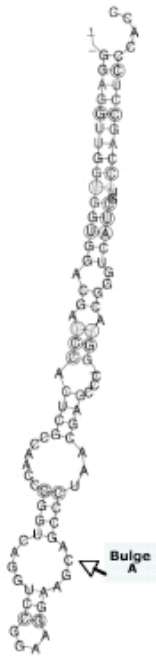
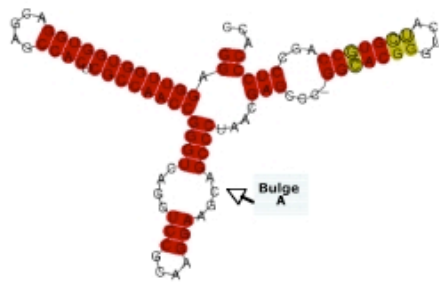


**Predicted structure of alpha-proteobacteria conserved *sra* genes**

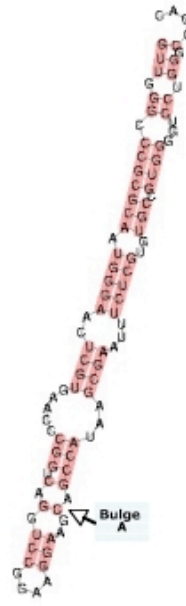
*sra05* = *ffs*



*Alifod prediction*

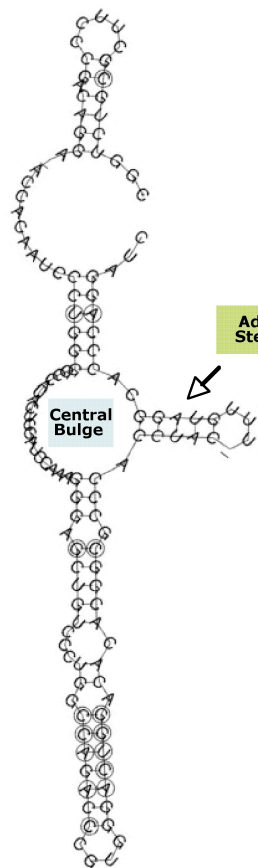


*RNAz prediction*

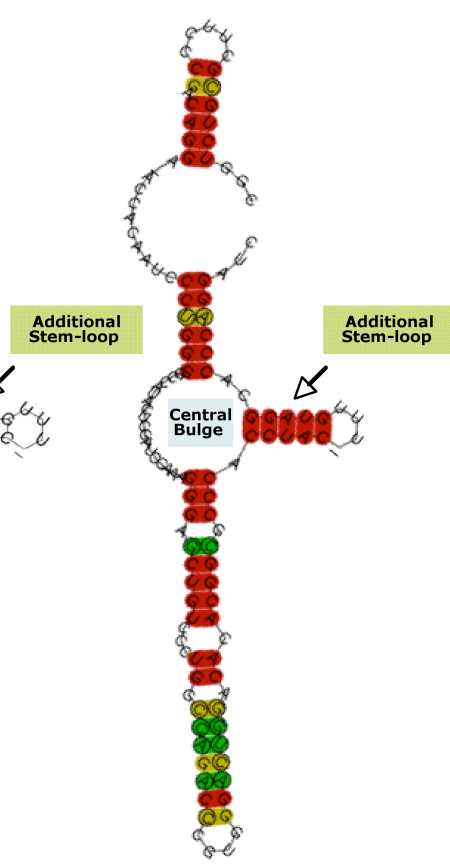


*RFAM*

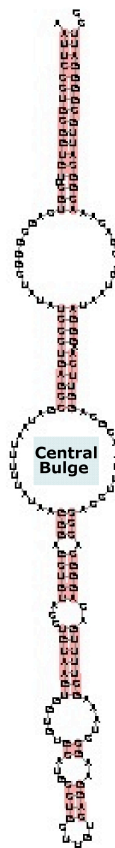
*sra56* = *6S*



*Alifod prediction*

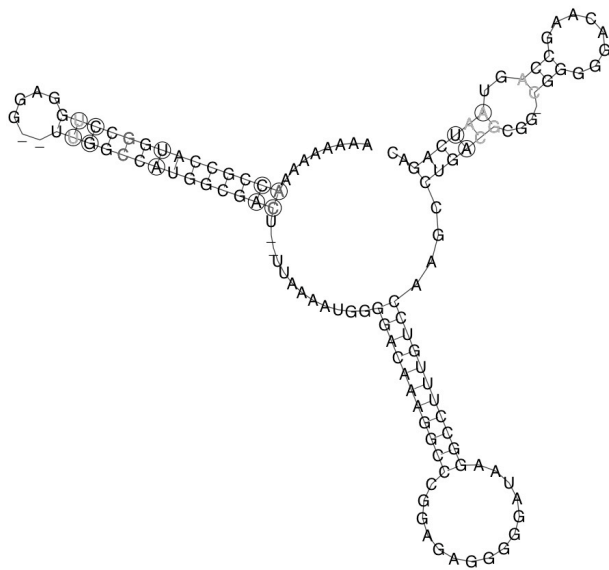


*RNAz prediction*

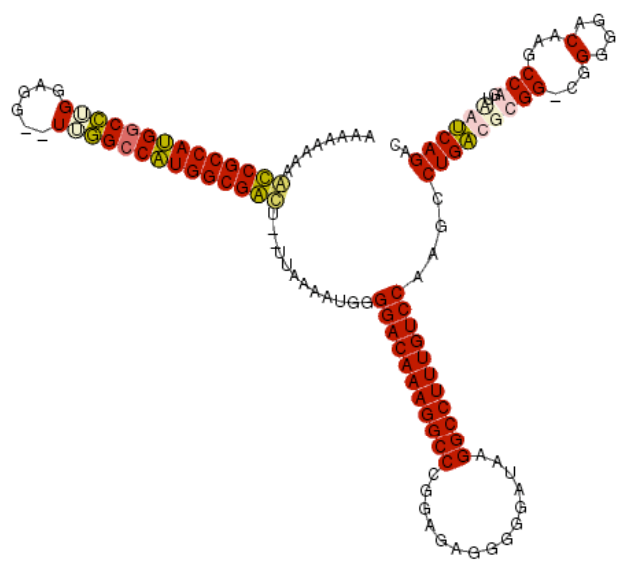


*RFAM*

sra03

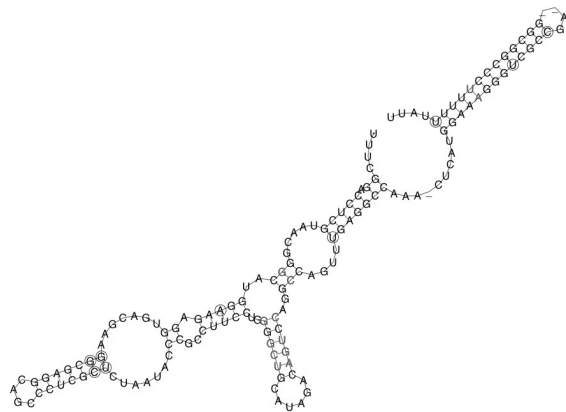


*Alifod prediction*

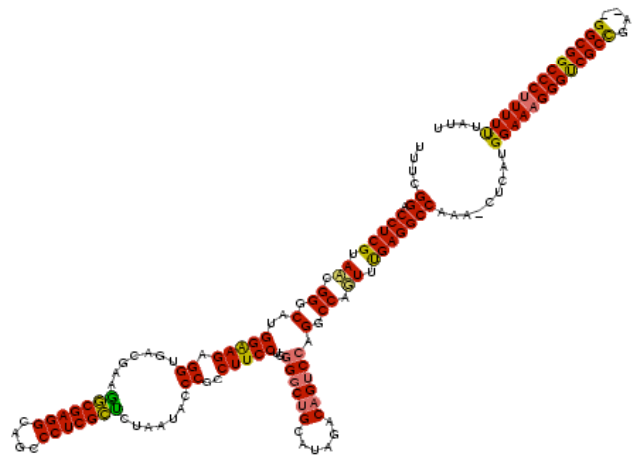


*RNAz prediction*

sra25

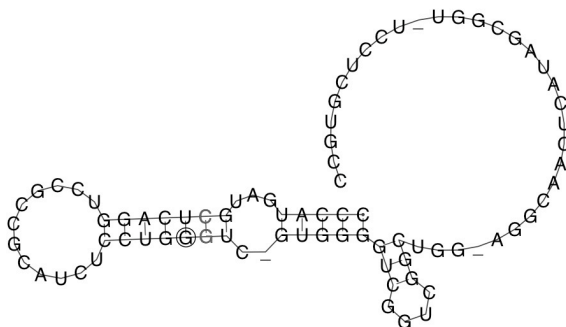


*Alifod prediction*



*RNAz prediction*

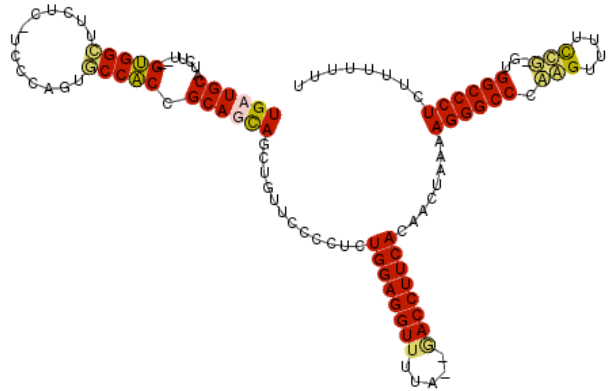
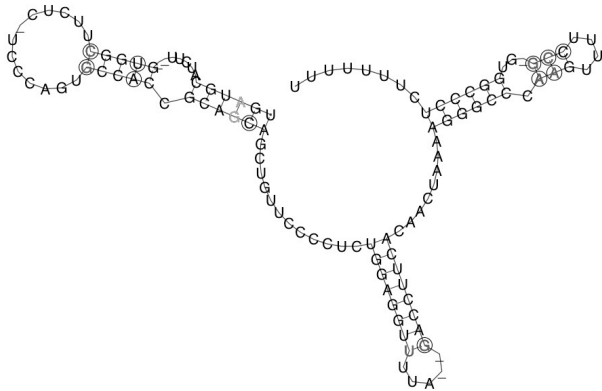
sra33



*Alifod prediction*

*No RNAz prediction ( $P > 0.5$ )*

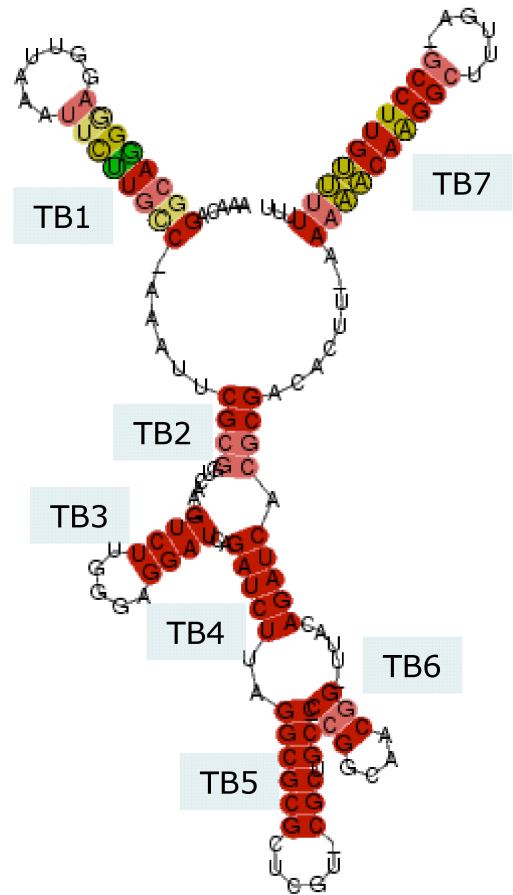
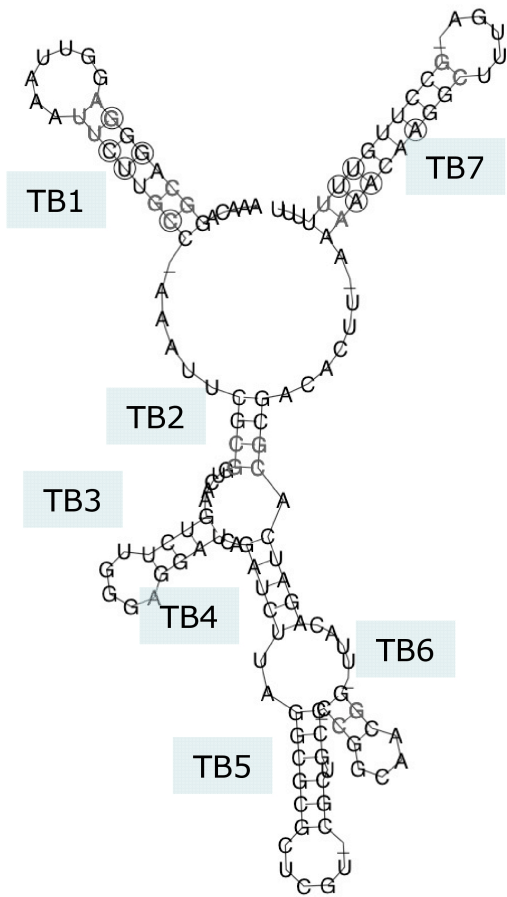
*sra41*



*Alifod prediction*

*RNAz prediction*

*sra32*



*Alifod prediction*

*RNAz prediction*