

Grand Challenges in Medical Image Analysis

October 29, 2018

Benchmarking Algorithm Performance for Research
Amsterdam

Bram van Ginneken

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Disclosures

- Developer CAD4TB (Delft Imaging Systems): royalties & funding
- Co-founder and CSO Thirona: stock, royalties & funding
- Developer Veolity (MeVis Medical Solutions) & DynaCAD Lung (InVivo): royalties & funding
- ScreenPoint is a spin-off from my group, led by Nico Karssemeijer; my group receives royalties
- Group funding: Canon Medical, Siemens Healthineers, Philips Medical, Sectra, Novartis

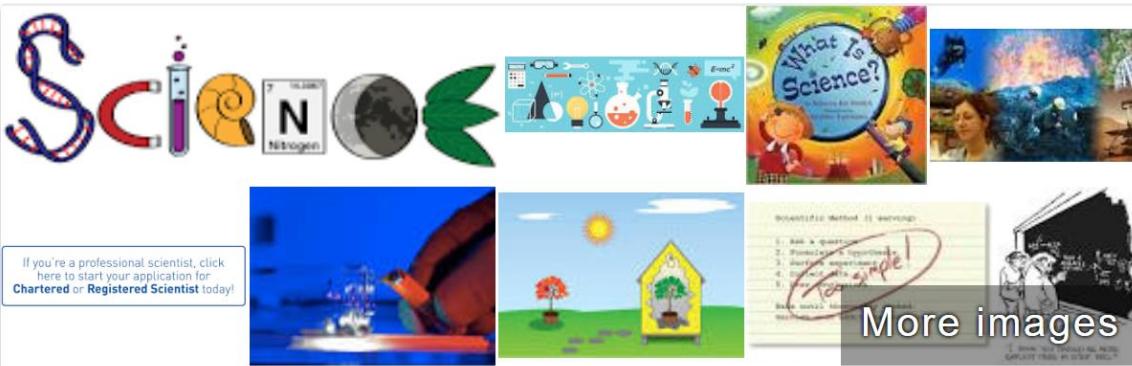
Tutorial on Benchmarking Algorithm Performance

October 29th at the 14th IEEE
International Conference on eScience
2018, Amsterdam, the Netherlands

[View the Project on GitHub](#)
NLeSC/IEEE-eScience-Tutorial-Designing-
Benchmarks

About

One of the currently most well-known benchmarks for algorithm performance is [ImageNet](#). Many challenges have been organized using this database, with the latest challenge now running on [Kaggle](#). In various scientific disciplines there is a growing interest to benchmark algorithm performance on research data. Many algorithms are proposed in the literature, but there is a growing need to compare them on the same data, using the same metrics and ground truth to compare their performance for a specific task. Organizing these open online benchmarks, will not only increase insight into which algorithms perform best for a given task, but open up these tasks for a wider audience to test their algorithms on, which could lead to new breakthroughs in the field. This tutorial shows two research fields with a longer history in benchmarking algorithm performance, such as medical image analysis (Bram van Ginneken) and multimedia information retrieval (Maria Eskevich). Mike Lees will talk about how benchmarking is being introduced for slum detection on satellite images, a field with strong restrictions on data sharing, and Kasper Marstal



Science



Science is a systematic enterprise that builds and organizes knowledge in the form of testable explanations and predictions about the universe. The earliest roots of science can be traced to Ancient Egypt and Mesopotamia in around 3500 to 3000 BCE. [Wikipedia](#)

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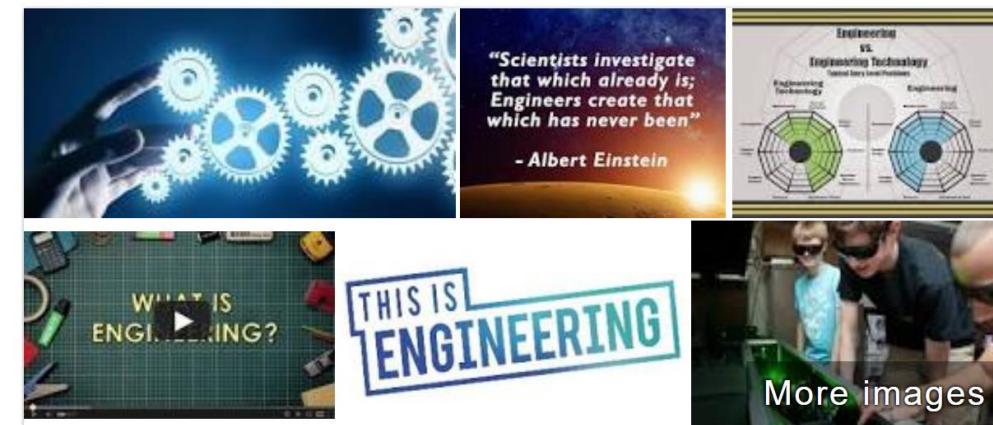


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Engineering



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Engineering is the creative application of science, mathematical methods, and empirical evidence to the innovation, design, construction, operation and maintenance of structures, machines, materials, devices, systems, processes, and organizations for the benefit of humankind. [Wikipedia](#)

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Why did I get involved in organizing challenges?

- Frustration about the typical output in my field: a paper where you present a new method and always claim it outperforms previously proposed methods. After almost a decade, I read too many of those papers that I did not believe.

Table 2. Segmentation error in comparison with a standard ASM and numbers from previous work (customized ASM [22] and deformable simplex mesh [5]), given as $\mu \pm \sigma$

Segmentation method	$D_{avg} [mm]$	$D_{RMS} [mm]$	$V_D [\%]$
Deformable model	1.6 ± 0.5	3.3 ± 1.2	5.1 ± 1.4
Active Shape Model	2.9 ± 1.1	5.2 ± 2.3	8.9 ± 2.4
Lamecker et al. [22]	2.3 ± 0.3	3.1 ± 0.5	7.0 ± 1.8
Soler et al. [5]	$2 \pm ?$	n.a.	n.a.

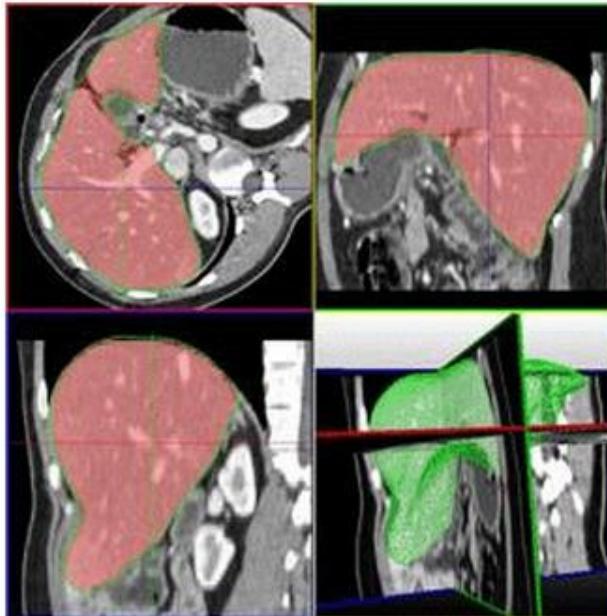


Fig. 5. Transversal, sagittal and coronal slices for the image with median average surface error. The result of the deformable model is displayed in white, the manually traced reference contour in dark gray.

SLIVER07

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Segmentation of the Liver 2007



Welcome to the website of the Segmentation of the Liver Competition 2007 (SLIVER07). The goal of this competition is to compare different algorithms to segment the liver from clinical 3D computed tomography (CT) scans.

This competition started as part of the workshop **3D Segmentation in the Clinic: A Grand Challenge**, on October 29, 2007 in conjunction with **MICCAI 2007**. Teams that participated in

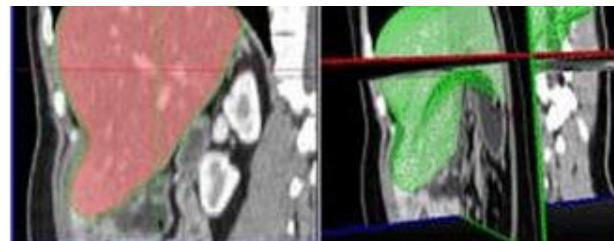
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SLIVER07

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Segmentation of the Liver 2007

The workshop is over, but through this website, the liver segmentation competition continues. You can browse the **results** of various systems, and read papers and descriptions about the methods that have been applied to the SLIVER07 data set. If you want to join the competition, you can **register** a team, **download** training and test data, and **submit** the results of your own algorithms, provided you adhere to and agree with **the rules**. More information is available in the answers to **frequently asked questions**.



Welcome to the website of the Segmentation of the Liver Competition 2007 (SLIVER07). The goal of this competition is to compare different algorithms to segment the liver from clinical 3D computed tomography (CT) scans.

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Results

For a table with more detailed results, click on the rank of the submission you are interested in.

Rank	Team	System	Submission Date	Avg Total Score
1	LME Erlangen	Semi-automatic	2010-01-14	84.6
2	Niki-Lab	Semi-automatic	2012-11-02	84.5
3	LiverPlanner	Interactive	2008-02-25	82.1
4	Afifi	Semi-automatic	2011-11-24	81.8
5	liver sirD	Semi-automatic	2011-11-30	80.5
6	liver sirA	Semi-automatic	2011-11-02	80.0
7	Niki-Lab	Semi-automatic	2011-08-23	79.7
8	Eccet	Interactive	2007-10-29	77.9
9	ZIB-Charite	Automatic	2007-10-22	77.3
10	Eccet	Interactive	2007-10-29	77.0
11	LME Erlangen	Automatic	2009-03-11	76.8
12	LiverV	Semi-automatic	2007-10-26	76.4
13	Eccet	Interactive	2007-10-29	76.3
14	LivSpl	Automatic	2012-06-22	76.2
15	Eccet	Interactive	2007-10-24	76.1
16	amcmil	Semi-automatic	2007-10-22	75.5
17	Lara	Interactive	2007-10-19	75.1
18	Eccet	Interactive	2007-10-24	74.7
19	LiverPlanner	Interactive	2008-02-25	74.2
20	LME Erlangen	Automatic	2008-12-22	74.0
21	Eccet	Interactive	2007-10-24	73.8
22	VAVlab	Semi-automatic	2012-04-04	72.9
23	VMGPU	Interactive	2008-02-20	72.9
24	VMGPU	Interactive	2008-02-20	72.9
25	KMSeg	Automatic	2012-07-28	72.4
26	Afifi	Semi-automatic	2011-09-05	72.3
27	ABE_UPM	Semi-automatic	2011-12-22	70.5
28	vislib	Semi-automatic	2009-07-27	70.0
29	LRV	Automatic	2010-03-05	69.9
30	KMSeg	Automatic	2012-06-06	69.5
31	LME Erlangen	Semi-automatic	2008-01-24	69.4
32	Dieter Seghers	Automatic	2008-09-01	68.9
33	RITS_UT	Semi-automatic	2009-11-19	68.5
34	Team FFAL	Automatic	2010-03-01	68.4
35	HUJI-liver	Semi-automatic	2008-03-03	67.9

A Shape-Guided Deformable Model

Table 2. Segmentation error in comparison with a standard ASM and number of previous work (customized ASM [22] and deformable simplex mesh [5]), given

Segmentation method	$D_{avg}[\text{mm}]$	$D_{RMSE}[\text{mm}]$	$V_D[\%]$
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Soler et al. [5]	$2 \pm ?$	n.a.	n.a.

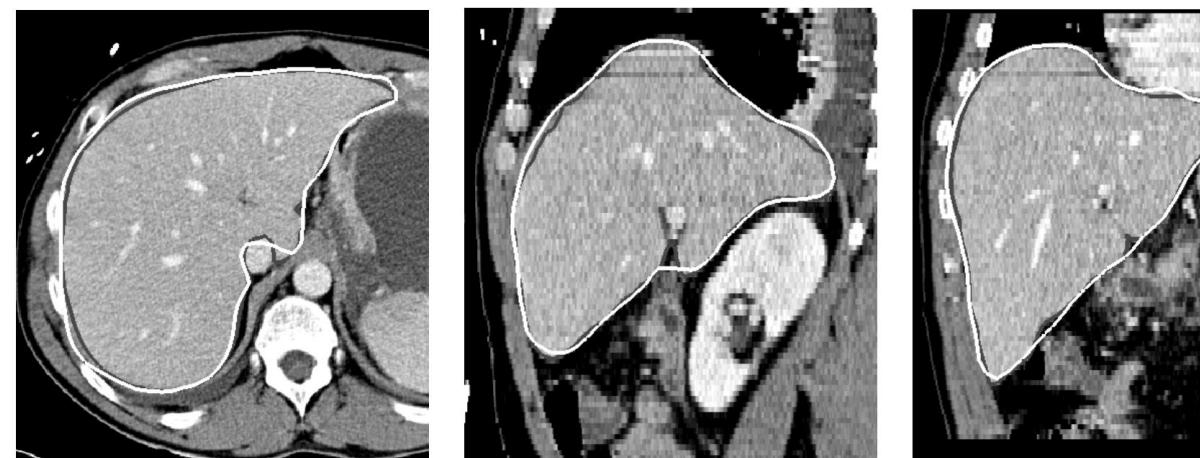
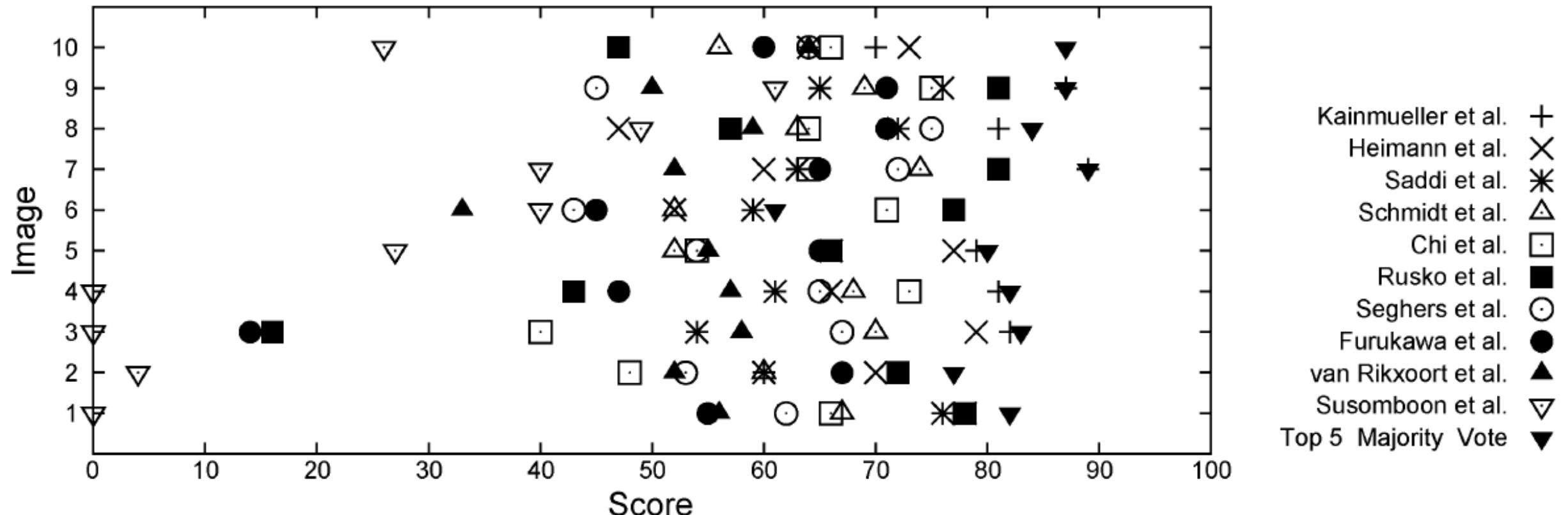


Fig. 5. Transversal, sagittal and coronal slices for the image with median surface error. The result of the deformable model is displayed in white, the traced reference contour in dark gray.

Comparison and Evaluation of Methods for Liver Segmentation From CT Datasets

Tobias Heimann*, Bram van Ginneken, *Member, IEEE*, Martin A. Styner, *Member, IEEE*, Yulia Arzhaeva, Volker Aurich, Christian Bauer, Andreas Beck, Christoph Becker, Reinhard Beichel, *Member, IEEE*, György Bekes, Fernando Bello, *Member, IEEE*, Gerd Binnig, Horst Bischof, *Member, IEEE*, Alexander Bornik, Peter M. M. Cashman, Ying Chi, Andrés Córdova, Benoit M. Dawant, Márta Fidrich, Jacob D. Furst, Daisuke Furukawa, Lars Grenacher, Joachim Hornegger, *Member, IEEE*, Dagmar Kainmüller, Richard I. Kitney, Hidefumi Kobatake, Hans Lamecker, Thomas Lange, Jeongjin Lee, Brian Lennon, Rui Li, Senhu Li, Hans-Peter Meinzer, *Member, IEEE*, Gábor Németh, Daniela S. Raicu, Anne-Mareike Rau, Eva M. van Rikxoort, Mikaël Rousson, László Ruskó, Kinda A. Saddi, Günter Schmidt, Dieter Seghers, Akinobu Shimizu, *Member, IEEE*, Pieter Slagmolen, Erich Sorantin, Grzegorz Soza, Ruchaneewan Susomboon, Jonathan M. Waite, Andreas Wimmer, and Ivo Wolf



Kainmueller et al. +
Heimann et al. X
Saddi et al. *
Schmidt et al. Δ
Chi et al. □
Rusko et al. ■
Seghers et al. ○
Furukawa et al. ●
van Rikxoort et al. ▲
Susomboon et al. ▽
Top 5 Majority Vote ▼



Bram van Ginneken

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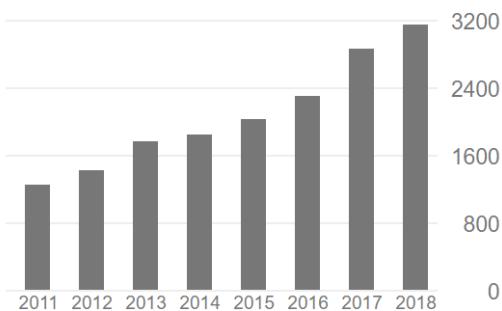
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Medical Image Analysis Medical Imaging Deep Learning Computer-Aided Diagnosis
Machine Learning

<input type="checkbox"/>	TITLE		⋮	CITED BY	YEAR
<input type="checkbox"/>	Ridge-based vessel segmentation in color images of the retina			2060	2004
	J Staal, MD Abràmoff, M Niemeijer, MA Viergever, B Van Ginneken IEEE transactions on medical imaging 23 (4), 501-509				
<input type="checkbox"/>	Reflectance and texture of real-world surfaces			1597	1999
	KJ Dana, B Van Ginneken, SK Nayar, JJ Koenderink ACM Transactions On Graphics (TOG) 18 (1), 1-34				
<input type="checkbox"/>	Comparative study of retinal vessel segmentation methods on a new publicly available database			684	2004
	M Niemeijer, J Staal, B van Ginneken, M Loog, MD Abramoff Medical Imaging 2004: Image Processing 5370, 648-657				
<input type="checkbox"/>	Comparison and evaluation of methods for liver segmentation from CT datasets			660	2009
	T Heimann, B Van Ginneken, MA Styner, Y Arzhava, V Aurich, C Bauer, ... IEEE transactions on medical imaging 28 (8), 1251-1265				
<input type="checkbox"/>	A survey on deep learning in medical image analysis			628	2017
	G Litjens, T Kooi, BE Bejnordi, AAA Setio, F Ciompi, M Ghafoorian, ... Medical image analysis 42, 60-88				
<input type="checkbox"/>	Active shape model segmentation with optimal features			577	2002
	B Van Ginneken, AF Frangi, JJ Staal, BM ter Haar Romeny, MA Viergever Medical Imaging, IEEE Transactions on 21 (8), 924-933				
<input type="checkbox"/>	Computer analysis of computed tomography scans of the lung: a survey			535	2006
	I Sluimer, A Schilham, M Prokop, B Van Ginneken IEEE transactions on medical imaging 25 (4), 385-405				
<input type="checkbox"/>	Computer-aided diagnosis in chest radiography: a survey			522	2001
	B Van Ginneken, BMTH Romeny, MA Viergever IEEE Transactions on medical imaging 20 (12), 1228-1241				
<input type="checkbox"/>	Automatic detection of red lesions in digital color fundus photographs			476	2005
	M Niemeijer, B Van Ginneken, J Staal, MSA Suttorp-Schulten, ... IEEE Transactions on medical imaging 24 (5), 584-592				

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Citations	21324	14042
h-index	70	57
i10-index	228	192

[Co-authors](#)[EDIT](#)

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What happened next?

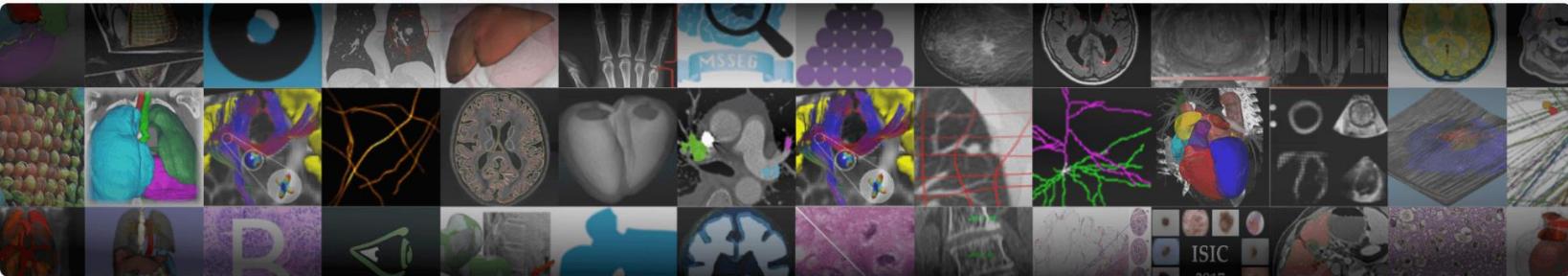
- I urged others to organize challenges. It was fun and useful.
- They asked for our website scripts. They copied bits and pieces.
- There were more and more challenges organized
- I figured we needed an easy way to set up a challenge, and reuse components
- So I hired an enthusiastic programmer to build a general platform
- And we started to keep an overview of challenges on a website: grand-challenge.org

grand-challenges - Home x +

← → C 🔒 https://grand-challenge.org

BramVanGinneken

Grand-Challenges ALL CHALLENGES



WHY CHALLENGES?

HOST A CHALLENGE

CONTRIBUTORS

ADMIN ▾

Grand Challenges in Biomedical Image Analysis

Every year, thousands of papers are published that describe new algorithms to be applied to medical and biomedical images, and various new products appear on the market based on such algorithms. But few papers and products provide a fair and direct comparison of the newly proposed solution with the state-of-the-art. We believe that such comparisons can help the research community and industry to develop better algorithms. We support the organization of these comparative studies and the dissemination of their results.

Organizing and participating in challenges is not the only way to facilitate better comparisons between new and existing solutions. If it were easy to publish and share your data, and the code you used to evaluate your algorithm's performance on that data, and possibly the algorithm itself, others could directly compare their approach to yours, using the same test data and the same evaluation metrics. With this site we provide tools to make it as easy as possible for you to publish your data and your evaluation for any paper you've written.

[Why Challenges?](#) describes the rationale for organizing grand challenges, provides advice for those who want to organize such events, and discusses where we hope the field will move to next.

[All Challenges](#) provides an overview of all previous, ongoing and upcoming challenges in biomedical image analysis that we are aware of. Drop us a note if you want your event listed on this overview.

[Create your own challenge](#) explains how you can set up your own challenge site in a matter of minutes.

Grand challenges in medical image analysis would not be possible without the support and enthusiasm from all [contributors](#).

Filters

[Modality](#)[Task type](#)[Structure](#)

Displaying 173 of 173

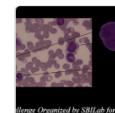
All Challenges

Here is an overview of all challenges that have been organized within the area of medical image analysis that we are aware of. If you know any study that would fit in this overview, or want to advertise your challenge, please send an email to support@grand-challenge.org and we will add the challenge to the list on this page.

[+ Add an external challenge](#)[Edit external challenges](#)

Active filters: 0

2019



C-NMC 2019

An effort to build an automated classifier for classification of normal vs malignant cells in B-ALL white blood cancer microscopic images that will overcome the problems associated with deploying sophisticated high-end machines with recurring reagent ...

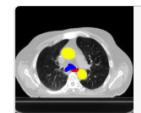
Participants: 4
Workshop: April 8, 2019
Associated with: ISBI 2019
Hosted on: grand-challenge.org



EAD2019

Endoscopic Artefact Detection (EAD) is a core problem and needed for realising robust computer-assisted tools. The EAD challenge has 3 tasks: 1) Multi-class artefact detection, 2) Region segmentation, 3) Detection generalisation.

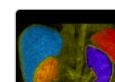
Participants: 23
Workshop: April 8, 2019
Associated with: ISBI2019
Hosted on: grand-challenge.org



SegTHOR

Segmentation of THoracic Organs at Risk in CT images

Workshop: April 8, 2019
Associated with: ISBI 2019
Hosted on: grand-challenge.org



CHAOS

BreastPathQ	HC18	PROSTATEX	Combined Radiology and Pathology Classification	Medical Segmentation Decathlon	3-D Validation of Tractography Using Experimental MRI	Multiple Sclerosis Segmentation (MSSEG)	Circuit Reconstruction from Electron Microscopy Images (CREMI)	xVertSeg	ANATOMY3	Longitudinal Multiple Sclerosis Lesion Segmentation Challenge	NEATBrainS15	Cephalometric X-Ray Landmark Detection	Liver CT Annotation Task	CETUS	MICCAI 2012 DTI Tractography Challenge	Brain Tumor Image Segmentation Challenge 2012	MS Lesion Segmentation Challenge 2008		
<small>MICCAI 2012 Dataset: Head Cancer</small>	<small>Automated measurement of head circumference using 2D ultrasound images</small>	<small>Classification of prostate feature using morphometric ROI data</small>	<small>Classification of breast malignancy based on a set of visual features</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>			
PAVES	Peripheral Artery/Vein Multi-Organ Segmentation (PAVES)	Multi-Organ Nuclei Segmentation Challenge	Medical Segmentation Decathlon	ROCC	LCTSC	TADPOLE	Computational Precision Medicine	Single Molecule Localization Microscopy Challenge	Gland Segmentation for Intravascular Imaging and Compt	Cephalometric X-ray image analysis Challenge	Lung nodule classification challenge	Brain Tumor Image Segmentation Challenge 2014	Bone Texture Characterization	Challenge US Biometric Measurements from fetal ultrasound	NeoBrainS12	RV Segmentation Challenge in Cardiac MRI	CAUSE07		
<small>Peripheal Artery/Vein Multi-Organ Segmentation (PAVES) is the challenge focused on providing the most comprehensive segmentation of the peripheral arteries and veins.</small>	<small>The challenge tests the generalizability of multi-organ segmentation methods to 10 different anatomical regions.</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Associated with MICCAI 2016</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
REFUGE	Correction of Brainshift with Intra-Operative Ultrasound	The Continuous Registration Challenge	CATARACTS	RETOUCH	CARELYON17	Ultrasound Nerve Segmentation	BigNeuron	The Second Overlapping Cervical Cytology Segmentation Chal	ImageCLEFmed: Medical classification	3D Deconvolution Microscopy	LOLA11	Cardiac Motion Estimation Challenge 2011	DTI Segmentation Challenge in IVUS	SLIVER07	2007	2007	2007		
<small>The goal of the REFUGE dataset is to evaluate the performance of automated registration and computer-aided algorithms for guidance and surgical registration using a common dataset of patient brain images.</small>	<small>Submit your method for lung and brain registration!</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
LUMIC	IDRID	ICBIS2018-Challenge	Skin Lesion Analysis Towards Melanoma Detection	Cervical Cancer Screening	ENIGMA: Cerebellum	Pediatric Bone Age Challenge	Tumor Proliferation Assessment Challenge (TUPAC)	Statistical Atlases and Computational Modelling of the Heart - S	Low-Dose CT Grand Challenge	Cell Tracking Challenge	Shape 2015 - Statistical Shape Model Challenge	GRASS	Localization Microscopy Challenge	LOLA11	Cardiac Motion Estimation Challenge 2011	DTI Segmentation Challenge in IVUS	SLIVER07		
<small>The LUMIC challenge tests the accuracy of automated segmentation methods in CT lung images for segmenting lung CT with radiographic digitalization.</small>	<small>This challenge evaluates automated techniques for analysis of fundus photographs. The target categories include fundus photograph, macula, maculopathy, and hemorrhage and includes a test set of 10 images. Each image will be graded by fundus experts according to the severity level of fundus lesions.</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
EndoVis	MURA: Bone X-Ray Deep Learning Competition	RSNA Pneumonia Challenge	Multi-shell diffusion MRI harmonization and enhancement	Ischemic Stroke Lesion Segmentation 2017	ProstateC-2 Challenge	Multi-Modality Whole Heart Segmentation	HVSMP 2016	MITOS-ATYPIA-14	Digital Mammography DREAM Challenge	Cell Tracking Challenge (2nd Edition)	OVERLAPPING CERVICAL CYTOLOGY SEGMENTATION CHALLENGE	AMIDAT3	DIADEM	EMPIRE10	SK10	2007	2007	2007	
<small>An as endoscopic vision (CV) challenge to develop a deep learning framework for extracting the content of the field of view, gather features for training and testing, and segmenting vascular structures.</small>	<small>Automatically detect a visual signal for pneumonia in chest X-rays.</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
ImageCLEF Tuberculosis 2018	Multi-shell diffusion MRI harmonization and enhancement	Dermoscopic Skin Lesion Analysis Towards Melanoma Detection	The skin of the challenge is to evaluate the performance of automated methods for the detection of melanoma lesions from 2D images.	ACDC	LITS - Liver Tumor Segmentation	Digital Mammography DREAM Challenge 2017	C11	MITOS-ATYPIA-14	Skin Lesion Analysis Towards Melanoma Detection	Cell Tracking Challenge (2nd Edition)	OVERLAPPING CERVICAL CYTOLOGY SEGMENTATION CHALLENGE	ANODE09	EXACT09	ANODE09	EXACT09	ANODE09	EXACT09		
<small>Assess the probability of 10 patient having residual lung lesions based on 2D chest X-ray images.</small>	<small>The aim of the challenge is to evaluate the performance of automated methods for the detection of melanoma lesions from 2D images.</small>	<small>The goal of the challenge is to help participants develop image analysis tools to facilitate the automated detection of melanoma lesions from dermoscopic images.</small>	<small>The skin of the challenge is to help participants develop image analysis tools to facilitate the automated detection of melanoma lesions from dermoscopic images.</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
Intervertebral Disc Segmentation Challenge	MR Brain Image Segmentation	18F-FDG PET Radiomic Risk Stratifiers in Head and Neck Cancer	Tissue Microarray Analysis for Throat Cancer	TruCED	WNI Segmentation Challenge	Brain Tumor Image Segmentation Challenge 2016	C11	MITOS-ATYPIA-14	Cell Tracking Challenge (2nd Edition)	OVERLAPPING CERVICAL CYTOLOGY SEGMENTATION CHALLENGE	ANODE09	EXACT09	ANODE09	EXACT09	ANODE09	EXACT09	ANODE09	EXACT09	
<small>The goal of MICCAI 2018 Challenge on Automatic Localization and Segmentation of Intervertebral Disc Segmentation involves two datasets (train and test). The test dataset consists of 100 images, and the test set consists of 10 images. The test set consists of 10 images, and the test set consists of 10 images. The test set consists of 10 images, and the test set consists of 10 images.</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
Brain Tumor Segmentation Challenge 2018	Pancreatic Cancer Segmentation Challenge 2018	Intervertebral Disc Segmentation Challenge	Combined Radiology and Pathology Classification	PROSTATEX	Coronary Artery Segmentation Challenge	6-month infant brain MRI segmentation	Biting Radiography Segmentation Challenge	OPTIMA Retinal Cyst Segmentation Challenge	Ischemic Stroke Segmentation Challenge 2015	CLUST14	orCaScore	VISCEL Benchmark 2	PROMISE12	VESSEL12	High angular resolution diffusion imaging 2012	ANODE09	EXACT09	ANODE09	EXACT09
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<small>Create an algorithm to automatically detect a given tissue of interest from a given set of histology and fluorescence images.</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
HC18	PROSTATEX	Combined Radiology and Pathology Classification	Medical Segmentation Decathlon	3-D Validation of Tractography Using Experimental MRI	Multiple Sclerosis Segmentation (MSSEG)	Circuit Reconstruction from Electron Microscopy Images (CREMI)	xVertSeg	ANATOMY3	Longitudinal Multiple Sclerosis Lesion Segmentation Challenge	NEATBrainS15	Cephalometric X-Ray Landmark Detection	Liver CT Annotation Task	CETUS	MICCAI 2012 DTI Tractography Challenge	Brain Tumor Image Segmentation Challenge 2012	MS Lesion Segmentation Challenge 2008	2007		
<small>Automated measurement of head circumference using 2D ultrasound images</small>	<small>Classification of prostate feature using morphometric ROI data</small>	<small>Classification of breast malignancy based on a set of visual features</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>			
PAVES	Multi-Organ Nuclei Segmentation Challenge (PAVES)	Medical Segmentation Decathlon	ROCC	LCTSC	TADPOLE	Computational Precision Medicine	Single Molecule Localization Microscopy Challenge	Gland Segmentation for Intravascular Imaging and Compt	Cephalometric X-ray image analysis Challenge	Lung nodule classification challenge	Brain Tumor Image Segmentation Challenge 2014	Bone Texture Characterization	Challenge US Biometric measurements from fetal ultrasound	NeoBrainS12	RV Segmentation Challenge in Cardiac MRI	CAUSE07	2007		
<small>Peripheral Artery/Vein Multi-Organ Segmentation (PAVES) is the challenge focused on providing the most comprehensive segmentation of the peripheral arteries and veins.</small>	<small>The challenge tests the generality of multi-organ segmentation methods to 10 different anatomical regions.</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Associated with MICCAI 2016</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
REFUGE	Correction of Brainshift with Intra-Operative Ultrasound	The Continuous Registration Challenge	CATARACTS	RETOUCH	CARELYON17	Ultrasound Nerve Segmentation	BigNeuron	The Second Overlapping Cervical Cytology Segmentation Chal	ImageCLEFmed: Medical classification	3D Deconvolution Microscopy	LOLA11	Cardiac Motion Estimation Challenge 2011	DTI Segmentation Challenge in IVUS	SLIVER07	2007	2007	2007		
<small>The goal of the REFUGE dataset is to evaluate the performance of automated registration and computer-aided algorithms for guidance and surgical registration using a common dataset of patient brain images.</small>	<small>Submit your method for lung and brain registration!</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
LUMIC	IDRID	ICBIS2018-Challenge	Skin Lesion Analysis Towards Melanoma Detection	Cervical Cancer Screening	ENIGMA: Cerebellum	Pediatric Bone Age Challenge	Tumor Proliferation Assessment Challenge (TUPAC)	Statistical Atlases and Computational Modelling of the Heart - S	Low-Dose CT Grand Challenge	Cell Tracking Challenge	Shape 2015 - Statistical Shape Model Challenge	GRASS	Localization Microscopy Challenge	LOLA11	Cardiac Motion Estimation Challenge 2011	DTI Segmentation Challenge in IVUS	SLIVER07		
<small>The LUMIC challenge tests the accuracy of automated segmentation methods in CT lung images for segmenting lung CT with radiographic digitalization.</small>	<small>This challenge evaluates automated techniques for analysis of fundus photographs. The target categories include fundus photograph, macula, maculopathy, and hemorrhage and includes a test set of 10 images. Each image will be graded by fundus experts according to the severity level of fundus lesions.</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
EndoVis	MURA: Bone X-Ray Deep Learning Competition	RSNA Pneumonia Challenge	Multi-shell diffusion MRI harmonization and enhancement	Ischemic Stroke Lesion Segmentation 2017	ProstateC-2 Challenge	Multi-Modality Whole Heart Segmentation	HVSMP 2016	MITOS-ATYPIA-14	Digital Mammography DREAM Challenge	Cell Tracking Challenge (2nd Edition)	OVERLAPPING CERVICAL CYTOLOGY SEGMENTATION CHALLENGE	AMIDAT3	DIADEM	EMPIRE10	SK10	2007	2007	2007	
<small>An as endoscopic vision (CV) challenge to develop a deep learning framework for extracting the content of the field of view, gather features for training and testing, and segmenting vascular structures.</small>	<small>Automatically detect a visual signal for pneumonia in chest X-rays.</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
ImageCLEF Tuberculosis 2018	Multi-shell diffusion MRI harmonization and enhancement	Dermoscopic Skin Lesion Analysis Towards Melanoma Detection	The skin of the challenge is to evaluate the performance of automated methods for the detection of melanoma lesions from 2D images.	ACDC	LITS - Liver Tumor Segmentation	Digital Mammography DREAM Challenge 2017	C11	MITOS-ATYPIA-14	Digital Mammography DREAM Challenge 2017	Cell Tracking Challenge (2nd Edition)	OVERLAPPING CERVICAL CYTOLOGY SEGMENTATION CHALLENGE	ANODE09	EXACT09	ANODE09	EXACT09	ANODE09	EXACT09		
<small>Assess the probability of 10 patient having residual lung lesions based on 2D chest X-ray images.</small>	<small>The aim of the challenge is to evaluate the performance of automated methods for the detection of melanoma lesions from 2D images.</small>	<small>The goal of the challenge is to help participants develop image analysis tools to facilitate the automated detection of melanoma lesions from dermoscopic images.</small>	<small>The skin of the challenge is to help participants develop image analysis tools to facilitate the automated detection of melanoma lesions from dermoscopic images.</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>	<small>Open for submissions</small>			
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<small>Automated measurement of head circumference using 2D ultrasound images</small>	<small>Classification of prostate feature using morphometric ROI data</small>	<small>Classification of breast malignancy based on a set of visual features</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>	<small>Data download</small>			
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<small>Peripheral Artery/Vein Multi-Organ Segmentation (PAVES) is the challenge focused on</small>																			



MRBrainS

Evaluation framework for **MR Brain** Image **Segmentation**



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Welcome to the MRBrainS website. On September 26th, 2013 we organized the **Grand Challenge on MR Brain Image Segmentation workshop** at the **MICCAI** in Nagoya, Japan, where we launched this evaluation framework. The aim of the MRBrainS evaluation framework is to compare (semi-)automatic algorithms for segmentation of grey matter, white matter and cerebrospinal fluid on multi-sequence (T1-weighted, T1-weighted inversion recovery and FLAIR) 3 Tesla MRI scans of the brain. More information about the workshop and the results of the workshop challenge can be found [here](#).

Evaluation framework: The Challenge Continues

The MRBrainS website will remain open as an online evaluation framework for new segmentation algorithms or new versions of already applied algorithms. So feel free to register, download the training and test data and submit your results. Any team, whether from academia or industry, can join! You are welcome to submit your new unpublished method, but we would also like to encourage teams with well established published methods to register and submit their results. In this way, the MRBrainS challenge will present an overview of the performance of a wide variety of available brain structure segmentation methods.

How does it work?

The organisation of this challenge is similar to that of previous challenges described on **Grand Challenges in Medical Image Analysis**. Five datasets (MRI scans with manual segmentations) are provided for training and fifteen datasets (only MRI scans) are provided for testing. Teams can **register** on this website, sign the confidentiality agreement, and **download** the data. They can train their segmentation algorithm on the training data, apply it to the test data and **submit** the obtained results as well as a description of their algorithm. Each submission will be evaluated against the reference standard (manual segmentations) and the evaluation results will be sent to the team contact person by e-mail and published in the **results** section of this website.

More information can be found on the **details** page. If you have additional questions, feel free to send an email to mrbrains13@isi.uu.nl.

Rank	Team name	Submission name	Date	Sum Scores	Sequences used	Speed	Doc
1	TailHot	Multi-modality aggregation network³	13-04-18	33	T1; T1_IR; FLAIR	~13 sec	 
2	LIVIA_ETS	HyperDenseNet²	06-02-18	59	T1; T1_IR; FLAIR	~4 min	 
3	CU_DL2	3D Deep Learning; voxnet2	28-06-16	69	T1; T1_IR; FLAIR	~2 min	 
4	CU_DL	3D Deep Learning; voxnet1³	16-06-16	71	T1; T1_IR; FLAIR	~2 min	 
5	MSL-SKKU	Deep Convolutional Neural Network	19-06-17	78	T1; T1-IR; FLAIR	~1.5 min	 
6	LRDE	Fully Convolutional Network	20-12-16	78	T1	~2 sec	 
7	MDGRU	Multi-Dimensional Gated Recurrent Units³	27-07-16	100	T1; T1_IR; FLAIR	~2 min	
8	FBI/LMB Freiburg	U-Net (3D)	01-05-16	105	T1-1mm; T1-IR; FLAIR	~2 min	 
9	PyraMiD-LSTM2	NOCC with rounds³	23-05-16	109	T1; T1-IR; FLAIR	~2 min	
10	AOC	Atlas of Classifiers	24-12-17	117	T1	~6 sec	 
11	IDSLA	PyraMiD-LSTM	05-06-15	124	T1; T1_IR; FLAIR	~2 min	 
12	WTA2	3D Cascade convolutional architecture - Method 2	23-05-18	128	T1; T1_IR; FLAIR	~2 min	 
13	STH	Hybrid ANN-based Auto-context method²	03-06-16	139	T1; T1-IR; FLAIR	~ 5 min	 
14	ISI-Neonatology	Multi-stage voxel classification	31-05-14	143	T1	~1.5 hours	 
15	UNC-IDEA	LINKS:Learning-based multi-source integration	09-02-15	151	T1; T1_IR; FLAIR	~3 min	 
16	BCH_CRL_IMAGINE	3D patch-wise DenseNet and Patch Fusion²	24-05-18	169	T1; T1-IR; FLAIR	~2 min	 
17	KSOM GHMF	ASeTs: MAP-Based with Manifold learning	13-05-14	173	T1; T1_IR; FLAIR	~23 min	 
18	MNAB2	Random Forests	21-02-14	173	T1; T1_IR; FLAIR	~25 min	
19	WTA	3D Cascade convolutional architecture - Method 1³	15-05-18	190	T1; T1_IR; FLAIR	~5 min	 
20	vicorob UdG T1_F	MSSEG using T1 + FLAIR (T1-IR skull)	14-01-16	193	T1; IR; FLAIR	~2 min	 
21	VBM12	VBM12_r738 with WMHC=2	07-10-15	196	T1	~6 min	 
22	BIGR2	Multi-Feature SVM Classification	26-09-13	197	T1; T1_IR; FLAIR	~35 min	 
23	vicorob UdG T1	MSSEG using only T1 (T1-IR skull)	21-01-16	212	T1; IR	~2 min	 
24	UofL BioImaging	MAP-Based Framework	26-09-13	222	T1	~6 sec	 
25	CMIV	Model-guided Level Sets and Skeletons	26-09-13	241	T1; FLAIR	~3 min	 
26	UB VPML Med	Multi-Atlas with Multiway Cut	26-09-13	246	T1; T1_IR; FLAIR	~30 min	 
27	Narsil	Segmentation using Ensemble Trees	26-09-13	254	T1; FLAIR	< 2 min	 
28	VIBOT10 M-T1+F	Convolutional Neural Network	27-02-17	259	T1; FLAIR	~3 min	 
29	CVSS	Multi-atlas segmentation based on supervoxels	14-02-17	260	T1; T1_IR; FLAIR	~2 min	 
30	bigr_neuro	Automatically Trained kNN Classifier	26-09-13	265	T1; FLAIR	~2 hours	 
31	drai	VGG based Fully Convolutional Neural Network	20-01-17	266	T1	~2 sec	 

Screenshot
made a few
months ago.
As of today
They rank
#30

Rank	Team name	Submission name	Date	Sum Scores	Sequences used	Speed	Doc
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1	Team 1	VGG based Fully Convolutional Neural Network	20-01-17	266	T1	~2 sec

Automated Brain-Tissue Segmentation by Multi-Feature SVM Classification

Annegreet van Opbroek¹, Fedde van der Lijn¹ and Marleen de Bruijne^{1,2}

¹ Biomedical Imaging Group Rotterdam, Departments of Medical Informatics and Radiology, Erasmus MC - University Medical Center Rotterdam, the Netherlands

² Department of Computer Science, University of Copenhagen, Denmark

Abstract. We present a method for automated brain-tissue segmentation through voxelwise classification. Our algorithm uses manually labeled training images to train a support vector machine (SVM) classifier, which is then used for the segmentation of target images. The classification incorporates voxel intensities from a T1-weighted scan, an IR scan, and a FLAIR scan; features to encode the voxel position in the image; and Gaussian-scale-space features and Gaussian-derivative features at multiple scales to facilitate a smooth segmentation.

An experiment on data from the MRBrainS13 brain-tissue-segmentation challenge showed that our algorithm produces reasonable segmentations in a reasonable amount of time.

Rank	Team name	Submission name	Date	Sum Scores	Sequences used	Speed	Doc
1	TailHot	Multi-modality aggregation network³	13-04-18	33	T1; T1_IR; FLAIR	~13 sec	 
2	LIVIA_ETS	HyperDenseNet²	06-02-18	59	T1; T1_IR; FLAIR	~4 min	 
3	CU_DL2	3D Deep Learning; voxnet2	28-06-16	69	T1; T1_IR; FLAIR	~2 min	 
4	CU_DL	3D Deep Learning; voxnet1³	16-06-16	71	T1; T1_IR; FLAIR	~2 min	 
5	MSL-SKKU	Deep Convolutional Neural Network	19-06-17	78	T1; T1-IR; FLAIR	~1.5 min	 
6	LRDE	Fully Convolutional Network	20-12-16	78	T1	~2 sec	 
7	MDGRU	Multi-Dimensional Gated Recurrent Units³	27-07-16	100	T1; T1_IR; FLAIR	~2 min	
8	FBI/LMB Freiburg	U-Net (3D)	01-05-16	105	T1-1mm; T1-IR; FLAIR	~2 min	 
9	PyraMiD-LSTM2	NOCC with rounds³	23-05-16	109	T1; T1-IR; FLAIR	~2 min	
10	AOC	Atlas of Classifiers	24-12-17	117	T1	~6 sec	 
11	IDSIA	PyraMiD-LSTM	05-06-15	124	T1; T1_IR; FLAIR	~2 min	 
12	WTA2	3D Cascade convolutional architecture - Method 2	23-05-18	128	T1; T1_IR; FLAIR	~2 min	 
13	STH	Hybrid ANN-based Auto-context method²	03-06-16	139	T1; T1-IR; FLAIR	~ 5 min	 
14	ISI-Neonatology	Multi-stage voxel classification	31-05-14	143	T1	~1.5 hours	 
15	UNC-IDEA	LINKS:Learning-based multi-source integration	09-02-15	151	T1; T1_IR; FLAIR	~3 min	 
16	BCH_CRL_IMAGINE	3D patch-wise DenseNet and Patch Fusion²	24-05-18	169	T1; T1-IR; FLAIR	~2 min	 
17	KSOM GHMF	ASeTs: MAP-Based with Manifold learning	13-05-14	173	T1; T1_IR; FLAIR	~23 min	 
18	MNAB2	Random Forests	21-02-14	173	T1; T1_IR; FLAIR	~25 min	
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22	BIGR2	Multi-Feature SVM Classification	26-09-13	197	T1; T1_IR; FLAIR	~35 min	 
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24	UofL BioImaging	MAP-Based Framework	26-09-13	222	T1	~6 sec	 
25	CMIV	Model-guided Level Sets and Skeletons	26-09-13	241	T1; FLAIR	~3 min	 
26	UB VPML Med	Multi-Atlas with Multiway Cut	26-09-13	246	T1; T1_IR; FLAIR	~30 min	 
27	Narsil	Segmentation using Ensemble Trees	26-09-13	254	T1; FLAIR	< 2 min	 
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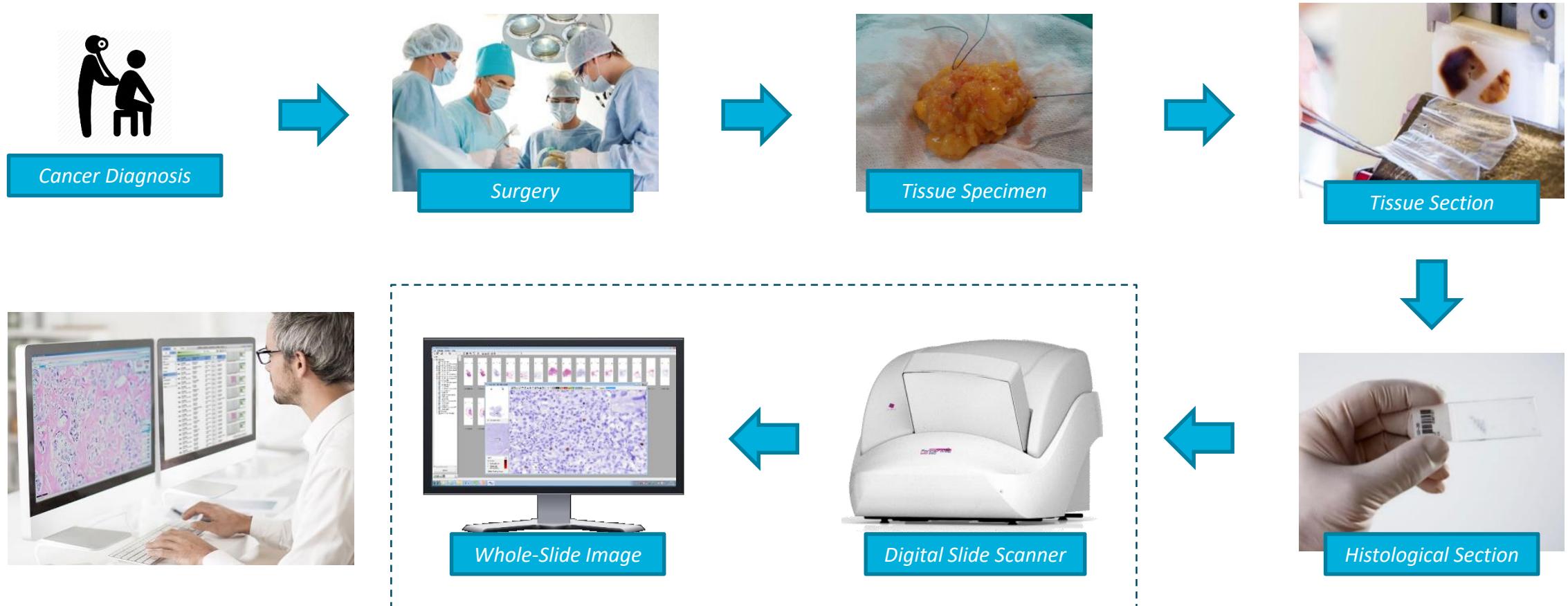
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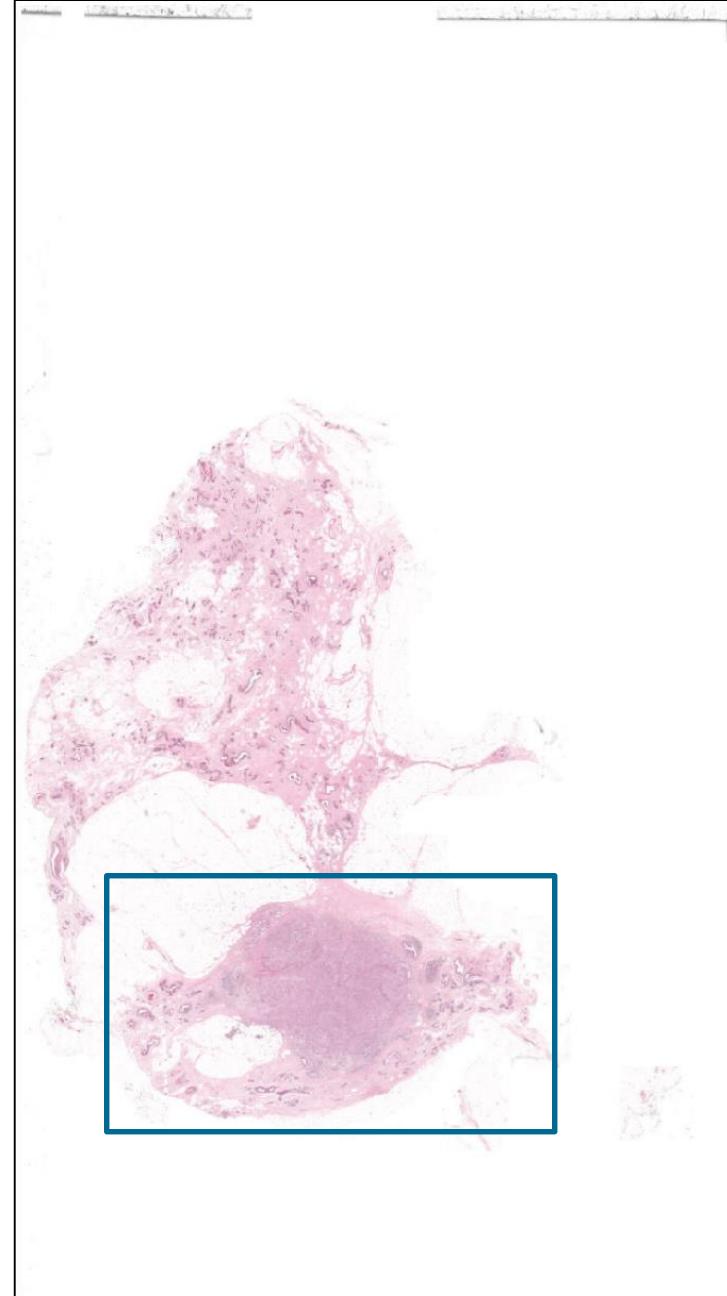
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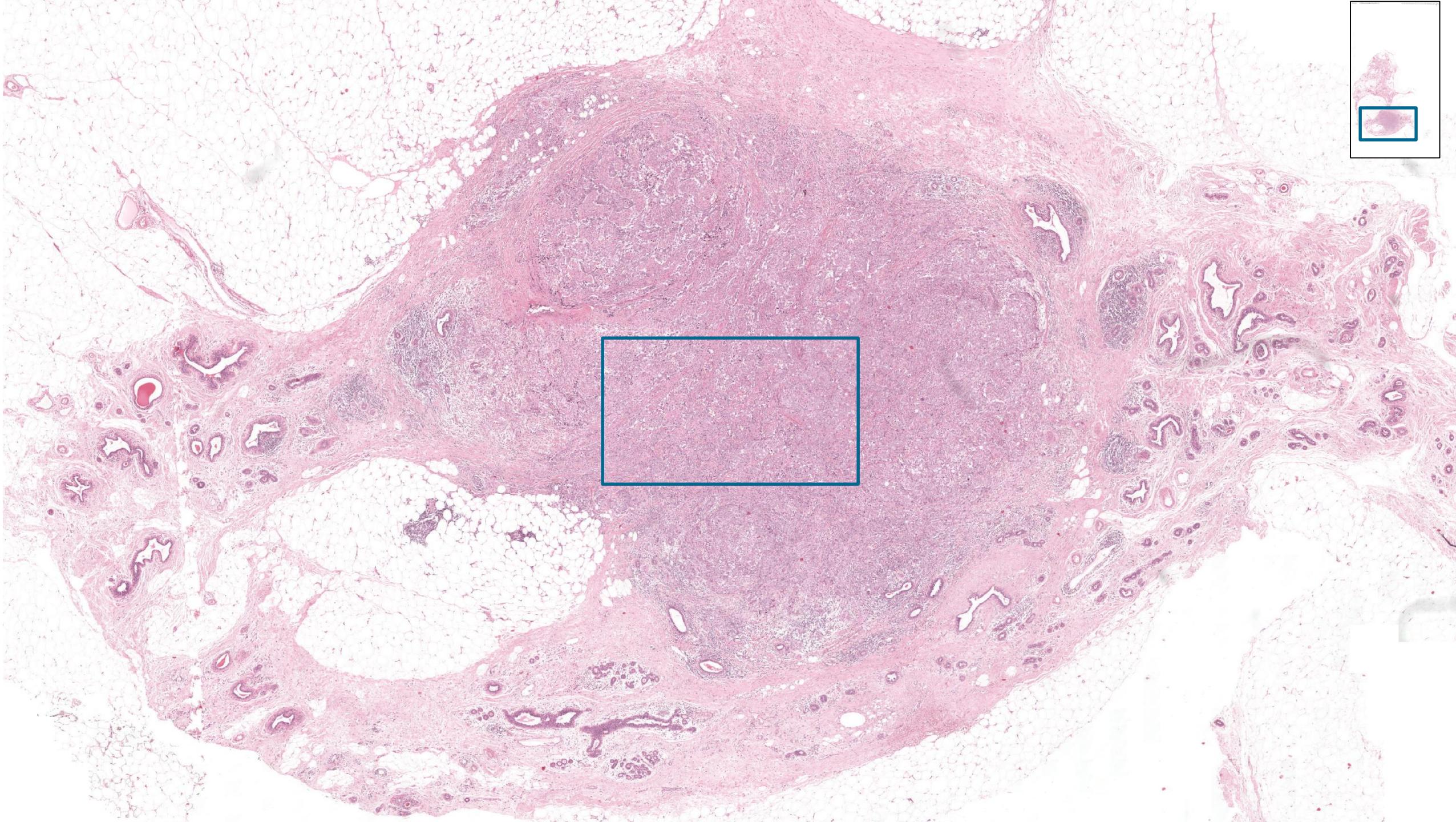
Good challenges run for years

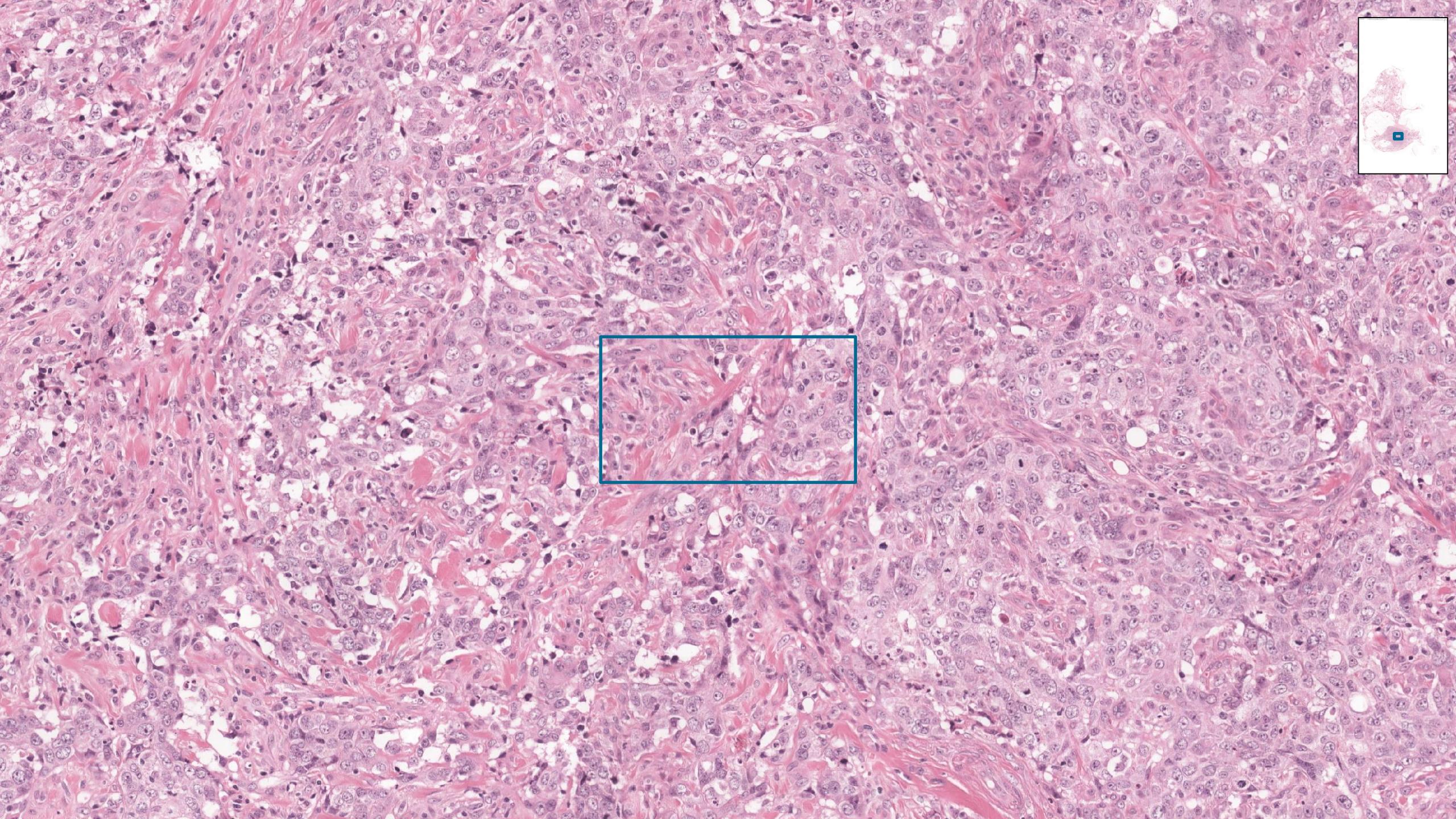
- ...if the dataset is good or exceptional and/or the problem is very important and not solved
- Maintaining challenges is not easy
- grand-challenge.org provides automated evaluation (via docker and support of a metric library evalutils). More later.
- The leaderboard and overview articles provide a ‘topical review’ of a field
- The leaderboards of a group of challenges show what are popular/powerful techniques in a field

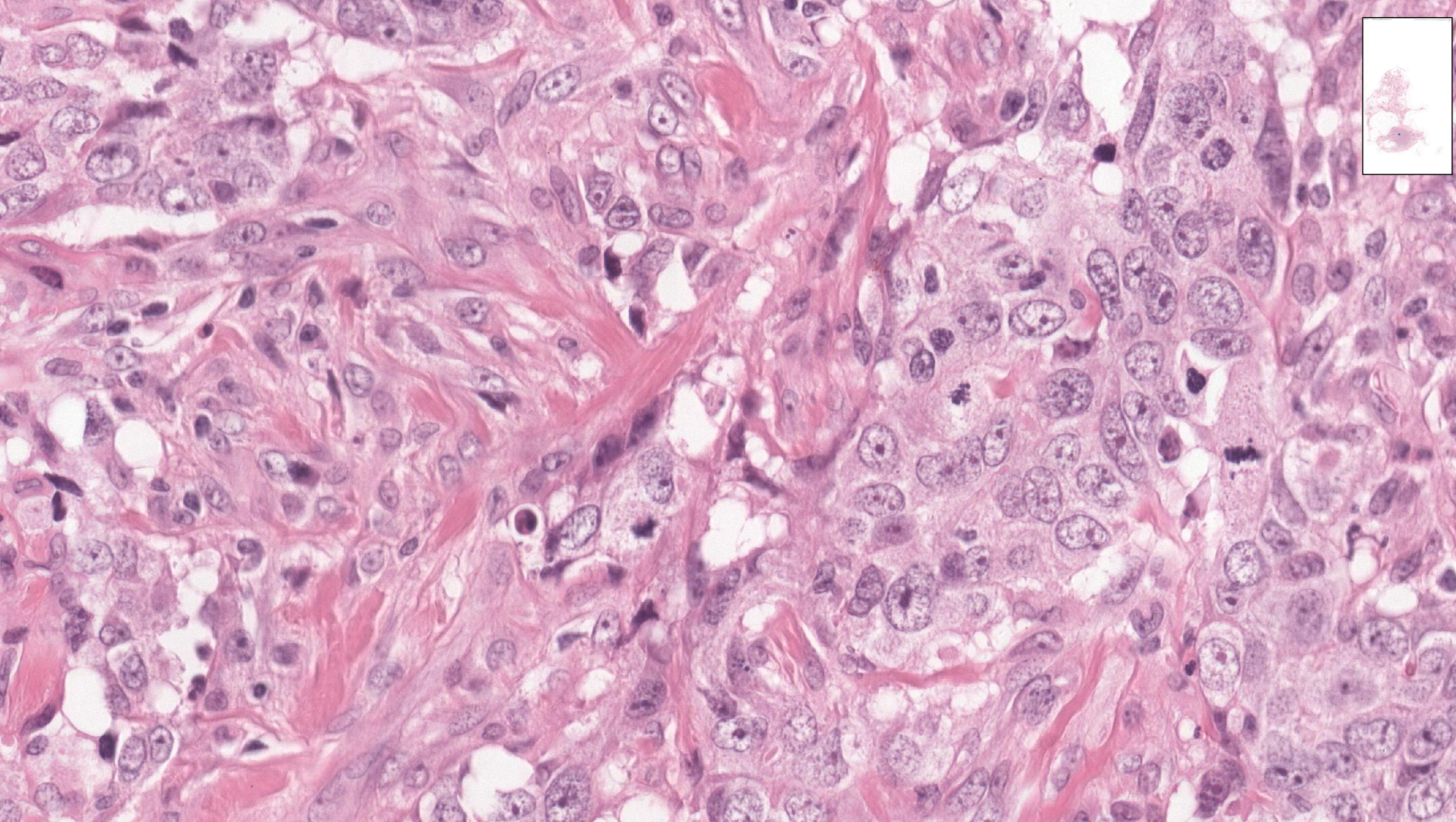
Digital Pathology

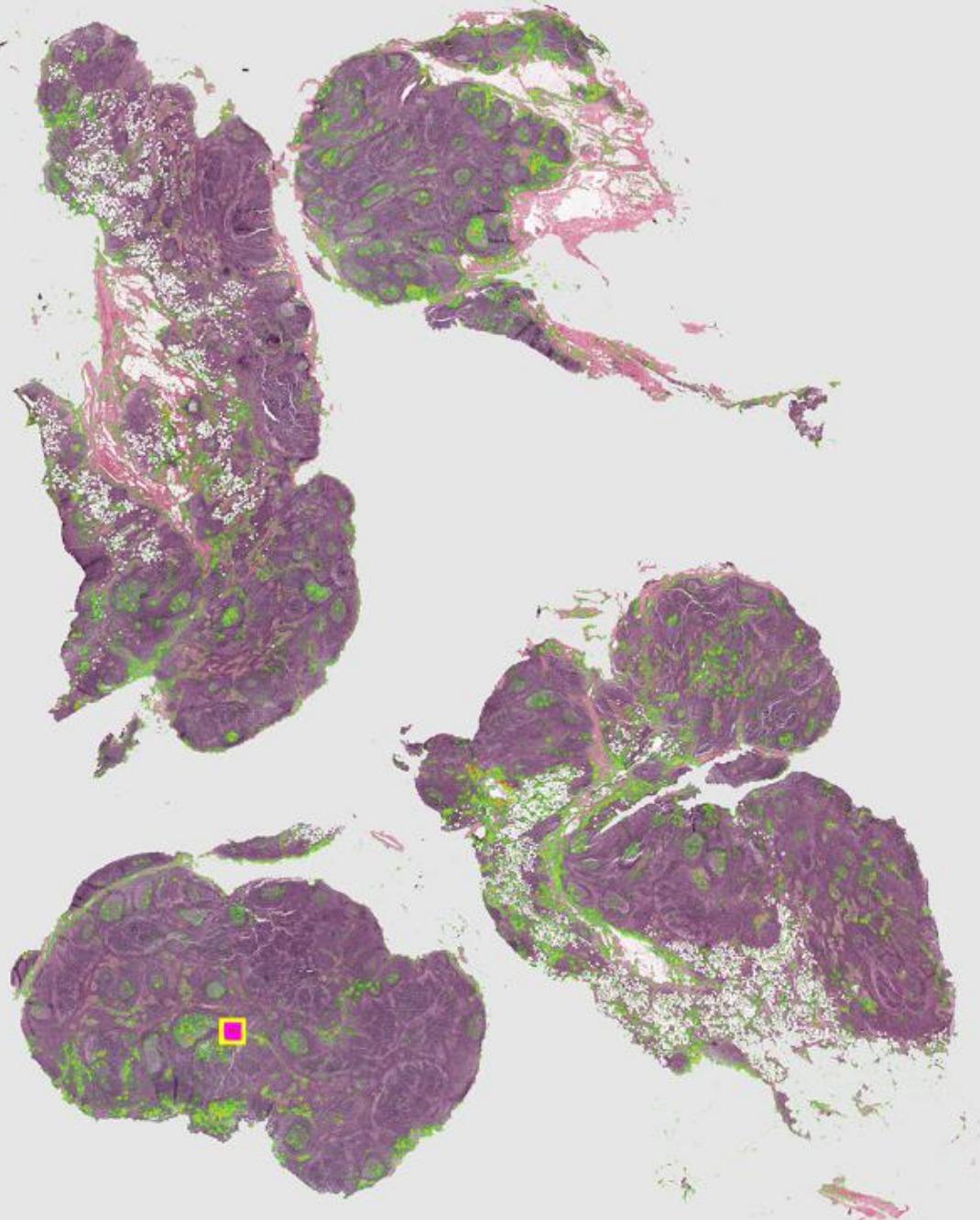












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Deep learning as a tool for increased accuracy and efficiency of histopathological diagnosis

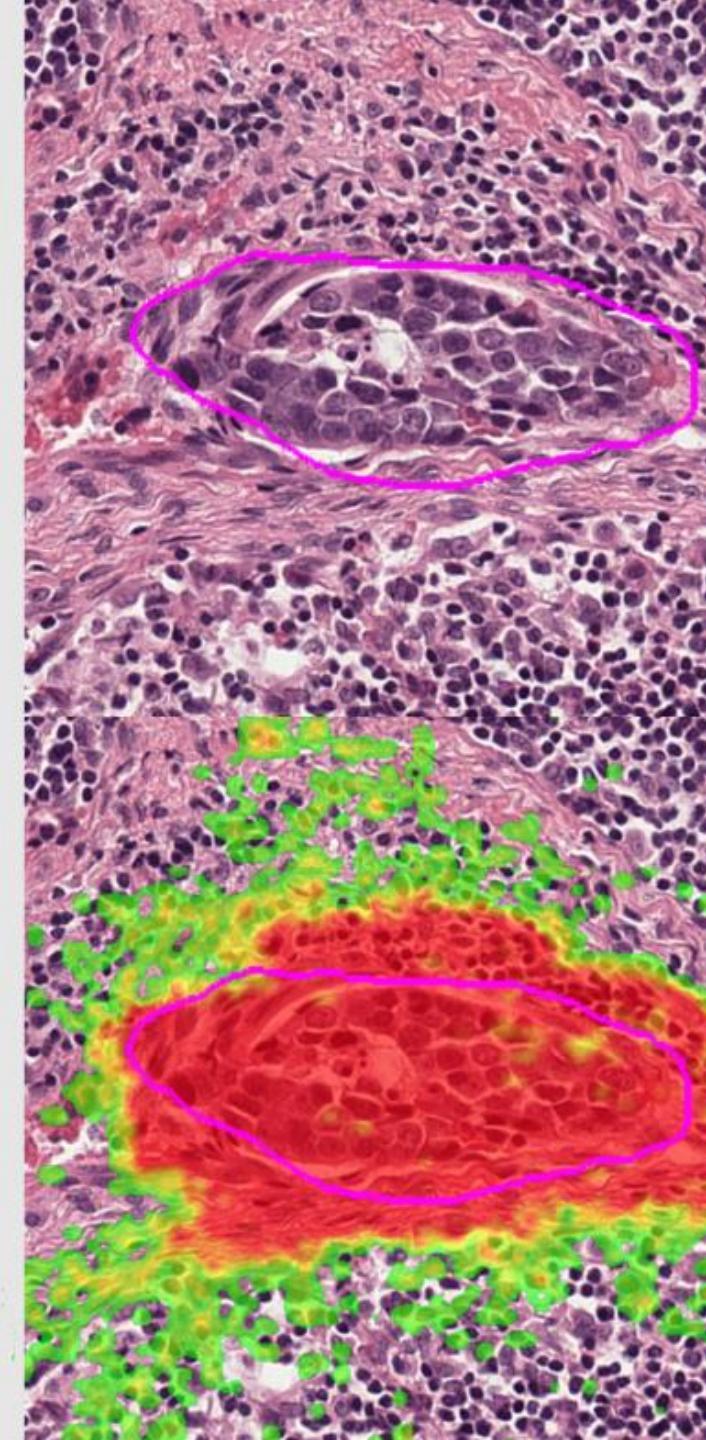
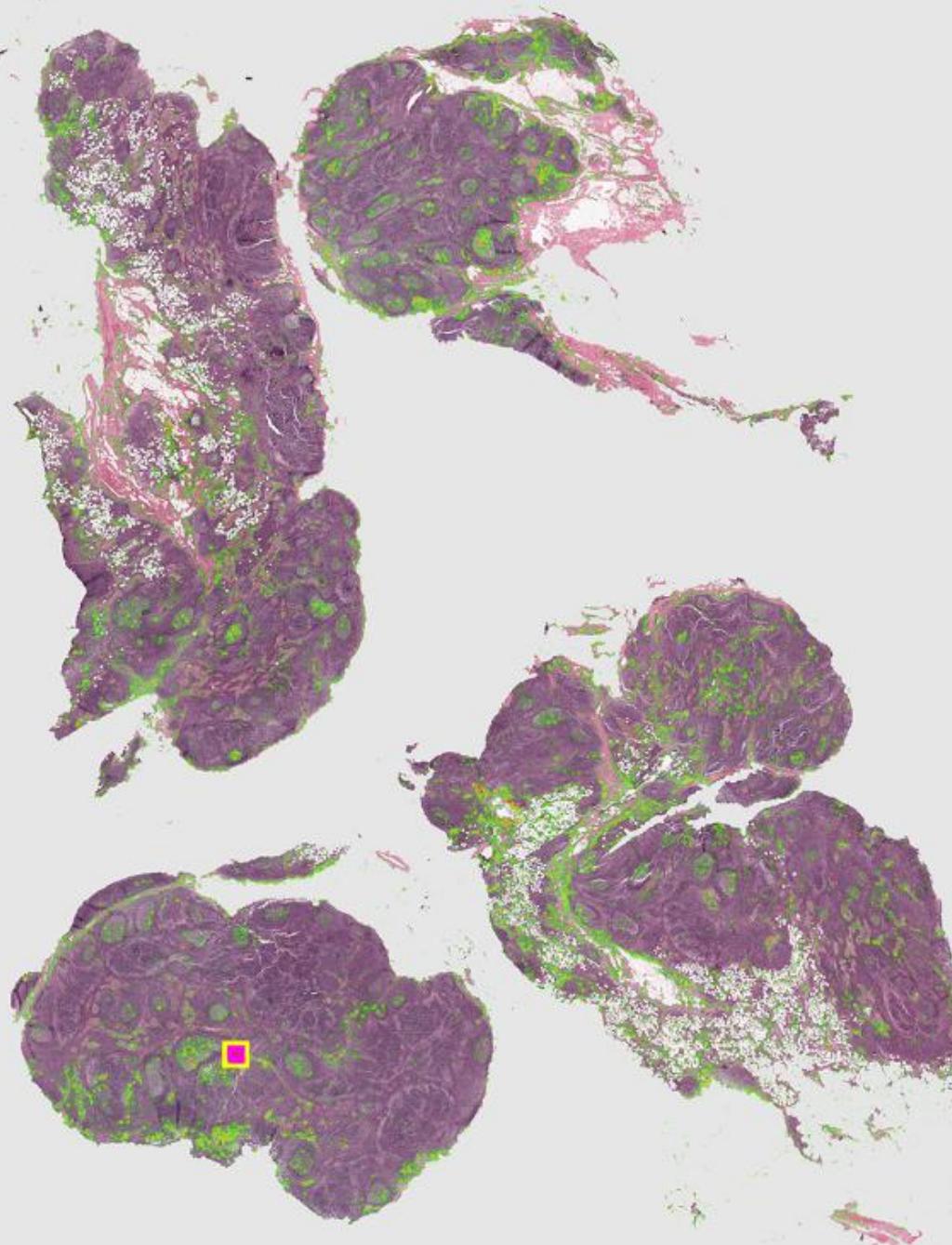
Received: 28 January 2016

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Geert Litjens¹, Clara I. Sánchez², Nadya Timofeeva¹, Meyke HermSEN¹, Iris Nagtegaal¹, Iringo Kovacs³, Christina Hulsbergen - van de Kaa¹, Peter Bult¹, Bram van Ginneken² & Jeroen van der Laak¹

Radboudumc



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SCIENTIFIC REPORTS

Using AI as a tool for
accuracy and efficiency
in pathological diagnosis

Nadya Timofeeva¹, Meyke HermSEN¹, Iris Nagtegaal¹,
Jeroen - van de Kaa¹, Peter Bult¹, Bram van Ginneken² &

Radboudumc



CAMELYON16



ISBI Challenge on cancer metastases detection in lymph node

Babak Ehteshami Bejnordi



Pathologist

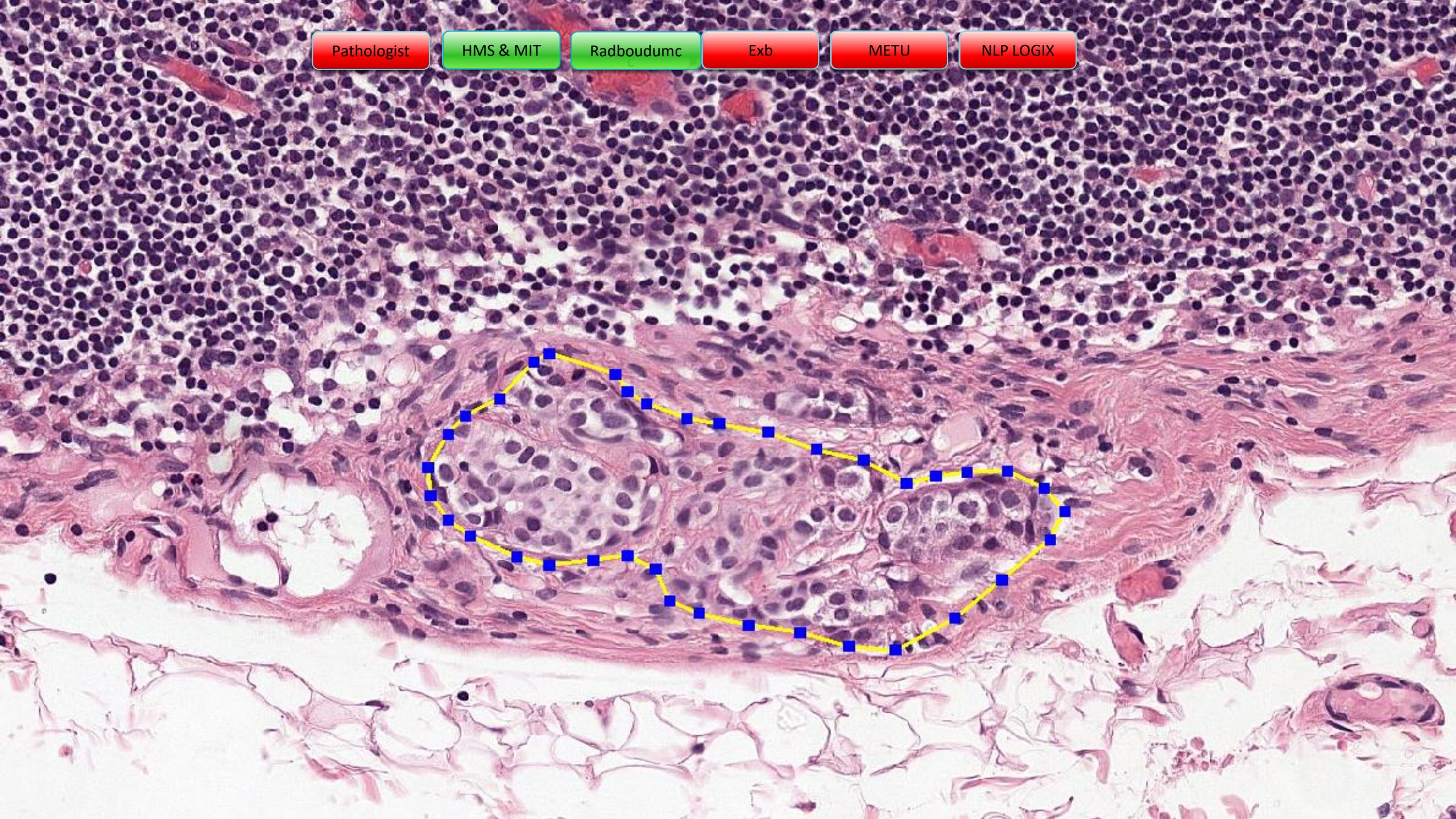
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METU

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THE NATIONAL ARTIFICIAL INTELLIGENCE RESEARCH AND DEVELOPMENT STRATEGIC PLAN

National Science and Technology Council

Networking and Information Technology
Research and Development Subcommittee

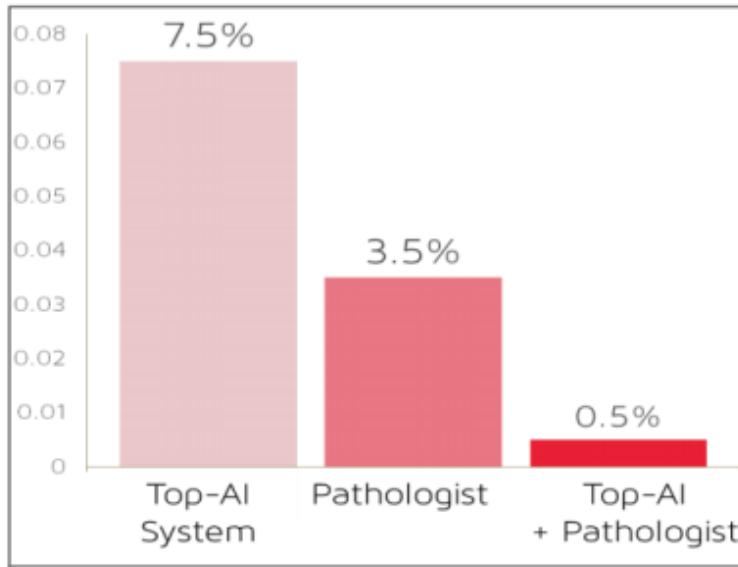
October 2016



National Institutes of Health (NIH) grants-supported research

ARTIFICIAL INTELLIGENCE FOR COMPUTATIONAL PATHOLOGY

Image interpretation plays a central role in the pathologic diagnosis of cancer. Since the late 19th century, the primary tool used by pathologists to make definitive cancer diagnoses is the microscope. Pathologists diagnose cancer by manually examining stained sections of cancer tissues to determine the cancer subtype. Pathologic diagnosis using conventional methods is labor-intensive with poor reproducibility and quality concerns. New approaches use fundamental AI research to build tools to make pathologic analysis more efficient, accurate, and predictive. In the 2016 Camelyon Grand Challenge for metastatic cancer detection,⁶⁹ the top-performing entry in the competition was an AI-based computational system that achieved an error rate of 7.5%.⁷⁰ A pathologist reviewing the same set of evaluation images achieved an error rate of 3.5%. Combining the predictions of the AI system with the pathologist lowered the error rate to down to 0.5%, representing an 85% reduction in error (see image).⁷¹ This example illustrates how fundamental research in AI can drive the development of high performing computational systems that offer great potential for making pathological diagnoses more efficient and more accurate.



AI significantly reduces pathologist error rate in the identification of metastatic breast cancer from sentinel lymph node biopsies.

JAMA | Original Investigation

Diagnostic Assessment of Deep Learning Algorithms for Detection of Lymph Node Metastases in Women With Breast Cancer

Babak Ehteshami Bejnordi, MS; Mitko Veta, PhD; Paul Johannes van Diest, MD, PhD; Bram van Ginneken, PhD; Nico Karssemeijer, PhD; Geert Litjens, PhD; Jeroen A. W. M. van der Laak, PhD; and the CAMELYON16 Consortium

IMPORTANCE Application of deep learning algorithms to whole-slide pathology images can potentially improve diagnostic accuracy and efficiency.

OBJECTIVE Assess the performance of automated deep learning algorithms at detecting metastases in hematoxylin and eosin-stained tissue sections of lymph nodes of women with breast cancer and compare it with pathologists' diagnoses in a diagnostic setting.

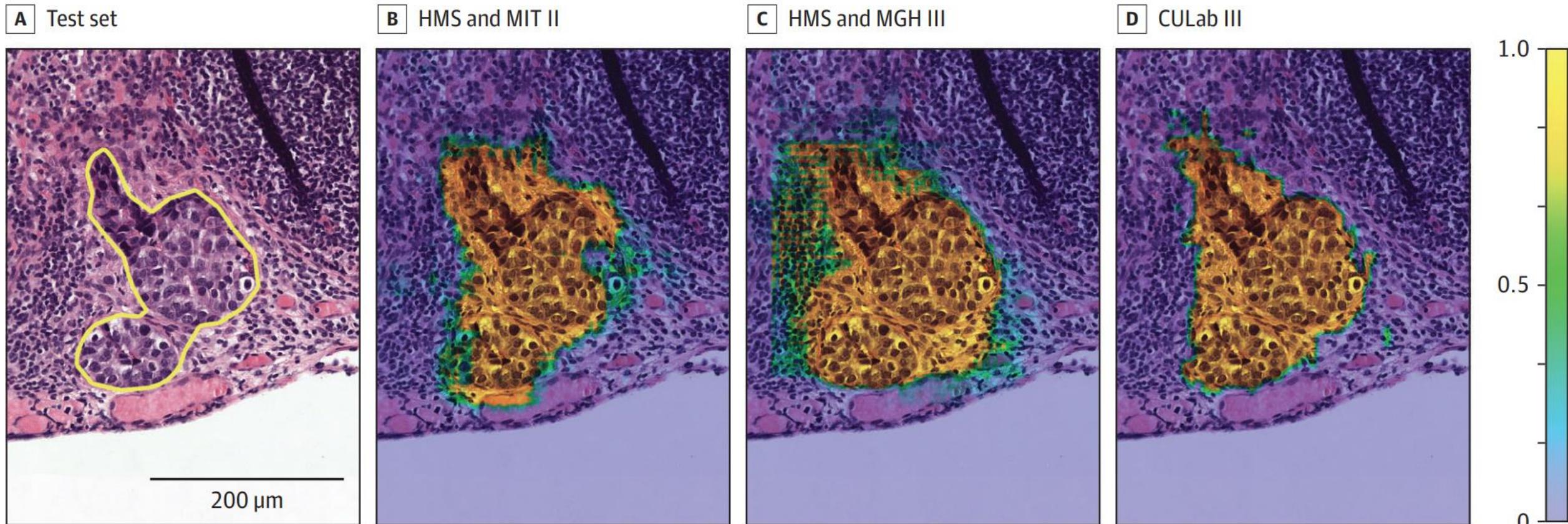
 Editorial page 2184

 Related articles page 2211 and page 2250

 Supplemental content

 CME Quiz at jamanetwork.com/learning and CME Questions page 2252

CONCLUSIONS AND RELEVANCE In the setting of a challenge competition, some deep learning algorithms achieved better diagnostic performance than a panel of 11 pathologists participating in a simulation exercise designed to mimic routine pathology workflow; algorithm performance was comparable with an expert pathologist interpreting whole-slide images without time constraints. Whether this approach has clinical utility will require evaluation in a clinical setting.



Advantages of challenges

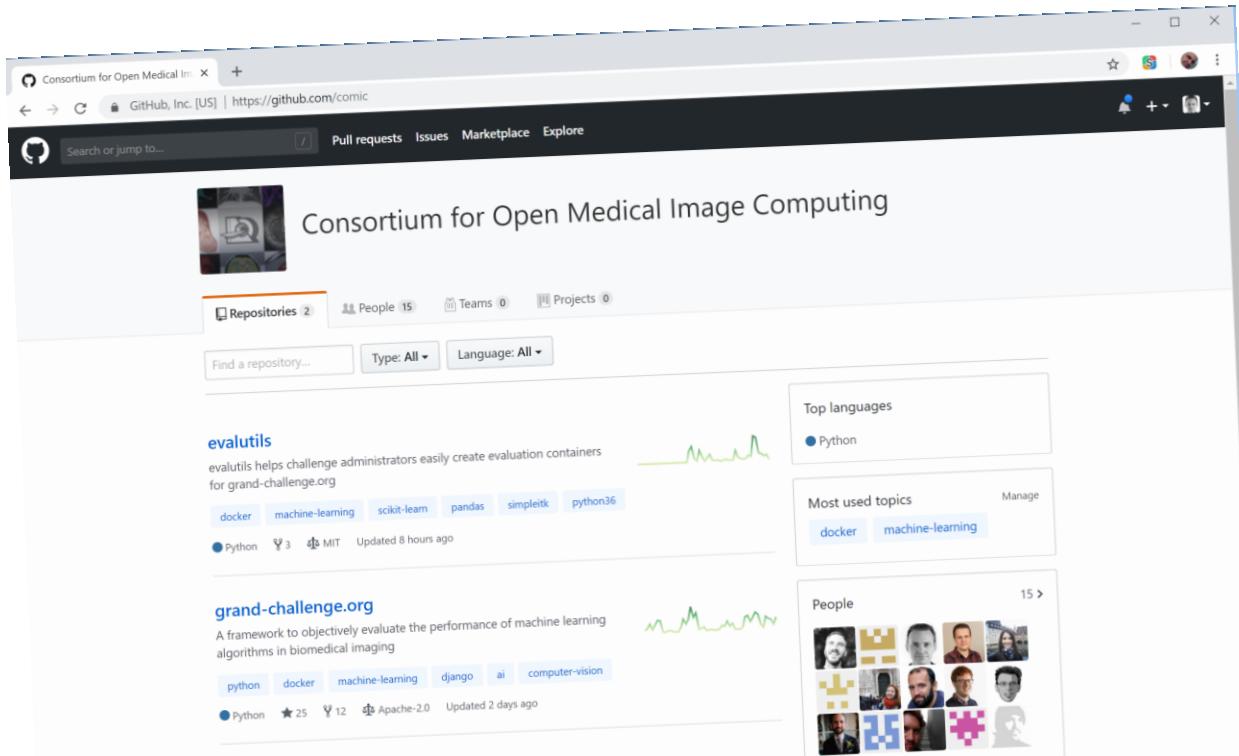
- **Compare** algorithms from the same task in a **fair and consistent** way
(same test data, same metric(s), same training data (if desired))
- Forces you to **think carefully** about aspects that relate to algorithm performance
- Leaderboard provides an **overview of techniques** that work well (are popular) for a task
- You can **combine** algorithms to see if they are complementary (even if you do not have access)
- Write an **overview article** to give a topical review enriched with real data
- Link the challenge to a **workshop** at a conference
- It is a way to **collaboratively solve an open research problem** and quickly move the field forward
- Challenges like ImageNet and Camelyon16 can have huge **impact**

Every paper a challenge?

- Why not create a challenge around every algorithm paper you write?
- Allows new techniques to be tested on your data
- Will generate citations to your paper
- You're pretty much doing everything already
 - Posing a question
 - Gathering data and annotations
 - Splitting this into train/valid/test data
 - Deciding on and implementing evaluation metrics
- Drawback: It's a lot of work
 - Creating a site
 - Participant management
 - Data sharing
 - Metrics definition
 - Submission gathering and continuous evaluation
 - Visualizing and comparing results

Simplifying the process with reusable code

- Groups in Nijmegen, Bremen, Rotterdam, Leuven, London set up the Consortium for Open Medical Image Computing (**comic**) in 2012
- Goal: make it easy to set up challenge websites with reusable components
- No funding, slow progress
- Became a priority when we saw it could be our internal platform in most research projects (and became a success when James Meakin started to work on it)



Open science

- Share data and annotations, evaluation code, algorithm code

The screenshot shows a web browser displaying the Zenodo website at <https://zenodo.org>. The page features a blue header with the Zenodo logo, a search bar, and navigation links for 'Upload' and 'Communities'. A 'Log in' and 'Sign up' button are also present. The main content area is titled 'Recent uploads' and lists three datasets:

- SemFi - Finnish Semantic Database with Syntactic Relations** (October 16, 2018, v2.1) by Mika Hämäläinen. It is a Dataset with Open Access. Description: SemFi is a semantic database for Finnish in which the words are linked to each other by the syntactic relations and their frequency in a big corpus. SemFi is based on the syntactic bigrams of The Finnish Internet Parsebank provided by Turku University. The semfi.db file is an SQLite database and...
Uploaded on October 16, 2018
2 more version(s) exist for this record
- sjPlot - Data Visualization for Statistics in Social Science.** (October 15, 2018, v2.8.1) by Daniel Lüdecke. It is Software with Open Access. Description: General Removed defunct functions. Deprecated sjt.lm(), sjt.glm(), sjt.lmer() and sjt.glmer() are now deprecated. Please use tab_model() instead. Changes to functions Arguments dot.size and line.size in plot_model() now also apply to marginal effects and diagnostic plots. plot_model() now uses a...
Uploaded on October 15, 2018
3 more version(s) exist for this record
- OpenAIRE's DOIBoost - Boosting CrossRef for Research** (October 1, 2018, v2.0) by La Bruzzo, Sandro; Manghi, Paolo; Mannocci, Andrea. It is a Preprint with Open Access. Description: Research in information science and scholarly communication strongly relies on the availability of openly accessible datasets of scholarly entities metadata and, where possible, their relative payloads. Since such metadata information is scattered across diverse, freely accessible, online...
Uploaded on October 11, 2018
2 more version(s) exist for this record

A sidebar on the right provides information about Zenodo's features:

- Zenodo now supports usage statistics!** (Read more about it, in our newest blog post.)
- Using GitHub?** (Just Log in with your GitHub account and click here to start preserving your repositories.)
- Zenodo in a nutshell**
 - Research. Shared.** – all research outputs from across all fields of research are welcome! Sciences and Humanities, really!
 - Citable. Discoverable.** – uploads gets a Digital Object Identifier (DOI) to make them easily and uniquely citable.
 - Communities** – create and curate your own community for a workshop, project, department, journal, into which you can accept or reject uploads. Your own complete digital repository!
 - Funding** – identify grants, integrated in reporting lines for research funded by the European Commission via OpenAIRE.
 - Flexible licensing** – because not everything is under Creative Commons.
 - Safe** – your research output is stored safely for the future in the same cloud infrastructure as CERN's own LHC research data.

Read more about Zenodo and its features.

Open science

- Share data and annotations, evaluation code, algorithm code

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Uploaded on October 11, 2018
2 more version(s) exist for this record
- Transfer of bacteriophages between the faeces of volunteers and water or saline** (October 10, 2018, v1.0) by [redacted]. Description: [redacted]

On the right side of the page, there are three boxes: "Zenodo now supports usage statistics!", "Using GitHub?", and "Zenodo in a nutshell".

The screenshot shows a GitHub repository page for "neuro-intracerebral-haemorrhage-segmentation" (Private). The repository has 3 pull requests, 0 issues, 0 projects, and 0 wiki pages. The master branch has 1 commit by AjayPatel86: "Code clean up". The commit details are as follows:

File	Description	Age
ipynb_checkpoints	Modified main file	3 months ago
data.py	Bug fixes in main file	3 months ago
data_augmentation.py	Bug fixes in main file	3 months ago
data_handling.py	Bug fixed in data handling	3 months ago
data_sampling.py	Bug fixes in main file	3 months ago
main.ipynb	Optimised evaluation	3 months ago
network_utils.py	Bug fixes in main file	2 months ago
prepare_data.py	Updated project files	3 months ago
stats.py	Optimised evaluation	3 months ago

At the bottom of the page, there are links for Contact GitHub, Pricing, API, Training, Blog, and About.

Open science

- Share data and annotations, evaluation code, algorithm code

The screenshot shows two side-by-side web pages. On the left is the Zenodo homepage, featuring a blue header with 'zenodo' and a search bar. It displays a message 'Zenodo now supports usage statistics!' with a chart icon. Below this is the Code Ocean homepage, which has a blue background with scientific illustrations and the text 'Discover & Run Scientific Code'. It claims 'Code Ocean is a cloud-based computational reproducibility platform'. A 'UPLOAD YOUR CODE' button is visible. At the bottom, there's a search bar and several featured projects: 'SOCIAL SCIENCES' (Oct 2018), 'ENGINEERING' (Jun 2018), 'MEDICAL SCIENCES' (May 2018), and 'SOCIAL SCIENCES' (Aug 2018). Each project has a brief description and a link.

The screenshot shows a GitHub repository page for 'neuro-intracerebral-haemorrhage-segmentation'. The top navigation bar includes 'Pull requests', 'Issues', 'Marketplace', and 'Explore'. The main area shows a list of files in the 'notebooks' directory, with commit history for each. The commits are as follows:

- ipynb_checkpoints: Modified main file, 3 months ago
- data.py: Bug fixes in main file, 3 months ago
- data_augmentation.py: Bug fixes in main file, 3 months ago
- data_handling.py: Bug fixed in data handling, 3 months ago
- data_sampling.py: Bug fixes in main file, 3 months ago
- main.ipynb: Optimised evaluation, 3 months ago
- network_utils.py: Bug fixes in main file, 2 months ago
- prepare_data.py: Updated project files, 3 months ago
- stats.py: Optimised evaluation, 3 months ago

At the bottom, there are links for 'Contact GitHub', 'Pricing', 'API', 'Training', 'Blog', and 'About'.

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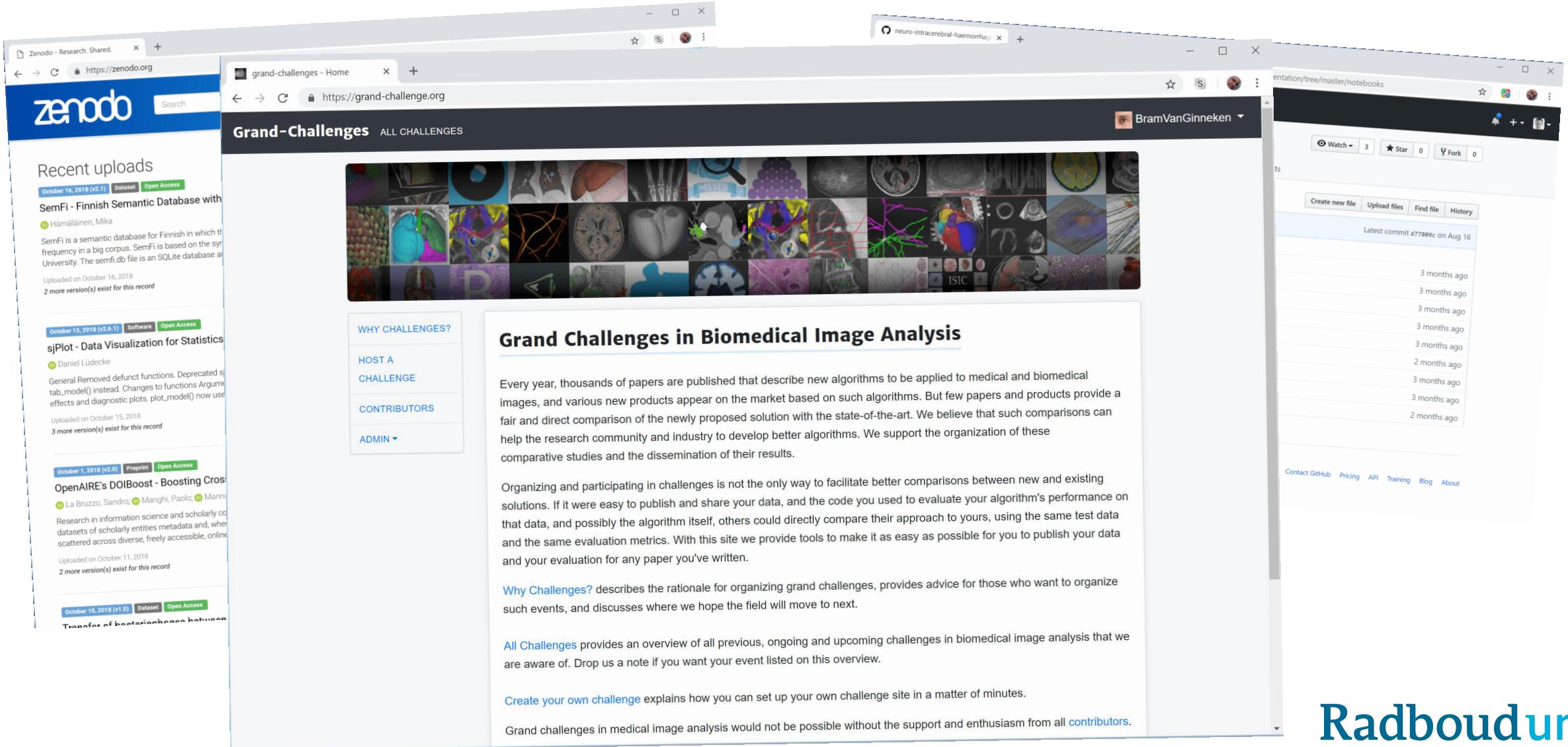
The screenshot displays two web pages side-by-side. On the left is the Zenodo website, featuring a blue header with 'zenodo' and a search bar. It shows a message 'Zenodo now supports usage statistics!' and a 'SIGN UP' button. Below this is the Code Ocean homepage with a blue background and the tagline 'Discover & Run Scientific Code'. It features a 'SIGN UP' button and a 'UPLOAD YOUR CODE' button. At the bottom, there's a search bar and several thumbnail previews of projects: 'SOCIAL SCIENCES' (Oct 2018), 'ENGINEERING' (Jun 2018), 'MEDICAL SCIENCES' (May 2018), and 'SOCIAL SCIENCES' (Aug 2018). Each preview includes a title and a brief description.

The screenshot shows the EnvoyAI for Developers website. The top navigation bar includes links for 'ABOUT', 'PLATFORM', 'PARTNERS', 'EXCHANGE', 'CONTACT', and 'BLOG'. The main heading is 'EnvoyAI for Developers' with the subtitle 'Add A Machine To The Exchange To Commercialize And Distribute Your Work To A Large Audience'. Below this is a large dark banner with a swirling pattern. The central section features the heading 'Empowering Innovation' and a paragraph explaining how to use the platform. It also highlights three key features: 'FREE CLOUD ACCOUNT', 'RAPID DISTRIBUTION', and 'GENEROUS REVENUE SHARING'. Each feature has a corresponding icon and a brief description.

Radboudumc

Open science

- Share data and annotations, evaluation code, algorithm code



Radboudumc



Statistics x +

https://grand-challenge.org/stats

Challenges

Public challenges: 35
Hidden challenges: 78
Public challenge with the most participants: LUNA16 (5,585 Participants)
Latest public challenge: ALL-challange

Registrations to public challenges in the past 30 days (top 10)

Yellow bar chart showing the number of registrations in the past 30 days for top 10 challenges.

Challenge	# of Registrations (past 30 days)
LUNA16	205
ICIP2018-Challenge	100
PROMISE12	75
IDRIID	70
REFUGE	65
HC18	58
CHAOS	55
PAVES	45
PROSTATEx	38
decathlon	35

Evaluations

Challenges using automated evaluation: 45
Total submissions: 1,959
Total submissions (last 30 days): 177
Public challenge with the most submissions: decathlon (197 Submissions)
Latest public result: Result for ICIAR2018-Challenge, created 11 hours ago, 126th position on leaderboard.

Evaluation

Long term sustainability of challenges depends on automating the evaluation

- Dealing with user submissions
- Generating and updating the results tables
- Running the same evaluation code after n years

Automated Evaluation - our approach

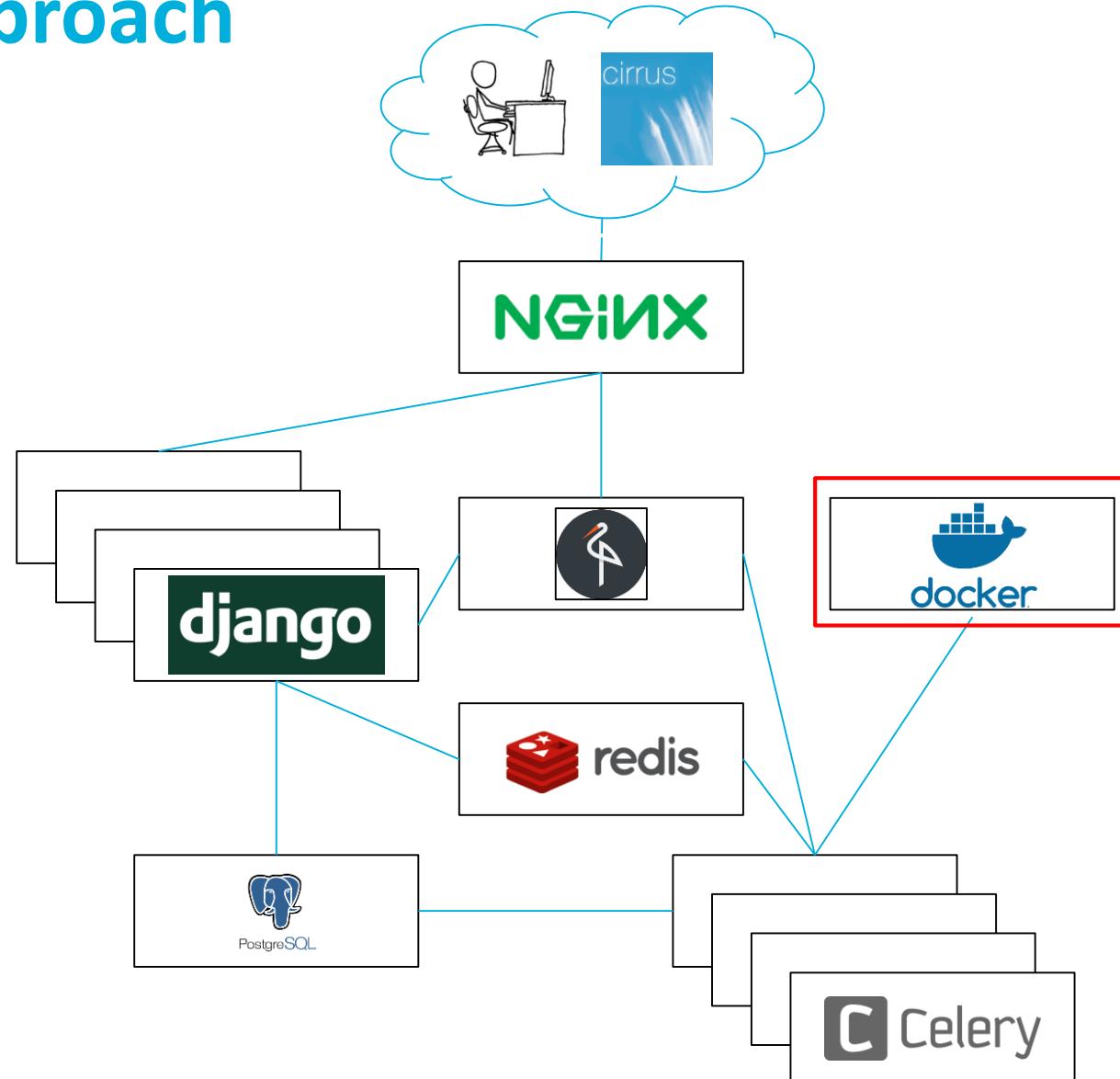
Use containers

- Challenge administrators can upload arbitrary code
- They are archived, and can run long term
- They are lightweight and can scale

Results stored in the database and tables dynamically updated

Define a clear API

- <https://grand-challengeorg.readthedocs.io/en/latest/evaluation.html#evaluation-container-requirements>
- The container will be executed on each submission, and only have access to this 1 submission
- The submission is available to the container on /input
- The container must produce /outputs/metrics.json
- Errors will be emailed to the user for them to fix



Radboudumc

Problem

Now every challenge administrator needs to create an evaluation container

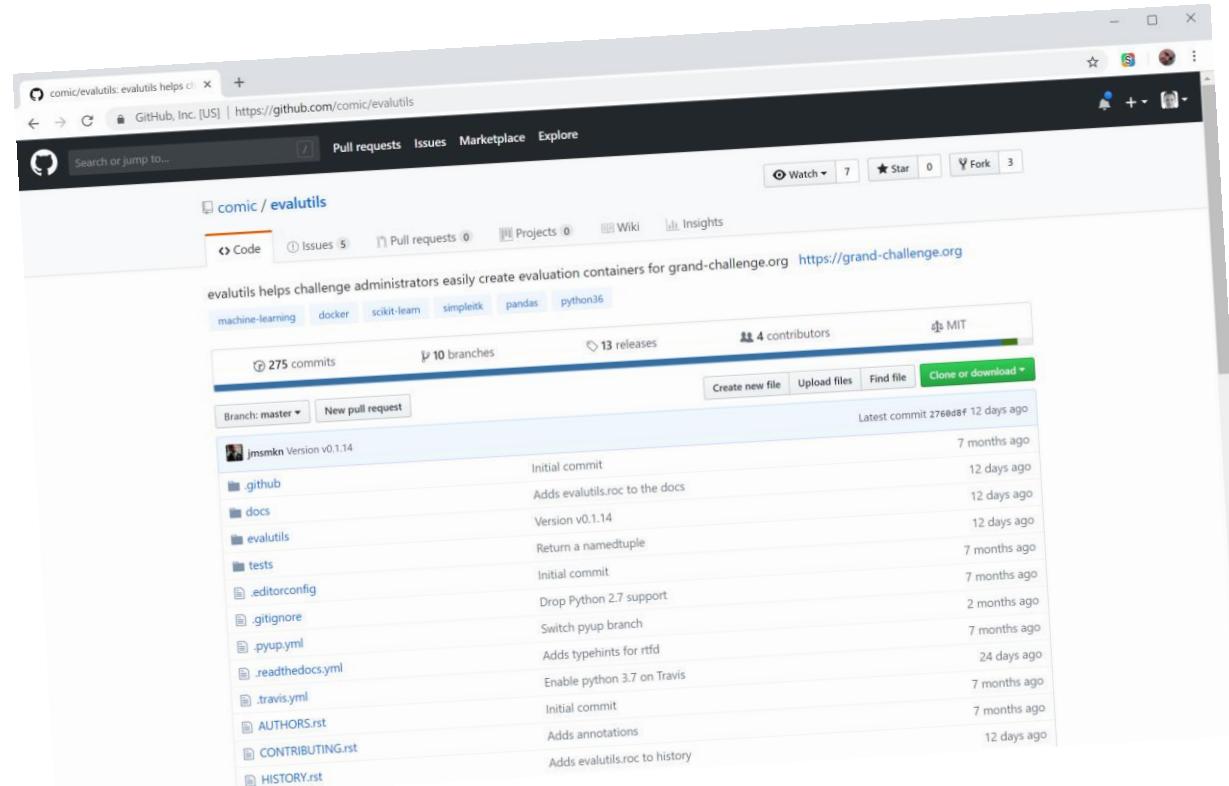
- Write a correct & maintainable dockerfile
- Load data
- Handle invalid submissions
- Repeated use of the same metrics across challenges
- Write the correct metrics to json

Evalutils

A tool to generate evaluation containers

- Boilerplate project generation
- Handles Classification, Segmentation and Detection tasks
- Pip installable
- IO module
- Validation module
- Metrics module

<https://github.com/comic/evalutils>



Grand-Challenges ALL CHALLENGES

SPLEEN18

HOME

EVALUATION

JOIN

TEAMS

SUBMIT

RESULTS

ADMIN ▾

Evaluation Methods

[+ Add a new method](#)

Show 10 entries

Search:

ID	Created	User	SHA
----	---------	------	-----

41162292-f53c-49be-a3cb-c719931d889e	Aug. 10, 2018, 12:53 p.m.	ghumpire	sha256:e34f8d7f26c5643fb6f02ee0934ca85d62eed59e9b26b0212aaeb33effb2b
e5270bcf-97a0-4aa3-b870-a725e9b8ebde	Aug. 10, 2018, 1:44 p.m.	ghumpire	sha256:2a50d0b17c88363fa0c9725c55599068e9e3dc43d3737f0be24feb68c474c

Showing 1 to 2 of 2 entries

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Grand-Challenges ALL CHALLENGES**SPLEEN18**[HOME](#)[EVALUATION](#)[JOIN](#)[TEAMS](#)[SUBMIT](#)[RESULTS](#)[ADMIN ▾](#)

Results

Show 10 entries

Search:

#	User (Team)	Created	Mean Dice	Max dice	Min dice	25pc dice	50pc dice	75pc dice	Mean jaccard	Mean volume similarity	Mean hausdorff distance	Comment
1st	ghumpire	10 Aug. 2018	0.9517	0.9792	0.7535	0.9492	0.9608	0.9703	0.9103	-0.0317	7.2430	

Showing 1 to 1 of 1 entries

[Previous](#) [1](#) [Next](#)

Grand-Challenges ALL CHALLENGES**SPLEEN18**[HOME](#)[EVALUATION](#)[JOIN](#)[TEAMS](#)[SUBMIT](#)[RESULTS](#)[ADMIN ▾](#)

Annotations

[+ Add images to this annotation set](#)[+ Set the labels on this annotation set](#)[Update the images used in this annotation set](#)**Kind**

Prediction

Dataset

ImageSet object (1e68a48d-3e65-4e73-9102-23f5a9bd86ba)

Upload Sessions[Upload Session <4b8ce781>, \(stopped\)](#)

Matched Annotations

Show [10](#) entriesSearch: **Key****Base Image**

(23922000369116261372423319636142128464583060', 633, 412, 412)	Image 1.2.392.200036.9116.2.6.1.37.2423319636.1421284645.83060 [633, 412, 412]
(23922000369116261372423319636143571331610597', 601, 410, 410)	Image 1.2.392.200036.9116.2.6.1.37.2423319636.1435713316.10597 [601, 410, 410]
(23922000369116261372423319636144487632458827', 713, 400, 400)	Image 1.2.392.200036.9116.2.6.1.37.2423319636.1444876324.58827 [713, 400, 400]

[←](#) [→](#) [⟳](#)https://apps.diagnijmegen.nl/Applications/CIRRUSWeb_master_98d13770/#!/?workstation=BasicWorkstation&grand_challenge_image=0cb7c22c-88b3-4544-969c-cc10c636acba&grand_chall...

Display



Maximum

1 mm

Abdomen

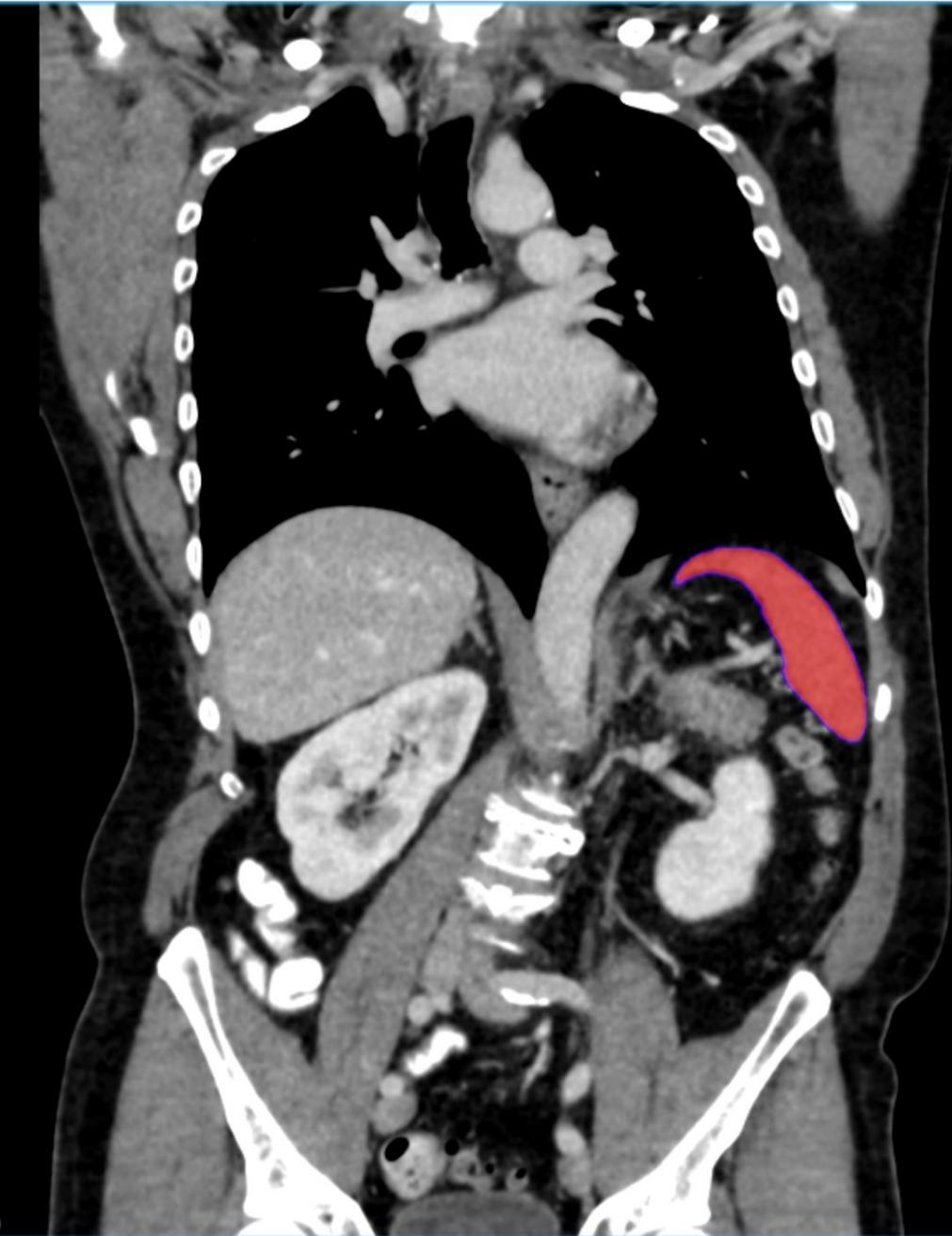
360

60

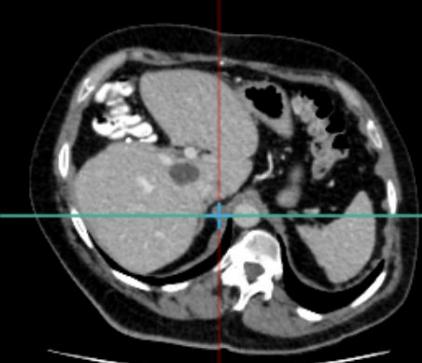
Orientation:

O A C S

Linking



200 mm



Toolbox: Scroll

Slice: 206 coronal
(142.4, -0.1, 1734.2): 120

Statistics x +

https://grand-challenge.org/stats

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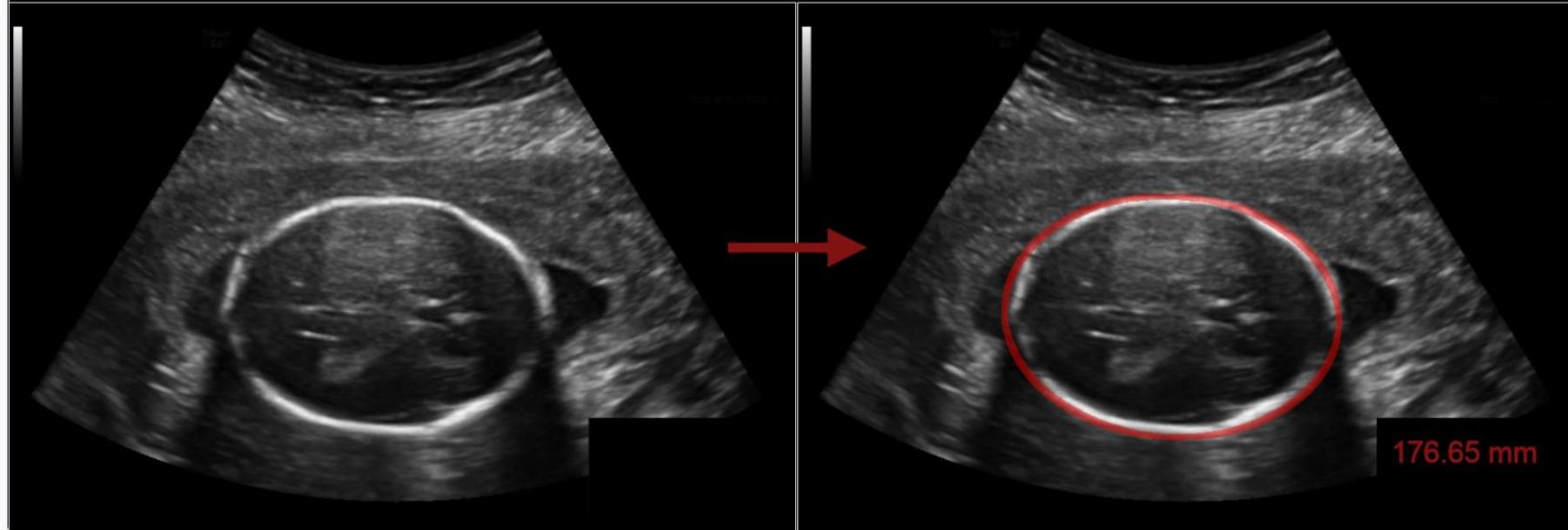
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Automated measurement of fetal head circumference

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Background

During pregnancy, ultrasound imaging is used to measure fetal biometrics. One of these measurements is the fetal head circumference (HC). The HC can be used to estimate the gestational age and monitor growth of the fetus. The HC is measured in a specific cross section of the fetal head, which is called the standard plane. The dataset for this challenge contains a total of 1334 two-dimensional (2D) ultrasound images of the standard plane that can be used to measure the HC. This challenge makes it possible to compare developed algorithms for automated measurement of fetal head circumference in 2D ultrasound images.

[Detailed description of the data](#)

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Search:

#	User	Created	Mean absolute difference (mm)	Std DICE (%)	Mean DICE (%)	Std difference (mm)	Mean difference (mm)	Std Hausdorff distance (mm)	Mean Hausdorff distance (mm)	Std absolute difference (mm)
1st	dra_v	9 Oct. 2018	2.3075	5.3681	95.2104	4.3083	0.1124	2.2359	2.4679	3.6378
2nd	thomasvandenheuvel	1 Oct. 2018	2.8332	2.7299	97.1010	4.2133	0.5603	1.6030	1.8319	3.1647
3rd	music	12 Oct. 2018	2.9370	7.2021	94.7254	5.5405	-1.6006	2.4486	2.6138	4.9614
4th	thomasvandenheuvel	1 Oct. 2018	3.0197	2.8001	97.0528	4.7317	0.7603	1.8327	1.9277	3.7179
5th	dra_v	1 Oct. 2018	3.4086	5.7906	94.5911	5.8620	0.3248	2.3650	2.8475	4.7765
6th	thomasvandenheuvel	1 Oct. 2018	3.6332	1.2309	97.2191	3.7182	2.8219	1.0795	1.7608	2.9278
7th	thomasvandenheuvel	1 Oct. 2018	4.1924	8.0644	95.4231	8.9749	1.3605	2.8286	2.3524	8.0484
8th	shenzexu	24 Oct. 2018	5.4308	10.0279	93.4925	10.0083	2.2773	3.6123	3.2561	8.7055
9th	631498444	11 Oct. 2018	6.3326	9.3412	94.6314	11.5555	4.8193	4.6598	3.2785	10.798
10th	631498444	9 Oct. 2018	6.4401	16.5091	87.8221	14.6160	1.2314	7.3371	7.4234	13.173

Showing 1 to 10 of 15 entries

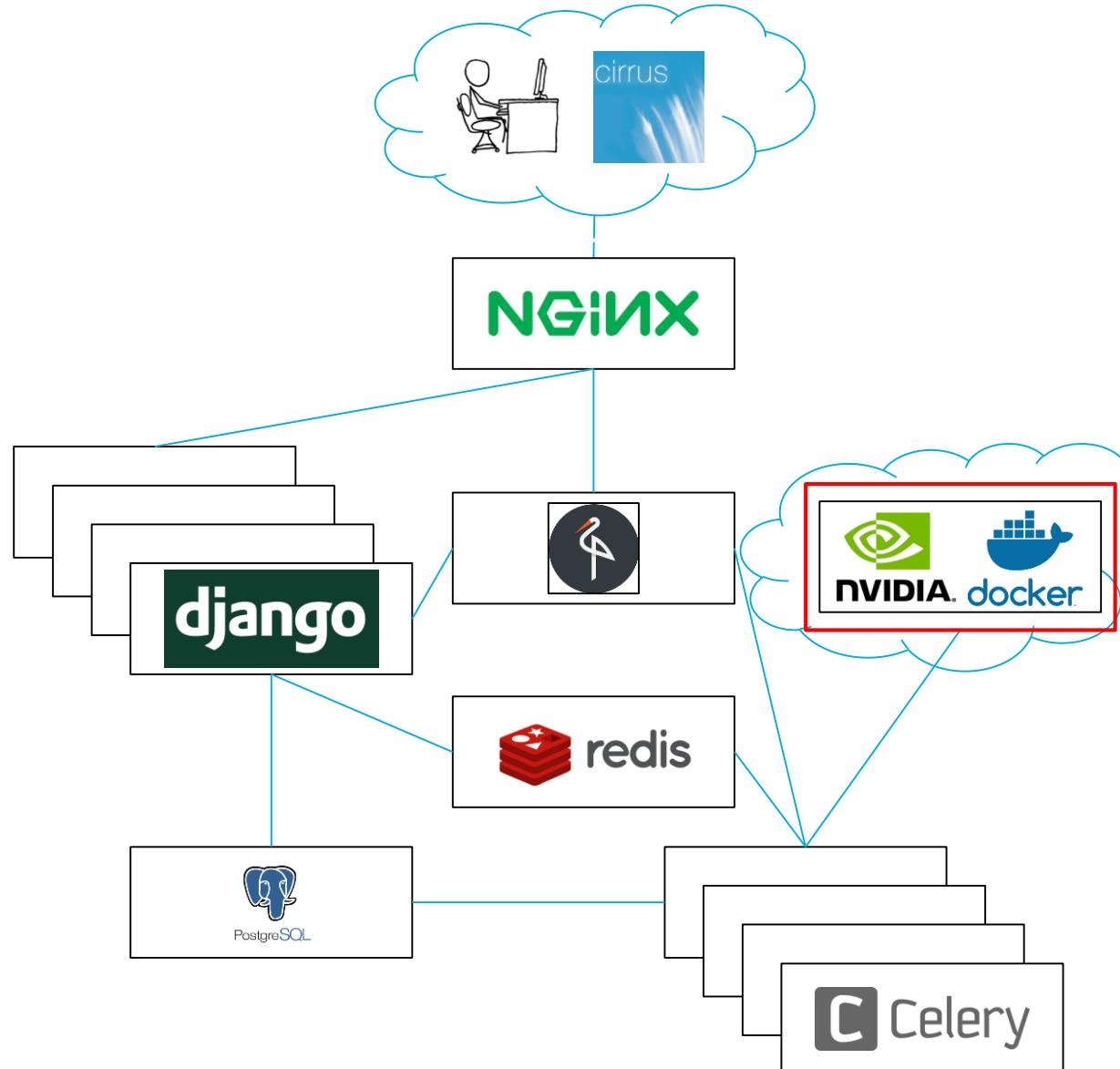
Previous **1** 2 Next

Next steps

- We can use containers not only for the evaluation, but also for the algorithms
- Users/teams upload a docker container with an algorithm (and a license)
- Algorithm is applied to secret test data → leaderboard entry created
- Teams could also upload containers to **train** systems on secret training data and download trained model afterwards (e.g. Mammography DREAM challenge used such a set up)
- External user could upload data and run algorithms on it, possibly for a fee (app store)
- These cloud based algorithms/apps could be embedded in a cloud based viewing environment
- External users could also assist in annotating data; crowd-sourcing platform that integrates data storage, annotation tools, algorithms

Algorithm execution

- Run on DIAG infrastructure or AWS
 - Choice of CPU only, K80 or V100
- Improvements
 - Separate queues for evaluations and algorithm executions
 - Speed
 - Docker registry on AWS
 - Data replica on AWS/apps.diagnijmegen.nl
 - Scaling and cost: spot instances w/ cost tracking?
 - Algorithm permissions: who can run it?
 - Service based architecture for integration in apps?
 - Results interface



LOGGLY

Radboudumc

Grand-Challenges ALL CHALLENGES

Algorithms

[+ Add a new algorithm](#)

Existing Algorithms

Show entriesSearch: **Title****Created****Creator**[Cardiomegaly Detection](#)

Sept. 13, 2018, 1:31 p.m.

[jamesmeakin.diag](#)

Showing 1 to 1 of 1 entries

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Dataset

3 different datasets are used in the experiment. In particular:

1. In the training phase, [SCR Dataset](#) and [JSRT](#) are utilized. Both of the dataset are publicly available and can be downloaded. JSRT dataset consists of 247 PA chest x-rays. SCR dataset provides the segmentation result for the lung fields, heart and clavicles on 247 images taken from JSRT dataset.
2. In the evaluation phase, publicly available the [ChestX-ray14 dataset](#) is utilized. It comprises 112,120 frontal-view chest X-ray images of 30,805 unique patients with 14 disease labels.

[Try out this algorithm](#)

Jobs for this algorithm

Show entriesSearch:

Created	Creator	Image	Status	Result
Oct. 25, 2018, 3:08 p.m.	pkcakeout248130100a1c4c02	Image cardiomegaly.mhd [1024, 1024]	The task executed successfully	{'out.mhd': {'cardiomegaly': {'errors': [], 'classification': True}}}
Sept. 14, 2018, 11:48 a.m.	pkcakeout248130100a1c4c02	Image 00000089_000.mhd [1024, 1024]	The task executed successfully	{'out.mhd': {'cardiomegaly': {'errors': [], 'classification': False}}}
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Sept. 14, 2018, 8:53 a.m.	jamesmeakin.diag	Image 00000008_002.mhd [1024, 1024]	The task executed successfully	{'out.mhd': {'cardiomegaly': {'errors': [], 'classification': True}}}
Sept. 14, 2018, 8:53 a.m.	jamesmeakin.diag	Image 00000008_001.mhd [1024, 1024]	The task executed successfully	{'out.mhd': {'cardiomegaly': {'errors': [], 'classification': False}}}



Run an algorithm

Select the images that you would like to run the algorithm on.

Image files*

[Choose Files](#)

Upload images for creating a new archive

[Submit](#)

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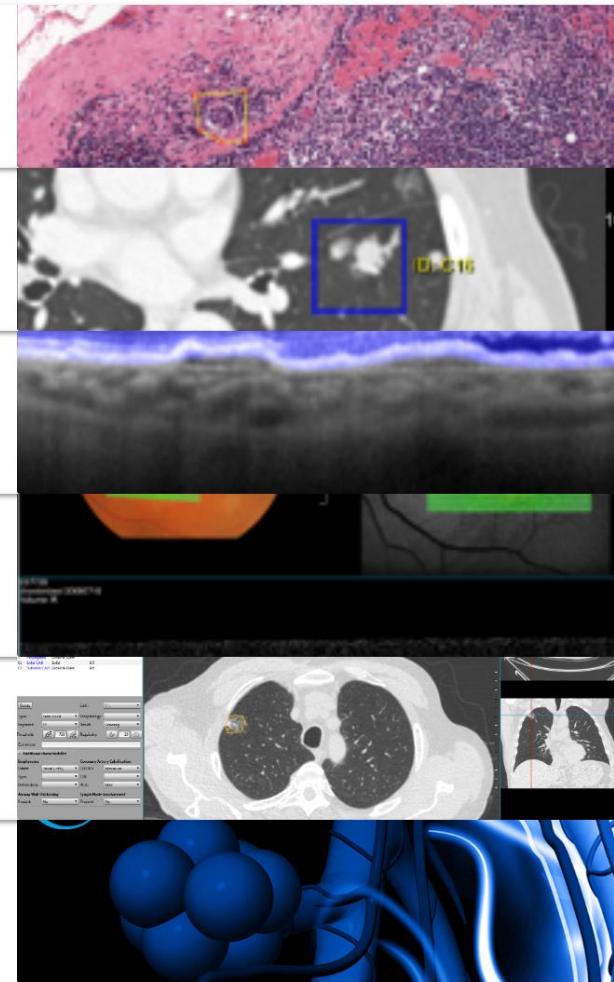
[Try out this algorithm](#)

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DIAG Applications

[CIRRUS Pathology](#)[CIRRUS Oncology](#)[AMI Ophthalmology AMD Workstation](#)[AMI Ophthalmology Multimodal Workstation](#)[CIRRUS Lung Screening](#)[CIRRUS Lung Screening \(Veolity branding\)](#)

Summary

- Benchmarking / challenges / competitions are useful for many reasons
- A paper or research project in which some algorithm is developed is a kind of challenge where the author is the only participant (and possibly some own implementations of prior work are shown to be outperformed)
- A challenge platform can be used for research projects (and for courses, etc etc)
- Excellent open source components exist today to build such a platform in a modular fashion
- Containers/dockers make sharing of running code easier; this will become more and more common
- Other use cases: app stores, could/web software for viewing and processing medical images, joint annotation and processing
- Please join and use/develop open source tools for benchmarking and reproducible science!

Table 2. Segmentation error in comparison with a standard ASM and numbers from previous work (customized ASM [22] and deformable simplex mesh [5]), given as $\mu \pm \sigma$

Segmentation method	$D_{avg}[\text{mm}]$	$D_{RMS}[\text{mm}]$	$V_D[\%]$
Deformable model	1.6 ± 0.5	3.3 ± 1.2	5.1 ± 1.4
Active Shape Model	2.9 ± 1.1	5.2 ± 2.3	8.9 ± 2.4
Lamecker et al. [22]	2.3 ± 0.3	3.1 ± 0.5	7.0 ± 1.8
Soler et al. [5]	$2 \pm ?$	n.a.	n.a.