The Effect of Triazole on Cysteine Biosynthesis in *Salmonella typhimurium*

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**SUMMARY**

The inhibition by triazole of the growth of wild-type *Salmonella typhimurium* is reversed by serine, methionine or cysteine. Two auxotrophs which responded to any one of these three compounds were hypersensitive to triazole and were shown by genetical analysis and enzyme assays to have a *cysE* (serine transacetylase) deficiency. Triazole hypersensitivity of the mutants was reversed by sulphite and sulphide and triazole prevented induction of sulphate permease and activating enzymes by O-acetyl serine (OAS) or serine in the wild-type. It is probable that the inhibitory effect of triazole is due to this interference with the induction of cysteine biosynthetic enzymes.

**INTRODUCTION**

In *Salmonella typhimurium* the formation of cysteine from sulphide and O-acetylserine (OAS) involves the convergence of two pathways (Smith, 1971; Fig. 1). Sulphate is taken up and reduced in four steps to give sulphide and OAS is formed from serine and acetyl CoA (serine transacetylase). Cysteine is involved in methionine synthesis: it reacts with O-succinylhomoserine to give cystathionine, which is then hydrolysed to give homocysteine, the immediate precursor of methionine. Regulation of cysteine synthesis results from feedback inhibition of the sulphate transport system and serine transacetylase and non-co-ordinate repression of all the other cysteine enzymes by cysteine. There is also evidence for induction control of cysteine enzymes for which OAS and the *cysB* (regulatory) gene product are necessary (Jones-Mortimer, 1969b).

Boguslawski, Walczak & Klopotowski (1967) reported that 1,2,4-triazole inhibited the growth of *Salmonella typhimurium*, and that this inhibition could be overcome by adding serine, methionine or cysteine to the growth media. We have isolated mutants which grow poorly unless provided with serine, methionine or cysteine. We have shown that these mutants have lowered serine transacetylase activity due to a mutation in the *cysE* gene. Triazole is shown to interfere with the induction process possibly due to a reduction in the intracellular level of the inducer OAS.

**METHODS**

*Organisms.* All bacteria used were derivatives of *Salmonella typhimurium* strain LT2 (Table 1). Map locations of their markers are included in Fig. 2. The transducing phage was the L4 mutant of phage P22 (Smith, 1968).
Fig. 1. Outline of the synthesis of cysteine and its relation to methionine synthesis in *Salmonella typhimurium*. APS, adenosine-5'-sulphatophosphate; PAPS, adenosine-3'-phosphate 5'-sulphatophosphate; OAS, O-acetylserine. cysA, sulphate permease; cysC, ATP-adenylylsulphate 5'-phosphotransferase (APS kinase); cysD, ATP sulphate adenyl transferase (ATP sulphurylase); cysE, serine transacetylase; cysH, adenosine 3'-phosphate 5'-sulphatophosphate reductase (PAPS reductase); cysGIJ, hydrogen sulphide NADP oxidoreductase (sulphite reductase). ------- Feedback inhibition: ······· induction; x - x - x - repression (after Smith, 1971).

Table 1. *Strains of Salmonella typhimurium*

The genotype designations are in accordance with Sanderson (1970): *arg*, arginine; *cys*, cysteine; *his*, histidine; *ilv*, isoleucine-valine; *lac*, lactose; *met*, methionine; *pur*, purine; *pyr*, pyrimidine; and *trp*, tryptophan.

<table>
<thead>
<tr>
<th>Strain designation</th>
<th>Genotype</th>
<th>Source</th>
</tr>
</thead>
<tbody>
<tr>
<td>wild-type LT2</td>
<td></td>
<td>P. E. Hartman</td>
</tr>
<tr>
<td>argE116</td>
<td></td>
<td>K. E. Sanderson</td>
</tr>
<tr>
<td>cysE396</td>
<td></td>
<td>K. E. Sanderson</td>
</tr>
<tr>
<td>pyrE125</td>
<td></td>
<td>K. E. Sanderson</td>
</tr>
<tr>
<td>SMC-1</td>
<td>cysE1354 hisCBHAFIE3501 HfrK5</td>
<td>K. Krajewska-Grynkiewicz</td>
</tr>
<tr>
<td></td>
<td>metK731(P22)</td>
<td>(DES* mutagenesis of metK731 (P22))</td>
</tr>
<tr>
<td>SMC-2</td>
<td>cysE1354 hisCBHAFIE3501 HfrK5</td>
<td>K. Krajewska-Grynkiewicz</td>
</tr>
<tr>
<td></td>
<td>metK731(P22)</td>
<td>(DES* mutagenesis of metK731 (P22))</td>
</tr>
<tr>
<td>TR117</td>
<td>trpC107 F'lac+</td>
<td>J. R. Roth</td>
</tr>
<tr>
<td>TW293</td>
<td>purE145 hisCBHAFIE3501 HfrK5</td>
<td>W. Walczak</td>
</tr>
<tr>
<td></td>
<td>hisCBHAFIE3501 HfrK5</td>
<td>Transduction of TW293 with wild type.</td>
</tr>
<tr>
<td>TW338</td>
<td>cysE1353 ilv-666 metK731(P22)</td>
<td>NG* mutagenesis of SMC-1</td>
</tr>
<tr>
<td>TW339</td>
<td>cysE1353 ilv-666 metK731(P22) F'lac+</td>
<td>Episome transfer from TR117 to TW338</td>
</tr>
<tr>
<td>TW340</td>
<td>cysE1353</td>
<td>—</td>
</tr>
</tbody>
</table>

* DES, diethyl sulphate, NG, N-methyl-N'-nitro-N-nitrosoguanidine.

**Culture media.** Sulphur-free minimal medium (BS) was the medium C of Vogel & Bonner (1956) in which magnesium sulphate was replaced by magnesium chloride. Ammonium sulphate or L-cysteine were added to final concentrations of 0.1 mM, and glucose to 0.2%. Medium L comprised medium C (pH 7) without citric acid, containing lactose at a final concentration of 0.5%. For the growth of auxotrophs, adenine (0.2 mM) with thiamin (0.01 mM) and amino acids (0.1 mM) except for L-serine (1 mM) were added. Solid media were made by the addition of 2% agar.

**Culture conditions.** Liquid cultures were grown at 37 °C with aeration and their growth followed turbidimetrically at 420 nm in a Unicam SP500 spectrophotometer. Extinction measurements were converted to μg dry wt.

**Phage lysates.** Log phase bacteria in Biomed (Poland) broth medium were infected with
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Fig. 2. Part of the linkage map of Salmonella typhimurium. Map positions, 1 to 138 min, are indicated at 20-min intervals. {, Transduction fragment; [ ], transduction fragment, orientation unknown; ( ), precise location unknown. Gene designation as in Table I with the following additions:
ar, aromatic; rfa, rough; ser, serine; str, streptomycin; trz, triazole resistance.

Phage L4 at a multiplicity of 0.1, and incubation continued for 8 to 18 h until lysis. Bacterial debris was removed by centrifuging at 6600 g, and phage sedimented by centrifuging for 50 min at 32000 g. Phage pellets were resuspended in C medium after soaking for 2 h; chloroform (one twentieth final volume) was then added to the suspensions which were kept at 4 °C.

Transduction. Quantities (0.1 ml) of overnight bacterial cultures in broth infected with phage at a multiplicity of at least 5 were spread on selective solid medium and incubated for 48 h at 37 °C.

Mutagenesis. Tenfold dilutions of overnight broth cultures of bacteria in fresh broth + N-methyl-N’-nitro-N-nitrosoguanidine (NG) at 25 μg/ml were incubated for 30 min at 37 °C and appropriate dilutions spread on nutrient agar to give single colonies. After incubation auxotrophy was tested by replica plating.

Assay of sulphate permease. The method was similar to that of Dreyfuss & Pardee (1966) except that cell separation was carried out on Sephadex columns rather than by membrane filtration. Log phase bacteria were washed twice and resuspended in BS medium. The
incubation mixture contained in a final volume of 0.5 ml, 25 nmol sodium sulphate, 15 μg chloramphenicol, 1 mg glucose, 5 mg dry wt of bacteria and 1.25 μCi of carrier-free [35S]sodium sulphate. After 5 min shaking at 21 °C this mixture was passed quickly through Sephadex G50, coarse mesh; the bed volume was 6 ml. Bacteria usually passed through the column in less than 30 s. Unabsorbed [35S]sulphate never appeared earlier than in the sixth drop. The first three drops of turbid bacterial suspension were collected, diluted and their extinction measured in the spectrophotometer at 420 nm before conversion to μg dry weight of cells. Radioactivity was measured in a Nuclear Chicago Biospan gas flow windowless counter.

**Assay of sulphate-activating enzymes.** This was based on the method of Pasternak (1962). Log phase bacteria were washed and resuspended in 100 mM-tris-HCl buffer (pH 8.0) containing 5 mM-EDTA and disrupted by sonication. After centrifuging at 10000 g for 1 h the extract was passed through a Sephadex G50 column equilibrated with tris buffer and further centrifuged at 105 000 g. The incubation mixture contained in a volume of 0.5 ml: sodium ATP, 3 μmol; magnesium chloride, 3.7 μmol; potassium sulphate, 0.3 μmol; carrier-free [35S]sodium sulphate, 10 μCi; tris-HCl buffer, pH 8.0, 25 μmol; and 0.12 to 1.25 mg extract protein. Reactions were stopped after 30 min of incubation at 21 °C by boiling for 2 min. For controls either ATP or the bacterial extract was omitted. Following centrifuging at 6600 g the supernatants were mixed with 90 mg of activated charcoal in 2 ml of 100 mM-sodium acetate buffer (pH 5.8) containing 30 mM-sodium sulphate. (It was found that under these conditions 10 μmol of ATP as a model adenosine compound could be adsorbed by the charcoal.) After 10 min the suspensions were passed through membrane filters (Biomed Coli 5, average pore size 0.5 μm), the deposited charcoal washed extensively with acetate buffer and the radioactivity on the filters counted with a thin mica window Geiger–Muller tube. The c.p.m. obtained were from an infinitely thick sample layer. A linear relationship between the radioactivity and quantity of the samples was obtained over the range 0.12 to 1.2 mg protein.

In an alternative method the enzymes were assayed in bacteria which had been washed with tris buffer and shaken with 10 μl toluene/ml for 15 min at 37 °C.

**Assay of serine transacetylase.** The method of Kredich & Tomkins (1966) was used.

**Protein determination.** The protein in cell-free extracts was determined by the method of Lowry, Rosebrough, Farr & Randall (1951).

**Chemicals.** O-acetyl-L-serine (OAS) was synthesized by the method of Wiebers & Garner (1967). All other chemicals used were commercial products; disodium ATP (Sigma Chemical Co. St Louis, Missouri, U.S.A.), acetyl-Co-enzyme A (Mann Research Laboratories, New York, U.S.A.), 1-amino acids and 1,2,4-triazole (Calbiochem, Los Angeles, California, U.S.A.), 3-amino-1,2,4-triazole (Fluke, Buchs, Poland), carrier-free [35S]sodium sulphate (Instytut Badan Jadrowych, Swierk, Poland), and chloramphenicol (Polfa, Warsaw, Poland).

**RESULTS**

**Origin and properties of serine, methionine or cysteine (SMC) mutants.**

Two mutants requiring either serine, methionine or cysteine for growth, SMC-1 and SMC-2 (Table 1) have been studied. SMC-1 grows slowly whereas SMC-2 does not grow at all on unsupplemented minimal agar medium. In each case the growth on this medium is stimulated by either serine, methionine or cysteine. The growth characteristics of SMC-1 and SMC-2 in supplemented and unsupplemented BS medium appear in Fig. 3 and 4. SMC-1 can grow on sulphate-supplemented medium but only after a lag lasting 2 to 3 h but the
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Addition of serine to the medium completely eliminates this lag, which is therefore perhaps due to inability to convert exogenous sulphate into endogenous sulphite because of slow accumulation of OAS necessary for induction of the cysteine enzymes. In contrast *SMC-2* cannot utilize sulphate without serine; residual growth in the presence of sulphate was not very different from that in unsupplemented medium. This residual growth probably results from utilization by the bacteria of their internal pool of cysteine. Further growth following the addition of serine suggested that BS medium may contain traces of a sulphur compound or that endogenous pools of these compounds were present. The latter assumption seems more plausible as residual growth even occurred in BS medium which had already been used for the growth of bacteria.

*SMC-1* and *SMC-2* also differed from each other in their growth responses to sulphite and sulphide; *SMC-1* grew on sulphite or sulphide, but *SMC-2* also required serine to give similar growth responses. This suggested that *SMC-2* was deficient in cysteine biosynthesis and, in fact, the specific activity of sulphite reductase in cell-free extracts of *SMC-2* was less than 1% of that in wild-type extracts.
The presence of cysteine or methionine in BS medium permitted the same rate of growth of both strains but total growth with cysteine was greater in both cases. It may be that growth on methionine as a source of sulphur only results from the sparing of cysteine. The addition of sulphate to BS + methionine medium did not stimulate further growth of either mutant. Stimulation of growth of the SMC mutants by cysteine and its precursors suggests that these strains are deficient in the biosynthesis of this amino acid.

**Hypersensitivity of SMC-1 to triazole**

The reversal of triazole inhibition by serine, methionine or cysteine (Boguslawski et al. 1967) suggests that the inhibitor and the SMC defect both affect cysteine biosynthesis and that if so the cysteine deficiency of SMC mutants might result in hypersensitivity to triazole. Thus the triazole sensitivity of SMC-1 (Fig. 5a) was compared with that of wild-type (Fig. 5b). It can be seen that 0.2 mM-triazole caused inhibition of the growth of SMC-1 for 9 h but did not noticeably affect the growth of wild-type. However, this increased triazole sensitivity of SMC-1 does not necessarily mean that triazole affected the same cytoplasmic product which is defective in the mutant. It is possible that the SMC mutations and triazole affect two different factors involved in the same process.
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Table 2. Serine transacetylase activities

<table>
<thead>
<tr>
<th>Strain</th>
<th>Specific activity (μmol/mg protein/min)</th>
<th>Relative specific activity</th>
</tr>
</thead>
<tbody>
<tr>
<td>Wild-type</td>
<td>13.3</td>
<td>100</td>
</tr>
<tr>
<td>SMC-1 (cysE353)</td>
<td>2.4</td>
<td>19</td>
</tr>
<tr>
<td>SMC-2 (cysE354)</td>
<td>0.1</td>
<td>6</td>
</tr>
<tr>
<td>cysE356</td>
<td>0.0</td>
<td>1</td>
</tr>
</tbody>
</table>

Genetic map location of SMC mutations

We have mapped the SMC mutations to see whether they coincide with any of the serine, cysteine and methionine structural genes already located on the Salmonella typhimurium linkage map (Fig. 2; Sanderson 1970). The mutation in SMC-1 was identified by its hypersensitivity to triazole. As SMC-1 was lysogenic for P22 conjugation was used for genetical analysis. Strain tw338 derived from SMC-1 carried an NG-induced ilv mutation (Table 1). It was infected with the F' lac+ episome (strain tw339). The episome can integrate into any region of the S. typhimurium chromosome and initiate transfer to F- strains (Anton, 1968). Strain tw339 was then crossed on serine-supplemented plates with a range of auxotrophic recipients three of which gave prototrophic recombinants with the SMC phenotype when tested by replica to plates containing 2 mM-triazole; all cysE356+, 80% pyrE125+ and 20% argE116+ recombinants were triazole hypersensitive. One of the triazole-hypersensitive pyrE+ recombinants (tw340) was isolated, purified by several single colony re-isolations and studied in some detail. It grew poorly on BS medium unless serine, methionine or cysteine was added. The similarity of growth pattern between this strain, other transductants and SMC-1 indicated that the metK mutation of SMC-1 did not influence the SMC phenotype.

Strain tw340 was then used as a donor to transduce pyrE125 and cysE596. Recombinants were selected on BS + serine medium and the triazole hypersensitivity of 200 recombinants of each cross tested. No pyrE+ recombinants had the SMC phenotype but 197 cysE+ recombinants did, which indicated 96.5% co-transduction of SMC-1 with cysE. In a similar cross between cysE356 and SMC-2 when 100 recombinants were screened, 98% co-transducibility was obtained.

These results strongly suggested that the SMC phenotype could result from mutation in the cysE gene which codes for serine transacetylase (Kredich, Becker & Tomkins, 1969).

The serine transacetylase activity of SMC mutants

Extracts of wild-type, SMC-1, SMC-2 and a representative cysE mutant were assayed (Table 2). The wild-type specific activity (13.3 nmol acetyl-CoA hydrolysed/mg protein/min) was lower than that reported for Escherichia coli b by Kredich & Tomkins (1966). However, in another experiment a Michaelis constant for serine of 0.5 mM was obtained with a crude cell-free extract of wild-type. This was close to the value of 0.56 mM obtained by Kredich & Tomkins with their purified preparation. Both SMC strains had low serine transacetylase specific activities. In SMC-1 there was an 80% and in SMC-2 a 90% reduction. The Michaelis constant for serine as determined in cell-free extracts of SMC-1 and SMC-2 was 1 mM, slightly higher than that for wild-type.
Fig. 6. Reversal of triazole inhibition of smc-1 (cysE1353) by sulphite or sulphide. BS medium+ sulphate was inoculated with washed bacterial suspension to a final density of 10 µg/ml. The concentration of triazole was 2 mM and of sulphite and sulphide 0.1 mM. T, triazole; C, control without triazole.

These results in conjunction with the genetical data permit the designation of smc-1 and smc-2, cysE1353 and 1354 respectively.

Reversal of triazole inhibition by sulphite and sulphide

The growth requirements of smc-1 (cysE1353), its hypersensitivity to triazole and its cysE gene deficiency suggested that triazole could interfere with cysteine biosynthesis. An attempt was therefore made to relieve triazole inhibition with cysteine intermediates. It is evident from Fig. 6 that triazole inhibition of smc-1 (cysE1353) can be reversed by either sulphite or sulphide. This could have been the result of interaction between these anions with the triazole ring. If this had occurred the inhibition of growth of wild-type Salmonella typhimurium by 3-amino-1,2,4-triazole which interferes with histidine and purine biosynthesis (Hilton, Kearney & Ames, 1965; Hulanicka, Klopotowski and Bagdasarion, 1969), should also have been reversed by these anions. No such reversal was obtained suggesting that the effect of the two sulphur compounds was more likely to have been directly concerned with cysteine biosynthesis. Triazole could inhibit a step in the cysteine pathway prior to sulphite formation or interfere with cysteine regulation.
Table 3. Prevention of the induction of cysteine enzymes of wild-type bacteria by triazole

Log phase bacteria were grown in BS medium + Na₂SO₄ (1 mM) and OAS (1 mM), serine (1 mM) or triazole (2 mM) as indicated.

<table>
<thead>
<tr>
<th>Growth medium supplement</th>
<th>Sulphate permease (nmol[^35S] sulphate mg dry wt/min)</th>
<th>Sulphate-activating enzymes (c.p.m.)</th>
</tr>
</thead>
<tbody>
<tr>
<td>None</td>
<td>0.42</td>
<td>785</td>
</tr>
<tr>
<td>OAS</td>
<td>0.70</td>
<td>7 708</td>
</tr>
<tr>
<td>Triazole</td>
<td>0.35</td>
<td>707</td>
</tr>
<tr>
<td>OAS + triazole</td>
<td>0.51</td>
<td>1 136</td>
</tr>
<tr>
<td>None</td>
<td>---</td>
<td>285</td>
</tr>
<tr>
<td>Serine</td>
<td>---</td>
<td>1 548</td>
</tr>
<tr>
<td>Serine + triazole</td>
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<td>484</td>
</tr>
</tbody>
</table>

Table 4. The activity of the cysteine enzymes of wild-type bacteria in the presence of triazole or selenate

Bacteria were grown in BS medium + Na₂SO₄ (1 mM). Triazole (10 mM) and Na₂SeO₄ (2 mM) were added to the reaction mixtures which contained protein at a final concentration of 0.25 mg/0.5 ml. The activities quoted are the averages of two independent assays each carried out in duplicate.

<table>
<thead>
<tr>
<th>Enzyme activities</th>
<th>Sulphate permease (nmol[^35S] sulphate dry wt/min)</th>
<th>Sulphate-activating enzymes (c.p.m.)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Addition</td>
<td></td>
<td></td>
</tr>
<tr>
<td>None</td>
<td>0.30</td>
<td>1 350</td>
</tr>
<tr>
<td>Triazole</td>
<td>0.29</td>
<td>1 150</td>
</tr>
<tr>
<td>Na₂SeO₄</td>
<td>0.06</td>
<td>150</td>
</tr>
</tbody>
</table>

Prevention of the induction of cysteine enzymes by triazole

In preliminary experiments the growth inhibitory effect of triazole could be reversed by O-acetyl-serine (OAS) which suggested that the inhibitor could interfere with induction of cysteine biosynthetic enzymes (Jones-Mortimer, 1969 b). Identical log phase cultures of wild-type were supplemented with OAS, serine, or either compound + triazole. After 2 h the bacteria were harvested and assayed for sulphate permease and the sulphate-activating enzymes (Table 3). OAS-induced sulphate permease nearly twofold but triazole added with the inducer more than halved the induction. A more striking effect was obtained with the sulphate-activating enzymes. OAS or serine was used as an inducer; OAS caused about a tenfold increase of the enzymes and triazole almost completely prevented the induction. Induction with serine was less than with OAS but again triazole almost eliminated it.

Triazole only slightly inhibited the activity of the enzymes assayed (Table 4), whereas sodium selenate, a substrate analogue known to inhibit the cysD (ATP sulphurylase) enzyme (Pasternak, 1962; Qureshi, 1971), inhibited both enzymes by about 80%. This would indicate that the triazole effect upon Salmonella typhimurium is unlikely to be due to inhibition of an enzyme and so supports the deduction that it interferes with the regulation of synthesis of cysteine enzymes.
GROWTH REQUIREMENTS FOR SERINE, METHIONINE OR CYSTEINE IN STRAINS OF *Salmonella typhimurium*

Growth requirements for serine, methionine or cysteine in strains of *Salmonella typhimurium* can result from leaky *cysE* mutations or the presence of triazole in the growth medium of wild-type organisms and it is of interest to explore the elements common to both phenomena.

The mutants *cysE*1353 and *E*1354 have lower intracellular levels of serine transacetylase (Table 2) and may thus contain less OAS. OAS is required to induce the enzymes of sulphate assimilation, sulphate permease, two sulphate activating enzymes and sulphite reductase. It is also an intermediate in cysteine biosynthesis (Jones-Mortimer, 1969a, b). These two processes could require different concentrations of OAS. Since *cysEr*353 has a higher serine transacetylase activity than *cysE*1354 and can utilize sulphite and sulphide (Fig. 3, 4) it is inferred that *cysE*1353 has enough OAS for cysteine synthesis but not enough to induce the cysteine enzymes. This limits growth of the mutant. In contrast, *cysE*1354 is unable to grow on sulphite suggesting that it has insufficient OAS for both synthesis or induction. The behaviour of both mutants suggests that induction of the sulphate assimilation enzymes can only occur when OAS transsulphurylase is saturated with OAS.

Both mutants can utilize sulphate in the presence of serine (Fig. 3; 4). This is probably due to an increased intracellular concentration of serine, the substrate for serine transacetylase. This is supported by the observation that addition of serine increased the specific activities of OAS-inducible enzymes (Table 3). The growth responses of *cysE*1353 and *E*1354 to methionine in the absence of other sulphur compounds (Fig. 3; 4) is probably partly due to a sparing effect upon cysteine utilization. As it is generally assumed (Roberts *et al.* 1963) that methionine cannot serve as the sole source of sulphur for enteric bacteria, the extent of the growth response of these mutants to methionine is puzzling.

Growth inhibition by triazole is transient (Fig. 5; 6) and reversed by sulphite or sulphide (Fig. 6). This resembles the growth response pattern of *cysE*1353 implying that triazole limits the rate of serine transacylation or inhibits the use of OAS in induction. Certainly triazole has no inhibitory effect upon the activity of the cysteine enzymes in crude cell-free preparations (Table 4). It has been found (T. Klopotowski, unpublished) that triazole hydrolyses acetyl-CoA non-enzymatically and if such a reaction reduced substantially the intracellular concentration of acetyl CoA, cysteine biosynthesis could be affected. On the other hand the observation that triazole was able to prevent induction of sulphur assimilation enzymes by OAS (Table 3) argues against the idea of the main inhibitory effect of triazole resulting from hydrolysis of acetyl-CoA or from some other effect on OAS synthesis.

The mechanism of OAS induction is unknown but since *cysB* mutants are repressed for all cysteine enzymes, a *cysB* gene product could be essential for OAS induction (Jones-Mortimer, 1969a; b). Kredich *et al.* (1969) found that OAS activated a serine transacetylase/OAS transsulphurylase complex. They considered that OAS could act as an inducer by combining with this complex. Triazole interference is not obviously related to either of these explanations, but triazole could be used for obtaining more information on how the enzymes of sulphur assimilation are induced.

Boguslawski *et al.* (1967) suggested that triazole inhibition of wild-type *Salmonella typhimurium* resulted from interference with C₁ metabolism on the basis of their observation that glycine potentiated triazole inhibition. One important difference between *cysE*1353 and triazole inhibited wild-type bacteria is that glycine has no effect on the mutant. Thus the present experiments appear to have no direct bearing on involvement with C₁ metabolism. Although it is felt that the triazole effect is concerned with cysteine biosynthesis, the
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precise cause of cysteine deficiency in triazole-inhibited *S. typhimurium* remains to be determined.

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