

Progress Report: Genome-wide hypothesis generation for single-cell expression via latent spaces of deep neural networks

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Abstract

We wrote an application for the [Chan Zuckerberg Initiative's Collaborative Computational Tools RFA](#). Our application was recommended for funding. We are writing our progress report as we go. This repository contains the report. Please feel free to file a [GitHub Issue](#) to ask a question. Some elements of this report are expected to also be written up via a published manuscript. In the event that we write a manuscript, we will begin from this report. Authorship will be determined in accordance with [ICMJE guidelines](#).

References
