Benjamin L. Moore

Bioinformatics PhD and data scientist

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Currently

Final year PhD Bioinformatics student at the University of Edinburgh, due to finish around September 2015.

Education

^{2012–15} **University of Edinburgh** Bioinformatics PhD (supervisors: Colin Semple and Stuart Aitken)

^{2011–12} **Imperial College London** MSc Bioinformatics and Theoretical Systems Biology (Distinction)

^{2008–11} **University of York** BSc Biology (First class honours)

Presentations

Tutorials 2014 WikiProject Computational Biology half-day tutorial (ISMB, Boston, USA)

Talks 2015 EdinbR: The Edinburgh R usergroup (various)

²⁰¹⁴ Unravelling higher order chromatin structure (4th Edinburgh Bioinformatics

meeting, Edinburgh, UK)

blogR: pop data analysis and R for the web (Psychology R-users, Edinburgh, UK)

Digital Economies 2014 conference (London, UK; declined invitation)

Posters 2015 Keystone Epigenomics (Keystone CO, USA)

2014 ISMB (Boston MA, USA)

Genome informatics (Cambridge, UK)

Chromatin: From nucleosomes to chromosomes (Cambridge, UK)

²⁰¹³ Genome informatics (CSHL NY, USA)

EpiGeneSys (Cambridge, UK)

Publications

Moore BL et al. Integrative modelling reveals the principles of multi-scale chromatin boundary formation in human nuclear organization. (In review)

Moore BL et al. High–quality protein backbone reconstruction from alpha carbons using Gaussian mixture models. Journal of computational chemisty, 34(22):1881-9.

Gardner PP, Daub J, Tate J, **Moore BL** et al. Rfam: Wikipedia, clans and the "decimal" release. Nucleic acids research, 39(S1):D141-5.

Technical skills

R Python Data visualisation
Statistics Machine learning Mediawiki
LaTeX Bash Cluster computing
Git / Mercurial Emacs HTML and CSS
UNIX Inkscape Javascript

Projects

blog

I author a data science blog at blm.io where I apply R and Python to open datasets and write-up the results. My posts have been featured by sites including FiveThirtyEight, BuzzFeed, AVclub, io9 and more, and also led to being an invited blogger at the Huffington Post. My original wordpress blog (benjaminImpore wordpress com) has received around 150 000 total views from

(benjaminlmoore.wordpress.com) has received around 150,000 total views from 180 countries.

EdinbR

I co-founded a usergroup for the R programming language in Edinburgh called EdinbR. I help to organise our monthly meetings and built our website: edinbr.org. Our meetings attract 30-50 statisticians, data scientists and developers for talks and discussion about the R language and its applications. We're sponsored by Revolution Analytics and have an organisational github account at github.com/EdinbR.

datarea

I entered Imperial College's Summer Data Challenge competition, where entrants analysed a given dataset and then proposed a startup idea based on their results. I developed a modelling technique to select housing areas for investment and was awarded third place: £2,000 and startup support from Imperial Create Lab. My entry can be seen at blm.io/datarea and the code is shared on my github account.

Awards

MRC Capacity building scholarship (fees + enhanced stipend) for my PhD MRC Full postgraduate scholarship (fees + stipend) for MSc ISCB computational biology Wikipedia competition 2013

(doi:10.1371/journal.pcbi.1003242; 2nd place)

Imperial Create Lab's "Summer data challenge" (summerdatachallenge.com; 3rd place)

Individual engagement grant from the Wikimedia Foundation to attend ISMB 2014

Training accreditation for Wikimedia UK, the national chapter of the Wikimedia Foundation

Front cover design selected for J. Comp. Chem. (34:22) (doi:10.1002/jcc.23388)

Placements

^{2010–11} **Wellcome Trust Sanger Institute** (Cambridge, UK) Worked for two summers on the Rfam database (rfam.xfam.org) in the Bateman group

Links

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References

Available on request.