



EvoEvo Deliverable 6.2

Guillaume Beslon, Santiago F. Elena, Dominique Schneider

► **To cite this version:**

Guillaume Beslon, Santiago F. Elena, Dominique Schneider. EvoEvo Deliverable 6.2: Project Communication Media. [Research Report] INRIA Grenoble - Rhône-Alpes. 2014. <hal-01577183>

HAL Id: hal-01577183

<https://hal.archives-ouvertes.fr/hal-01577183>

Submitted on 25 Aug 2017

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.



EvoEvo Deliverable 6.2

Project Communication Media

Due date: M3 (February 2014)
Person in charge: Guillaume Beslon
Partner in charge: INRIA
Workpackage: WP6 (Management)
Deliverable description: Project communication media: Leaflet and slideshow describing the project objectives.

Revisions:

Revision no.	Revision description	Date	Person in charge
1.0	First release of the project Leaflet and Slideshow	24/01/14	G. Beslon (INRIA)
1.1	Correction of the Leaflet by Santiago Elena and Guillaume Beslon	28/01/14	G. Beslon (INRIA)
1.2	Corrections by Dominique Schneider	29/01/14	G. Beslon (INRIA)
1.3	Correction of a typo in the leaflet	12/12/14	G. Beslon (INRIA)



Table of Contents

1. Description of the communication media.....	3
1.1. Introduction.....	3
1.2. Description of the leaflet.....	3
1.3. Project slideshow	4



1. Description of the communication media

1.1. Introduction

The communication media of the project are to be used by project members to present the project and to disseminate the concepts and results. They are thus complementary from the project website (see deliverable 6.1). The first version of the project leaflet and of the project slideshow are available on the Digital Asset Management (DAM) of the EvoEvo Intranet (media directory). Both files will be updated throughout the project to disseminate project results.

1.2. Description of the leaflet

The Leaflet is organized as an A4 double side sheet that must be folded twice vertically. It is to be printed by the partners that would like to use it.

Link to the project leaflet: http://www.evoevo.eu/download/2_-_media/leaflet_A4_V1.4.pdf

PROJECT SUMMARY

Evolution is the major source of complexity on Earth at the origin of all the species we can observe, interact with or breed. On a smaller scale, evolution is at the heart of the adaptation process for many species, in particular micro-organisms (e.g. bacteria, viruses...). Microbial evolution results in the emergence of the species itself, and it also contributes to the organisms' adaptation to perturbations or environmental changes. These organisms are not only organised by evolution, they are also organised to evolve.

The EvoEvo project will develop new evolutionary approaches in information science and will produce algorithms based on the latest understanding of molecular and evolutionary biology. Our ultimate goal is to address open-ended problems, where the specifications are either unknown or too complicated to express, and to produce software able to operate in unpredictable, varying conditions.

We will start from experimental observations of micro-organism evolution, and abstract this to reproduce EvoEvo, in biological models, in computational models, and in application software. Our aim is to observe EvoEvo in action, to model EvoEvo, to understand EvoEvo and, ultimately, to implement and exploit EvoEvo in software and computational systems.

The EvoEvo project will have impact in ICT, through the development of new technologies. It will also have impact in biology and public health, by providing a better understanding of micro-organism adaptation (such as the emergence of new pathogens or the development of antibiotic resistances).

CONTACT
<http://www.evoevo.eu>
contact@evoevo.eu

EvoEvo is an Information and Communication Technologies initiative funded by the European Commission under FP7.

Project name: Evolution of Evolution
Project acronym: EvoEvo
Project reference: 610427
Programme acronym: FP7-ICT
Subprogramme area: ICT-2013.9.6
Contract type: Collaborative project (generic)
Call: EVLIT (Evolving Living Technologies)

Logos: Inria, Université Joseph Fourier, CSIC, The University of York, EvoEvo, European Union, Seventh Framework Programme.

Figure 1: first page of the project Leaflet as it is visible from the website











WORK PROGRAM	KEY CONCEPTS	PARTNERS
<p>EvoEvo will achieve its main objective through the achievement of four scientific and technological objectives organized to bridge biological knowledge to ICT applications.</p>	<p>Evolution of Evolution is a process that acts through modification of organisms genotype-to-phenotype mapping. In the project, four characteristics of mapping will be studied in real organisms, modelled through computational evolution and used in a real application.</p>	  
<p>(1) Biological domain</p> 	<p>1) BIOLOGICAL DOMAIN Observe, quantify and characterize EvoEvo in <i>Escherichia coli</i> and <i>Tobacco etch Virus</i> through experimental evolution.</p>	<p>Guillaume Beslon, INRIA Grenoble Rhône-Alpes, Beagle Team, Lyon, France INRIA is associated with INSA-Lyon, Université Claude Bernard-Lyon 1 and the LIRIS Laboratory.</p>
<p>(2) Biological models</p> 	<p>2) COMPUTATIONAL BIOLOGY Simulate "evolution of evolution" by mean of individual-based models and <i>in silico</i> experimental evolution.</p>	
<p>(3) Computational Framework</p> 	<p>3) COMPUTATIONAL FRAMEWORK Design a computational evolutionary platform to exploit EvoEvo in applicative software.</p>	<p>Dominique Schneider, Université Joseph Fourier Grenoble 1, Laboratoire Adaptation et Pathogénie des Microorganismes, Grenoble, France.</p>
<p>(4) Application domain</p> 	<p>4) APPLICATION DOMAIN Apply EvoEvo to real ICT problems and propose proofs of concept for the approaches developed in the project.</p>	 <p>Paulien Hogeweg, Utrecht University, Theoretical Biology and Bioinformatics Group, Utrecht, Nederland.</p>
	<p>SYSTEMS OF INTEREST</p>	<p>THE UNIVERSITY of York</p>
	<ul style="list-style-type: none"> • MICRO-ORGANISMS Two model micro-organisms will be studied: the bacterium <i>E. coli</i> and <i>Tobacco etch virus</i>. Both models will be studied from the genomic to the phenotype and population levels. • COMPUTATIONAL MODELS Two different simulation frameworks ("aevo" and "pearls-on-a-string") will be used and merged to create an integrated model. • APPLICATIONS Two different applications will be used as proof of concept: on-line data stream clustering and evolution of a personal companion. 	<p>Susan Stepney, University of York, York Centre for Complex Systems Analysis, York, UK</p>
		 <p>Santiago Elena, Consejo Superior de Investigaciones Científicas, Instituto de Biología Molecular y Celular de P...</p>

Figure 2: second page of the Leaflet as it is visible from the website

1.3. Project slideshow

The project slideshow is available on the website in pdf format. It presents the general concepts of the project, its organization and its expected impact.

Link to the project slideshow: http://www.evoevo.eu/download/2_-_media/EvoEvo.pdf