

NAME: WENRUI DUAN

CURRENT ADDRESS:

6715 Stillhouse LN,
Dublin, OH 43016

E-mail: wduan@fiu.edu

CITIZENSHIP: USA

ACADEMIC EXPERIENCE:

October 2016 –

Associate Professor, Faculty of Herbert Wertheim College of Medicine
Department of Human & Molecular Genetics, the Florida International University,
Miami, Florida.

January 2010- October 2016

Assistant Professor- Research, Faculty of Medical School
Division of Medical Oncology, Department of Internal Medicine and
Comprehensive Cancer Center, The Ohio State University, Columbus, Ohio.

July 2004 – December 2009

Research Scientist, Principal Investigator, Comprehensive Cancer Center, The Ohio
State University, Columbus, Ohio.

September 1999 – June 2004

Post-doc/research associate, Ohio State University Medical Center, The Ohio State
University, Columbus, Ohio.

September 1992 – August 1999

Graduate Research /Teaching Associate, College of Biological Science, The Ohio State
University, Columbus, Ohio.

December 1987 - June 1991

Lecturer: Department of Biology, Inner Mongolia Normal University. Huhhot, China.

1982 – 1987

Teaching Associate: Department of Biology, Inner Mongolia Normal University.
Huhhot, China.

EDUCATION:

Ph. D. Molecular Genetics, December, 1999

Department of Molecular Genetics, the Ohio State University, Columbus, OH

M. S. Zoology, December, 1994

College of Biological Science, the Ohio State University, Columbus, OH

M. S. Biology, July 1989
Inner Mongolia Normal University, Huhhot, China

B. S. Biology, 1982
Inner Mongolia University, Huhhot, China

PUBLICATIONS:

Original Reports

1. Kalvala A, Gao L, Aguila B, Dotts K, Rahman M, Nana-Sinkam P, Zhou X, Wang QE, Amann J, Otterson GA, Villalona-Calero MA, **Duan W*** Rad51C-ATXN7 fusion gene expression in colorectal tumors. *Mol Cancer*. 15(1):47. In press 2016, PMID: 27296891
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4. Kalvala A, Gao L, Aguila B, Reese T, Otterson GA, Villalona-Calero MA, **Duan W***. Overexpression of Rad51C splice variants in colorectal tumors. *Oncotarget*. 6(11):8777-87. 2015. PMID: 25669972
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7. Singh M, Leasure JM, Chronowski C, Geier B, Bondra K, **Duan W**, Hensley LA, Villalona-Calero M, Li N, Vergis AM, Kurmasheva RT, Shen C, Woods G, Sebastian N,

- Fabian D, Kaplon R, Hammond S, Houghton P, Palanichamy K, Chakravarti A. FANCD2 is a Potential Therapeutic Target and Biomarker in Alveolar Rhabdomyosarcoma Harboring the PAX3/FOXO1 Fusion Gene. *Clin Cancer Res.* 20(14):3884-95. 2014.
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 9. Shilo K, Wu X, Sharma S, Welliver M, **Duan W**, Villalona-Calero M, Fukuoka J, Sif S, Baiocchi R, Hitchcock CL, Zhao W, Otterson GA. Cellular localization of protein arginine methyltransferase-5 correlates with grade of lung tumors. *Diagn Pathol.* 8(1):201, 2013.
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Note: * corresponding author

Abstracts (Peer Reviewed, Published).

1. **Duan W**, Ding H, Subler M, Windle J, Otterson G, Villalona-Calero M. Lung specific expression of mutant p53 (273H) in mice. *Proceedings of the American Association for Cancer Research* (42) 2001, P 221 (abstr 1185)

2. Ding H, **Duan W**, Zhu W, Ju R, Subler M, Otterson G, Villalona-Calero, M. Mechanisms of regulation of CHK2 expression after exposure to the topoisomerase II inhibitor genistein. *Proceeding of AACR-NCI-RORTC International Conference*. 2001, P 139 (abstr 708).
3. Kolesar J, Miller J, **Duan W**, Drengler R, Felton S, Schaaf L, Von Hoff D, Kuhn J, Villalona-Calero M. TI Induction in PBMC Predicts Response to MMC/CPT-11. *Proceedings of American Society of Clinical Oncology* 2002, P115a (abstr 456)
4. Zhu W, Dai Z, Srinivasam K, Ding H, **Duan W**, Villalona M, Plass C, Otterson G. Methylated Sp1 binding sites are responsible for p21 silencing. *Proceedings of the American Association for Cancer Research* (43) 2002, P 1115 (abstr 5527).
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8. **Duan W**, Gao L, Zhu W, Otterson G, and Villalona-Calero M. In vivo expression of human mutant p53-273H does not regulate the expression of p21^{WAF1/CIP1} in p53 null and heterozygous mice. *Proceedings of the American Association for Cancer Research Annual Meeting* (44) 2003, P660.
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10. Panico K, **Duan W**, Otterson G, Kleiber B, Hindman K, Shah M, Young D, Wu W, Kuhn J, Villalona-Calero M. Phase I and pharmaco-biological study of thalidomide and celecoxib as modulators of irinotecan's anti-cancer activity. *Proceedings of the AACR-NCI-EORTC International Conference, Molecular Targets and Cancer Therapeutics*, 2003, p6197 (abstr B248).
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Proceedings of the 95th Annual Meeting of the American Association for Cancer Research, 2004 (abstr 1871).

12. Gao L, **Duan W**, Zhu W, Otterson G, Villalona-Calero M. Gene dosage and aging in lung tumorigenesis in p53 (273H) transgenic mice. *Proceedings of the 95th Annual Meeting of the American Association for Cancer Research*, 2004 (abstr 2553).
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14. Villalona-Calero M, Sawada T, Fukino K, **Duan W**, Brena RM, Williams N, Plass C, Eng C. Variable responses to EGFR tyrosine kinase inhibitors in non-small cell lung cancer (NSCLC) may be due to variable somatic EGFR mutational spectra. *The third international Chicago Symposium on malignancies of the Chest and Head and Neck*. 2004.
15. **Duan W**, Gao L, Wu X, Zhang Y, Otterson G, and Villalona-Calero M. Differences between the Pirh2 and Mdm2 in Response to DNA Damage. *Proceedings of the 96th Annual Meeting of the American Association for Cancer Research*, 2005 (abstr 3639).
16. **Gao L**, Duan W, Zhang Y, Wu X, Otterson G, and Villalona-Calero M. Establishment and characterization of a line of transgenic mice harboring a lung specific human type II p53(175H) mutation. *Proceedings of the 96th Annual Meeting of the American Association for Cancer Research*, 2005 (abstr 1055).
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18. Villalona-Calero M, **Duan W**, Gao L, Zhang Y, Wu X, Otterson G. Establishment and characterization of a line of transgenic mice harboring a lung specific human type II p53(175H) mutation: a potential model of human disease. *Proceedings of the 11th World Conference on Lung Cancer*, 2005 (abstr P-128)
19. Villalona-Calero M, **Duan W**, Gao L, Wu X, Otterson G. Establishment of Lung Cancer Transgenic Animal Models with Type Specific P53 Mutants for Experimental Therapeutics Studies. *4th International Chicago Symposium on Malignancies of the Chest and Neck & Head*, 2006 (abstr 4.04.04), pp909
20. **Duan W**, Gao L, Wu X, Otterson G, and Villalona-Calero M. Expression of a mutant p53 in lung specific transgenic mice cause an age-related demographic shift in lung tumor

incidence. *Proceedings of the 98th Annual Meeting of the American Association for Cancer Research*, 2007 (abstr 3875).

21. Gao L, **Duan W**, Wu X, Otterson G, and Villalona-Calero M. Human mutant p53 lung cancer cells respond differently to PRIMA-1 induced apoptosis. *Proceedings of the 98th Annual Meeting of the American Association for Cancer Research*, 2007 (abtr 4879).
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23. **Duan W**, Gao L, Wu X, Wang L, Hade EM, Gao GX, Barsky SH, Otterson GA, Villalona-Calero M. Mutant p53 collaborating with K-ras gene mutations and p16INK4a promoter methylation result in an age-related demographic shift in spontaneous lung tumor formation in transgenic mice. *Proceedings of the 99th Annual Meeting of the American Association for Cancer Research*, 2008 (abstr 2960).
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26. **Duan W**, Gao L, Wu X, Cohn D, O'Malley D, Dial L, Otterson G, and Villalona-Calero M. Analysis of FANCD2 protein monoubiquitin status and nuclear foci formation in ovarian cancers. *Proceedings of the 100th Annual Meeting of the American Association for Cancer Research*, 2009 (abstr 1019).
27. Gao L, **Duan W**, Wu X, Otterson G, and Villalona-Calero M. PRIMA-1 regulates microRNA 34A to promote apoptosis in human lung cancer cells. *Proceedings of the 100th Annual Meeting of the American Association for Cancer Research*, 2009 (abstr 5586).
28. Cohn DE, **Duan W**, Leon M, Resnick K, Zhao W, Gao L, Villalona M. Assessment of the Fanconi Anemia/BRCA pathway in epithelial ovarian cancer: An exploratory analysis providing insight into the use of PARP inhibitors. *Proc Soc Gynecol Oncol* 2010.
29. Resnick K, Leon M, **Duan W**, Zhao W, Salani R, Cohn DE, Villalona-Calero M. Identification of candidates for poly(ADP-ribose) polymerase inhibition: A pilot analysis in patients with cervical cancer. *Proc Soc Gynecol Oncol* 2010.

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31. Zhao W, **Duan W**, Leon ME, Chen AP, Sofletea G, Thurmond J, Ramaswamy B, O'Malley D, Bekaii-Saab TS, Villalona MA. Targeting fanconi anemia (FA) repair pathway deficiency for treatment with PARP inhibitors. *Proceedings of American Society of Clinical Oncology*, 2010 (abstr 43623)
32. **Duan W**, Gao L, Vereb G, Wu X, Otterson G, and Villalona-Calero M. Regression of spontaneous non-small cell lung cancers in P53 mutant transgenic mice following exposure to PRIMA-1. *Cancer Res* 2011;71(8 Suppl):Abstract nr 610
33. Li Gao **Duan W**, Wu X, Otterson G, and Villalona-Calero M. Inhibition of poly(ADP-ribose) polymerase in lung cancer cells with defective Fanconi anemia (FA) pathway. *Proceedings of the 102 Annual Meeting of the American Association for Cancer Research*, 2011 (abstr 5500). 04/2011
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CURRENT RESEARCH PROJECTS

Project 1: Targeting cancer cells containing mutant p53

p53 mutations are reported in 50-60% of non-small cell lung cancers and in up to 90% of small cell tumors, thus p53 represent a common mutation in this malignancy. Studies have shown that restoring wild type p53 function leads to regression of cancers, supporting the notion of treating human cancers by way of pharmacological reactivation of p53. The novel small molecule PRIMA-1 (p53-dependent reactivation and induction of massive apoptosis) has been shown to induce apoptosis in human tumor cells containing mutant p53. We are investigating the effects of PRIMA-1 in apoptosis using human lung cancer cell lines containing wildtype, mutant and null p53. Because most normal lung cells contain wild type p53, PRIMA-1 is expected to be much less toxic to normal lung cells compared to cancer cells. We have developed two lines of lung specific mutant p53 transgenic mice, which develop spontaneous non-small cell lung cancers. We plan to treat these lung tumor bearing mice with PRIMA-1 either as a single agent or in combination with cisplatin in order to uncover the potential role of PRIMA-1 in inducing apoptosis in spontaneous non small cell lung cancers. We are especially interested in the relationship between PRIMA-1 and microRNAs in apoptosis. This project was funded by Joan's Legacy and LUNgevity Foundation.

Project 2: Functional Analysis of the Fanconi Anemia (FA) pathway to select cancer treatment

The Fanconi Anemia (FA) pathway is essential for human cells to maintain integrity following DNA damage. Twenty genes are involved in the homologous recombination repair mediated by this pathway. More recent studies provide evidences linking disruption of FA cascade in sporadic cancers. These disruptions of FA pathway may involve epigenetic silencing of the FA-core complex or mutations of one or several FA genes. Cancer cells with these defects have been shown to be more sensitive to treatment with DNA interactive chemotherapeutic agents, as well as to agents which disrupt other repair mechanisms. However, the complexity to evaluate each individual gene for genetic or epigenetic changes have prevented to take advantage of the presence of these defects to select the patients most likely to benefit from DNA breaking treatment. The development of assays that can evaluate functionality of the whole pathway would be critical to make this a practical approach. Monoubiquitylation of FancD2 protein is a critical and gene-action converging step in the activation of this pathway, leading to the repair of DNA interstrand cross-link agents induced damage. Foci on chromatin can be appreciated when the function of this pathway is spared. We are conducting research in solid tumor specimens from patients undergoing surgical procedures to evaluate for an all inclusive method assessing the functionality of the Fanconi Anemia pathway. This test could potentially lead to the identification of patients most likely to respond to DNA breaking agents or to agents inhibiting compensatory mechanisms of repair. This project was funded by NCI research grant.

Project 3: *Genetic and epigenetic alterations in human lung cancer*

A major mechanism of DNA repair related to homologous recombination is the Fanconi Anemia (FA) pathway. FA genes collaborate with BRCA genes to form foci of DNA repair on chromatin following DNA damage or during S phase of cell cycle. Lack of repair foci and BRCA gene (BRCA1 or BRCA2) deficiency have been identified as predictors of the cytotoxicity of DNA breaking agents respectively. Although the numbers of cancer patients with germ line BRCA deficiency are low, BRCA genes are two of many genes that collaborate in the FA pathway. More recent studies provide evidence linking disruption of FA/BRCA cascade in sporadic cancers. However, more than 20 genes are actively involved in the FA pathway and both genetic and epigenetic alterations can occur. The aggregate contribution of these genes results first in activation and FANCD2 followed by the formation of FANCD2 repair foci in the nucleus of proliferating cells. We have recently developed a FA triple staining immunofluorescence based method (FATSI) to evaluate FANCD2 foci formation, which is capable of evaluating the presence or absence of FANCD2 foci using formalin fixed paraffin embedded (FFPE) tumor samples, and have generated preliminary data showing somatic deficiency of this pathway in tumors across several organ sites that include lung cancers. However, the specific genetic and epigenetic alterations that lead to inactivation of FA pathway in these sporadic tumors and the potential implication of these changes related to treatment are still unknown. We hypothesize that a substantial proportion of lung tumor harbors somatic genetic or epigenetic alterations resulting in defective FA homologous recombination repair. In order to provide molecular evidence that will help us to understand FA pathway inactivation in lung cancer, we propose to accomplish the following research aims: 1) To identify FANCD2 foci defective tumor in lung cancer patients by analyzing human lung tumor and matching non-tumor samples; 2) To perform a functional genomics analysis of FANCD2 foci defective tumors identified in aim 1 through analysis of the RNA transcriptome associated with the FA pathway in the tumor and adjacent non-tumor tissues, using next generation sequencing (RNAseq), 3) To investigate promoter methylation of FA genes in FANCD2 foci defective tumors identified in aim 1. This project was supported by NCI research grant and American Cancer Society research grant.