



Suppl. Fig. 7. Tracking co-evolution of somatic nuclear and mitochondrial DNA mutations in CLL and AML using single cell DNA and protein sequencing (Tapestri). A Pseudobulk coverage profile of the mitochondrial genome using mitochondrial DNA (mtDNA) panel for scDNAseq. B Expression of CD3, CD4 and CD8 (T cells), CD56 (NK cells), CD19 and CD5 (CLL), CD33 and CD14 (monocytes) in CLL4 (n=5,596).