

## Appendix 7.

### Distribution of distances between start codon and transcription start site for putative strong promoters of *E. coli*

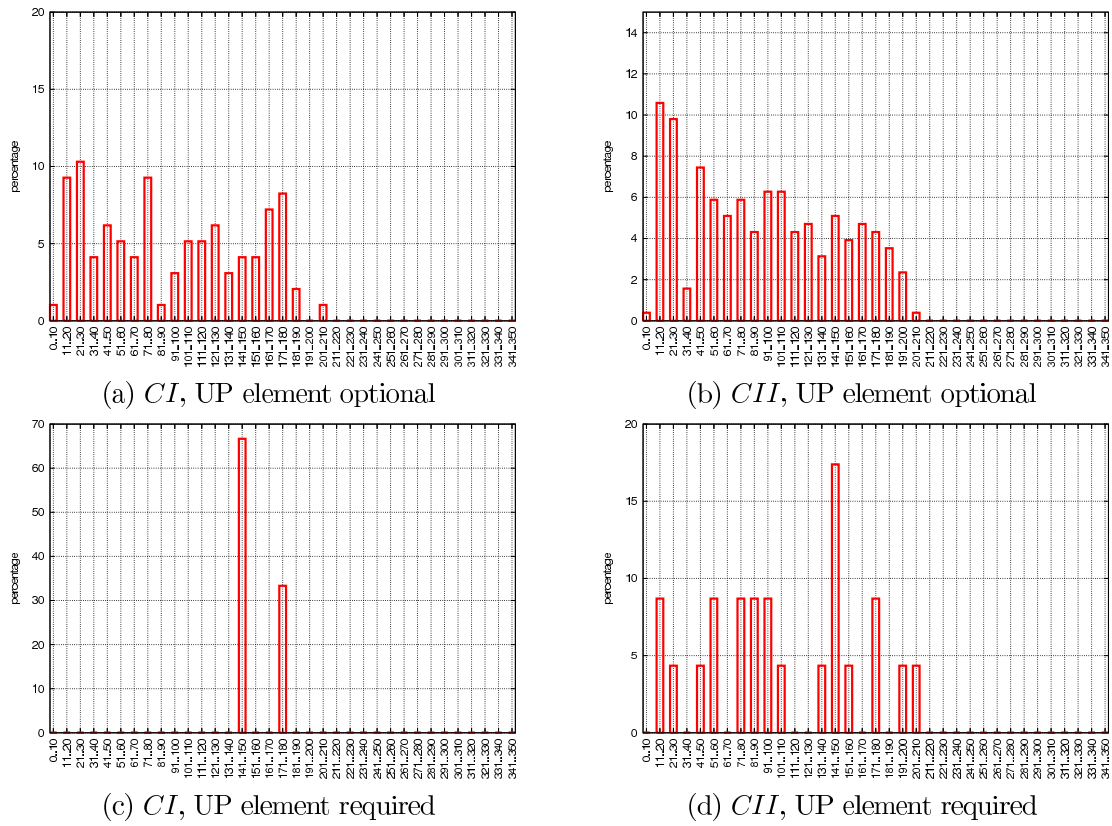


Figure 7.1. Distribution of distances between start codon and transcription start site for strong promoters identified by BACTRANS<sup>2</sup> in *E. coli* genome, under four constraint sets. See text, Subsection "Genome analysis upon request" for the definition of *CI* and *CII* constraints.