

A general kernel boosting framework to integrate pathways for cancer genomic data analysis

Li Zeng¹, Zhaolong Yu², Yiliang Zhang¹ and Hongyu Zhao^{1,2,3,*}

¹ Department of Biostatistics, Yale University, New Haven, CT
06511, USA

² Interdepartmental Program in Computational Biology and
Bioinformatics, Yale University, New Haven, CT 06511, USA

Abstract

Keywords:

1 Introduction