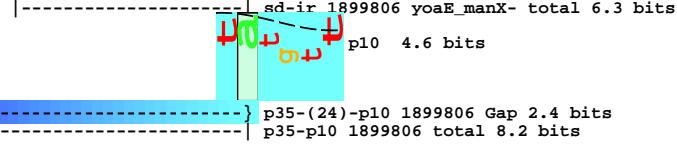
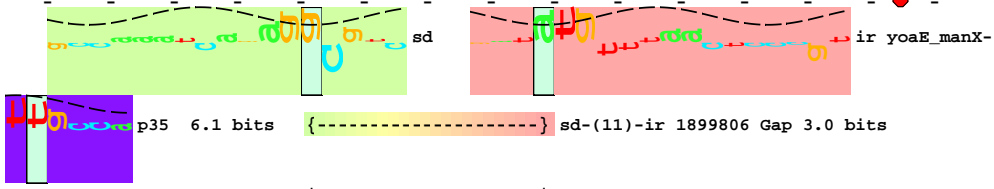
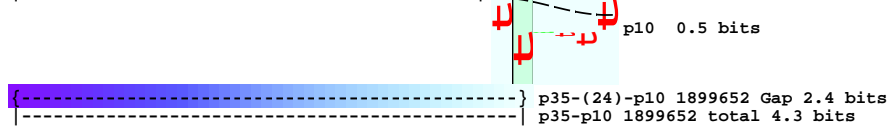
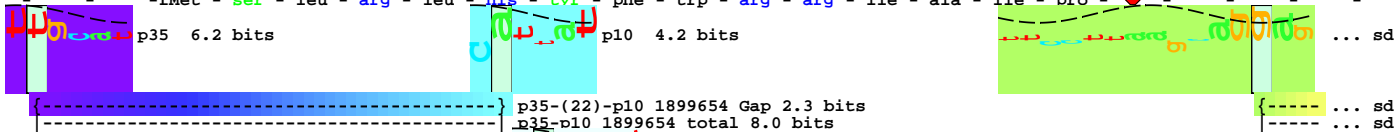


*1899850 *1899840 *1899830 *1899820 *1899810 *1899800 *1899790 *1899780
 5' c a t t c g t t t g c a a a g g a a g g t a a a t c t t t g c c a a a t c a g a g g c g t c t c t g a t a t g t t t a a c t c c c g t t t a a c a a c c a t g g a 3'
 - his - ser - phe - ala - lys - glu - gly - lys - ser - leu - pro - asn - gln - arg - arg - leu - asp - met - phe - asn - ser - arg - leu - thr - thr - met - glu -



*1899770 *1899760 *1899750 *1899740 *1899730 *1899720 *1899710 *1899700
 5' g t a t a g g g c a g t a g c c c g c a g t a t g g a t c g t c a c c g a c g t c a t t t c a g c a t c a g g c c t t t t a a c g c c t g c c t t t c t g g c a c 3'
 - tyr - arg - ala - val - ala - arg - ser - met - asp - arg - his - arg - arg - his - phe - ser - ile - arg - pro - phe - asn - ala - cys - leu - ser - gly - thr -

*1899690 *1899680 *1899670 *1899660 *1899650 *1899640 *1899630 *1899620
 5' t c t a t g c c g c a c c t t t c g t t g c a t t t g t c g t t a c g c t g c a t t t c t g g c g t c g a a t a g c t a t t c c t t a a g c a g g a g 3'
 -fMet - pro - his - leu - ser - phe - ala - phe - cys - arg - tyr - ala - cys - ile - ile - ser - gly - val - glu -
 - leu - cys - arg - thr - phe - arg - leu - his - phe - val - val - thr - pro - ala - leu - phe - leu - ala - ser - asn - ser - tyr - ser - leu - ser - arg - ser -



*1899610 *1899600 *1899590
 5' c t t g t c a t g g a a t t c t t a a t g g a c c c 3'
 -fMet - glu - phe - leu - met - asp -
 - leu - ser - trp - asn - ser - - - -fMet

... sd [-----] ... NC_000913.yoaE

