Research Software Directory

Dr. Jason Maassen Netherlands eScience Center

netherlands Science center

> ESCAPE-OSSR Meeting 15/06/2023



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Science Park Amsterdam





~270 projects (on many different topics)

Astronomy: fast radio bursts

In collaboration with ASTRON & UvA

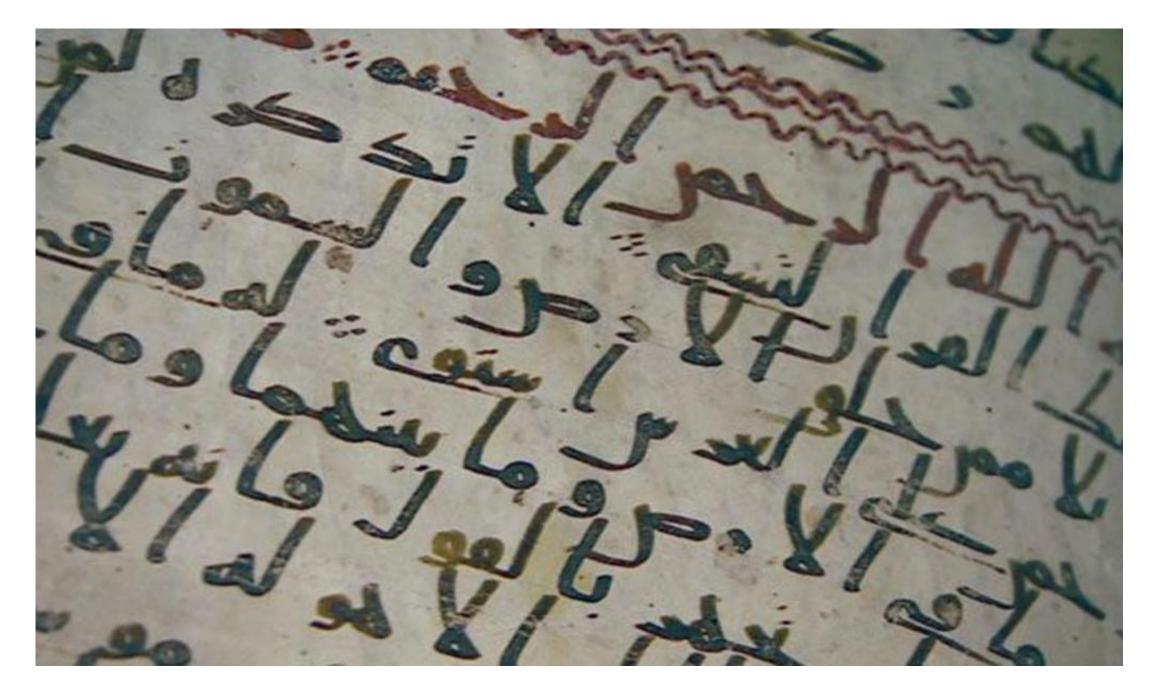




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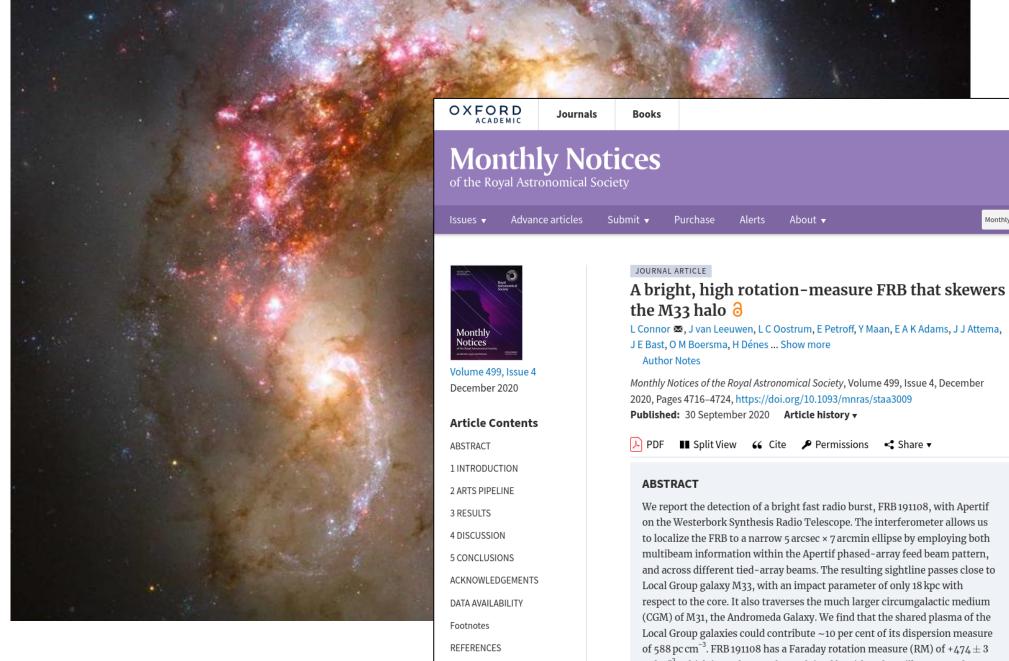




~270 projects (on many different topics)

Astronomy: fast radio bursts

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Author notes

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respect to the core. It also traverses the much larger circumgalactic medium (CGM) of M31, the Andromeda Galaxy. We find that the shared plasma of the Local Group galaxies could contribute ~10 per cent of its dispersion measure of 588 pc cm⁻³. FRB 191108 has a Faraday rotation measure (RM) of +474 \pm 3 rad m⁻², which is too large to be explained by either the Milky Way or the intergalactic medium. Based on the more moderate RMs of other extragalactic sources that traverse the halo of M33, we conclude that the dense magnetized plasma resides in the host galaxy. The FRB exhibits frequency structure on two scales, one that is consistent with quenched Galactic scintillation and broader spectral structure with $\Delta\nu \approx 40$ MHz. If the latter is due to scattering in the shared M33/M31 CGM, our results constrain the Local Group plasma environment. We found no accompanying persistent radio sources in the Apertif imaging survey data.

Keywords: fast radio bursts Issue Section: Article

1 INTRODUCTION

Fast radio bursts (FRBs) are extragalactic radio pulses, of which approximately 110 have been discovered to date (Lorimer et al. 2007; Petroff et al. 2016). They are short duration (μ s-ms), bright (0.01–100 Jy peak flux density), highly dispersed, and relatively common (~10³ sky⁻¹ d⁻¹ above 1 Jy, Cordes & Chatterjee 2019; Petroff Hessels & Lorimer 2010). The most pressing questions in FRB science fall



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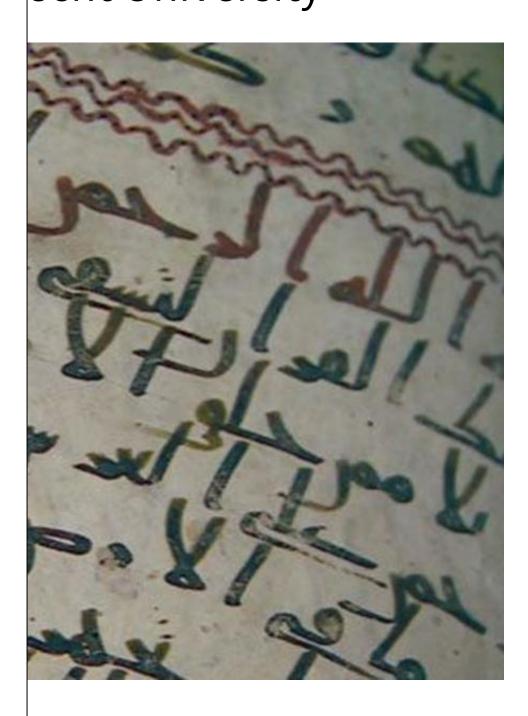
The statistical properties of stars at redshift, z = 5, compared with the present epoch

The complex organic molecular content in the L1517B starless core

The Nature of 500 Micron Risers II: Multiplicities and Environments of Sub-mm Faint Dusty Star-Forming Galaxies

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Our role in the Research Software landscape

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Broad impact, re-use, sustainability and proper credit and recognition of research software is very important to us!

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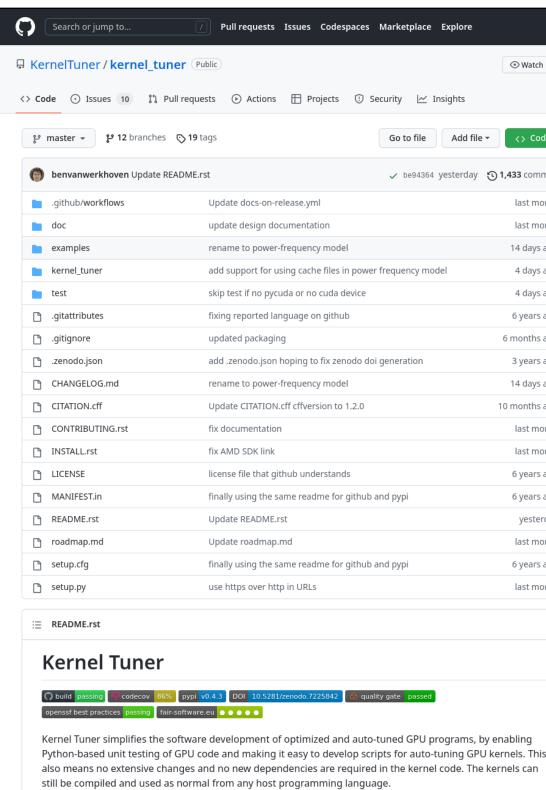
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Research software is an essential output of research projects



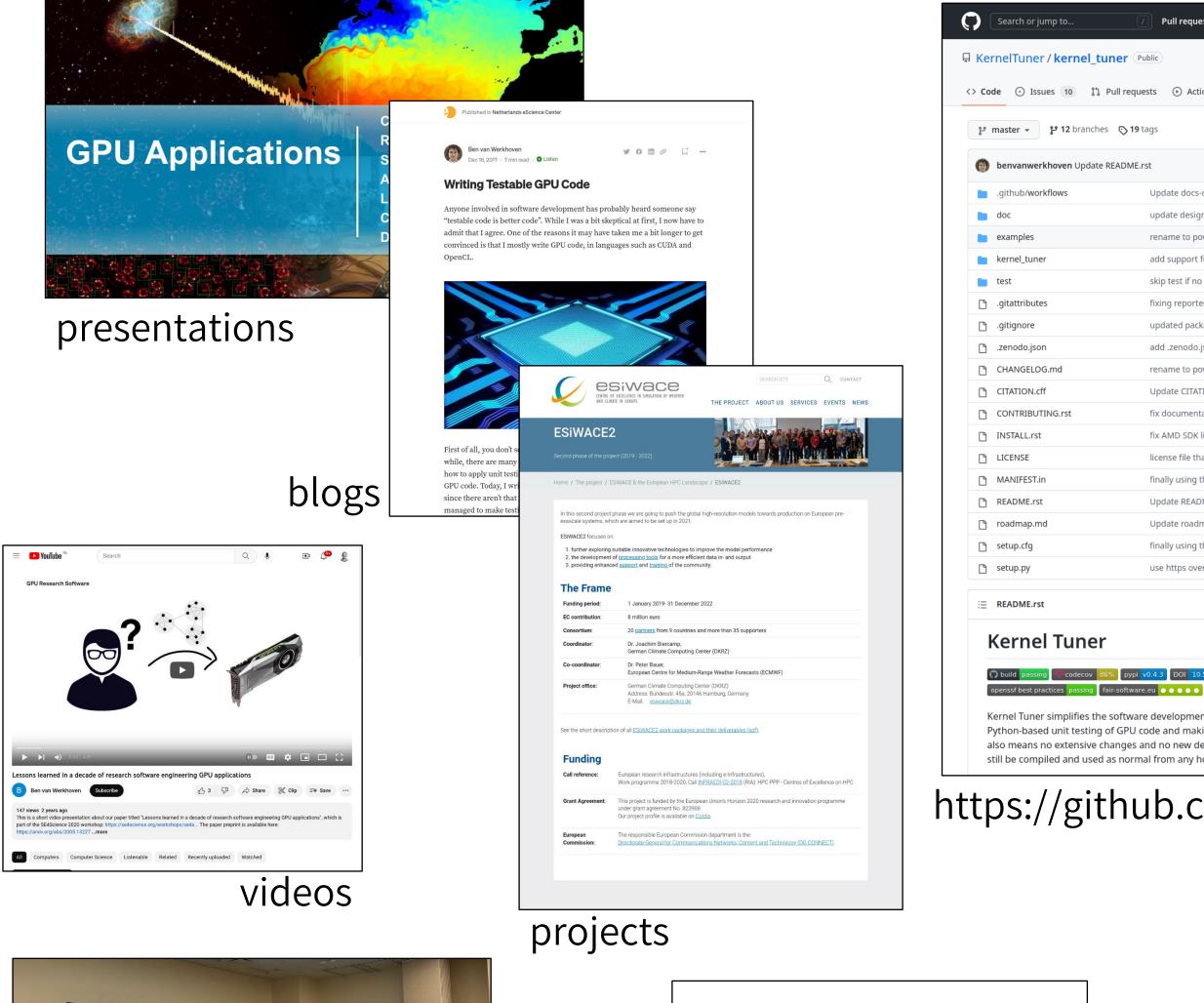


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re development of optimized and auto-tuned GPU programs, b code and making it easy to develop scripts for auto-tuning GPU and no new dependencies are required in the kernel code. The	l kernels. This	Environments 2

https://github.com/KernelTuner/kernel_tuner

🔗 dev_environment 🛛 Active

Research software has context





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code auto-tuner

Future Generation Computer Systems Volume 90, January 2019, Pages 347-358

Kernel Tuner: A search-optimizing GPU



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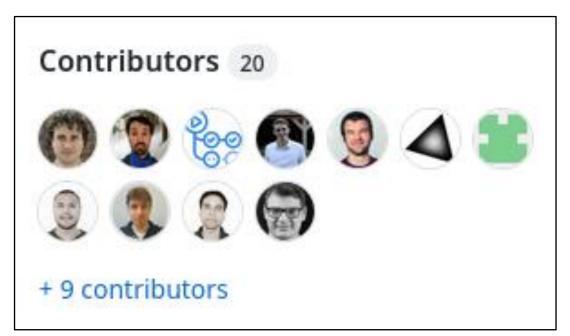
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Kernel Tuner simplifies the software development of optimized and auto-tuned GPU programs, by enabling Python-based unit testing of GPU code and making it easy to develop scripts for auto-tuning GPU kernels. This also means no extensive changes and no new dependencies are required in the kernel code. The kernels can still be compiled and used as normal from any host programming language.

https://github.com/KernelTuner/kernel_tuner



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3D particle averaging and detection of macromolecular symmetry in localization microscopy

Hamidreza Heydarian, Maarten Joosten, Adrian Przybylski, Florian Schueder, Ralf Jungmann, Ben van Werkhoven, Jan Keller-Findeisen, Jonas Ries, Sjoerd Stallinga, Mark Bates & Bernd Rieger 🖂

Nature Communications 12, Article number: 2847 (2021) Cite this article 3441 Accesses 9 Citations 19 Altmetric Metrics

A <u>Publisher Correction</u> to this article was published on 26 May 2021

• This article has been <u>updated</u>

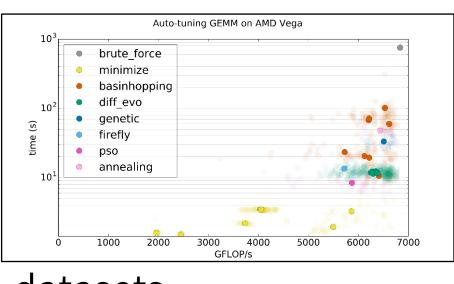
Abstract

Single molecule localization microscopy offers in principle resolution down to the molecular level, but in practice this is limited primarily by incomplete fluorescent labeling of the structure. This missing information can be completed by merging information from many structurally identical particles. In this work, we present an approach for 3D single particle analysis in localization microscopy which hugely increases signal-to-noise ratio and resolution and enables determining the symmetry groups of macromolecular complexes. Our method does not require a structural template, and handles anisotropic localization uncertainties. We demonstrate 3D reconstructions of DNA-origami tetrahedrons, Nup96 and Nup107 subcomplexes of the nuclear pore complex acquired using multiple single molecule localization microscopy techniques, with their structural symmetry deducted from the data.

Introduction

Single molecule localization microscopy (SMLM) is one of the most widely applied types of

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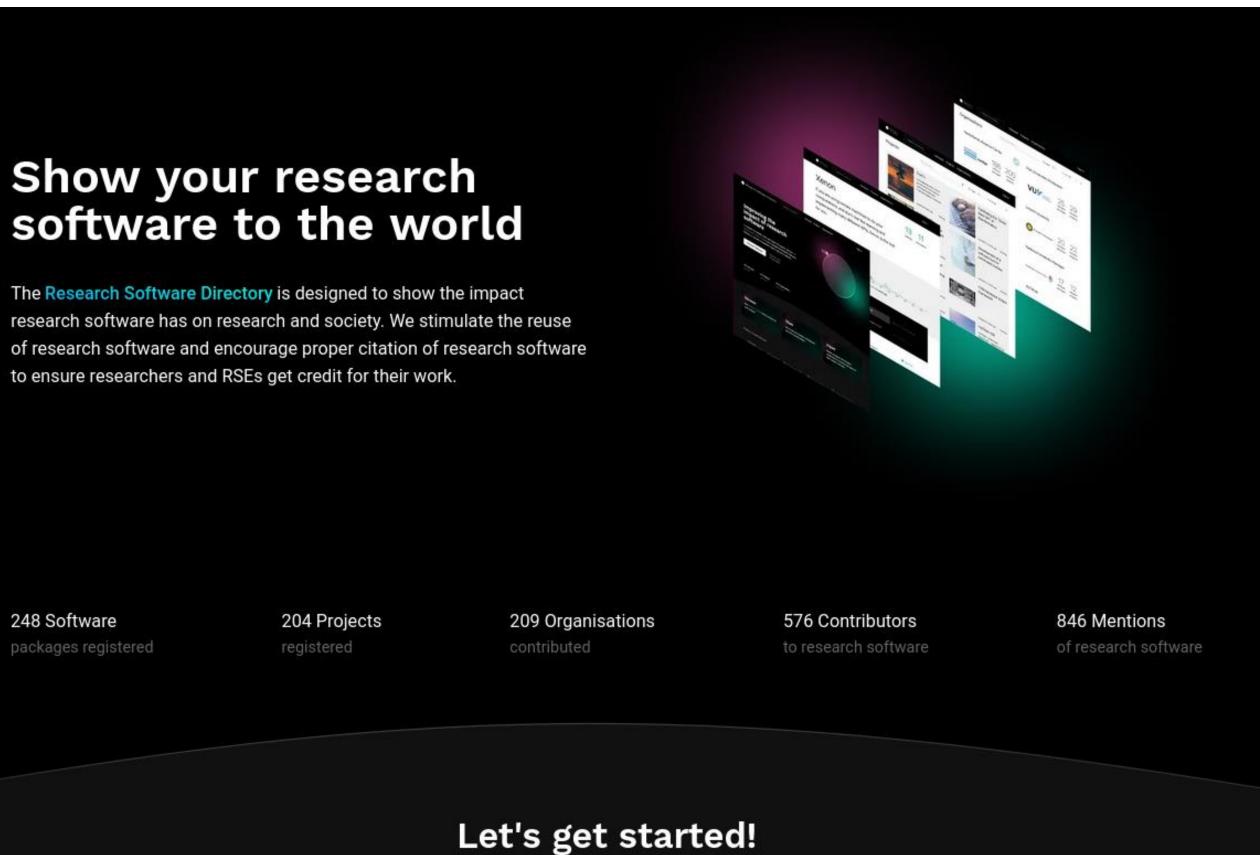
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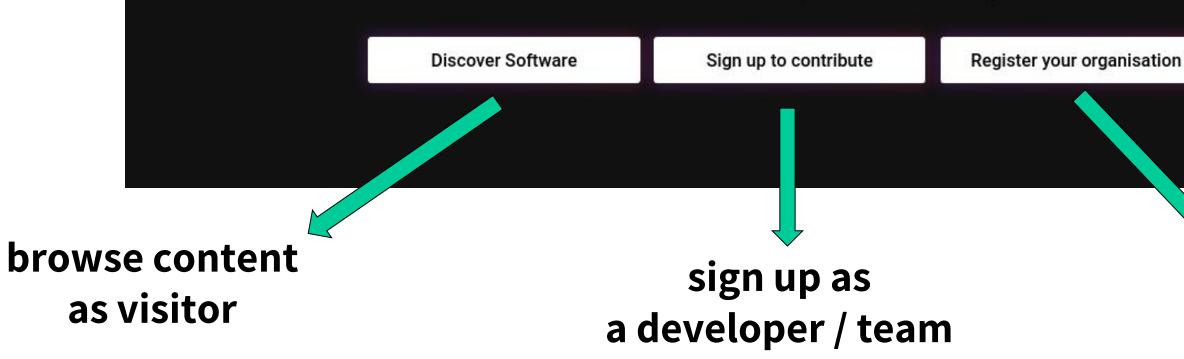
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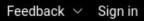


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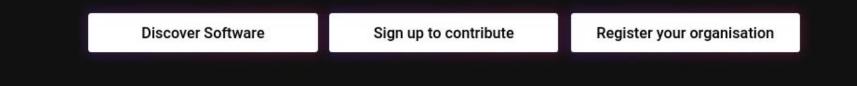
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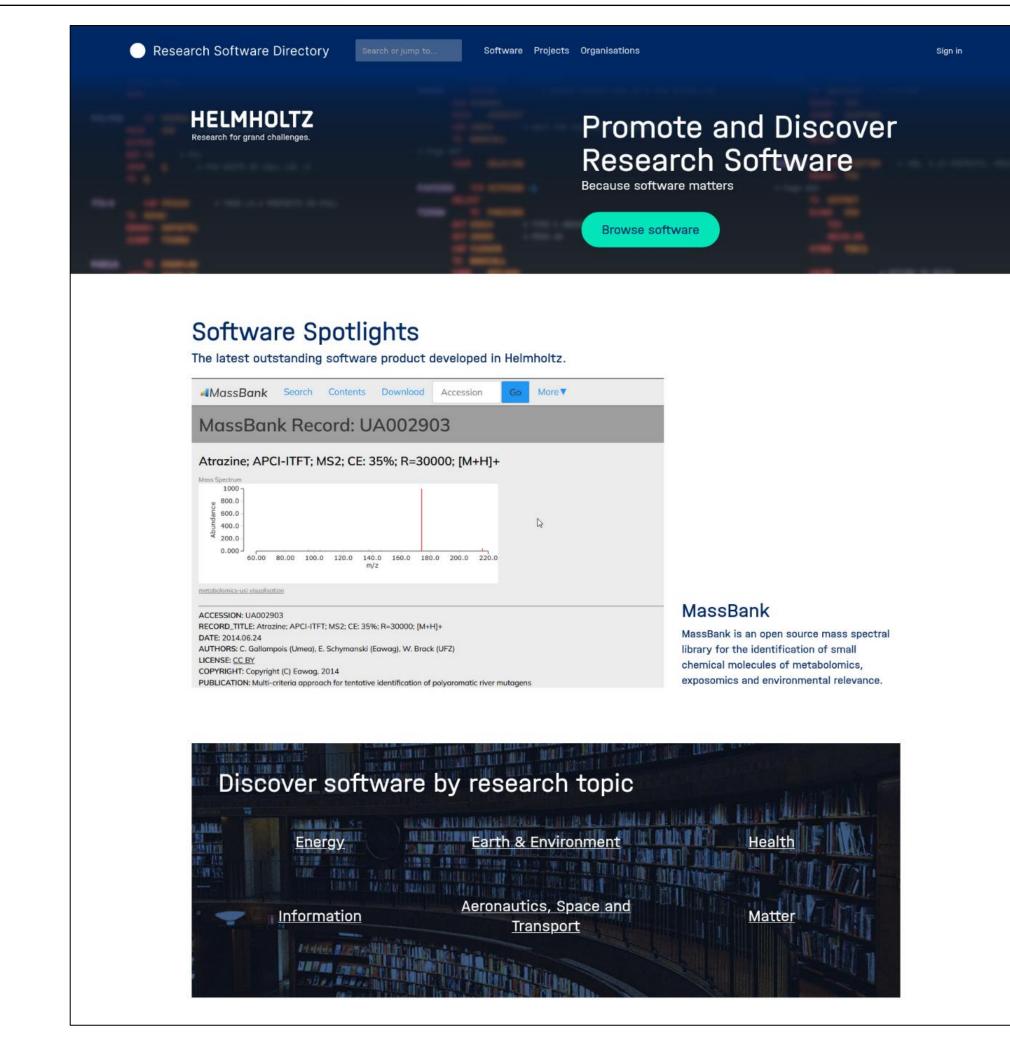
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open source and shared codebase at: https://github.com/research-software-directory/RSD-as-a-service



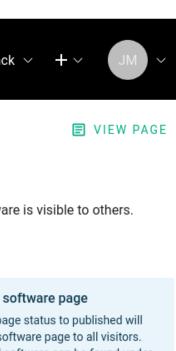
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		 * Allows developers to easily unit test and auto-tune GPU code * Generic auto-tuning of user-defined parameters for CUDA, OpenCL, and C kernels * Supports more than 20 different search optimization methods to speedup tuning * Successfully used in 10+ different eScience projects, across various disciplines 	Find or add keyword Select from top 30 list or start typ	
		 Kernel Tuner simplifies the development of efficient GPU programs, or _kernels It does by making kernels written in C/C++, OpenCL, or CUDA accessible from Python, while taking care of the required synchronization between data kept in host memory and data kept in device memory. This has a number of advantages. First, it simplifies _auto-tuning_ of the kernel 	so IMPORT KEYWORD Licenses What licenses do apply to can also import licenses	o your software? You
		parameters. In fact, Kernel Tuner comes standard with a variety of strategies for efficiently searching the parameter space, leading to greatly improved performance of tuned kernels. Second, it allows for unit testing of GPU code from within Python.	Apache-2.0	
		Kernel Tuner does not add any additional dependencies to the kernel code, and does not require extensive code changes. Furthermore, it is noteworthy that kernels tuned by Kerne Tuner do not require any changes after tuning to make them production readytuned kernel can be used as-is from any host programming language.		
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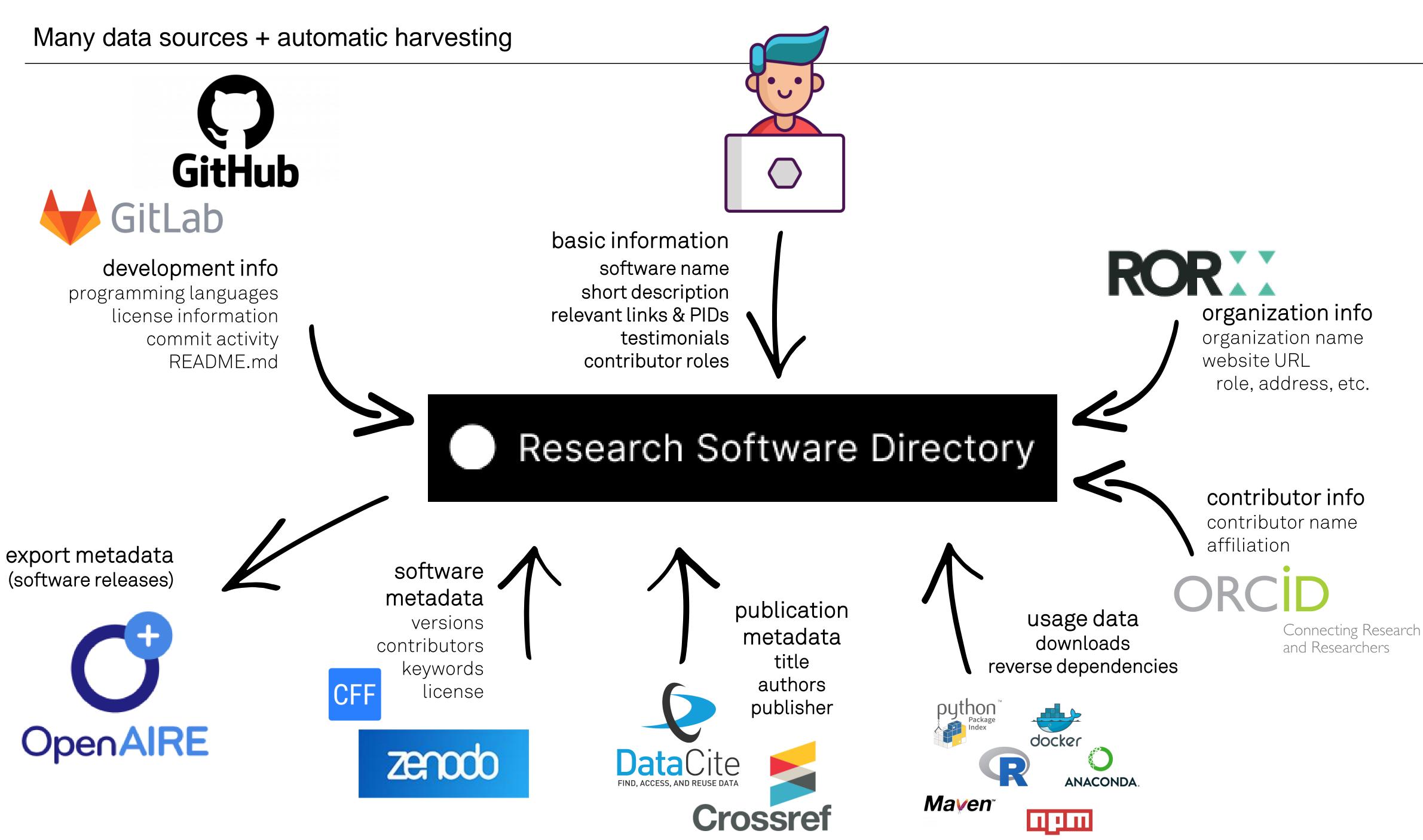
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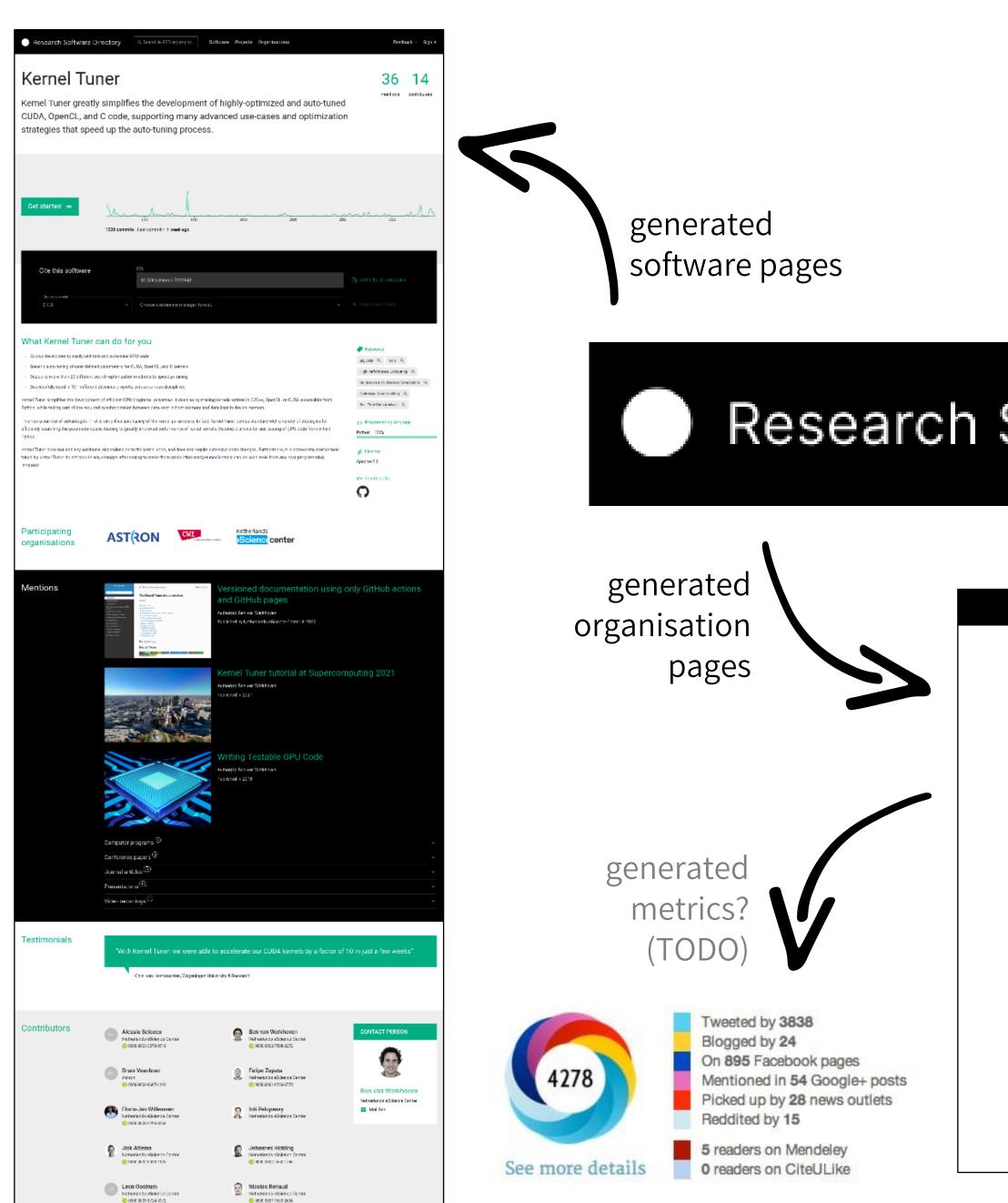
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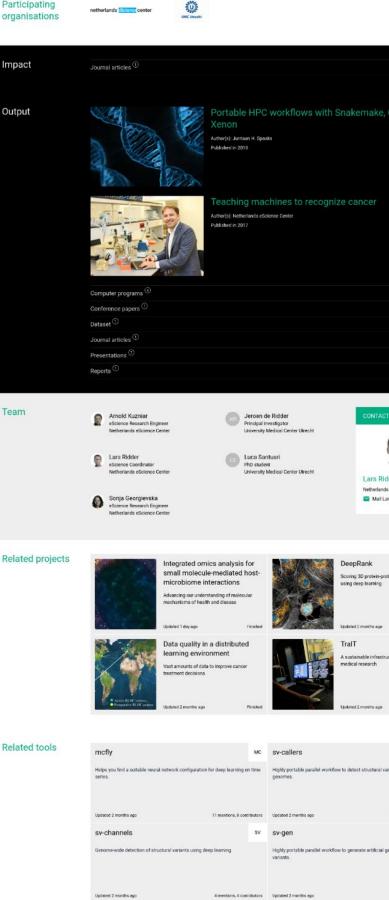
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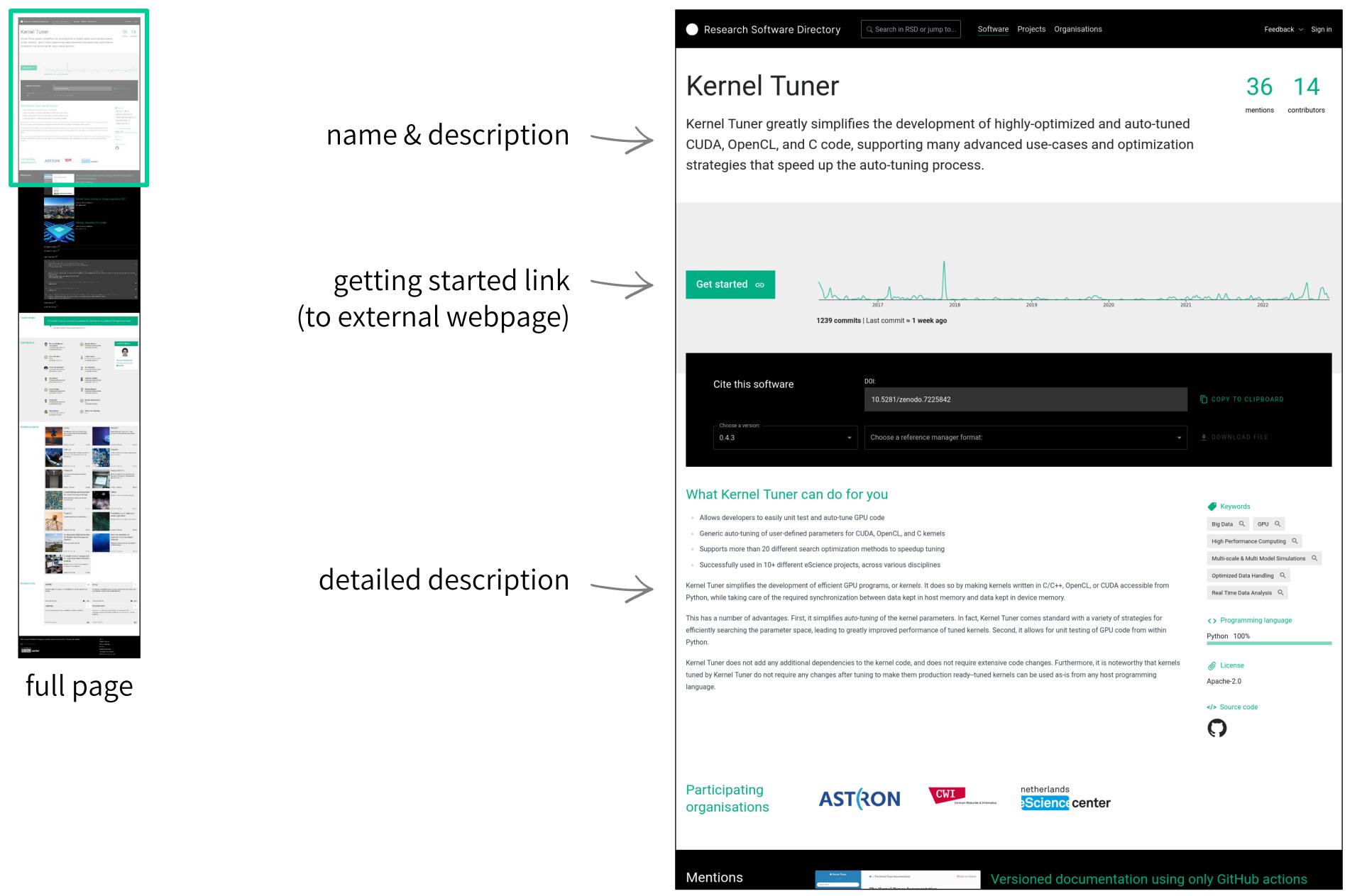


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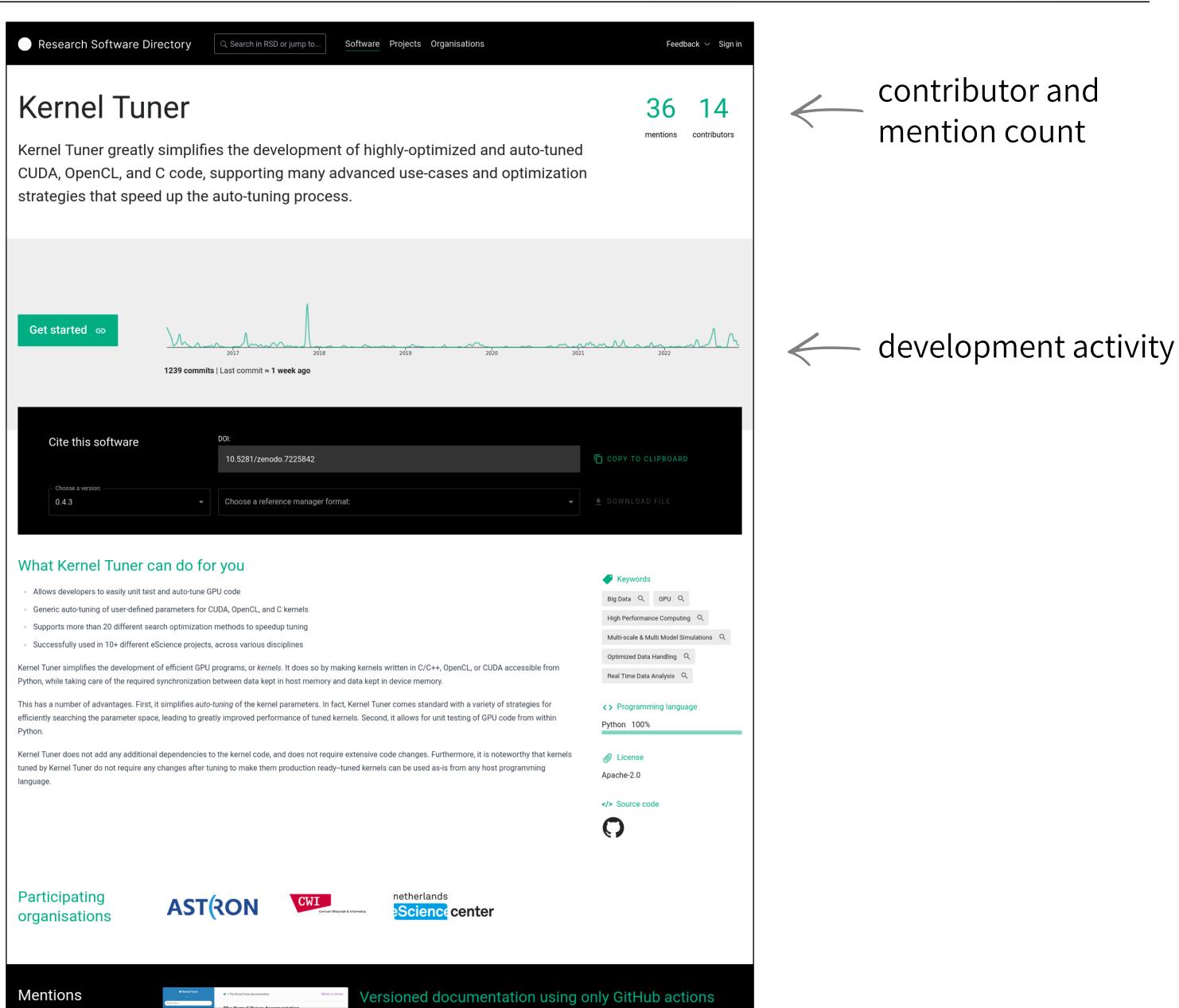
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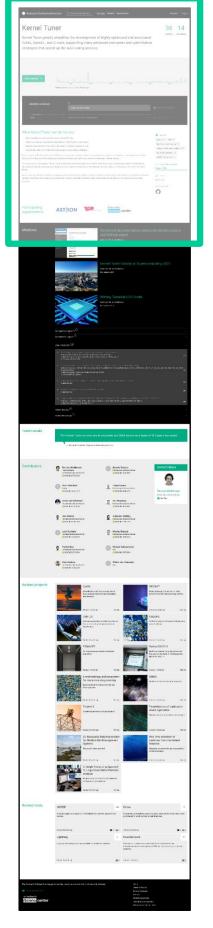
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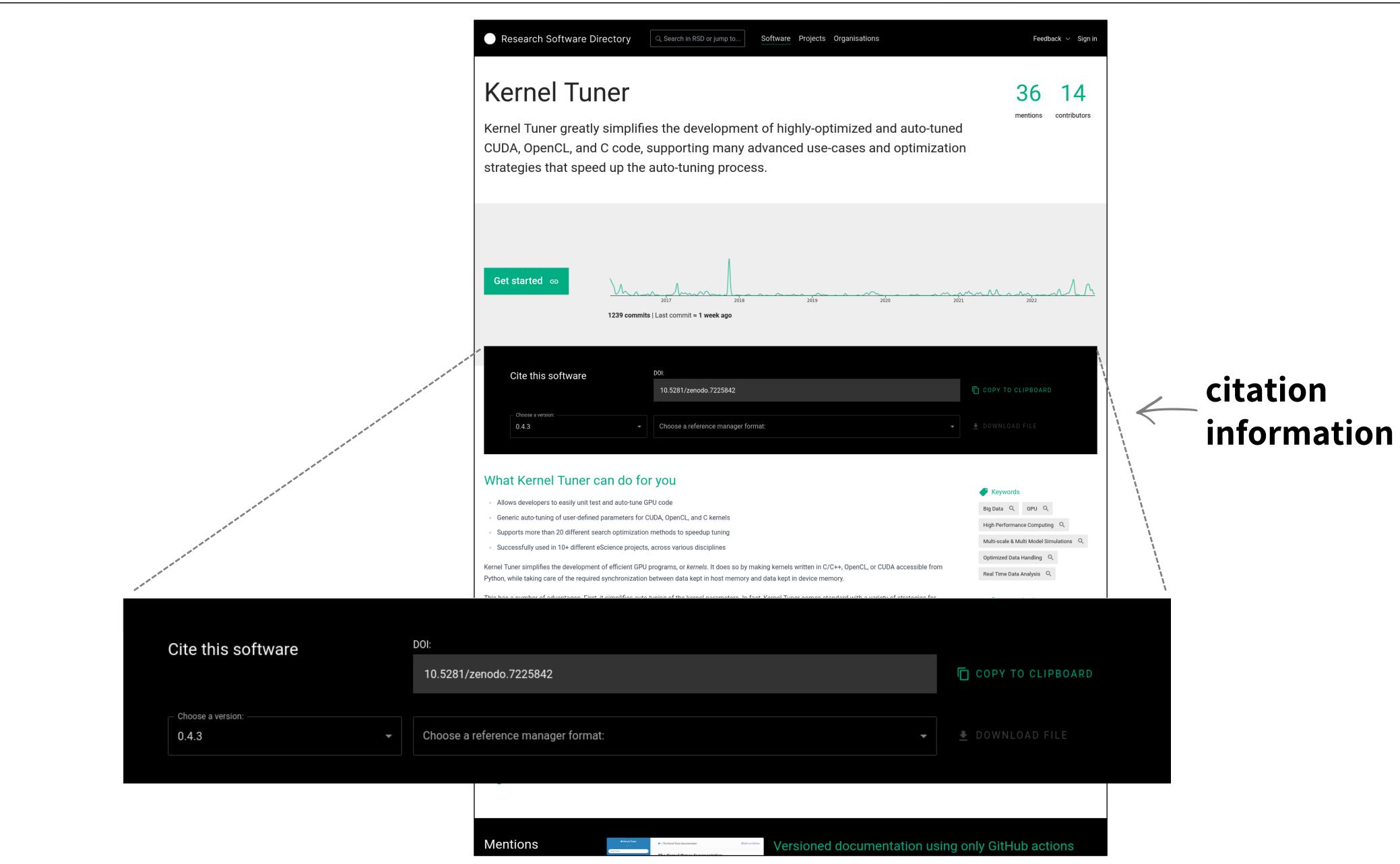
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Citation information

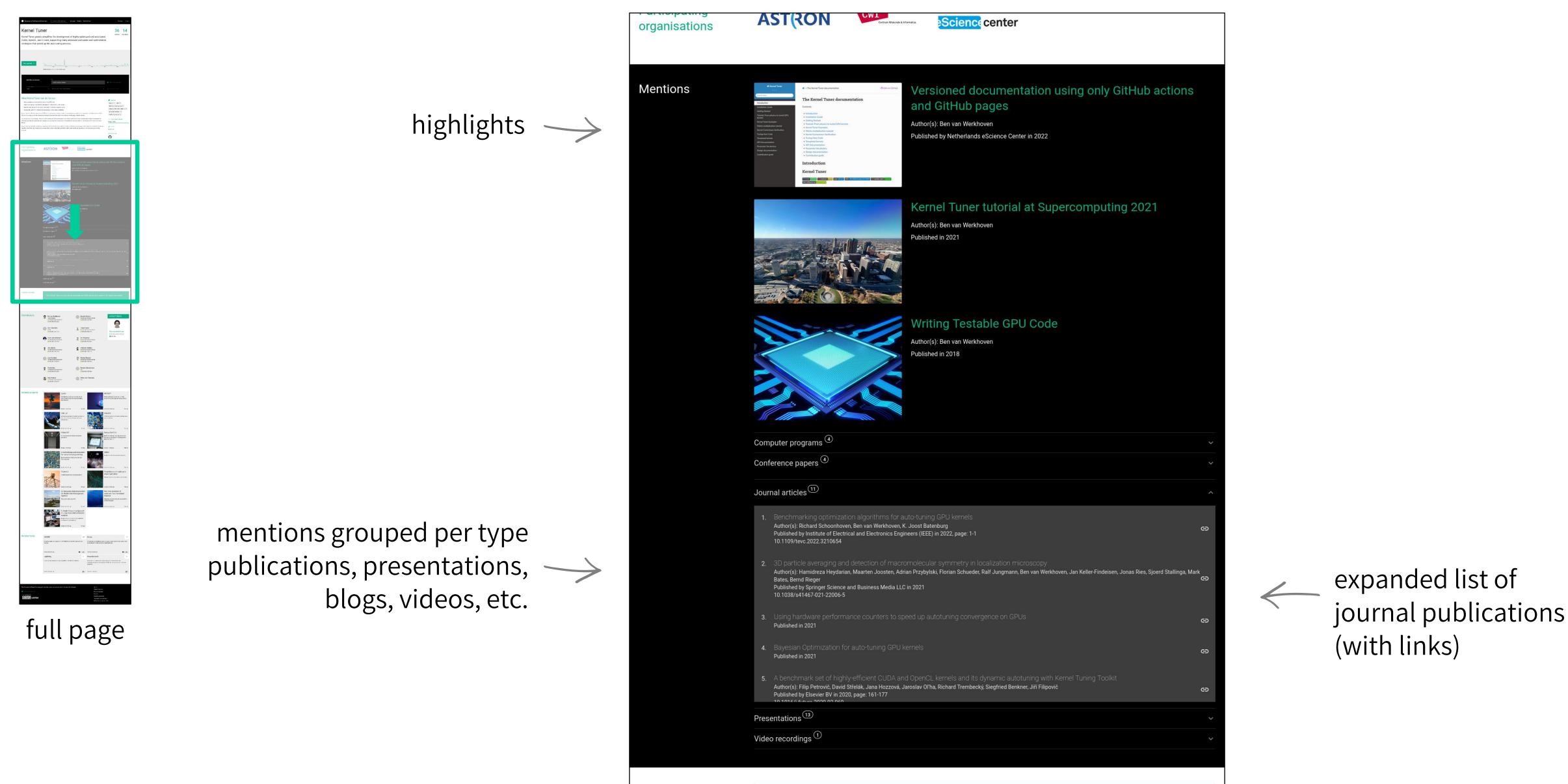


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https://research-software-directory.org/software/kernel-tuner

Mentions: related research output



Testimonials

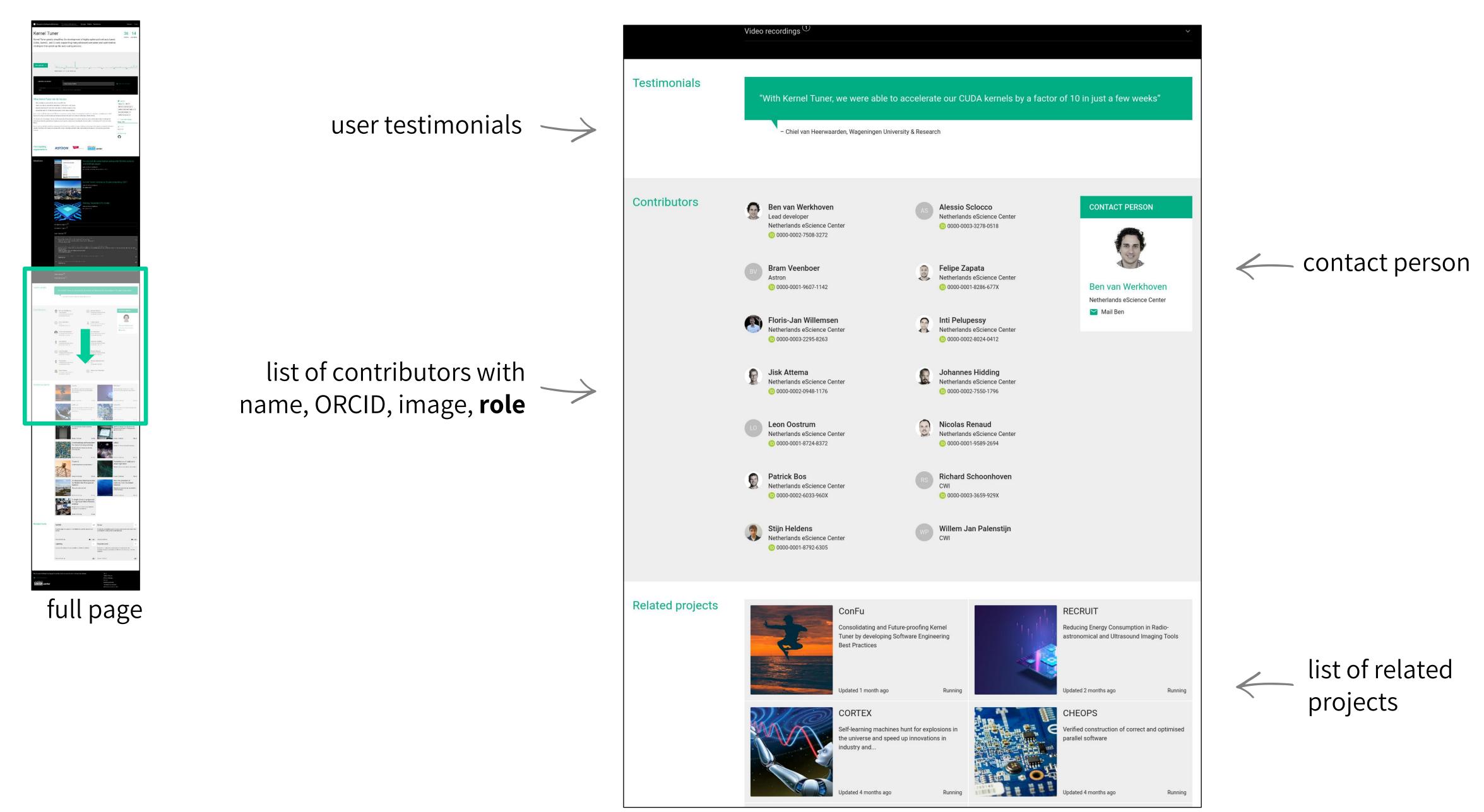


"With Kernel Tuner, we were able to accelerate our CUDA kernels by a factor of 10 in just a few weeks"

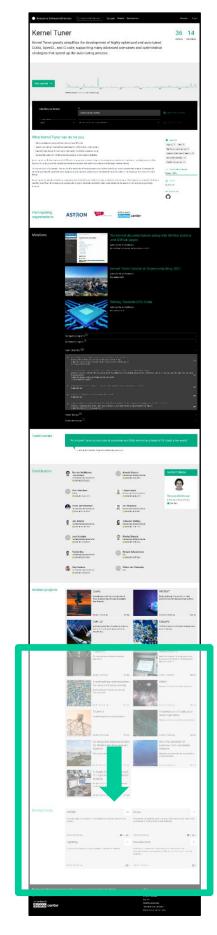
https://research-software-directory.org/software/kernel-tuner



Contributors, testimonials, related projects

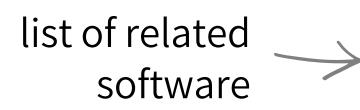


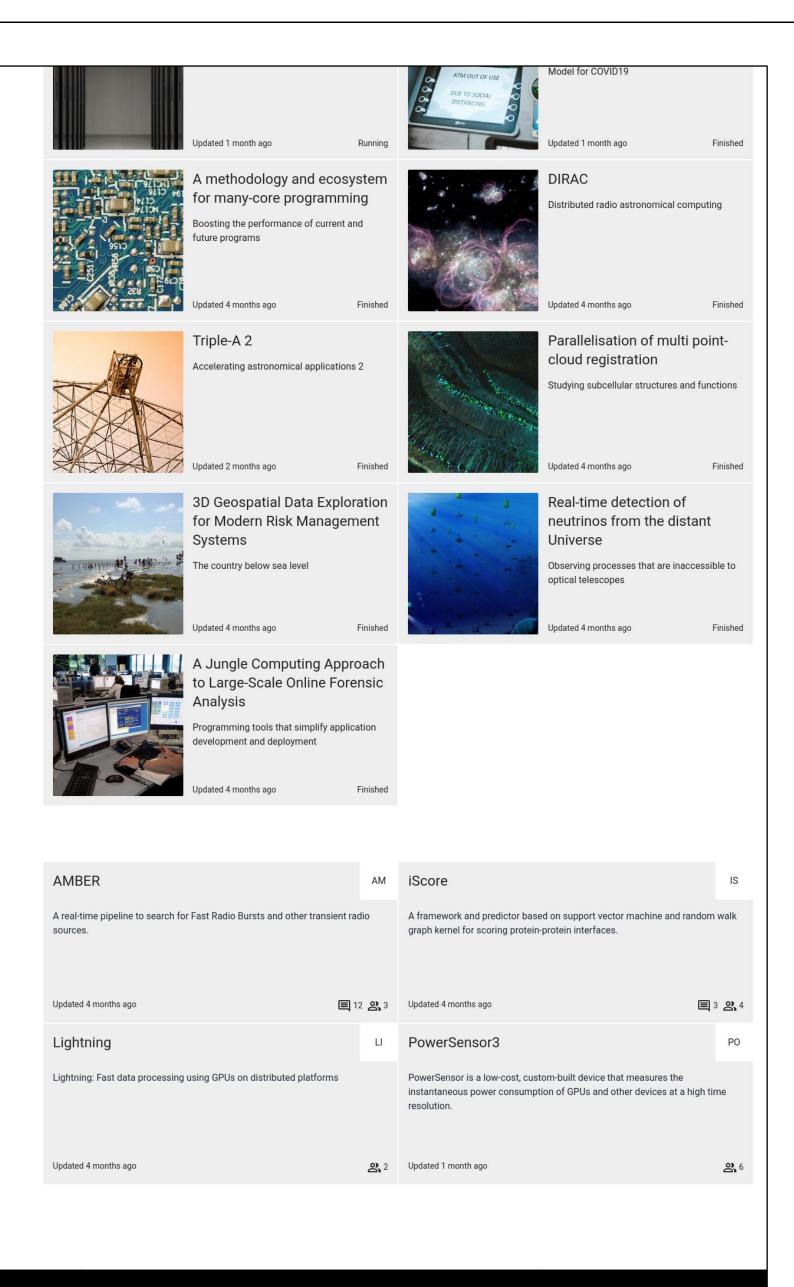
Related projects and software



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Related tools





list of related \leftarrow projects (continued)

Plans for 2023



Research Software Directory is actively being maintained and developed. Our plans for this year are:

- link to package managers, container registries, software heritage, etc., and use as data sources.
- add additional APIs to share our data with external services \checkmark (REST, OAI-PMH, etc.) + connect to OpenAIRE
- improve look and feel of software pages, software highlights, etc. U
- improve search, organizational overview, insight, etc. U
- harvest various indicators on software impact and quality (and figure out how to display them)
- research community curated software collections
- federated search between different research software directories







Questions?



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