Curriculum Vitae

Yen-Yi Ho

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Education

2009 Ph.D., Biostatistics, Johns Hopkins University,

Thesis Title: Gene Association Networks and Higher-Order Interactions:

Algorithms and Statistical Models.

Committee: Dr. Giovanni Parmigiani (Advisor), Dr. Leslie Cope,

Dr. Pien-Chien Huang, Dr. Thomas Louis

2001 M.S., Epidemiology, National Taiwan University

1999 B.S., Double major in Medical Technology & Public Health, National Taiwan University

Employment

July 2011 –	Faculty, Biostatistics and Bioinformatics Shared Resource
	Masonic Cancer Center, University of Minnesota

- July 2011 Assistant Professor, Division of Biostatistics, School of Public Health, University of Minnesota
- 2009 2011 Postdoctoral Research Fellow

McKusic-Nathans Institute of Genetic Medicine Johns Hopkins University School of Medicine

- 2006 2007 Statistical Consultant, Biostatistics Consulting Center Department of Biostatistics, Johns Hopkins University
- 2005 2009 Statistical Consultant, School of Nursing Johns Hopkins University
- 2001 2003 Research Associate, Bioresource Collection and Research Center Hsinchu, Taiwan

Honors and Awards

2003 – 2005 Government Funded Scholarship for Studying Abroad,

Taiwanese Ministry of Eduction

1999 Research Creativity Award, National Science Council, Taiwan

Awarded for outstanding undergraduate research

Teaching Experience

2011 – Member, Statistical Genetics and Bioinformatics Curriculum Committee

Division of Biostatistics, University of Minnesota

Fall 2012 – Present Co-Instructor, Statistics for Human Genetics and Molecular Biology (PUBH 7445)

Division of Biostatistics, University of Minnesota

Professional Memberships and Service

American Statistical Association Institute of Mathematical Statistics Referee, American Journal of Epidemiology Referee, Turkish Journal of Medical Sciences

Software

- 1. Ho, Y.-Y. (2009). LiquidAssociation: R/Bioconductor package for estimating liquid association using the conditional normal model. Available at http://www.bioconductor.org
- 2. Gunderson T.* (2014). fastLiquidAssociation: R/Bioconductor package for exploring liquid association on a genome-wide scale. Available at http://www.bioconductor.org. * Package was developed under Ho's supervision as the author's thesis advisor.

Publications

- 1. Chen, W.J., Liu P.-H., **Ho, Y.-Y.**, and Chien, K.-L., et al. (2003). Sibling recurrence risk ratio analysis of the metabolic syndrome and its components over time. *BMC Genetics* 4, S33-S28.
- 2. Lee, W.-C., Ho, Y.-Y. (2003). Potential for gene-gene confounding bias in case-parental control studies. *Annals of Epidemiology* 13, 261-266.
- **3. Ho, Y.-Y.**, Parmigiani, G., Louis, T.A., Cope, L.M. (2010). Modeling Liquid Association. *Biometrics* **67**, 133-141. doi: 10.1111/j.1541-0420.2010.01440.x.
- **4.** Jiang, Q., **Ho, Y.-Y.**, Hao L, Nichols Berrios C, Chakravarti, A. (2011). Copy number variants in candidate genes are genetic modifiers of Hirschsprung disease. *PLoS One* **6**, e21219.

5. Ho, Y.-Y., Matteini A.M., Beamer B., and Fried L., et al. (2011). Exploring biologically relevant pathways in frailty. *Journal of Gerontology A Biological Sciences and Medical Sciences* **66**, 975-979.

- 6. Shen A., Baker J., Scott G., Davis Y., Ho Y.-Y., Siliciano R. (2013). Endothelial Cell Stimulation Overcomes Restriction and Promotes Productive and Latent HIV-1 Infection of Resting CD4+ T Cells. *Journal of Virology* 87, 9768-79. doi: 10.1128/JVI.01478-13.
- 7. Terrell A.N., Huynh M., Grill A., Kovi R.C., O'Sullivan M.G., Guttenplan J.B., Ho, Y.-Y., Peterson L.A. (2014). Mutagenic activity of furan in female Big Blue B6C3F1 mice. Mutation Research/Genetic Toxicology and Environmental Mutagenesis 770, 46-54.
- 8. Ho, Y.-Y., Cope, L.M., Parmigiani, G. (2014). Modular network construction using eQTL data: an analysis of computational costs and benefits. Frontiers in Genetics 5, 40. doi: 10.3389/fgene.2014.00040
- 9. Ho, Y.-Y., Baechler E.C., Ortmann W., Behrens T.W., Graham R.R., Bhangale T.R., Pan W. (2014). Using Gene Expression to Improve the Power of Genome-Wide Association Analysis. *Human Heredity* 78, 94-103. doi: 10.1159/000362837
- 10. Abbott, K., Nyre, E., Abrahante, J., Ho, Y.-Y., Isaksson, R.V., Starr, T. (2014) The Candidate Cancer Gene Database: a database of cancer driver genes from forward genetic screens in mice. Nucleic Acids Research 43, D844-8. doi: 10.1093/nar/gku770.
- 11. Gunderson T.*, Ho, Y.-Y.* (2014) An efficient algorithm to explore liquid association on a genome-wide scale. *BMC Bioinformatics* 15, 371. *Ho was the corresponding author and the first author's thesis advisor.
- **12.** Gupta M., McCauley J., Farkas A., Gudeloglu A., Neuberger M.M., **Ho Y.-Y.**, Yeung L., Vieweg J, Dahm P. (2014) Clinical Practice Guidelines on Prostate Cancer: A Critical Appraisal. *The Journal of Urology* pii: S0022-5347(14)04811-3. doi: 10.1016/j.juro.2014.10.105.
- 13. Nho R., Im J., Ho, Y.-Y., Hergert P. (2014) MicroRNA-96 inhibits FoxO3a function in IPF fibroblasts on type I collagen matrix. *American Journal of Physiology-Lung Cellular and Molecular Physiology* 307, L632-42. doi: 10.1152/aj-plung.00127.2014.
- **14.** Gavin K., Linde J.A., Pacanowski C.R., French S.A., Jeffery R.W., **Ho Y.-Y.** (2015) Weighing frequency among working adults: cross-sectional analysis of two community samples. *Preventive Medicine Reports* **2**, 44-46. doi: 10.1016/j.pmedr.2014.12.005.
- 15. Ho, Y.-Y., O'Connell M., Guan W., Basu S. (2015) Powerful Association Test Combining Rare Variant and Gene Expression Using Family Data from Genetic

Analysis Workshop 19. Genetic Analysis Workshop 19 Proceedings 2015;9 Suppl 8:S33.

Book Chapter

1. Ho, Y.-Y., Cope, L., Dettling, M., and Parmigiani, G. (2007). Statistical methods for identifying differentially expressed gene combinations. *Methods in Molecular Biology* 408, 171-191.

Submitted

- 1. Ho, Y.-Y., LaRue R.S., Largaespada D.A. Individual-oriented gene set analysis using insertional mutation data. [Under review]
- 2. Fagan D.H., Fettig L.M., Avdulov S., Peterson M.S., **Ho, Y.-Y.**, Polunovsky V.A, and Yee D. Acquired tamoxifen resistance in MCF-7 breast cancer cells requires hyperactivation of eIF4F-mediated translation. [Under review]
- **3. Ho, Y.-Y.***, Vo T.N.*, Chu H., LeSage M.G., Luo X., Le C.T. A Bayesian hierarchical model for estimating the demand curve. [Under review] *These authors contributed equally to this work
- **4.** Abbott K, **Ho**, **Y.-Y.**, Erickson J.. Automatic health record review to identify gravely ill Social Security disability applicants. [Under review]
- 5. Arsoniadis E.G., **Ho, Y.-Y.**, Melton G.B, Madoff R.D., Le C., Kwaan M. African Americans and Short Term Outcomes After Surgery for Crohns Disease: An ACS-NSQIP Analysis. [Under review]

Presentations

Oral Presentation

- 1. Statistical Methods for identifying differentially expressed gene combinations (2007), International Biometric Society Eastern North American Region (ENAR).
- 2. Exploring Liquid Association in Non-Gaussian Multivariate Distributions (2008), Joint Statistical Meetings.
- **3.** nPARS: A Comprehensive Search Algorithm for Constructing Bayesian Networks Using Large-Scale Genomic Data (2011), Joint Statistical Meetings, Miami.
- 4. Using Gene Expression to Improve the Power of Genome-Wide Association Analysis (2014). International Biometric Society Eastern North American Region (ENAR).
- 5. Ho, Y.-Y., O'Connell M., Guan W., Basu S. (2015) Powerful Association Test Combining Rare Variant and Gene Expression Using Family Data from Genetic Analysis Workshop 19. Genetic Analysis Workshop 19, Vienna, Austria [Presented by O'Connell M]

Grants

1. Models for Tobacco Products Evaluation

Principal investigator: Dorothy Hatsukami, PhD

Mechanism: U19 (NIH/FDA)

2. Transdisciplinary Collaborative Center for Research African American Mens Health

Principal investigator: Selwyn Vickers, PhD

Mechanism: U54

3. Constituent Yields and Biomarkers of Exposure for Tobacco Product Regulation

Principal Investigator: Irina Stepanov, PhD

Mechanism: R01

4. Inducing NK cells to remember and fight cancer

Principal Investigator: Jeff Miller, MD

Mechanism: R01

5. Minnesota Obesity Center

Principal Investigator: Allen Levine, PhD

Mechanism: P30

6. Interactions between tobacco smoke constituents in rodent tumor models

Principal Investigator: Lisa Peterson, PhD

Mechanism: R01

7. Molecular mechanisms of leukemia stem cell persistence in AML relapse

Principal Investigator: Zohar Sachs, PhD

Mechanism: CTSI/KL2

8. Integrating Somatic Mutation and Gene Expression Data to Identify Active Driver Pathways Associated with Cancer Survival

Principal Investigator Yen-Yi Ho, PhD

Mechanism: Grant-In-Aid, Office of the Vice President for Research, University of Minnesota