# CURRICULUM VITAE ~ Chi Wang

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Date Prepared: August 29, 2022

## CURRICULUM VITAE

Chi Wang, PhD Professor, with tenure, Regular Title Series **Department of Internal Medicine Division of Cancer Biostatistics University of Kentucky College of Medicine** 



### I. GENERAL INFORMATION

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Lexington, KY, 40536

chi.wang@uky.edu **Email** 

**Telephone** 859-323-2045 859-323-2074 Fax

Personal Website http://sweb.uky.edu/~cwa236/

#### II. EDUCATION

<u>Undergraduate</u> [oldest at top, newest at bottom]

**Peking University** 

Beijing, China

09/1997-06/2001 BS, Statistics

**Professional/Graduate** [oldest at top, newest at bottom]

**Peking University** 

Beijing, China

09/2001-06/2003 MS, Statistics

Johns Hopkins Bloomberg School of Public Health

Baltimore, MD, USA

09/2004-12/2009 PhD, Biostatistics

# **III. PROFESSIONAL EXPERIENCES** [oldest at top, newest at bottom]

Not applicable.

# IV. ACADEMIC APPOINTMENTS [specify tenure/non-tenure track, academic/nonacademic, full-time or part-time]

Faculty [oldest at top, newest at bottom]

	University of California, Riverside Riverside, CA
07/2009-06/2010	Assistant Professor, Department of Statistics, tenure-track, full-time
	University of Kentucky
07/2010-06/2016	Lexington, KY Assistant Professor, Division of Cancer Biostatistics, Department of Biostatistics, College of Public Health and Biostatistics and
	Bioinformatics Shared Resource Facility, Markey Cancer Center, Regular Title Series, tenure-track, full-time
07/2016-12/2019	Associate Professor, Division of Cancer Biostatistics, Department of Biostatistics, College of Public Health and Biostatistics and
	Bioinformatics Shared Resource Facility, Markey Cancer Center, Regular Title Series, with tenure, full-time
01/2020-06/2022	Associate Professor, Division of Cancer Biostatistics, Department of Internal Medicine, College of Medicine and Biostatistics and Bioinformatics Shared Resource Facility, Markey Cancer Center, Regular
	Title Series, with tenure, full-time
05/2021-06/2022	Associate Professor, Department of Statistics, College of Arts and Sciences, Joint (Secondary) Appointment
07/2017-	Associate Director for Bioinformatics, Biostatistics and Bioinformatics Shared Resource Facility, Markey Cancer Center, full-time
07/2022-	Professor, Division of Cancer Biostatistics, Department of Internal Medicine, College of Medicine and Biostatistics and Bioinformatics Shared Resource Facility, Markey Cancer Center, Regular Title Series, with tenure, full-time
07/2022-	Professor, Department of Statistics, College of Arts and Sciences, Joint (Secondary) Appointment

# V. HOSPITAL or CLINICAL APPOINTMENTS [oldest at top, newest at bottom; specify full-time or part-time]

Not applicable.

VI.	<b>CONSULTING ACTIVITIES</b> [oldest at top, newest at bottom in each section]
Not ap	pplicable.

VII. TEACHING ACTIVITIES [oldest at top, newest at bottom in each section; use for students, residents, fellows, continuing education programs]

#### **University Faculty**

#### Course Instructor

I was the instructor of the following courses. For most courses, I prepared course materials, gave a series of lectures throughout the semester, made and graded homework and exams.

Particularly, I have developed a new course "Sequencing Data Analysis" on introducing state-ofthe-art statistical and bioinformatics methods for next-generation sequencing data analysis, and have taught the course twice.

University	of	California	, Riverside
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Riverside, CA

09/2009-12/2009 Probability and Statistics for Science and Engineering [STAT 155]

> This is an introductory course for undergraduate students from science and engineering departments. It covers basic probability and statistics concepts and methods with an emphasis on techniques and applications that are useful in engineering, computer science, and the physical sciences.

Discrete Data Analysis [STAT 205] 01/2010-03/2010

> This course is designed for PhD students from Department of Statistics to introduce statistical theory and methods for categorical data analysis.

General Statistical Models [STAT 171] 03/2009-05/2010

> This course is designed for MS students from Department of Statistics to provide an applied introduction of statistical methods for categorical data

analysis.

**University of Kentucky** 

Lexington, KY

01/2012-05/2012 Biometrics II [BST 676]

> This course, the second of a two-semester sequence in biometrics, is designed for students in the joint Epi-Bio PhD program. It develops numerous techniques for constructing and rigorously evaluating point estimators, hypothesis testing procedures, and interval estimators.

01/2013-05/2013 Biometrics II [BST 676] 01/2014-05/2014 Biostatistics I [CPH 580]

> This course, the first of a two-semester sequence in biostatistics, is an introductory course for graduate students from various colleges. It covers basic statistics concepts and methods used in medicine, public health, and the biological sciences. It also demonstrates using SAS to perform basic

statistical analysis.

01/2015-05/2015 Biostatistics I [CPH 580] 08/2016-12/2016 Doctoral Seminar [CPH 786]

> This course is designed for PhD students from Department of Biostatistics to enhance their research ability. Students are required to read research

	articles, give presentations, participate in paper discussions, and attend colloquium sessions
01/2017-05/2017	Sequencing Data Analysis [CPH 738-001]
	This is <i>a new course that I developed</i> for graduate students from various
	colleges. It introduces state-of-the-art statistical and computational
	methods for processing and analyzing genomic data generated by next
	generation sequencing, and provides hands-on genomic data analysis
	experience.
01/2018-05/2018	Doctoral Seminar [CPH 786]
01/2019-05/2019	Sequencing Data Analysis [CPH 738-002/STA715-005]

# **Guest Lectures**

I was invited to give guest lectures (1 to 1.5 hours each) for the following courses from various departments in College of Medicine and College of Public Health.

	University of Kentucky
	Lexington, KY
04/2013	Biology and Therapy of Cancer [MI/MED/PHA 616, topic: Cancer
	Biostatistics for Basic and Translational Research]
09/2013	Drug Discovery, Development, and Translation [PHS 760, topic:
	Bioinformatics for Translational Discoveries and Targeted Clinical
	Studies]
10/2013	Practical Statistics [IBS 611, topic: Bioinformatics I & II]
11/2013	Introduction to Bioinformatics [CPH 738-007, topic: Next-Generation
	Sequencing]
04/2014	Introduction to Bioinformatics [CPH 738-007, topic: Next-Generation
	Sequencing]
12/2014	Practical Statistics [IBS 611, topic: Introduction to Bioinformatics]
09/2015	The 3 <sup>rd</sup> annual Markey Cancer Center Clinical Trials Boot Camp [topic:
	Bioinformatics for NGS data @ Biostatistics and Bioinformatics
	Shared Resource Facility]
04/2017	Introduction to Bioinformatics [BMI 633, topic: Next-Generation
	Sequencing]
08/2017	2017 Markey Cancer Center Clinical Trials Boot Camp [topic: Genomics
	and Bioinformatics in Clinical Studies]
12/2017	Introduction to Bioinformatics [BMI 633, topic: Next-Generation
	Sequencing]
03/2019	Introduction to biomedical image informatics [BMI 734, topic: Cancer
	Somatic Mutation Analysis based on Next Generation Sequencing]
03/2019	Introduction to biomedical image informatics [BMI 734, topic: Cancer
	Somatic Mutation Analysis based on Next Generation Sequencing]
02/2020	Introduction to Bioinformatics [BMI 633, topic: Cancer Somatic Mutation
	Analysis based on Next Generation Sequencing]
10/2020	Instrumental Techniques In Forensic Chemistry [TOX 920, topic: Basic
	Probability and Statistics Review]

10/2020	Forensic and Analytical DNA [TOX 910, topic: Basic Probability and
	Statistics Review]
11/2020	Introduction to Bioinformatics [BMI 633, topic: Cancer Genomics with
	A Focus on Somatic Mutation Analysis]
04/2021	Introduction to Bioinformatics [BMI 633, topic: Cancer Genomics with
	A Focus on Somatic Mutation Analysis
10/2021	Forensic and Analytical DNA [TOX 910, topic: Basic Probability and
	Statistics Review]
12/2021	Introduction to Bioinformatics [BMI 633, topic: Cancer Genomics with
	A Focus on Somatic Mutation Analysis]

# **Professional Course/Program Faculty**

American College of Clinical Pharmacy (ACCP) Foundation Mentored Research Investigator Training (MeRIT) program

Lexington, KY

06/2019 Basic Statistics Review

# VIII. ADVISING ACTIVITIES [oldest at top, newest at bottom in each section]

#### Advisor/Co-Advisor

I have served as advisor/co-advisor for the following eight PhD students from University of Kentucky. Six of those students have already graduated. During the entire period of time that a student worked on the dissertation, I spent one to three hours per week with the student to have a one-to-one meeting, provide guidance on research topic selection, literature review, statistical methods derivation, simulation studies and real data analyses, encourage participation and presentation in conferences, and revise manuscripts and dissertation.

Supervision	Role	Student's	Department	Current
time period		name		position
06/2013-	Co-Advisor	Hong Wang*	Statistics	Sanofi
07/2016				
06/2015-	Co-Advisor	Li Xu	Statistics	Wells Fargo
07/2020				
06/2016-	Advisor	Zhengyan Huang	Biostatistics	Everest Clinical
07/2019				Research
06/2016-	Co-Advisor	Tingting Zhai	Statistics	Novartis
07/2020				
06/2016-	Co-Advisor	Xu Zhang	Statistics	Gilead Sciences
07/2020				
06/2018-	Co-Advisor	Tiantian Zeng	Statistics	Merck
05/2022				
06/2016-	Co-Advisor	Menghan Wang	Statistics	
present		-		
06/2020-	Advisor	Kun Liu	Statistics	
present				

<sup>\*</sup> Winner of the ENAR 2016 Distinguished Student Paper Award

### **Dissertation Committee Member**

I have served as a member on dissertation committees for students from a variety of departments at the University of Kentucky, where I joined the committees at either early or late stage of students' dissertation development depending on their needs. My duties include attending committee meetings and final defense and providing statistical advice on students' dissertation works.

Time period	Student's	Department	Degree
	name		
01/2011-11/2011	Zhenyu Huang	Biostatistics	MS
03/2012-10/2013	Yin Hu	Computer Science	PhD
01/2014-06/2014	Jinpeng Liu	Computer Science	MS
11/2014-12/2014	Yan Huang	Computer Science	PhD
01/2015-11/2016	Hongyuan Wang	Statistics	PhD
06/2015-07/2015	Adam Parrish	Communication	PhD
09/2017-11/2018	Jin Xie	Statistics	PhD
09/2017-05/2019	Chenlu Ke	Statistics	PhD

09/2017-03/2018	Xinan Liu	Computer Science	PhD
02/2018-05/2018	Xiaoli Kong	Statistics	PhD
08/2018-02/2020	Yuntong Li	Statistics	PhD
10/2018-07/2020	Yue Cui	Statistics	PhD
11/2018-05/2020	Xue Ding	Statistics	PhD
01/2019-present	Xiaofei Zhang	Computer Science	PhD
03/2019-08/2019	Eugene Hinderer	Molecular & Cellular Biochemistry	PhD
05/2019-06/2019	Yi Zhang	Computer Science	PhD
06/2019-11/2020	Aisaku Nakamura	Statistics	PhD
08/2019-10/2020	Matthew Rutledge	Statistics	PhD
08/2019-present	Pei Wang	Statistics	PhD
11/2019-present	Jinpeng Liu	Computer Science	PhD
02/2020-09/2020	Ting Zeng	Statistics	PhD
03/2020-09/2021	Justin Wayne Gorski	CCTS	PhD
04/2020-present	Sheng Yuan	Statistics	PhD
10/2020-present	Nan Lin	Pharmacy	PharmD
04/2021-present	Leon Su	Statistics	PhD
10/2021-present	Percy Yeh	Statistics	PhD
08/2021-present	Brien Washington	Medical Physics	PhD

# IX. ADMINISTRATIVE ACTIVITIES & UNIVERSITY SERVICE [oldest at

top, newest at bottom in each section]

# **College**

# **University of Kentucky**

Lexington, KY

#### Administration & Clinical Operations

Member, College of Public Health Faculty Council 07/2015-06/2016

#### **Education & Research**

Member, College of Public Health MPH Admissions Committee 07/2011-06/2014 Member, College of Public Health Joint PHD Program Qualify Exam 07/2011-06/2013 Committee

Member, College of Public Health Research Committee 07/2017-06/2019

#### **Medical Center**

# **Markey Cancer Center**

Lexington, KY

# **Administration & Clinical Operations**

07/2017-present Associate Director for Bioinformatics, Biostatistics and Bioinformatics

**Shared Resource Facility** 

#### **Education & Research**

07/2017-present	Member, Biospecimen Procurement and Translational Pathology Shared
	Resource Facility Scientific Advisory Committee
10/2017-05/2019	Member, Oncogenomics Shared Resource Facility Scientific Advisory
	Committee
06/2019-present	Chair, Oncogenomics Shared Resource Facility Scientific Advisory
-	Committee
10/2020-05/2021	Member, Committee for External Genomics Vendor, Oncogenomics
	Shared Resource Facility

# X. SPECIAL ASSIGNMENTS

Not applicable.

# XI. HONORS & AWARDS [specify nature/meaning of each; academic, professional, honorary, not grants; oldest at top, newest at bottom]

10/2000	First Prize, the National College Mathematics Contest in Modeling, China
09/2000	Bao Jie Scholarship, Peking University, Beijing, China
09/2001	Second Prize, the 9th Challenge Cup Contest of Peking University,
	Beijing, China
09/2002	An Tai Scholarship, Peking University, Beijing, China
09/2004-06/2009	Graduate Scholarships, Johns Hopkins Bloomberg School of Public
	Health, Baltimore, MD
02/2008	Distinguished Student Paper Award, International Biometric Society
	Eastern North American Region (ENAR)
06/2010	Member of the Delta Omega Alpha Chapter
06/2010	Member of the Phi Beta Kappa Society

# XII. PROFESSIONAL ACTIVITIES, PUBLIC SERVICE & PROFESSIONAL DEVELOPMENT [oldest at top, newest at bottom in each section]

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07/2009-present 12/2020-present 11/2021-present	International Biometric Society International Chinese Statistical Association International Society for Computational Biology
<b>Review Panels</b>	
03/2013-04/2014	National Science Foundation Reviewer, Methodology, Measurement, and Statistics (MMS) Program
12/2010 12/2010	Oncology Research Information Exchange Network (ORIEN)
12/2018-12/2019 05/2021-05/2023	Member (Biostatistician reviewer), ORIEN Scientific Review Committee Member (Biostatistician reviewer), ORIEN Scientific Review Committee
	National Cancer Institute
06/2017-06/2017	Reviewer, Special Emphasis Panel focus on Clinical & Translational R21 & Omnibus R03: ZCA1 TCRB-J (O1)
04/2018-04/2018	Reviewer, Special Emphasis Panel focus on Clinical and Translational Exploratory/Developmental Studies ZCA1 TCRB-J (M1) S
07/2019-07/2019	Reviewer, Special Emphasis Panel on Collaborative Research at the NIH Clinical Center ZCA1 TCRB-J (O3)
10/2021-10/2021	Reviewer, NCI SPORE (P50) Review II ZCA1RPRB-6 (J1)
03/2022-03/2022	Reviewer, NCI Pancreatic Cancer Detection Consortium (U01/U24) Review ZCA1 RPRB-8 (M2)
	American Heart Association Institute for Precision Cardiovascular Medicine
10/2019-09/2021 07/2020-12/2020	Member, Data Science Study Section AHA COVID-19 Data Challenge phases I and II

#### Internal grant reviews

Internal grant review	<u>ws</u>
	University of Kentucky
12/2011	Kentucky Lung Cancer Research Program
05/2013	American Cancer Society Institutional Research Grant
12/2013	American Cancer Society Institutional Research Grant
01/2014	Kentucky Lung Cancer Research Program
08/2014	American Cancer Society Institutional Research Grant
12/2014	American Cancer Society Institutional Research Grant
11/2015	Markey Cancer Center Cancer Center Support Grant Pilot Program
12/2015	Kentucky Lung Cancer Research Program
08/2016	American Cancer Society Institutional Research Grant
06/2017	American Cancer Society Institutional Research Grant

06/2017 Center for Clinical and Translational Science Pilot and Innova	tion
Research Program	
11/2017 Peter and Carmen Lucia Buck Clinical Translational Research	Award
03/2018 American Cancer Society Institutional Research Grant	
06/2018 American Cancer Society Institutional Research Grant	
12/2018 American Cancer Society Institutional Research Grant	
01/2019 Markey Cancer Center Cancer ORIEN Award	
05/2019 American Cancer Society Institutional Research Grant	
10/2019 Markey Cancer Center Cancer Center Support Grant Pilot Prog	gram
12/2019 Center for Clinical and Translational Science Pilot and Innova	tion
Research Program	
03/2020 American Cancer Society Institutional Research Grant	
05/2020 Markey Cancer Center Cancer Center Support Grant Pilot Prog	gram
09/2020 American Cancer Society Institutional Research Grant	
03/2021 American Cancer Society Institutional Research Grant	
03/2022 American Cancer Society Institutional Research Grant	

# **Editorial Boards**

06/2014-present	Biometrics & Biostatistics International Journal, MedCrave network
07/2014-present	Journal of Biometrics & Biostatistics, Hilaris

# Journal Peer-Reviewing

03/2008-06/2018	Statistica Sinica
03/2010-04/2019	BMC Bioinformatics
06/2010-06/2022	PLoS ONE
06/2011-01/2012	The International Journal of Biostatistics
06/2011-06/2011	Journal of Classification
06/2011-08/2017	Statistical Applications in Genetics and Molecular Biology
10/2011-04/2022	Statistics in Medicine
01/2012-05/2015	Journal of Causal Inference
02/2012-04/2012	Journal of Statistical Planning and Inference
04/2012-04/2012	BMC Genomics
01/2013-01/2013	Journal of Biopharmaceutical Statistics
02/2013-02/2013	International Journal of Computational Biology and Drug Design
02/2013-04/2013	Computational Statistics and Data Analysis
03/2013-03/2013	Computational Biology and Chemistry
12/2014-06/2016	Biometrics & Biostatistics International Journal
04/2014-09/2017	Scientific Reports
01/2015-06/2018	Annals of Applied Statistics
06/2016-03/2020	Bayesian Analysis
07/2016-03/2017	Canadian Journal of Statistics
07/2016-10/2018	Briefings in Functional Genomics
07/2017-10/2017	Bioinformatics
07/2017-10/2018	Journal of the American Statistical Association

11/2018-09/2019	Cancer Medicine
04/2020-04/2020	Nucleic Acids Research
09/2020-11/2021	American Journal of Epidemiology
12/2021-12/2021	Journal of the Royal Statistical Society: Series C
02/2022-02/2022	Scandinavian Journal of Statistics

XIII. SPEAKING ENGAGEMENTS [Invited lectureships, panel sessions; oldest at top, newest at bottom in each section]

<u>Local</u>	
	University of California, Los Angeles
03/2010	Los Angeles, CA Department of Statistics: "Exponential Tilt Models for Time-to-Event Outcomes"
	University of Louisville Louisville, KY
02/2016	Department of Bioinformatics and Biostatistics: "Estimation of the Average Causal Effect while Accounting for Uncertainty in Confounder
11/2018	and Effect Modifier Selection" Department of Bioinformatics and Biostatistics: "Differential Abundance Analysis for Proteomic and Metabolomic Data"
	Peking University
05/2017	Beijing, China Department of Statistics: "Causal effect estimation while accounting for uncertainty in confounder and effect modifier selection"
	Jilin University
06/2017	Changchun, Jilin, China Translational Medical Science Institute, The First Bethune Hospital of
06/2017	Jilin University: "Bioinformatics Methods for Cancer 'Omics Studies" Department of Statistics: "NanoStringDiff: A Novel Statistical Method for Differential Expression Analysis Based on NanoString nCounter Data"
	University of Kentucky Lexington, KY
02/2018	Markey Research Seminar: "Probabilistic and Statistical Modeling for Cancer Omics Data"
03/2021	Department of Statistics: "Statistical Methods for Complex Cancer Data"
	University of Pittsburgh
02/2022	Pittsburgh, PA
03/2022	Hillman Cancer Center "Statistical Methods for Cancer Omics Data"
State/Regional	
	21st Century Cures
03/2018	Knoxville, TN Southeast Conference, Using Big Data to Overcome Health Disparities: "Mutational Characterization of Squamous Cell Lung Cancers from

Appalachian Kentucky: Moving Closer to Personalized Treatment"

	University of Kentucky
	Lexington, KY
03/2013	Inaugural Breast Cancer Symposium: "Bioinformatics Support from
	MCC's Biostatistics Shared Resource Facility (BSRF)"
03/2015	10 <sup>th</sup> Annual CCTS Spring Conference: "Statistical and Bioinformatics
	Methods for Genomic Data Analysis"
03/2016	4th Annual Breast Cancer Symposium: "Bioinformatics Data Processing
	and Analysis"
03/2017	5 <sup>th</sup> Annual Breast Cancer Symposium: "Bioinformatics methods for breast
	cancer genomic data"
10/2018	2 <sup>nd</sup> Commonwealth Computational Summit: "Bioinformatics Methods for
	Next Generation Sequencing-Based Cancer Studies"
	Southern Regional Council on Statistics
	Carrollton, KY
06/2019	Southern Regional Council on Statistics 2019 Conference: "Statistical
	Methods for Cancer Somatic Mutation Analysis"

# National/International

The Ohio State University	The	Ohio	State	Univ	ersity
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Columbus, OH

11/2019 Mathematical Biosciences Institute Workshop on Evolutionary Dynamics

in Cancer: "A probabilistic method to estimate the temporal order of pathway mutations during carcinogenesis by leveraging intra-tumor

phylogenies and functional annotations"

#### **International Chinese Statistical Association**

virtual conference

12/2020 ICSA 2020 Applied Statistics Symposium: "A Statistical Framework for

Genome-Scale Mutual Exclusivity Analysis of Cancer Mutations"

### **Oncology Research Information Exchange Network (ORIEN)**

03/2022 Tampa, FL

> 3rd Annual ORIEN Scientific Retreat: "Radiogenomics Consortium and Biomarker Development for Non-Small Cell Lung Cancer (NSCLC)

Patients"

#### XIV. RESEARCH & INTELLECTUAL CONTRIBUTIONS

A. PUBLICATIONS [oldest at top, newest at bottom in each section; number each within each section; published or accepted for publication/in press; NOT in preparation]

#### Statistical and Bioinformatics Methodological Papers

 $\overline{(^*: corresponding \ or \ co-corresponding \ author, ^\dagger: \ student/GRA)}$ 

- 1. Yang J, Wang C and Yang Y. The grouping of DNA sequences model. Journal of Mathematics in Practice and Theory 31(1):31-38, 2001. (in Chinese)
- 2. Geng Z, Wang C and Zhao O. Decomposing a moral graph to search for v-structures. Journal of Multivariate Analysis 96(2): 282-294, 2005.
- 3. Dominici F, Wang C, Crainiceanu C and Parmigiani P. Model selection and health effect estimation in envionmental epidemiology. Epidemiology 19:558-560, 2008.
- 4. Irizarry RA, Wang C, Zhou Y and Speed TP. Gene set enrichment analysis made simple. Statistical Methods in Medical Research 18:565–575, 2009.
- 5. Chen S, Wang C, Caffo BS, Eberly LE and Schwartz BS. Adaptive control of the false discovery rate in voxel-based morphometry. Human Brain Mapping 30:2304-2311, 2009.
- 6. Wang C\*, Tan Z and Louis TA. exponential tilt models in the presence of censoring. Journal of Statistical Planning and Inference 141:1102-17, 2011.
- 7. Wang C\*, Parmigiani G and Dominici F. Bayesian effect estimation accounting for adjustment uncertainty (with discussion). Biometrics, 68(3): 661-86, 2012.
- 8. Wu H<sup>‡</sup>, Wang C<sup>‡</sup> and Wu Z. A new shrinkage estimator for dispersion improves differential expression detection in RNA-seq. Biostatistics, 14(2): 232-43, 2013. ‡Authors with equal contribution.
- 9. Wang C\*, Tan Z and Louis TA. An exponential tilt mixture model for time-to-event data to evaluate treatment effect heterogeneity in randomized clinical trials. Biometrics & Biostatistics International Journal, 1(2):00006, 2014.
- 10. Wang C\*, Tan Z and Louis TA. An exponential tilt model for quantitative trait loci mapping with time-to-event data. Journal of Bioinformatics Research Studies, 1(2):2, 2014.
- 11. Tian S, Chang HH, Wang C, Jiang J, Wang X and Niu J. Multi-TGDR, a multi-class regularization method, identifies the metabolic profiles of hepatocellular carcinoma and cirrhosis infected with hepatitis B or hepatitis C virus, BMC Bioinformatics, 15:97, 2014.
- 12. Tian S, Wang C and An MW. Test on existence of histology subtype-specific prognostic signatures among early stage lung adenocarcinoma and squamous cell carcinoma patients using a Cox-model based filter. Biology Direct, 10:15, 2015.
- 13. Wang C\*, Dominici F, Parmigiani G and Zigler CM. Accounting for uncertainty in confounder and effect modifier selection when estimating average causal effects in generalized linear Models. Biometrics, 71(3):654-65, 2015.
- 14. Chen L, Wang C, Qin ZS and Wu H. A novel statistical method for quantitative comparison of multiple ChIP-seq datasets. Bioinformatics, 31(12):1889-96, 2015.
- 15. Wu H, Wang C and Wu Z. PROPER: Comprehensive power evaluation for differential expression using RNA-seq. Bioinformatics, 31(2):233-41, 2015.
- 16. Wang H<sup>†</sup>, Horbinski C, Wu H, Liu Y, Sheng S, Liu J, Weiss H, Stromberg A, Wang C\*. NanoStringDiff: A novel statistical method for differential expression analysis based on NanoString nCounter data. Nucleic Acids Research, 44(20): e151, 2016.

- 17. Wang C, Liu J and Fardo DW. Causal effect estimation in sequencing studies: A Bayesian method to account for confounder adjustment uncertainty. BMC Proceedings, 10(7): 411-415, 2016.
- 18. Tian S, Chang HH, Wang C. Weighted SAMGSR: combining significance analysis of microarray-gene set reduction algorithm with pathway topology-based weights to select relevant genes, Biology Direct, 11(1):50, 2016.
- 19. Tian S, Wang C, Chang HH, Sun J. Identification of prognostic genes and gene sets for early-stage nonsmall cell lung cancer using bi-level selection methods. Scientific Reports, 7:46164, 2017.
- 20. Huang  $Z^{\dagger}$ , Chen L and Wang  $C^{*}$ . Classifying lung adenocarcinoma and squamous cell carcinoma using RNA-Seq Data. Cancer Studies and Molecular Medicine. 2017 Sep; Volume 3: Issue 2.
- 21. Tian S, Wang C, Chang HH. A longitudinal feature selection method identifies relevant genes to distinguish complicated injury and uncomplicated injury over time. BMC Medical Informatics and Decision Making, 18(5):115, 2018.
- 22. Tian S, Wang C, Chang HH. To select relevant features for longitudinal gene expression data by extending a pathway analysis method. F1000Research. 2018;7.
- 23. Li Y<sup>†</sup>, Fan TWM, Lane AN, Kang WK, Arnold SM, Stromberg AJ, Wang C<sup>\*</sup>, Chen Li. SDA: A semi-parametric differential abundance analysis method for metabolomics and proteomics data. BMC Bioinformatics, 2019 Dec 1;20(1):501.
- 24. Wang M<sup>†</sup>, Yu T, Liu J, Chen L, Stromberg AJ, Villano JL, Arnold SM, Liu C, Wang C\*. A probabilistic method for leveraging functional annotations to enhance estimation of the temporal order of pathway mutations during carcinogenesis. BMC bioinformatics. 2019 Dec;20(1):1-2.
- 25. Tian S, Wang C\*. Feature Selection for Longitudinal Data by Using Sign Averages to Summarize Gene Expression Values over Time. Biomed Res Int. 2019;2019:1724898. doi: 10.1155/2019/1724898. eCollection 2019.
- 26. Tian S, Wang C, Wang B. Incorporating Pathway Information into Feature Selection towards Better Performed Gene Signatures. Biomed Res Int. 2019;2019:2497509. doi: 10.1155/2019/2497509. eCollection 2019. Review.
- 27. Huang Z<sup>†</sup>, Lane AN, Fan TW, Higashi RM, Weiss HL, Yin X, Wang C\*. Differential Abundance Analysis with Bayes Shrinkage Estimation of Variance (DASEV) for Zero-Inflated Proteomic and Metabolomic Data. Scientific Reports. 2020 Jan 21;10(1):1-2.
- 28. Tian S, Wang C, Suarez-Farinas M. GEE-TGDR: A longitudinal feature selection algorithm and its application to lncRNA expression profiles for psoriasis patients treated with immune therapies. BioMed Research International, 2021;2021:8862895.
- 29. Tian S, Wang C\*. An ensemble of the iCluster method to analyze longitudinal lncRNA expression data for psoriasis patients. Human Genomics 2021 Apr 20;15(1):23.
- 30. Liu S<sup>†</sup>, Liu J, Xie Y, Zhai T, Hinderer EW, Stromberg AJ, Vanderford NL, Kolesar JM, Moseley HNB, Chen L, Liu C, Wang C\*. MEScan: a powerful statistical framework for genome-scale mutual exclusivity analysis of cancer mutations. Bioinformatics. 2021 Jun 9;37(9):1189-1197.
- 31. Huang Z, Wang C. A Review on Differential Abundance Analysis Methods for Mass Spectrometry-Based Metabolomic Data. Metabolites. 2022 Apr;12(4):305.

## **Collaborative Papers** (\*: corresponding or co-corresponding author)

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#### XIV. RESEARCH & INTELLECTUAL CONTRIBUTIONS - continued

**B.** ABSTRACT PRESENTATIONS [specify type: Podium, Poster, Exhibit, Electronic, Educational Exhibit, Scientific Exhibit; oldest at top, newest at bottom in each section]

#### Local/State/Regional Meetings

- 1. 03/2007. Wang C, Z Tan and T Louis. Exponential Tilt Models in the presence of Censoring. International Biometric Society Eastern North American Region (ENAR) Spring Meeting. Atlanta, Georgia.
- 2. 03/2008. Wang C, F Dominici and G Parmigiani. A Bayesian Approach to Effect Estimation Accounting for Adjustment Uncertainty. International Biometric Society Eastern North American Region (ENAR) Spring Meeting. Arlington, Virginia. Distinguished Student Paper Award.
- 3. 03/2015. Wang C, F Dominici, G Parmigiani and C Zigler. Accounting for Uncertainty in Confounder Selection when Estimating Average Causal Effects in Generalized Linear Models. International Biometric Society Eastern North American Region (ENAR) Spring Meeting. Miami, Florida.

#### National/International

- 1. 08/2008. Wang C, Z Tan and T Louis. Exponential Tilt Models in the presence of Censoring. American Statistical Association Joint Statistical Meetings. Denver, CO.
- 2. 10/2014. Wang C, F Dominici and G Parmigiani. Bayesian Estimation of Average Causal Effect with Adjustment for Confounding. 3<sup>rd</sup> International Conference & Exhibition on Biometrics & Biostatistics. Baltimore, MD.
- 3. 08/2016. Wang C, F Dominici, G Parmigiani and C Zigler. Accounting for Uncertainty in Confounder and Effect Modifier Selection When Estimating Average Causal Effects in Generalized Linear Models. American Statistical Association Joint Statistical Meetings. Chicago, IL.
- 4. 03/2017. Wang H, Horbinski C, Wu H, Liu Y, Sheng S, Liu J, Weiss H, Stromberg A, Wang C. A Novel Statistical Tool for Differential Expression Analysis of NanoString nCounter Data. Statistical Practice in Cancer Conference, Tampa, FL.
- 5. 08/2018. Liu S, Liu J, Xie Y, Zhai T, Hinderer EW, Stromberg AJ, Canderford NL, Kolesar JM, Moseley HNB, Chen L, Liu C, Wang C. A New Statistical Method for Genome-Scale Mutual Exclusivity Analysis of Tumor Mutations. Vancouver, Canada.
- 6. 08/2020. Huang Z, Lane AN, Fan TW, Higashi RM, Weiss HL, Yin X, Wang C. Differential Abundance Analysis with Bayes Shrinkage Estimation of Variance (DASEV) for Zero-Inflated Proteomic and Metabolomic Data. American Statistical Association Joint Statistical Meetings. Virtual conference.

## XIV. RESEARCH & INTELLECTUAL CONTRIBUTIONS - continued

### C. SPONSORED RESEARCH PROJECTS, GRANT & CONTRACT ACTIVITIES

- A total of 48 funded grants
- PI/contact MPI from the National Cancer Institute:

1 R21 grant (\$382,304)

2 R03 grants (around \$145,000 each)

- Co-I of 21 R01/P01/P20/P30 grants from National Institutes of Health
- PI/contact MPI of 2 grants (\$150,000 each) from Kentucky Lung Cancer Research Program.

#### Active

**Project Title:** Statistical Methods for Cancer Progression Delineation and

Subtype Identification

**Project Number:** R03 CA259670

**Principal Investigator(s):** Wang, C

**Role in Project:** ΡI 10% Effort:

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 07/2021-06/2023

Total Award: \$149,804

**Grant Number:** R03 CA259670

**Project Title:** Characterization of Squamous Cell Lung Cancers from

Appalachian Kentucky

PO2 415 1600001032 **Project Number:** 

**Principal Investigator(s):** MPIs: Wang, C [contact]; Arnold, S; Liu, C

**Role in Project:** contact MPI

Effort: 2%

**Institution/University:** University of Kentucky

**Source of Funding:** Kentucky Lung Cancer Research Program

07/2018-06/2023 **Duration of Project:** 

**Total Award:** \$150,000

**Grant Number:** PO2 415 1600001032

**Project Title:** A Novel Genomic Approach for High-Throughput Drug Screening

**Project Number:** 

**Principal Investigator(s):** MPIs: Wang, C [contact]; Liu, C

**Role in Project:** contact MPI

no effort requested Effort: University of Kentucky **Institution/University:** 

**Source of Funding:** UK CCTS High Impact Pilot Award

**Duration of Project:** 11/2021-11/2022 **Total Award:** \$50,000 **Grant Number:** No ID

Aberrant Glycogen Modulates Cerebral Glucose Metabolism in **Project Title:** 

Aging and Alzheimer's Disease

**Project Number:** R01 AG066653

**Principal Investigator(s):** Sun. R.

**Role in Project:** Co-Investigator

**Effort:** 5%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 04/2020-03/2025

**Total Award:** \$1,912,500 **Grant Number:** R01 AG066653

**Project Title:** Targeting Translation Dependence in Colorectal Cancer

Progression

R01 CA175105 **Project Number:** 

**Principal Investigator(s):** She, OB

**Role in Project:** Co-Investigator

2% **Effort:** 

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 04/2013-11/2024

**Total Award:** \$1,816,875 **Grant Number:** R01 CA175105

**Project Title:** Non-destructive optical spectroscopic assay for high-throughput

metabolic characterization of in vitro tumor models and patient-

derived organoids

R21 EB032515 **Project Number:** 

**Principal Investigator(s):** Zhu, C

**Role in Project:** Co-Investigator

2% **Effort:** 

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 08/2022-04/2025

**Total Award:** \$600,473

**Grant Number:** R21 EB032515

**Project Title:** G Protein Signaling in Brain Injury

**Project Number:** R56 NS124707 **Principal Investigator(s):** Andres, D

**Role in Project:** Co-Investigator

**Effort:** 

**Institution/University:** University of Kentucky **Source of Funding:** NIH

**Duration of Project:** 07/2022-06/2023

**Total Award:** \$535,500

**Grant Number:** R56 NS124707

**Project Title:** Targeting Epigenetic Heterogeneity to Improve Lung Cancer

Immunotherapy Response (CII or MONC)

**Project Number:** R01 CA237643 **Principal Investigator(s):** Brainson, C **Role in Project:** Co-Investigator

2% **Effort:** 

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 07/2019-06/2024 **Total Award:** \$1,749,940 **Grant Number:** R01 CA237643

**Project Title:** Mechanistic and Pharmacologic Studies of Selective Mithramycin

Analogues Targeting EWS-FLI1 in Ewing Sarcoma

**Project Number:** R01 CA243529

**Principal Investigator(s):** MPIs: Leggas, M [contact]; Thorson, J; Tsodikov, O

**Role in Project:** Biostatistician

Effort: 1%

**Institution/University:** University of Kentucky

**Source of Funding: NIH** 

**Duration of Project:** 06/2020-05/2025 **Total Award:** \$3,010,556 R01 CA243529 **Grant Number:** 

**Project Title:** Integrin Alpha6beta4 Regulation of Cancer Epigenetics

R01 CA223164 **Project Number: Principal Investigator(s):** O'Connor, K **Role in Project:** Co-Investigator

1% **Effort:** 

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 01/2019-12/2023 **Total Award:** \$2,330,265 **Grant Number:** R01 CA223164

Ceramide and Acute Phase Proteins Elevation During Aging **Project Title:** 

**Project Number:** R01 AG019223

**Principal Investigator(s):** Nikolova-Karakashian, M

**Role in Project:** Co-Investigator

**Effort:** 

University of Kentucky **Institution/University:** 

**Source of Funding:** NIH

**Duration of Project:** 08/2002-05/2023 **Total Award:** \$2,049,176 **Grant Number:** R01 AG019223

**Project Title:** Altered Lipid Metabolism as a Novel Target for Colon Cancer

Treatment

**Project Number:** R01 CA208343

**Principal Investigator(s):** MPI: Evers, BM [contact]; Gao, T

**Role in Project:** Co-Investigator

**Effort:** 10%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 08/2017-07/2023 **Total Award:** \$2,113,800 **Grant Number:** R01 CA208343

**Project Title:** Mechanistic Impact of PI3K/mTOR Signaling on Intestinal

Homeostasis

**Project Number:** R01 DK048498 **Principal Investigator(s):** Evers, BM **Role in Project:** Co-Investigator

**Effort:** 5%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 03/1996-06/2025 **Total Award:** \$3,648,051 **Grant Number:** R01 DK048498

RNA Surveillance and Protein Translation in FTD **Project Title:** 

**Project Number:** R01 NS115507

**Principal Investigator(s):** Zhu, H

**Role in Project:** Co-Investigator

**Effort:** 3%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 07/2020-06/2025 **Total Award:** \$2,754,540 **Grant Number:** R01 NS115507

**Project Title:** RIT1-mediated Protection Following Traumatic Brain Injury

**Project Number:** R01 NS102196 **Principal Investigator(s):** Andres, D **Role in Project:** Co-Investigator

**Effort:** 2%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH **Duration of Project:** 02/2018-01/2023 **Total Award:** \$2,544,995 **Grant Number:** R01 NS102196

**Project Title:** University of Kentucky Markey Cancer Center – Cancer Center

**Support Grant** 

P30 CA177558 **Project Number: Principal Investigator(s):** Evers, BM

**Role in Project:** Faculty Biostatistician

**Effort:** 5%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 07/2013-06/2023 **Total Award:** \$10,763,550 **Grant Number:** P30 CA177558

**Project Title:** Methionine Metabolism and Lung Cancer Lineage Fate

**Project Number:** RSG-19-081-01-TBG

**Principal Investigator(s):** Brainson, C **Role in Project:** Co-Investigator

Effort: 1%

**Institution/University:** University of Kentucky **Source of Funding:** American Cancer Society

**Duration of Project:** 07/2019-06/2023

**Total Award:** \$792,000

**Grant Number:** RSG-19-081-01-TBG

**Project Title:** University of Kentucky Center for Cancer and Metabolism

**Project Number:** P20 GM121327

**Principal Investigator(s):** MPI: St. Clair, D [contact]; Zhou, B.P.

**Role in Project:** Biostatistician

**Effort:** 2.5%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 03/2017-12/2021 **Total Award:** \$11,212,250 **Grant Number:** P20 GM121327

COVID-19: Evaluation of SARS-CoV-2 Positivity, Genetic Risk **Project Title:** 

Factors & Outcomes in Patients Enrolled on TCC

**Project Number:** No ID **Principal Investigator(s):** Kolesar, J **Role in Project:** Co-Investigator

**Effort:** 2.5%

University of Kentucky **Institution/University:** 

**Source of Funding:** M2Gen

**Duration of Project:** 10/2020-10/2025 **Total Award:** \$375,512 **Grant Number:** No ID

## XIV. RESEARCH & INTELLECTUAL CONTRIBUTIONS - continued

#### C. SPONSORED RESEARCH PROJECTS, GRANT & CONTRACT ACTIVITIES continued

#### **Completed**

**Project Title:** Statistical Detection and Biochemical Classification of Cancer

Driven Mutation Patterns in Biological Networks

**Project Number:** R21 CA205778

**Principal Investigator(s):** MPI: Wang, C [contact]; Moseley, H

**Role in Project:** Contact MPI

9 % Effort:

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 08/2016-01/2019

**Total Award:** \$382,304

**Grant Number:** R21 CA205778

**Project Title:** Differential Abundance Methods for Large Heterogeneous-

Featured Metabolomics Datasets

**Project Number:** R03 CA211835

**Principal Investigator(s):** MPI: Wang, C [contact]; Flight, R

**Role in Project:** Contact MPI

**Effort:** 9 %

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 09/2016-08/2017

**Total Award:** \$145,716

**Grant Number:** R03 CA211835

Harnessing Advanced Genomic and Bioinformatics Technologies **Project Title:** 

for in-depth Molecular Characterization of Lung Adenocarcinoma

in KY

PO2 415 1400004000 1 **Project Number:** 

**Principal Investigator(s):** Wang, C **Role in Project:** Ы 4 % **Effort:** 

University of Kentucky **Institution/University:** 

**Source of Funding:** The Kentucky Lung Cancer Research Program

**Duration of Project:** 07/2015-06/2018

**Total Award:** \$150,000

**Grant Number:** PO2 415 1400004000 1

**Project Title:** A Probabilistic Model to Identify Unique Tumorigenesis Process

of Lung Cancer in Appalachian Kentucky

**Project Number:** No ID

**Principal Investigator(s):** Wang C, Arnold S, and Liu C

**Role in Project:** contact MPI

**Effort:** 0%

University of Kentucky **Institution/University:** 

UK CCTS High Impact Pilot Award **Source of Funding:** 

**Duration of Project:** 04/2019-03/2021

**Total Award:** \$37,000 **Grant Number:** No ID

**Project Title:** Development of a Model-based Bioinformatics Method for

> Comparing Somatic Mutation Patterns between Groups, with Application to Squamous Cell Lung Cancer Data in Appalachian

Kentucky

No ID **Project Number: Principal Investigator(s):** Wang, C **Role in Project:** PΙ

0 % **Effort:** 

University of Kentucky **Institution/University:** 

**Source of Funding:** UK CCTS Junior Investigator Award

**Duration of Project:** 01/2015-07/2016

**Total Award:** \$25,000 **Grant Number:** No ID

**Project Title:** Genomics in Cancer for Appalachian Kentucky

**Project Number:** No ID

**Principal Investigator(s):** MPI: Arnold, S [contact] and Wang, C

**Role in Project: MPI** 0 % **Effort:** 

**Institution/University:** University of Kentucky

UK CCTS **Source of Funding:** 

**Duration of Project:** 10/2013-03/2015

**Total Award:** \$225,000 **Grant Number:** No ID

**Project Title:** Systems Biochemistry in Lung Cancer Toward a Mechanistic

Understanding of NSCLC

P01 CA163223 **Project Number:** 

**Principal Investigator(s):** Lane, A

**Role in Project:** Co-Investigator

8 % **Effort:** 

**Institution/University:** University of Kentucky

**Source of Funding: NIH** 

**Duration of Project:** 03/2013-02/2020 **Total Award:** \$4,795,060 **Grant Number:** P01 CA163223

**Project Title:** Distinct Redox Mechanism in Normal and Cancer Cells as a Novel

Therapeutic Target

**Project Number:** R01 CA205400

**Principal Investigator(s):** MPI: St Clair, D [contact]; St Clair W

**Role in Project:** Co-Investigator

**Effort:** 10%

University of Kentucky **Institution/University:** 

**Source of Funding:** NIH

**Duration of Project:** 04/2016-03/2021 **Total Award:** \$1,697,812 **Grant Number:** R01 CA205400

**Project Title:** Peripheral Blood Exosome Lipids as Biomarkers of Disease

Activity in Crohn's Disease

R21 DK118954 **Project Number:** 

**Principal Investigator(s):** Barrett, T

**Role in Project:** Co-Investigator

**Effort:** 2%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 09/2018-08/2021

**Total Award:** \$431,478

**Grant Number:** R21 DK118954

**Project Title:** Carryover: UK – Pediatric Brain and Central Nervous System

Tumors C2417

PON2 728 1900003176 **Project Number:** 

**Principal Investigator(s):** Durbin, E. **Role in Project:** Co-Investigator

**Effort:** 5%

University of Kentucky **Institution/University:** 

**Source of Funding:** Kentucky Pediatric Cancer Research Trust Fund

07/2020-06/2021 **Duration of Project:** 

**Total Award:** \$161,806

**Grant Number:** PON2 728 1900003176

**Project Title:** A Pilot Study of Molecular Profile Differences Between Long and

Short Term Lung Cancer Survivors

PO2 415 1600001032 **Project Number:** 

**Principal Investigator(s):** Rangnekar, V **Role in Project:** Co-Investigator

**Effort:** 1%

**Institution/University:** University of Kentucky

Kentucky Lung Cancer Research Program **Source of Funding:** 

**Duration of Project:** 07/2018-06/2021

\$150,000 **Total Award:** 

**Grant Number:** PO2 415 1600001032 **Project Title:** A Novel Peptide to Inhibit Rictor-amplified Lung Tumorigenesis

**Project Number:** PO2 415 1600001032

**Principal Investigator(s):** Yang, H-S **Role in Project:** Co-I **Effort:** 1.5%

**Institution/University:** University of Kentucky

**Source of Funding:** Kentucky Lung Cancer Research Program

07/2018-06/2021 **Duration of Project:** 

**Total Award:** \$150,000

**Grant Number:** PO2 415 1600001032

**Project Title:** A Phase I Dose Escalation Study on the Safety of Lapatinib with

Dose-Dense Paclitaxel in Patients with Platinum-Resistant Ovarian

Cancer

**Project Number:** P30 CA177558 Pilot Project

**Principal Investigator(s):** Ueland, F **Role in Project:** Biostatistician

Effort: 1%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 07/2020-06/2021

**Total Award:** \$68,550

**Grant Number:** P30 CA177558 Pilot Project

**Project Title:** Utility of ctDNA in Personalized Therapy for Non-Small-Cell

Lung Cancer

**Project Number:** PO2 415 1400004000 1

**Principal Investigator(s):** Kolesar, J **Role in Project:** Co-I **Effort:** 1%

**Institution/University:** University of Kentucky

**Source of Funding:** Kentucky Lung Cancer Research Program

**Duration of Project:** 07/2017-06/2021

**Total Award:** \$150,000

**Grant Number:** PO2 415 1600001032

**Project Title:** Natural Product-Based Modulators of 4E-BP1 Phosphorylation

**Project Number:** R01 CA203257

**Principal Investigator(s):** MPI: Thorson, J; She, Q-B

**Role in Project:** Co-Investigator

**Effort:** 1%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 04/2016-03/2021 **Total Award:** \$1,933,963 **Grant Number:** R01 CA203257

**Project Title:** Selection of Personalized Cancer Therapies by Evaluating Intra-

tumoral Heterogeneity and Phylogentic Analysis

**Project Number:** No ID **Principal Investigator(s):** Kolesar, J **Role in Project:** Co-Investigator

**Effort:** 0 %

**Institution/University:** University of Kentucky

CCSG pilot grant **Source of Funding: Duration of Project:** 07/2018-06/2019

**Total Award:** \$50,000 **Grant Number:** No ID

**Project Title:** Latexin function in the maintenance and regeneration of the

hematopoietic system

**Project Number:** R01 HL124015

**Principal Investigator(s):** Liang, Y

**Role in Project:** Co-Investigator

Effort: 5 %

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 01/2016-12/2020 **Total Award:** \$1,250,000 R01 HL124015 **Grant Number:** 

**Project Title:** Novel pRNA Nanoparticle Delivery as Directed Therapy for

Colorectal Cancer Metastasis

R01 CA195573 **Project Number:** 

**Principal Investigator(s):** MPI: Evers, BM [contact]; Guo, P; Thorson, J

**Role in Project:** Biostatistician

5 % **Effort:** 

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 09/2015-08/2020 **Total Award:** \$1,384,065

**Grant Number:** R01 CA195573

Regulation of Snail in breast cancer progression and metastasis **Project Title:** 

**Project Number:** R01 CA125454

**Principal Investigator(s):** Zhou, B

**Role in Project:** Co-Investigator

5 % (effort only in 2016-2018) **Effort:** 

University of Kentucky **Institution/University:** 

**Source of Funding:** NIH

**Duration of Project:** 12/2006-04/2018

**Total Award:** \$921,710 **Grant Number:** R01 CA125454

**Project Title:** Role of Tcl1 and par-4 in regulation of chronic lymphocytic

leukemia

R01 CA165469 **Project Number:** 

**Principal Investigator(s):** MPI: Bondada, S; Rangnekar V

**Role in Project:** Co-Investigator

**Effort:** 3 %

**Institution/University:** University of Kentucky

**Source of Funding:** 

**Duration of Project:** 02/2013-01/2018 **Total Award:** \$1,386,095 **Grant Number:** R01 CA165469

**Project Title:** A role for c-Abl/Arg in Melanoma Progression

**Project Number:** R01 CA166499 **Principal Investigator(s):** Plattner, R **Role in Project:** Co-Investigator

**Effort:** 5 %

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 09/2012-08/2017 **Total Award:** \$1,062,435 **Grant Number:** R01 CA166499

**Project Title:** Assessing the Mechanism of Drug Resistance in Lung Cancer

PO2 415 1600001032 **Project Number:** 

**Principal Investigator(s):** Kolesar, J **Role in Project:** Co-Investigator

**Effort:** 1%

**Institution/University:** University of Kentucky

**Source of Funding:** Kentucky Lung Cancer Research Program

07/2018-06/2022 **Duration of Project:** 

\$150,000 **Total Award:** 

**Grant Number:** PO2 415 1600001032

Genomic Architecture of a Key Alzheimer's Disease mimic: **Project Title:** 

**CARTS** 

R56 AG057191 **Project Number:** 

**Principal Investigator(s):** Fardo, D

**Role in Project:** Co-Investigator

**Effort:** 5%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 05/2019-02/2022

**Total Award:** \$1,079,062 **Grant Number:** R56 AG057191

**Project Title:** Targeting Epigenetic Heterogeneity to Improve Lung Cancer

Immunotherapy Response (CII or MONC)

R21 AA026787 **Project Number:** 

**Principal Investigator(s):** Chen, G

**Role in Project:** Co-Investigator

**Effort:** 2.5%

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 02/2019-01/2022

**Total Award:** \$401,626

**Grant Number:** R21 AA026787

**Project Title:** Roles of RORalpha in Breast Cancer Development and

Progression

R01 CA215095 **Project Number:** 

**Principal Investigator(s):** Xu, R

**Role in Project:** Co-Investigator 5% Yrs 3 and 5 only **Effort:** University of Kentucky **Institution/University:** 

NIH **Source of Funding:** 

**Duration of Project:** 07/2017-06/2022 **Total Award:** \$1,726,893 **Grant Number:** R01 CA215095

**Project Title:** Identification of Lung Adenocarcinoma Subtypes using

Radiogenomics and Deep Learning

**Project Number:** PO2 415 1600001032

**Principal Investigator(s):** Chen, J **Role in Project:** Co-I **Effort:** 1%

**Institution/University:** University of Kentucky

Kentucky Lung Cancer Research Program **Source of Funding:** 

**Duration of Project:** 07/2018-06/2022

**Total Award:** \$150,000

**Grant Number:** PO2 415 1600001032

**Project Title:** Translational Control in CR(VI) Carcinogenesis

R21 ES031712 **Project Number:** 

**Principal Investigator(s):** She, QB

**Role in Project:** Co-Investigator 1% Yr1, 2% Yr2 **Effort:** 

**Institution/University:** University of Kentucky

**Source of Funding:** NIH

**Duration of Project:** 06/2020-05/2022

\$688,500 **Total Award:** 

**Grant Number:** R21 ES031712

**Project Title:** Germline and Environmental Factors Associated with Pediatric

Brain and Central Nervous System Tumors in Kentucky

**Project Number:** No ID **Principal Investigator(s):** Durbin, E

**Role in Project:** Co-Investigator

**Effort:** 5%

**Institution/University:** University of Kentucky

Pediatric Cancer Research Trust Fund **Source of Funding:** 

**Duration of Project:** 07/2020-06/2022

**Total Award:** \$499,715 **Grant Number:** No ID

**Project Title:** Novel Antibody-Enzyme Fusion Therapy Targeting Ewing's

Sarcoma

No ID **Project Number: Principal Investigator(s):** Sun, R

**Role in Project:** Co-Investigator

**Effort:** 1%

**Institution/University:** University of Kentucky

**Source of Funding:** V Foundation **Duration of Project:** 07/2020-06/2022

**Total Award:** \$200,000 **Grant Number:** No ID

#### XIV. RESEARCH & INTELLECTUAL CONTRIBUTIONS - continued

**E. OTHER CREATIVE ACTIVITIES** [oldest at top, newest at bottom; include innovative materials, clinical protocols, institutional packages, modules, computer programs, innovative teaching materials, patented and copyrighted intellectual property; describe where work used and by whom]

#### **Software**

bacr: an R package that implements the Bayesian Adjustment for Confounding (BAC) method for estimating the average causal effect of a treatment on an outcome from cohort studies. The software package is available at CRAN.

NanoStringDiff: an R package to perform differential expression analysis based on gene expression data generated from the NanoString nCounter system. The software package is available at Bioconductor.

**SDAMS**: an R package that implements a semiparametric method for differential abundance/expression analysis of proteomic, metabolomic and scRNAseq data. The software package is available at Bioconductor.

**PATOPA**: a bioinformatics software to delineate the temporal order of driver mutations during carcinogenesis by leveraging functional annotation and pathway information. The software is available at GitHub.

**DASEV**: an R package that implements a two-part model with Bayesian shrinkage estimation of variance for differential abundance analysis of proteomic and metabolomic data. The software package is available at http://sweb.uky.edu/~cwa236/DASEV.html.

MEScan: a bioinformatics software to identify cancer driver mutations by genome-wide screen of mutually exclusive mutation patterns. The software is available at GitHub.

**XV. OTHER ACTIVITIES** [oldest at top, newest at bottom; writing board examinations, curricular design committees]

Not applicable.

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