Figure 6. Comparison between bulk transcriptomic data and single cell pig data. A) Alluvial plot comparing the assigned specificity categories based on bulk data and single cell data. Note that the classification for the bulk data was calculated solely on data composed of brain, retina, kidney, liver, adipose tissue, large intestine, lung and lymphoid tissue data. **B)** Hypergeometric test, based on matching elevated genes between tissues and cell types. **C)** Comparison between Tau scores computed using both datasets.