Andrew L. Beers

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https://github.com/AndrewBeers
https://github.com/QTIM-Lab

Education

Brown University, 3.76 GPA, Magna Cum Laude
- Concentration: Environmental Studies
- *Best Research Project 2015*, Institute at Brown for Environment and Society.
- *Library Innovation Prize 2015*, Brown University.
- *Explore Grant 2013*, Social Innovation Initiative (SII) at Brown University.

Research

Quantitative Tumor Imaging Lab, Center for Machine Learning @ the MGH/HST Martinos Center for Biomedical Imaging
Boston, MA, Research Assistant (2016-2017), Programmer (2017)
- Design deep learning models for 3D image segmentation, image synthesis, image classification, regression, denoising, and superresolution of medical imaging data using Tensorflow and Keras.
- Develop open-source software packages (3D Slicer, Glue, DeepNeuro) for both clinical and academic users.
- Create in-house machine learning pipelines for diagnosis, prognosis, and treatment planning for brain tumors. Facilitate the usage of this pipeline by clinicians testing new treatments in ongoing clinical trials.
- Design curricula and give lectures for machine learning classes at MIT and the Martinos Center.

Cedar Creek Ecosystem Science Reserve, NSF Long-Term Ecological Research Site
East Bethel, MN, Research Intern (2015)
- Formalized and enacted a protocol for surveying diseased trees in Cedar Creek's forests, in support of a larger project on oak wilt epidemiology.

Brown University Center for Environmental Studies
Providence, RI, Research Assistant (2014-15)
- Made a website to visualize 134 years of iceberg data using Javascript, and d3.js. Modeled iceberg observer behavior to determine unrecorded changes in observers in the historical record.

American Civil Liberties Union
Boston, MA, Researcher (2014)
- Contributed to a rebuttal for an expert witness in an upcoming state-level reproductive justice case, and critiqued statistical methods in the opposition's supporting epidemiological literature.

Employment

American Civil Liberties Union
New York, NY, Online Production Assistant (2015-16)
- Ran statistical analysis on the ACLU's email fundraising campaign, and coded responsive donation pages.

Ceres
Boston, MA, Insurance Intern (2014)
- Created a tool with VBA and Bloomberg to track eco-friendly "green bonds" in the global market.

Planned Parenthood Federation of America
Washington, DC, Digital Fundraising Intern (2013)
- Created a tool in R to evaluate and rank the effectiveness of PPFA's national fundraising campaigns.

TIC
Washington, DC, Programming Counselor (Summers 2011/2012)
- Created and taught a programming curriculum using LOGO for game creation aimed at children aged 6-14.

Publications

Semi-Automated Pulmonary Nodule Interval Segmentation using the NLST data. Yoganand Balagurunathan, Andrew Beers, Jayashree Kalpathy-Cramer, Michael McNitt-Gray, Lubomir Hadjiiski, Bensheng Zhao, Jiangguo Zhu, Hao Yang, Stephen S.F. Yip, Hugo J.W.L. Aerts, Sandy Napel, Dimitry Cherenov, Kenny Cha, Heang-Ping Chan, Carlos Flores, Alberto Garcia, Robert Gillies, Dimitry Goldgof. Accepted for publication at Medical Physics (2017).
- We compare state of the art lung CT nodule segmentation methods from several institutions, evaluate their effectiveness at estimating nodule growth between two time points, and grade their ability to separate malignant tumors from benign nodules using nodule growth statistics. I contribute all statistical analysis, a visualization platform for the data, and figures.

Residual Convolutional Neural Network for Determination of IDH Status in Low- and High-grade Gliomas from MR Imaging. Chang Ken, Harrison X Bai, Hao Zhou, Chang Su, Wenya Linda Bi , Ena Agbodza, Vasileios K. Kavouridis, Joeky T. Senders, Alessandro Boaro, Andrew L. Beers, […], Elizabeth Gerstner, Paul J. Zhang, Bruce Rosen, Li Yang, Raymond Y Huang. Accepted for publication in Clinical Cancer Research (2017).
- Gliomas with the IDH genotype present different visual phenotypes on MR imaging scans. The presence of IDH in glioblastoma is one of the best predictive biomarkers available to the clinician for patient survival, so there is much interest in developing a fast and reliable classifier for IDH prediction in MR images. We present such a method using "2.5D" convolutional neural networks, and show physician-level performance on both high- and low grade tumors derived from clinical scans. Contributed software support for deep learning, writing.

Sequential 3D U-Nets for Biologically-Informed Brain Tumor Segmentation. Beers, Andrew, Ken Chang, James Brown, Emmett Sartor, C. P. Mammen, Elizabeth Gerstner, Bruce Rosen, and Jayashree Kalpathy-Cramer. Accepted for poster presentation at SPIE 2018. https://arxiv.org/abs/1709.02967.
- When treating glioblastoma, a particularly-malignant form of brain cancer, clinicians spend an inefficient amount of time manually demarcating different tissues of interest on MRI scans. To relieve clinicians of this time burden, many have attempted to create fully-automatic 3D tumor segmentation algorithms. I contribute to the field with a deep learning software package that takes into account prior knowledge about the spatial distribution of different tissues of interest. Derivatives of this algorithm are currently in use in clinical trials in my lab. Submitted to the MICCAI BraTS tumor segmentation competition, rankings not yet released.

Anatomical DCE-MRI phantoms generated from glioma patient data. Beers, Andrew, Ken Chang, James Brown, Emmett Sartor, Elizabeth Gerstner, Bruce Rosen, and Jayashree Kalpathy-Cramer. Accepted for presentation at SPIE 2018.
- DCE-MRI is a 4D imaging modality -- 3D images + time -- that radio-oncologists use to extract information about tumor blood-flow and vessel functioning. DCE-MRI data is extremely noisy, and many have developed denoising algorithms that extract blood-flow signal from the noise to create predictive and reproducible cancer biomarkers. In support of our own efforts to de-noise DCE-MRI using deep learning, I create a public dataset of patient-derived noiseless "phantom" data for the purpose of evaluating denoising algorithms. No such phantom exists derived from in-vivo patient data, particularly for the case vessel structure of brain tumors.

Institutionally Distributed Deep Learning Networks. Chang, Ken, Niranjan Balachandar, Carson K. Lam, Darvin Yi, James M. Brown, Andrew Beers, Bruce R. Rosen, Daniel L. Rubin, and Jayashree Kalpathy-Cramer. https://arxiv.org/abs/1709.05929 Under review at *Scientific Reports*.
- Researchers at medical institutions want to use deep learning algorithms on their imaging data, but patient privacy concerns and prohibitions on data-sharing mean that no one institution has enough data to train an effective model. We compare the effectiveness of different methods by which institutions can share model weights, but not their underlying training data, in order to circumvent this need for large, shared datasets. I contribute software support for machine learning aspects, data analysis, and writing.

A Multi-Institutional Comparison of Dynamic Contrast-Enhanced Magnetic Resonance Imaging Parameter Calculations. Ger, Rachel B., Abdallah SR Mohamed, Musaddiq J. Awan, Yao Ding, Kimberly Li, Xenia J. Fave, Andrew L. Beers, […], Brandon Driscoll, Hesham Elhalawani, and David A. Hormuth. Nature Scientific Reports (September 2017).
- DCE-MRI is a 4D imaging modality -- 3D images + time -- wherein a high intensity contrast agent is injected into a patient's blood-stream, and the time-course of that contrast agent is monitored via MRI imaging. Vascular permeability is a biomarker useful for benchmarking brain tumor development, and it can be extracted from DCE-MRI by fitting its time-curves to a standard model. I developed a fitting algorithm for determining BBB permeability that outperformed all other submissions in a multi-institution study on such algorithms.

Repeatability of ktrans derived from DCE-MRI in newly diagnosed glioblastoma across multiple baseline images and processing methods. Andrew Beers, Yi-Fen Yen, Kyrre Eeg Emblem, Elizabeth R Gerstner, Bruce Rosen, and Jayashree Kalpathy-Cramer. Presented at ISMRM (April 2017).

- *Ktrans*, commonly interpreted as blood brain barrier permeability, is a valuable imaging biomarker when treating glioblastoma. I show with with multiple no-treatment baseline scans that its statistical repeatability is lower than previously thought, and very much subject to the particular method by which *ktrans* is calculated.

Multimodal Imaging of Vascularity and Drug Delivery in GBM Patients Treated with Anti-angiogenesis Inhibitor. Yi-Fen Yen, Jayashree Kalpathy-Cramer, Ciprian Catana, Xiao Da, Yangming Ou, Andrew L. Beers, Jacob Hooker, Bruce Rosen, Tracy Batchelor, and Elizabeth R. Gerstner. Presented at ISMRM (April 2017).

- In this abstract, members of our lab attempt to find proxies in MR imaging for chemotherapy drug delivery, in order to assess treatment and guide further research. I contribute data collation and image pre-processing software.

Making sense of large data sets without annotations: analyzing age-related correlations from lung CT scans. Yashin Dicente Cid; Artem Mamonov; Andrew Beers; Armin Thomas; Vassili Kovalev; Jayashree Kalpathy-Cramer; Henning Müller. Proc. SPIE 10138, Medical Imaging 2017: Imaging Informatics for Healthcare, Research, and Applications (March 13, 2017).

- 12,000+ lung CT scans from 9,000+ patients in Belarus' national lung screening program are available, but are bereft of almost any clinically-useful information. We show that one can use random forests to predict patient age from imaging features extracted from automatically-segmented lungs in these scans. I contribute code for machine learning and statistical analysis. Won Best Student Paper award on Medical Informatics track at SPIE.

The Grand Banks Iceberg Mapper. Andrew Beers. Brown Digital Repository. (2015). doi:10.7301/Z0HT2M7B

- Iceberg sightings off the coast of Newfoundland are both valuable and problematic indicators of climate change in the North Atlantic Ocean. I merged iceberg sightings from shipside, aerial, radar, and satellite sighting databases into a single, unprecedented dataset spanning from 1880 to the present day. The Iceberg Mapper uses d3.js to create a interactive web platform for users to investigate these sightings, and better understand the difficulty of reconciling differences in sighting methods between different eras.

Fully-Automatic Low- and High-grade Glioma Volumetric Segmentation in MRI using Deep Learning. Chang, Ken (co-author), Andrew Beers (co-author), [...], Elizabeth Gerstner, Bruce Rosen, Jayashree Kalpathy-Cramer. Submitted at *Radiology*.

DeepNeuro: Open Source, Validated Deep Learning Applications for Neuroimaging. Beers, Andrew, Ken Chang, James Brown, Bruce Rosen, Jayashree Kalpathy-Cramer. In preparation for *Neuroinformatics*.

Open-Source Software

DeepNeuro / qtim_gbmSegmenter - https://github.com/QTIM-Lab/DeepNeuro

- Open-source deep learning Python package for medical imaging. DeepNeuro is an open-source, extensible framework for all of our labs deep learning projects that includes qtim_gbmSegmenter. qtim_gbmSegmenter is a one-off application of a deep learning segmentation model for glioblastoma currently in use in our lab's clinical trials. Sole creator.

Segmentation Wizard (3D Slicer) - https://github.com/QTIM-Lab/SlicerSegmentationWizard

- I contribute a module to 3D-Slicer, an open-source platform for medical imaging software. Includes tools for drawing tumor annotations via intensity thresholding via differences between treatment time points. Used for clinical studies at MGH before being supplanted by DeepNeuro. Sole creator.

glue-viz :: medical - https://github.com/glue-viz/glue-medical

- Glue is a visualization Python package for astronomical data, but nothing in principle prevents it from being a medical imaging visualization tool too. I am working with the creator of Glue to extend glue to medical image data formats -- beta releases available.

qtim_preprocessor - https://github.com/QTIM-Lab/qtim_PreProcessor

- A dockerized a Python-runnable version of our labs internal preprocessing pipeline for brain MRIs. Has been used by our collaborators in Tata Memorial Hospital. Now included in DeepNeuro. Sole creator.

Advocacy

Lucy Parsons Center – Organizing Collective Member (2016-Present)
- Promote events and order books for a progressive community center and bookstore in Jamaica Plain.

Bluestockings Magazine – Art Director (2014-15), Designer (2013-14)
- Designed three 100-page print issues, illustrated 1-3 articles per week, and managed a staff of 10 illustrators for the feminist magazine's website, which reached 10,000+ hits on featured pieces.

Brown/RISD Design for America Chapter – Project Lead (2013-14), Project Member (2012-13)
- Developed the Empathy Box Project, which sought to spread the stories of those with autism via social sharing. Applied for and won grant funding ($1400+).

Brown University Social Action Housing Group – President (2013-14), Head of Publicity (2012-13)
- Organized events and interviewed prospective members in a 45-person residential housing group for those doing social justice work.

Press

"A story in time: Icebergs & Climate Change": Rhode Island NSF Epscor
- http://web.uri.edu/rinsfepscor/2015/06/15/a-story-in-time-icebergs-climate-change/.

"At Loyola HS: 'Empathy boxes' to raise awareness of autism": Angelus News
- https://angelusnews.com/content/at-loyola-hs-empathy-boxes-to-raise-awareness-of-autism

Technical Skills

Programming: Experienced with Python (Tensorflow / Keras / Theano), R (Shiny), Matlab, Javascript (d3.js), C++,. Comfortable with CUDA, Linux, Docker, GPU Programming, ArcGIS, LaTeX, SPSS, Adobe Suite, and VBA.