Abstraction for dealing with the Multiple Realizability of Evolution

Artem Kaznatcheev

Department of Computer Science, University of Oxford
Department of Translational Hematology & Oncology Research, Cleveland Clinic

<u>egtheory.wordpress.com</u>

Use computer...

... control experiments

... visualize data

simulate experiments

genetic algorithms

computer programs

build artificial biologies

bioinformatics

Use computer...

... control experiments

... visualize data

simulate experiments

genetic algorithms

computer programs

build artificial biologies

bioinformatics

Practical skills from CS applied to the outputs of field X

Use computer...

... control experiments

... visualize data

simulate experiments

genetic algorithms computer programs

build artificial biologies

bioinformatics

Practical skills from CS applied to the outputs of field X

abstraction and multiple realizability

algorithms

Theorems, lemmas, and proofs

conceptual analysis

Mathematical techniques from CS applied to the conceptual grounding of field X

Use computer...

.. control experiments

... visualize data

simulate experiments

genetic algorithms

computer programs

Computational-X

build artificial biologies

bioinformatics

Practical skills from CS applied to the outputs of field X

abstraction and multiple realizability

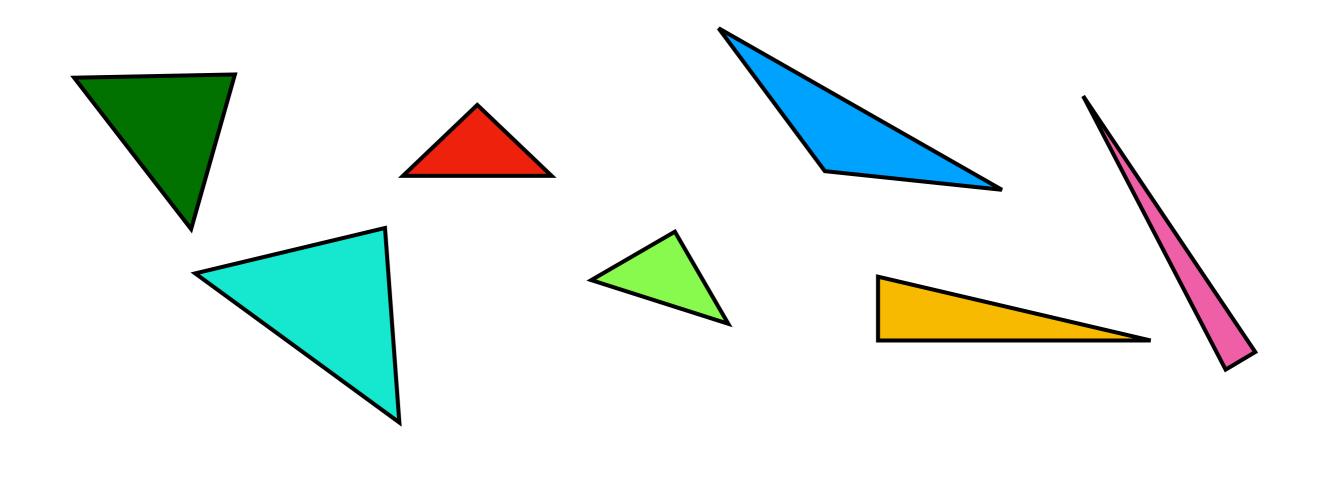
algorithms

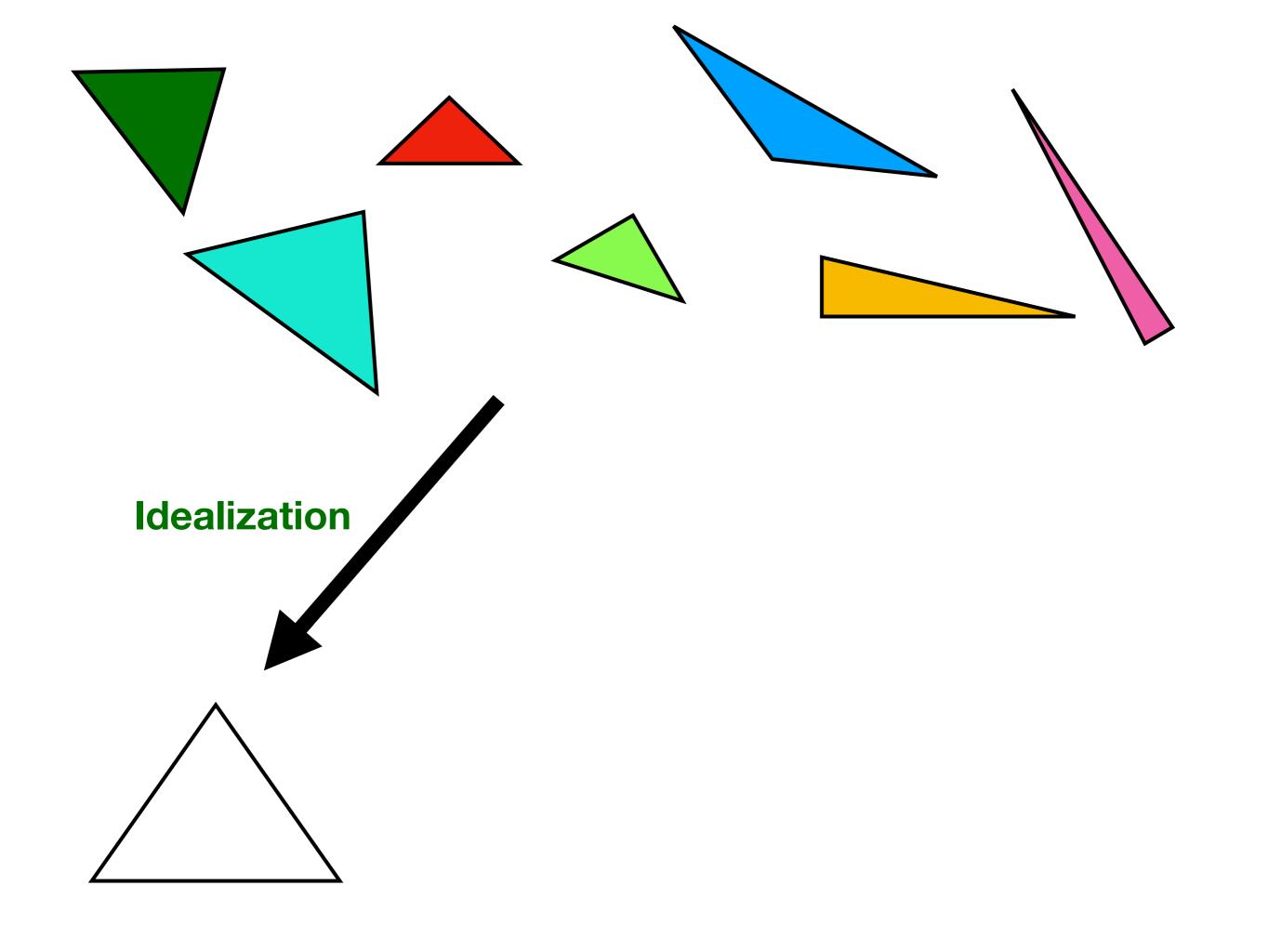
Theorems, lemmas, and proofs

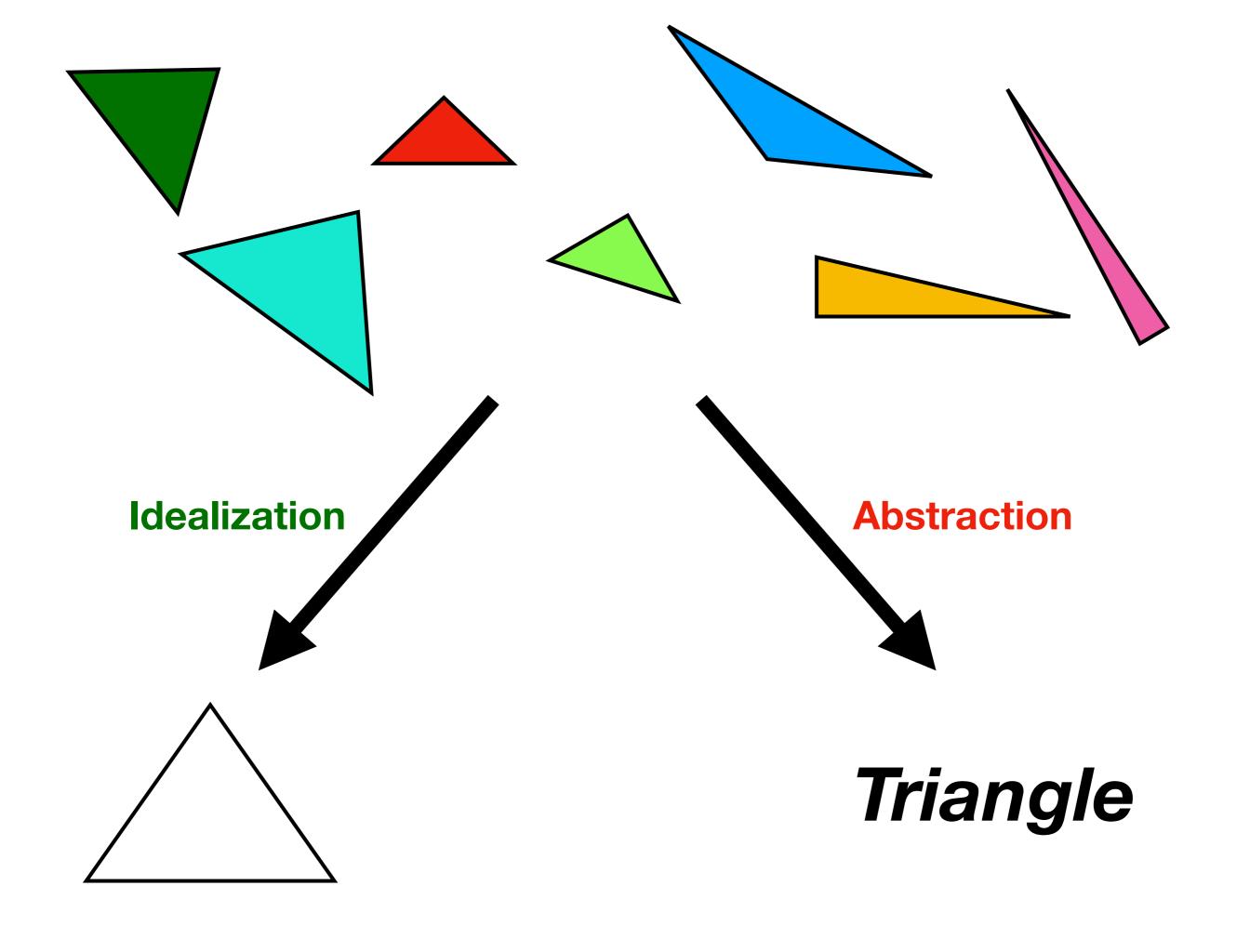
Algorithmic-X

conceptual analysis

Mathematical techniques from CS applied to the conceptual grounding of field X









ldealization

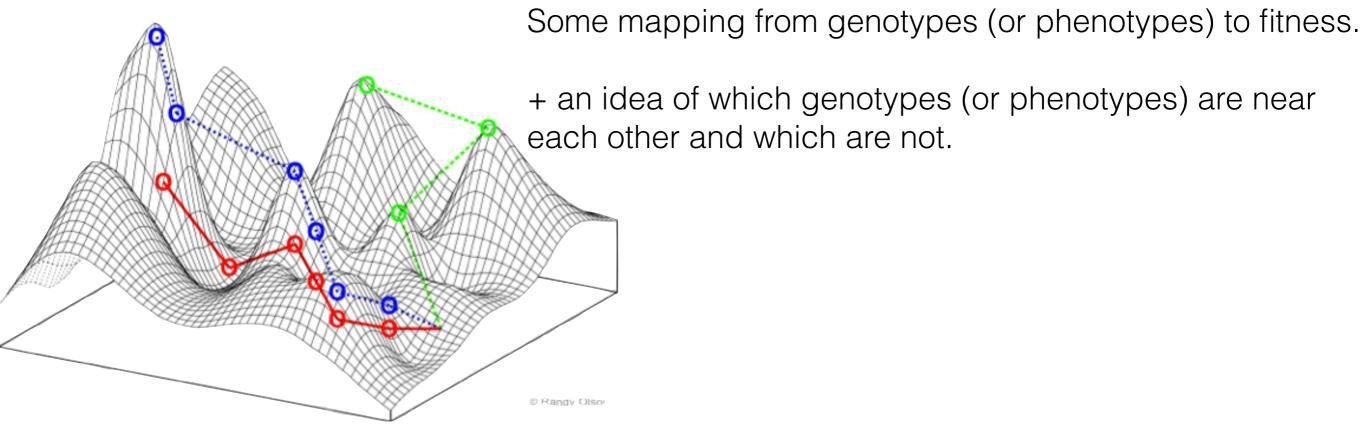
Theoretical Abstraction

Abstraction

Triangle

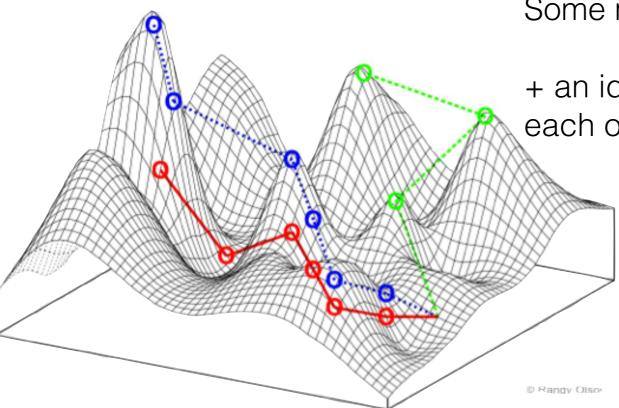
Kaznatcheev, A. (2019)

Computational complexity as an ultimate constraint on evolution **Genetics**, 302000.2019



"In a rugged field of this character selection will easily carry the species to the nearest peak"

- Wright (1932)



Some mapping from genotypes (or phenotypes) to fitness.

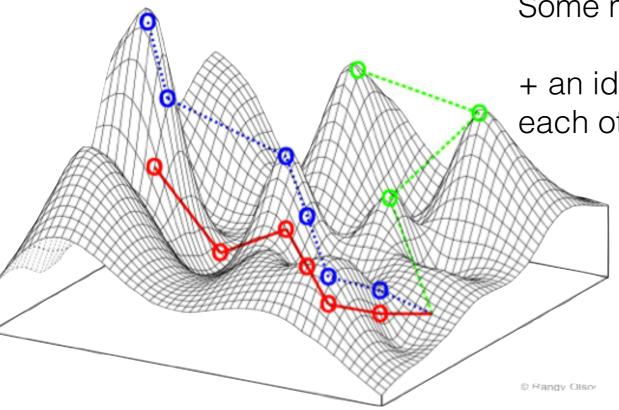
+ an idea of which genotypes (or phenotypes) are near each other and which are not.

A genotype is a **local fitness peak** if all nearby genotypes are of the same or lower fitness

A **constraint** is anything that prevents evolution from finding a local fitness peak

"In a rugged field of this character selection will easily carry the species to the nearest peak"

- Wright (1932)



Some mapping from genotypes (or phenotypes) to fitness.

+ an idea of which genotypes (or phenotypes) are near each other and which are not.

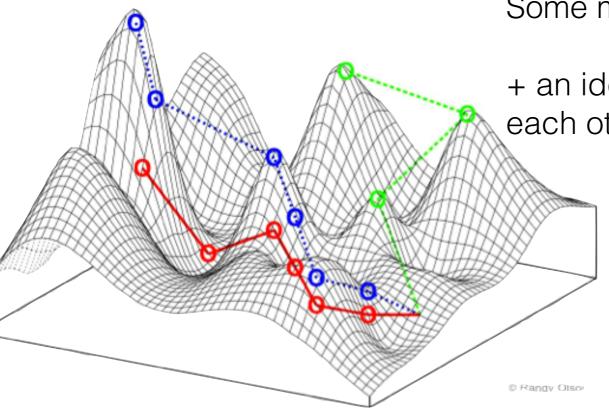
A genotype is a **local fitness peak** if all nearby genotypes are of the same or lower fitness

A **constraint** is anything that prevents evolution from finding a local fitness peak

Algorithms and Problems

Different population structures, developmental structures, trait co-variants, standing variation, etc... can produce different evolutionary dynamics and correspond to **different algorithms**

Families of different fitness landscapes correspond to **different problems**



Some mapping from genotypes (or phenotypes) to fitness.

+ an idea of which genotypes (or phenotypes) are near each other and which are not.

A genotype is a **local fitness peak** if all nearby genotypes are of the same or lower fitness

A **constraint** is anything that prevents evolution from finding a local fitness peak

Algorithms and Problems

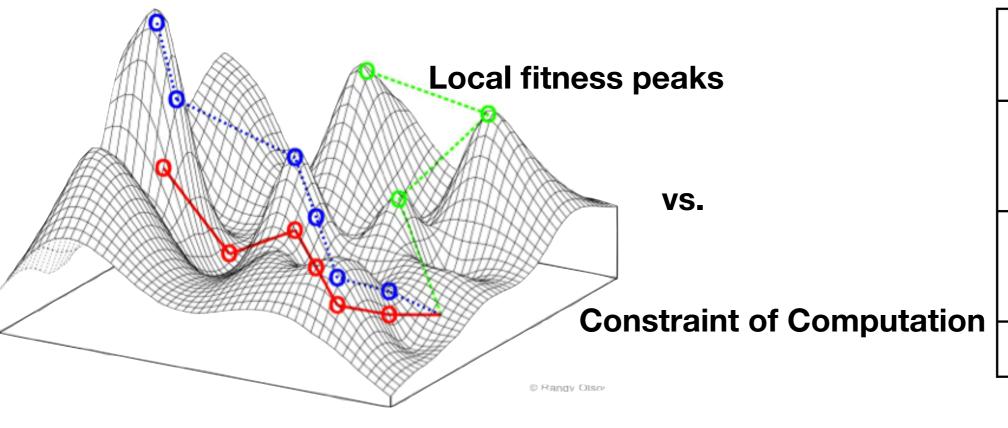
Different population structures, developmental structures, trait co-variants, standing variation, etc... can produce different evolutionary dynamics and correspond to **different algorithms**

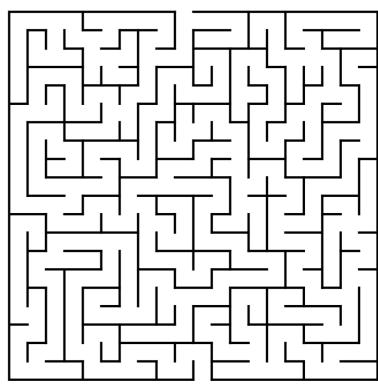
proximal constraints

Families of different fitness landscapes correspond to **different problems**

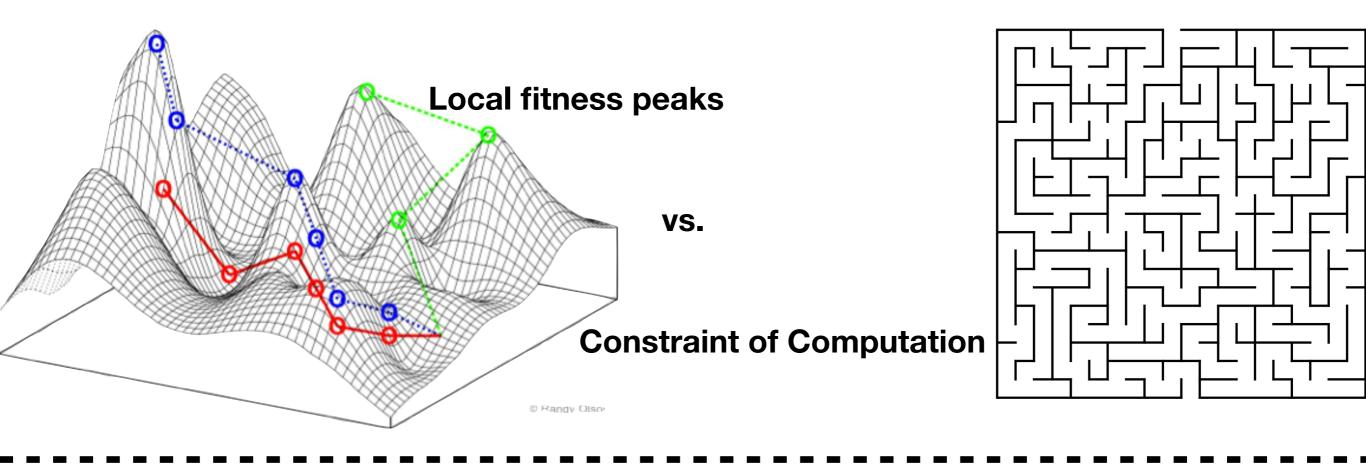
ultimate constraints

Fitness Landscapes and Constraint of Computation





Fitness Landscapes and Constraint of Computation



Kaznatcheev, A. (2019)

Computational complexity as an ultimate constraint on evolution **Genetics**, 302000.2019

Now, for any probability of failure $0 < \delta < 1$, let $m_{\delta} = \frac{\log \frac{1}{\delta}}{2 - \log 3}$ (where log is base 2).

Theorem 24. There exist semismooth fitness landscapes on $2nm_{\delta}$ loci that with probability $1-\delta$, take 2^n or more fittest mutant steps to reach their fitness peak from a starting genotype sampled uniformly at random.

Theorem 27. Finding a local optimum in the NK fitness landscape with $K \ge 2$ is PLS-complete.



Theorem 35. If $PLS \neq P$ and $log(f_{max}/f_{\delta}) \in O(n^k)$ then (for NK-model with $K \geq 2$) a local s-approximate peak cannot be found in time polynomial in n and $log(f_{max}/f_{\delta}) \in O(n^k)$ then (for NK-model with $K \geq 2$) a local s-approximate peak cannot be found in

Landscape type	Max allowed epistasis type	Hardness of reaching local optima
smooth	AB Ab aB ab magnitude	Easy for all strong-selection weak-mutation (SSWM) dynamics
semismooth	AB Ab aB ab sign	Hard for SSWM with random fitter mutant or fittest mutant dynamics
rugged	Ab aB	Hard for all SSWM dynamics: initial genotypes with all adaptive paths of exponential lengths Hard for all evolutionary dynamics (if FP != PLS) Easy for finding approximate local peaks with moderate optimality gap: selection coefficient can drop-off as power law Hard for approximate local peaks with small optimality gap: selection coefficient cannot drop-off exponentially



ldealization

Empirical Abstraction

Abstraction

Kaznatcheev, A. (2017)

Two conceptions of evolutionary games: reductive vs effective.

BioRxiv: 231993.

Kaznatcheev, A., Peacock, J., Basanta, D., Marusyk, A. & Scott, J.G. (2019)

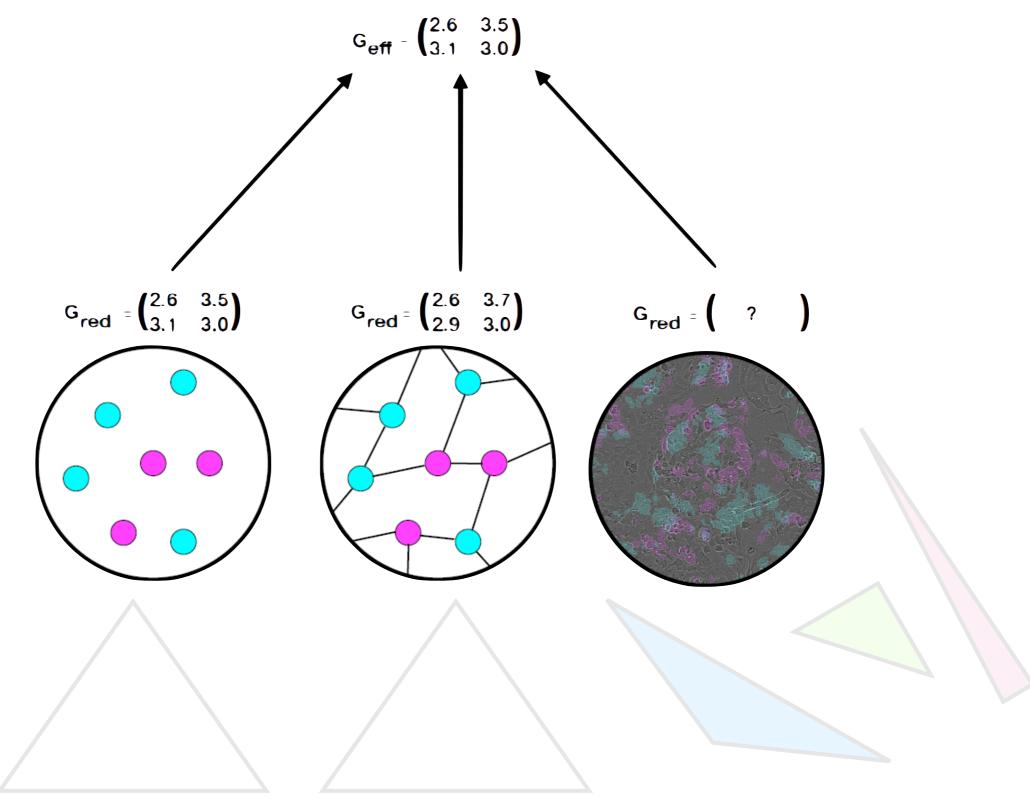
Fibroblasts and Alectinib switch the evolutionary games played by non-small cell lung cancer

Nature Ecology & Evolution 12(108): 20150154.

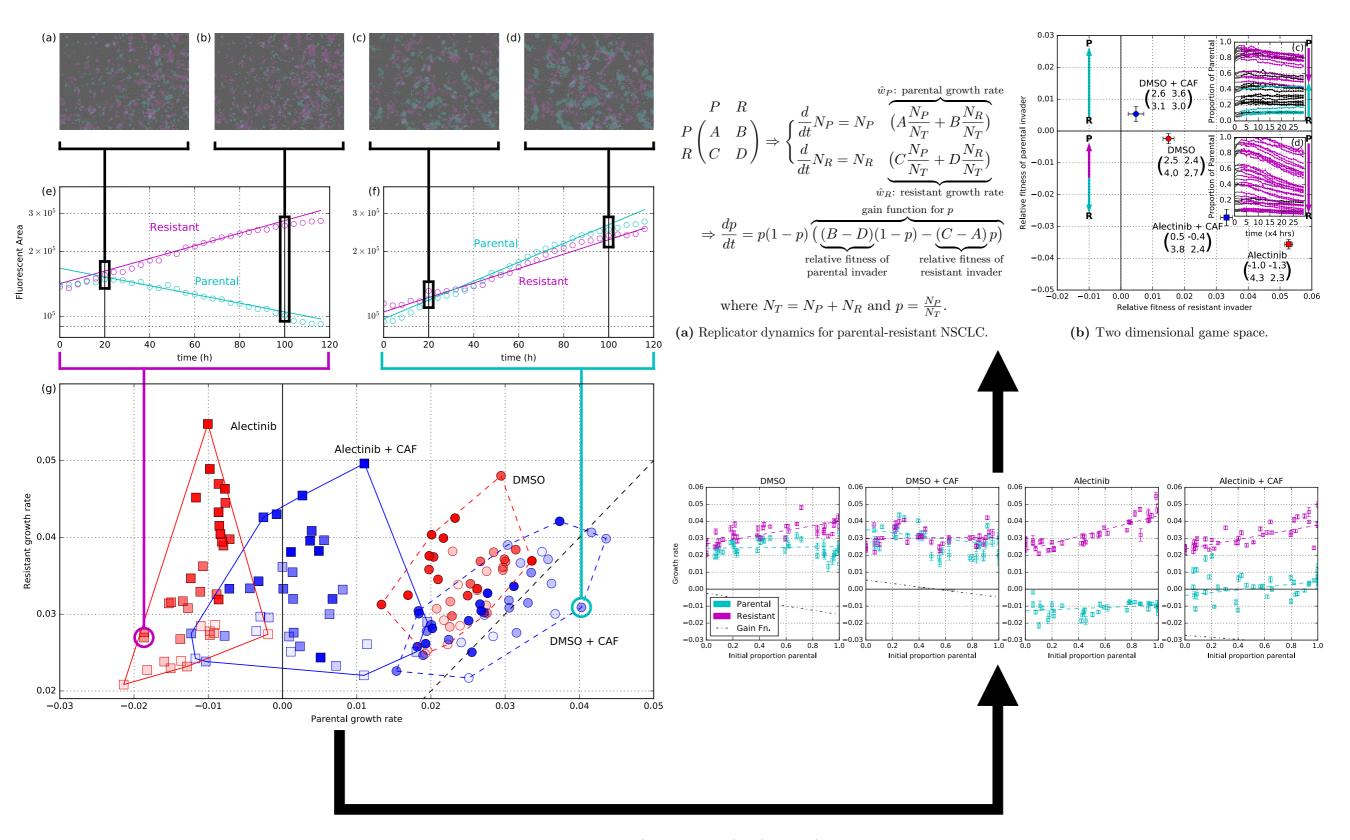
Triangle

Reductive vs effective games (in cancer)





Reductive vs effective games (in cancer)



Kaznatcheev, A., Peacock, J., Basanta, D., Marusyk, A. & Scott, J.G. (2019) Fibroblasts and Alectinib switch the evolutionary games played by non-small cell lung cancer **Nature Ecology & Evolution** 12(108): 20150154.



David Basanta



Jeffrey Peacock

Andriy Marusyk



Cleveland Clinic

Jacob G Scott





Kaznatcheev, A. (2019)

Computational complexity as an ultimate constraint on evolution **Genetics**, 302000.2019

Kaznatcheev, A. (2017)

Two conceptions of evolutionary games: reductive vs effective.

BioRxiv: 231993.

Kaznatcheev, A., Peacock, J., Basanta, D., Marusyk, A. & Scott, J.G. (2019) Fibroblasts and Alectinib switch the evolutionary games played by non-small cell lung cancer Nature Ecology & Evolution 12(108): 20150154.