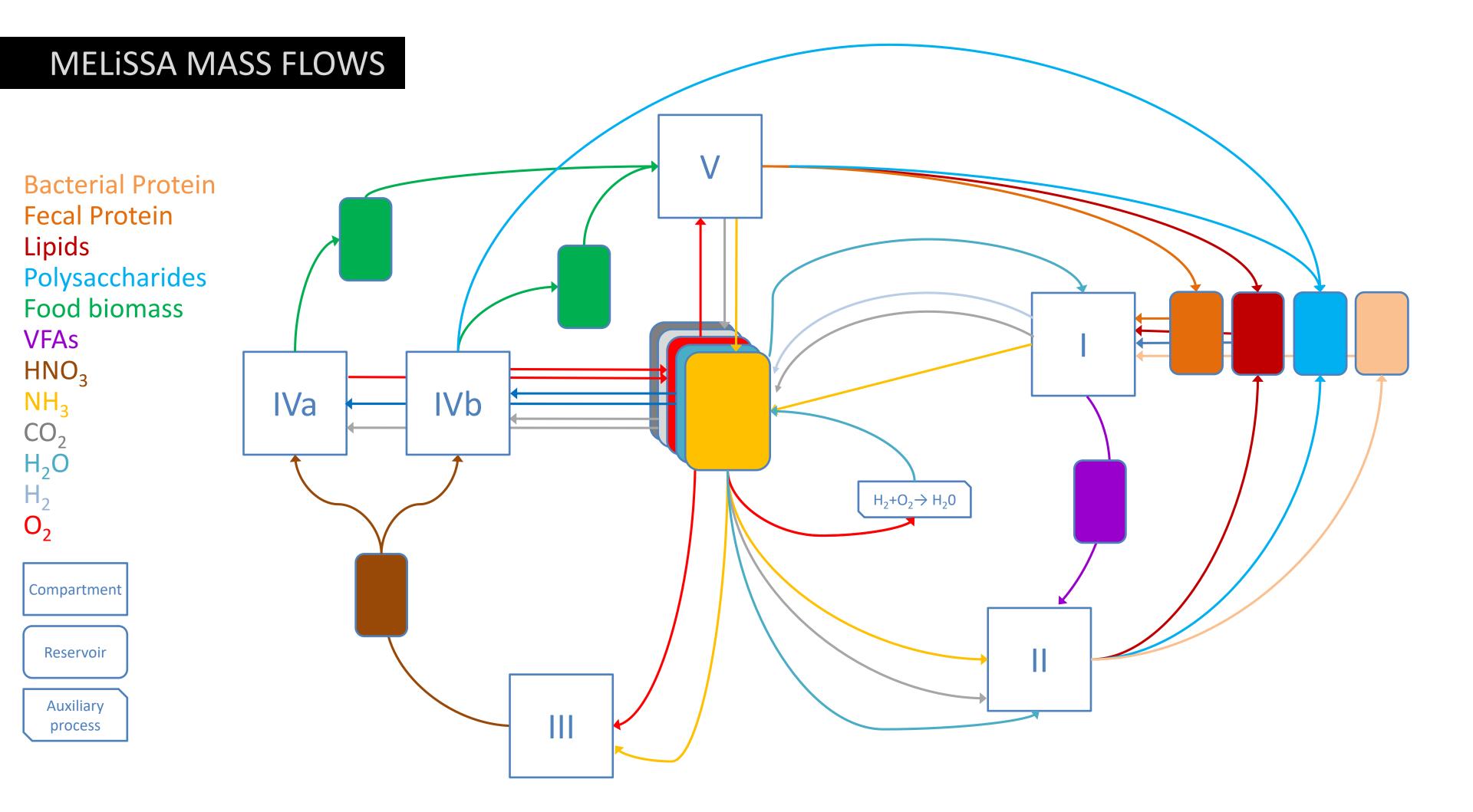
DEFINITION

- Works with agents and ticks
- Focus on interactions and emergent patterns
- High granularity and ontological correspondence

```
61⊟ to setup
        import-drawing "ABMbackground2.png"
                                 bac.prot fec.prot poly lipids food.p food.a vfa.ace vfa.but hno3 nh3 co2 h2o o2 h2
     set reservoir array:from-list [5861 528 3106 826 1440 400 8710 1278 1025 1502 5949 7514 5349 0] ;1019 in the same order as the previous
     set survey matrix:from-row-list [[ 0 0 0 0 -1 -1 -1 -1 -1 -1 -1 0 -1 -1];compartment I
                                          -1 -1 -1 -1 -1 -1 0 0 -1 0 0 -1 -1 -1]; compartment II
                                          -1 -1 -1 -1 -1 -1 -1 -1 -1 0 -1 -1 0 -1];compartment III
                                          -1 -1 -1 -1 -1 -1 -1 -1 0 -1 0 0 -1 -1];compartment IVa
                ➤ initial model v0985 - NetLogo
                                                                                                                                              File Edit Tools Zoom Tabs Help

✓ view updates

                                 *abc Button ▼
        set pr
                                                                C I: Thermophilic Anaerobic Bacteria
                                           C V: Human Crew
                                                                                                                               Plants for cons. [dry weight]
        set to
        set he
        set st
        set con
                                                                                                     VFAs - HNO3 - HN3
                                                                 C II: Photoheterotrophic Bacteria
        set sh
        set si
                                          C IV: Hydroponic Farm
                                           C IVa: Higher Plants
        set yo
        set xc
                                                                                                     CO2 - O2 - H2 - H2O
                                                                   C III: Nitrifying Bacteria
     let px 1
     let py 1
     foreach
          set
         set
                {{array: 5861.006 529.668 3111.38 826.619 1440 399.9976 8627.698 1265.923 1257.9262 1452.883 6414.9527 7717.4284 4843.1239 -1.0E-4}}
     create-bacteria 150
       set compartment 2;25 lts per bioreactor
        set biomass 1
                        ; as in 0=no, 1=yes
       set requirements array:from-list [58.0644 8.5189 8.1694 0.1620]; acet, but, nh3, co2
                          array:from-list [39.0720 9.5040 4.2240 22.1147]; bac.prot, pl, lp, h2o
        set products
                           array:from-list [58.0644 8.5189 8.1694 0.1620]
        set needs
                          array:from-list [0 0 0 0]
        set intake
```



MELISSA STOICHIOMETRY

COMPARTMENT I

Fecal protein

$$3,2CH_{1.76}O_{0.239}N_{0.239} + 3,035H_2O = C_2H_4O_2 + 0,1C_4H_8O_2 + 2.3H_2 + 0.76NH_3 + 0,8CO_2$$

Bacterial protein

$$3,2CH_{1.4697}O_{0.34}N_{0.2807} + 2,712H_2O = C_2H_4O_2 + 0,1C_4H_8O_2 + 1,3162H_2 + 0.8982NH_3 + 0,8CO_2$$

Polysaccharides

$$3,199CH_{1.667}O_{0.833} + 1,134H_2O = 1C_2H_4O_2 + 0,1C_4H_8O_2 + 1,4H_2 + 0,8CO_2$$

Lipids

$$C_{16}H_{32}O_2 + 13,0278H_2O = 6,5278C_2H_4O_2 + 0,6528C_4H_8O_2 + 0,3333CO_2 + 13,3611H_2$$

MELISSA STOICHIOMETRY

COMPARTMENT I

Fecal protein

$$\frac{3.2C}{1.76}H_{1.76}O_{0.239}N_{0.239} + 3.035H_{2}O = \frac{C_{2}H_{4}O_{2}}{0.239} + \frac{0.1C_{4}H_{8}O_{2}}{0.239} + 2.3H_{2} + 0.76NH_{3} + \frac{0.8C}{0.8C}O_{2}$$

Bacterial protein

$$3,2CH_{1,4697}O_{0,34}N_{0,2807} + 2,712H_2O = C_2H_4O_2 + 0,1C_4H_8O_2 + 1,3162H_2 + 0.8982NH_3 + 0,8CO_2$$

Polysaccharides

$$3,199CH_{1.667}O_{0.833} + 1,134H_2O = 1C_2H_4O_2 + 0,1C_4H_8O_2 + 1,4H_2 + 0,8CO_2$$

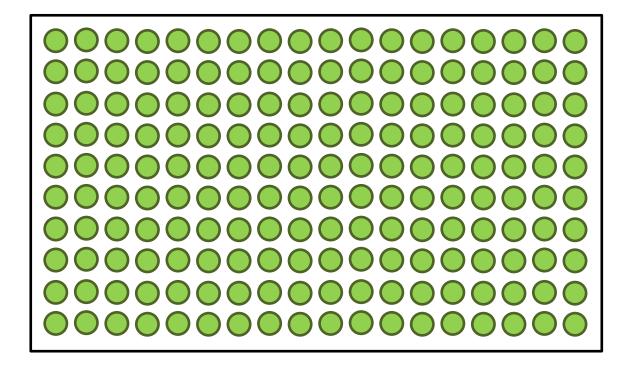
Lipids

$$C_{16}H_{32}O_2 + 13,0278H_2O = 6,5278C_2H_4O_2 + 0,6528C_4H_8O_2 + 0,33333CO_2 + 13,3611H_2$$

MELISSA MASS BALANCE

Compound	Consumed (g)	Produced (g)	Flow conservation	Delta
Bacterial Protein	13828	13828	100.0%	0.00
Fecal Protein	1148	1148	100.0%	0.00
Polysaccharides	7486	7486	100.0%	0.00
Lipids	1912	1912	100.0%	0.00
Food - higher plants	3600	3600	100.0%	0.00
Food - algae	400	400	100.0%	0.00
Acetate	20501	20501	100.0%	-0.01
Butyrate	3008	3008	100.0%	-0.01
HNO ₃	2228	2228	100.0%	-0.07
NH ₃	3487	3494	99.8%	6.90
CO ₂	13703	13788	99.4%	85.92
H ₂ O	17602	17564	100.2%	-37.95
O ₂	12186	12186	100.0%	0.00
H ₂	1018	1027	99.2%	8.18
Total	102107	102170	99.9%	62.95

1 AGENT



ATTRIBUTES

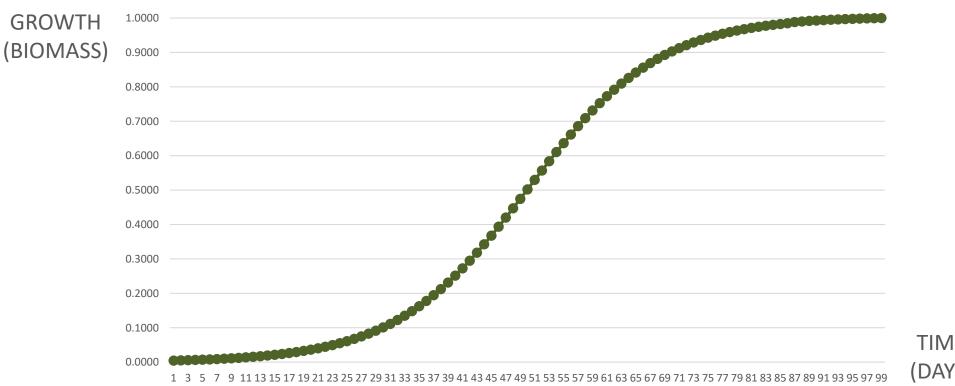
- Ideal plant: 100 day growth cycle, 40g dry weight,
 60kcal, 0.5 harvest index
- Plant plot agent: 180 plants, 1 plant plot provides enough nutrients for a crew of 6 for 1 day
- 100 day production line: 100 plant plots

BEHAVIOR

Input-output: stoichiometry

STATES

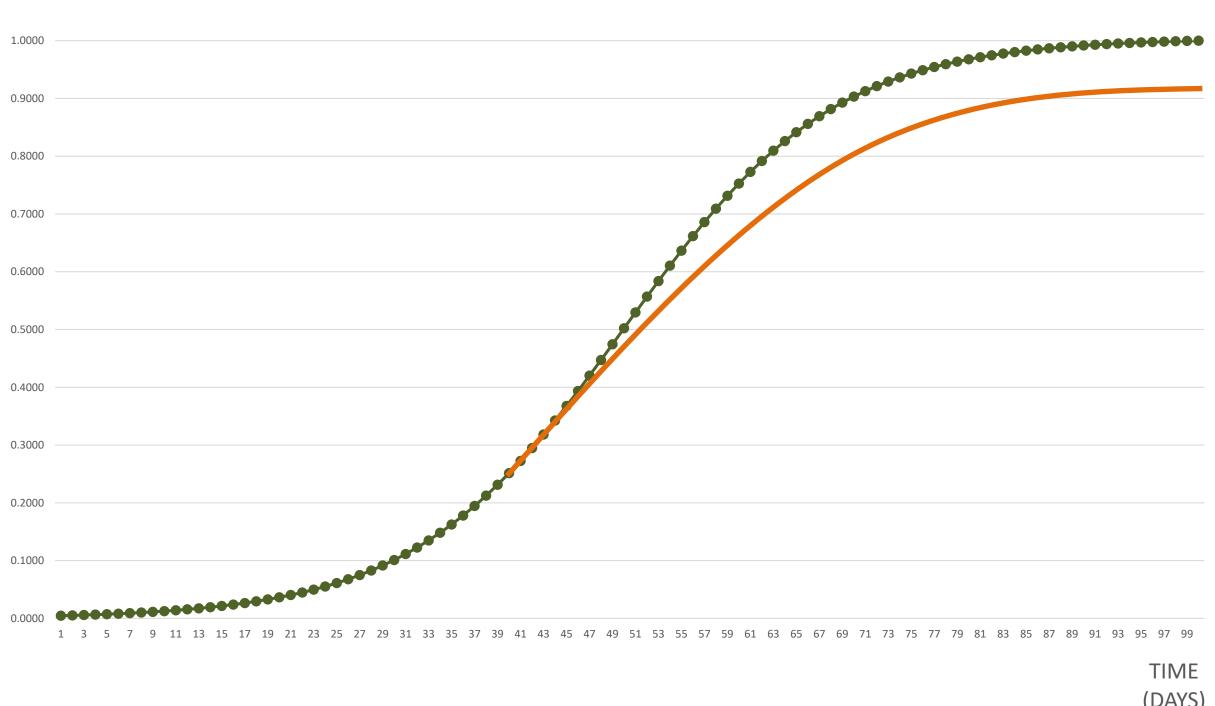
- Growth follows a sigmoid curve
- Reaching 40g in 100 days (10% first and 10% last week)
- For each day there's a specific biomass increase, and hence, the corresponding necessary input can be deduced according to the plant plot's stoichiometry



TIME (DAYS)

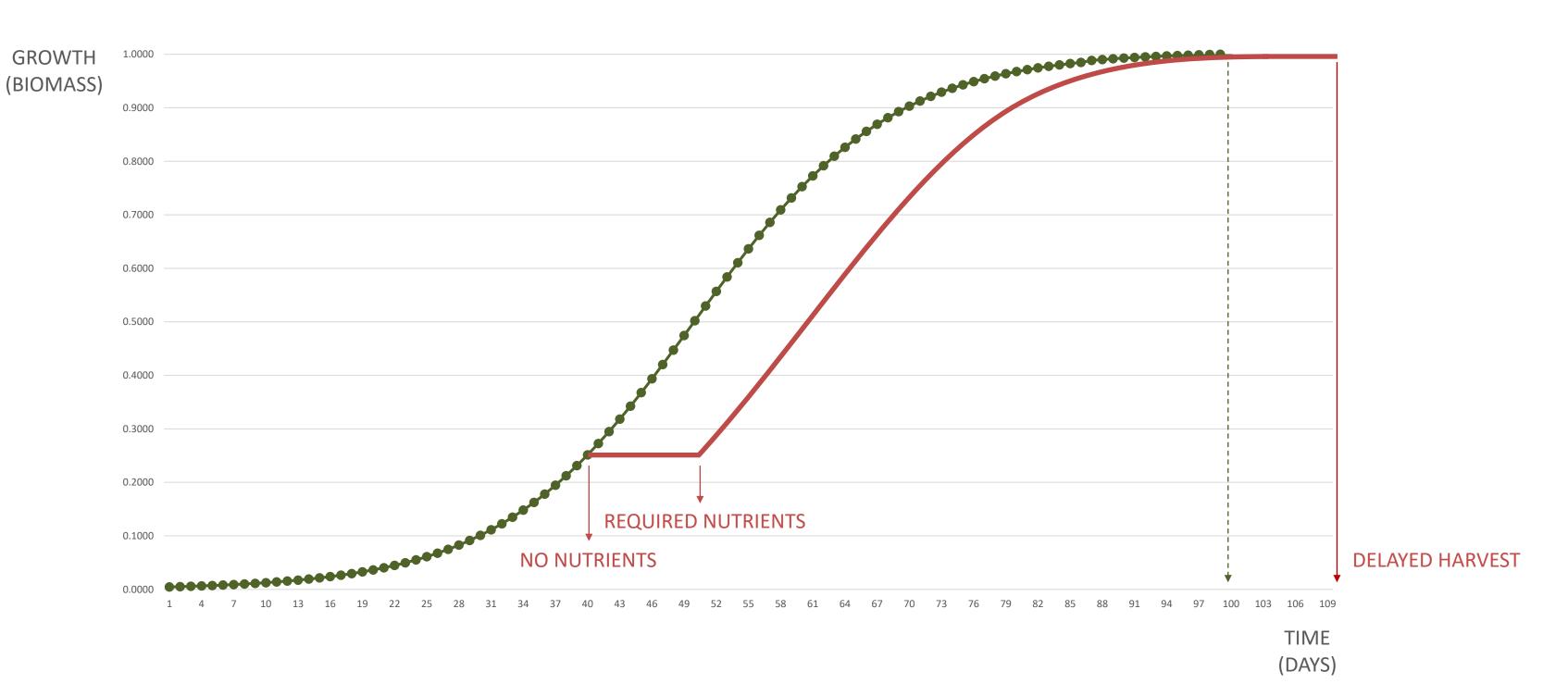
GROWTH

(BIOMASS)



REDUCED HARVEST

(DAYS)

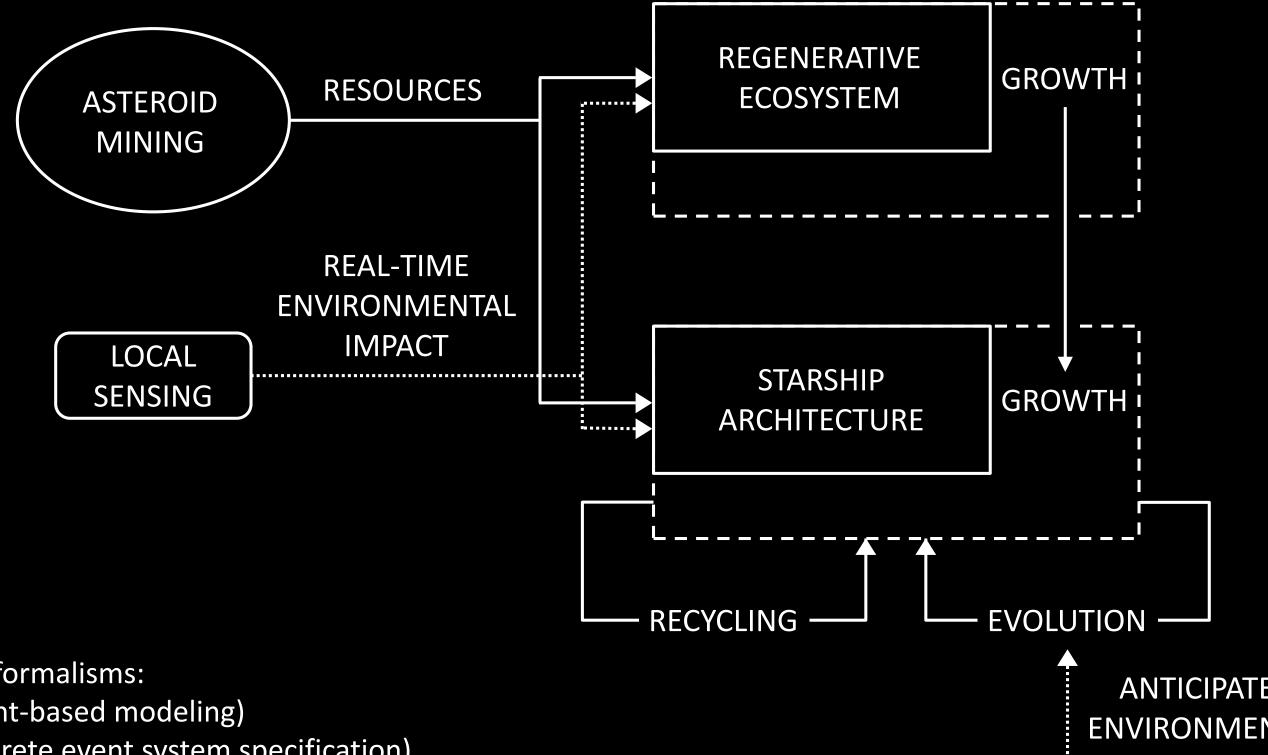


GROWTH



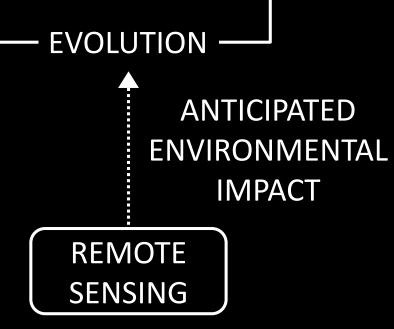
TIME (DAYS)

SIMULATION OVERVIEW



Simulation formalisms:

- ABM (agent-based modeling)
- DEVS (discrete event system specification)
- SD (system dynamics)
- EA (evolutionary algorithms)



CONCLUSIONS

- CAS approach to create a robust system
- Exploratory modeling, not predictive modeling
- Consequences: co-evolution, intractability





