$E:/AD_PD_HD_MS_computation_9-14-19/HuAgeDis_MSNDsplitRnaseq (recoded) _v7.71_minMod100 (7-18-19) _Human_Gene_Atlas_Gene_Targets_Comparison (recoded) _v7.71_minMod100 (recoded) _v7.7$ 150 **-**100 -Number of genes overlap P.value 0.002 0.001 50 **-**BDCA4+_DendriticCells -CD14+_Monocytes -CD19+_BCells(neg._sel.) -CD33+_Myeloid 7 CD56+_NKCells WholeBlood -CD8+_Tcells -Module names based on data from enrichR