Overview of Colon CFR Resource

1997 - 2007 2007 - 2012

PROGRESS TO DATE (Phases I & II)

RESEARCH PLAN

Supporting

PROPOSED CORE ACTIVITIES

Case ascertainment/recruitment

 Phase-I
 Phase-II
 Phase-III

 1998-2002
 2002-2007
 2007-2012

Population-based, no sampling Population-based, early onset

Population-based, positive family history

Population-based, minority

Clinic-based

Enrollment	Pop-based	Clinic-based
Probands	7,252	1,117
Family members	15,920	5,2209
Controls (P-B & spouse)	5,091	

Data Collection

Family history: 12,178 Epi/risk factors: 28,263 Follow-up: 15,711 Pathology records: 6,012 Fresh frozen tissue: 272 EBV lines: 4,100

Tumor Phenotyping

MSI 4,000 IHC 4,000 Methylation of *MLH1* 2,469

Germline Testing for MMR (MLH1, MSH2, MSH6)

DNA sequence variance testing (via dHPLC) 1,747
Deletions & duplications (via MLPA) 1,747

Multidisciplinary Research-ongoing

- Germline MMR testing (MLH1, MSH2, MSH6, PMS2)
- Tumor methylation (*MLH1*)
- Mapping of new susceptibility genes
 - Genetic Linkage in CRC Families
 - Fine mapping/positional cloning
 - TGFB
- Etiologic pathways (candidate gene)
 - NSAIDs & COX/PG Metabolism
 - Folate, vitamin D, and Calcium
 - Metabolic Syndrome, Obesity, and IGF
- MYH association studies
- Research on Minority Populations
- Intervention study to promote colonoscopy screening
- Numerous studies under grant review
- Numerous studies initiated by outside investigators

Etiologic Research (incl. proposals under review)

- MMR Gene Research
- Base excision repair
- Mapping new CRC genes, e.g.:
 - Syndrome X Families
 - LD Mapping in Hispanic Families
 - Whole Genome Association study
- Candidate gene studies, e.g.:
 - Alpha-1-antitrypsin
 - Lipid peroxidation
 - HCAs, PAHs, and Nitrosamines
- Genome-wide association studies
- Epigenetic epidemiology, e.g.:
 - CIMP
 - BRAF
- Incorporating emerging technology (e.g., CGH)
- Incorporating methodological research

Clinical Research

- MSI & prognosis
- HNPCC & heterogeneity of cancer risks
- Pathology and clinical research
- Pharmacogenetics

Behavioral Research

- Screening & CRC risk factor knowledge gaps
- Education and counselilng (including newsletters)

Prevention & Screening

- NSAIDS
- Vitamin D with/without Calcium
- Other screening studies

Collaborations with Other Initiatives

- EDRN
- NHGRI Medical Sequencing Program
- The Cancer Genomic Atlas (TCGA)
- CAPP2

Phase III Recruitment Core

- Population-based cases <50 yrs or + family history
- Clinic-based cases
- Multiple-case Newfoundland families
- Minority recruitment
- CRC-affected first-degree relatives (FDR)
- FDR of mutation positive probands
- FDR of Lynch Syndrome & Amsterdam II probands.
- FDRs of Syndrome X probands

Retention & Follow-up Core (current & proposed) Active Follow-up Data Collection

- Family cancer history
- Personal health history and CRC screening
- Epidemiology/risk factors data collection
- Quality of Life (QOL)

Passive Follow-up Data Collection

- Vital status and address updates
- Newsletters, other participant information

Biospecimen Core (current & proposed, proposed)

- Biospecimen (blood/buccal, tissue) collection
- DNA extraction (from blood/buccal & tissue)
- Tracking and reporting
- Quality Assurance and Quality Control
- Central repository and dispatch
- EBV transformation &/or WGA
- Tissue Micro Arrays (TMA)
- Virtual tissue repository

Mol. Char. Core (current & proposed, proposed)

- IHC testing
- MMR testing (*DNA sequencing*, MLPA)
- MYH mutation testing
- BRAF mutation testing
- MLH1 methylation
- New gene testing

Clinical Outcomes Core (proposed)

- Baseline diagnosis & treatment data collection
- Treatment & outcomes data collection
- Recurrences & new primaries

Bioinformatics & Biostatistics Core

Administration Core