

Location of the *hisGDCBHAFI* Operon on the Physical Map of *Escherichia coli*

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The *his* operon, located at 44 min on the genetic map of *Escherichia coli* (1), encodes the eight enzymes concerned with the biosynthesis of the amino acid histidine (8). The *his* cistrons were shown by complementation assays of different auxotrophic strains to reside on two contiguous *Hind*III fragments, of 5.3 and 4.5 kb, derived from strain W3110 (2, 3, 5). The structural organization of the *his* operon was

restriction sites for *Eco*RV that generate two fragments of less than 100 bp which probably went undetected during construction of the physical map (7) (cf. Fig. 1B and C). These data establish the exact location of the *his* operon on the Kohara physical map. Moreover, the relative distance from the assigned *his* operon to the neighboring marker of the Kohara physical map, *sbcB* (Fig. 1A), appears to be

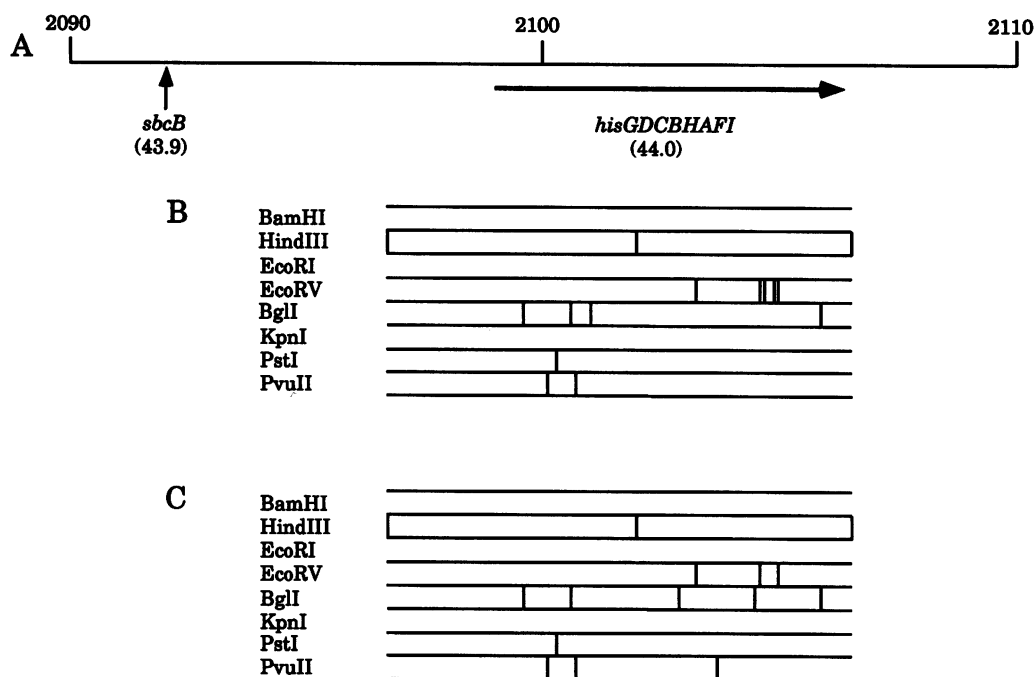


FIG. 1. Correlation of the physical map location of the *E. coli his* operon with its genetic map location and alignment with the physical map of *E. coli*. (A) The physical map location of *sbcB* in kilobase pairs was derived from reference 7. The genetic map locations of *sbcB* and *hisGDCBHAFI* in minutes are indicated by the numbers in parentheses as assigned in reference 1. The direction of transcription of the *his* operon is indicated by the arrow. (B) Restriction map of the DNA fragment containing the *hisGDCBHAFI* operon. (C) A section of the Kohara physical map (7) around kb 2100.

revealed by sequencing an 8-kb DNA region (4) derived from plasmid pHC9800 (6). The restriction pattern of the inserted *E. coli* DNA fragment deduced from the DNA sequence and experimentally verified matches the corresponding section of the Kohara restriction map (7) around kb 2100 for five of the eight enzymes (*Bam*HI, *Hind*III, *Eco*RI, *Kpn*I, and *Pst*I) (Fig. 1). The only exceptions are that the sequence lacks a *Bgl*II and a *Pvu*II site and that there are two additional

consistent with that derived from the published linkage map of *E. coli* K-12 (1, 7).

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