

Erratum

A Transcriptional Barrier in the Regulatory Region of the Tryptophan Operon of *Escherichia coli*: Its Role in the Regulation of Repressor-independent RNA Synthesis

Peter H. Pouwels and Hans Pannekoek

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Table 1 should read as follows:

Table 1. The effect of the bacterial growth rate on the synthesis of *trp*RNA

Bacterial strain	Carbon source	Tryptophan (µg/ml)	Bacteria per ml	Divisions per h	<i>trp</i> RNA (counts/min) per 20 µl	[³ H]RNA (ml)	Yield of RNA (%)	Genome equivalents per bacterium	<i>trp</i> RNA/genome (counts/min × 10 ⁵)
<i>HfrH, trpR</i>	1 glucose	—	2.8 × 10 ⁸	1.0	510	1.50	60	2.15	2.0
	2 glycerol	—	3.0 × 10 ⁸	0.52	91	2.50	66	1.6	0.74
	3 proline	—	5.6 × 10 ⁸	0.24	152	1.50	61	1.4	0.47
	—	—	9.6 × 10 ⁸	0.22	130	2.10	35	1.4	0.57
	4 glucose	50	1.1 × 10 ⁹	0.84	338	1.50	63	1.95	1.9
W3110 <i>ΔtrpED102 trpR</i>	5 proline	50	2.1 × 10 ⁹	0.48	114	1.50	57	1.6	0.45
	6 glucose	25	6.6 × 10 ⁸	0.84	812	1.70	31	1.95	3.4
	25	2.8 × 10 ⁸	0.62	750	2.50	71	1.7	5.4	
	7 glycerol	25	3.6 × 10 ⁸	0.29	536	2.50	68	1.5	3.7
	8 proline	25	2.0 × 10 ⁸	0.11	255	2.50	57	1.3	4.3
DX-I	25	5.6 × 10 ⁸	0.22	448	1.80	40	1.4	3.55	
	9 glucose	—	2.0 × 10 ⁸	0.44	536	1.50	55	1.6	2.9
	10 proline	—	4.2 × 10 ⁸	0.12	82	1.50	51	1.3	0.43

P.H. Pouwels