★★★★★

5/5 Stars

Please share details on your experience with it.

> " I regularly use IPA's canonical pathway enrichment, upstream regulator, and causal network analysis with both bulk and single cell RNA-Seq data. It's absolutely awesome since I typically work with data from a wide range of immune cell types.

Would you recommend our software to others, and if so, why?

> " I definitely recommend your software since it's very practical to use and provides in-depth analysis of genomic data using manually curated biological findings from the literature.

Source: Bioinformaticist, Federal Government

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