

The Structure of Diphenazithionin, a Novel Antioxidant from *Streptomyces griseus* ISP 5236

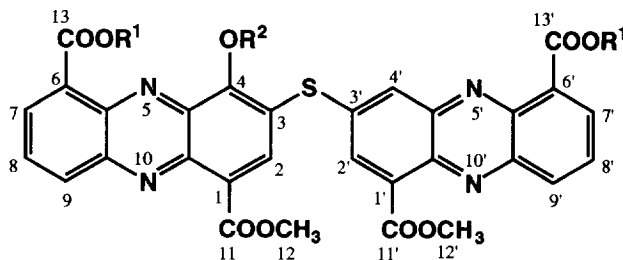
Yoshiko Hosoya, Hayamitsu Adachi, Hikaru Nakamura, Yoshio Nishimura,
 Hiroshi Naganawa, Yoshiro Okami* and Tomio Takeuchi

Institute of Microbial Chemistry, 3-14-23 Kamiosaki, Shinagawa-ku, Tokyo 141, Japan

Abstract: A novel member of the phenazine group of metabolites, diphenazithionin, has been isolated as an inhibitor of lipid peroxidation from *Streptomyces griseus* ISP 5236. It possesses a unique structure composing of two different phenazinecarboxylic acids connected by a sulfur linkage.
 Copyright © 1996 Elsevier Science Ltd

Free radicals are involved in causation of several significant human diseases including inflammation, ischemia reperfusion injury, rheumatism, etc.¹ Moreover, lipid peroxidation possibly plays an important role in inducing carcinogenesis or aging of cells.² Vitamins such as vitamin C or E are well-known agents to inhibit oxidation, but other potent and stable antioxidizing drugs can still be expected from nature. Actinomycetes have provided many kinds of potent antioxidants with low molecular weight.³ We have investigated physiological responses and inducible secondary metabolites of organisms under oxidizing conditions, and designed a new screening methodology⁴ using oxidizing agents added to the fermentation medium for actinomycetes. As a result of the screening, we isolated diphenazithionin (1), an inhibitor of lipid peroxidation which has a novel structure with the phenazine skeleton, from a culture filtrate of *Streptomyces griseus* ISP 5236. Here we report the structure elucidation by an X-ray crystallographic experiment.

Figure 1

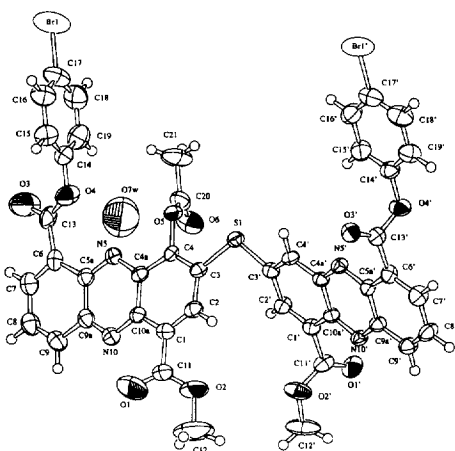


1: R¹=R²=H (Diphenazithionin)

2: R¹=*p*-BrC₆H₄, R²=COCH₃

Compound 1 [dark green powder, HR-FAB MS (M+H)⁺ *m/z* 611.0878, C₃₀H₁₉N₄O₉S (Calcd. 611.0873)] has two methyl esters and two carboxylic acid groups (δ_{H} 4.22 and 4.24, δ_{C} 168.5, 169.0, 170.2 and 170.6)⁵. Only 15 protons, including the two methyl groups, were also observed in the ¹H NMR spectrum⁵. The UV spectrum (λ_{max} : 260, 305, 375 and 442) and the characteristic low-field chemical shifts of ¹H NMR spectrum showed a typical feature of the phenazine

skeleton similar to phencomycin⁶, a carboxylated phenazine compound. These facts suggested that the major skeletal structure was composed of two phenazine components.

Figure 2. X-Ray crystal structure of 2.

and the non-hydrogen atoms were anisotropically refined, including the hydrogen atoms in the calculated position. The final cycle of full-matrix least-squares refinement was based on 4446 observed reflections ($I > 2.00 \sigma(I)$) and converged with $R = 0.077$. Crystallographic parameters have been deposited with the Cambridge Crystallographic Data Center. An ORTEP drawing of **2** is shown in Fig. 2.

As shown in Fig. 2, the present study clarified the structure of diphenazithionin as 4-hydroxy-3,3'-thio-di(1-carbomethoxyphenazine-6-carboxylic acid).

The phenazine compounds have often been isolated from streptomycetes as antibiotics⁹ and also as antioxidants, but we have reported in this paper the first discovery of diphenazithionin, a dimeric phenazinecarboxylic acid linked by a sulfur atom, showing strong antioxidant activity.

REFERENCES AND NOTES

- (a) Abe, K.; Yoshida, S.; Watson, B.; Busto, R.; Kogure, K.; Gingsberg, M. *Brain Res.* **1983**, *273*, 166-169. (b) Kuzuya, T.; Hoshida, S.; Nishida, M.; Kim, Y.; Fuji, H.; Kitabatake, A.; Kaneda, T.; Tada, M. *Cardiovasc. Res.* **1989**, *23*, 323-330. (c) Cashman, J.R. *Pharmac. Res.* **1985**, *6*, 253-261.
- (a) Diplock, A.T.; Rice-Evans, A.; Burdon, R.H. *Cancer Research* **1994**, *54*, 1952s-1956s. (b) Hockenbery, D.M. *Cell* **1993**, *75*, 241-251. (c) Stadtman, E.R. *Science* **1992**, *257*, 1220-1224.
- Shin-ya, K.; Shimazu, A.; Hayakawa, Y.; Seto, H. *J. Antibiot.* **1992**, *45*, 124-125. (b) Mo, C.; Shin-ya, K.; Furihata, K.; Furihata, K.; Shimazu, A.; Hayakawa, Y.; Seto, H. *J. Antibiot.* **1990**, *43*, 1337-1339.
- Hosoya, Y.; Fujita, Y.; Hotta, K.; Okami, Y. *J. Antibiot.* in preparation
- ¹H-NMR (TFA-d, 500 MHz) δ 9.52 (s, 2-H), 9.30 (d, $J = 7.2$ Hz, 7'-H), 8.47 (dd, $J = 7.2, 6.1$ Hz, 8'-H), 8.96 (d, $J = 7.2$ Hz, 9'-H), 8.83 (brs, $J = 0.8$ Hz, 2'-H), 8.36 (brs, $J = 0.8$ Hz, 4'-H), 9.24 (d, $J = 6.1$ Hz, 7-H), 8.79 (dd, $J = 7.2, 6.1$ Hz, 8-H), 9.21 (d, $J = 7.2$ Hz, 9-H), 4.22 (3H, s, 12-H), 4.24 (3H, s, 12'-H); ¹³C-NMR (TFA-d, 125 MHz) δ 56.0 (12'), 56.4 (12), 114.3 (3), 120.7, 121.4 (4'), 128.5 (9), 130.4, 130.8 (6), 134.6 (9a), 134.9 (4), 134.9 (8'), 135.6, 135.7, 137.6 (9'), 138.8, 139.1 (2'), 141.2 (7), 142.3 (10'a), 143.0 (8), 143.2 (6'), 145.7 (7'), 153.4 (2), 154.8, 165.2 (1), 168.5 (11'), 169.0 (11), 170.2 (13'), 170.6 (13).
- Gilpin, M.L.; Fulston, M.; Payne, D.; Cramp, R.; Hood, I. *J. Antibiot.* **1995**, *48*, 1081-1085.
- Illi, V.O. *Tetrahedron Letters* **1979**, 2431-2432.
- Castro, B.; Evin, G.; Selve, C.; Seyer, R. *Synthesis* **1977**, 413.
- (a) Osato, T.; Maeda, K.; Umezawa, H. *J. Antibiot.* **1954**, *7A*, 15. (b) Tipton, C.D.; Rinehart, K.L. *J. Am. Chem. Soc.* **1970**, *92*, 1425.
- Shin-ya, K.; Furihata, K.; Hayakawa, Y.; Seto, H. *Tetrahedron Letters* **1991**, *32*, 943-946.