Transition-state stabilization in *Escherichia coli* ribonuclease P RNA-mediated cleavage of model substrates

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ABSTRACT

We have used model substrates carrying modified nucleotides at the site immediately 5' of the canonical RNase P cleavage site, the -1 position, to study Escherichia coli RNase P RNA-mediated cleavage. We show that the nucleobase at -1 is not essential but its presence and identity contribute to efficiency, fidelity of cleavage and stabilization of the transition state. When U or C is present at -1, the carbonyl oxygen at C2 on the nucleobase contributes to transition-state stabilization, and thus acts as a positive determinant. For substrates with purines at -1, an exocyclic amine at C2 on the nucleobase promotes cleavage at an alternative site and it has a negative impact on cleavage at the canonical site. We also provide new insights into the interaction between E. coli RNase P RNA and the -1 residue in the substrate. Our findings will be discussed using a model where bacterial RNase P cleavage proceeds through a conformational-assisted mechanism that positions the metal(II)-activated H₂O for an in-line attack on the phosphorous atom that leads to breakage of the phosphodiester bond.

INTRODUCTION

Transfer RNA (tRNA) genes are transcribed as precursors with extra residues at their 5'- and 3'-ends that have to be removed to generate functional tRNAs. The endoribonuclease—RNase P—is responsible for removing the extra 5' residues, i.e. the 5' leader. In Bacteria, RNase P consists of one protein and one RNA subunit, referred to as the C5 protein and RNase P RNA (RPR), respectively; in Archaea and Eukarya, the number of proteins is expanded. Irrespective of origin, the catalytic activity resides in the RNA moiety and the RNA alone can mediate cleavage in the absence of the protein (1-3). However, recent data suggest the existence of RNase P-like activities based solely on protein (4,5). Several determinants in the tRNA precursor substrate influence binding and cleavage efficiency [for recent reviews see (6-8)]. For one of these, the residue at -1, genetic and biochemical data suggest that it interacts with a conserved A at position 248 in Escherichia coli (Eco) RPR_{wt}. This interaction is referred to as the N_{-1}/A_{248} interaction [(6,9); see Figure 1A]. The recently published structure of RNase P in complex with tRNA positions A_{248} close to the 5' termini of the tRNA (11); however, it is still unclear if and how N_{-1} in the substrate interacts with A_{248} .

The identity of the -1 residue varies in tRNA precursors. In *E. coli*, the majority ($\approx 57\%$) carry a uridine at -1, whereas in others, e.g. *Mycobacterium tuberculosis*, a cytidine is more frequent (12,13). In a tRNA precursor context, in vitro data suggest that substrates with U at -1 bind to RPR with higher affinity compared with substrates with other -1 residues (9,14). Moreover, for substrates with C at -1, which can pair with a guanosine at position +73 [the discriminator base position; (15)], in vitro and in vivo studies show that cleavage occurs at the correct site between positions -1 and +1 (the canonical site or the +1 site), as well as at an alternative site between residues at -1 and -2, i.e. at -1 (6–8,16). Cleavage of substrates with C_{-1}/G_{+73} and U_{-1}/G_{+73} pairs has furthermore demonstrated that optimal cleavage of the former substrate requires higher Mg²⁺concentrations (14). The rate of cleavage for Eco RPR_{wt} as a function of the identity of the residue at -1, however, does not change to any significant extent in a tRNA precursor context (9,14). In contrast, for model hairpin loop substrates (see Figure 1B), the -1 identity influences the

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Figure 1. Structures of the model hairpin loop substrate pMini3bp and structure of uridine, cytidine, iso-cytidine, guanosine and adenosine. (A) Illustration of the $\dot{N}_{-1/248}$ - and the <u>RCC</u>A-RPR interactions [interacting residues underlined (6,9,10)]. Substrate residues are marked in black and residues marked in grey represent the RPR. Black and grey arrows mark the canonical and the alternative cleavage sites, respectively. ? indicate if and how residues -1 and 248 interact (see the main text). (B) Structure of pMini3bpUG and its derivatives. The black arrow marks the canonical cleavage site at +1 and the grey arrow marks the alternative cleavage site -1. X and Y indicate where base changes were introduced as indicated on the right side. Abasic, deletion of the base; IsoC, iso-cytidine; Ino, inosine; 2AP, 2-amino purine; DAP, 2;6-diamino purine; Pu, purine. (C) Structure of the bases uridine (U), cytidine (C), iso-cytidine (IsoC), Abasic, guanosine (G) and adenosine (A). Chemical groups marked in small grey circles refer to the groups that were substituted or deleted as indicated. The ring numbering for U and G is as indicated. For details see text.

rate of cleavage (6,7,13,14,17). Hence, the effect of the -1 residue in a pre-tRNA context is likely to be obscured by other elements affecting RPR-mediated cleavage (14). Moreover, there is no information about the contribution

of the chemical groups on the -1 base to catalysis. Therefore, we decided to reinvestigate the impact of the identity of the -1 residue on cleavage using a model substrate. Our specific objectives were to identify chemical groups of the -1 nucleobase that contribute to catalysis and to determine if the nucleobase at -1 is absolutely necessary for cleavage. As available data suggest that the protein subunit of RNase P does not interact with the residue at -1 (6–8), we studied cleavage mostly in the absence of the protein. Our results show that the nucleobase at the -1 position is not necessary for cleavage in the context of a model hairpin loop substrate. However, the presence, identity and specific chemical groups at -1 influence cleavage site selection and the kinetics of cleavage. The -1 position contributes to the stabilization of the transition state for cleavage at the canonical cleavage site. We will discuss our data in view of the recently solved structure of RNase P in complex with tRNA (11) and a model that emphasizes the importance of the positioning of Mg(II) at and near the cleavage site for ensuring cleavage at the correct site.

MATERIALS AND METHODS

Preparation of substrates, RPRs and C5 protein

The different pMini3bp derivatives except pMini3bpAbasicG and pMini3bpIsoCG were purchased from Dharmacon (Lafayette, CO. USA). The pMini3bpAbasicG and pMini3bpIsoCG were synthesized in-house essentially according to Wincott et al. (18). The rSpacer and isoC phosphoramidites were purchased from MedProbe AS, Norway (Glen Research, USA) and ChemGenes, USA, respectively. All the RNA substrates were purified on 15% denaturing polyacrylamide gels and extracted overnight using a Biotrap device following the manufacturer's procedures (Schleicher and Schuell, GmbH, Germany; Elutrap in USA and Canada). This was followed by phenol-chloroform extraction according to standard procedures. The different substrates were labeled with ³²P at the 5'-end with $[\gamma$ -³²P]ATP as described elsewhere (14).

The construction of the Eco RPR_{G248} encoding gene has been reported elsewhere (19). The Eco RPR_{wt} and Eco RPR_{G248} were generated as T7 RNA polymerase run-off transcripts (20), whereas the C5 protein was purified from an E. coli BL21(DE3) strain harbouring the plasmid pET33b carrying the His6-C5 gene (N-terminal fusion). Briefly, the pET33b plasmid with the His6-C5 gene behind an isopropyl β-D-1-thiogalactopyranoside-inducible promoter was transformed into E. coli BL21(DE3). The cells were grown at 37°C in LB liquid medium supplemented with 50 μ g/ml kanamycin. At OD₆₀₀ = 0.6, isopropyl β-D-1-thiogalactopyranoside was added to a final concentration of 1.5 mM and the culture was incubated for another 4h. The His6-C5 protein was purified essentially as described by Feltens et al. (21), and the protein concentration was determined using the standard Bradford protein assay.

Assay conditions—cleavage by RPR alone and in the presence of the C5 protein

The RPR alone reactions were conducted in buffer C [50 mM Mes (pH 6.1 at 37°C), 0.8 M NH₄Cl] and indicated Mg(OAc)₂ concentrations. Before adding the preheated (37°C) substrate, the RPRs were pre-incubated at 37°C in buffer C and Mg(OAc)₂ for at least 10 min to allow proper folding. In the Mg²⁺ titration experiments, the concentrations of substrates were 0.02 μ M while the concentration of RPRs varied between 0.8 and 5.2 μ M [the concentration, see Figure legend 3; (22)].

All the reactions with the His6-C5 protein were carried out in buffer A [50 mM Tris–HCl (final pH 7.2), 5% (w/v) PEG 6000, 100 mM NH₄Cl] supplemented with 10 mM MgCl₂. The RPR was pre-incubated in buffer A at 37°C for 10 min. The His6-C5 was added, and incubation was continued for an additional 10 min followed by addition of preheated (37°C) substrate. The concentrations of RPR and His6-C5 were 0.004 μ M and 0.21 μ M [empirically determined; see also e.g. (23)], respectively, and the concentration of substrate was 0.02 μ M.

To terminate the reactions, double volumes of stop solution were added (10 M urea, 100 mM EDTA).

Determination of the kinetic constants under single-turnover conditions

The kinetic constants k_{obs} and k_{obs}/K^{sto} (= k_{cat}/K_m) were determined under saturating single-turnover conditions in buffer C at pH 6.1 and 800 mM Mg(OAc)₂. At this pH, the chemistry of cleavage is rate-limiting (24). The final concentration of the substrate was 0.02 μ M, and the concentration of RPRs varied between 0.8 and 29 μ M, depending on the substrate–RPR combination. Given that the substrates were labeled at the 5'-end, we used the 5' cleavage fragments in our activity measurements. While calculating the rate, the incubation times for each substrate and RPR combination were adjusted to ensure that the velocity measurements were in the linear range, i.e. $\leq 40\%$ of the substrate had been consumed.

The k_{obs} and k_{obs}/K^{sto} values were obtained by linear regression from Eadie–Hofstee plots as described elsewhere [(25); see also (26,27)].

RESULTS

To test the contribution of residue -1 to cleavage, we used the short model substrate, pMini3bp [Figure 1; (12)], in which cleavage relies on residues near the cleavage site, mainly the N₋₁/A₂₄₈ and the <u>RCC</u>A-RPR interactions (interacting residues underlined, Figure 1A; 6,9,10). For this purpose, we generated pMini3bp variants carrying changes at -1 (Figure 1B). These substrates were studied with respect to (i) cleavage site recognition, where cleavage between -1 and +1 is referred to the canonical or correct site (Figure 1) and (ii) Mg²⁺ dependence. We also determined the kinetic constants k_{obs} and k_{obs}/K^{sto} (k_{obs}/K^{sto} = k_{cat}/K_m; see later in the text) for cleavage under saturating-single turnover conditions at pH 6.1. At this pH, previous data have suggested that





the cleavage of other model hairpin loop substrates is rate-limiting [see e.g. Ref. (24)]. In the simplified scheme the kinetic constant kobs reflects the rate of cleavage as indicated while the rate constant $k_{obs}/K^{sto} = k_{+1}$ (Scheme 1). Under these conditions, we argue that $K^{sto} \approx K_d$ because $k_{-1} >> k_{obs}$ in the cleavage of pMini3bpUG by Eco RPR_{wt}, where k_{-1} corresponds to the rate of dissociation (Supplementary Figure S1; for experimental details see Supplementary data and Refs (25,28,29); we assume that this is the case for all the pMini3bp variants). Going from ES^1 to ES^2 involves breaking the -1/+73 base pair in the substrate (e.g. as when C is at -1 and G at +73; see Figure 1B) and positioning of the Mg²⁺ responsible for generating the nucleophile. Furthermore, the ES^1 to ES^2 route results in cleavage at the canonical cleavage site +1, whereas ES¹ to ES^{1*} describes the pathway (and e.g. does not involve breaking of the -1/+73 pair if present) that gives cleavage at the alternative site -1 [see Figure 1; see also (19)].

The nucleobase at -1 is not essential for cleavage

To understand if the nucleobase at -1 is necessary for cleavage, we prepared a substrate in which the base had been deleted, pMini3bpAbasicG (Figure 1B). Eco RPR_{wt} cleaved this substrate at two positions, at the canonical site +1 and at -1, with low efficiency compared with cleavage of pMini3bpUG where the base is present (Figure 2A). Cleavage of pMini3bpAbasicG at +1 and at -1 occurred throughout the entire concentration range of Mg^{2+} tested (Figure 3A); however, we noted that there was less cleavage at -1 at lower concentrations of Mg² (Supplementary Figure S2). The Mg²⁺ requirement for optimal cleavage was found to be comparable with that for cleavage of pMini3bpUG (Figure 3B). However, compared with cleavage of precursor tRNAs and other longer model substrates [cf. Figures 5 and 6 in Ref. (14)], cleavage of the substrate lacking the nucleobase at -1 and the other pMini3bp variants (see later in the text) required a higher concentration of Mg^{2+} to reach optimal cleavage.

Next, we determined the kinetic rate constants at saturating Mg^{2+} concentration (800 mM). As shown in Table 1, deleting the nucleobase at -1 (pMini3bp*AbasicG*) resulted in a 4200-fold reduction in both k_{obs} and k_{obs}/K^{sto} for cleavage at +1 compared with cleavage of pMini3bpUG at the same position.



Figure 2. Cleavage of pMini3bp variants with *Eco* RPR. (A) pMini3bpUG and pMini3bp*Abasic*G; the reaction was performed at 37°C in buffer C containing 800 mM Mg²⁺ (see 'Materials and Methods'). Lanes 1 (pMini3bp*Abasic*G) and 2 (pMini3bpUG), controls incubation without *Eco* RPR_{wt}; lanes 3 (pMini3bp*Abasic*G) and 4 (pMini3bpUG) incubation with *Eco* RPR_{wt} for 1179 min and 20 s, respectively. The concentrations of *Eco* RPR_{wt} and substrate were ≈ 0.8 and 0.02 μ M, respectively. The 5'CL frags indicate the 5'

Given that pMini3bp*Abasic*G is cleaved at +1 and at -1, we compared the rate constants for cleavage at these positions. The k_{obs} values for cleavage at +1 and -1 were essentially the same, whereas k_{obs}/K^{sto} was slightly higher for cleavage at -1. Following the argument that $K^{sto} \approx K_d$ (see earlier in text), we calculated the K_d values using the data for cleavage at +1. This revealed that there was no significant effect on binding comparing pMini3bp*Abasic*G and pMini3bpUG.

Taken together, these data show that the nucleobase at -1 is not essential for cleavage of the model substrate used here, but it does affect both the kinetics and accuracy of *Eco* RPR-mediated cleavage.

The identity of residue -1 contributes to catalysis

We previously reported that the kinetic constant k_{obs} (see above Scheme 1) decreased 100-fold when U was replaced with C at the -1 position in the pMini3bp model

Figure 2. Continued

cleavage fragments generated after cleavage at +1 and -1. The difference in migration of the +1 cleavage fragments in lanes 3 and 4 is likely because of the absence of the -1 nucleobase in the 5' cleavage product after cleavage of pMini3bpAbasicG at +1. For experimental details see 'Materials and Methods'. (B) Frequency of cleavage of pMini3bpAbasicG at the alternative site -1 with Eco RPR_{wt} at 800 mM^{2+} . For the calculations of the percentage of cleavage, we used the 5' cleavage fragments and the data are the mean and experimental errors of at least three independent experiments. The error bars indicate experimental errors and the frequencies of cleavage at -1 were calculated as described previously (30). For experimental details, see figure legend 2A. (C) Cleavage of pMini3bpUG and pMini3bpIsoCG with Eco RPR_{wt} and Eco RPR_{G248}; the reaction was performed at 37°C in buffer C containing 800 mM Mg²⁺ (see 'Materials and Methods'). Lanes 1 (pMini3bpUG) and 2 (pMini3bpIsoCG), controls incubation without Eco RPR for 90 min; lanes 3 and 4, pMini3bpUG incubated with Eco RPR_{wt} and Eco RPR_{G248}, respectively, for 20s; lanes 5 and 6, pMini3bpIsoCG incubated with Eco RPR_{wt} (60 min) and Eco RPR_{G248} (90 min), respectively. The concentrations of Eco RPR and substrate were ≈ 0.8 and $0.02 \,\mu$ M, respectively. The 5'CL frags indicate the 5' cleavage fragments generated after cleavage at +1 and -1. (D) Frequency of cleavage of pMini3bpIsoCG at the alternative site -1 (mean and experimental errors of at least three independent experiments) with Eco RPR_{wt} and Eco RPR_{G248} at 800 mM Mg²⁺. For experimental details and calculations, see figure legends 2B and 2C. (E) Cleavage of pMini3bpGG and pMini3bpAG with Eco RPR_{wt} in the absence and in the presence of C5. Lanes 1 (pMini3bpGG) and 2 (pMini3bpAG), controls incubation without Eco RPR_{wt} and C5 for 180 min; lanes 3 (pMini3bpGG) and 4 (pMini3bpAG) cleavage without C5 for 180 min; lanes 5 (pMini3bpGG) and 6 (pMini3bpAG) cleavage with C5 for 1 min. 5' CL Frags +1 and -1 mark the migrations of the 5' cleavage fragments. The reactions were carried out at 37° C and 800 mM Mg^{2+} and 10 mM Mg^{2+} as described in 'Materials and Methods'. In the absence of C5, the concentration of Eco RPR_{wt} was $3.2 \,\mu\text{M}$, whereas it was $0.002 \,\mu\text{M}$ in its presence. The substrate concentration was $0.02 \,\mu$ M. (F) Percentage of cleavage at -1 for substrates carrying 2;6-diaminopurine (DAP), 2-aminopurine (2AP) or guanosine (G) at -1 as indicated. The reactions were performed in the absence of C5 in buffer C at 800 mM Mg²⁺ as described earlier in text and in 'Materials and Methods'. The calculations were done as described in figure legend 2B. (G) Percentage of cleavage at -1 for substrates carrying purine (Pu), adenosine (A) or inosine (Ino) at -1 as indicated. The reactions were performed in the absence of C5 in buffer C at $800\,mM~Mg^{2+}$ as described earlier in text and in 'Materials and Methods'. The calculations were done as described in figure legend 2B.



Figure 3. Cleavage of the pMini3bp variants by *Eco* RPR as a function of Mg²⁺ concentration. Panels A–E show the Mg²⁺ profiles for *Eco* RPR_{wt} (A–D) and *Eco* RPR_{G248} (E). The experiment was performed under single-turnover conditions at 37°C as described in 'Materials and Methods' and the concentration of substrates was 0.02 μ M, whereas the concentration of *Eco* RPR (in parenthesis) varied according to the following: Panel A pMini3bp*AbasicG* ($\Delta_{-1}G_{+73}$, 3.2 μ M);

substrate. These earlier experiments were done at 160 mM Mg^{2+} , a non-optimal concentration [Table 1; cf. cleavage of pMini3bpUG and pMini3bpCG Figures 5 and 6 (14)]. Hence, to investigate the impact of the identity of the residue at -1, we now determined the optimal Mg^{2+} concentration for cleavage of these two substrates. A comparison of the Mg^{2+} profiles for cleavage of pMini3bpUG (Figure 3B) and pMini3bpCG (Figure 3C) revealed that cleavage of the latter required a higher Mg^{2+} concentration for optimal cleavage. This is consistent with our previous findings where we used pre-tRNA and longer model substrates [cf. Figures 5 and 6 in (14)].

Next, we determined k_{obs} and k_{obs}/K^{sto} for cleavage of pMini3bpUG and pMini3bpCG at 800 mM Mg²⁺, a concentration that resulted in optimal cleavage for both substrates (see earlier in text). As shown in Table 1, we detected an almost 10-fold reduction in both these constants due to substitution of U with C at -1. However, it was not possible to determine whether the reduced cleavage of the C₋₁ variant was due to the identity of residue -1 or due to pairing between C₋₁ and G₊₇₃ (Figure 1B). Hence, we generated two new variants, pMini3bpUA and pMini3bpCA, where pairing between residues -1 and +73 was expected to be weaker and absent, respectively. This would allow us to study if k_{obs} and k_{obs}/K^{sto} did differ for substrates with U or C at the -1 position.

An analysis of the cleavage as a function of Mg^{2+} concentration suggested that pMini3bpUA was cleaved more efficiently at position +1 (compared with pMini3bpCA) with no detectable cleavage at -1 (Figure 3B). The determination of k_{obs} and k_{obs}/K^{sto} at two Mg^{2+} concentrations (160 and 800 mM; Table 1) revealed a 3-fold higher k_{obs} value for pMini3bpUA irrespective of Mg^{2+} concentrations. At saturating Mg^{2+} (800 mM), k_{obs}/K^{sto} was also increased by a factor of three for pMini3bpUA suggesting that the K_d is not changed relative to pMini3bpCA. Rather the difference comparing U and C at -1 is related to a change in k_{obs} (Scheme 1).

In conclusion, the identity of residue -1 in *Eco* RPRmediated cleavage contributes to catalysis with model hairpin loop substrate (see also later in text). U carries oxygen at position 4 (O4), whereas C has an exocyclic amine (N4) at this position (Figure 1B). Thus, the O4 of U at -1 likely confers only a minor advantage in RPRmediated catalysis. This is consistent with our previous

Figure 3. Continued

pMini3bpGG ($G_{-1}G_{+73}$, 5.2 µM); pMini3bp*Iso*CG ($IsoC_{-1}G_{+73}$, 3.2 µM); Panel B pMini3bpUG ($U_{-1}G_{+73}$, 0.8 µM); pMini3bpUA $(U_{-1}A_{+73}, 3.1 \,\mu\text{M});$ pMini3bpCA $(C_{-1}A_{+73}, 3.1 \,\mu\text{M});$ Panel C pMini3bp2APG (2AP₋₁G₊₇₃, 5.2 μ M); pMini3bpCG (C₋₁G₊₇₃, 1.6 μ M); pMini3bpInoG (Ino₋₁G₊₇₃, 5.2 μ M); Panel D pMini3bpAG pMini3bpPuG $(A_{-1}G_{+73},$ 5.2 μM); $(Pur_{-1}G_{+73},$ 5.2 µM): pMini3bpDAPG (DAP₋₁ \hat{G}_{+73} , 5.2 μ M); Panel E (cleavage with *Eco* RPR_{G248}) pMini3bpUG $(U_{-1}G_{+73}, 0.8 \,\mu M;$ left v-axis); pMini3bpIsoCG (IsoC-1G+73, 0.8 µM; right y-axis). For the calculations, we used the 5' cleavage fragments and the data are the mean of at least three independent experiments. The bars indicate the experimental errors.

Substrate/[Mg ²⁺]	k _{obs} (min ⁻¹)	$k_{obs}/K^{sto}~(min^{-1}\mu M^{-1})$	$K_d \; (\mu M)$
pMini3bpUG 160 mM 800 mM	0.86 ± 0.12 4.2 ± 0.48	$\begin{array}{c} 0.31 \pm 0.12 \\ 1.6 \pm 0.5 \end{array}$	2.8 2.6
pMini3bpAbasicG 800 mM +1	0.0010 ± 0.000006	0.00038 ± 0.000053	2.6
-1 pMini3bpCG 160 mM 800 mM	$\begin{array}{c} 0.0012 \pm 0.00004 \\ 0.0065 \pm 0.0005 \\ 0.6 \pm 0.3 \end{array}$	0.0010 ± 0.00014 0.0006 ± 0.00012 0.19 ± 0.12	1.2 11 3.2
pMini3bpUA 160 mM 800 mM	$\begin{array}{c} 0.8 \pm 0.06 \\ 4.8 \pm 0.95 \end{array}$	$\begin{array}{c} 0.11 \pm 0.01 \\ 3.7 \pm 0.4 \end{array}$	7.3 1.3
pMini3bpCA 160 mM 800 mM	$\begin{array}{c} 0.29 \pm 0.033 \\ 1.6 \pm 0.2 \end{array}$	$\begin{array}{c} 0.095 \pm 0.015 \\ 1.3 \pm 0.2 \end{array}$	3.1 1.2
pMini3bp <i>Iso</i> CG 160 mM 800 mM	$\begin{array}{c} 0.0046 \pm 0.0003 \\ 0.062 \pm 0.014 \end{array}$	$\begin{array}{c} 0.0013 \pm 0.0003 \\ 0.009 \pm 0.001 \end{array}$	3.5 6.9
pMini3bpAG 800 mM	0.11 ± 0.061	0.034 ± 0.01	3.2
800 mM	0.0088 ± 0.0036	0.0012 ± 0.0002	7.3
800 mM pMini3bp2APG	0.22 ± 0.056	0.099 ± 0.026	2.2
800 mM pMini3bpDAPG 800 mM	0.016 ± 0.007 0.025 ± 0.0048	0.0088 ± 0.003 0.013 ± 0.002	1.8 1.9
pMini3bpPuG 800 mM	0.029 ± 0.0094	0.023 ± 0.007	1.3

Table 1. The kinetic constants kobs and kobs/Ksto for cleavage of the various substrates by wild type Eco RPR

The experiments were performed under single-turnover conditions at different Mg^{2+} concentrations at pH 6.1 as described in 'Materials and Methods'. The final concentration of substrate was >10 nM. The concentration of *Eco* RPR was varied between 0.8 and 29 μ M dependent on the substrate. For details regarding the calculation of K_d, see the main text. The data represent mean and experimental errors calculated from at least three independent experiments. ND = not determined.

finding that the kinetic rate constants for cleavage of other model hairpin substrates with U_{-1}/A_{+73} and C_{-1}/A_{+73} under multiple turnover conditions are similar (13). Moreover, the replacement of U with C at -1 (pMini3bpUA and pMini3bpCA) resulted in a 3-fold effect on the kinetic constants, whereas almost a 10-fold effect was observed comparing cleavage of pMini3bpUG and pMini3bpCG. This indicates that the possibility to form a strong -1/+73 pair affects the kinetics of cleavage.

The oxygen at C2 of U_{-1} and C_{-1} contributes to catalysis

Both U and C carry oxygen at C2 (O2) on the nucleobase (Figure 1C). To understand the influence of O2 on cleavage, we generated a pMini3bp derivative pMini3bp*Iso*CG in which O2 of U (or C) at position -1 is substituted with an exocyclic amine, 2NH₂. *Eco* RPR_{wt} cleaved pMini3bp*Iso*CG preferentially (\approx 97%; Figure 2D) at the canonical site +1 irrespective of Mg²⁺ concentration but at a reduced rate relative to cleavage of pMini3bpUG. Moreover, as observed for the other pMini3bp variants,

cleavage of pMini3bp*Iso*CG required high Mg²⁺ concentration for optimal cleavage (Figure 3A).

Determination of the kinetic constants at 800 mM Mg²⁺ revealed an \approx 70- and >150-fold reduction in k_{obs} and k_{obs}/K^{sto}, respectively, compared with when pMini3bpUG was used. Using these data, K_d was calculated to be just \approx 3-fold higher relative to the K_d value for pMini3bpUG (Table 1). These data suggest that O2 of U (or C) at -1 contributes significantly to *Eco* RPR-mediated catalysis of a model substrate and can, therefore, be considered to be a positive determinant.

Presence of the exocyclic amine $(2NH_2)$ of G at -1 has a negative impact on cleavage at the canonical site +1

Bacteria also have pre-tRNAs with purines at -1; *E. coli* carry $\approx 13\%$ and 8% A and G at -1, respectively (13). Processing studies with *Eco* RPR_{wt} (with and without the C5 protein) and yeast pre-tRNA^{Ser} derivatives show that cleavage between pyrimidines and purines is preferred to cleavage between two purines (31). Therefore, we decided to study the influence of a purine at -1 on substrate

cleavage (all substrates have G at the +1 position; Figure 1). We were particularly interested to understand the impact of the exocyclic amine at C2 on the nucleobase (2NH₂; Figure 1C) in RPR-mediated cleavage. We predicted that the 2NH₂ at -1 (when present) would have a negative impact on cleavage by affecting the kinetic rate constants. Hence, six pMini3bp variants with different purines at -1 were generated (see Figure 1B) and subjected to cleavage by *Eco* RPR_{wt}. As with the substrates discussed earlier in the text, we studied cleavage site recognition and Mg²⁺ dependence and determined the kinetic constants k_{obs} and k_{obs}/K^{sto} at 800 mM Mg²⁺.

As seen with the other pMini3bp variants, cleavage of the substrates with the different purine derivatives at -1required high Mg²⁺ concentration for optimal cleavage (Figure 3A, C and D). Moreover, introduction of G at -1 resulted in substantial cleavage at the alternative site -1 (Figure 2E and F). However, the frequency of cleavage at -1 depended on the Mg²⁺ concentration; there was less cleavage at -1 at lower Mg²⁺ concentrations (Supplementary Figure S2). When A was present at -1, cleavage occurred preferentially at the correct site +1 (Figure 2E and G). The substrate carrying inosine at -1was also cleaved mainly at +1 ($\approx 2\%$ cleavage at -1; Figure 2G). As pMini3bpAG, this substrate lacks the 2NH₂ group (compared with G at -1; pMini3bpGG versus pMini3bpInoG and pMini3bpAG; see Figure the pMini3bp2APG 1C). In contrast, and pMini3bpDAPG (with 2-AminoPurine and 2:6-DiAminoPurine at -1, respectively, Figure 1B and C) substrates were both cleaved more frequently at the alternative site -1 (Figure 2F; For pMini3bp2APG and pMini3bpDAPG there was no variation in cleavage site selection as a function of Mg^{2+} with the possible exception in the cleavage of pMini3bp2APG at high Mg²⁺, Supplementary Figure S2). As in the substrate with G at -1, both pMini3bp2APG and pMini3bpDAPG carry the 2NH₂ group on the nucleobase. Moreover, a slight increase in the frequency of cleavage at -1 was detected when comparing cleavage of pMini3bpAG and pMini3bpPuG (Figure 2G). Cumulatively these data indicate that deletion of $6NH_2$ at -1 in a model substrate context marginally affect cleavage site recognition, whereas an exocylic amine $(2NH_2)$ at C2 on the purine base has a negative impact on the selection of the cleavage site.

Determination of the kinetic constants, k_{obs} and k_{obs}/K^{sto} (Table 1) revealed that both were lower for substrates with A or G at -1 relative to those with U or C (cf. pMini3bpAG or pMini3bpGG versus pMini3bpUG or pMini3bpCG). This is particularly apparent when G is present at -1 (pMini3bpGG). In this case, there is an almost 500- and >1000-fold reduction (for cleavage at +1) in k_{obs} and k_{obs}/K^{sto} , respectively, compared with cleavage of pMini3bpUG. For the substrate with A at -1 (pMini3bpAG versus pMini3bpUG), k_{obs} was reduced ≈ 40 -fold and $k_{obs}/K^{sto} \approx 50$ -fold. When the 2NH₂ of G₋₁ (pMini3bpInoG versus pMini3bpGG) was deleted both k_{obs} and k_{obs}/K^{sto} increased. By contrast, introduction of an exocyclic amine at C2 (pMini3bp2APG and pMini3bpDAPG) resulted in a decrease compared with cleavage of the substrate with A at -1 (pMini3bpAG; Table 1). Deletion of 6NH₂ from the A at -1 (Figure 1C) decreased mainly the k_{obs} (3- to 4-fold), whereas k_{obs}/K^{sto} changed only marginally (cf. cleavage of pMini3bpAG and pMini3bpPuG; Table 1). A comparison of K_d values revealed only a modest increase (2- to 3-fold) for the substrate with G at -1 relative to the substrates with U or C at -1. These data emphasize that the presence of 2NH₂ on the purine base at -1 has a negative impact on the kinetics of cleavage, in agreement with our prediction. The presence of the exocyclic amine at C6 (6NH₂) of A at -1, however, appears to have only a modest influence on cleavage efficiency.

In conclusion, RPR-mediated cleavage between purines in a model substrate context is unfavourable. An exocyclic amine at C2 on the purine base has a negative impact on cleavage both with respect to accuracy and kinetics of cleavage at +1. From our data, it also appears that a chemical group at C6 contributes to catalysis.

Influence of the protein subunit of *Eco* RNase P

The RNase P protein, C5, binds to the 5' leader of the substrate, and it affects binding and affinity of catalytic important Mg²⁺ (32-37). Also comparing cleavage with and without C5 of yeast pre-tRNA^{Ser} with G at -1 indicates that the presence of the C5 protein affects cleavage site selection (31). Therefore, the C5 protein is a positive factor for cleavage and selection of the cleavage site. Hence, we inquired whether addition of C5 to the reaction also influenced the choice of cleavage site when cleaving the model hairpin loop substrate with G (or A) at -1. Figure 2E shows that the reconstituted *Eco* RNase P holoenzyme cleaved pMini3bpGG at +1 with no apparent cleavage at -1. No change in choice of cleavage site was observed for cleavage of the substrate having A at -1 with RPR or with the reconstituted *Eco* RNase P holoenzyme. However, the substrate with A at -1 was cleaved more efficiently compared with cleavage of the substrate with G at -1 also in the presence of the C5 protein (cf. lanes 5 and 6 Figure 2E).

These data show that the C5 protein suppressed cleavage at the alternative site -1 also in the case of model substrates carrying G at -1. This further indicates that C5 has a strong positive effect on the cleavage site recognition process.

Interaction between residue -1 and 248 in the RPR

The A at position 248 in the RPR (Figures 1A and 4) has been suggested to interact with the -1 residue in the substrate (9). In the crystal structure of RNase P in complex with tRNA, which represents a post cleavage state, A₂₄₈ is positioned near the 5' termini of the tRNA [(11); Figure 5A]. Nucleotide analogue-modification interference studies suggest that the Hoogsteen surface of A₂₄₈ plays an important role in the interaction with the substrate (38). As shown here, substitution of O2 of U (or C) at -1 with NH₂ affects the kinetics of cleavage (see earlier in text; pMini3bp*Iso*CG). Previous data reveal that the 2'OH of residue -1 contribute to catalysis [(7) and



Figure 4. Secondary structure model of *Eco* RPR and a model of the N_{-1}/A_{248} interaction. (A) Secondary structure of *Eco* RPR_{wt} according to Massire *et al.* (39). Residue A_{248} that was changed to G is indicated in grey. (B) A putative model of the N_{-1}/A_{248} interaction. Black and grey residues mark substrate and RPR residues, respectively, and the dashed arrow marks the scissile bond.

references therein]. The O2 and 2'OH of residue -1 are exposed on the same surface (Figure 5B; see later in text). Together this opens for the possibility that N7 and 6NH₂ of A₂₄₈ are hydrogen-bonded to the 2'OH and O2 of

residue -1, respectively, in the transition state (Figure 4B). If this hypothesis is valid, changing of A_{248} to G in the RPR would restore the interaction between IsoC at -1and 248 by forming hydrogen bonding between 2NH₂ of IsoC and O6 of the G at 248. As a consequence, the rate of cleavage of pMini3bpIsoCG should increase using an RPR variant with G at 248 compared with cleavage with RPR_{wt}. To test this prediction, we generated Eco RPR_{G248}. Structural probing revealed no significant overall structural difference compared with Eco RPR_{wt} [Supplementary Figure S3: see also (19). We note the appearance of additional cleavages, located between well-established lead(II)-induced sites IV and V, when Eco RPR_{G248} was subjected to RNase T1 cleavage. This might indicate a slight change in the structure in this region].

As shown in Figure 2C and D, Eco RPR_{G248} cleaved pMini3bpUG and pMini3bpIsoCG preferentially at the correct site, +1. For pMini3bpIsoCG, we also observed $\approx 9\%$ cleavage at -1, which was higher at lower Mg²⁺ (Supplementary Figure S2). Moreover, compared with *Eco* RPR_{wt} , the G₂₄₈ variant showed similar Mg²⁺ pMini3bpUG profiles in cleavage of and pMini3bpIsoCG (Figure 3E). Determinations of the apparent rate constants (k_{app}) at 800 mM Mg²⁺ revealed that *Eco* RPR_{G248} cleaved these two substrates with slightly reduced rates relative to the wild type (Table 2). Importantly, the rate of cleavage did not improve as predicted. Rather, the rate was lower with Eco RPRG248 compared with Eco RPR_{wt}. These findings, therefore, suggest that 6NH₂ of A₂₄₈ is not engaged in hydrogen bonding with O2 of U (or C) at -1 or at least not in the context of a model substrate (see also the 'Discussion').

DISCUSSION

Here we provide data that the nucleobase, the identity and specific chemical groups of the nucleobase at position -1in a model substrate contribute significantly to catalysis in the E. coli RPR alone reaction. In particular, our data suggest that the carbonyl oxygen (O2) at C2 on C and U at -1 is a positive factor, whereas a purine (A or G) at -1 and the presence of an exocyclic amine at C2 (2NH₂) have a negative impact on the kinetics and accuracy of cleavage. The nucleobase at -1 is, however, not essential for cleavage although it affects the kinetic rate constants, k_{obs} and k_{obs}/K^{sto} . It contributes to catalysis by stabilizing the transition state, at least in the context of a model substrate, and this corresponds to ≈ 5.1 kcal [using k_{obs} values for cleavage of pMini3bpAbasicG and pMini3bpUG at +1 and $\Delta\Delta G = -RTlnk_{obs}(\Delta_{-1})/2$ $k_{obs}(U_{-1})$; (43)]. Of these 5.1 kcal, the contribution of the O2 (with U at -1) amounts to 2.6 kcal $[\Delta \Delta G = -RTlnk_{obs}(IsoC_{-1})/k_{obs}(U_{-1})]$, which corresponds roughly to one hydrogen bond. In accordance with this is that the presence of $2NH_2$ on a purine at -1destabilizes the transition state with up to 2 kcal (using kobs values for cleavage of pMini3bpInoG and pMini3bpGG at +1). Interestingly, Baidya et al. (44) used modified pyrimidines at position 17 (5' residue of



Figure 5. The structure of the RPR active site and model of the RNase P cleavage site. (A) The active site of RPR from the PDB structure 30KB (11). Residues G_{+1} of the 5' matured tRNA and A_{248} (E. coli numbering) are marked as indicated. The grey spheres corresponds to the Mg^{2+} ions observed in the structure, and the blue structure represents the 5' leader that was soaked into the crystal. (B) Model of the RNase P cleavage site with C_{-1}/G_{+73} and G_{+1}/C_{+72} . The model is part of the SRP RNA structure [PDB code 1LNT; (40)]. The phosphorous atom to be attacked is marked with the grey dashed double arrow. The light blue sphere represents Mg^{2+} bound in the minor groove and is positioned at a distance of 6.7Å relative to phosphorous atom I, and 7.27 Å relative to phosphorous atom II; I and II are referred to as P(+1) and P(-1) in the main text. This distance allows for a nucleophilic attack on P(+1) by an activated H₂O coordinated to a divalent metal ion (41,42). The boxed hairpin structure shows the sequence of SRP RNA that mimics the RNase P cleavage site. The blue arrow marks the cleavage site. (C) Model of the RNase P cleavage site showing the conformational change in the substrate that facilitates the nucleophilic attack on P(+1). The model was prepared using the

the cleavage site) in the hammerhead ribozyme and showed that an abasic residue at position 17 almost abolished cleavage. They also suggested that O2 of C17 makes an important contribution to catalysis by stabilizing the transition-state structure. The recently solved structure of the Schistosoma mansoni hammerhead ribozyme suggests hydrogen bonding between O2 of C17 at the cleavage site and 6NH₂ of A13 that presumably helps to stabilize the transition state (45). However, the recent crystal structure of bacterial RNase P in complex with tRNA (11) does not reveal any structural information about the nature of the interaction between residue -1 in the substrate and the RPR (see later in text). We conclude that the O2 (in the case of U or C) at position -1 in a model substrate acts as a positive determinant in Eco RPR-mediated catalysis whereas 2NH₂ (in the cases of U, C or purines) at this position has a negative impact.

Comparing cleavage of pre-tRNA versus model hairpin loop substrates

Studies where pre-tRNA^{Asp} was used (9,46) show modest changes in both k_{obs} and K_d (except for the variant with C at -1 where K_d increased 230-fold, possibly owing to the C at -1 being paired with the G at +73 in the substrate). In contrast, cleavage of model substrates (this report) caused an almost 500-fold change in k_{obs} (for G at -1) with only small changes in K_d . Moreover, model substrates with G at -1 (or purines with $2NH_2$ or where the nucleobase had been deleted at -1) resulted in significant cleavage at -1 whereas the other -1 variants were cleaved mainly at +1 (Figure 2 and Supplementary Figure S2). This is in contrast to the U at -1 substitution in pretRNA^{Asp}, which did not change the cleavage site.

Table 2. Apparent rates, k_{app} , for cleavage of pMini3bpUG and pMini3bpIsoCG with *Eco* RPR_{wt} and *Eco* RPR_{G248}

Substrate	Eco RPR variant		
	Wt	G ₂₄₈	
pMini3bpUG +1	1.5 ± 0.03	0.98 ± 0.06	
pMini3bp <i>Iso</i> CG +1 -1	$\begin{array}{c} 0.026 \pm 0.002 \\ \text{ND} \end{array}$	$\begin{array}{l} 0.011 \pm 0.0004 \\ 0.001 \pm 0.00007 \end{array}$	

 $k_{app} = pmol$ cleaved per min. The experiments were performed in buffer C containing 800 mM Mg²⁺ at 37°C. The concentrations of RPRs and substrates were 4 and 0.02 μ M, respectively. The data represent mean and experimental errors calculated from at least three independent experiments.

ND = not determined.

Figure 5. Continued

structure shown in panel B and rotating the P-O5' phosphoester bond (marked with a blue arrow) 50°. The rotation displaces the pro-Sp nonbridging oxygen and exposes the P(+1)-atom to an inline hypothetical nuclephilic attack by a Mg^{2+} -activated H₂O (grey dashed arrow). The grey arrows mark the 2'OH and O2 of C₋₁, and I and II refer to the changes in cleavage rates as a result of changing O2 to 2NH₂ and O4 to 4NH₂, respectively. However, cleavage of pre-tRNA^{Asp} at the alternative site -1 was observed when residue -1 was changed together with replacement of the 2'OH with 2'H at -1, i.e. two cleavage site determinants had been changed (9,46). A comparison of the structures of pre-tRNA^{Asp} and pMini3bpUG reveals striking similarities at and near their respective cleavage sites

$$\bullet \bullet \bullet A_{-3}A_{-2}U_{-1}G_{+1}G_{+2} \bullet \bullet \bullet C_{+71}C_{+72}G_{+73}C_{+74}C_{+75}A_{+76}\bullet$$

[black circles correspond to residues that are not shown and the canonical cleavage site is between U_{-1} and G_{+1} ; Figure 1 and (46)]. The T-loop/stem, known to affect site selection (14,22), is absent in the pMini3bp model substrates. This makes them dependent on the remaining determinants, e.g. the -1 residue (Figure 1A). Together this argues that the impact of the -1 residue in a pre-tRNA (all ribo) context is obscured by the presence of several residues and regions that interact with the RPR on formation of the RPR-substrate complex. Nevertheless, for both model and pre-tRNA substrates, the base at -1 as well as its identity influences cleavage efficiency and site recognition but it is not essential for cleavage. However, as shown in this report using model substrates, specific chemical groups on the -1 nucleobase influence both transition-state stabilization and accuracy in Eco RPRmediated cleavage.

We have reported elsewhere that the presence of an exocyclic amine $(2NH_2)$ on the nucleobase at +1 plays an important role for cleavage at the correct (+1) site (17). Our present data show that replacement of U (or C) at -1 with purines with $2NH_2$ in a model substrate had a negative impact on the kinetic constants with significant cleavage at the alternative site -1. On the basis of these data we suggest that $2NH_2$ (in the case of G) at -1functions as a positive determinant for cleavage at -1, i.e. at the site immediately 5' of a purine carrying $2NH_2$ (see Figure 1). At the same time, it acts as a negative factor for cleavage at +1. This would explain why the efficiency of cleavage between two G residues is low (31). Most bacterial tRNA^{His} precursors [and pre-tRNA^{SeCys}; Ref (47)] carry G at both -1 and +1, and this model also provides one reason to why bacterial RPR cleaves pre $tRNA^{His}$ and pre-tRNA^{SeCys} at -1 and not between two G-residues [reviewed in (6)]. However, we have to consider that addition of the C5 protein, which interacts with the substrate 5' leader (see earlier in text), resulted in cleavage mainly at +1 even with G at -1 in the model substrate (Figure 2E). It is conceivable that this is due to G_{-1} in the model substrate is not being paired with a C at +73, in contrast to pre-tRNA^{His} and pre-tRNA^{SeCys} in which the G at -1 is paired to a C. The presence of a G_{-1}/C_{+73} pair in different precursor substrates plays an important role for cleavage at -1 (6). Also, E. coli pre-tRNA(valU) and pre-tRNA(thrW) carry G at -1 (48) but in those two cases G_{-1} is unlikely to pair with the discriminator base (which is A in both cases) at position +73. Another factor to consider is that the difference in cleavage site selection of the G_{-1} model substrate with and without the C5 protein may be an effect of the higher Mg²⁺ concentration used in the 'RPR alone reaction' (Supplementary Figure S2). Evidently, the role of the C5 protein in the cleavage site recognition process in the processing of different precursors requires further studies. The present study together with our previous data (17), however, indicates the importance of the presence of G (and the $2NH_2$) 3' of the scissile bond at least in the '*Eco* RPR alone reaction'. In this context, we note the importance of the exocyclic amine of a guanosine marking the cleavage site in the reaction catalyzed by the group I ribozyme (49), indicating similarities with group I RNA and RPR-mediated cleavage.

Interaction between the -1 residue in the substrate and *Eco* RPR and positioning of Mg²⁺ at the cleavage site

A₂₄₈ has been suggested to play a key role in the interaction with residue -1; this interaction is referred to as the N_{-1}/A_{248} interaction [(9,38,46,50); see also (6) and Figure 1A]. The crystal structure of the RNase P-tRNA complex, which represents the post-cleavage state, does not provide any information about the interaction between the RPR and the -1 residue (11). Although it cannot be excluded that the Watson-Crick surface of A_{248} is involved in pairing with residue -1 as suggested elsewhere (9,46), we consider this unlikely. The reasons are: (i) as mentioned earlier in the text, the identity of residue -1 varies and must be taken into account; (ii) N3-methyl-U at -1 in a model substrate did not affect the kinetic rate constants, k_{obs} and k_{obs}/K^{sto} , to any significant extent compared with a substrate with an unmodified U at -1 (51); and (iii) a model substrate with C_{-1}/G_{+73} is cleaved with increased frequency at -1 (and not a decrease in cleavage at the alternative site -1, as would be predicted if it was base pairing between -1and 248) when A₂₄₈ in Eco RPR_{wt} was substituted with a G (19). Alternatively, given that the Hoogsteen surface of A₂₄₈ is facing the tRNA 5'-end in the crystal structure (Figure 5A) it is possible that N7 and $6NH_2$ interact with specific groups of the -1 residue. This would be consistent with nucleotide analogue-modification interference studies, which suggest that the Hoogsteen surface of A₂₄₈ plays an important role on interaction with the substrate (38). Here a possibility is that $6NH_2$ and N7 of A_{248} interact with O2 (in the case of U and C) and the 2'OH (Figure 4B; see also below) of residue -1, respectively. The lack of rescue in cleavage of pMini3bpIsoCG with RPR_{G248} argues against this alternative; however, we cannot conclusively exclude this possibility (see later in the text).

RPR catalysis depends on divalent metal ions, preferentially Mg^{2+} , and two Mg^{2+} are positioned close to the tRNA 5'-end in the RNase P-tRNA complex (Figure 5A). In our model, the Mg^{2+} that activates the nucleophilic water is positioned 6.7 Å from the phosphorous between -1 and +1 [P(+1) in Figure 5B] in the substrate (7,52,53). The distance between this Mg^{2+} and the phosphorous between -1 and -2 [P(-1) in Figure 5B] is just 0.57 Å longer. Hence, changing the identity of residues or chemical groups in the vicinity of the cleavage site can affect the charge distribution. This would then shift the positioning of Mg^{2+} that influences the rate of cleavage and site selection. This model is consistent with our data where substitutions of the nucleobase and chemical groups at -1 or at position 248 in the RPR affected the rate constants and site selection [see also (19); it also raises the possibility that the change in charge distribution owing to the presence of G at 248 is why we did not observe rescue of cleavage of pMini3bp*Iso*CG with *Eco* RPR_{G248}]. In addition, deletion of the exocyclic amine (2NH₂) of the G at +1 in a model substrate influenced the charge distribution at the cleavage site (17). Substitutions in the T-loop also influenced both the rate and site of cleavage in metal(II)-ion induced hydrolysis of yeast tRNA^{Phe} (54,55).

A model for displacement of the -1 residue in the *Eco* RPR-substrate complex

For bacterial RPR, the 2'OH at -1 plays an important role for catalysis (6–8). It contributes 2.3 kcal to the stabilization of the transition state in cleavage of a hairpin loop model substrate using the same formula discussed earlier in text and data from Brännvall and Kirsebom (51). As reported here, the contribution of the carbonyl oxygen O2 of C (or U) at -1 in a model substrate to transition-state stabilization was 2.6 kcal (see earlier in text). The 2'OH and O2 of C at -1 in a model of the cleavage site were exposed on the same surface (marked with grey arrows in Figure 5C). We cannot exclude that one or both these groups form hydrogen bonds with the RPR in the transition state. However, available data suggest that the 2'OH at -1 acts as an outer (or inner) sphere ligand for Mg^{2+} [(24,56–59); but see also (46) for an alternative interpretation]. On the basis of this, it is therefore conceivable that also the carbonyl oxygen of C (or U) at -1 is involved in Mg²⁺ binding although the crystal structure does not confirm this. Nevertheless, on RPRsubstrate complex formation, where residue +73 pairs with U294 in the RPR, the C_{-1}/G_{+73} (when present) opens. This conformational change is likely accompanied with displacement of the C at -1 and positioning of A_{248} such that it stacks over the $G_{\pm 1}/C_{\pm 72}$ base pair (Figure 5A). We propose that displacement of the -1residue is the result of a 50-degree rotation (Figure 5C) such that O2 (in the case of U or C at -1) and the 2'OH are facing away from the scissile phosphate. As a consequence, the Mg^{2+} -activated H_2O bound to the substrate in the vicinity of the phosphorus [P(+1)] to be attacked is positioned for an in-line attack [Figure 5; Ref (7)]. At the same time, the 2'OH at -1 is prevented from attacking the phosphorus, which would lead to cleavage products with faulty ends, i.e. a 5'OH and a 2'; 3' cyclic phosphate (24). This model is consistent with a conformationalassisted mechanism of cleavage but needs to be confirmed by a structure of RNase P in complex with its precursor substrate.

SUPPLEMENTARY DATA

Supplementary Data are available at NAR Online.

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