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Supplementary Material

Bismuth(III) Thiophosphinates: Understanding How Small Atomic Change Influences Antibacterial Activity and Mammalian Cell Viability

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Scheme S1. Synthesis of diphenylphosphinothioic acid, HOP(=S)Ph₂/HSP(=O)Ph₂ and diphenylphosphinodithioic acid, HSP(=S)Ph₂.



Fig. S1. ¹H NMR of [Bi(SP(=O)Ph₂)₃] (**1**) in DMSO-d₆.



Fig. S2. ¹H NMR of [BiPh(SP(=O)Ph₂)₂], (**2**) in DMSO-d₆. ****ortho and *meta Bi-Ph protons from complex 3**.



Fig. S3. ¹H NMR of [BiPh₂(SP(=O)Ph₂)] (3) in DMSO-d₆.

7.75 7.75 7.73 7.73 7.53 7.51 7.47 7.46 7.45 7.45 7.45

Fig. S4. ¹H NMR of $[Bi(SP(=S)Ph_2)_3]$ (4) in DMSO-d₆.

Fig. S6. ³¹P NMR of [BiPh(SP(=O)Ph₂)₂] (2) in DMSO-d₆.

Fig. 57. ³¹P NMR of [BiPh₂(SP(=O)Ph₂)] (3) in DMSO-d₆.

Fig. S8. ³¹P NMR of $[Bi(SP(=S)Ph_2)_3 \cdot DMSO]$ (4) in DMSO-d₆.

	2	3	4·DMSO
Formula <i>M</i> r Crystal size [mm]	C ₃₀ H ₂₅ BiO ₂ P ₂ S ₂ 752.54 0.23 x 0.16 x 0.09	C ₂₄ H ₂₀ BiOPS 596.41 0.21 x 0.17 x 0.10	C ₃₈ H ₃₆ BiOP ₃ S ₇ 1034.98 0.24 x 0.16 x 0.10
Crystal system	Monoclinic	Monoclinic	Monoclinic
Space group a [Å]	<i>P2₁/c</i> 10.7908(4)	<i>P2₁/n</i> 9.6557(4)	<i>P2₁/n</i> 9.29900(10)
<i>b</i> [Å]	11.0293(4)	10.5962(4)	21.7360(2)
<i>c</i> [Å]	23.3960(6)	21.2320(9)	20.4159(2)
α [°]	90	90	90
в [°]	94.434(2)	93.243(3)	100.7700(10)
γ [°]	90	90	90
V [Å ³]	2776.14(16)	2168.85(15)	4053.84(7)
Ζ	4	4	4
T [K]	123(2)	123(2)	123(2)
ρ calcd. [gcm ⁻¹]	1.801	1.827	1.696
$\mu [{\rm mm}^{-1}]$	15.175	8.311	13.269
Independent reflections collected	5740	6685	8349
Rint	0.1301	0.0548	0.0473
R1 [$I > 2\sigma(I)$]	0.0587	0.0261	0.0281
wR2 (all data)	0.1611	0.0486	0.0733
GoF	1.015	1.009	1.095

Table S1. Crystallographic data and structure refinement for complexes 2, 3 and 4. CCDC numbers 1947001-1947003.

Fig. S9. Comparison of the powder X-ray diffraction pattern of bulk crystalline product (black, top) and the calculated pattern from the single crystal X-ray diffraction studies (red, bottom) of [BiPh(SP(=O)Ph₂)₂] (**2**).

Fig. S10. Comparison of the powder X-ray diffraction pattern of bulk crystalline product (black, top) and the calculated pattern from the single crystal X-ray diffraction studies (red, bottom) of [BiPh₂(SP(=O)Ph₂)] **(3)**.

Fig. S11. Comparison of the powder X-ray diffraction pattern of bulk crystalline product (black, top) and the calculated pattern from the single crystal X-ray diffraction studies (red, bottom) of [Bi(SP(=S)Ph₂)₃·DMSO] **(4)**.

Compound	MRSA	VRE	E. coli	P. aeruginosa
1 [Bi(SP(=O)Ph ₂) ₃]	/	/	/	/
2 [BiPh(SP(=O)Ph ₂) ₂]	9.2	9.2	2.3	1.1
3 [BiPh ₂ (SP(=O)Ph ₂)]	2.5	2.5	1.3	0.30
4 [Bi(SP(=S)Ph ₂) ₃]	0.49	0.49	/	/

Table S2: Selectivity indices of compounds 1 - 4, where SI = IC₅₀ / MIC. IC₅₀ of complexes against human fibroblast cells.