```
16  GUCCACCAGCAGU---UGCUGCAAAG  38  dnr mRNA
    :||   |||:|| |     ::|||||| |
56  UAGACGGUUGUGAUUUGUGACGUUUC  31  ErsA
```

Figure S2. Model of base-pairing interaction between ErsA and *dnr* mRNA by IntaRNA software. The region from nt +31 to +56 of ErsA was predicted to bind within the coding sequence of *dnr* mRNA, from nt 16 to nt 38 downstream the translation start site (ATG). Sequence coordinates refer to the +1 translational start site for *dnr* mRNA.