## Biosynthesis of Pyocyanin, a Phenazine Microbial Metabolite

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Summary Tracer experiments have shown that phenazinel-carboxylic acid and its 5-methyl quaternary salt are incorporated into pyocyanin by *Pseudomonas aeruginosa* by decarboxylative hydroxylation.

NATURAL products with structures based on the phenazine ring system now number about thirty:<sup>1</sup> they are elaborated exclusively by micro-organisms. The best known of them is pyocyanin (1),<sup>2</sup> a metabolite of *Pseudomonas aeruginosa*, other strains of which alternatively produce aeruginosins A  $(4)^3$  and B (5).<sup>4</sup> Biosynthetic studies on the origins of the phenazine nucleus implicate shikimic acid,<sup>5</sup> and at least one molecule of anthranilic acid has been shown to be involved in the production of chlororaphin, a phenazine-1-carboxamide pigment.<sup>6</sup> With regard to intermediates involved at later stages, it has been shown that iodinin (6) may be derived in high efficiency from phenazine-1,6-diol and the corresponding 5-oxide.5a On the other hand, 1-hydroxyphenazine (7) is not a precursor of pyocyanin.<sup>7</sup>

We have already shown that 5-methylphenazinium-1carboxylate (2) can be converted by aqueous ammonia into aeruginosin A (4)<sup>8</sup> [which with sulphite ion gives aeruginosin B (5)<sup>9</sup>] and by photo-oxidation into pyocyanin,<sup>10</sup> prompting the suggestion that it may represent a common intermediate in the biosynthesis of these metabolites.<sup>10</sup> We now report on the testing of this compound along with 1-carboxyphenazine (8), also a microbial metabolite, and phenazine methosulphate (3) as precursors of pyocyanin in the living system.

When 1-carboxy-6,7,8,9-tetradeuterio-5-methylphenazinium chloride was administered over the pigment producing period to growing cultures of an appropriate strain of Ps. aeruginosa, a 2.4% incorporation into pyocyanin was found. The product was most easily assayed by conversion into 1-hydroxyphenazine (7), the mass spectrum of which is simple in the high mass region where, for the undeuteriated material, two prominent peaks are found at m/e 196 (M<sup>+</sup>) and 168 (M<sup>+</sup> - CO).<sup>†</sup> The mass spectrum of the 1-hydroxyphenazine from the feeding experiment showed corresponding peaks of the  ${}^{2}H_{4}$  species but not of the <sup>2</sup>H<sub>3</sub>, indicating specific hydroxylation at C-1 with concomitant decarboxylation. A similar result was obtained with 1-carboxy-6,7,8,9-tetradeuteriophenazine (1.4%) in-1,2,3,4-Tetradeuteriophenazine methosulcorporation). phate was also incorporated into pyocyanin (0.7%) with approximately equal quantities of the  ${}^{2}H_{3}$  and  ${}^{2}H_{4}$  species being produced.

The highly specific hydroxylation of (2) and (8) and the respective levels of incorporation strongly support the biosynthetic sequence  $(8) \rightarrow (2) \rightarrow$  pyocyanin (1). On the other hand, the significantly lower level of incorporation of





phenazine methosulphate (3), which, if involved, ought to be the immediate precursor of pyocyanin, is in keeping with it being a non-specific substrate rather than a true biosynthetic precursor.

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† The hydroxyphenazine from the three feeding experiments contained 0.4-2.6% of deuteriated species. Measurements were made at both  $M^+$  and  $M^+$  – CO to avoid error from impurity in the sample. The contribution of ions at m/e 200, 199, 172, and 171 due to the undeuteriated material was allowed for; reasonable accuracy was assured as they were of low intensity relative to those due to deuteriated species.

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