


Name	Mikyeong Lee	
Country	USA	
Organization	National Institutes of Health/National Institute of Environmental Health Sciences (NIH/NIEHS)	
Current Position	Staff Scientist	

Educational Background

2011 PhD in Public Health, Graduate School of Public Health, Seoul National University, South Korea
 2005 MPH in Public Health, Graduate School of Public Health, Seoul National University, South Korea
 2003 BS in Mathematics and Statistics, Pusan National University, South Korea

Professional Experiences

2016 – current: Genomics and the Environment in Respiratory and Allergic Health Group, Epidemiology Branch, NIEHS
 2016: Research Professor, Institute of Medical Science, Kangwon National University, South Korea
 2011-2015: Research Assistant Professor, Institute of Health and Environment, Seoul National University, South Korea
 2008-2009: Senior Researcher, Genomic Medicine Institute, Seoul National University and PSOMA Therapeutics, Inc., South Korea
 2005-2007: Researcher, Genomic Medicine Institute, Seoul National University and Macrogen, Inc., South Korea

Professional Organizations

American Thoracic Society
 American Society of Human Genetics
 International Genetic Epidemiology Society

Main Scientific Publications

Lee *et al.* Pulmonary Function and Blood DNA Methylation: A Multiancestry Epigenome-Wide Association Meta-analysis. *Am J Respir Crit Care Med.* 2022
 Lee *et al.* House dust microbiota in relation to adult asthma and atopy in a US farming population. *J Allergy Clin Immunol.* 2021
 Lee *et al.* Genome-wide DNA methylation and long-term ambient air pollution exposure in Korean adults. *Clin Epigenetics.* 2019
 Reese *et al.* Epigenome-wide meta-analysis of DNA methylation and childhood asthma. *J Allergy Clin Immunol.* 2019
 Lee *et al.* Exposures Related to House Dust Microbiota in a U.S. Farming Population. *Environ Health Perspect.* 2018
 Wyss *et al.* Multiethnic meta-analysis identifies ancestry-specific and cross-ancestry loci for pulmonary function. *Nat Commun.* 2018
 Lee *et al.* Epigenome-wide association study of chronic obstructive pulmonary disease and lung function in Koreans. *Epigenomics.* 2017
 Lee *et al.* DNA methylation and smoking in Korean adults: epigenome-wide association study. *Clin Epigenetics.* 2016
 Kim *et al.* Genome-wide association studies identify locus on 6p21 influencing lung function in the Korean population. *Respirology.* 2014