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Fig. 25-4. Comparison of the promoter regions of the late operons in phages P2, 186 and P4. The transcriptional start sites are indicated by the arrows. The common sequences are indicated in bold, and their consensus sequences are indicated below the promoter sequences and compared to conserved -10 and -35 regions of the σ^{70} promoter.

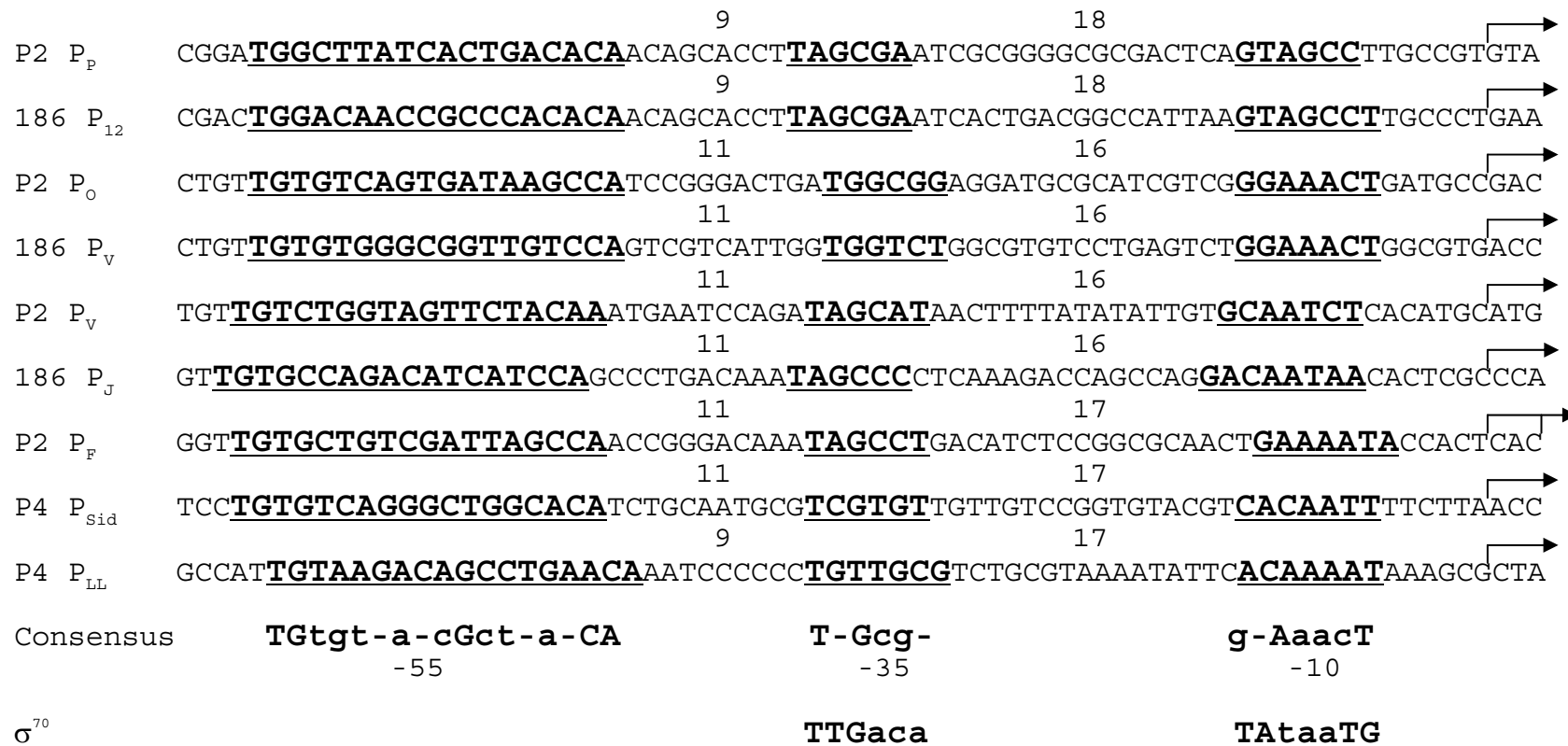


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