

Nucleotide sequence of the HU-1 gene of *Salmonella typhimurium*

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Bacterial cells contain a class of small, basic, heat stable "histone-like" proteins which are thought to condense the chromosome into a tightly folded nucleoid structure (1). The most abundant of these proteins, HU, is present in *E. coli* and *Salmonella* as a heterotypic dimer whose subunits are encoded by the hupA and hupB genes (2,3). We report here the nucleotide sequence of the hupB gene of *Salmonella typhimurium* and its comparison with the *E. coli* sequence. The protein predicted by the *Salmonella* sequence is highly conserved and differs from the *E. coli* protein only at amino acid position 8 despite the presence of 20 other conservative changes in the sequence.

TTCATGGGT	CCTGCCACAT	TAACGGGT	TGTAAGGCC	TGGCAGGCT	GATAAACIG	1-60
		G	G	T		
CCCCCGGCG	GTAATCGAA	GATTCAAGG	CGATATAAAT	TATAAAGAGC	AAGAGAGA	61-119
T	T	C T A G	A		G	
M N K S Q L I E* K I A A G A D I S K A A						
GTC AAT AAA TCT CAA CTG ATC GAA AAA ATT GCT GCA GGG GCT GAT ATC TCT TGT AGG GCT GCG						120-179
T	C	G			A	
A G R A L D A I I A S V T E S L K E G D						
GCT GGA CGT GCG TTA GAT CGG ATT ATT GCT TCT GGT ACC GAA TCT CTG AAA GAA GGG GAT						180-239
C			T	C A T		
D V A L V G F G T F A V K E R A A R T G						
GAC GTT GCA CTG GTA GCC TTT GGT ACT TTT GCT GTT AAA GAG CGT CCT GCC CGT ACT GGT						240-299
T A		T		C		
R N P Q T G K E I T I A A A K V P S F R						
CCC AAC CGG CAA GGT AAA GAG ATC ACC ATC GCC GCT GCG AAA GIG CGG ACT TTC CGT						300-359
G C			T T	A C		
A G K A L K D A V N						
GCA GGT AAA CGG CTG AAA GAC CGG GTC AAC TAACGGT						360-419
A			TG	AGT	G	
AGTACAAGG	CCCATCAACT	GATGTGCGT	TITIATGGC	GATTCGGACT	TTCGTGCG	420-478
T	T		T - T	TGA	C T C	

References:

1. Drlica, K., and Rouviere-Yaniv, J. Microbiol Rev. 51,301-319. (1987)
2. Kano, Y. et al., Mol. Gen. Genet. 201, 360-362 (1985).
3. Kano, Y. et al., Mol. Gen. Genet. 209, 408-410 (1987).