The Structure of the Pyoverdin Isolated from Various *Pseudomonas syringae* Pathovars[§]

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From seven different pathovars of *Pseudomonas syringae* representing various genetic subgroups, and one strain of *Pseudomonas viridiflava* the same pyoverdin siderophore (1) was isolated, probably identical with the pyoverdin whose amino acid composition (but not their sequence) had been reported before. 1 is the first pyoverdin where two of the ligands for Fe^{3+} are β -hydroxy Asp units. Its remarkably high complexing constant for Fe^{3+} at pH 5 as compared with other pyoverdins offers a definite advantage in plant infection. The structure elucidation of 1 will be described and the taxonomical implications regarding pyoverdins with different structures ascribed previously to *P. syringae* strains will be discussed.

§ Part CV of the series "Bacterial constituents". For part CIV see Hohlneicher et al. (2001).

Abbreviations: Common amino acids, 3-letter code; aThr, allo Thr; OHAsp, β-hydroxy-Asp; AcOHOrn, N⁵-acetyl-N⁵-hydroxy-Orn; cOHOrn, cyclo-N⁵-hydroxy-Orn (3-amino-1-hydroxy-piperidone-2); TAP, N/O-trifluoroacetyl (amino acid) isopropyl ester; Chr, pyoverdin chromophore; Suc, succinic acid side chain; Suca, succinamide side chain; ESI, electrospray ionization; FAB, fast atom bombardment; CA, collision activation; u, mass units based on the 12 C scale; COSY, correlated spectroscopy; DEPT, distortionless enhancement by polarization transfer; HMBC, heteronuclear multiple bond correlation; HMQC, heteronuclear multiple quantum coherence; NOE, nuclear Overhauser effect; ROESY, rotating frame nuclear Overhauser and exchange spectroscopy; TOCSY, total correlation spectroscopy; WATERGATE, water suppression by gradient-tailored excitation; DSS, 2,2-dimethyl-2-silapentane-5-sulfonate; TMS, tetramethylsilane; pv, pathovar; ATCC, American Type Culture Collection; CFBP, Collection Française de bactéries phytopathogènes; LMG, Laboratorium voor Microbiologie van Ghent, Belgian Coordinated Collection of Microorganisms.