Q&A: David Altshuler on how to share millions of human genomes

The following is an edited excerpt.

To fully understand the role of human genetic variation and its role in disease, researchers need to pool DNA and clinical data from millions of people.

Earlier this week, more than 70 research, health care, and patient advocacy organizations, including big players such as the U.S. National Institutes of Health and the United Kingdom's Wellcome Trust Sanger Institute, announced a plan to do just that.

ScienceInsider spoke with geneticist David Altshuler of the Broad Institute in Cambridge, Massachusetts, who has led planning for the alliance, about its aims.

Read the full story here: Q&A: David Altshuler on How to Share Millions of Human Genomes