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Thioredoxín Reductase Inhibitíon

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The inside cover picture shows the X-ray structure of human thioredoxin reductase dimer (TrxR; PDB: 2J3N; ribbon representation), with the catalytic site residues Sec and Cys highlighted (cyan surface). Enzyme activity assays and biochemical studies identified the Sec–Cys dyad as essential for TrxR inhibition by water-soluble Au^l-phosphine complexes. For more details, see the Full Paper by M. Laguna, P. J. Dyson, et al. on p. 96 ff.

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Inside Cover

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