Next Steps
Next steps: Genotyping

- Add representatives from each study to the Genotyping Group – establish the plan for QA for each study (trios?)
- Nail down the schedule of DNA shipping and genotype production for all six studies
- Get genotypes from both platforms for HapMap samples
- Special genotyping topics: MHC, mitochondria, CNVs
- Establish the plan for data quality checking (NCBI + Study PI) for each study – should include tracking any correlation of success rate with MAF
- Post-data release: ideally, PIs need to be prepared to check the most significant results on an independent platform
- Decide how to handle misattributed paternity in trios
Next steps: Consortium Function

- Assemble a complete set of all slides shown at this meeting
- Distribute detailed summary of quality standards adopted by GAIN
- Form an Analysis Working Group
- Other Working Groups?
- Each study PI needs to initiate steps to get approval from the Data Access Committee – GAIN staff to provide background to help IRBs interpret this need
- Study PIs are encouraged to assemble their crack analytic teams, and to plan for replication now
- Conference call early January, and then monthly
- Meeting July 18 – 19, 2007
Next steps: Communication About GAIN to Users

- Clear description of each study, including usage limitations based on consent
- Post OHRP written decision
- Display names and project titles of GAIN data users?
Thanks for a GREAT meeting!

And have a safe trip home

The fog has lifted – both literally and figuratively

And the lights are on!