



S2 Fig: LC-MS analysis of T4 DNA composition.

(A-B) Extracted ion chromatogram overlay of nucleotide mass traces in T4 WT DNA (front) and NgTET-treated T4 DNA (back). Data exemplifies differences in relative abundances of cysteine nucleotides (dC, 5hmdC, 5fdC, 5cadC) **(B)** while dG, dT and dA **(A)** show similar trends. Signal intensities have been normalized against the overall nucleotide signal in the respective sample to correct for differences in sample concentration and injection volume. **(C-D)** Extracted ion chromatogram of presumed 5ghmdC **(C)**. The fragmentation pattern of presumed 5ghmdC corresponds to the predicted one **(D)**. **(E)** Relative abundance of cytosine derivatives in T4 phage isolated from different strains (two-sided Student's t-test, * - $P_{\text{signif}} < 0.05$, ** - $P_{\text{signif}} < 0.025$; n.d.: not detected). $n = 3$ biological replicates.