TABLE 1. mRNA half-life determination in S1-depleted cells

		half-life (min) <sup>a</sup>					
	pn	pnp <sup>+b</sup>		Δ <i>pnp-751</i> <sup>b</sup>			
mRNA	ara	glu	ara	glu			
pnp <sup>c</sup> glnS	1.8 2.3	6.4 2.3	5.5 2.1	6.4 2.2			

<sup>&</sup>lt;sup>a</sup>Calculated as detailed in Materials and Methods; the reported results are the average of two independent determinations.

TABLE 2. mRNA half-life in S1-overexpressing cells

		pnp <sup>+a</sup>			Δpnp-751 <sup>a</sup>		
		half-life	e (min) <sup>c</sup>		half-life (min) <sup>c</sup>		
mRNA	R.A. <sup>b</sup>	-IPTG	+IPTG	R.A. <sup>b</sup>	-IPTG	+IPTG	
pnp <sup>d</sup>	2.2	2.3	27.9	0.6	4.3	10.7	
pnp-deaD <sup>e</sup>	10.8	3.4	17.1	1.8	2.2	23.0	
cspE	1.0	4.0	29.8	0.8	4.9	6.7	
glnS	1.1	3.1	37.6	0.7	1.9	3.4	
glyA	1.1	4.6	>48	0.3	4.2	10.8	
rpsO <sup>f</sup>	1.9	2.8	20.6	1.5	2.3	3.7	

<sup>&</sup>lt;sup>a</sup>Cultures of C-1a  $(pnp^+)$  and C-5691  $(\Delta pnp-751)$  with plasmids pQE31-S1 and pREP4 grown and experiment performed as detailed in Figure 7 legend.

<sup>&</sup>lt;sup>b</sup>Cultures of C-5698 ( $pnp^+$ ) and C-5707 ( $\Delta pnp$ -751) grown as detailed in Materials and Methods and in the legends of Figures 4 and 5.

<sup>°</sup>For  $pnp^+$ , the sum of 2.25- and 2.5-kb mRNA and for  $\Delta pnp$ -751 strain, the sum of 1.3-, 0.7-, and 0.3-kb signals were considered for pnp half-life calculation.

<sup>&</sup>lt;sup>b</sup>Relative abundance, calculated as the ratio between mRNA amounts in induced and noninduced cultures 60 min after IPTG addiction.

<sup>&</sup>lt;sup>c</sup>Calculated as detailed in Materials and Methods, the reported half-lives are the average of at least two independent determinations in all cases but the cspE and glyA mRNA half-lives in  $pnp^+$  cells without IPTG, which are the results of a single determination. <sup>d</sup>For  $pnp^+$ , the sum of 2.25- and 2.5-kb mRNA and, for  $\Delta pnp-751$ , the sum of 1.3-, 0.7-, and 0.3-

For  $pnp^+$ , the sum of 2.25- and 2.5-kb mRNA and, for  $\Delta pnp$ -751, the sum of 1.3-, 0.7-, and 0.3-kb signals were considered for half-life calculation.

<sup>&</sup>lt;sup>e</sup>Signals corresponding to 5.4- and 3.4-kb-long RNAs (terminating at the terminator *deaDt*; Zangrossi et al. 2000) were considered for half-life calculation in the  $pnp^+$  and  $\Delta pnp-751$ , respectively.

<sup>&</sup>lt;sup>f</sup>The sum of the two signals detected (see Figure 7) was considered for half-life calculation in both the  $pnp^+$  and  $\Delta pnp-751$ .