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