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### DABS (DATA AND BIOINFORMATION STUFF)

Posted on January 8, 2021 - [<https://www2.hshsl.umaryland.edu/hslupdates/?p=4555>] by J.P. Courneya - [<https://www2.hshsl.umaryland.edu/hslupdates/?author=16>]

The Center for Data and Bioinformation Services (CDABS) is the University of Maryland Health Sciences and Human Services Library hub for, data and bioinformation learning, services, resources, and communication.

This is the first issue of our blog. It contains links worth your time from all over the web focused on what we do best – data and bioinformation. We will provide links to tutorials, training opportunities, software packages, interviews with experts, data sources and more!

1. Wired magazine interview with Dr. Eric Green, head of the National Human Genomics Research Institute (NHGRI), about the future of Genomics. (8 minute read): <https://www.wired.com/story/30-years-since-the-human-genome-project-began-whats-next/> - [<https://www.wired.com/story/30-years-since-the-human-genome-project-began-whats-next/>]
2. The Missing Semester of Your CS Education. Learn how to master the command-line, use a powerful text editor, use fancy features of version control systems, and much more! (11 lectures + 11 Hours YouTube course): <https://missing.csail.mit.edu/> - [<https://missing.csail.mit.edu/>]
3. Propel Postdoctoral Scholars Program new @Stanford School of Medicine aims to 1) attract and train talented postdocs who would bring diversity (broadly defined) to both their training program and to the professoriate in the life sciences, including from backgrounds or perspectives typically underrepresented in the life sciences, 2) provide postdoctoral funding support and professional development programs that will prepare individuals to assume leadership roles as faculty in academia, and, in so doing, 3) champion a culture of belonging and inclusion in science and academia, within and beyond Stanford. The application deadline is February 8, 2021. (10 minute read): <https://propel.stanford.edu/> - [<https://propel.stanford.edu/>]
4. Retrieve genome data by BioProject using the `datasets` and `dataformat` command-line tools by providing a BioProject accession. (5 minute read): <https://www.ncbi.nlm.nih.gov/datasets/docs/command-line-start/> - [<https://www.ncbi.nlm.nih.gov/datasets/docs/command-line-start/>]

1/26/2021

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