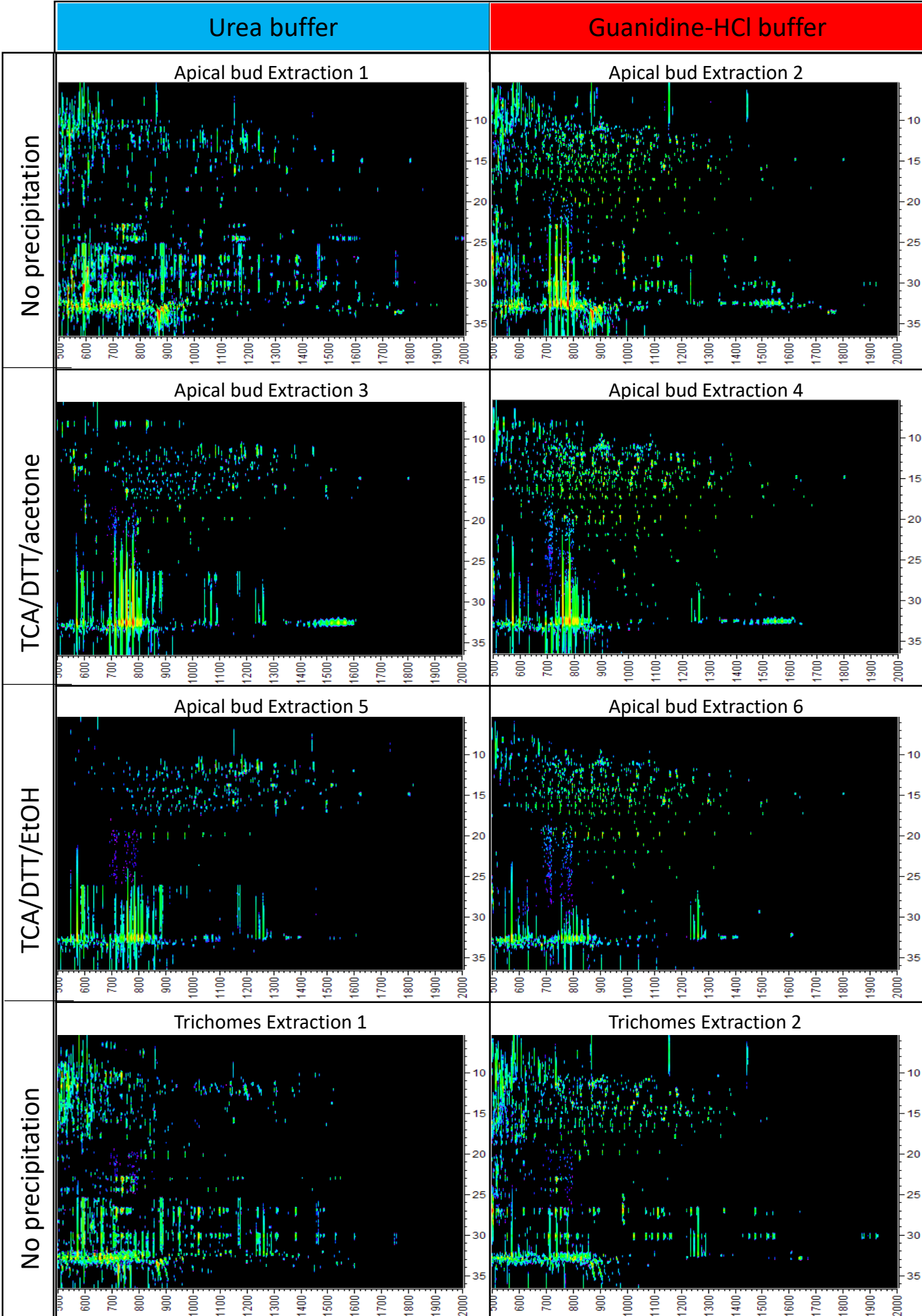
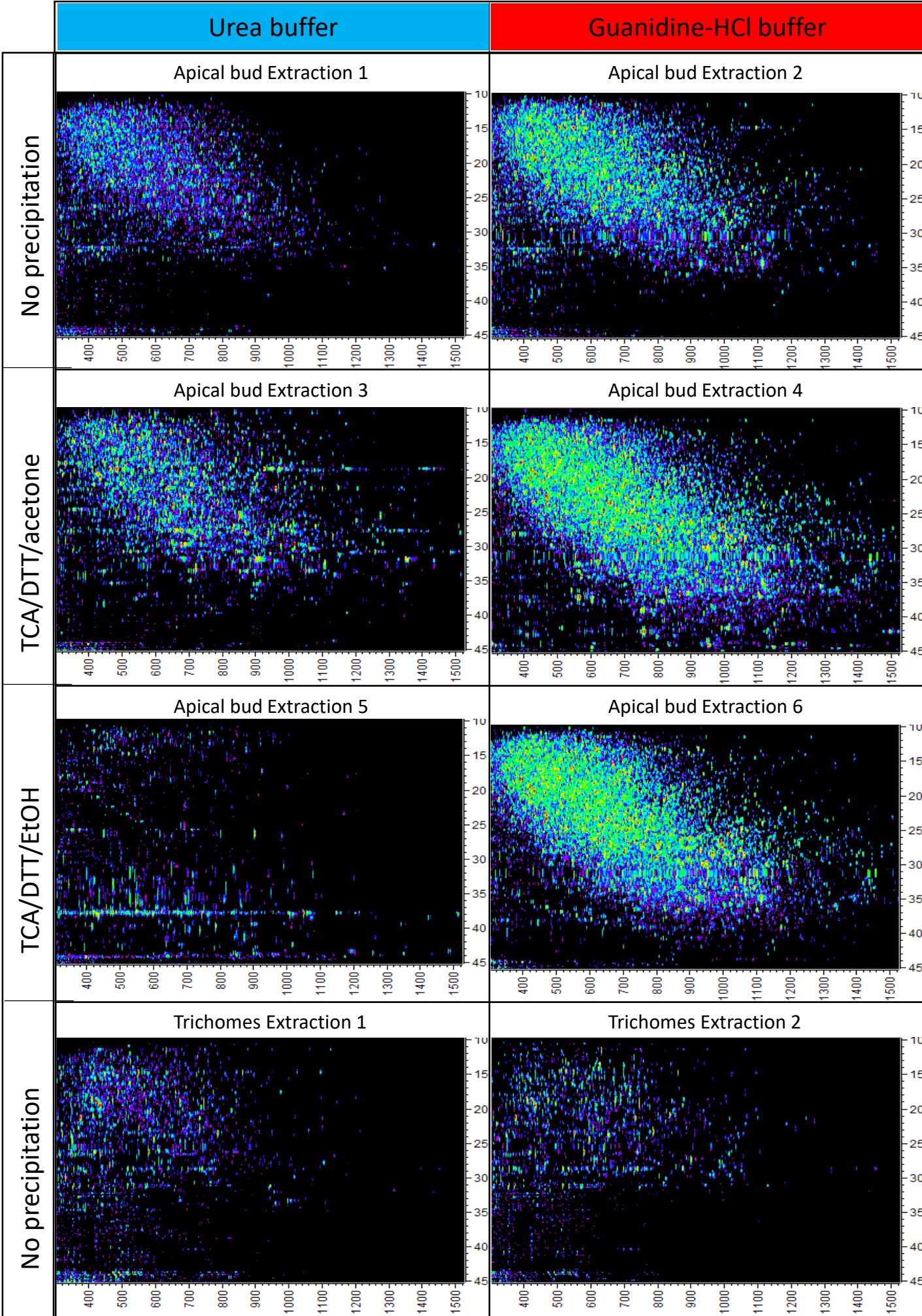


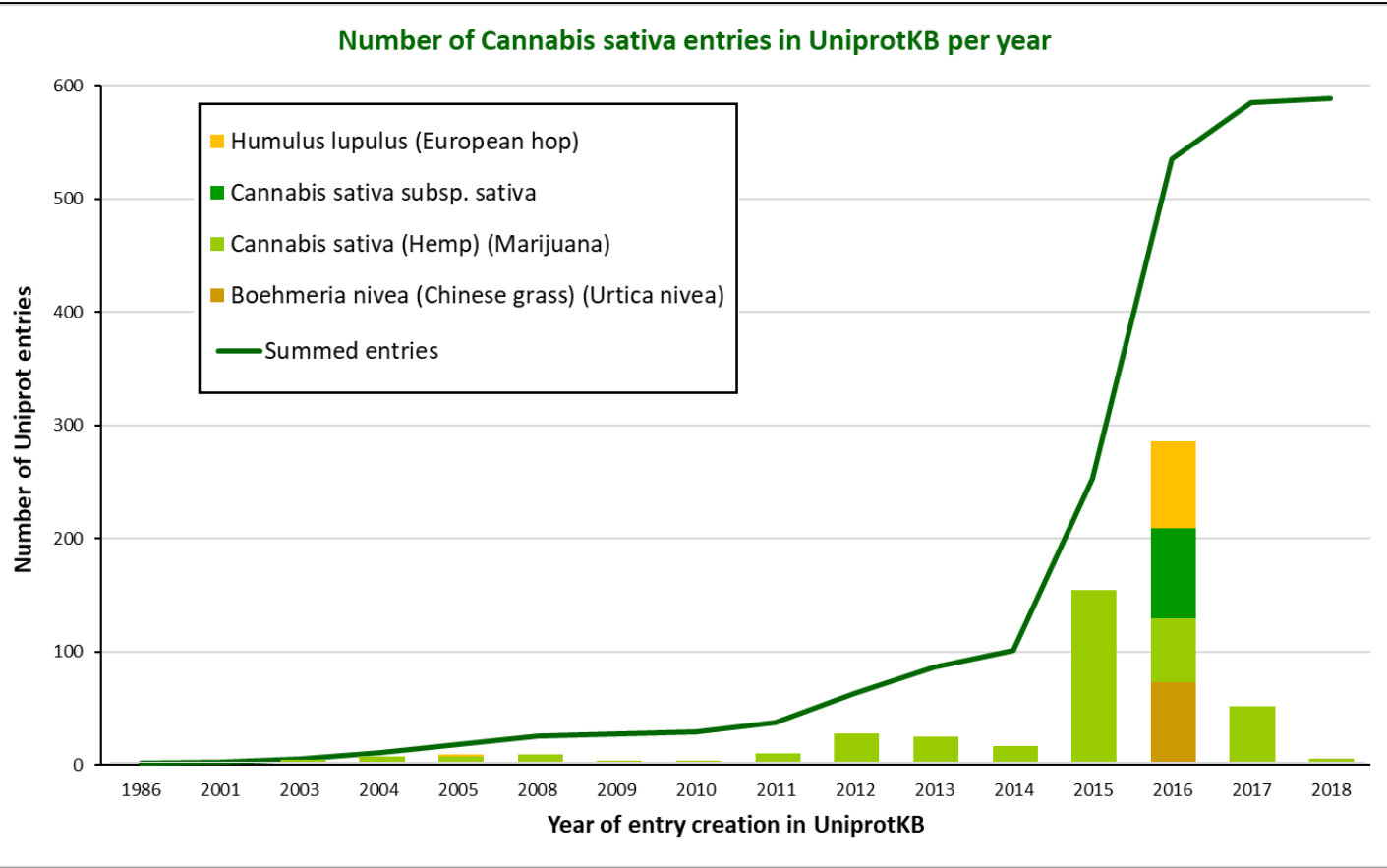
Supplementary Figure S1: LC-MS maps of intact proteins.



Supplementary Figure S2: nLC-MS maps of tryptic peptides.



Supplementary Figure S3: Distribution of Cannabis sativa entries in UniprotKB from 1986 to 2018.



Supplementary Figure S4: Quantitative comparison of the six extraction methods.

Values obtained number of intact proteins quantified by UPLC-MS and number of tryptic peptides identified by nLC-MS/MS were converted into percent and coefficient of variation (CV). Resulting percentages are plotted in the histogram below and vertical bars denote CVs.

