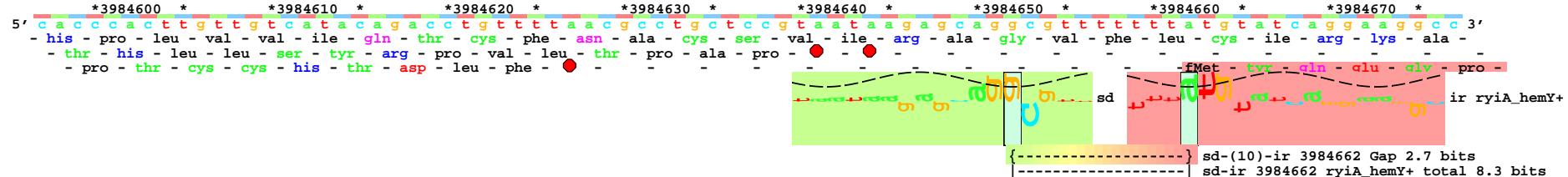


- 1 -

piece 1, NC_000913, ryaA_hemY+, config: linear, direction: +, begin: 3984597, end: 3984728



Sequence alignment diagram showing the alignment of six cDNA sequences (3984680, 3984690, 3984700, 3984710, 3984720) against a reference sequence. The reference sequence is at the top, and the cDNA sequences are below it. A red dot highlights a difference at position 3984700.



<----- ... NC_000913.hem



{ p35-(22)-p10 3984711 Gap 2.3 bits
| p35-p10 3984711 total 7.0 bits }

| ~~foo~~ fit 3301/11 total 700 bytes