

piece 1, NC\_000913, dnaA\_rpmH-, config: linear, direction: -, begin: 3882388, end: 3881733

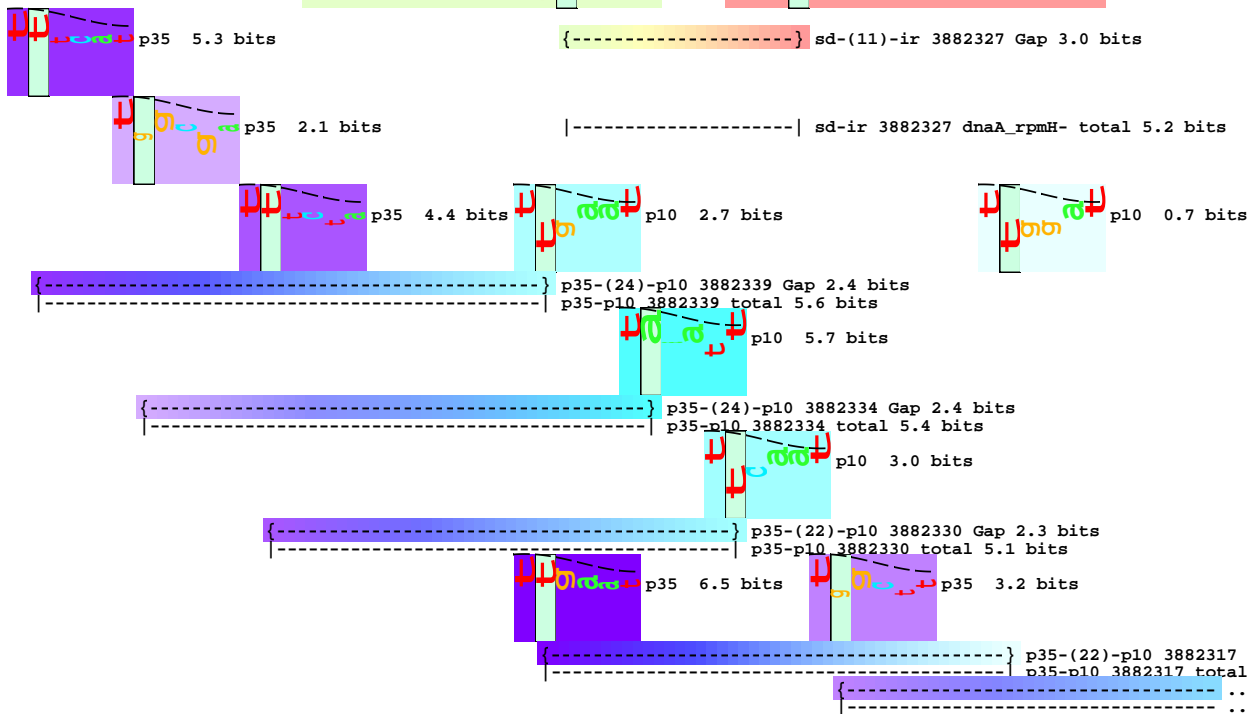
5' <sup>\*</sup>c a g t a c a g a c g g t t g a a a a g t g c g t t c a t g g c g a t t t c t a c c t a a a c t t g a a t a a a t t c a a t g g c t t t a t t g g a t a t c c g 3'

- gln - tyr - arg - arg - leu - lys - ser - ala - phe - his - gly - asp - phe - tyr - leu - asn - leu - asn - lys - phe - asn - gly - phe - ile - gly - tyr - pro -

- ser - thr - asp - gly - fMet - arg - phe - met - ala - ile - ser - thr - fMet - ala - leu - leu - asp - ile - arg -

- val - gln - thr - val - glu - lys - cys - val - ser - trp - arg - phe - leu - pro - lys - leu - glu -

...-----] NC\_000913.rpmH sd ir dnaA\_rpmH-

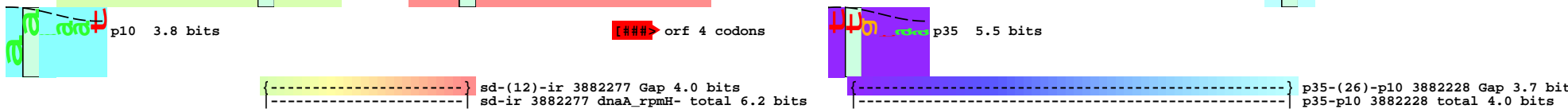


5' <sup>\*</sup>c o g a a a a t g a a a c g a t g g a c a c c g a a g c o a t g g g t g a t t a a a g a a c t t g a a t a a a t t g t a a t a a t t g t a c a c t c c g g a g t c a a t 3'

- pro - lys - asn - glu - thr - met - asp - thr - glu - ala - met - gly - asp -

- arg - lys - met - lys - arg - trp - thr - pro - lys - pro - trp - val - ile - lys - glu - ala - gly - leu - fMet - tyr - thr - pro - glu - ser - ile -

sd ir dnaA\_rpmH- orf 26 codons ... p10



... p35-(22)-p10 3882303 Gap 2.3 bits

... -----| p35-p10 3882303 total 4.7 bits

5' <sup>\*</sup> <sup>\*3882220 \*</sup> <sup>\*3882210 \*</sup> <sup>\*3882200 \*</sup> <sup>\*3882190 \*</sup> <sup>\*3882180 \*</sup> <sup>\*3882170 \*</sup> <sup>\*3882160 \*</sup> <sup>\*3882150 \*</sup> 3'

- leu - phe - pro - tyr - leu - pro - arg - phe - ser - ala - pro - phe - arg - arg - glu - asn - val - arg - pro - his - thr - thr - ser - gly - asn - gln - his - gly -  
 -fMet - tyr - asp - leu - thr - pro - val - glu - thr - ser - met - ala -

... p10 2.2 bits

5' <sup>\*</sup> <sup>\*3882140 \*</sup> <sup>\*3882130 \*</sup> <sup>\*3882120 \*</sup> <sup>\*3882110 \*</sup> <sup>\*3882100 \*</sup> <sup>\*3882090 \*</sup> <sup>\*3882080 \*</sup> <sup>\*3882070 \*</sup> 3'

- ala - pro - gly - gly - gly - leu - tyr - gly - leu - met - gly - lys - ala - gln - gly - ser - ser - trp - ile - phe - ile - arg - ser - ile - lys - pro - ile -  
 - arg - arg - val - glu - asp - tyr - thr - gly -

p35 4.8 bits

... p35-(23)-p10 3882063 Gap  
... p35-p10 3882063 total 4.1

5' <sup>\*</sup> <sup>\*3882060 \*</sup> <sup>\*3882050 \*</sup> <sup>\*3882040 \*</sup> <sup>\*3882030 \*</sup> <sup>\*3882020 \*</sup> <sup>\*3882010 \*</sup> <sup>\*3882000 \*</sup> <sup>\*3881990 \*</sup> 3'

- phe - val - tyr - gly - his - ser - met - val - ile - lys - phe - ser - asn - met - arg - arg - lys - fMet - pro - ala - ser - arg - gln - asp - arg - leu - his - leu - ala - ser -  
 -fMet - ser - met - val - ile - lys - phe - ser - asn - met - arg - arg - lys - ser - cys - pro - pro - arg - gly - arg - ile - val - tyr - thr -

p10 0.7 bits

p10 3.4 bits

... --} p35-(23)-p10 3882063 Gap 1.4 bits  
... --| p35-p10 3882063 total 4.1 bits

p35 5.6 bits

p35-(23)-p10 3882038 Gap 1.4 bits  
p35-p10 3882038 total 7.6 bits

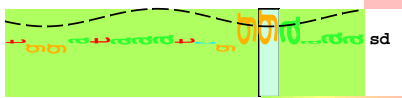
5' <sup>\*</sup> <sup>\*3881980 \*</sup> <sup>\*3881970 \*</sup> <sup>\*3881960 \*</sup> <sup>\*3881950 \*</sup> <sup>\*3881940 \*</sup> <sup>\*3881930 \*</sup> <sup>\*3881920 \*</sup> <sup>\*3881910 \*</sup> 3'

- ser - gly - lys - ser - cys - gly - fMet - arg - ser - leu - gly - tyr - asp - pro - arg -  
 -fMet - asp - lys - ser - gly - lys - ser - val - arg - asn - arg - arg - ser - leu - ala - gln - phe - arg - leu - fMet - ile - arg - gly -

p35 4.3 bits

ir dnaA\_rpmH-

###> orf 13 codons



sd  
sd-(8)-ir 3881948 Gap 2.4 bits  
sd-ir 3881948 dnaA\_rpmH- total 9.1 bits

p10 3.6 bits

p35-(24)-p10 3881955 Gap 2.4 bits  
p35-p10 3881955 total 5.4 bits

5' <sup>\*</sup> <sup>\*3881900 \*</sup> <sup>\*3881890 \*</sup> <sup>\*3881880 \*</sup> <sup>\*3881870 \*</sup> <sup>\*3881860 \*</sup> <sup>\*3881850 \*</sup> <sup>\*3881840 \*</sup> <sup>\*3881830 \*</sup> 3'

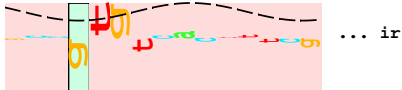
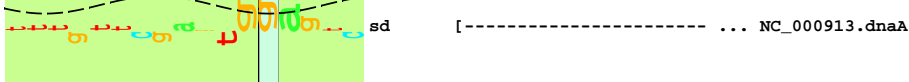
- ser - arg - ser - phe - cys - arg - ile - leu - ile - gly - his - ile - thr - ala - asp - ser - gly - ser - cys - val - thr - leu - lys - gln - gly - leu - phe -  
 - pro - asp - arg - phe - ala - gly - ser - fMet - gln - asp - leu - asp - arg - ala - tyr - asn - arg - arg - gln - arg - phe - val - arg - his - pro - gln - ala - gly - ser - phe - arg -

p35 5.2 bits

p10 4.1 bits

{-----} p35-(21)-p10 3881869 Gap 3.3 bits  
|-----| p35-p10 3881869 total 6.0 bits

\*3881820 \*            \*3881810 \*            \*3881800 \*            \*3881790 \*            \*3881780 \*            \*3881770 \*            \*3881760 \*            \*3881750 \*  
5' g a c g t a c g t c a a c a a t c a t g a a t g t t t c a g c c t t a g t c a t t a t c g a c t t t t g t t o g a g t g g a g t c c g c c g t g t c a c t t t o g 3'  
- asp - val - arg - gln - gln - ser - ● -fMet - phe - gln - pro - - - -fMet - phe - glu - trp - ser - pro - pro - cys - his - phe - arg -  
- arg - thr - ser - thr - ile - met - asn - val - ser - ala - leu - val - ile - ile - asp - phe - cys - ser - ser - gly - val - arg - arg - val - thr - phe - ala -



{-----} sd-(9)-ir 3881752 Gap 2.3 bits  
|-----| sd-ir 3881752 dnaA\_rpmH- total 5.2 bits

\*3881740 \*  
5' c t t t g g c a 3'  
- leu - trp -  
- phe - gly -  
- leu - ala -

... ----- ... NC\_000913.dnaA

... ir dnaA\_rpmH-

