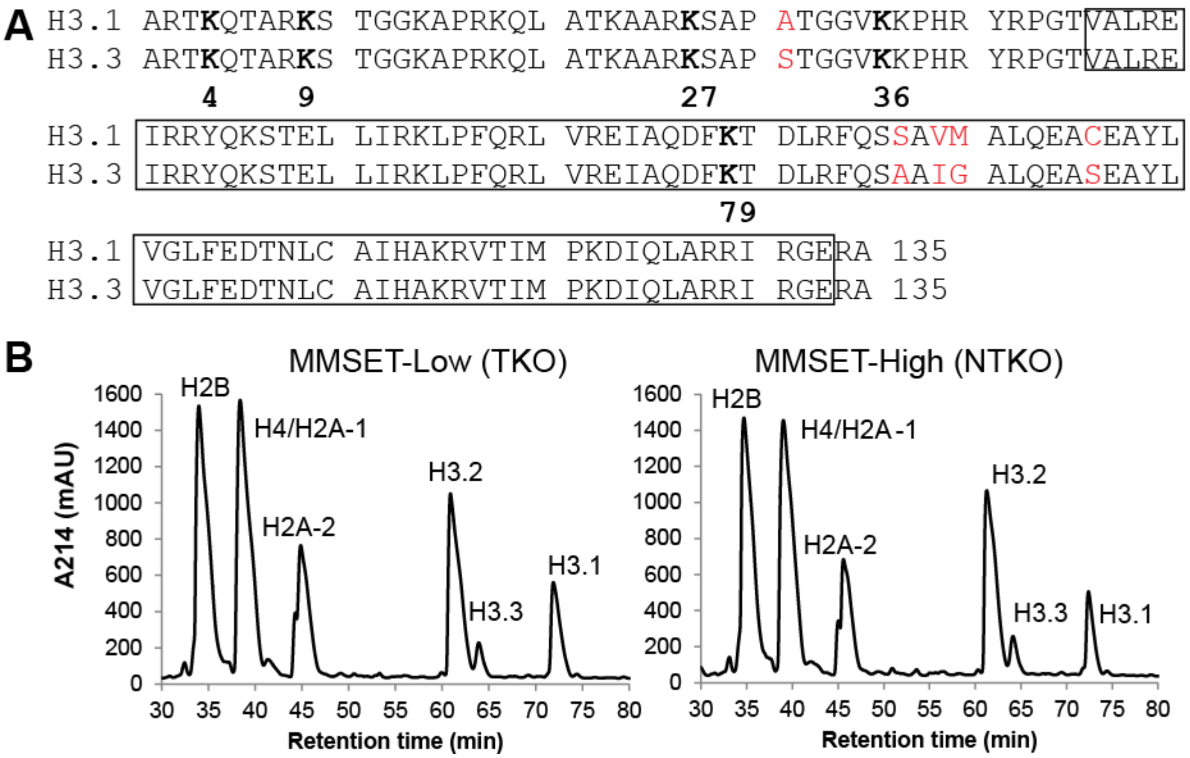

Fig S1. (A). The alignment of H3.1 and H3.3 variants. The different residues are labeled as red and the globular domain is indicated within a box. (B) UV trace of reverse phase HPLC separation of acid extracted total histones from MMSET-Low and MMSET-High cells.