

# Phenotypic Noise and the Cost of Complexity

Charles Rocabert, Guillaume Beslon, Carole Knibbe, Samuel Bernard

## ▶ To cite this version:

Charles Rocabert, Guillaume Beslon, Carole Knibbe, Samuel Bernard. Phenotypic Noise and the Cost of Complexity. EvoLyon 2019, Nov 2019, Lyon, France. hal-02402443

## HAL Id: hal-02402443 https://hal.science/hal-02402443v1

Submitted on 10 Dec 2019

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



# Phenotypic Noise and the Cost of Complexity

Charles Rocabert<sup>1,2</sup>, Guillaume Beslon<sup>1,3</sup>, Carole Knibbe<sup>1,4</sup>, Samuel Bernard<sup>1,5</sup>

1. Inria, France

Evolvable phenotypic noise

for two phenotypic characters

- 2. Synthetic and Systems Biology Unit, BRC, Szeged, Hungary
- 3. CNRS UMR5205 LIRIS, INSA Lyon, France
- 4. INSERM U1060 CarMeN, INSA Lyon, France

5. CNRS UMR5208 Institut Camille Jordan, Univ Lyon 1, France

This work was supported by the European Commission 7th Framework Programmes 659 (FP7-ICT-2013.9.6 FET Proactive: Evolving Living Technologies), and EvoEvo Project 660 (ICT-610427)

#### BACKGROUND

Theory predicts that phenotypic noise is positively selected under directional selection because it increases the mean fitness value, and counter-selected under stabilizing selection (1).

It has been suggested that under directional selection, the fitness gain provided by phenotypic noise also promotes adaptive evolution (2), while the link is unclear. Evolution on multiple phenotypic characters suffers from the cost of complexity (3). The impact of phenotypic noise in multidimensional phenotypes is less understood.

#### METHODS

We used a quantitative model to study the adaptive evolution of organisms with multiple phenotypic traits under selection and evolvable phenotypic noise (4) in a generalized fitness landscape.

#### RESULTS

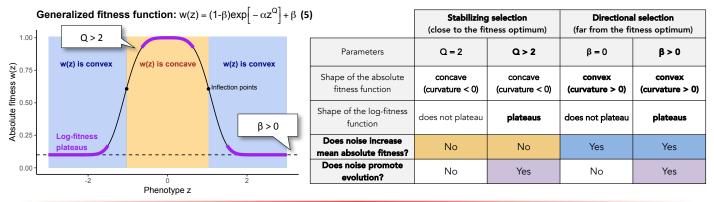
Phenotypic noise promotes adaptative evolution under directional and/or stabilizing selection if the logarithmic fitness plateaus.

For multiple phenotypic characters under selection, the phenotypic noise evolves to a one- dimensional noise aligned with the direction of the fitness optimum.

#### CONCLUSION

Phenotypic noise can decrease the cost of complexity and promotes adaptive evolution in flat regions of the fitness landscape.

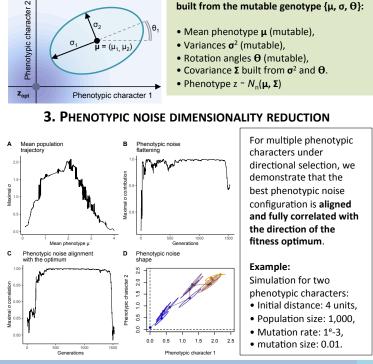
#### 1. PHENOTYPIC NOISE EFFECT ON FITNESS DEPENDS ON THE SHAPE OF THE FITNESS LANDCAPE (SINGLE CHARACTER)



#### 2. A MODEL FOR PHENOTYPIC NOISE EVOLUTION

Multi-dimensional phenotypic noise is

### 4. PHENOTYPIC NOISE PROMOTES ADAPTIVE EVOLUTION AND DECREASES THE COST OF COMPLEXITY

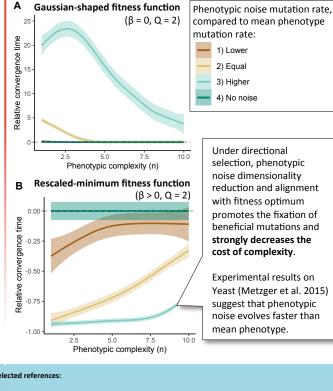


• Phenotypic noise: variability in phenotypes of isogenic organisms in constant environment, aka developmental noise, phenotypic heterogeneity, cellular noise, biological noise, intra-genotypic variability,

• Directional selection: selection far from fitness optimum characterized by a convex (positive second derivative) fitness landscape.

• Stabilizing selection: selection close to fitness optimum characterized by a concave fitness landscape. • Adaptive evolution: capacity of increasing mean population fitness as measured by the rate of

increase of the log-fitness with respect to the mean phenotype. • Cost of complexity: Reduction of the fraction of beneficial mutations when the number of phenotypic characters under selection increases.



#### Selected references:

(1) Lande (1980); Pal (1998); Kawecki (2000); Paenke et al (2007) ; Zhang et al (2009) Mol Sys Biol 5:299; Bruijning et al (2019) (2) Bódi et al (2017) PLOS Biol 15:e2000644; Duveau et al (2018) Elife 7:e37272

(3) Orr (2000) Evolution 54:13-20; Martin & Lenormand (2006) Evolution 60:893-907 (4) Ito et al (2009) Mol Sys Biol 5:264; Pelabon et al (2010) Evolution 64:1912-1925; Viñuelas et al (2012) Prog Biophys Mol Biol 110:44-53; Shen et al (2012) PLOS Genetics 8:e1002839; Metzger et al (2015) Nature 521:344-347

(5) Zhang et al (2009) Mol Sys Biol 5:299; Draghi et al (2019);