

**Multi-IC Symposium on Application of Genomic Technologies to
Population-Based Studies**

June 5-6, 2006

Bethesda Marriott

SYMPOSIUM GOALS AND OUTPUT

Symposium Goals:

1. To identify common, critical issues that have been encountered in applying genomic technologies to population studies at NIH and creative approaches to solving them;
2. To develop approaches for prioritizing and conducting population studies using genomic technologies for use by individual ICs as desired; and
3. To identify new tools for genomics, categorization of phenotypes, and database standardization required for genome-wide association and sequence-based studies.

Symposium Output: Series of recommendations to the various ICs for issues to consider in soliciting, funding, and implementing these studies, as well as recommendations for new genomic tools needed and approaches for facilitating their use.