

Efficient methods of collecting
phenotype and exposure data
and
Research in clinical care systems
or small clinical practices

Efficient methods of collecting phenotype and exposure data

- Centralized databases and standard data structure
- Developing ontologies and consistent methods for capturing phenotypes—Gene and Environment Initiative
 - OBO / NCBO phenotype effort
 - SNoMed/Disease Ontology discussions
 - Lifestyle ontology??
- increasing implementation of electronic health records
- opportunity for feedback to EHRs to improve quality and consistency of data capture

Efficient methods of collecting phenotype and exposure data

- social economic status
 - Privacy issue
- environmental exposures
 - Air Quality System
(<http://www.epa.gov/ttn/airs/airsaqs/index.htm>)
 - Data still limited
- Geographic Information Systems (GIS)
 - EPA Environmental Geospatial Access Project
(http://www.epa.gov/enviro/geo_data.html)

Research in clinical care systems or small clinical practices

- Traditional Epidemiological Cohorts vs. Clinical Care Systems
 - completeness limited data vs incomplete broad data
 - enrollment bias
 - efficiency
- Clinic-hospital based cohorts
 - Clinical and Translational Science Award network
- Network of EMR-linked Biobanks
 - Mayo, Vanderbilt, Marshfield Clinic, Group Health Seattle, Northwestern U.

Research in clinical care systems or small clinical practices

- Practice Based Research Networks
 - improvement of primary care
 - expanding into quality assessment
 - applications for genomics research
 - Agency for Healthcare Research and Quality (AHRQ)
 - <http://www.ahrq.gov/research/pbrn/pbrnfact.htm>
 - <http://www.aafp.org/online/en/home/clinical/research/fpbrn.html>
 - <http://jama.ama-assn.org/cgi/content/full/297/4/403>
 - <http://www.ajhp.org/cgi/content/abstract/64/19/2044>

Goals for the Future

- Should we maximize use of Electronic Health Records for phenotype development and how?
- How to support ontology/vocabulary development for phenotypes and software and data mining tools for electronic medical records?
- Should we develop “best practices” repository for phenotype datasets?
- How do we integrate with CTSA network to maximize data harmonization?

Goals for the Future

- How do we support integration of phenotypes with environmental data from EPA and other databases such as Census track data?
- Should we maximize use of practice based research networks for cohort development, and if so how?
- How can we develop and test methods for sample collection and management across PBRNs?