Barrett's Esophagus Translational Research Network (BETRNet) Renewal Concept

Division of Cancer Biology

Division of Cancer Prevention

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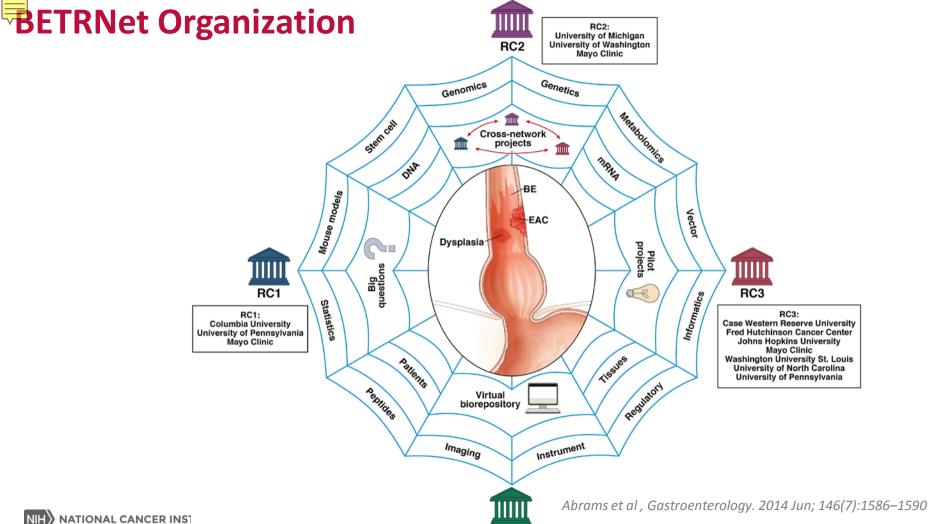
Barrett's Esophagus Translational Research Network (BETRNet)

- DCB/DCP multi-disciplinary and multi-institutional program
- Collaboration among multiple centers, with basic and clinical researchers and resource sharing, to tackle a rare cancer
- Significant oversight and active participation by NCI staff allows real-time modifications to the program based on scientific needs and accomplishments
- Enable a deeper understanding of EA biology and its putative precursor lesion,
 Barrett's esophagus (BE), provide strategies for EA prevention, and better define individuals at risk
- Funded by NCI in 2011; 3 Research Centers and 1 Coordinating Center

Esophageal Adenocarcinoma

- Highly morbid and highly lethal cancer
- Unexplained dramatic rise in incidence and mortality
- BE confers increased risk
 - ~ 11-30 fold greater than average risk
 BUT
 - 80-90% of EA cases have no prior diagnosis of BE





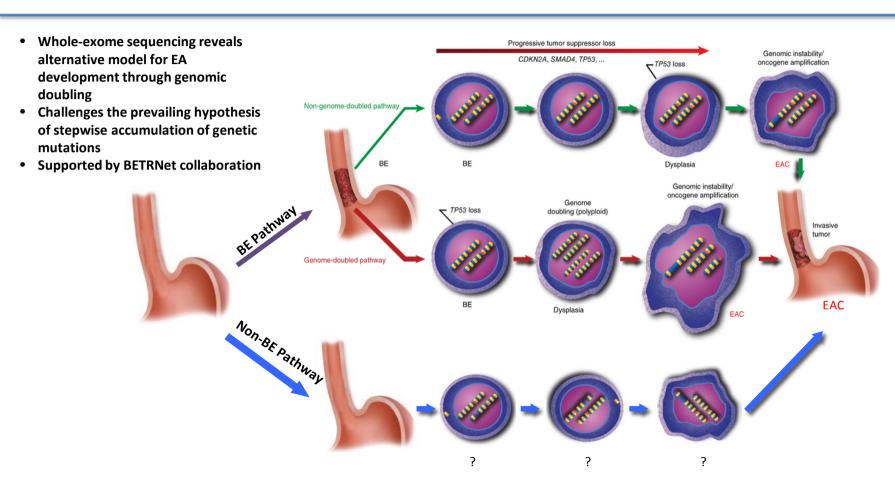
From Discovery to Validation Through Team Science

- First transgenic (L2-IL-1β) mouse model of Barrett's-like metaplasia with multiple tumors at 12 months - developed by RC1
 - BE/EA may arise from a stem/progenitor cell in the gastric cardia; this model challenges the dominant thinking in the field about the origin of BE/EA
 - BE gene signature more closely resembles cardia than squamous epithelium
- Accelerated validation of the mouse model through the consortium
 - Testing of chemopreventives (e.g., statins, obeticholic acid) in RC2
 - Validation of imaging markers for high grade dysplasia and EA (e.g., EGFR, Transglutaminase 2) in RC3

From Discovery to Validation Through Team Science (cont'd.)

- Protective role of GSTT2 gene in African American esophagus
 - Aimed to explain the increased prevalence of EA in Caucasians versus African Americans (ratio 4-6:1)
 - GSTT2 enzyme has a protective role against oxidative damage; however, in Caucasian GSTT2 is less expressed
- Study made possible by combining rare esophageal cancer specimens from multiple BETRNet institutions
- Study completed and manuscript in preparation

From Discovery to Validation Through Team Science (cont'd.)



BETRNet Coordinated Outreach

- Pilot projects with flexible funds to exploit emerging ideas advertised to outside investigators
- Cross-BETRNet focused sessions at national and international meetings included outside investigators
- Leading experts invited to BETRNet Steering Committee meetings
- Publications in major journals

Patient Registry-Virtual Biorepository

- Access to specimens available through collaboration with BETRNet
- Review of proposals by an access committee
- 3901 unique specimens available as of November 2015
- Current revision to the PR-VR "public face" for easy access by the larger scientific community



BETRNet Program Evaluation

(Performed During Year 4)

- RFA provided the necessary infrastructure to break down silos and support collaborative research to synergize science and accelerate opportunities for translational studies
- BE could be a model for understanding the development of other adenocarcinomas with unknown or inaccessible precursors
- Noted Significant Accomplishments
 - EGFR probe with great potential for clinical use
 - Epigenetics and exome sequencing- an important foundation for future application
 - Registry-tissue repository exceptionally valuable for a wide range of research studies
 - Cross-Center collaborations
 - Interactions with other NCI-supported entities such as the EDRN and CISNET

BETRNet Program Renewal (RFA Reissuance)

Research Centers (U54)

Multi-institutional, trans-disciplinary research groups to support:

- Well-defined and NCI coordinated platform to advance research on a rare cancer
- Collaborative accrual of sufficient cohorts/specimens that could not be collected at a single institution
- Collaborative approach to new research directions and expertise sharing to permit adequate hypothesis testing (lesson learned)

Coordinating Center (U24)

Administration of the consortium and continued expansion of the virtual biorepository

NCI Staff

- Active role in program organization, management, and performance evaluation to maximize integration and coordination of the complex U54 research structure with diverse scientific areas
- Partnership with BETRNet investigators to prioritize and support research and outreach activities

Research Questions - New RFA

Cell of Origin

- Biology and development of esophageal stem cells and whether they are slow dividing cells or proliferative progenitors
- Effects of acid and inflammation on esophageal stem cells and their modulation by preventive agents
- Role of esophageal or other stem cells in the development of EA

Genomics of EA

- Impacts of spatial distribution of genetic alterations in BE on the evolution of EA
- The genomics of tumor heterogeneity and tumor plasticity and their impacts on the progression of EA
- The genomics of obesity and their effects on the risk of EA to inform molecular targeting for chemoprevention

Research Questions (continued)

Microbiota

 Alterations in BE and EA microbiota, their role in BE-EA transition and potential modulation for cancer prevention

Technologies and Models

- New technology for minimally invasive screening and surveillance
- Models such as engineered mice, PDX models, cell lines, and organoids that could provide a platform for understanding human esophageal epithelial transformation

Proposed Budget

Set aside request: \$5.5 million in total costs (TC) per year for 5 years

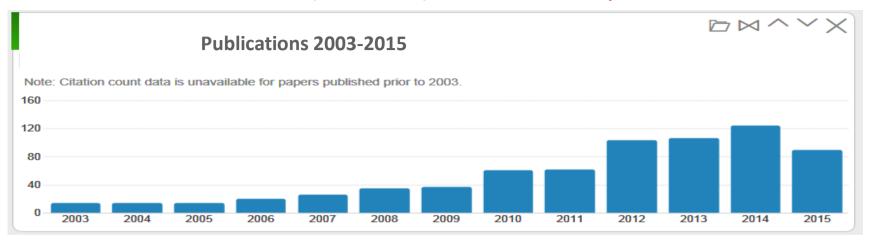
- 4 Research Centers
 - \$1.28 million TC per year each
- 1 Coordinating Center
 - \$375,896 TC per year
- Invited speakers/small workshops
 - \$15,000 per year (administered by DCB DCP)

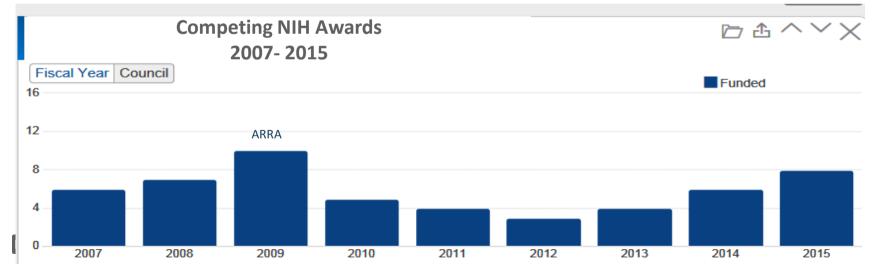
- Total requested set aside for entire project period:
 - \$27.6 million

Portfolio Analysis - Fiscal Year 2014

- NCI Extramural: ~ \$17.2 million in total costs (TC)
 - BETRNet, R01s, U01s and P01
- NCI Intramural: \$670,804 TC
- Other NIH institutes: \$2.2 million TC
 - K awards, some R01s
- No existing multi-disciplinary collaborative programs

Correlation of BETRNet (2011-2105) With the Landscape of EA Research







Program Evaluation Metrics (New Program)

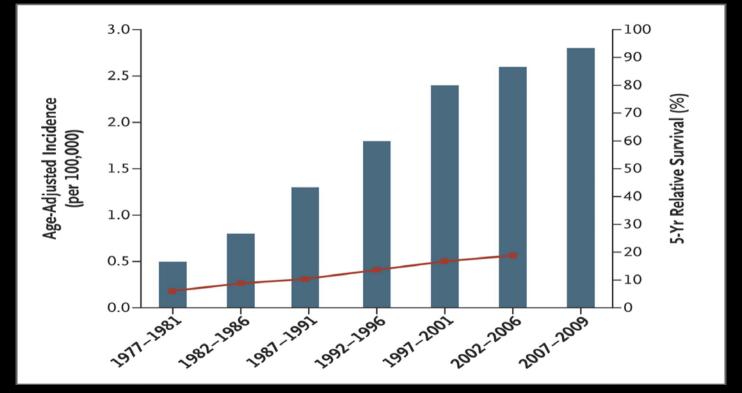
- Detailed scientific understanding of EA formation (years 1-4)
 - Validation of the initial cell of origin findings in animal models and humans
 - Validation of the genomic doubling hypothesis underlying the molecular risk for EA formation
 - Sharing of scientific expertise and valuable specimens to tackle these research questions
 - Publication of the team research findings
- Improved PR-VR availability to the larger scientific community (years 1-5)
 - Enhanced PR-VR accessibility for review by the larger scientific community (year 1)
 - Promotion of specimens availability for collaborative research studies (years 1-5)
 - Invitation of outside investigators to contribute specimens for collaborative research studies (years 1-5)
- New research tools and technologies for patient management (years 1-5)
 - Development of non-endoscopic and imaging devices for screening and surveillance
 - Development and validation of biomarker for risk stratification
 - Development of chemopreventive approaches
- Model organisms, GWAS, data and resource sharing (according to NIH policies) (years 3-5)
 - Links to data public sites available through the Coordinating Center website
 - Links to publications from outside investigators utilizing the resources



•Questions?



Temporal Trends in Incidence Rates and Survival Rates for Esophageal Adenocarcinoma



Additional BETRNet Accomplishments

- First-in-human trial
 - Image guided endoscopic biopsy using fluorescent peptides to bind with cell surface markers (e.g., epidermal growth factor receptor)
 - Aberrant vimentin methylation common in upper GI neoplasia
 - Novel non-endoscopic device study is underway

Additional Program Evaluation Criteria

- Collaboration and participation in new pilot studies within the network and outside the network
- Sharing of human specimens between centers to answer collaborative research questions
- New grant applications from cross-BETRNet studies
- New models and resources to be shared with the scientific community
- New research tools & applications for patient management
- Active participation in Steering Committee meetings, focused scientific committees, and other BETRNet activities
- Publication of collaborative research findings