

Debian: The ultimate platform for neuroimaging research

Michael Hanke & Yaroslav O. Halchenko

DebConf10, New York City

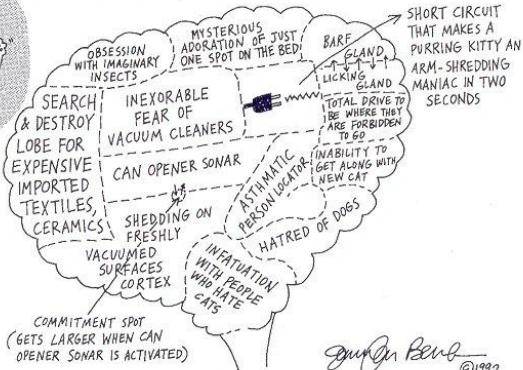




n the beginning there was just Debian. . .

. . . then there was need for two PhD degrees

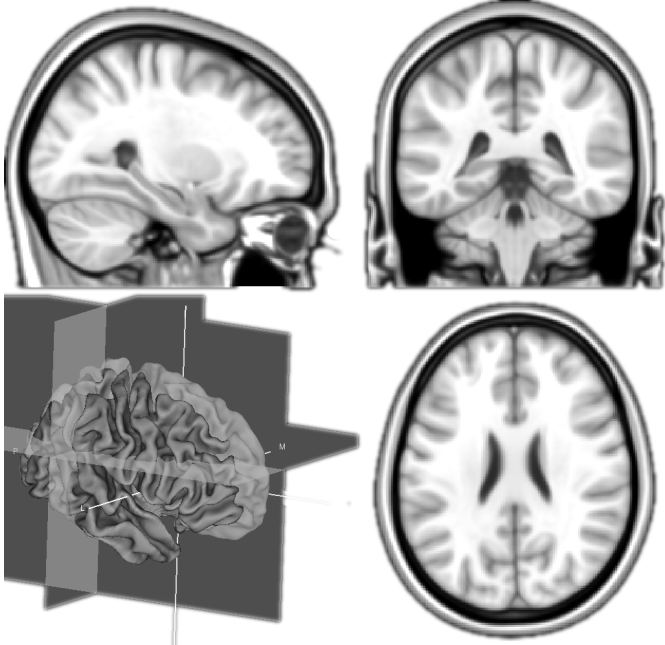
Mapping of the CAT BRAIN

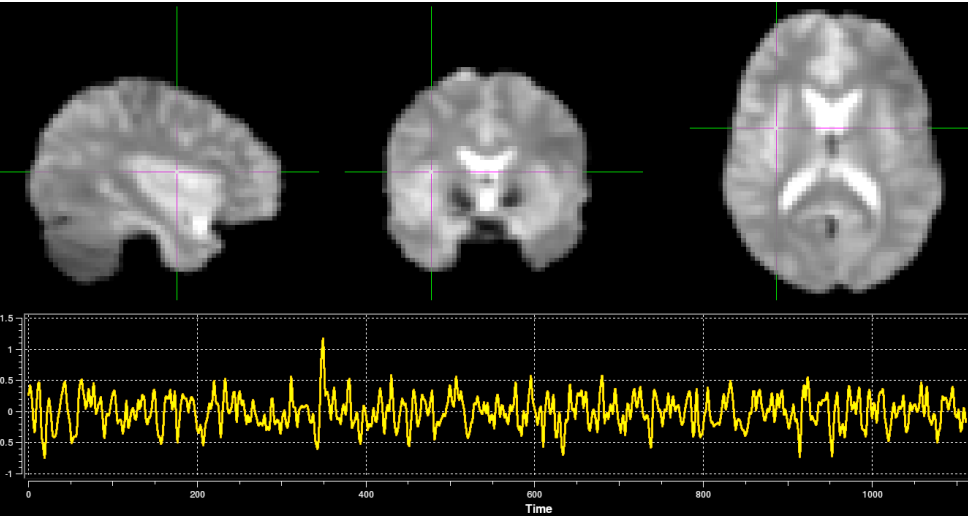


Jim Ben
©1992

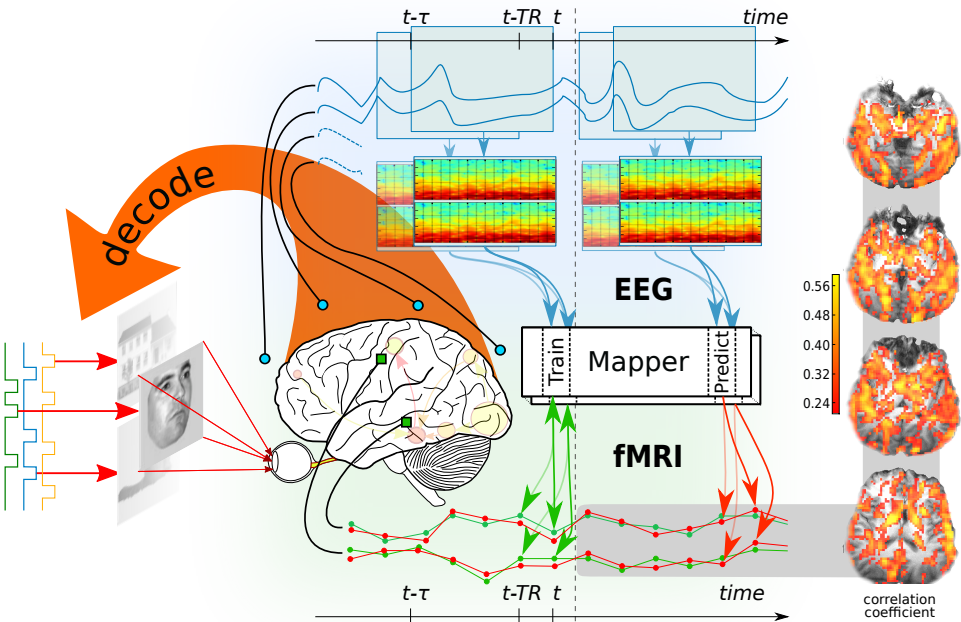


Image (C) 2008 by Jan Ainali





```
% du -h bold_run1.nii  
418M    bold_run1.nii  
% fslview bold_run1.nii &
```



Neuro-imaging in Debian: 2005



Neuro-imaging in Debian: 2005



Neuro-imaging in Debian: 2005



MINC

DICOM



Neuro-imaging in Debian: 2005



MINC

DICOM



MedCon



Praat



```
michael@meiner ~ % sudo apt-get install easy-phd  
Reading package lists... Done  
Building dependency tree  
Reading state information... Done  
E: Couldn't find package easy-phd
```



```
michael@meiner ~ % sudo apt-get install easy-phd
Reading package lists... Done
Building dependency tree
Reading state information... Done
E: Couldn't find package easy-phd
```

WNPP

- ▶ Software for data collection
(psychophysical/behavioral experiments)
- ▶ Software for data analysis (MRI file format support, algorithms, ...)
- ▶ Software for visualization (MRI-capable 3D/4D viewer, ...)

```
michael@meiner ~ % sudo apt-get install easy-phd
Reading package lists... Done
Building dependency tree
Reading state information... Done
E: Couldn't find package easy-phd
```

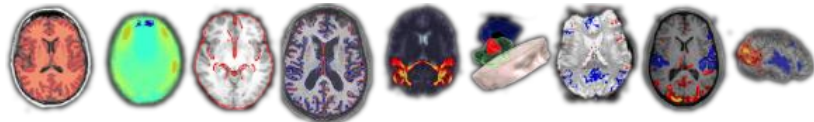
WNPP

- ▶ Software for data collection
(psychophysical/behavioral experiments)
- ▶ Software for data analysis (MRI file format support, algorithms, ...)
- ▶ Software for visualization (MRI-capable 3D/4D viewer, ...)

There is free and/or open-source software for everything

Case study: FSL

- ▶ Comprehensive analysis suite
- ▶ Well documented
- ▶ Open-source
- ▶ Downloadable for free, but non-commercial license
- ▶ Multiple developers
- ▶ Heterogeneous code
- ▶ Languages: C++, C, Tcl/Tk, sh, csh
- ▶ Public mailing list
- ▶ No public bugtracker
- ▶ No public VCS



FSL: What's in the tarball?

FOSS Visualization tool FSLView depends on Qt3
(25k lines of code)

FSL: What's in the tarball?

FOSS Visualization tool FSLView depends on Qt3
(25k lines of code)

Non-free Core analysis tools (160k lines of code)

FSL: What's in the tarball?

FOSS Visualization tool FSLView depends on Qt3
(25k lines of code)

Non-free Core analysis tools (160k lines of code)

Binary Contributed tool (only temporarily present)

FSL: What's in the tarball?

FOSS Visualization tool FSLView depends on Qt3
(25k lines of code)

Non-free Core analysis tools (160k lines of code)

Binary Contributed tool (only temporarily present)

3rd-party libiconv libgd libgdc libpng newmat newran
nifticlib zlib tcl tk (mostly outdated)

FSL: What's in the tarball?

FOSS Visualization tool FSLView depends on Qt3
(25k lines of code)

Non-free Core analysis tools (160k lines of code)

Binary Contributed tool (only temporarily present)

3rd-party libiconv libgd libgdc libpng newmat newran
nifticlib zlib tcl tk (mostly outdated)

Datasets Huge amount of atlases, templates, models
(approx. 1 GB)

FSL: What's in the tarball?

FOSS Visualization tool FSLView depends on Qt3
(25k lines of code)

Non-free Core analysis tools (160k lines of code)

Binary Contributed tool (only temporarily present)

3rd-party libiconv libgd libgdc libpng newmat newran
nifticlib zlib tcl tk (mostly outdated)

Datasets Huge amount of atlases, templates, models
(approx. 1 GB)

Preconditions for packaging

ITP/RFP: newmat, nifticlib, vtk5

FSL: Journey into Debian

Oct 2005 Preliminary core packages ready; first contact with upstream (very positive)

FSL: Journey into Debian

- Oct 2005 Preliminary core packages ready; first contact with upstream (very positive)
- May 2006 *newmat* uploaded to Debian. Thanks to Philippe Coval

FSL: Journey into Debian

- Oct 2005 Preliminary core packages ready; first contact with upstream (very positive)
- May 2006 *newmat* uploaded to Debian. Thanks to Philippe Coval
- Aug 2006 *libvtk5* becomes available in Debian. Thanks to A. Maitland Bottoms

FSL: Journey into Debian

- Oct 2005 Preliminary core packages ready; first contact with upstream (very positive)
- May 2006 *newmat* uploaded to Debian. Thanks to Philippe Coval
- Aug 2006 *libvtk5* becomes available in Debian. Thanks to A. Maitland Bottoms
- Sep 2006 *nifticlib* uploaded to Debian
dicomnifti uploaded to Debian

FSL: Journey into Debian

- Oct 2005 Preliminary core packages ready; first contact with upstream (very positive)
- May 2006 *newmat* uploaded to Debian. Thanks to Philippe Coval
- Aug 2006 *libvtk5* becomes available in Debian. Thanks to A. Maitland Bottoms
- Sep 2006 *nifticlib* uploaded to Debian
dicomnifti uploaded to Debian
- Aug 2007 *fslview* uploaded to Debian
Upstream discontinues support for Windows

FSL: Journey into Debian

- Oct 2005 Preliminary core packages ready; first contact with upstream (very positive)
- May 2006 *newmat* uploaded to Debian. Thanks to Philippe Coval
- Aug 2006 *libvtk5* becomes available in Debian. Thanks to A. Maitland Bottoms
- Sep 2006 *nifticlib* uploaded to Debian
dicomnifti uploaded to Debian
- Aug 2007 *fslview* uploaded to Debian
Upstream discontinues support for Windows
- Dec 2007 *fsl* uploaded to Debian. Upstream recommends to use the package

FSL: Journey into Debian

Oct 2005 Preliminary core packages ready; first contact with upstream (very positive)

May 2006 *newmat* uploaded to Debian. Thanks to Philippe Coval

Aug 2006 *libvtk5* becomes available in Debian. Thanks to A. Maitland Bottoms

Sep 2006 *nifticlib* uploaded to Debian
dicomnifti uploaded to Debian

Aug 2007 *fslview* uploaded to Debian
Upstream discontinues support for Windows

Dec 2007 *fsl* uploaded to Debian. Upstream recommends to use the package

Still not in Debian

FSL's data packages (data.debian.org)
Regression test-suite

Intermediate conclusions

1. Need to have a repository!

Staging area for packages that are technically ready

Essential backports researchers need latest scientific software on stable platforms

Derivatives Most neuroscientific Debian users run Ubuntu

Intermediate conclusions

1. Need to have a repository!

Staging area for packages that are technically ready

Essential backports researchers need latest scientific software on stable platforms

Derivatives Most neuroscientific Debian users run Ubuntu

But also need to push packages into Debian!

- ▶ Quality/Manpower
- ▶ Infrastructure
- ▶ Visibility

Neuro-imaging in Debian: 2010



MINC

DICOM



MedCon



Praat



Neuro-imaging in Debian: 2010



MINC

DICOM



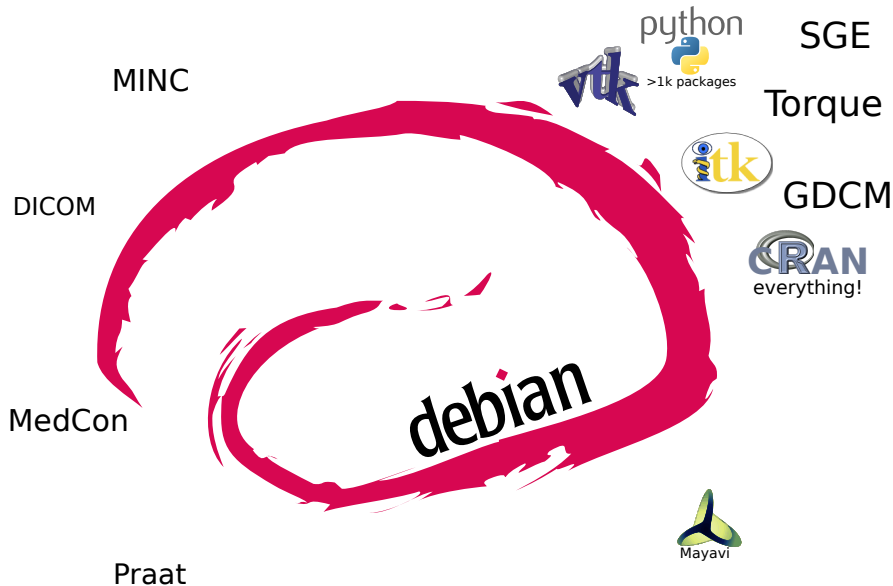
MedCon

debian



Praat

Neuro-imaging in Debian: 2010



Neuro-imaging in Debian: 2010



NIFTI-1

MINC



python



>1k packages

SGE

Torque



GDCM



debian



Mayavi

GIFTI

MGH

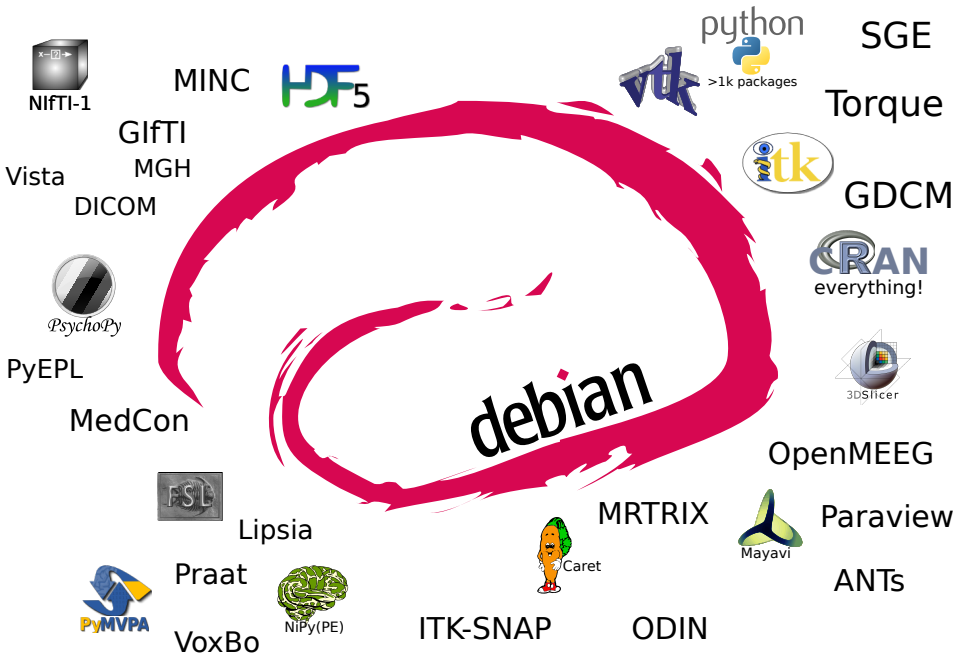
DICOM

Vista

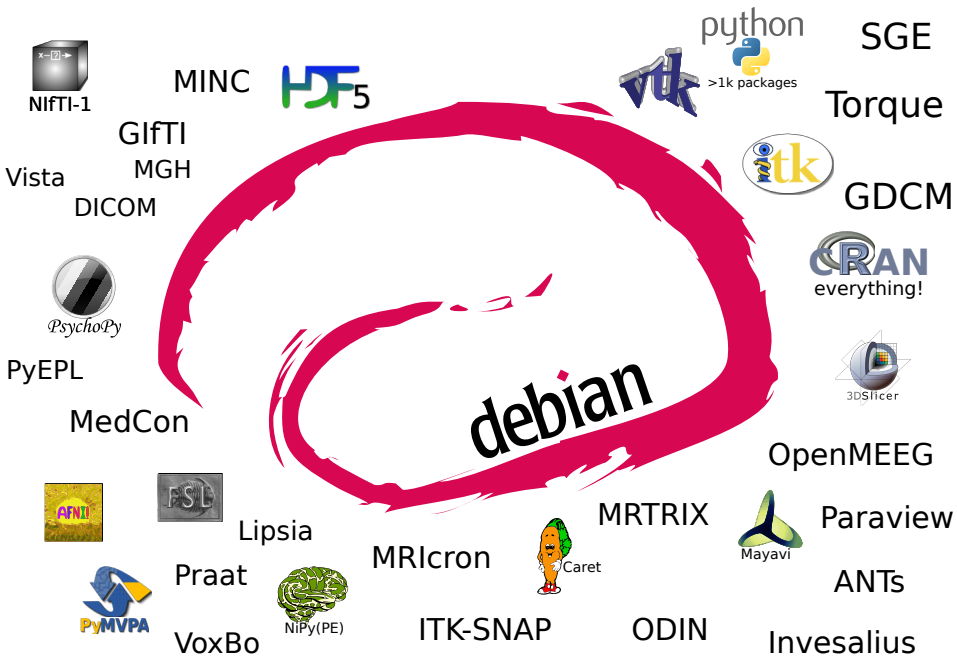
MedCon

Praat

Neuro-imaging in Debian: 2010



Neuro-imaging in Debian: 2010



Intermediate conclusions

1. Need to have a repository!

Intermediate conclusions

1. Need to have a repository!

2. Need for tailored presentations!

Availability What is available for **my** field?

Activity Are people actively working on the stuff that **I** care about?

Seriousness Do they acknowledge necessities in the scientific world (justification of development resources, proper referencing)?

Summary

Imaging

Cross-platform for visualizing, processing and analysing of bioimages

GoFigure2 is an open-source, cross-platform application for visualizing, processing and analyzing of multidimensional microscopy data. Users can visualize, segment and track cells through time, detect cell-division and ultimately generate lineages.

[Translate description](#)

The list to the right includes various software projects which are of some interest to the Debian Med Project. Currently, only a few of them are available as Debian packages. It is our goal, however, to include all software in Debian Med which can sensibly add to a high quality Debian Pure Blend.

For a better overview of the project's availability as a Debian package, each head row has a color code according to this scheme:

- ☐ Official Debian packages with high relevance
- ☐ Official Debian packages with lower relevance
- ☐ Debian packages in contrib or non-free
- ☐ Packaging has started and developers might try the packaging code in VCS
- ☐ Unofficial packages built by somebody else
- ☐ No known packages available but some record of interest (WNPP bug)
- ☐ No known packages available

Debian Med Imaging packages

Amide

Software pro obrazy v lékařství

<http://amide.sourceforge.net/>

Maintainer: Debian Med Packaging Team (Dominique Belhachemi)

Popcorn: 26 users (15 upd.)

Versions and Archs

Debtags

License: DFSG free

Official Debian package

Svn

Fix translated description

AMIDE: (Amide's a Medical Imaging Data Examiner - Amide slouží ke zkoumání lékařských obrazů) AMIDE je nástroj pro prohlížení a analýzu souborů obrázků v lékařství. Mezi jeho možnosti patří zpracovávání mnoha souborů dat zároveň, import z mnoha různých formátů, spojování obrazů, kreslení a analýza 3D oblastí podle zájmu a porovnávání.



Fsl

analysis tools for FMRI, MRI and DTI brain imaging

<http://www.fmrib.ox.ac.uk/fsl/>

Maintainer: NeuroDebian Team (Michael Hanke)

Popcorn: 21 users (21 upd.)

Versions and Archs

Debtags

License: non-free

Debian package in contrib/non-free

Git

FSL is a comprehensive library of image analysis and statistical tools for fMRI, MRI and DTI brain imaging data. The suite consists of various commandline tools, as well as simple GUIs for its core analysis pipelines. Among others, FSL offers implementations of standard GLM analysis, white matter tractography, tissue segmentation, affine and non-linear co-registration, and independent component analysis. This is a meta-package that depends on the latest FSL version.

Please register by following [this link](#) if you are using fsl.

Please cite: S.M. Smith, M. Jenkinson, M.W. Woolrich, C.F. Beckmann, T.E.J. Behrens, H. Johansen-Berg, P.R. Bannister, M. De Luca, I. Drobnjak, D.E. Flitney, R. Niazy, J. Saunders, J. Vickers, Y. Zhang, N. De Stefano, J.M. Brady, P.M. Matthews *Advances in functional and structural MR image analysis and implementation as FSL*. NeuroImage, 23:208-219 (2004)

Upload screenshot

Mni-colin27-nifti

Talairach stereotaxic space template

<http://packages.bic.mni.mcgill.ca/tgz/>

Responsible: NeuroDebian Team

License: custom, DFSG-compliant

Unofficial Debian package

This template MRI volume was created from 27 T1-weighted MRI scans of a single individual that have been transformed into the Talairach stereotaxic space. The anatomical image is complemented by a brain and a head mask. All images are in 1x1x1 mm resolution.

This package provides the template in NIFTI format.

Please cite: C.J. Holmes, R. Hoge, L. Collins, R. Woods, A.W. Toga, A.C. Evans: *Enhancement of MR images using registration for signal averaging*. J Comput Assist Tomogr, 22: 324-333 (1998)

Remark of Debian Med team: This package is waiting for the Debian data package archive to become available

[About](#)

[Social Contract](#)[Free Software](#)[Partners](#)[Donations](#)[Contact Us](#)

[News](#)

[Project News](#)[Events](#)

[Getting Debian](#)

[CD vendors](#)[CD ISO images](#)[Network install](#)[Pre-installed](#)

[Debian Packages](#)

[Documentation](#)

[Release Info](#)[Installation manual](#)[Debian Books](#)

[Support](#)

[Debian International](#)[Security Information](#)[Bug reports](#)[Mailing Lists](#)[Mailing List Archives](#)[Ports/Architectures](#)

[Miscellaneous](#)

[Help Debian](#)

[Developers' Corner](#)

[Site map](#)[Search](#)

What is Debian?

[Debian](#) is a [free](#) operating system (OS) for your computer. An operating system is the set of basic programs and utilities that make your computer run. Debian uses the [Linux](#) kernel (the core of an operating system), but most of the basic OS tools come from the [GNU project](#), hence the name GNU/Linux.

Debian GNU/Linux provides more than a pure OS: it comes with over 25000 [packages](#), precompiled software bundled up in a nice format for easy installation on your machine.

[Read more...](#)

Getting Started

The [latest stable release of Debian](#) is 5.0. The last update to this release was made on June 26th, 2010. Read more about [available versions of Debian](#)

If you'd like to start using Debian, you can easily [obtain a copy](#), and then follow the [installation instructions](#) to install it.

If you're upgrading to the latest stable release from a previous version, please read [the release notes](#) before proceeding.

To get help in using or setting up Debian, see our [documentation](#) and [support](#) pages.

Users that speak languages other than English should check the [international](#) section.

People who use systems other than Intel x86 should check the [ports](#) section.

News

Visibility Issues

Unnecessarily hard to discover how huge and diverse Debian actually is.

Visibility Issues

Unnecessarily hard to discover how huge and diverse Debian actually is.

But “raw” information is easily available

- ▶ Ultimate Debian Database and Debian Data Export
- ▶ Debian Pure Blends taskfiles

Visibility Issues

Unnecessarily hard to discover how huge and diverse Debian actually is.

But “raw” information is easily available

- ▶ Ultimate Debian Database and Debian Data Export
- ▶ Debian Pure Blends taskfiles

Feasible to remix and generate an overview tailored towards a particular audience

Visibility Issues

Unnecessarily hard to discover how huge and diverse Debian actually is.

But “raw” information is easily available

- ▶ Ultimate Debian Database and Debian Data Export
- ▶ Debian Pure Blends taskfiles

Feasible to remix and generate an overview tailored towards a particular audience – without the need to maintain additional information elsewhere



Table Of Contents

Welcome to the Debian Neuroscience Repository

- [News](#)
- [How to use this repository](#)
 - [Package authentication](#)
- [Installation](#)
 - [Virtual Machine](#)
 - [Debian](#)
- [The Team](#)
- [Contact](#)

Next topic

[Frequently Asked Questions](#)

Quick search

Go



Welcome to the Debian Neuroscience Repository

This repository provides mostly neuroscience-related packages to be used on [Debian](#) systems (or Debian-derivatives like [Ubuntu](#)). It contains both unofficial or prospective packages which are not (yet) available from the main [Debian](#) archive, as well as backported or simply rebuilt packages also available elsewhere. Please see the [Frequently Asked Questions](#) for more information about the goals of this project.

This service is provided “as is”. There is no guarantee that a package works as expected, so use them at your own risk. If you encounter problems, please [report](#) them.

Please [spread the word about NeuroDebian](#), if you like it.



News



NeuroDebian Project
NeuroDebian

Packages for the new gifticlib release 1.0.9 available: <http://neuro.debian.net/...> (and others)
6 days ago

New nifticlib 2.0.0 – fixed handling of large compressed NIFTI images: <http://neuro.debian.net/...>
6 days ago

Fresh FSL 4.1.6 is available: <http://neuro.debian.net/...> (and other packages)
18 days ago

New PyNIFTI bugfix release 0.20100607.1: <http://neuro.debian.net/...>
18 days ago



[Join the conversation](#)



Table Of Contents

fsl – analysis tools for FMRI, MRI and DTI brain imaging

- Package Details
 - Package popularity
- Binary packages
 - NeuroDebian
 - Debian
 - Ubuntu
- Packages for the complete FSL suite
- Report bugs
- Additional information
 - Usage information
 - Upgrading from FSL 3.x
 - Building binary packages yourself
 - Advanced: Arch-dependent compiler flags
 - FSL benchmarks

Previous topic

fiji – The Fiji image processing suite (based on ImageJ)

Next topic

fsl-atlases – MN152 standard space brain atlases

Quick search



fsl – analysis tools for FMRI, MRI and DTI brain imaging

FSL is a comprehensive library of image analysis and statistical tools for fMRI, MRI and DTI brain imaging data. The suite consists of various commandline tools, as well as simple GUIs for its core analysis pipelines. Among others, FSL offers implementations of standard GLM analysis, white matter tractography, tissue segmentation, affine and non-linear co-registration, and independent component analysis.

This is a meta-package that depends on the latest FSL version.

External links:



Citable reference:

S.M. Smith, M. Jenkinson, M.W. Woolrich, C.F. Beckmann, T.E.J. Behrens, H. Johansen-Berg, P.R. Bannister, M. De Luca, I. Drobnjak, D.E. Flitney, R. Niazy, J. Saunders, J. Vickers, Y. Zhang, N. De Stefano, J.M. Brady, P.M. Matthews(2004). *Advances in functional and structural MR image analysis and implementation as FSL NeuroImage*, 23:208-219. (DOI)

Note: The software authors ask users to [register](#). Available user statistics might be helpful to acquire funding for this project and therefore foster continued development in the future.

Package Details

Package popularity

- Debian [1]: 170 ([more info](#))
- Ubuntu[1]: 458
- NITRC[2]: 1051

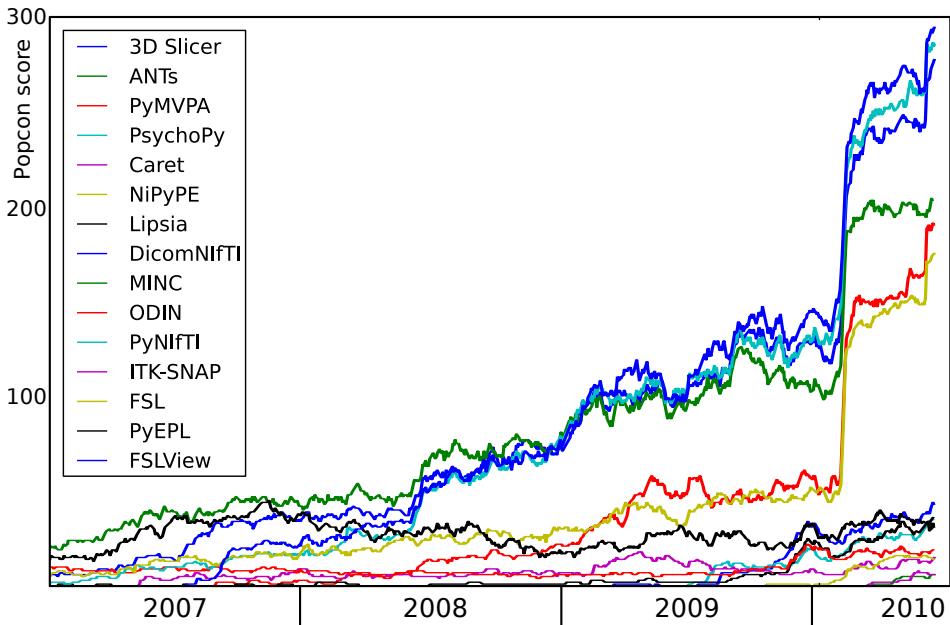
[1] (1, 2) Due to the nature of this data, the reported number can only be considered a conservative estimate of the lower bound of the true number of installations.

[2] This is the total number of downloads from NITRC for this software, comprising all releases for all platforms – typically not Debian packages.

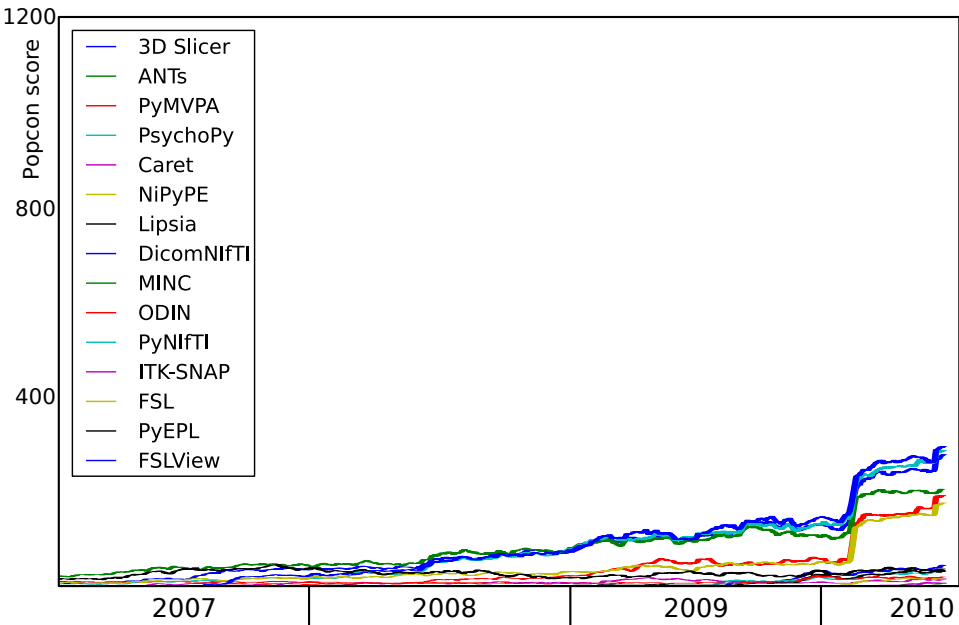
Binary packages

NeuroDebian

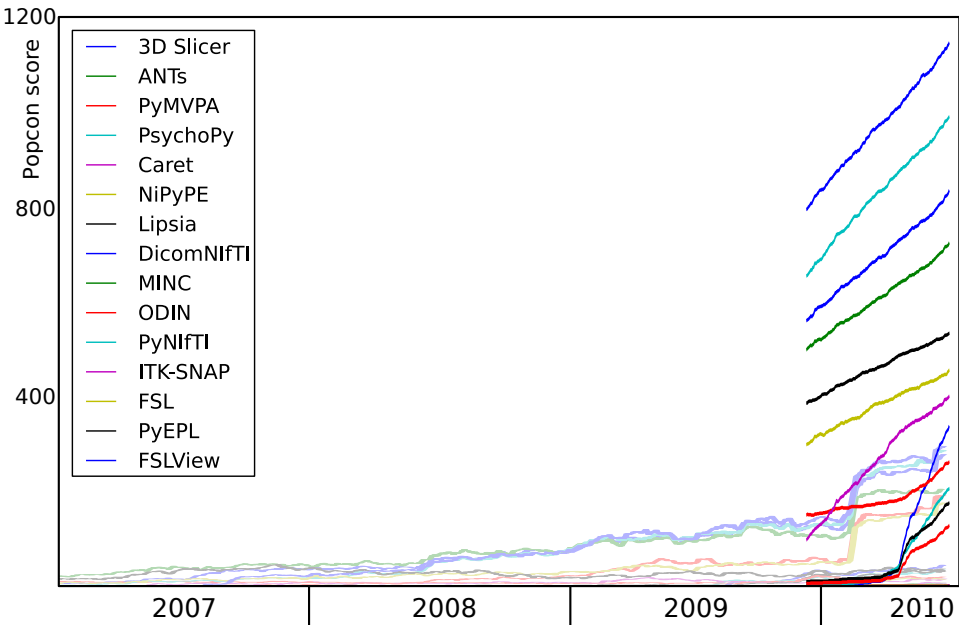
Do people use it? (Debian)



Do people use it? (Debian)



Do people use it? (Ubuntu)



The truth and nothing but...

Popcon: 1% of all users are do neuroimaging stuff

NITRC: 15% of neuroimaging researchers use
GNU/Linux

50% of those use Debian(-derivatives)

The truth and nothing but...

Popcon: 1% of all users are do neuroimaging stuff

NITRC: 15% of neuroimaging researchers use
GNU/Linux

50% of those use Debian(-derivatives)

That is not enough!

Intermediate conclusions

1. Need to have a repository!
2. Need for tailored presentations!

Intermediate conclusions

1. Need to have a repository!
2. Need for tailored presentations!
3. Need to let people know!
 - ▶ Participate in scientific mailing lists
 - ▶ Present at "native" conferences

Intermediate conclusions

1. Need to have a repository!
2. Need for tailored presentations!
3. Need to let people know!
 - ▶ Participate in scientific mailing lists
 - ▶ Present at "native" conferences
 - ▶ NeuroDebian poster at CNS2010 in Montreal

Intermediate conclusions

1. Need to have a repository!
2. Need for tailored presentations!
3. Need to let people know!
 - ▶ Participate in scientific mailing lists
 - ▶ Present at "native" conferences
 - ▶ NeuroDebian poster at CNS2010 in Montreal
 - ▶ Upcoming Debian booth at SfN2010 in San Diego

Intermediate conclusions

1. Need to have a repository!
2. Need for tailored presentations!
3. Need to let people know and try!
 - ▶ Participate in scientific mailing lists
 - ▶ Present at "native" conferences
 - ▶ NeuroDebian poster at CNS2010 in Montreal
 - ▶ Upcoming Debian booth at SfN2010 in San Diego
 - ▶ Virtual machine image with Debian lenny and some neuro-tools

Intermediate conclusions

1. Need to have a repository!
2. Need for tailored presentations!
3. Need to let people know and try!

- ▶ Participate in scientific mailing lists
- ▶ Present at "native" conferences

▶ NeuroDebian poster at CNS2010 in Montreal

brilliant! ... NeuroDebian lets you try out all the cool toys of neuroscience research with a very straightforward ease of use (esp. with the **virtual machine**). very nice...keep up the good work!

Best part for me is that I can run it in a **VM** without having to worry about replacing my native OS.

–Anonymous reviewers at www.nitrc.org

Intermediate conclusions

1. Need to have a repository!
2. Need for tailored presentations!
3. Need to let people know and try!
 - ▶ Participate in scientific mailing lists
 - ▶ Present at "native" conferences
 - ▶ NeuroDebian poster at CNS2010 in Montreal
 - ▶ Upcoming Debian booth at SfN2010 in San Diego
 - ▶ Virtual machine image with Debian lenny and some neuro-tools

Intermediate conclusions

1. Need to have a repository!
2. Need for tailored presentations!
3. Need to let people know and try!
4. **Need to work together!**

Debian



Developers



Well-engineered software

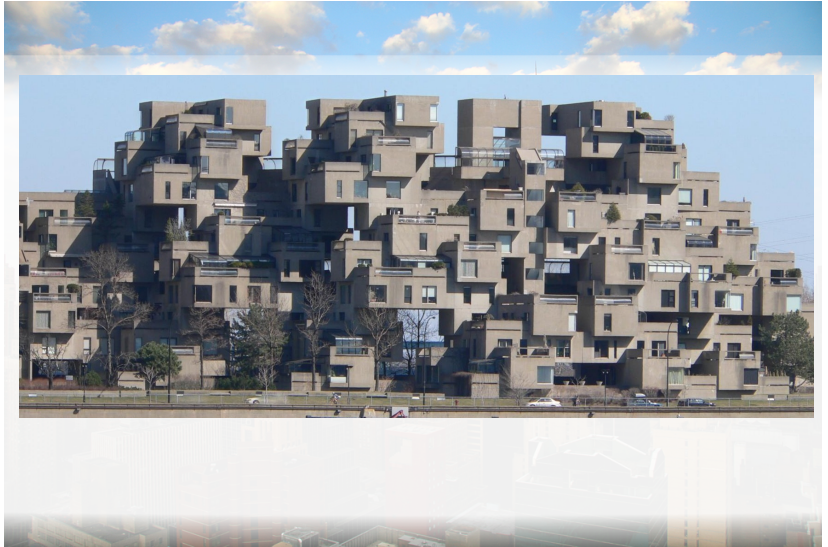


Image by Lewis Hine, 1930



Image by Ad Meskens

Software



Debian Community



Debian



Frequent pitfalls of scientific FLOSS

- ▶ Absent software engineering background or experience
- ▶ 1-man show or
1-adviser show with 2-year rotation students
- ▶ Ivory tower development model
- ▶ Absent consistency in coding, structuring, versioning/releases
- ▶ Absent unit- and/or regression testing
- ▶ Illiteracy or confusion in licensing
- ▶ Lack of man-power for proper support

The Naked CIO: Is open source dead?

“Open source lacks **true and defined standards, best-of-breed capabilities, fully functional integration and knowledgeable staff to support it cost-effectively.**”

Debian in CIO eyes

Open-source lacks **Debian provides** true and **clearly** defined **standards**, best-of-breed **capabilities**, fully functional **integration**, robust delivery and maintenance **infrastructure**, and knowledgeable staff **community** to **support** it cost-effectively.

Debian as an ecosystem

Debian provides **an ecosystem** with clearly defined **standards**, best-of-breed **capabilities**, fully functional **integration**, robust delivery and maintenance **infrastructure**, and knowledgeable community to **support** it cost-effectively.

Standards

*The Debian packaging is rather more involved.
You have to read whole books on Debian
policies.*

–Jon Peirce (PsychoPy)

Standards

*The Debian packaging is rather more involved.
You have to read whole books on Debian
policies.*

–Jon Peirce (PsychoPy)

*Debian New Maintainers' Guide: Chapter 1 -
Getting started The Right Way
The following is the very important
documentation which you should read along
with this document:*

- ▶ *Debian-policy*

Standards

The Debian packaging is rather more involved. You have to read whole books on Debian policies.

–Jon Peirce (PsychoPy)

*Debian New Maintainers' Guide: Chapter 1 -
Getting started The Right Way
The following is the very important
documentation which you should read along
with this document:*

- ▶ *Debian-policy*
- ▶ *Developers-reference*

Standards

The Debian packaging is rather more involved. You have to read whole books on Debian policies.

–Jon Peirce (PsychoPy)

*Debian New Maintainers' Guide: Chapter 1 -
Getting started The Right Way
The following is the very important
documentation which you should read along
with this document:*

- ▶ *Debian-policy*
- ▶ *Developers-reference*
- ▶ *Autotools Tutorial*

Standards

*The Debian packaging is rather more involved.
You have to read whole books on Debian
policies.*

–Jon Peirce (PsychoPy)

*Debian New Maintainers' Guide: Chapter 1 -
Getting started The Right Way
The following is the very important
documentation which you should read along
with this document:*

- ▶ *Debian-policy*
- ▶ *Developers-reference*
- ▶ *Autotools Tutorial*
- ▶ *GNU-standards*

Standards

- ▶ Constitution, Social Contract and DFSG
 - ▶ Robust democracy
 - ▶ Legal assurance clearinghouse

Standards

- ▶ Constitution, Social Contract and DFSG
 - ▶ Robust democracy
 - ▶ Legal assurance clearinghouse
- ▶ Debian Policy
 - ▶ Compliance with common standards (FHS, GNU Coding)
 - ▶ Uniform and robust deployment

Standards

- ▶ Constitution, Social Contract and DFSG
 - ▶ Robust democracy
 - ▶ Legal assurance clearinghouse
- ▶ Debian Policy
 - ▶ Compliance with common standards (FHS, GNU Coding)
 - ▶ Uniform and robust deployment
- ▶ Developer Reference(s) and Guides
 - ▶ Coherent development
 - ▶ Efficient infrastructure utilization
- ▶ HOWTOs, Wiki, ...

Standards

- ▶ Constitution, Social Contract and DFSG

- ▶ Robust democracy
- ▶ Legal assurance clearinghouse

- ▶ Debian Policy

ANTS-2 will follow ITK coding style and meet
(Neuro)Debian distribution standards

- ▶ Uniform and robust deployment

–Brian Avants (ANTs)

- ▶ Developer Reference(s) and Guides

- ▶ Coherent development
- ▶ Efficient infrastructure utilization

- ▶ HOWTOs, Wiki, ...

Standards Capabilities



NIFTI-1

MINC



python
>1k packages

SGE

Torque



GDCM

CRAN
everything!



3DSlicer

OpenMEEG

Paraview

ANTs

MRTRIX



Mayavi



Caret

ITK-SNAP

ODIN



NiPy(PE)

Praat

VoxBo



Lipsia



PyMVPA

MedCon

PyEPL



PsychoPy

DICOM

GIFTI

MGH

Vista



debian

Standards Capabilities



NIFTI-1

MINC



python



>1k packages

SGE

Torque



GDCM



If you use Debian or Ubuntu, NeuroDebian is indispensable. Many of these tools are a huge hassle to install and maintain by hand.

—Anonymous reviewer at www.nitrc.org

debian



OpenMEEG

MRTRIX



Paraview

ANTs



Caret

ITK-SNAP

ODIN



NiPy(PE)

Praat

VoxBo



GIFTI

MGH

DICOM

Vista

PyEPL

MedCon



Lipsia

Psychopy

Standards Capabilities Integration Infrastructure

Debian is not just the largest software archive, it is an ecosystem (symbiotic human/software being) regulated by the standards and common practices

- ▶ Uniform deployment mechanisms (installation, upgrades)
- ▶ Distribution-wide QA, transitions, BTS

Debian is the largest deployment facility with transparent facilities to reach the target audience in seconds

- ▶ archiving, delivery, transitions, snapshotting
- ▶ i18n
- ▶ buildd (diverse architectures), rebuilds

Debian maintainers are deployment experts

- ▶ Taking deployment burden (and bugreports) away
- ▶ Eating their own ..., since often they are users as well

Debian maintainers are deployment experts

- ▶ Taking deployment burden (and bugreports) away
- ▶ Eating their own ..., since often they are users as well
- ▶ Facilitating expertise transfer

Debian maintainers are deployment experts

- ▶ Taking deployment burden (and bugreports) away
- ▶ Eating their own ..., since often they are users as well

I have always found my friends Debian developers to be pretty good at getting me do boring but useful stuff.

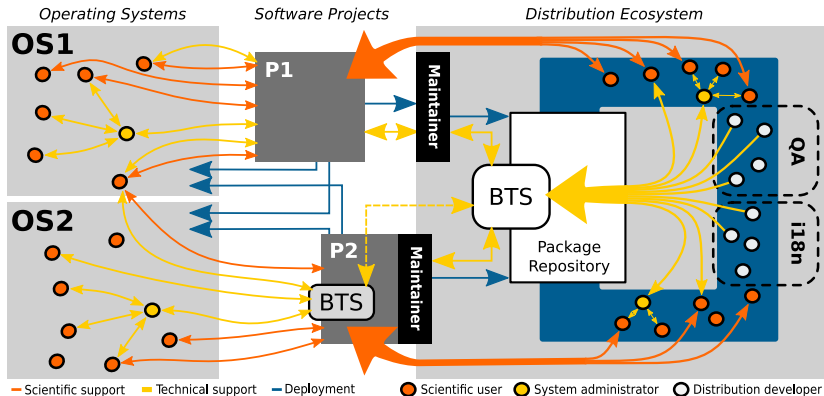
–Gael Varoquaux (mayavi2, joblib, ...)

Debian maintainers are deployment experts

- ▶ Taking deployment burden (and bugreports) away
- ▶ Eating their own ..., since often they are users as well
- ▶ Facilitating expertise transfer
- ▶ Science Deployment Guide:
<http://github.com/neurodebian/SciDeployGuide>

Standards Capabilities Integration Infrastructure Support

Standards Capabilities Integration Infrastructure Support



Conclusion

Debian is the ultimate platform for neuroimaging research because

Conclusion

Debian is the ultimate platform for neuroimaging research because

It got electrolytes

Conclusion

Debian provides an **ecosystem** with clearly defined **standards**, best-of-breed **capabilities**, fully functional **integration**, robust delivery and maintenance **infrastructure**, and knowledgeable community to **support** it cost-effectively.

Debian

- ▶ is rich and versatile
- ▶ is tailored toward neuroimaging research
- ▶ provides robust deployment infrastructure
- ▶ works together with upstream
- ▶ raises standards in scientific FLOSS

To become stronger we need to provide...

- ▶ Better coverage of neuroscience software

To become stronger we need to provide...

- ▶ Better coverage of neuroscience software
- ▶ Targetted web-presence and release-notes

To become stronger we need to provide...

- ▶ Better coverage of neuroscience software
- ▶ Targetted web-presence and release-notes
- ▶ Co-installability of multiple version

To become stronger we need to provide...

- ▶ Better coverage of neuroscience software
- ▶ Targetted web-presence and release-notes
- ▶ Co-installability of multiple version
- ▶ Convenience in referencing the works
(come back in 10 minutes)

To become stronger we need to provide...

- ▶ Better coverage of neuroscience software
- ▶ Targetted web-presence and release-notes
- ▶ Co-installability of multiple version
- ▶ Convenience in referencing the works
(come back in 10 minutes)
- ▶ Means for reproducible research
(Version-Control & VM Snapshotting)

To become stronger we need to provide...

- ▶ Better coverage of neuroscience software
- ▶ Targetted web-presence and release-notes
- ▶ Co-installability of multiple version
- ▶ Convenience in referencing the works
(come back in 10 minutes)
- ▶ Means for reproducible research
(Version-Control & VM Snapshotting)
- ▶ Large data packages

To become stronger we need to provide...

- ▶ Better coverage of neuroscience software
- ▶ Targetted web-presence and release-notes
- ▶ Co-installability of multiple version
- ▶ Convenience in referencing the works
(come back in 10 minutes)
- ▶ Means for reproducible research
(Version-Control & VM Snapshotting)
- ▶ Large data packages
- ▶ Extended QA: Infrastructure for regression and heavy testing

To become stronger we need to provide...

- ▶ Better coverage of neuroscience software
- ▶ Targetted web-presence and release-notes
- ▶ Co-installability of multiple version
- ▶ Convenience in referencing the works
(come back in 10 minutes)
- ▶ Means for reproducible research
(Version-Control & VM Snapshotting)
- ▶ Large data packages
- ▶ Extended QA: Infrastructure for regression and heavy testing
- ▶ Further dissemination of Debian
(Debian booths, papers)

Global Positioning Coordinates

Contact

<http://neuro.debian.net>
team@neuro.debian.net

Acknowledgements

Debian Med/Science
Pure Blends Developers
Stephen José Hanson
James V. Haxby
Stefan Pollman



Brain Download:

iz compltes.

ICANHASCHEEZBURGER.COM

