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Evolving Software Ecosystems

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informatique.umons.ac.be/genlog

Current Research Interests

- software evolution
- software quality
- model-driven software engineering
- empirical software engineering
- human-machine interaction
- Use of formal techniques to support the above
 - graph transformation
 - logic-based formalisms
 - model checking
 - statistical analysis
- Develop automated tool support

Ongoing Research Projects



ARC Project « Ecological Studies of Open Source Software Ecoysystems », 2012-2017

- Interdisciplinary research, using ideas from biological ecology to understand and improve evolution/maintenance of software ecosystems

FRFC Project « Data-Intensive Software System Evolution », 2013-2017

- Collaboration with University of Namur (A. Cleve)
- Empirical study of co-evolution of programs and database in data-intensive software systems

COST Action IC1404 « Multi-Paradigm Modelling for Cyber-Physical Systems », 2015-2018

FRIA PhD Scholarship « Executable Modeling of Gestural Interaction Applications », 2011-2015

- *Romuald Deshayes.* Using domain-specific modeling languages, model transformation, executable modelling through high-level Petri nets

Human-machine interaction

Executable Modeling of Gestural Interaction Applications





Controlling interactive applications (e.g. games, domotics, ...) using hand gestures



Human-machine interaction

Executable Modeling of Gestural Interaction Applications





Human-machine interaction

Executable Modeling of Gestural Interaction Applications





Research Context Software Evolution









Research Context Software Ecosystems



Chapter 10 Studying Evolving Software Ecosystems based on Ecological Models

Tom Mens, Maëlick Claes, Philippe Grosjean and Alexander Serebrenik

Research on software evolution is very active, but evolutionary principles, models and theories that properly explain why and how software systems evolve over time are still lacking. Similarly, more empirical research is needed to understand how different software projects co-exist and co-evolve, and how contributors collaborate within their encompassing software ecosystem.

In this chapter, we explore the differences and analogies between natural ecosystems and biological evolution on the one hand, and software ecosystems and software evolution on the other hand. The aim is to learn from research in ecology to advance the understanding of evolving software ecosystems. Ultimately, we wish to use such knowledge to derive diagnostic tools aiming to analyse and optimise the fitness of software projects in their environment, and to help software project communities in managing their projects better.



Research Context Software Ecosystems



- Study of macro-level software evolution
 - evolution of large collections or distributions of software projects or packages
 - E.g. forges like GITHUB, SourceForge, Savannah, Google Code

J.M. Gonzalez-Barahona et al. *Macro-level software evolution: a case study of a large software compilation.* Empirical Software Engineering 14(3): 262-285 (2009)

M. Caneill, S. Zacchiroli. *Debsources: Live and historical views on macro-level software evolution*. Int. Symp. ESEM 2014

Research Context Software Ecosystems



- Study of **macro-level** software evolution
 - evolution of large collections or distributions of software projects or packages
 - E.g. forges like GITHUB, SourceForge, Savannah, Google Code
- Study **socio-technical aspects** of the community of contributors (end-users, developers, debuggers, ...)
- Focus on coherent collections of projects or packages
 - a.k.a. software ecosystems
 - E.g. Debian, Ubuntu, GNOME, KDE, CRAN, Eclipse, ...

Software Ecosystems Definitions

- David Messerschmitt & Clemens Szyperski, 2003
 - *"a collection of software products that have some given degree of symbiotic relationships."*
- Mircea Lungu, 2008 [Ph.D. dissertation]
 - *"a collection of software projects that are developed and evolve together in the same environment."*
- Slinger Jansen et al., 2013
 - *"a set of actors functioning as a unit and interacting with a shared market for software and services, together with the relationships among them."*







Research Context



- Focus on **open source** software
- Free access to source code, defect data, developer and user communication
- Historical data available in open repositories
 - Observable communities
 - Observable activities
- Increasing popularity for personal and commercial use
- A huge range of community and software sizes





- Determine and improve the factors that drive success or failure of OSS projects within their ecosystem
- Investigate new techniques and mechanisms to predict and improve quality and survival of OSS projects
 - Inspired by research in *ecology, social network* analysis, ...

informatique.umons.ac.be/genlog/projects/ecos





 Specific questions depend on the software ecosystem under study









- Specific questions for CRAN
 - R package archive network
 - Which packages are more likely to cause, upon update, problems in dependent packages?
 - When and why is code cloned across packages?

Daniel German, Bram Adams et al. "The Evolution of the R Software Ecosystem", CSMR 2013

Maelick Claes, Tom Mens, Philippe Grosjean "On the Maintainability of CRAN Packages", CSMR-WCRE 2014 "maintaineR: web-based dashboard for maintainers of CRAN packages", ICSME 2014 "An empirical study of identical function clones in CRAN" [In preparation]



• **CRAN** (R package archive)

• R package description file format:

```
Package: pkgname
Version: 0.5-1
Date: 2004-01-01
Title: My First Collection of Functions
Authors@R: c(person("Joe", "Developer", role = c("aut", "cre"),
             email = "Joe.Developer@some.domain.net"),
          person("Pat", "Developer", role = "aut"),
          person("A.", "User", role = "ctb",
              email = "A.User@whereever.net"))
Author: Joe Developer and Pat Developer, with contributions from A. User
Maintainer: Joe Developer <Joe.Developer@some.domain.net>
Depends: R ( \geq 1.8.0 ), nlme
Suggests: MASS
Description: A short (one paragraph) description of what
  the package does and why it may be useful.
License: GPL (>= 2)
URL: http://www.r-project.org, http://www.another.url
BugReports: http://pkgname.bugtracker.url
```



irefox 🔻 🕅 🕼 CRAN Package Che	eck Results	-															
🗲 🔶 😏 😵 🔞 cran.r-p	project.org/web/c	hecks/check_sur	mmary.html														☆ 📾 🔹 🕹 🐇 🚺
Package	Version	r-devel Linux x86_64 (Debian Clang)	r-devel Linux x86_64 (Debian GCC)	r-devel Linux x86_64 (Fedora Clang)	r-devel Linux x86_64 (Fedora GCC)	r-devel OS X x86_64	<u>r-devel</u> <u>OS X</u> x86_64	<u>r-devel</u> <u>Windows</u> ix86+x86_64	r-patched Linux x86_64	<u>r-patched</u> <u>Solaris</u> <u>sparc</u>	r-patched Solaris x86	r-release Linux ix86	r-release Linux x86_64	r-release MacOS X <u>x86_64</u>	r-release Windows ix86+x86_64	<u>r-oldrel</u> <u>Windows</u> ix86+x86_64	Maintainer
<u>A3</u>	0.9.2	NOTE	NOTE	NOTE	NOTE	NOTE		NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	OK	Scott Fortmann-Roe
abc	1.8	NOTE	NOTE	NOTE*	NOTE*	NOTE*	NOTE	NOTE*	NOTE	NOTE*	NOTE	NOTE	NOTE	NOTE	NOTE*	WARN*	Michael Blum
abcdeFBA	0.4	NOTE	NOTE	NOTE	NOTE	NOTE	OK	NOTE	NOTE	NOTE	NOTE	<u>OK</u>	<u>OK</u>	OK	OK	OK	Abhilash Gangadharan
ABCExtremes	1.0	NOTE	NOTE	NOTE	NOTE	NOTE		NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	<u>OK</u>	
ABCoptim	0.13.11	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	WARN	OK	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	George Vega Yon
ABCp2	1.1	NOTE	NOTE	NOTE	NOTE	NOTE		NOTE	NOTE	<u>OK</u>	NOTE	<u>OK</u>	<u>OK</u>	<u>OK</u>	OK	OK	M. Catherine Duryea
abctools	0.2-2	NOTE	NOTE	NOTE	NOTE	NOTE	<u>OK</u>	NOTE	NOTE	NOTE	NOTE	<u>OK</u>	<u>OK</u>	<u>OK</u>	OK	OK	Matt Nunes
abd	0.2-5	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	Kevin M. Middleton
abf2	0.7-0	OK	<u>OK</u>	OK	OK	<u>OK</u>		OK	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	OK	Matthew Caldwell
abind	1.4-0	OK	OK	NOTE	NOTE	OK		OK	OK	<u>OK</u>	OK	OK	<u>OK</u>	OK	OK	OK	Tony Plate
abn	0.83	NOTE	NOTE	NOTE	NOTE	NOTE	OV	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	Fraser Lewis
abundant	1.0	NOTE	NOTE	NOTE	NOTE	NOTE	OK	NOTE	NOTE	OK	NOTE	OK	OK	OK	OK	OK	Adam J. Kothman
Accelerometry	2.0	OV	OV	OV	OV	NOTE	OK	INDIE	OV	OK	NOTE	OK	OK	OK	OK	UK MAD NI	Dane R. van Domeien
AcceptanceSampling	1.0-3	NOTE	NOTE	NOTE	NOTE	NOTE		NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	Andreas Kiermeier
accrued	1.0	OK	OK	OK	OK	OK		OK	OK	OK	OK	OK	OK	OK	OK	NOTE	Tulie Faton
ACD	1.5.3	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	Fabio Mathias Correa
Ace	0.0.8	OK	OK	NOTE	NOTE	OK	5215	OK	OK	OK	OK	OK	OK	OK	OK	OK	Brian Claggett
acepack	1.3-3.3	OK	OK	NOTE	NOTE	OK		OK	OK	OK	OK	OK	OK	OK	OK	OK	Ionathan Baron
acer	0.1.2	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	OK	OK	OK	OK	OK	Even Haug
aCGH.Spline	2.2	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	OK	Tom Fitzgerald
acm4r	1.0	NOTE	NOTE	NOTE	NOTE	NOTE	OK	NOTE	NOTE	OK	NOTE	OK	OK	OK	OK	OK	Andrea Benedetti
ACNE	0.7.0	OK	OK	OK	OK	OK	ERROR	OK	OK	OK	OK	OK	OK	OK	OK	OK	Henrik Bengtsson
acopula	0.9.2	<u>OK</u>	OK	NOTE	NOTE	OK		OK	OK	<u>OK</u>	OK	OK	<u>OK</u>	OK	OK	<u>OK</u>	Tomas Bacigal
aCRM	0.1.0	<u>OK</u>	OK	NOTE	NOTE	<u>OK</u>	OK	OK	<u>OK</u>	<u>OK</u>	OK	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	Michel Ballings
acs	1.2	OK	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	OK	<u>OK</u>	<u>OK</u>	OK	OK	OK	<u>OK</u>	OK	OK	Ezra Haber Glenn
ACTCD	1.0-0	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>		<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	Wenchao Ma
Actigraphy	1.2	NOTE	NOTE	NOTE	NOTE	NOTE		NOTE	NOTE	NOTE	NOTE	<u>OK</u>	<u>OK</u>	<u>OK</u>	OK	OK	Berkley Shands
actuar	1.1-6	NOTE	NOTE	NOTE	NOTE	NOTE		NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	Vincent Goulet
ActuDistns	3.0	NOTE	NOTE	NOTE	NOTE	NOTE		NOTE	NOTE	NOTE	NOTE	OK	OK	<u>OK</u>	OK	OK	Saralees Nadarajah
ada	2.0-3	<u>OK</u>	<u>OK</u>	NOTE	NOTE	OK		OK	OK	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	<u>OK</u>	OK	Mark Culp
adabag	3.2	OK	OK	OK	OK	OK	<u>OK</u>	OK	OK	OK	OK	OK	OK	OK	OK	OK	Esteban Alfaro
adagio AdapEnetClass	0.5.9	NOTE	NOTE	NOTE	NOTE	NOTE	OF	NOTE	NOTE	NOTE	NOTE	OK	OK	OK	OK	OK	Hans W. Borchers
AdaptFit	0.2.2	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	nasinur Kanaman Khan
AdaptFitOS	0.45	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	NOTE	OK	Manuel Wiesenfarth
AdaptiveSparsity	1.3	INOTE	HOTE	HOTE	NOTE	NOTE	NOTE		HOIL	NOTE	HOTE	TOTE	NOTE	NOTE	MOL	OK	Kristen Zvamunt
AdaptiveSparsity	14	NOTE	NOTE	NOTE	NOTE	NOTE	OK	NOTE	NOTE	OK	NOTE	NOTE	NOTE	OK	NOTE		Kristen Zygmunt
adantivetau	1.1-1	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	OK	WARN	Philin Johnson
1 MONO		Seas .	NOTE:	NOTE	Nome -	NOTE	ALCONT .	Series Series	NOTE	Seas	NOTE	NOTE:	100000	SCAS	NIOTE D	or	



- Tool support for CRAN packages: maintaineR
 - package dependency, conflict and clone analysis

Summary History Depende	ency list Dependency graph Namespace Clones					
Minimum clone AST size	Functions	Size	LOC	Packages	Hash	
10 ‡		1348	214	aplpack_1.2.7	da3f45d4292b4936	4d55f9c251285d7c
Minimum clone LOC 3 ‡ Sort packages by	<pre>{ if (missing(x)) return("bagplot, version 2012/12/05, peter wolf") if (transparency == TRUE) { return("bagphull = peter(cal bagshull = "00" = cal = "") </pre>			apipaon_1.2.9		
 Oldest first Alphabetical 	<pre>col.loophull = paste(col.loophull, "99", sep = "") col.baghull = paste(col.baghull, "99", sep = "") }</pre>		Name		Туре	Conflicts
Show only Last CRAN version RGENERATE	<pre>win <- function(dx, dy) { atan2(y = dy, x = dx) }</pre>		abc		function	gvcm.cat_1.6 forams_2.0-4 pomp_0.49-2
RCImMAWGE tsDyn het.test	RMAWGEN		cv4abo)	function	None
psychotree	Lev.		cv4pos	stpr	function	None
demography fxregime		rchange	expect	ed.deviance	function	None
PSAboot GetR	party	e en ange	postpr		function	None
mobForest RandForestGl MAclinical ndvits	JI Dfast					



- Specific questions for **Debian**
 - Open source Linux distribution
 - Which packages are more likely to cause future co-installation (CI) conflicts with other packages?
 - Can I upgrade a set of installed Debian packages without "breaking" my installation?
 - Based on a formalisation and SAT solving
 - Automated tooling coinst.irill.org

Jerome Vouillon and Roberto Di Cosmo "Broken Sets in Sotware Repository Evolution", ICSE 2013





- Specific questions for **Debian**
 - Open source Linux distribution
 - Which packages are more likely to cause future co-installation (CI) conflicts with other packages?
 - Can I upgrade a set of installed Debian packages without "breaking" my installation?
 - Based on a formalisation and SAT solving
 - Automated tooling <u>coinst.irill.org</u>
 - How do CI conflicts evolve over time?

Maelick Claes, Tom Mens, Roberto Di Cosmo "A historical analysis of Debian package co-installability conflicts", [Submitted]





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Debian historical evolution of package CI-conflicts (*testing* and *stable* distribution)





Debian historical evolution of package CI-conflicts Some results

- Ratio of CI-conflicting packages remains constant over time
- Occasional "jumps" correspond to introduction or removal of problematic packages that spread the problem to (in)direct depending packages
- The more often a package is CI-conflicting, the shorter it tends to live
- The longer it takes for a package to become CI-conflicting, the longer it tends to live
- The most likely causes of introduction or removal of CIconflicts are the introduction or removal of declared conflicts in Debian package control files

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- Specific questions for GNOME
 - Linux desktop environment
 - Which projects have a higher chance of survival?
 - How is workload distributed over different projects/contributors?
 - What is the "bus factor" risk? Who are the top contributors (for a specific activity type)?

B. Vasilescu, A. Serebrenik, M. Goeminne, T. Mens"On the variation and specialisation of workload:A case study of the GNOME ecosystem community" Empirical Software Engingeering journal, 2014.

- Ċ
- Gnome visualisation tool support
 E.g. Complicity (Neu et al., University of Lugano)



Data Extraction



Version control repositories store source code and other commits

E.g., Subversion, Git

Mailing lists for communication between developers and users

Issue tracking systems for recording bug reports and change requests

- E.g., Bugzilla, JIRA
- Question and Answer websites
 - E.g. StackOverflow

Data Extraction



Using open source MetricsGrimoire tool suite (<u>https://github.com/MetricsGrimoire</u>)

CVSAnalY

 extracts information from SVN or Git source code repository logs and stores it into relational database

MailingListStats

extracts mailing list information from mbox format



Bicho

 extracts information from issue tracking systems such as Bugzilla and JIRA



• The same contributor may use different aliases







Ordering	Rajesh Sola	Sola Rajesh				
Spelling: misspelling,	Rene Engelhard	Fene Engelhard				
diacritics, punctuation	Démurget	Demurget				
	J. A. M. Carneiro	J A M Carneiro				
Middle initials, patronyms,	Daniel M. Mueth	Daniel Mueth				
nicknames, additional surnames, incomplete names	Alexander Alexandrov Shopov	Alexander Shopov				
	Carlos Garnacho Parro	Carlos Garnacho				
	Jacob "Ulysses" Berkman	Jacob Berkman				
	A S Alam	Amanpreet Singh Alam				
Name variants:	Γιωργοσ	Georgios				
transliteration, diminutives	Mike Gratton	Michael Gratton				
Software-specific:	mrhappypants	Aaron Brown				
usernames, projects, tooling artefacts	Arturo Tena/libole2	Arturo Tena				
tooning arteraets	(16:06) Alex Roberts	Alex Roberts				
Mix	Any combina	Any combination of those				

Semi-automatic approach:

- eliminate specific quirks observed during extraction Example: "(16:06) Alex Roberts"
- compute similarity between each pair of aliases (based on Levenshtein distance)
- cluster together aliases with high similarity
- post-process manually
 rely on external information (websites)
 precise but labor-intensive

id = 17 { John Doe, Doe John, john@doe.org, john_doe@hotmail.com, john.doe@gmail.com }





- several merge algorithms exist
- the "noisier" the data, the worse they perform!
- simple algorithms have higher precision and recall than more complex ones

A Comparison of Identity Merge Algorithms for Software Repositories

Mathieu Goeminne*, Tom Mens*

Institut d'Informatique, Faculté des Sciences, Université de Mons

Science of Computer Programming 28(8), August 2013

Abstract

Software repository mining research extracts and analyses data originating from multiple software repositories to understand the historical development of software systems, and to propose better ways to evolve such systems in the future. Of particular interest is the study of the activities and interactions between the persons involved in the software development process. The main challenge with such studies lies in the ability to determine the identities (e.g., logins or e-mail accounts) in software repositories that represent the same physical person. To achieve this, different identity merge algorithms have been proposed in the past. This article provides an objective comparison of identity merge algorithms, including some improvements over existing algorithms. The results are validated on a selection of large ongoing open source software projects.

Keywords: software repository mining, empirical software engineering, identity merging, open source, software evolution, comparison

GNOME Characteristics

Dataset shared on

https://bitbucket.org/mgoeminne/sgl-flossmetric-dbmerge/downloads

FLOSSMetrics compliant MySQL database

Goeminne *et al.* "A historical dataset for GNOME contributors", MSR 2013



GNOME Characteristics



- 16 years of activity
- > 1.3M of commits
- (> 0.6M of code commits)
- > 12M of file touches
- (> 6M of code file touches)
- Mainly C, C++, Python

Bipartite contributor-project graph



GNOME Top Contributor Distribution



Who are the top GNOME contributors?

in the version repository



in the bug tracker







Top Contributor Distribution

Approach

- Analyse individual GNOME projects
- Identify core groups
 - Compute Venn diagrams of most active (top 20) persons per considered data source
 - Show % of activity attributable to each person
 - Take into account identity merges



GNOME Top Contributor Distribution





How is workload distributed over different authors and projects?





How is workload distributed over different authors and projects **per activity type**?







- Extract file information for each commit in the git repository of each GNOME project
- Associate a unique activity type *t* to each file
- Count the number of file touches





How is workload distributed over different authors and projects **per activity type**?

- Two dual views (cf. contributor-project graph)
- Distribution of workload over different *projects* per activity type
- Distribution of workload over different *authors* per activity type?





Basic Workload metric **APTW**(*a*,*p*,*t*)

number of file touches of an Author *a* for a given
Project *p* and activity Type *t*

Many derived metrics

• based on sum and Gini coefficient

Workload Metrics





GNOME Workload Metrics



Main findings

Workload is log-normally distributed over GNOME projects



GNOME Workload Metrics



Main findings

The majority of GNOME authors are involved in a very low number of file touches.



GNOME Workload Metrics





Main findings

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Highest workload is represented by coding activity, followed by activities of development documentation, translation/internationalisation, and build file creation.

Relative importance of activity types



What are the favourite activity types for GNOME?

Two dual views

- Relative importance of each activity type *per author*
- Relative importance of each activity type *per project*



Relative importance of activity types



What are the favourite activity types for GNOME?

Approach

- Use statistical tests to compare distributions
- Verify if a data set corresponding to an activity type tends to have higher values than a data set corresponding to another activity type





Relative importance of activity types

Examples of statistical comparison tests

- (Wilcoxon-)Mann–Whitney U test
- Kruskal-Wallis test

Problems with traditional statistical tests:

- Not robust to populations of unequal sizes
- Different tests can be inconsistent with each other
- Pairwise comparison of all activity types requires 78 different combinations (12 * 13 / 2)
- Traditional tests are not transitive



Relative importance of activity types

Solution:

- Use a single test that respects transitivity
- T procedure [Konietschke et al 2012]

		Pair Lower Upper p -value	
Activ	vity Developers	B-A -0.560 -0.444 0.000	A
type		C–A -0.503 -0.313 7.536e-10	\neg
A 2	$2\ 2\ 2\ 3\ 3\ 3\ 3\ 3\ 3\ 4\ 4\ 4\ 4\ 4\ 4\ 4\ 5\ 5$	D–A -0.320 -0.027 1.997e-02	
B 1	$1 \ 1 \ 1 \ 1 \ 1 \ 1 \ 1 \ 1 \ 1 \ 1 \$	C–B -0.014 0.242 9.742e-02	X
C 1	$1\ 1\ 1\ 1\ 1\ 1\ 1\ 1\ 1\ 2\ 2\ 2\ 2\ 2\ 3\ 3$	D-B 0.237 0.470 1.200e-06	, t
D 1	$1\ 1\ 2\ 2\ 2\ 2\ 2\ 2\ 3\ 3\ 3\ 3\ 3\ 4\ 4\ 4\ 4$	D–C 0.090 0.404 2.432e-03	(c)(















Heterogeneous communities



Does the relative importance of activity types differ between *frequent and occasional* authors?

Idea

Equally split the authors in two bins of more or less equal size, based on the author workload:

about 50% of all authors were involved in <14 file touches



Heterogeneous communities





Heterogeneous communities

Observations



Coders have a higher workload and are involved in less projects



Translators are less active but are involved in more projects

Can be explained in part by the use of *Damned Lies*, a Web application used to manage the localisation (I10n) activities of the GNOME project



Heterogeneous communities



Sylvia Neu *et al.* "Telling stories about GNOME with Complicity", VISSOFT 2011

Complicity is a web-based application supporting software ecosystem analysis by means of interactive visualizations.

Affectional bond view:

- size of rectangle = author's lifetime in days
- color = number of projects



Heterogeneous communities



Unverified assumptions:

- 1. Authors contributing a lot to few projects are likely to be *developers* (D)
- 2. Authors contributing less often to more projects are likely to be *translators* (T)
- 3. Authors tend to have an affectional bond to *either* development *or* translation work

Case Study: GNOME Heterogeneous communities





AW(a)

Case Study: GNOME Relative Workload



How strongly do authors focus on specific activities?

Basic measures:

- RATW(a,t)
 - = % of the total workload of author a
 dedicated to activity type t
- RAWS(a) = author specialisation
 - = Gini index of of inequality of RATW(a,t) aggregated over all activity types



Case Study: GNOME Relative Workload

How strongly do authors focus?







How strongly do authors focus?



Occasional authors tend to focus on a single activity type Case Study: GNOME Relative Workload



How strongly do authors focus? RAWS(a)



Frequent authors tend to focus on few activity types.

Case Study: GNOME Workload Distribution



Main observations for GNOME ecosystem:

- Workload is unevenly distributed over projects and authors
- Clear distinction between frequent and occasional authors
- Authors form heterogeneous subcommunities of coders and translators
- GNOME is code-centric: workload is concentrated code-related activities (coding, build files, development documentation)

GNOME Next steps



Observation: existing generic tool support does not take the specificities of the ecosystem into account, making the support suboptimal.

Having gained better understanding of the GNOME ecosystem specificities, we hope to come up with better change support mechanisms

- Dedicated to specific sub communities
 - e.g. Damned Lies application for translation community
- Estimation (of cost or effort) and prediction models (e.g. of defects) could be improved
- Tools should be able to focus on those activities/projects a contributor is interested in (based on his historic activity profile)