**Table 2d:** Single-cell data derived from other species

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Title** | **Algorithm** | **Species** | **Tissue** | **# Cells** | **SRA/GEO** | **Reference** |
| **Worm neuron cells**1 | scDeepCluster | C. elegans | Neuron | 4,186 | GSE98561 | [156] |
| **Cross species, stimulation with LPS and dsRNA** | scGen | Mouse, rat, rabbit, and pig | bone marrow-derived phagocyte | 5,000 to 10,000 /species | 13 accessions in ArrayExpress | [118] |

1. Processed data is available at <https://github.com/ttgump/scDeepCluster/tree/master/scRNA-seq%20data>