

A word cloud visualization showing terms related to bioinformatics workflows. The most prominent words are "author", "analysis", "gene", "section", "result", "Figure", "data", "tool", "workflow", "paper", "Table", "version", "article", "used", "may", "might", "cluster", "page", "reference", "seem", "will", "app", "useful", "using", "RNA seq", "pipeline", "provide", "quite", "one", "example", "read", "available", "biological", "make", "text", "Reviewer response", "many", "set", "cell", "file", "training", "better", "number", "information", "different method", "work", "refer", "might", "cluster", "page", "reference", "seem", "will", "app".