

# 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 | <b>University of Edinburgh</b> Bioinformatics PhD (supervisors: Colin Semple and Stuart Aitken)                   |
| 2011–12 | <b>Imperial College London</b> MSc Bioinformatics and Theoretical Systems Biology ( <a href="#">Distinction</a> ) |
| 2008–11 | <b>University of York</b> BSc Biology ( <a href="#">First class honours</a> )                                     |

## Presentations

|           |      |  |
|-----------|------|--|
| Tutorials | 2014 | WikiProject Computational Biology half-day tutorial ( <a href="#">ISMB</a> , Boston, USA)  |
| Talks     | 2015 | EdinbR: The Edinburgh R usergroup ( <a href="#">various</a> )  |
|           | 2014 | Unravelling higher order chromatin structure ( <a href="#">4th Edinburgh Bioinformatics meeting</a> , Edinburgh, UK)<br>blogR: pop data analysis and R for the web ( <a href="#">Psychology R-users</a> , Edinburgh, UK) |
| Posters   | 2015 | Unravelling higher order chromatin organisation ( <a href="#">Keystone Epigenomics</a> , Colorado, USA)  |
|           | 2013 | Integrative modelling of higher order chromatin structure ( <a href="#">Genome informatics</a> , CSHL, NY)<br>Unravelling higher order chromatin structure ( <a href="#">EpiGeneSys</a> , Cambridge, UK)                 |

## Publications

|      |   |
|------|---|
| 2015 | <b>Moore BL</b> et al. Integrative modelling reveals the principles of multi-scale chromatin boundary formation in human nuclear organization. ( <a href="#">In review</a> )              |
| 2013 | <b>Moore BL</b> et al. High-quality protein backbone reconstruction from alpha carbons using Gaussian mixture models. <a href="#">Journal of computational chemistry</a> , 34(22):1881-9. |
| 2011 | Gardner PP, Daub J, Tate J, <b>Moore BL</b> et al. Rfam: Wikipedia, clans and the “decimal” release. <a href="#">Nucleic acids research</a> , 39(S1):D141-5.                              |

## Technical skills

R  
Statistics  
LaTeX  
Git / Mercurial  
UNIX  
Python  
Machine learning  
Bash  
Emacs  
Inkscape  
Data visualisation  
Mediawiki  
Grid computing (SGE / PBS)  
HTML and CSS  
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, |
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and also led to being an invited blogger at the Huffington Post. My original wordpress blog (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](http://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](https://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](http://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  
Individual engagement grant from the Wikimedia Foundation to attend ISMB 2014  
ISCB computational biology Wikipedia competition: second place prizewinner  
Summer data challenge: third place prizewinner

## Placements

2010–11 **Wellcome Trust Sanger Institute** ([Cambridge, UK](#)) Worked for two summers on the Rfam database ([rfam.xfam.org](http://rfam.xfam.org)) in the Bateman group

## Links

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## References

Available on request.