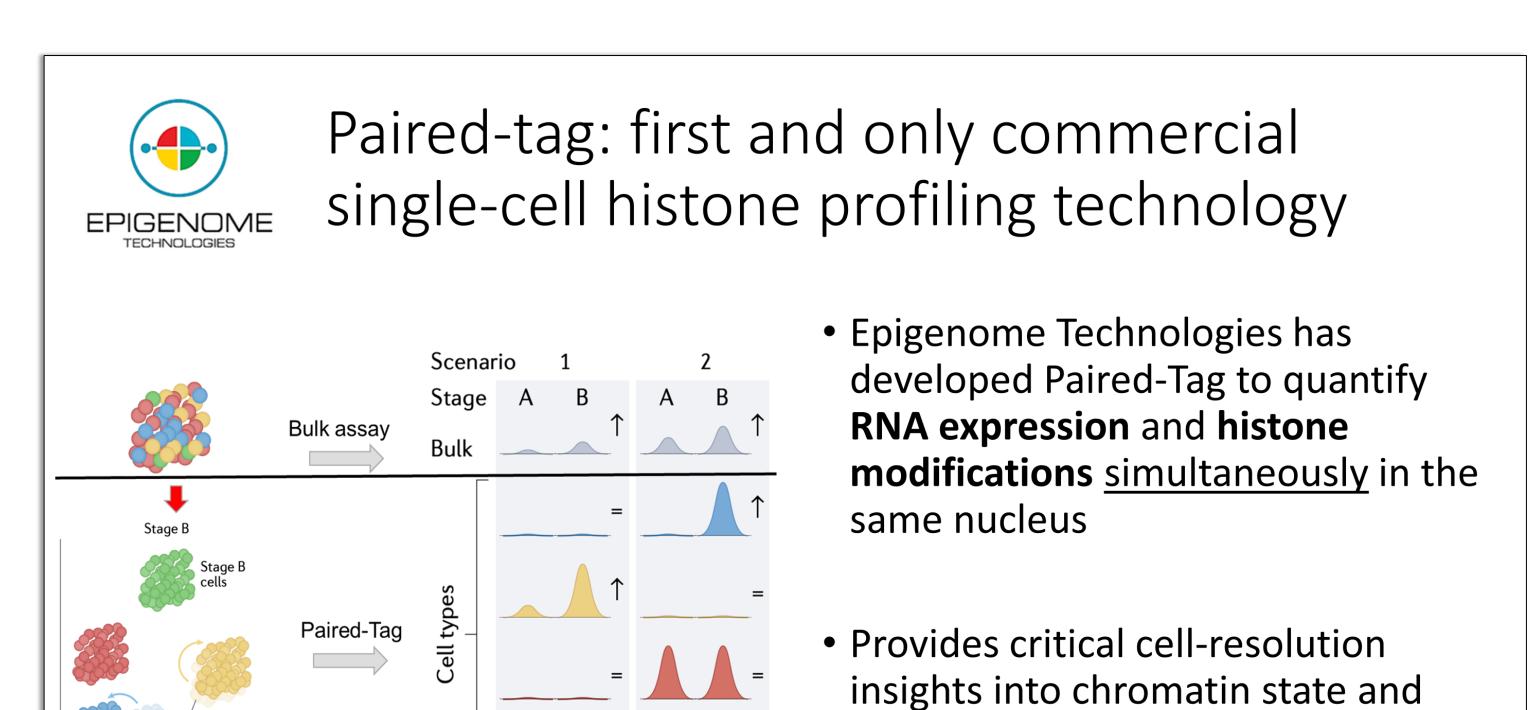
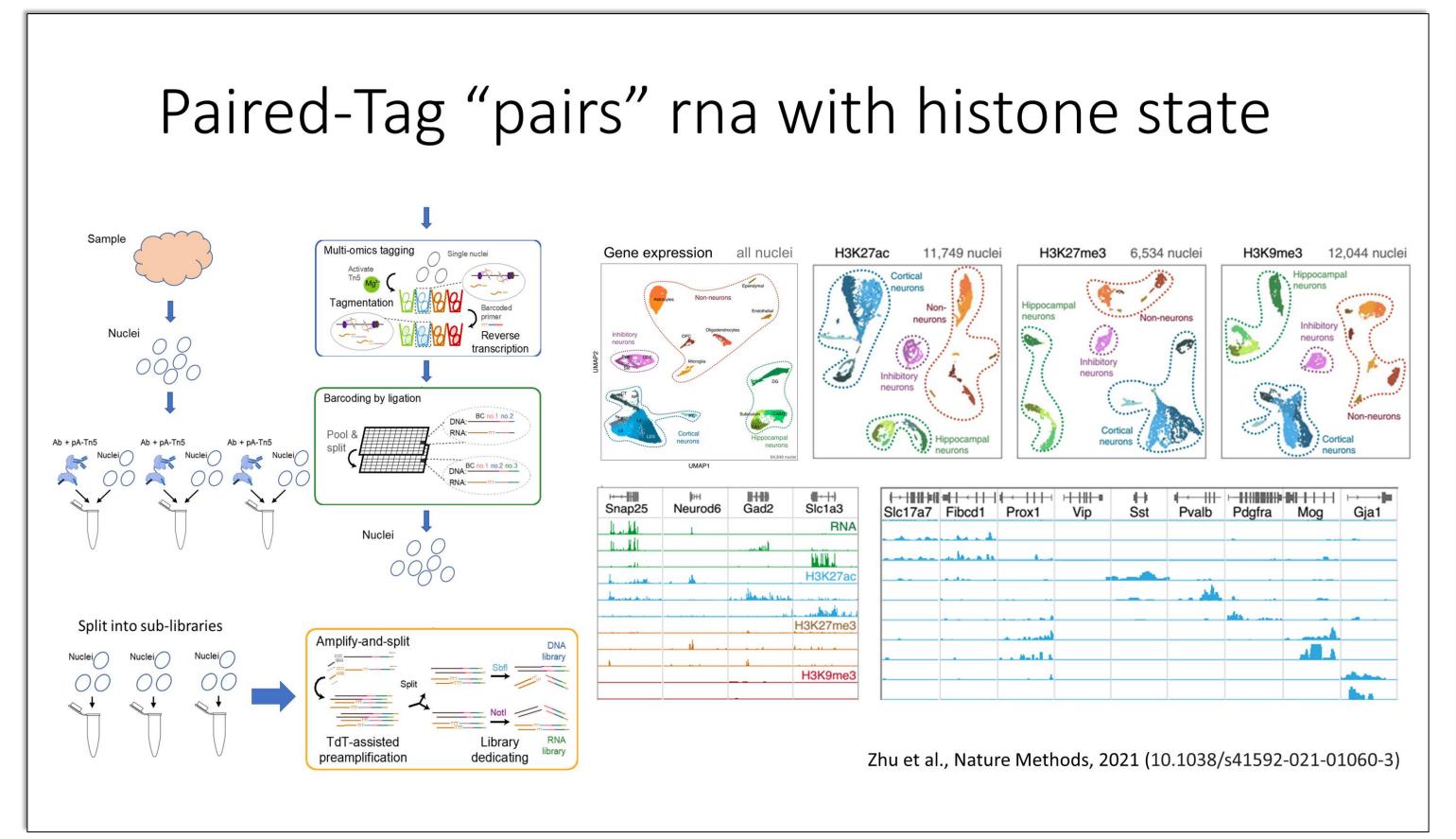
## Single-cell joint analysis of histone modification state and gene expression

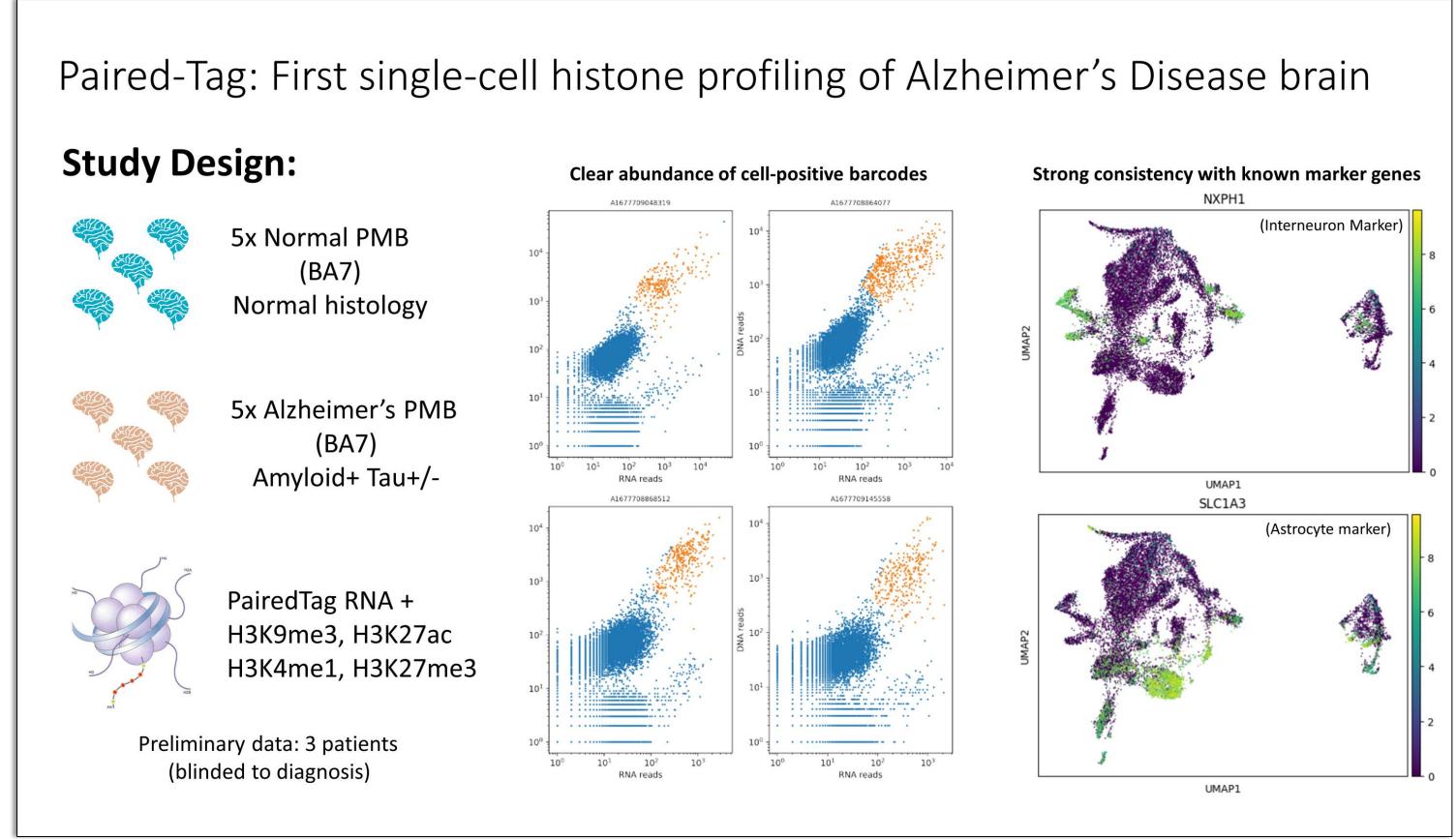
Christopher Hartl, Haruhiko Ishii, Hunki Lim, Pei Lin, Epigenome Technologies, Inc., San Diego, CA www.epigenome.us

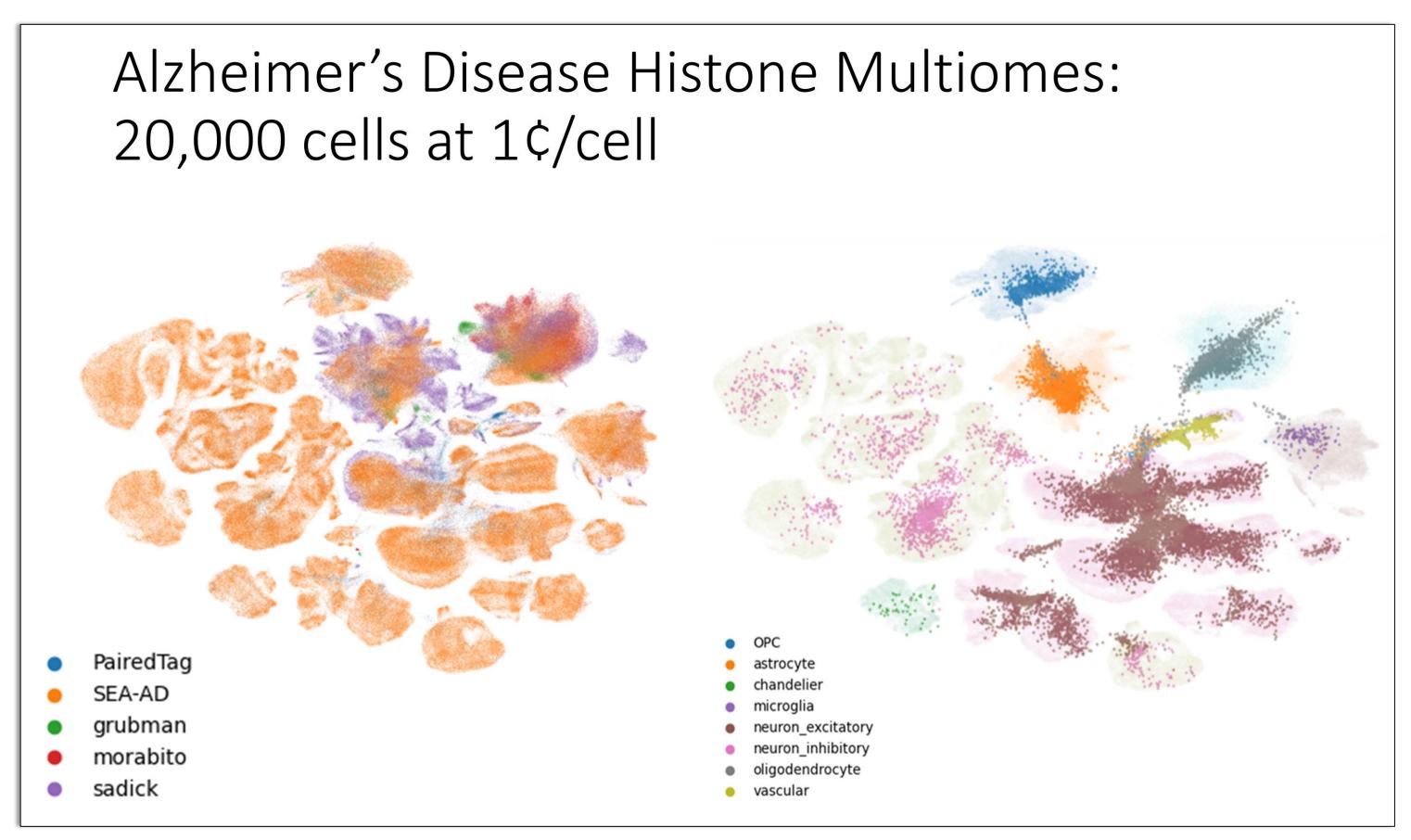


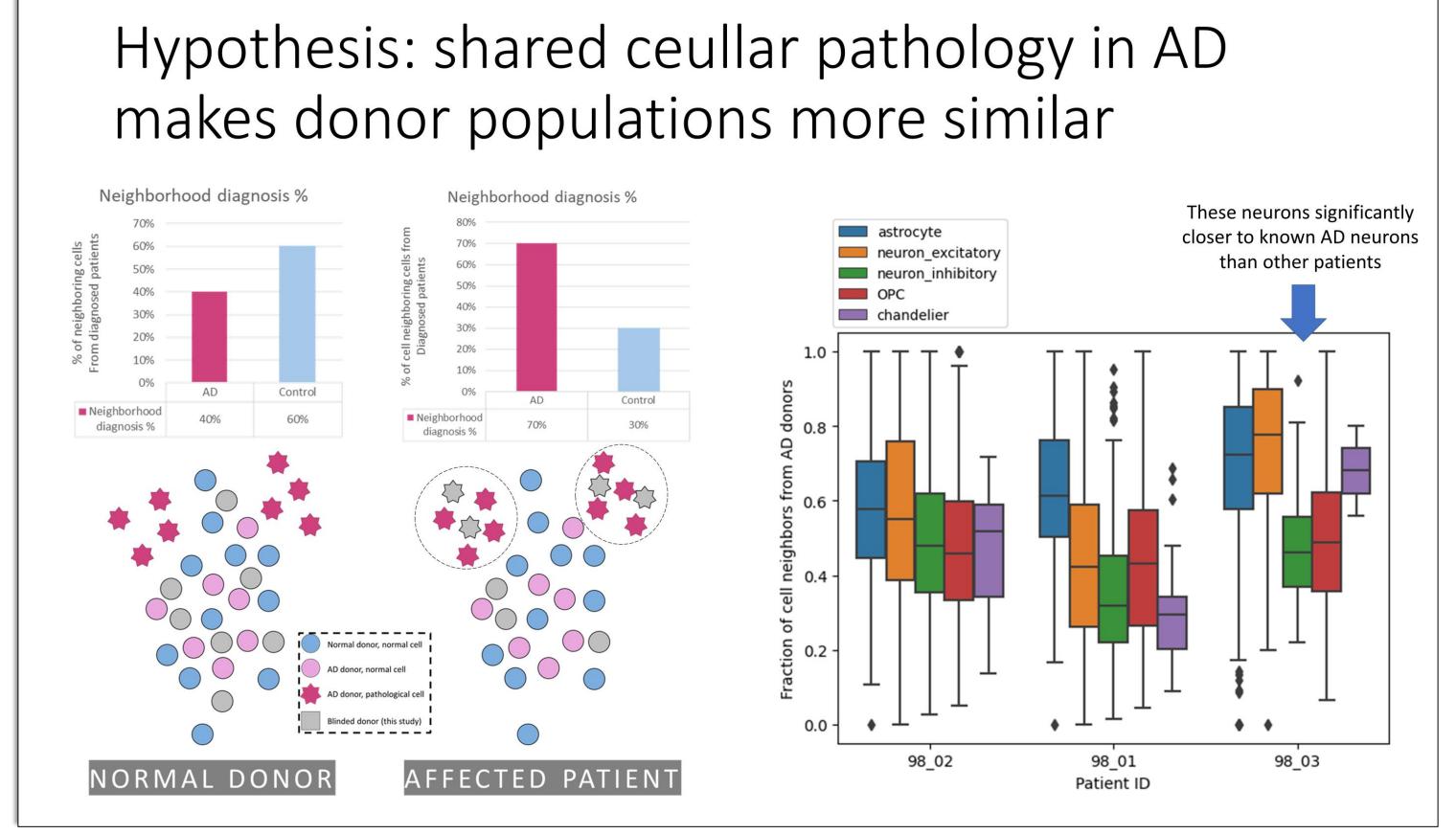
DNA-bound proteins such as

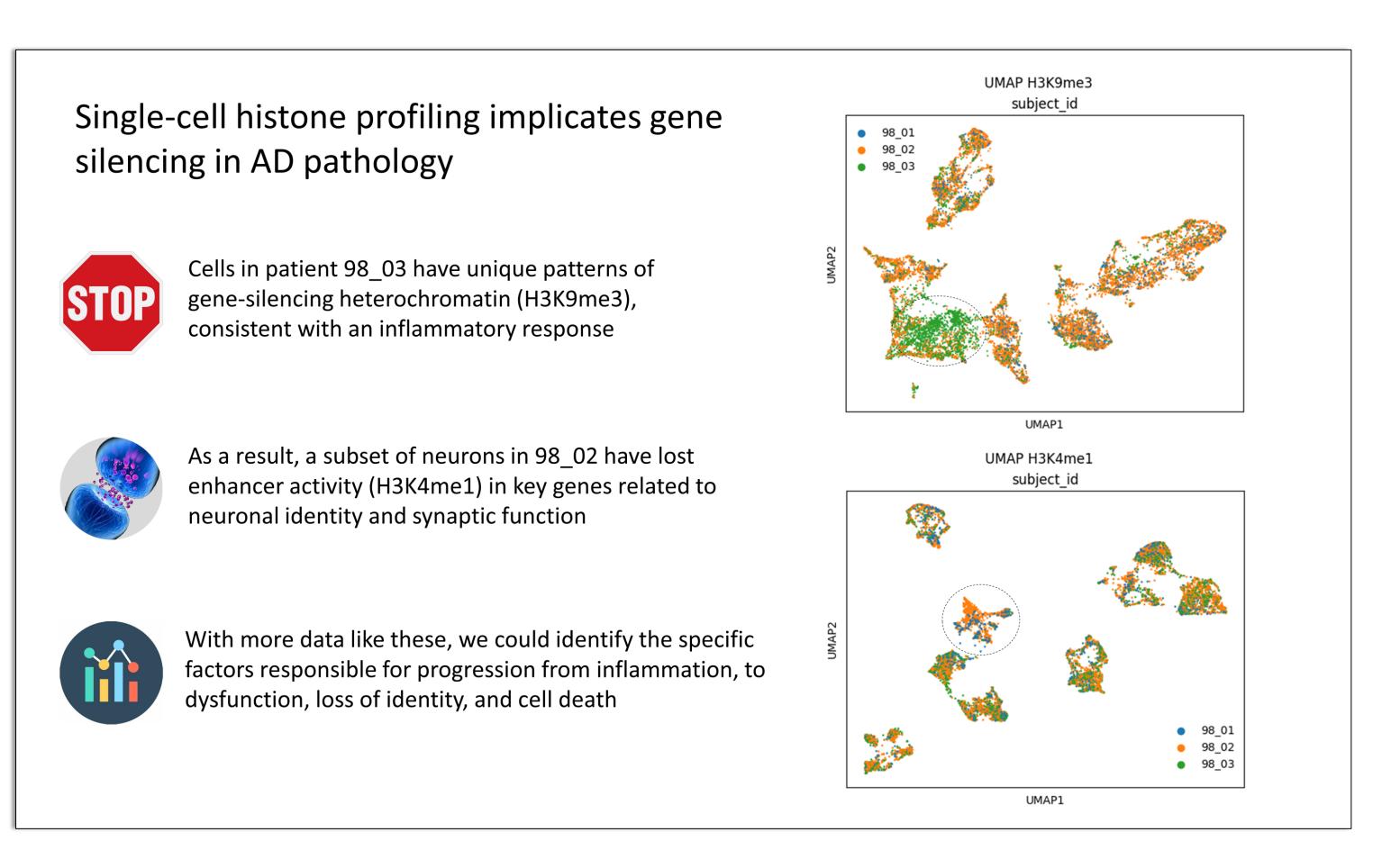
modified histones











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