

GRANT NEILSON

Bioinformatician with four years of industry experience analyzing NGS datatypes with a specialism in WGS. An expert in programming with R and Python, have extensively used both languages for analysis of high dimensional omics data as well as developing packages for both languages. A named contributor on the NF- Core Sarek pipeline. Experience working with AWS services such as EC2, S3 and ECR. Have extensively used version control software i.e., Gitlab and Docker to containerize analysis. An articulate and reliable individual with strong communication skills. This is demonstrated by my achievements and accomplishments to date.

EDUCATION

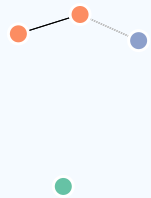
- 2019
|
2018
- **MSc Biology**
University of Southampton
N/A
 - **Undergraduate Researcher**
University of Southampton
N/A

INDUSTRY EXPERIENCE

- 2025
|
2020
- **Cosyne Therapeutics**
London
N/A

SELECTED PUBLICATIONS, POSTERS, AND TALKS

- 2015
- **Asymmetric Linkage Disequilibrium: Tools for Dissecting Multiallelic LD**
Journal of Human Immunology
• Authored with Richard Single, Vanja Paunic, Mark Albrecht, and Martin Maiers.

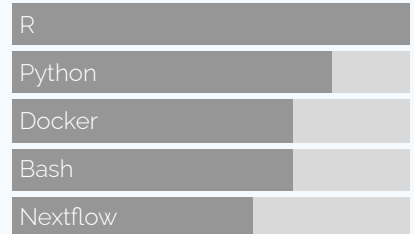


View this CV online with links at
grantneilson.me/cv/

CONTACT

✉ grant.neilson5@gmail.com
🐙 [github](#)
in [LinkedIn](#)

LANGUAGE SKILLS



Made with the R package [pagedown](#).

The source code is available on
github.com/nstrayer/cv.

Last updated on 2025-06-19.