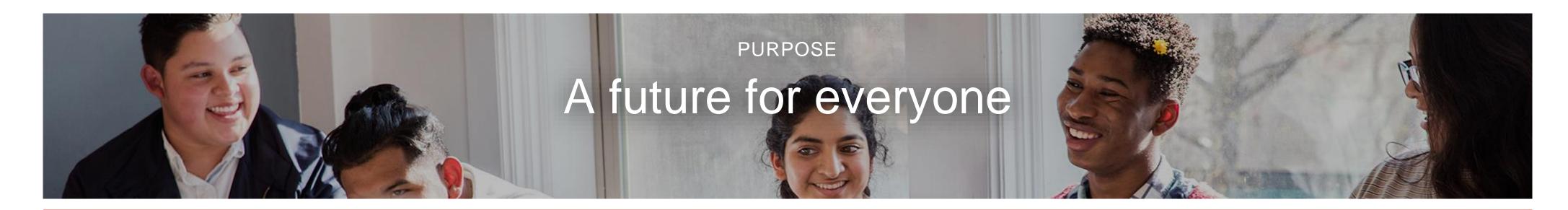
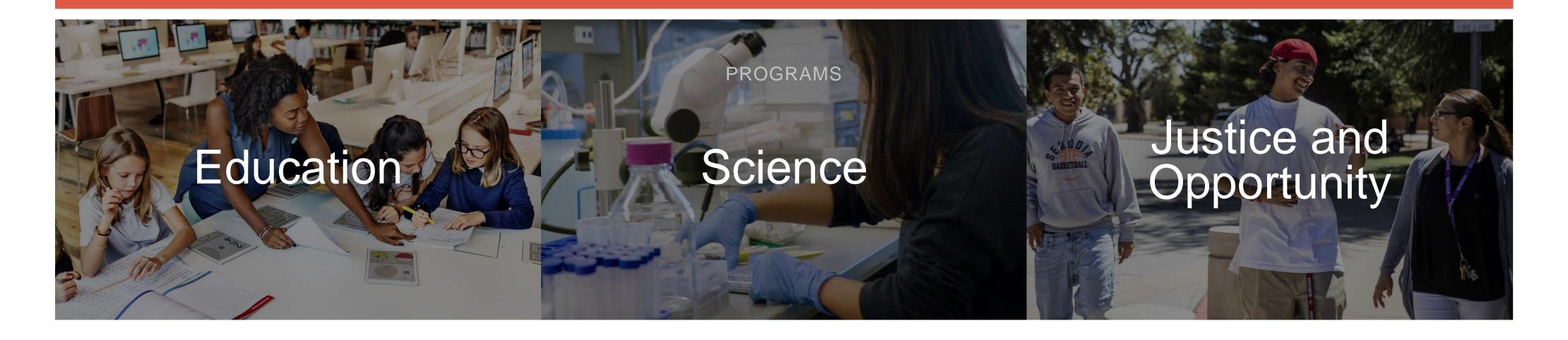


CHAN ZUCKERBERG INITIATIVE



APPROACH Change at scale





CHAN ZUCKERBERG SCIENCE

Supporting science and technology that will make it possible to cure, prevent, or manage all diseases by the end of the century

83 years

Accelerating biomedical science by developing new tools and technologies and supporting open, collaborative models of research

10 years



ACCELERATING BIOMEDICAL RESEARCH



Collaboration by experimental scientists, physicians, theorists, and engineers



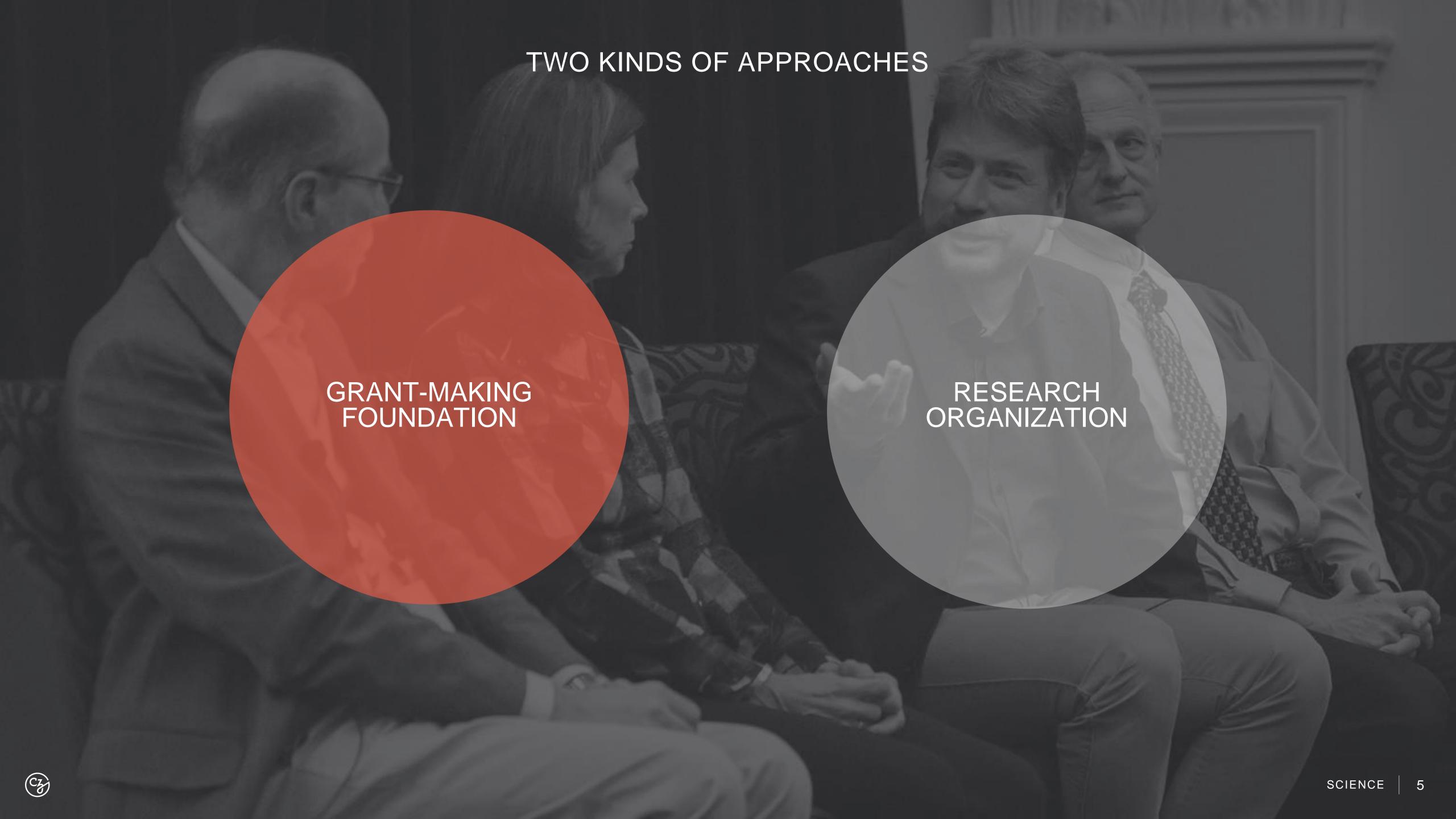
Enabling tools and technologies



Building support for science

People – Technology – Collaboration – Open Science





OUR APPROACH

Richard Sever

WE FUND

Grants and Partners

External institutions worldwide

GRANT-MAKING FOUNDATION

RESEARCH ORGANIZATION

WE BUILD

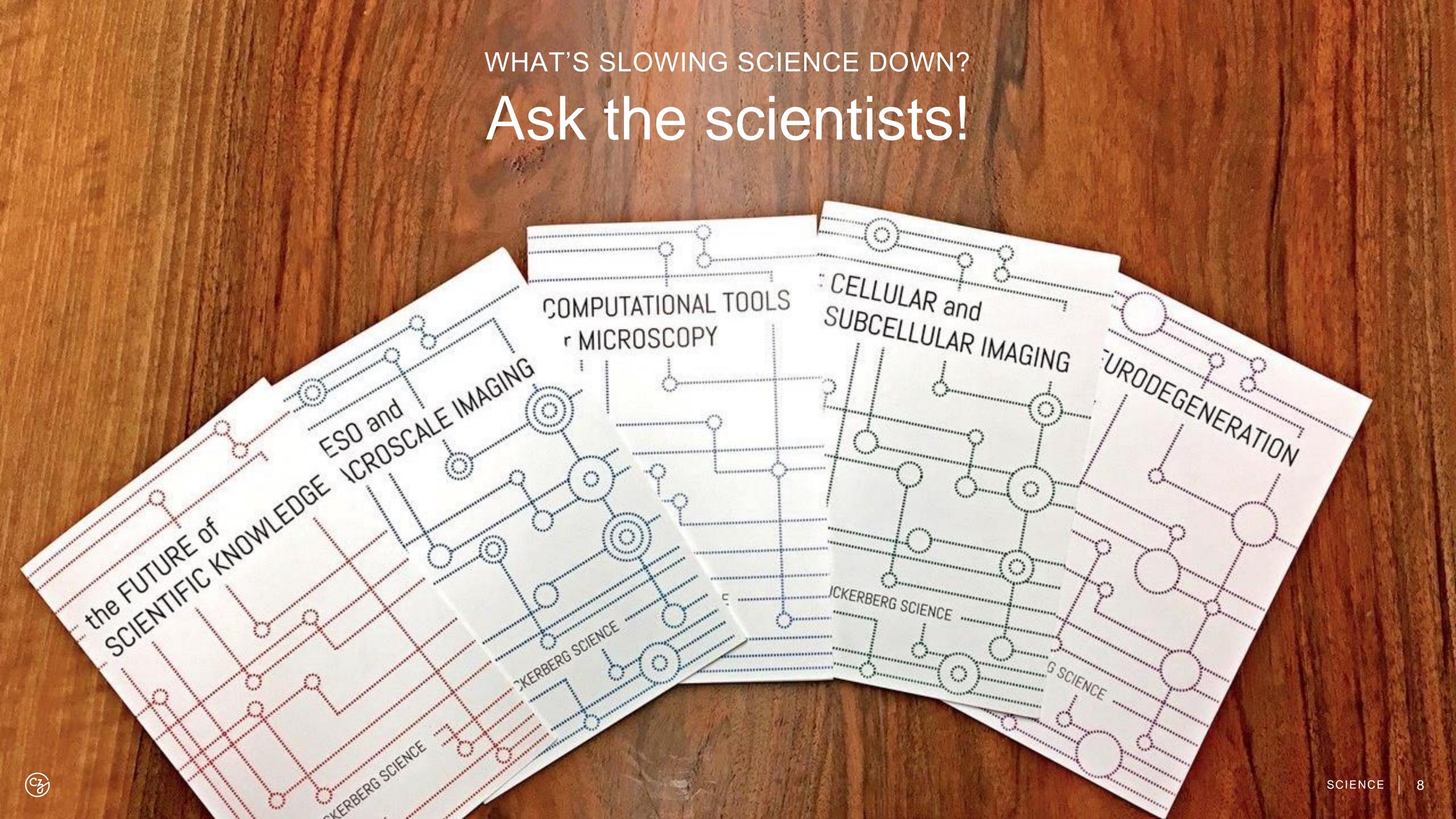
Open-source software, tools

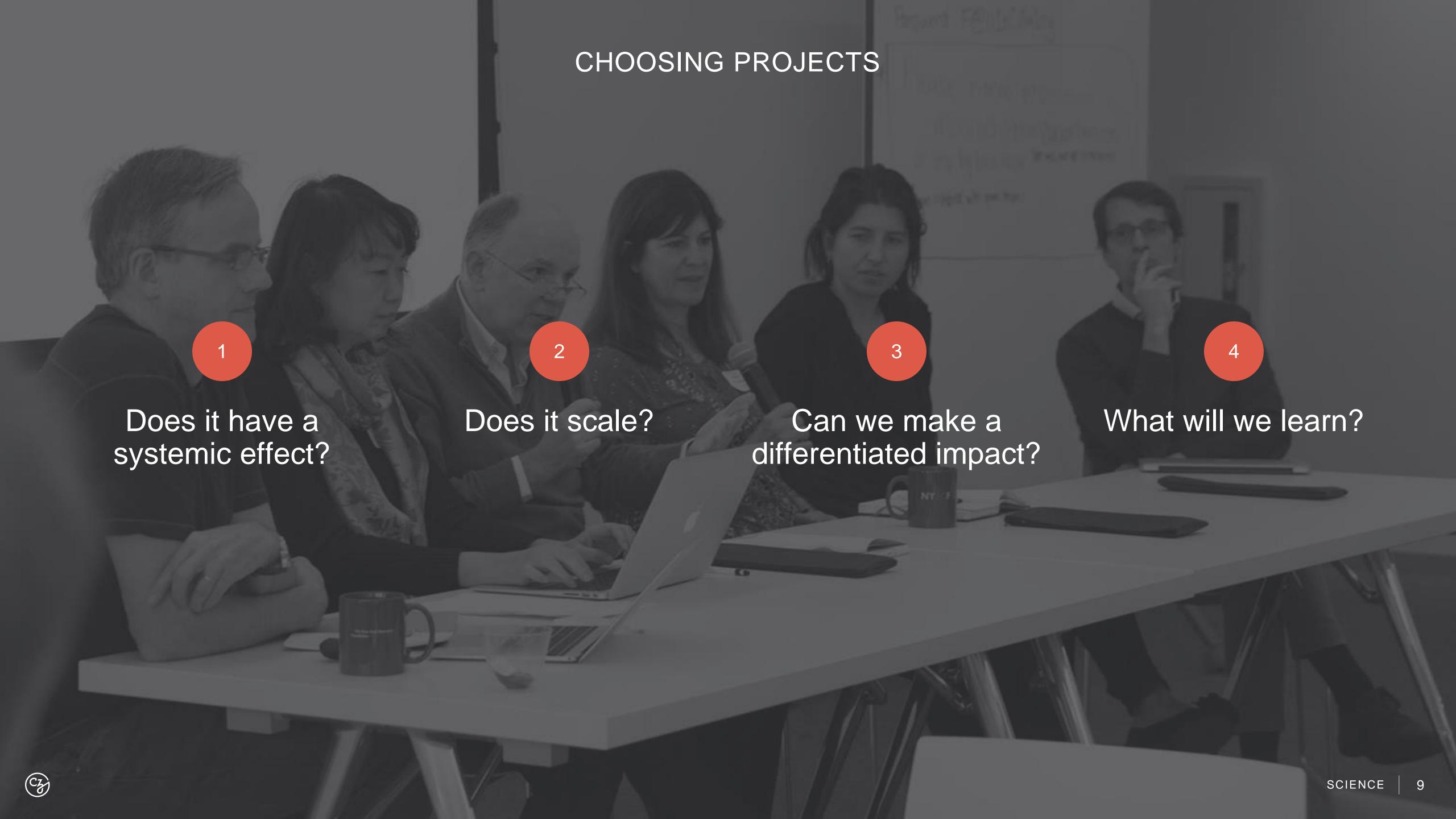
CZI engineers and computational biologists



how we choose projects









Tools that are ROBUST, RELIABLE, SCALABLE, and SHAREABLE



ACCELERATING KNOWLEDGE SHARING AND DISCOVERY



PREPRINTS FOR BIOLOGY

AI/ML FOR THE BIOMEDICAL LITERATURE





A free, open reference map of all cells in the healthy human body

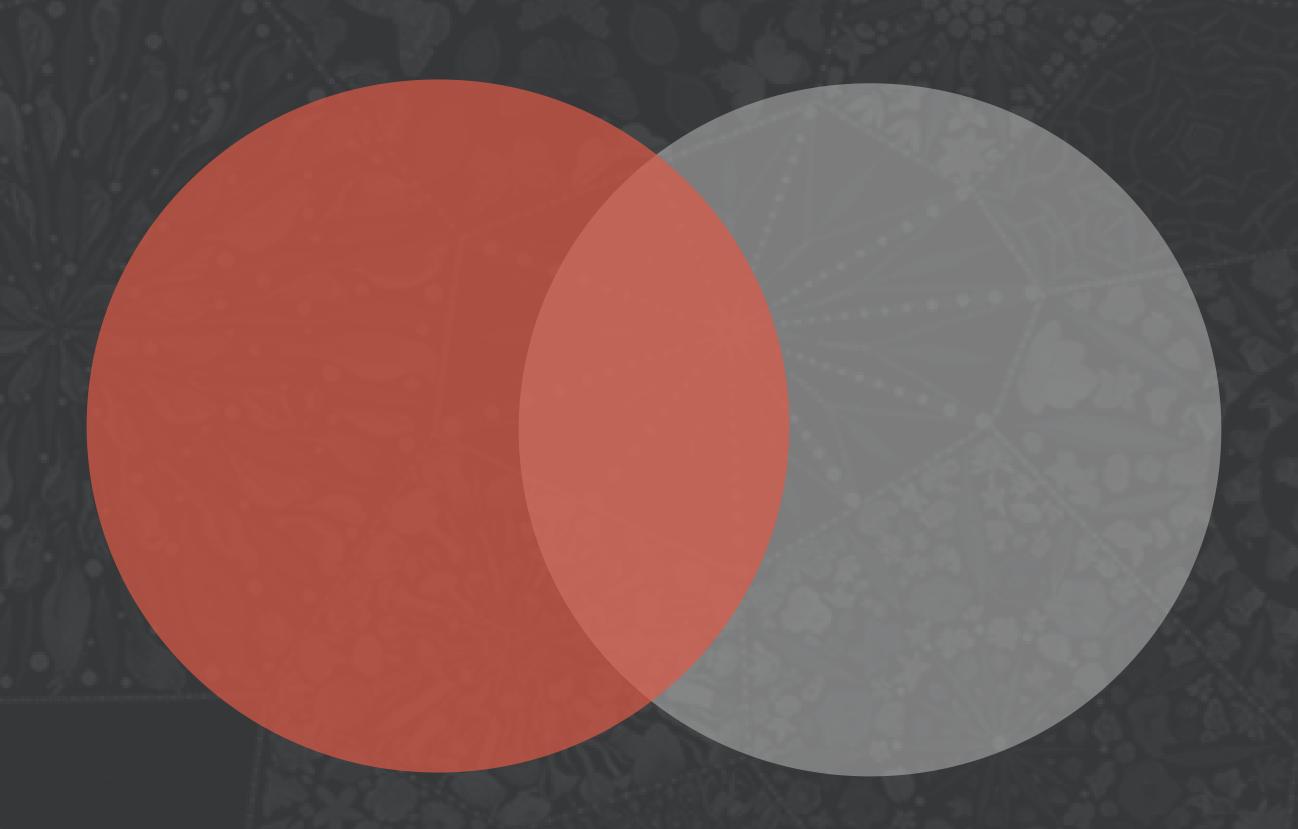
TYPES, NUMBERS, LOCATIONS, NEIGHBORS, MOLECULAR COMPOSITION





WE FUND

Researchers worldwide



WE (HELP) BUILD

Data coordination platform

ALSO ENGAGED

Wellcome, NIH, Klarman Foundation, EU, DFG, RIKEN, Wallenberg Foundation



WE FUND: OPEN REQUESTS FOR APPLICATIONS

REQUEST FOR APPLICATIONS

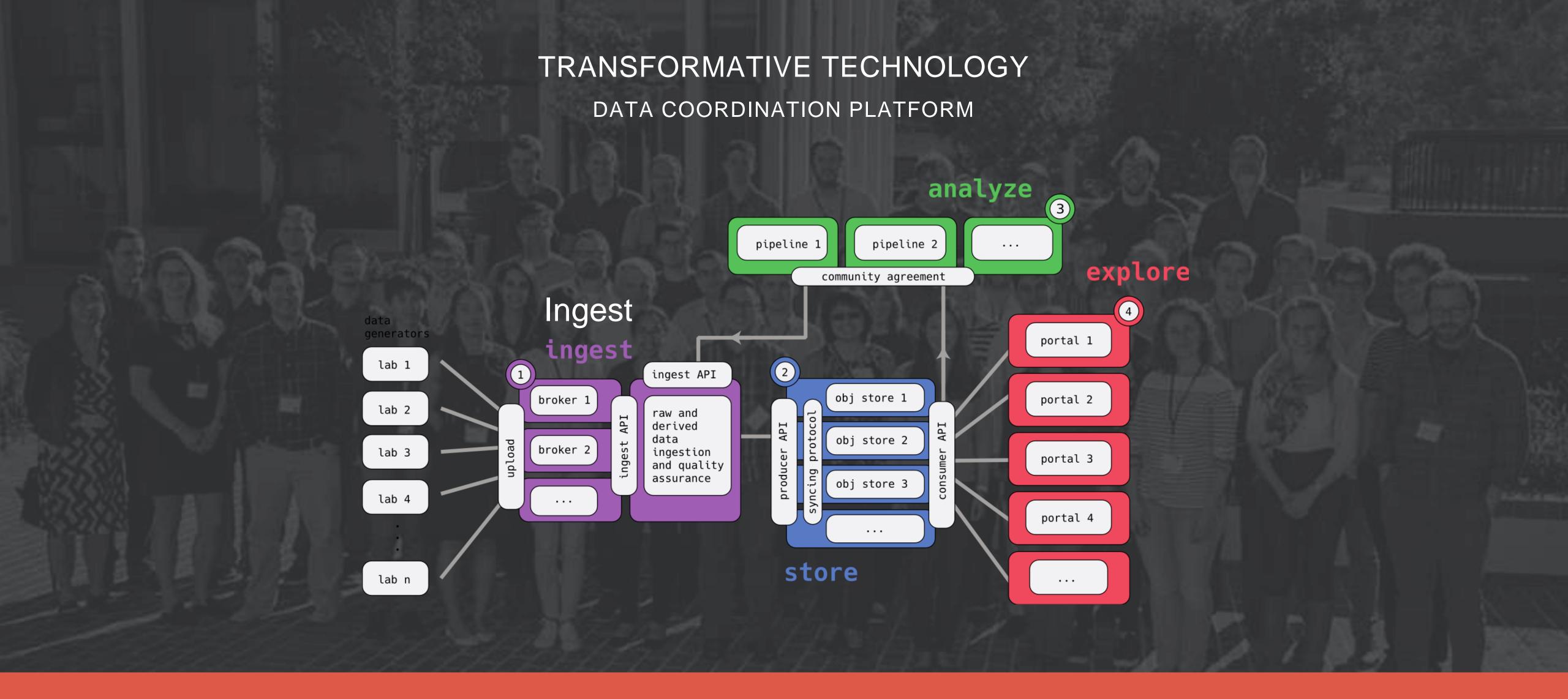
Pilot Projects for a Human Cell Atlas

The Human Cell Atlas is a global effort to create a reference map of all cells in the healthy human body as a resource for studies of health and disease.

The Chan Zuckerberg Initiative invites applications for one year pilot projects to develop technologies for the Human Cell Atlas, establish best practices in the field, and begin a common data archive for analysis and investigation. Full details here.

Collaborative computational tools for the Human Cell Atlas supporting open, collaborative development of new computational tools, algorithms, visualizations, and benchmark datasets

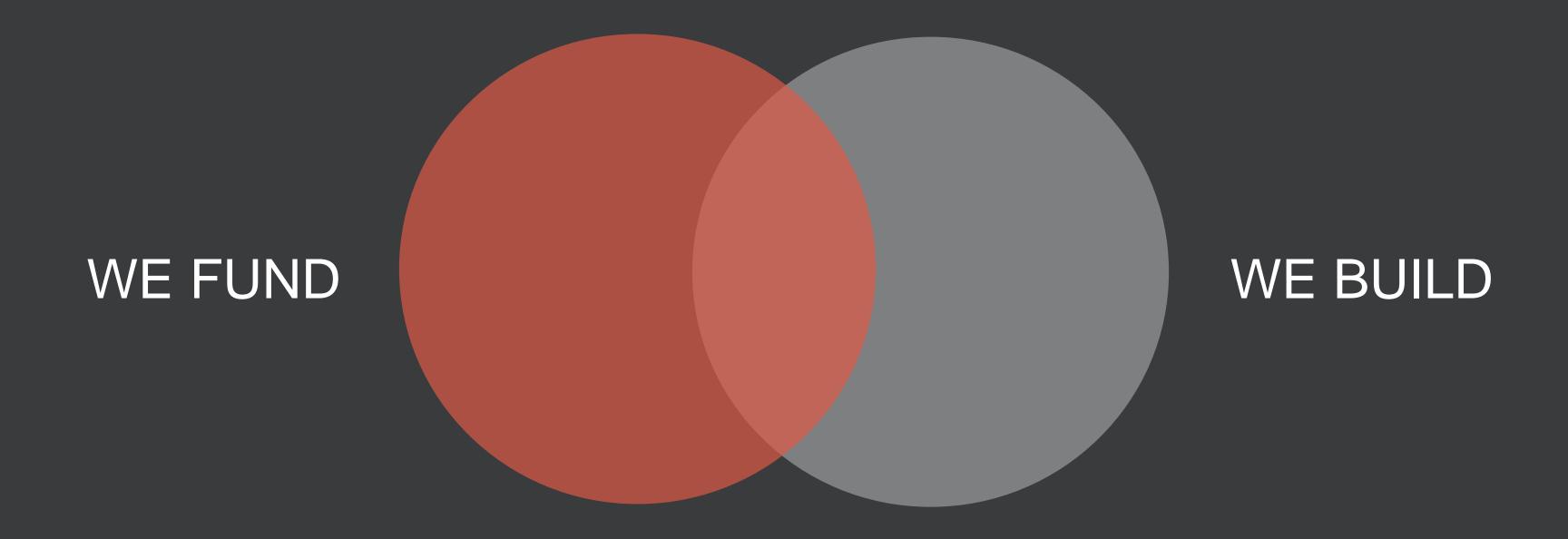




DEVELOPED JOINTLY BY

European Bioinformatics Institute, Broad Institute, UC Santa Cruz and, CZI





To stay updated about future opportunities

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