

“ I mainly use IPA, which has helped make sense of large gene expression data sets. This pathway analysis makes my data presentable and understandable.

— Alexa Ciminera, Graduate Student, City of Hope

Source: Alexa Ciminera, Graduate Student, City of Hope



TechValidate
by SurveyMonkey

✓ Validated

Published: Mar. 3, 2019 TVID: DF1-737-EF6