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Physical Locations of *bglA* and *serA* on the *Escherichia coli* K-12 Chromosome

KATHRYN O'DAY,[1] JANE LOPILATO,[2]* AND ANDREW WRIGHT[1]

Department of Molecular Biology and Microbiology, Health Sciences Campus, Tufts University School of Medicine, Boston, Massachusetts 02111,[1] and Department of Biology, Simmons College, 300 The Fenway, Boston, Massachusetts 02115[2]

We determined that the *bglA7* mutation was linked by P1 transduction to *serA*, which is located at 63 min on the *Escherichia coli* genetic map (details to be presented elsewhere). Previously *bglA*, which encodes phospho-β-glucosidase A, was mapped to 84 min (1, 3). The gene product of *serA* is phosphoglycerate dehydrogenase and is required for serine synthesis (4). The Kohara phages (2) were used in complementation-recombination tests to determine the physical locations of *bglA* and *serA* (Table 1).

TABLE 1. Physical mapping of *bglA* and *serA*

Mutation	Physical map location[a]	Phages[b]		Strain[c]
		Tested	Positive[d]	
bglA7	3050–3060	465–474	468, 469	JL630
serA27	3068–3073	465–474	470, 471	JL628

[a] The physical map locations are in kilobase pairs and were derived by using reference 2.

[b] The Miniset names of the Kohara phages have been used; they are listed in reference 2a.

[c] Strains JL628 (*bglR67 bglB*::λ*lacZ bglA*::Tn*10 serA27*) and JL630 (*bglR67 bglB*::λ*lacZ bglA7*) are derived from laboratory strain RV (F⁻ Δ*lacX74 thi*).

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* Corresponding author.