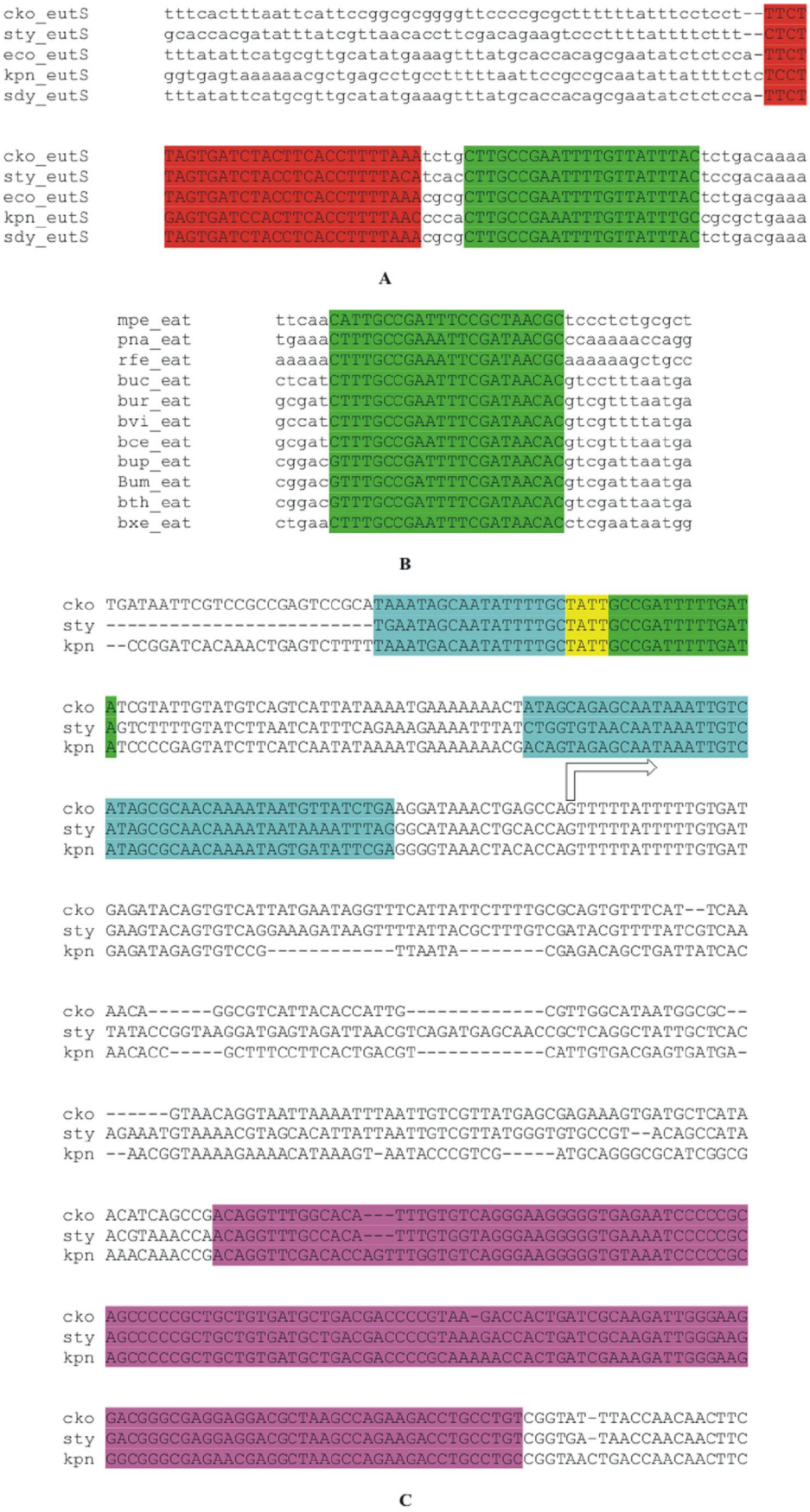

Figure S2. The regulatory elements in the upstream regions of various bacteria. A. The upstream region of the *eut* operon of the *Enterobacteriaceae*; B. The upstream region of the *eut* operon of Betaproteobacteria; C. The upstream region of the *cob* operon of the Entrobacteriales. Green – the putative EutR-binding site; red – the putative Crp-binding site; blue – the PocR-binding sites [8]; purple – the B12-riboswitch [25]; yellow – the EutR-binding site and the PocR-binding site overlap. The arrow indicates the transcription start site of *cbiA* [23]