

piece 1, NC_000913, ampH_sbmA-, config: linear, direction: -, begin: 395892, end: 395492

*395890 * *395880 * *395870 * *395860 * *395850 * *395840 * *395830 * *395820 *

5' t c c c g g c t t t g g g a a a a a g a c t t a a a c a t a t g c a c t c c c t g c t a a c t t t c t t a t c g t t t t a a c a a c t g c t g t g t a t t a a c c g 3'

- ser - arg - leu - trp - glu - lys - arg - leu - lys - his - met - his - ser - cys - fMet - tyr -

- pro - gly - phe - gly - lys - lys - asp - leu - asn - ile - cys - thr - pro - ala - asn - phe - leu - ser - phe -

- pro - ala - leu - gly - lys - lys - thr -

NC_000913.sbmA

*395810 * *395800 * *395790 * *395780 * *395770 * *395760 * *395750 * *395740 *

5' c a t g a c c g c t a c a a a g c g t c a g c c a a a t g a a a c c g c c c a t t g t g a c c c a t a g c a a g g c g c t t a g t t c g c c a g c c a g c c a t 3'

-fMet - thr - ala - thr - lys - arg - gln - pro - asn - glu - thr - ala - his -fMet - cys - asp - pro -

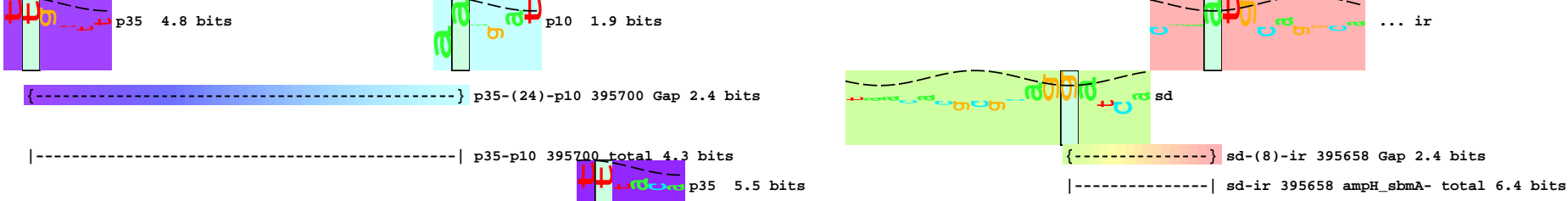
-fMet - lys - pro - pro - ile - val - thr - his - ser - lys - ala - leu - ser - ser - pro - ala -

*395730 * *395720 * *395710 * *395700 * *395690 * *395680 * *395670 * *395660 * *395650

5' a a g a a t t g t t t c g a c g g a t t t c a c a a a c c a a a g a t t g t t t a c a t a a g c g g a t t a a c a c g c g c a g g a t c a c a a a t g c a g g c a 3'

-fMet - phe - arg - arg - ile - ser - gln - thr - lys - asp -fMet - phe - thr -

-fMet - cys - leu - his - lys - arg - ile - asn - thr - arg - arg - ile - thr - asn - ala - gly - ile -



p10 1.8 bits

p35-(24)-p10 395668 Gap 2.4 bits

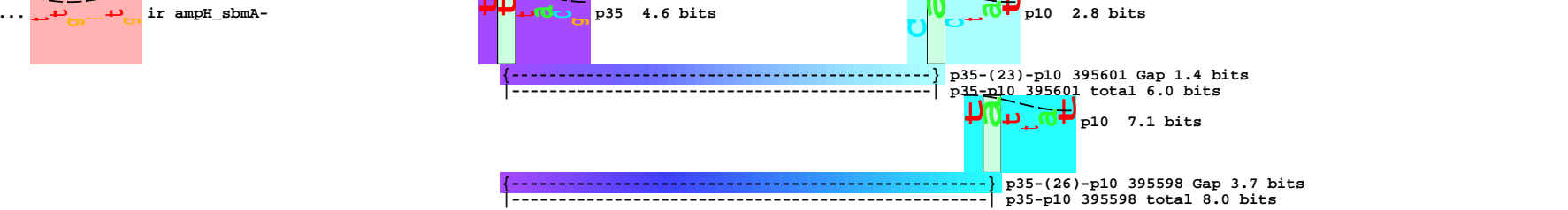
p35-p10 395668 total 4.9 bits

*395640 * *395630 * *395620 * *395610 * *395600 * *395590 * *395580 * *395570

5' t t g g t g a g c a g g a a a a g a t c g t g t t t a c g c g c c g t t a c g t t t t t a a c a c t a t t a t t g c a c t a a t t c g c c c a t t g c a a t c c t 3'

- leu - val - ser - arg - lys - lys - ile - val - phe - thr - arg - arg - tyr - val - phe - asn - thr - ile - ile - ala - leu - ile - arg - pro - leu - gln - ser -

- gly - glu - gln - glu - lys - asp - arg - val - tyr - ala - pro - leu - arg - phe -fMet - his -



*395560 * *395550 * *395540 * *395530 * *395520 * *395510 * *395500 *

5' a t c a a t g a g t a g t a t a a a t a c g c t c a g t t a c c t t c a t t c a a t c t a t g g a c a c c a c c g t t g a a a c g t a g t c t g c t t t t 3'

- ile - asn - glu -fMet - ser - ser - ile - asn - thr - leu - ser - tyr - leu - his - ser - ile - tyr - gly - his - his - arg -fMet - lys - arg - ser - leu - leu -

-fMet - ser - ser - ile - asn - thr - leu - ser - tyr - leu - his - ser - ile - tyr - gly - his - his - arg - val - glu - thr -

[###] orf 34 codons

... NC_000913.ampH