LIVE ONLINE EVENT - JANUARY 13 @ 2:30 pm ET
RWJF New Recommendations to Build a Healthier America

Building on its original work from 2009-which helped advance a national movement to address non-medical factors that affect our health - the RWJF Commission to Build a Healthier America will release new recommendations.

During the live event featuring Commission Co-Chairs Mark McClellan and Alice Rivlin, Commissioners will offer recommendations covering three key areas:

- Prioritizing investments in America’s youngest children
- Encouraging leaders in different sectors to work together to create communities where healthy decisions are possible
- Challenging health professionals and health care institutions to expand their focus from treating illness to helping people live healthy lives

Connect to LIVE ONLINE EVENT

SCIENTIFIC PUBLICATIONS

Public Health Reports
Applying Social Determinants of Health to Public Health Practice
**FUNDING OPPORTUNITY**

The NIH Common Fund’s Epigenomics program has released a new Funding Opportunity Announcement (FOA): **Computational Analyses Exploiting Reference Epigenomic Maps (RFA-RM-14-001)**. The goal of this FOA is to encourage community use of the reference epigenomic maps that have been developed as part of the Epigenomics program. We are particularly interested in applications that integrate information from reference epigenomic maps with other data types, either existing public datasets or investigator-generated data such as GWAS/genetic studies, eQTLs, etc.

Examples of potential projects might include (but are not limited to):

- Integrative analyses that combine reference epigenomic maps with other public (e.g. ENCODE, TCGA, IHEC, Brainspan, etc.) or investigator-generated data sets.
- Analyses that use reference epigenomic maps to aid in interpretation of the results of GWAS or other genetic studies.
- Analyses that use reference epigenomic data to identify specific features that distinguish cell types.
- Integration of reference epigenome maps with other -omic data sets.
- Cross-species comparative epigenomic analyses using human reference epigenome maps.
Approaches that exploit epigenomic mapping data along with other data sets to identify novel biological pathways, biomarkers, or therapeutic targets

Important information about this FOA:

- Uses the R01 mechanism
- Limited to $200K direct costs
- Limited to two years
- Letters of Intent due February 3, 2014

Receipt date March 3, 2014

The NIH Common Fund supports goal-driven, research networks in which investigators generate data, solve technological problems, and/or otherwise pilot resources and tools that will be stimulatory to the broader research community. Common Fund programs are designed to achieve their goals within a maximum of 10 years. More information about the programs currently supported by the Common Fund can be found at www.commonfund.nih.gov

RESOURCES

Chart Book: The War on Poverty at 50 Section 1: More Complete Poverty and Income Measures Show Progress Over the Last 50 Years

Visit the Mid-South TCC website (Resources tab) to find a list of Scientific Publications on Social Determinants of Health
www.uab.edu/midsouthtcc/resources

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