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Xifeng Wu

Present Title & Affiliation

Primary Appointment

Department Chair, Department of Epidemiology, Division of OVP, Cancer Prevention and Population Sciences, The University of Texas MD Anderson Cancer Center, Houston, TX

Director, Center for Translational and Public Health Genomics, The University of Texas MD Anderson Cancer Center, Houston, TX

Professor, Department of Epidemiology, Division of Cancer Prevention and Population Sciences, The University of Texas MD Anderson Cancer Center, Houston, TX

Dual/Joint/Adjunct Appointment

Professor, Division of Epidemiology, Human Genetics and Environmental Sciences, The University of Texas School of Public Health, Houston, TX

Bio Statement

My major area of interest is using a molecular epidemiological approach to develop and validate genetic biomarkers for cancer risk assessment and for clinical outcome prediction via a variety of epidemiologic and molecular techniques (including genetics, epigenetics, cytogenetics, and computational biology). I am the principle investigator of multiple large NCI funded epidemiologic projects. We are applying candidate gene and genome-wide scanning approaches to identify cancer susceptibility loci and germline genetic predictors of clinical outcomes. We have developed or adapted an array of phenotypic assays in population studies (including but not limited to mutagen sensitivity assay for latent genetic instability, Comet assay and host-cell reactivation assay for DNA repair capacity, interphase FISH to identify hotspots for DNA damage, quantitative FISH to measure telomere length, mitochondrial DNA deletion and copy number changes, and global hypermethylation). We are also applying a variety of high-throughput genome-wide approaches, such as SNP-array CGH, gene expression array, microRNA array, and methylation array, to profile genetic and biological changes in premalignant and malignant somatic tissues. The long term goal of my research is to incorporate epidemiological, clinical, and genetic (germline and/or somatic) information to develop personalized risk prediction models for cancer etiology and clinical outcomes, for example, my group developed the first basic and enhanced bladder cancer risk prediction model. My research team is multi-disciplinary with diverse expertise in molecular biology, genetics, epidemiology, and statistics. I have an excellent track record of training students, post-docs, clinical fellows, and junior faculties and many of them have got independent tenure-track faculty positions.

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Education & Training

| 1994 | The University of Texas Health Science Center at Houston School of Public Health, Houston, TX, PHD, Epidemiology |
|------|--|
| 1987 | Zhejiang University, Hangzhou, China, MS, Industrial Hygiene and Occupational Diseases |
| 1984 | Fudan University, Shanghai, China, MD, Medicine |

Experience/Service

Institutional Committee Activities

Member, Bladder Cancer SPORE Executive Committee, 2002-present

Honors and Awards

| 2012-present | Wall of Science Publication Gallery Inclusion, The University of Texas MD Anderson Cancer Center |
|--------------|---|
| 2011 | Robert M. Chamberlain Distinguished Mentor Award, The University of Texas MD Anderson Cancer Center |
| 2008-present | Senior Fellow of the MD Anderson Research Trust, The University of Texas MD Anderson Cancer Center |

Selected Publications

Peer-Reviewed Original Research Articles

- He Y, Gong Y, Lin J, Chang DW, Gu J, Roth JA, Wu X. Ionizing radiationinduced γ-H2AX activity in whole blood culture and the risk of lung cancer. Cancer Epidemiol Biomarkers Prev. e-Pub 1/8/2013. PMID: 23300022.
- Ye Y, Madison B, Wu X, Rustgi AK. A LIN28B polymorphism predicts for colon cancer survival. Cancer Biol Ther 13(14):1390-5, 12/1/2012. e-Pub 10/10/2012. PMCID: PMC3542229.
- 3. Zhang N, Wu X, Yang L, Xiao F, Zhang H, Zhou A, Huang Z, Huang S. FoxM1 Inhibition Sensitizes Resistant Glioblastoma Cells to Temozolomide by Downregulating the Expression of DNA-Repair Gene Rad51. Clin Cancer Res 18(21):5961-71, 11/1/2012. e-Pub 9/12/2012. PMID: 22977194.
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- Gu J, Wang Y, Wu X. MicroRNA in the Pathogenesis and Prognosis of Esophageal Cancer. Curr Pharm Des. e-Pub 9/25/2012. PMID: 23092349.
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- Lin J, Lu C, Stewart DJ, Gu J, Huang M, Chang DW, Lippman SM, Wu X. Systematic Evaluation of Apoptotic Pathway Gene Polymorphisms and Lung Cancer Risk. Carcinogenesis. e-Pub 8/2012. PMID: 22665367.
- Lin M, Gu J, Eng C, Ellis LM, Hildebrandt MA, Lin J, Huang M, Calin GA, Wang D, Dubois RN, Hawk ET, Wu X. Genetic Polymorphisms in MicroRNA-Related Genes as Predictors of Clinical Outcomes in Colorectal Adenocarcinoma Patients. Clin Cancer Res 18(14):3982-91, 7/2012. e-Pub 6/2012. PMID: 22661538.
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- Chang DW, Gu J, Wu X. Germline prognostic markers for urinary bladder cancer: Obstacles and opportunities. Urol Oncol 30(4):524-32, 7/2012. PMID: 22742565.
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- Majumdar A, Curley SA, Wu X, Brown P, Hwang JP, Shetty K, Yao ZX, He AR, Li S, Katz L, Farci P, Mishra L. Hepatic stem cells and transforming growth factor β in hepatocellular carcinoma. Nat Rev Gastroenterol Hepatol. e-Pub 6/2012. PMID: 22710573.
- Lin M, Eng C, Hawk ET, Huang M, Lin J, Gu J, Ellis LM, Wu X. Identification of polymorphisms in ultraconserved elements associated with clinical outcomes in locally advanced colorectal adenocarcinoma. Cancer. e-Pub 6/2012. PMID: 22673945.
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