Bio:

Cen Wu is an associate professor in statistics at Kansas State University. His current research has mainly been motivated by data contamination and heavy-tailed distributions that widely exist in disease phenotypes and multi-level omics measurements from cancers and other complex diseases. Tackling these problems in a high dimensional setting demands robust variable selection methods, within both the frequentist and Bayesian frameworks. His statistical methodological work includes Bayesian sparse learning, high/ultra-high dimensional robust variable selection, and integrative analysis of cancer genomics data from multiple platforms.