Reproducing a single-cell proteomics data analysis using QFeatures and scp

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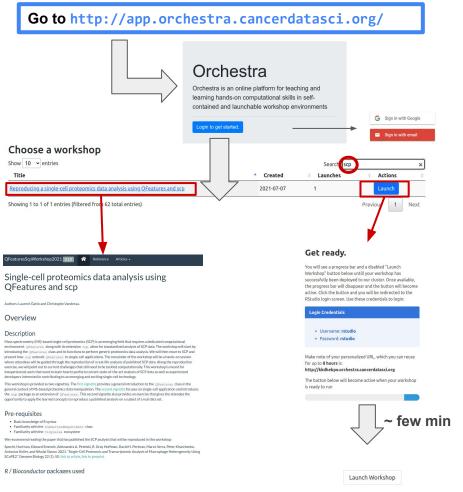
@lgatt0



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