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Stable albicidin resistance in *Escherichia coli* involves an altered outer-membrane nucleoside uptake system

ROBERT G. BIRCH,[1]* JOHN M. PEMBERTON[2] AND W. V. SHIROMI BASNAYAKE[1]

[1] *Department of Botany and* [2] *Department of Microbiology, University of Queensland, St Lucia 4067, Australia*

Albicidin blocked DNA synthesis in intact cells of a PolA⁻ EndA⁻ *Escherichia coli* strain, and in permeabilized cells supplied with all necessary precursor nucleotides, indicating a direct effect on prokaryote DNA replication. Replication of phages T4 and T7 was also blocked by albicidin in albicidin-sensitive (Alb^s) but not in albicidin-resistant (Alb^r) *E. coli* host-cells. All stable spontaneous Alb^r mutants of *E. coli* simultaneously became resistant to phage T6. The locus determining albicidin sensitivity mapped at *tsx*, the structural gene for an outer-membrane protein used as a receptor by phage T6 and involved in transport through the outer membrane of nucleosides present at submicromolar extracellular concentrations. Albicidin does not closely resemble a nucleoside in structure. However, Alb^s *E. coli* strains rapidly accumulated both nucleosides and albicidin from the surrounding medium whereas the Alb^r mutants were defective in uptake of nucleosides and albicidin at low extracellular concentrations. An insertion mutation blocking Tsx protein production also blocked albicidin uptake and conveyed albicidin resistance. Albicidin supplied at approximately 0·1 μM blocked DNA replication within seconds in intact Alb^s *E. coli* cells, but a 100-fold higher albicidin concentration was necessary for a rapid inhibition of DNA replication in permeabilized cells. We conclude that albicidin is effective at very low concentrations against *E. coli* because it is rapidly concentrated within cells by illicit transport through the *tsx*-encoded outer-membrane channel normally involved in nucleoside uptake. Albicidin resistance results from loss of the mechanism of albicidin transport through the outer membrane.

Introduction

Chlorosis-inducing isolates of *Xanthomonas albilineans*, the sugarcane leaf scald pathogen, produce a family of antibacterial compounds in culture. The major component, named albicidin, has been partially characterized as a novel, low-molecular-mass antibiotic which is rapidly bactericidal to a range of Gram-positive and Gram-negative bacteria at concentrations as low as 1 ng ml⁻¹ (Birch & Patil, 1985*b*). Inhibition of prokaryote DNA replication was identified as the primary mode of action of albicidin, based on studies of incorporation of labelled precursors (Birch & Patil, 1985*a*). A combination of ultrastructural, genetic and biochemical evidence indicates that a member of the albicidin family of toxins produced by *X. albilineans* in invaded xylem causes chlorosis in sugarcane leaf scald disease by blocking plastid DNA replication, resulting in blocked chloroplast differentiation (Birch & Patil, 1983, 1987*a, b*). Albicidins are, therefore, of interest as phytotoxins, potential clinical antibiotics, and as tools to study prokaryotic DNA replication. We are particularly

interested in bacterial genes encoding different mechanisms of albicidin resistance as candidates for transfer to plants, with the long-term aim of producing recombinant albicidin-resistant sugarcane cultivars.

We report here that stable albicidin resistance in *Escherichia coli* occurs by loss of a nucleoside transport mechanism involved in rapid, illicit intracellular accumulation of albicidin.

Methods

Bacterial and phage strains. Xanthomonas albilineans strains LS116, LS136 and LS155 were isolated from diseased sugarcane from Queensland, Australia. Strains of *E. coli* used are described in Table 1. Bacteriophages T4, T6 and T7 were obtained from Dr L. Sly, UQM culture collection (Dept of Microbiology, Univ. of Queensland).

Albicidin production. Albicidins produced in culture by *X. albilineans* were purified as described previously (Birch & Patil, 1985*a*), except that Merck Fractogel TSK HW-40(S) replaced Sephadex LH-20 for gel-filtration in methanol. This change was necessary because all recently tested batches of Sephadex LH-20 irreversibly adsorbed albicidin. The manufacturers of Sephadex (Pharmacia) indicated that the highly aromatic structure of albicidin may result in different degrees of adsorption to different batches of LH-20.

Table 1. *E. coli* strains used

All strains are *E. coli* K12, except HB101 and RRI which are *E. coli* K12 × *E. coli* B hybrids and UQM70 which is a local isolate used for albicidin assays.

Strain designation	Genotype	Reaction to: Albicidin*	T6	Source or reference†
BRE2050	F⁻metB ilv rpsL cytR9 deoR8 Δ(argF–lac) U169	S	S	Bremer et al. (1988)
CSH57A	F⁻ara leuB tonA lacY proC tsx-67 purE galK trpE his argG malA rpsL xyl mtl ilvA metA thi supE λ⁻	R	R	B. Bachmann
EMG2	F⁺, wild-type	S	S	B. Bachmann
GP4	BRE2050 Φ (tsx–lacZ) 1(Hyb)	R	R	Bremer et al. (1988)
HB101	F⁻hsdS(r⁻ m⁻) recA ara proA lacY galK rpoL xyl mtl supE λ⁻	S	S	Maniatis et al. (1982)
HB101 Albʳ		R	R	This study
KL251	F⁻recA ara leuB azi tonA lacZ proC tsx-67 purE trpE rpsL xyl mtl metE thi supE λ⁻	S	R	B. Bachmann
KL251 Albʳ		R	R	This study
KMBL1789	F⁻ argA pheA bio thyA deoB endA polA λ⁻	S	S	B. Bachmann
LE392	F⁻hsdR (r⁻ m⁺) lacY galK galT metB trpR supE supF λ⁻	S	S	Maniatis et al. (1982)
LE392 Albʳ		R	R	This study
P1694	F⁻ thr ara leu proA lacY galK rpsL xyl mtl argE thi supE λ⁻	S	S	⎫
P1744	tsx-206 ⎤	S	R	
P1807	tsx-211	Rᶜ	R	
P1808	tsx-212 ⎬ Mutants of strain P1694	Rᶜ	R	⎬ Mannings & Reeves (1978)
P1809	tsx-213	Rᶜ	R	
P1814	tsx-218	Rᶜ	R	
P1816	tsx-220 ⎦	Rᶜ	R	⎭
Q358	hsdR(r⁻ m⁺) sup E φ80ᴿ	S	S	Maniatis et al. (1982)
Q358 Rifʳ		S	S	This study
Q358 Albʳ		R	R	This study
RRI	recA⁺ derivative of HB101	S	S	Maniatis et al. (1982)
RRI Albʳ		R	R	This study
UQM 70	Prototrophic, Valᴷ wild-type	S	S	⎫
W3747	F′ (episome F13; argF lac tsx-69 purE) relA spoT metB Δ corresponding to F13	R	R	⎪ L. Sly, Dept of Microbiology, Univ. of Queensland
χ148	F⁻ ara leuB azi tonA lacY tsx-67 purE galK trpE rpsL xyl mtl thi supE λ⁻	R	R	⎬
χ342	Hfr proC relA spoT metB λ⁻	S	S	
χ478	lysA recA⁺ parent of KL251	R	R	⎭
294 RecA⁻	F⁻(?) hsdR(r⁻ m⁺) recA endA thi supE λ⁻	S	S	Uhlin et al. (1983)
294 RecA⁻Albʳ		R	R	This study

* Rᶜ denotes strains showing an intermediate level of Albʳ; see Fig. 1(b).

† Strains denoted B. Bachmann were kindly provided by Dr Barbara Bachmann, *E. coli* Genetic Stock Center, Yale University, CT, USA.

Unless stated otherwise, the mixture of albicidins obtained after HW-40 (S) chromatography was used in experiments reported here. To ensure that antibiotics produced by Australian isolates of *X. albilineans* were typical of albicidins described from Hawaiian isolates (Birch & Patil 1985a, b), we confirmed that albicidin-sensitive strains of *E. coli* were inhibited around colonies of *X. albilineans*, whereas mutants selected for resistance to pure albicidin were not inhibited, and vice versa.

Selection of albicidin-resistant mutants. Albicidin-resistant (Albʳ) mutants of *E. coli* strains were selected using a gradient-plate technique (Gerhardt, 1981), or by plating untreated cell suspensions onto plates of Z agar (1 g glucose, 10 g tryptone, 5 g yeast extract, 10 g NaCl, 0·4 g CaCl₂.2H₂O, 15 g agar per litre of water) containing 100 ng albicidin ml⁻¹.

Scoring albicidin and phage resistance. In mapping and cloning experiments, colonies were tested for resistance or sensitivity to albicidin by patching them, using sterile toothpicks, onto Z agar plates containing 100 ng albicidin ml⁻¹. Resistance to phage T6 was tested by

patching colonies onto Z agar plates spread with 0·1 ml of a T6 lysate (about 10¹⁰ p.f.u. ml⁻¹); the results were confirmed by cross-streaking bacteria against phage on Z agar plates (Gerhardt, 1981). Dose-response relationships to albicidin were determined as previously described (Birch & Patil, 1985a), except that basal layers were of Z agar rather than minimal medium.

Effect of albicidin on phage replication. Multiplication of phages T4 and T7 was tested in *E. coli* strains RRI, RRI Albʳ, Q358 and Q358 Albʳ, both in the presence and absence of albicidin. Bacterial cells from an overnight culture in Z broth were resuspended at 10⁶ cells ml⁻¹ in MC buffer (Miller, 1972) with 10⁴ p.f.u. ml⁻¹ of phage and allowed to adsorb for 5 min at 24 °C. Suspensions were then diluted 10-fold into Z broth or Z broth containing 100 ng albicidin ml⁻¹ and incubated at 37 °C with gentle shaking. Samples were withdrawn at intervals and plated in soft agar seeded with sensitive bacterial cells to detect any increase in phage titre, which would indicate replication and subsequent release after host-cell lysis. Inhibition of phage replication was indicated by absence of a burst of phage release within 3 h incubation in these one-step growth experiments.

Mapping and cloning of alb locus. The approximate location of the gene controlling albicidin sensitivity (*alb*) was determined by estimating the frequency of cotransfer of albicidin sensitivity with selected amino acid markers mobilized from *E. coli* Q358(pULB113) Rif[r] to the polyauxotrophic Alb[r] strain CSH57A, as described by Van Gijsegem & Toussaint (1982). Fine mapping was undertaken using the generalized transducing phage P1 *cm ts*, essentially as described by Miller (1972).

Cloning into plasmid pBR322 and into cosmid pHC79 used published protocols (Pühler & Timmis, 1984) and biochemicals from Boehringer-Mannheim.

Thymidine uptake. The rate of transport of [³H]thymidine into cells (Hankte, 1976) was compared for several *E. coli* strains and related Alb[r] mutants.

The effect of albicidin on thymidine uptake was determined using Alb[s] *E. coli* strains 294 RecA⁻ (Thy⁺) and KMBL1789 (Thy⁻), by the same procedure in the presence and absence of albicidin at 100 ng ml⁻¹ and 1 μg ml⁻¹. An additional control treatment in these experiments included 1 mM-adenosine, which is known to reduce thymidine uptake by competing for transport across the cytoplasmic membrane (Krieger-Brauer & Braun, 1980).

Albicidin uptake. *E. coli* strains were grown in Luria broth (Miller, 1972) with shaking at 37 °C for approximately 4 h to an OD_{520} of 2·0. Cells were harvested by centrifugation at 8000 *g* for 5 min, washed in minimal medium (6 g Na_2HPO_4, 3 g KH_2PO_4, 1 g NH_4Cl, 2·5 g $MgSO_4.7H_2O$, 2 g glucose per litre of water), then resuspended in 0·01 times the original culture volume of minimal medium containing 100 ng albicidin ml⁻¹. At intervals after addition of albicidin, 0·2 ml samples were removed and immediately centrifuged at 15 000 *g* for 1 min. Supernatants were removed from the cell pellets, and the cells were washed once by resuspending in 0·2 ml minimal medium and centrifuging at 15000 *g* for 1 min. Albicidin activity remaining in the supernatant and in the wash was determined for each sample as described previously (Birch & Patil, 1985*a*) except that *E. coli* UQM70 was used as the assay strain. Centrifugation rather than filtration must be used to separate cells from the surrounding medium because at these low albicidin concentrations substantial albicidin activity is lost by adsorption to glassfibre filters. A wash-step proved necessary to recover albicidins weakly adsorbed to the bacterial cell-surface. With Alb[r] *E. coli* cells this wash recovers substantial albicidin activity which may otherwise wrongly be assumed to have been taken up by cells.

DNA replication in permeabilized cells. The concentration of albicidin required to block replicative DNA synthesis was determined using intact cells of the PolA⁻ EndA⁻ *E. coli* strain KMBL1789, as described previously (Birch & Patil, 1985*a*), and was compared with the concentration required to inhibit DNA synthesis in toluene-permeabilized cells of the same strain prepared and assayed as described by Moses & Richardson (1970).

Results and Discussion

Albicidin-resistant mutants

Two classes of spontaneous albicidin-resistant mutants were obtained from gradient plates: (*a*) apparently unstable or inducible mutants in which a high proportion of the population became phenotypically albicidin-sensitive after a single subculture on non-selective medium; (*b*) stable mutants which remained resistant after many transfers through non-selective medium.

Only the stable mutant class was obtained from Z agar plates containing 100 ng albicidin ml⁻¹. These mutants showed an approximate 100-fold increase in albicidin resistance relative to their parent strains (Fig. 1). All subsequent experiments reported here were confined to stable mutants.

Albicidin-resistant *E. coli* strains carrying lysogenic phage λ showed many turbid plaques when plated on media containing albicidin, suggesting phage induction by albicidin treatment. All agents which are known to induce the lytic cycle of phage λ affect DNA replication (Roberts & Devoret, 1983). Results discussed below indicate that albicidin-resistant *E. coli* strains fail to accumulate a lethal intracellular antibiotic concentration. However, sufficient albicidin apparently penetrates such cells to induce prophage λ.

Approximate location and attempted cloning of the alb locus

After patch-mating *E. coli* Q358(pULB113) Rif[r] with *E. coli* CSH57A, transconjugants prototrophic for one of seven auxotrophic mutations of CSH57A were selected. One-hundred transconjugants of each type were patched to test for cotransfer of albicidin sensitivity. Cotransfer frequencies were 17% with *leu*, 91% with *proC*, 77% with *purE* and 0% with *ile*, *trp*, *his* and *met*. These data indicate that *alb* is located at 9 to 10 min on the *E. coli* chromosome.

To confirm this approximate location, *E. coli* W3747 (an Alb[r] strain carrying episome F13 and a corresponding chromosomal deletion from 6 min to 12 min) was patch-mated with *E. coli* strains KL251 (an Alb[s]ProC⁻ RecA⁻ strain) and RRI (an Alb[s]ProA⁻RecA⁺ strain), selecting for transconjugants not requiring proline. All Pro⁺ transconjugants of KL251 remained Alb[s] whereas 38 of 40 Pro⁺ transconjugants of RRI were Alb[r]. This suggested that the Alb[r] gene on F13 is recessive and confers albicidin resistance only in the RecA⁺ strain following recombination and segregation.

Genomic libraries of *E. coli* strains Q358 and Q358 Alb[r] were prepared by cloning fragments from partial *Hind*III and *Sau*3AI endonuclease digests of chromosomal DNA into plasmid pBR322 and cosmid pHC79, followed by transformation or transfection into suitable recipients. Although the Alb[s] gene was expected to be dominant, no Alb[s] clones were detected from several libraries screened. It was also expected that cloning into recombination-proficient recipients would allow transformants which received the recessive Alb[r] locus to be detected after recombination and segregation in some of the progeny, as described above for F13. Several Alb[r] transformants were obtained in this manner, but when the plasmids in these transformants
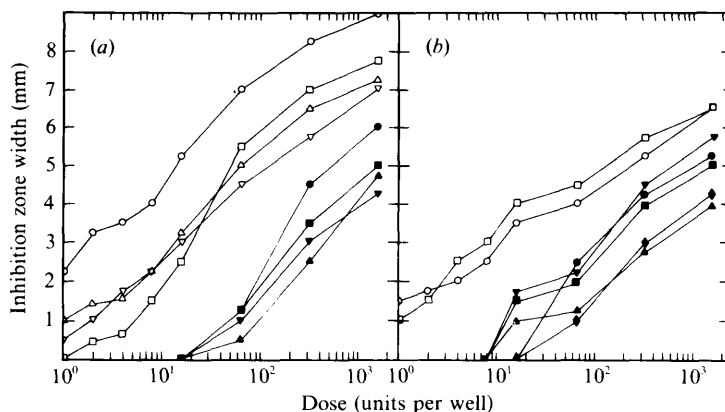
Fig. 1. Dose-response relationships for inhibition of growth of *E. coli* strains by albicidin. (*a*) Parent strains 294 (O), KL251 (□), Q358 (△) and LE392 (▽), and spontaneous mutants 294 Alb^r (●), KL251 Alb^r (■), Q358 Alb^r (▲) and LE392 Alb^r (▼). (*b*) Parent strain P1694 (O), and *tsx* mutants P1744 (□), P1807 (▲), P1808 (▼), P1809 (●), P1814 (■) and P1816 (◆). Plate-assay conditions are described in Methods. The results shown are from a typical experiment.

Table 2. *Linkage relationship among proC, lac, alb and tsx*

| Transduction | | Selected marker | No. tested | Frequency of unselected donor marker (%) | | |
P1 donor	Recipient			*lac*	*alb*	*tsx*
EMG2	χ478	*proC*$^+$	697	18	59	52
χ148	χ342	*proC*$^+$	600	12	59	59
Q358 Alb^r	χ342	*proC*$^+$	491	–	45	44

were subsequently re-isolated and transformed into *E. coli* Q358, the Alb^r gene was not transmitted with the plasmids. One possible explanation was instability of the high-copy-number plasmids carrying *alb* or adjacent genes.

It is known that albicidin selectively inhibits DNA synthesis without evidence of damage or binding to DNA, suggesting specific interaction with DNA polymerase or other DNA replication proteins (Birch & Patil, 1985*a*). Of the known *E. coli* genes for DNA replication proteins, only *dnaZ* and *dnaX* (coding for subunits of DNA polymerase III) map within 2 min of *alb*.

Phages T4 and T7 code for their own replicative polymerases, and do not require host DNA polymerase III (Nossal, 1983). We therefore tested sensitivity of these phages to albicidin. Replication of phages T4 and T7 was blocked by albicidin in Alb^s *E. coli* RRI and Q358, but not in Alb^r mutants of these strains. Accordingly, the Alb^r mutations could affect albicidin uptake, albicidin inactivation or an *E. coli* gene involved in chromosome and in T4 and T7 replication. No such gene is known to map near the *alb* locus.

Precise mapping of the alb locus at tsx

Results of P1 transduction experiments to map the *alb* locus more accurately are shown in Table 2. All Alb^r transductants in these experiments were observed to be T6^r. In contrast, a variable proportion of T6^r transductants remained Alb^s. However, a difficulty encountered in this work was the relatively high frequency at which spontaneous T6^r mutants arose; this may have contributed significantly to the apparent T6^r Alb^s transductants. Because of the very close linkage between *alb* and *tsx* in transduction experiments, we selected more than 430 spontaneous Alb^r mutants of strains 294 RecA$^-$, Q358, HB101, LE392 and RRI and tested their sensitivity to phage T6. All Alb^r mutants were found to have simultaneously become T6^r, showing only a few plaques on cross-streaked plates, while the parent strains were completely lysed by T6.

However, in a reciprocal experiment not all T6^r *E. coli* strains were Alb^r. We selected non-mucoid T6^r (i.e. *tsx*) mutants of Q358, HB101 and χ342, and found that a

variable proportion remained Alb^s. The proportion was dependent on the albicidin concentration in the test-plates; many of the spontaneous *tsx* mutants selected appeared to gain resistance to low albicidin concentrations.

To examine this phenomenon further, we determined the dose-response relationships to albicidin for a series of *E. coli tsx* mutants known to have alterations in the structure or amount of the outer-membrane receptor protein encoded by the *tsx* allele, as indicated by their ability to plate extended host-range mutants of phage T6 (Manning & Reeves, 1978). Strain P1744 has a *tsx* protein very similar to the parent strain P1694 (Manning & Reeves, 1978) and showed similar albicidin sensitivity (Fig. 1*b*), whereas the other mutants with more substantially altered *tsx* proteins showed decreased sensitivity to albicidin.

Relationship between albicidin resistance and nucleoside uptake

The *tsx* gene product is an outer-membrane protein first recognized as the receptor for phage T6 and colicin K, and now known to function in the transport of all nucleosides and deoxynucleosides, except cytidine and deoxycytidine, through the *E. coli* outer membrane (Krieger-Brauer & Braun, 1980). The *tsx* protein is most effective in transport of nucleosides present at sub-micromolar concentrations in the surrounding medium, and supplies nucleosides for transport across the cyto-plasmic membrane by two uptake systems termed *nupC* and *nupG*. These processes result in a 300- to 600-fold intracellular concentration of nucleosides within 1 min of addition to the surrounding medium (Munch-Petersen *et al.*, 1979).

Albicidin has a molecular mass of 842 Da, three to four times that of common nucleosides (Birch & Patil, 1985*b*). However, albicidin supplied at approximately 0·1 μM acts within 1 min to specifically block DNA replication. This is a very low concentration for effective antibiotic activity. One hypothesis we proposed was that albicidin was actively accumulated as a nucleoside analogue via the *tsx* protein outer-membrane channel, leading to inhibition of DNA replication at a higher intracellular concentration. Albicidin resistance in *E. coli* would, therefore, result from mutations at *tsx* which reduced albicidin uptake. This would account for the recessive Alb^r gene located at *tsx*, and the difficulty encountered in cloning the *alb* (= *tsx*) gene. Other genes for outer-membrane proteins are harmful in multiple copies (Stoker *et al.*, 1982), and cloning of the intact *tsx* gene has not been reported. Also consistent with this hypothesis was the observation that spontaneous Alb^r mutants of
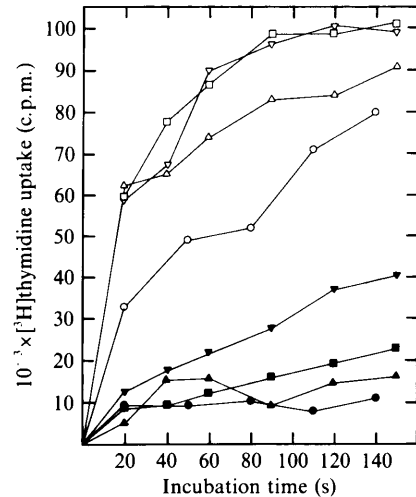

Fig. 2. Uptake of [³H]thymidine into *E. coli* parent strains 294 (O), RRI (□), Q358 (△), LE392 (▽), and into spontaneous mutants 294 Alb^r (●), RRI Alb^r (■), Q358 Alb^r (▲) and LE392 Alb^r (▼). Exponential phase cultures in minimal medium at OD$_{520}$ = 0·26 were supplied with 0·1 μM-[³H]thymidine [77 Ci mmol^{-1} (2·85 TBq ml^{-1})] at time-zero. Thymidine uptake was measured by determining radioactivity in washed cells as described in Methods. The results shown are from a typical experiment.

several strains tested all showed reduced nucleoside uptake (Fig. 2).

Different chemical groupings on the *tsx* protein are involved in receptor activities for T6 and colicin K (Hankte, 1976), and some changes in the conformation of the receptor can be matched by specific phage host-range mutants (Manning & Reeves, 1978; Riede *et al.*, 1985). It seems probable, therefore, that some *tsx* mutations may alter phage receptor activity without affecting albicidin-uptake activity, which would explain the existence of T6^{r}Alb^s mutants. The absence of the reciprocal Alb^{r}T6^s mutants might then indicate that albicidin transport places less constraint on conformation of the *tsx* protein channel than T6 receptor activity.

Relationship between albicidin resistance and albicidin uptake

Radioactive albicidin is not available for direct measurement of uptake by cells. Although albicidins can be extracted from washed albicidin-treated cells by 95% acetone, the dose-response relationship of the extracted activity differs from the applied activity, indicating alteration to albicidins in treated cells and precluding quantitative determination of uptake by this approach (data not shown). We therefore estimated albicidin uptake by following the removal of albicidin from the medium surrounding Alb^s and Alb^r *E. coli* cells, as described in Methods.
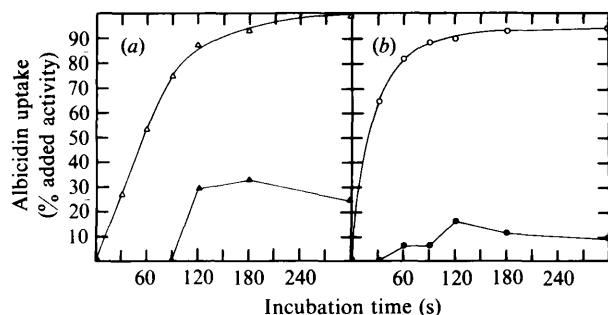
Fig. 3. Uptake of albicidin by *E. coli* strains. (*a*) Parent strain Q358 (△) and spontaneous mutant Q358 Alb[r] (▲). (*b*) Parent strain BRE2050 (○) and isogenic Φ (*tsx–lacZ*) *1* (Hyb) fusion derivative strain GP4 (●). Albicidin uptake was estimated by following the removal of albicidin from the medium surrounding cells, as described in Methods. Data points are means of two replicates. The results shown are from a typical experiment.

The albicidin-sensitive *E. coli* strain Q358 rapidly removed albicidin from the surrounding medium, whereas the spontaneous Alb[r] mutant of this strain did not (Fig. 3*a*). The difference in rates of albicidin removal closely resembled the difference in rates of nucleoside uptake by these strains (Fig. 2).

To provide a more rigorous test of our hypothesis that albicidin resistance in *E. coli* results from mutations at *tsx* which reduce albicidin uptake, we investigated an isogenic pair of *E. coli* strains differing only in one thoroughly characterized mutation which eliminates production of the Tsx protein. *E. coli* strain BRE2050 is Tsx[+]; strain GP4 is identical to BRE2050 except for an insertion mutation that generates a *tsx–lacZ* protein fusion close to the amino-terminus of Tsx, so that strain GP4 does not produce any Tsx protein (Bremer *et al.*, 1988). Strain BRE2050 is Alb[s] and rapidly removed albicidin from the surrounding medium; strain GP4 was Alb[r] and showed greatly reduced albicidin uptake (Fig. 3*b*).

Effect of albicidin on DNA replication in permeabilized cells

The hypothesis of rapid intracellular accumulation of albicidin also leads to the prediction that DNA replication in a permeabilized cell system will appear less-sensitive to albicidin than in intact cells of an albicidin-sensitive strain. This was indeed the case, as shown in Fig. 4. A 100-fold increase in albicidin concentration over that effective in intact cells was necessary to achieve rapid inhibition of DNA replication in permeabilized cells. Lower albicidin concentrations caused a slower, partial inhibition of DNA replication in permeabilized cells, presumably due to delay in contact
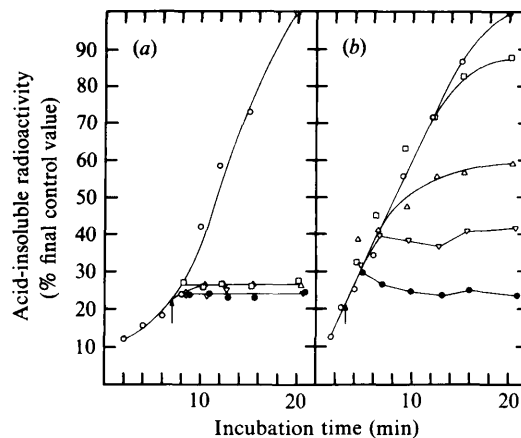

Fig. 4. Effect of albicidin on DNA synthesis in intact (*a*) and toluene-permeabilized (*b*) cells of *E. coli* KMBL1789. ○, Control; □, 100 ng albicidin ml⁻¹; △, 1 μg albicidin ml⁻¹; ▽, 10 μg albicidin ml⁻¹; ●, 3 mM-*N*-ethylmaleimide. Arrows indicate time of addition of antibiotic to cell suspensions previously supplied with the labelled precursors [³H]thymidine (*a*) and [³H]thymidine 5-triphosphate (*b*). Radioactivity incorporated into the acid-insoluble fraction was determined at intervals, as described in Methods. Final c.p.m. ml⁻¹ for control samples, plotted as 100 radioactivity units, were (a) 11.9×10^3 (*a*) and 15.2×10^3 (*b*). The results shown are from a typical experiment.

between albicidin and its molecular target at lower concentrations. These results with the PolA⁻EndA⁻ strain KMBL1789 provide direct evidence that replicative DNA synthesis is blocked by albicidin; use of the permeabilized cell system with all necessary deoxynucleoside triphosphates provided confirms that the target of albicidin is indeed the DNA replication machinery rather than precursor biosynthesis or uptake (Birch & Patil, 1985*a*).

Absence of competition between albicidin uptake and nucleoside uptake

Albicidin did not interfere with nucleoside uptake by *E. coli* (Table 3) and added nucleosides do not protect *E. coli* from albicidin (Birch & Patil, 1985*a*). This finding can be contrasted with that for the maleimide antibiotic showdomycin, which is transported across the cytoplasmic membrane by the *nupC* system but does not use the *tsx* channel. A wide variety of nucleosides protect *E. coli* from showdomycin by competitive inhibition of showdomycin uptake, and showdomycin-resistant mutants have an altered cell-membrane nucleoside transport system (Komatsu, 1981). The difference arises because nucleosides do not compete with each other for transport through the *tsx* channel, but do compete for transport across the cytoplasmic membrane (Krieger-Brauer &

Table 3. *Effect of albicidin on uptake of thymidine by E. coli*

Thymidine uptake was measured as c.p.m. ml^{-1} in washed cells 2 min after addition of [^{3}H]thymidine at 0·1 μM. Results are means of two replicates, expressed as a percentage of the water control. Control values were: 294 RecA$^-$, 9·9 × 10^5 c.p.m. ml^{-1}; KMBL1789, 8·0 × 10^5 c.p.m. ml^{-1}.

E. coli strain	H$_2$O control	Thymidine uptake in the presence of:		
		100 ng Alb ml^{-1}	1 μg Alb ml^{-1}	1 mM-Adenosine
294 RecA$^-$	100	127	138	15
KMBL1789	100	131	143	10

Braun, 1980). We have not obtained stable Alb^r mutants of *E. coli* which are unaltered at *tsx*, so the route of albicidin transport across the cytoplasmic membrane remains unknown.

Conclusion

We conclude that albicidin is effective at very low concentrations against *E. coli* because it is rapidly concentrated within cells by illicit transport mediated by the *tsx*-encoded outer-membrane channel normally involved in nucleoside uptake. Albicidin resistance in *E. coli* maps at or adjacent to *tsx*, is accompanied by resistance to phage T6 and results from loss of the mechanism of albicidin transport through the outer membrane.

Other bacteria sensitive to low concentrations of albicidin presumably also accumulate the antibiotic intracellularly as a nucleoside analogue, but mechanisms of resistance may differ to that in *E. coli*. We have, for example, recently investigated albicidin resistance in *Klebsiella oxytoca* and discovered a gene for an albicidin-binding protein which confers high-level albicidin resistance when expressed in *E. coli* (Walker *et al.*, 1988).

It may be possible to detect other mechanisms of spontaneous albicidin resistance in *E. coli* by plating a Thy$^-$ strain on media containing albicidin and a low concentration of the required nucleoside. Any resulting Alb^r mutants would be expected to have an altered nucleoside-uptake system capable of excluding albicidin while transporting thymidine, or to have a different mechanism such as an albicidin-resistant target. We are interested in such mechanisms as they may prove applicable to obtain albicidin-resistant plant cells by selection or genetic transformation.

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