Supplementary Materials

NSMB-A20845C

Title: Aminoacylation of tRNA with phosphoserine for synthesis of cysteinyl-tRNA^{Cys}

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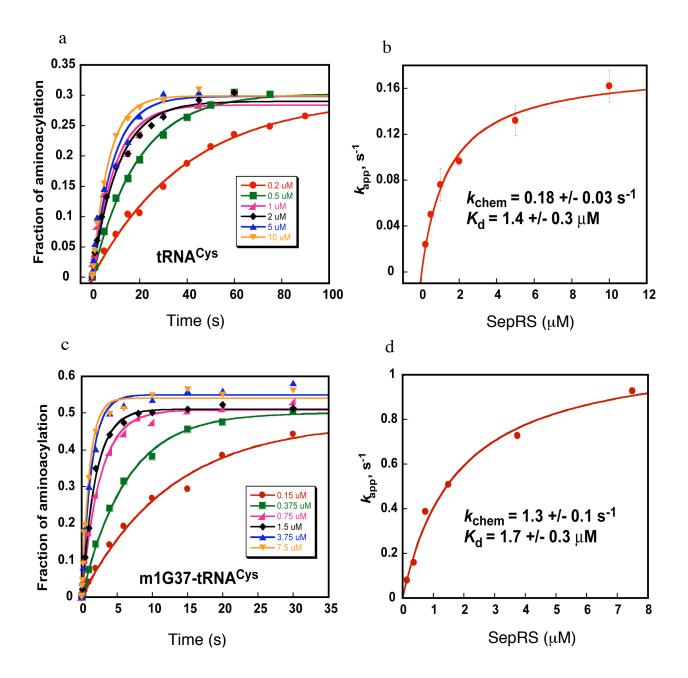


Figure 1 Single turnover kinetics of aminoacylation of *M. jannaschii* tRNA^{Cys} by *M. jannaschii* SepRS. (A) Time courses of aminoacylation of unmodified transcript by SepRS (0.2 to 10 μ M). (B) Replot of $k_{app}s$ vs SepRS concentrations to a hyperbolic equation. (C) Time courses of aminoacylation of m1G37-transcript by SepRS (0.15 to 7.5 μ M). (D) Replot of $k_{app}s$ vs SepRS concentrations to a hyperbolic equation. The derived k_{chem} and K_d were the average of 2 determinations. The ± term refers to standard deviations.

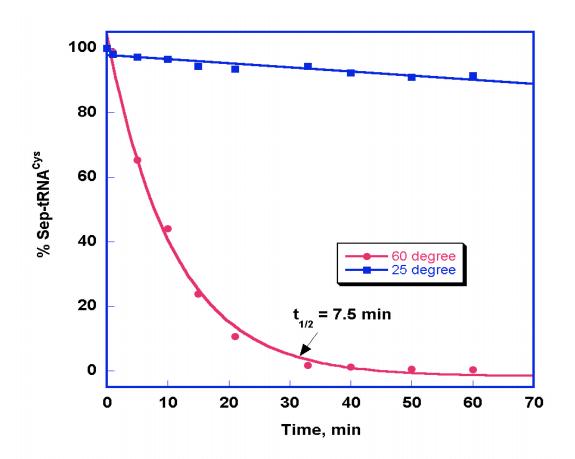


Figure 2 Stability of *M. jannaschii* Sep-tRNA^{Cys} measured at 60 °C (red) and at 25 °C (blue). The tRNA transcript was labeled with α -³²P-A76 (5 μ M) and was aminoacylated with Sep by SepRS. After phenol extraction and ethanol precipitation, Sep-tRNA^{Cys} was resuspended in 1 x aminoacylation buffer (30 mM NaAc, pH 6.0, 20 mM KCl, 10 mM MgCl₂, and 5 mM DTT) and incubated at the specified temperatures. At specific time points, an aliquot was removed and digested with P1 nuclease. Deacylation of Sep-tRNA^{Cys} was monitored by PEI cellulose plates.

	1	2	3	4	5
Mj tRNA (10 μ M)	+	-	+	-	-
Sep (2 mM)	+	+	+	-	-
ATP (10 mM)	+	+	+	-	+
ECysRS (15 μM)	+	+	+	+	+
SepCysS-His ₆ (15 μM)	+	+	-	+	+
ECysRS → SepCysS → -His ₆	•	-		-	•

Figure 3 Lack of pull-down of *E. coli* CysRS by *M. jannaschii* SepCysS analyzed by SDS-PAGE.

tRNA	M.j SepRS vs. $M.j$ tRNA ^{Cys(a)}				E.coli CysRS vs. E coli tRNA ^{Cys}		
	$K_{\rm m}$ (μ M)	$k_{ m cat} \ ({ m s}^{-1})$	$k_{\text{cat}}/K_{\text{m}}(\mu \mathbf{M}^{-1}\mathbf{s}^{-1})$	relative	$k_{ m cat}/K_{ m m} \ (\mu {f M}^{-1}{f s}^{-1})$	relative	
WT	0.97 ± 0.01	0.24 ± 0.04	0.25 ± 0.04	1	$0.99 \pm 0.06^{(b)}$	1	
U73G	1.3 ± 0.3	0.01 ± 0.001	0.008 ± 0.002	0.03	$(7 \pm 0.1) \times 10^{-6}$ (b)	7×10^{-6}	
G34C	19 ± 5	0.08 ± 0.01	0.004 ± 0.001	0.02	$(6 \pm 0.3) \times 10^{-4}$ (b)	6 × 10 ⁻⁴	
C35U	7.0 ± 0.4	0.10 ± 0.02	0.014 ± 0.002	0.06	(3 ± 0.1) × 10 ⁻³	3×10^{-3}	
A36G	22 ± 2	0.17 ± 0.01	0.008 ± 0.001	0.03	(2 ± 0.2) × 10 ⁻³	2 x 10 ⁻³	
G37A	11.4 ± 0.7	0.11 ± 0.01	0.01 ± 0.001	0.04	Not determined		
A38G	2.9 ± 0.2	0.04 ± 0.01	0.014 ± 0.004	0.06	Not determined		
G15C-C48G	3.7 ± 0.8	0.17 ± 0.03	0.05 ± 0.01	0.2	0.02 ± 0.02	0.02	

Table 1. Steady-state kinetics

(a) Transcripts of m1G37-*Mj* tRNA^{Cys} were used except G37A.
(b) Data from Liu *et al.* (2007).