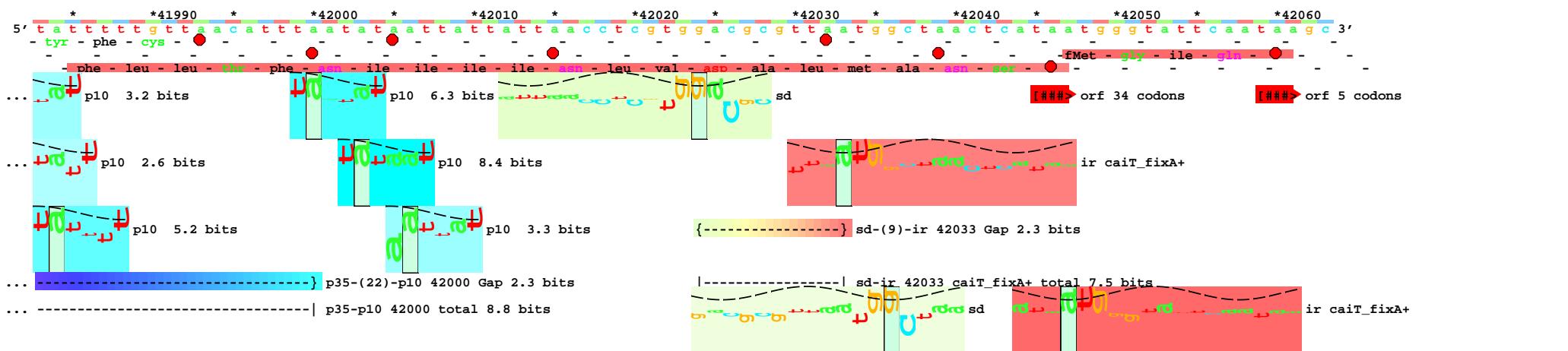
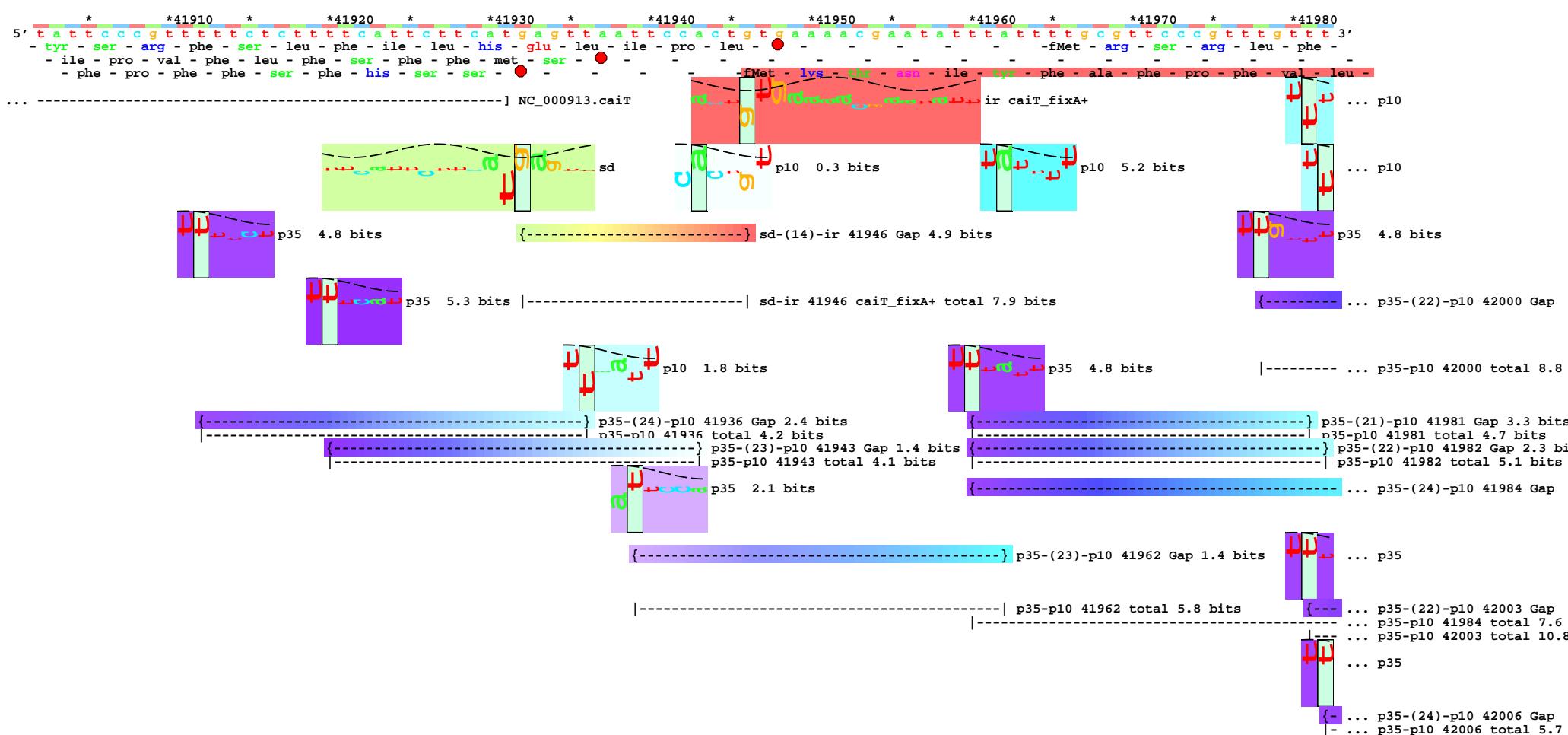
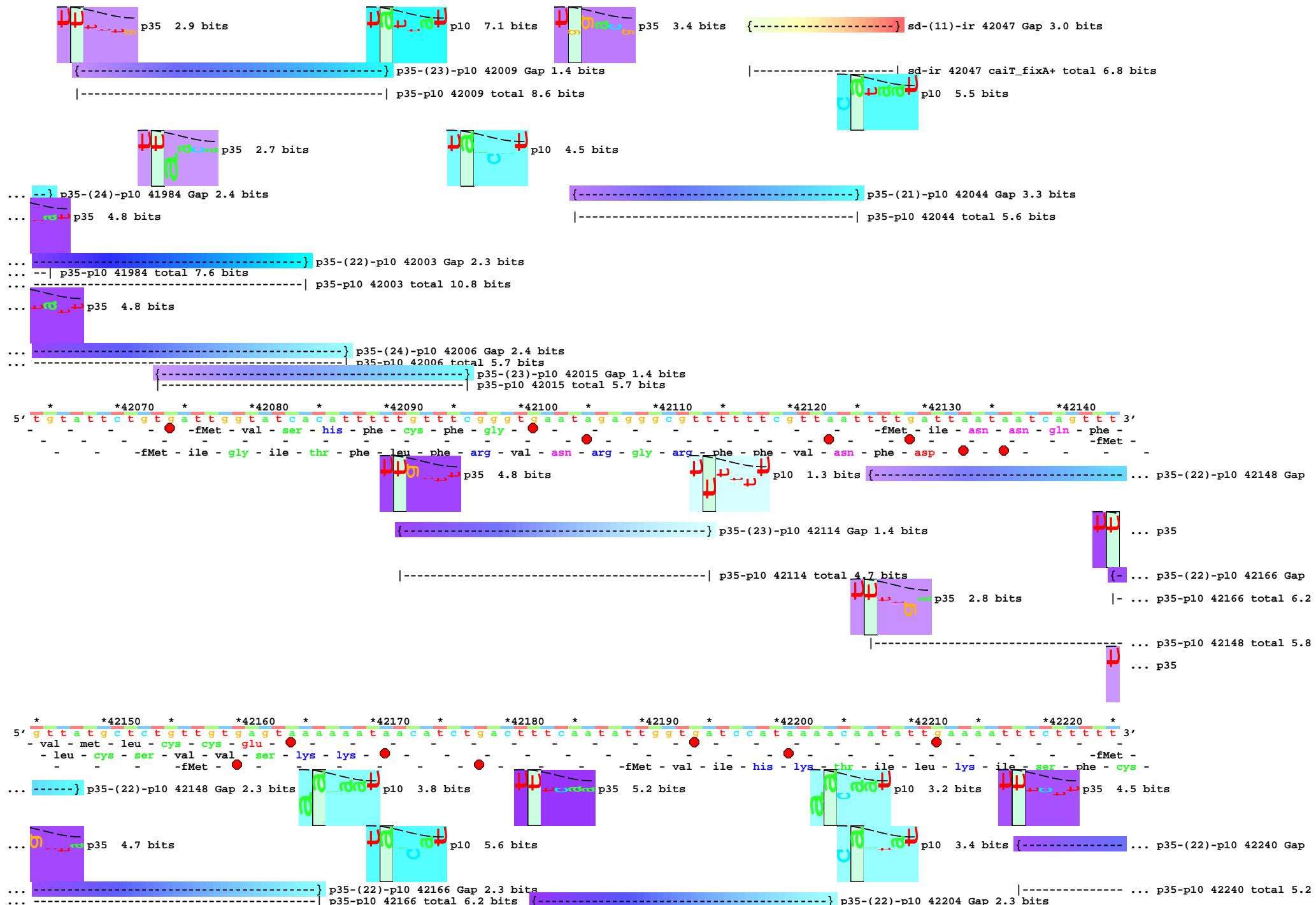


- 1 -  
piece 1, NC\_000913, caiT\_fixA+, config: linear, direction: +, begin: 41902, end: 42422





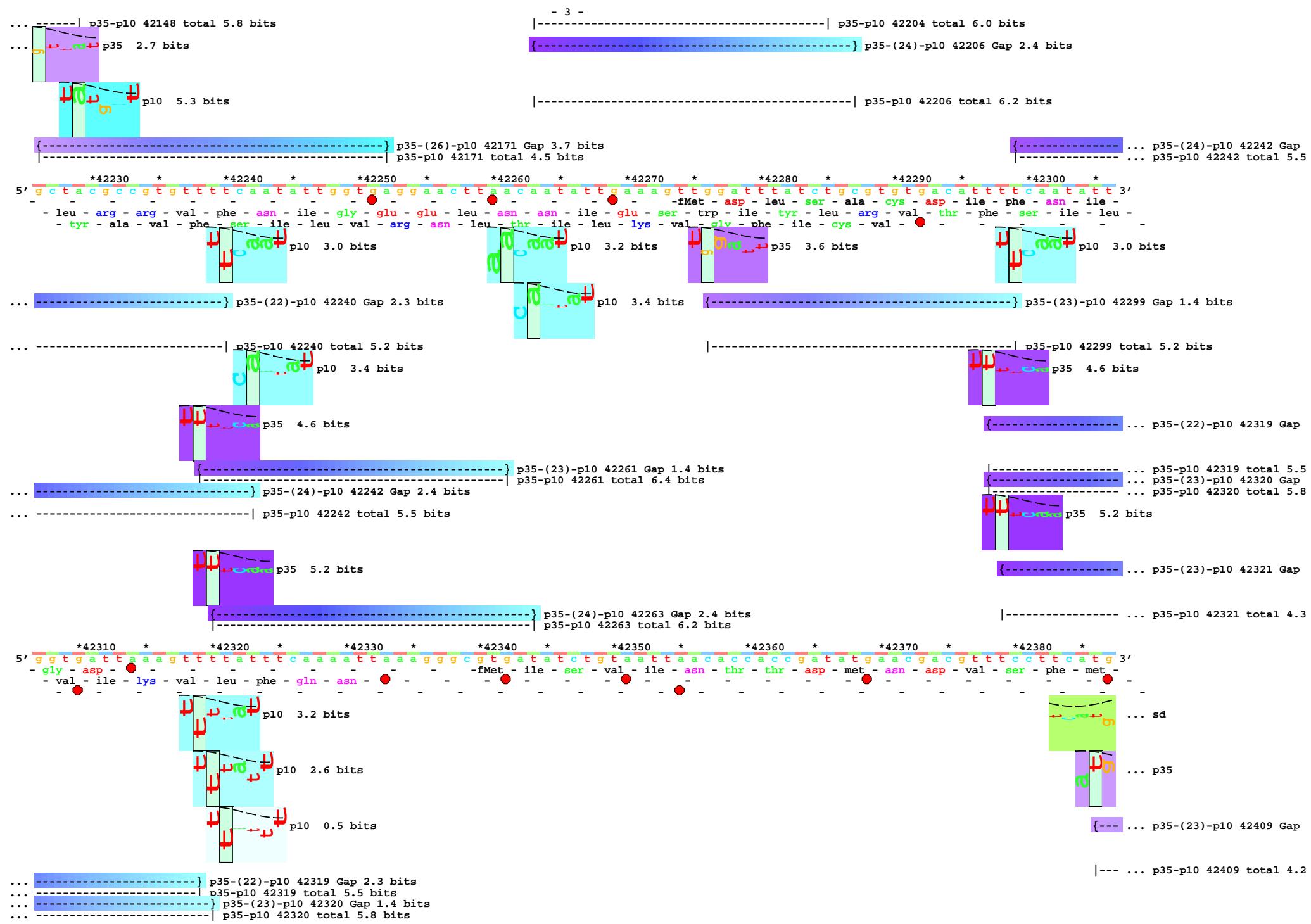


Diagram illustrating a bacterial promoter region (p35-p10) with various regulatory elements and transcription start sites (TSS). The sequence is shown as 5' to 3'.

- TSS:** \*42390, \*42400, \*42410, \*42420
- Regulatory Elements:**
  - sd:** sigma factor binding site (green box).
  - ir caiT\_fixA+:** inverted repeat (red box).
  - p35:** promoter element (purple box).
  - p10:** promoter element (cyan box).
  - sd-(8)-ir:** sigma factor binding site followed by an inverted repeat (green and red boxes).
- Protein Coding:** att, tct, ggaa, gatc, caat, ga, met, att, tact, ttt, gatc, a, ile, ser, gly, asp, ala, met, lys, ile, ile, thr, cys, fMet, gln, fMet, leu.
- Annotations:** p35-(23)-p10 42321 Gap 1.4 bits, p35-p10 42321 total 4.3 bits; p35-(23)-p10 42409 Gap 1.4 bits, p35-p10 42409 total 4.2 bits; sd-(8)-ir 42403 Gap 2.4 bits, sd-ir 42403 cait fixA+ total 14.7 bits; p10 3.0 bits.