





















Support Status for Scalable, Out-of-core Dask-based Analysis

 = released	 = awaiting release
 = PR open	 = not begun

	scanpy (CPU, multi-core, lazy)	rapids-single-cell (multi-GPU, lazy)
i/o (anndata)		
QC Metrics		
Scale, Filter, log1p, normalize		
highly_variable_genes (non-seurat v3 flavors)		
highly_variable_genes (seurat v3 flavours)		
Dense PCA		
Sparse PCA		
UMAP		
Leiden	