

Alignment and structure of GcvB.

This is a part of the 5' end of the Rfam alignment GcvB. The inner (green) stem usually has a single bulge loop. However, due to the one sequence that has a 3nt insertion (yellow column), the original RNAalifold treats all sequences as if they had a bulge loop there. Because of this, it prefers the large interior loop over two small bulges, which leads to the 3 missing base pairs (magenta). Treating gaps as not being there – instead of as “phantom” nucleotides – remedies that.

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U00096-2/2940718-2940864 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGGUGAA---CUUUGGCUUACG--GU
U29581-1/24251-24397 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGGUGAA---CUUUGGCUUACG--GU
U01030-1/175-29 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGGUGAA---CUUUGGCUUACG--GU
AB016765-1/195503-195649 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGGUGAA---CUUUGGCUUACG--GU
BX950851-1/1137282-1137144 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGGUGAA---CUUUGGCUUACG--GU
BX571861-1/50378-50242 ACUUC-CUAAGCCGGNACGAAAAGUGA-AUCAGCUUA---ACSCUAUGAAA---CUUUGGCUUUGUG--GU
AL627277-1/14855-15001 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGAUGGG---CUUUGGCUUACG--GU
AE008836-1/18189-18335 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGAUGGG---CUUUGGCUUACG--GU
AE016843-1/272852-272998 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGAUGGG---CUUUGGCUUACG--GU
AE016987-1/274185-274331 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGGUGAA---CUUUGGCUUACG--GU
AE013916-1/1147-1288 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGGUGAA---CUUUGGCUUACG--GU
AJ414146-1/50795-50654 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGAUGGG---CUUUGGCUUACG--GU
AB017137-1/228019-228160 ACUUC-CUGAGCCGGNACGAAAAGUUUUUACGGAUUGCGU-SUUCUGAUGGG---CUUUGGCUUACG--GU
U32799-1/164-25 UCCUU-ACAAGUACACUAUAUAG-AUUGGUAUUGCC-UUACUGAUAAG---UGAUAUUCAGACUUAAG---
AE006092-1/5387-5242 CCUCUAAAUGCGCCUAUCUUUAUG-AUUGGUAUUGCC-UUACUGAUAAG---UGAUAUUCAGACUUAAG---
AE004173-1/8619-8498 CGCAACGGCCGUGAUCGGCUGAUU-UCCU-----UGUAAAU---CAAAGGCCUUAAGU--GU
AE016797-1/298902-299024 CGCAACGGCCGUGAUCGGCUGAUU-UCCU-----UGUAAAU---CAAAGGCCUUAAGU--GU

GcvB reference ..... ((...)) ..... ((((((.....)))))) ..... (((((((.....)))))) .....
Originale ..... ((((((.....)))))) ..... (((((((.....)))))) ..... (((((((.....)))))) .....
New ..... ((((((.....)))))) ..... (((((((.....)))))) ..... (((((((.....)))))) .....
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