

## SUPPORTING INFORMATION

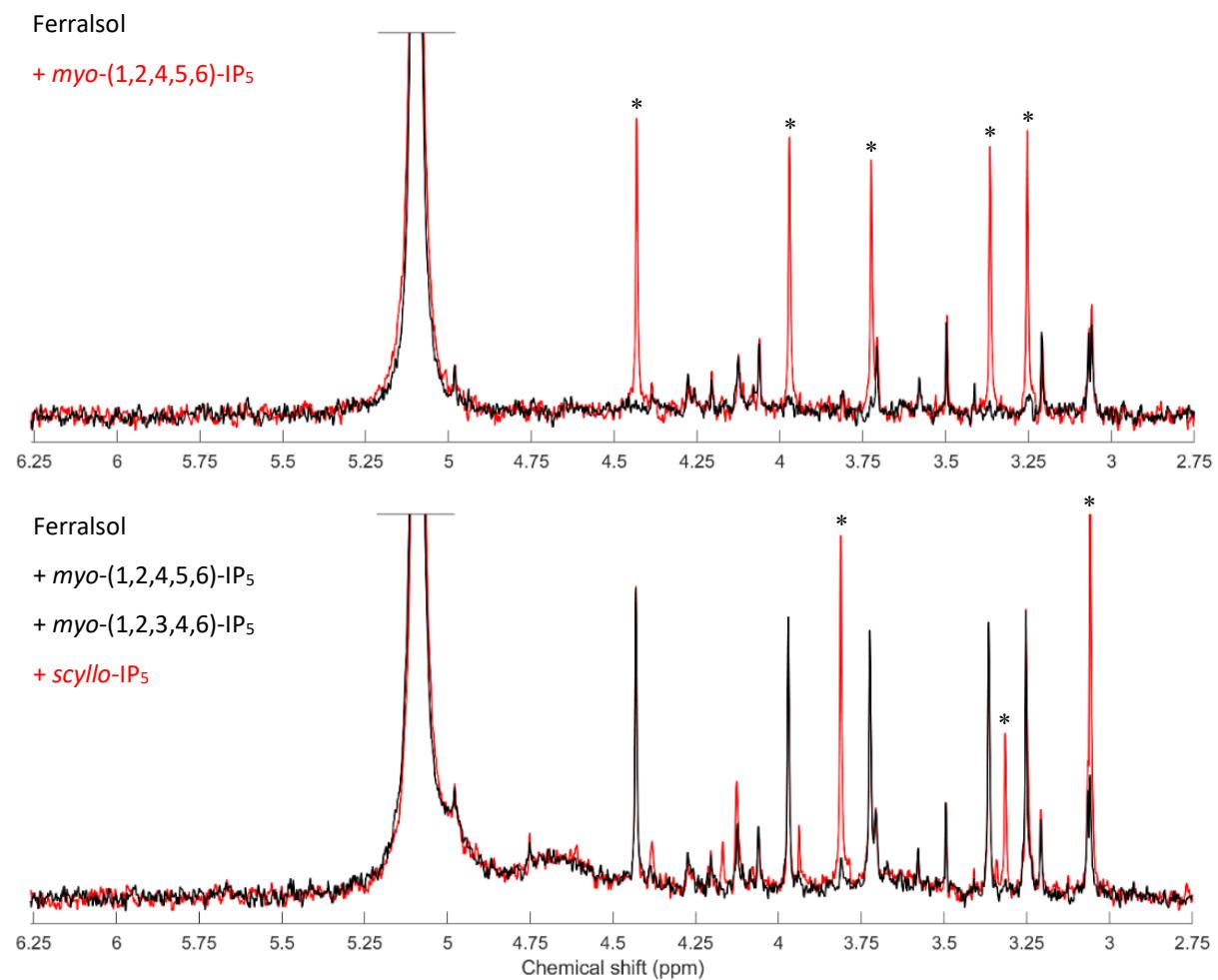
**NMR observability.** Measures of NMR observability were calculated for the untreated and the hypobromite oxidised extracts of all soils. Measures of NMR observability refer to the percentage of total P detected using NMR compared to that by ICP-OES. For the untreated soil extracts, measures of NMR observability ranged from 52 % (Gleysol) to 89 % (Ferralsol), with an average NMR observability of 66 %. For the hypobromite oxidised extracts, measures of NMR observability ranged from 58 % (Ferralsol) to 94 % (Cambisol), with an average value of 83 %.

### Inositol hexakisphosphate concentrations before and after hypobromite oxidation.

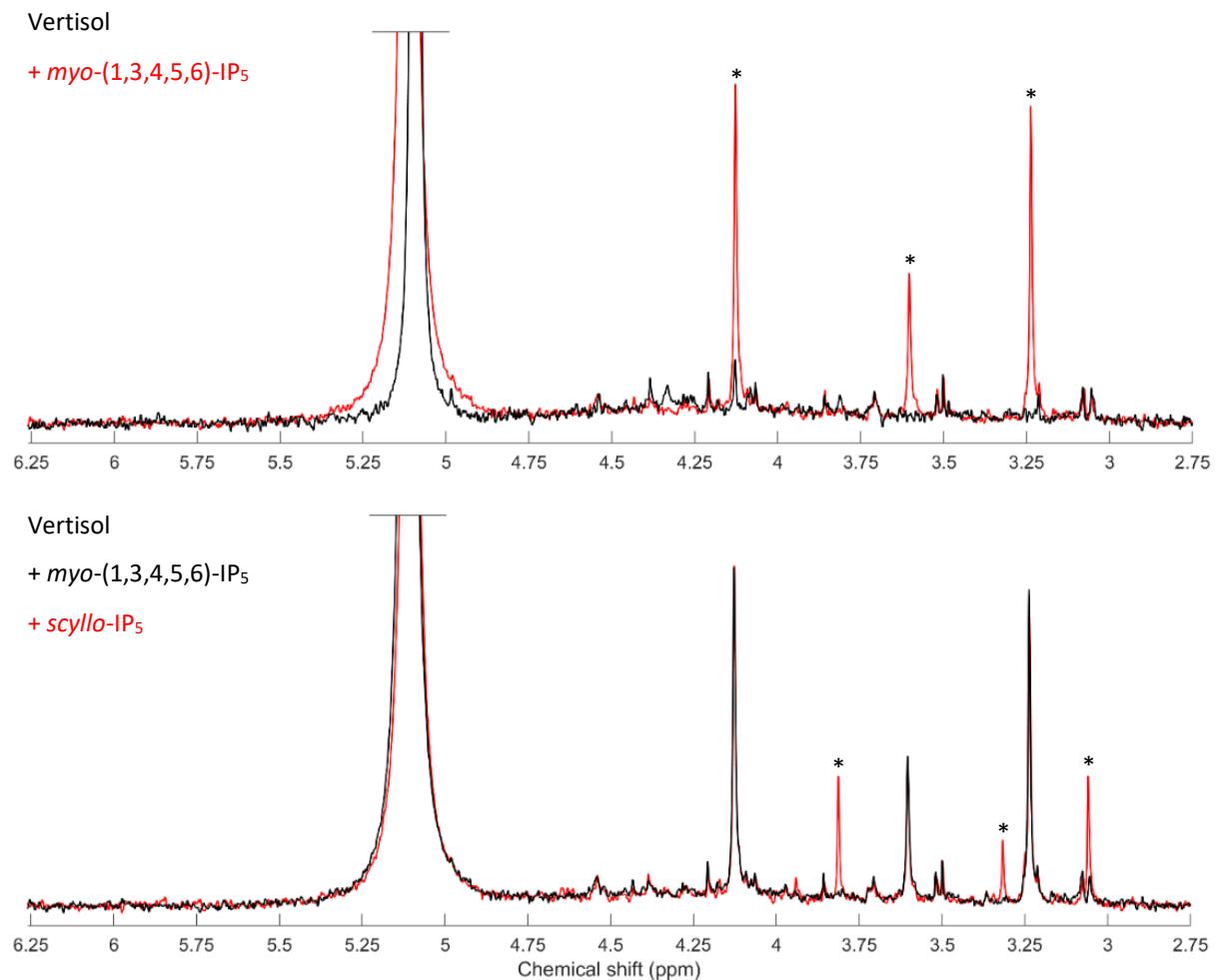
**Table SI1.** Concentrations of inositol hexakisphosphates in 0.25 M NaOH + 0.05 M EDTA soil extracts before and after hypobromite oxidation (HO). Quantification was based on spectral integration and deconvolution fitting of solution  $^{31}\text{P}$  NMR spectra. The proportion of P (%) detected in hypobromite oxidised extracts compared to that in untreated extracts is provided in brackets.

Concentrations (mg P/kg <sub>soil</sub> )		Ferralsol	Vertisol	Cambisol	Gleysol
<i>myo</i> -IP <sub>6</sub>	before HO	4.4	0.6	46.2	90.4
	after HO	1.1 (25)	0.6 (111)	26.3 (57)	85.0 (94)
<i>scyllo</i> -IP <sub>6</sub>	before HO	2.5	0.4	34.9	42.6
	after HO	0.4 (14)	0.3 (68)	15.6 (45)	41.1 (97)
<i>neo</i> -IP <sub>6</sub> 4-eq/2-ax	before HO	-	-	4.2	7.0
	after HO	-	-	1.4 (33)	8.8 (126)
<i>chiro</i> -IP <sub>6</sub>	before HO	-	-	7.2	6.7
	after HO	-	-	9.4 (130)	8.6 (128)

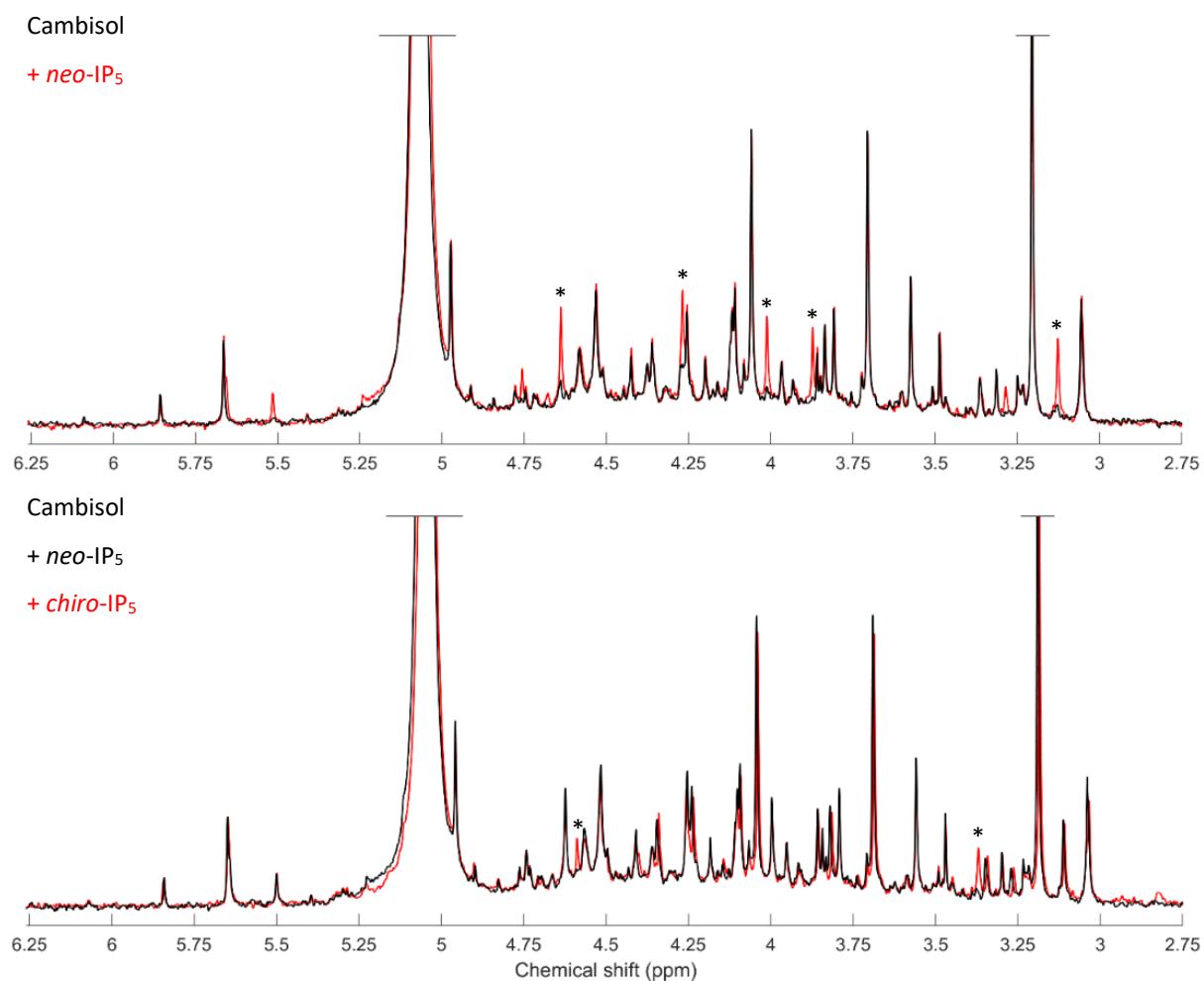
**Solution  $^{31}\text{P}$  NMR spectra of spiked hypobromite oxidised soil extracts**



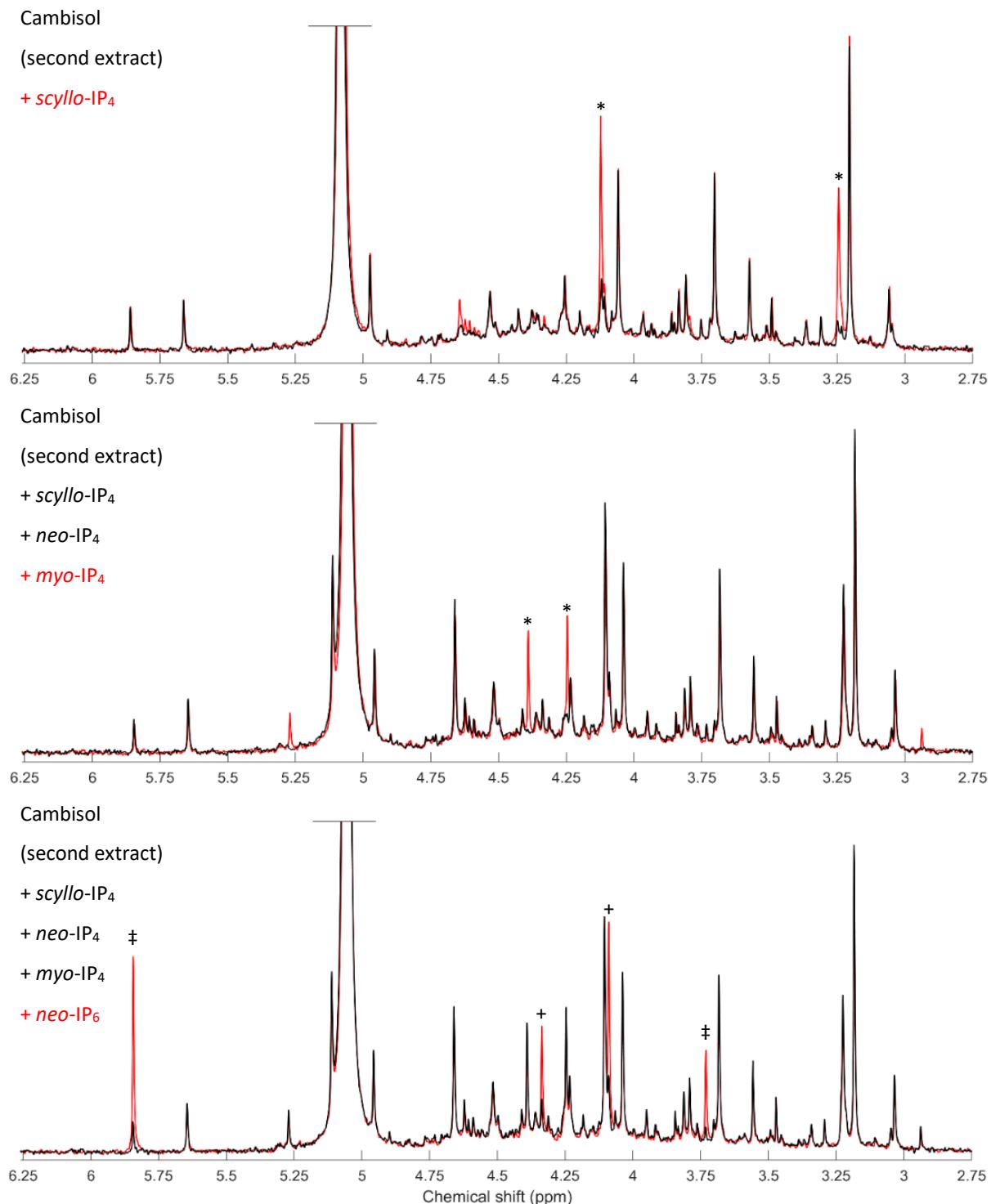
**Figure SI1.** Solution  $^{31}\text{P}$  NMR spectra of the orthophosphate and phosphomonoester region on Ferralsol extract following hypobromite oxidation (black trace), and also that following a spike with an IP standard (red trace). Peaks assigned to the IP standard marked with \*.



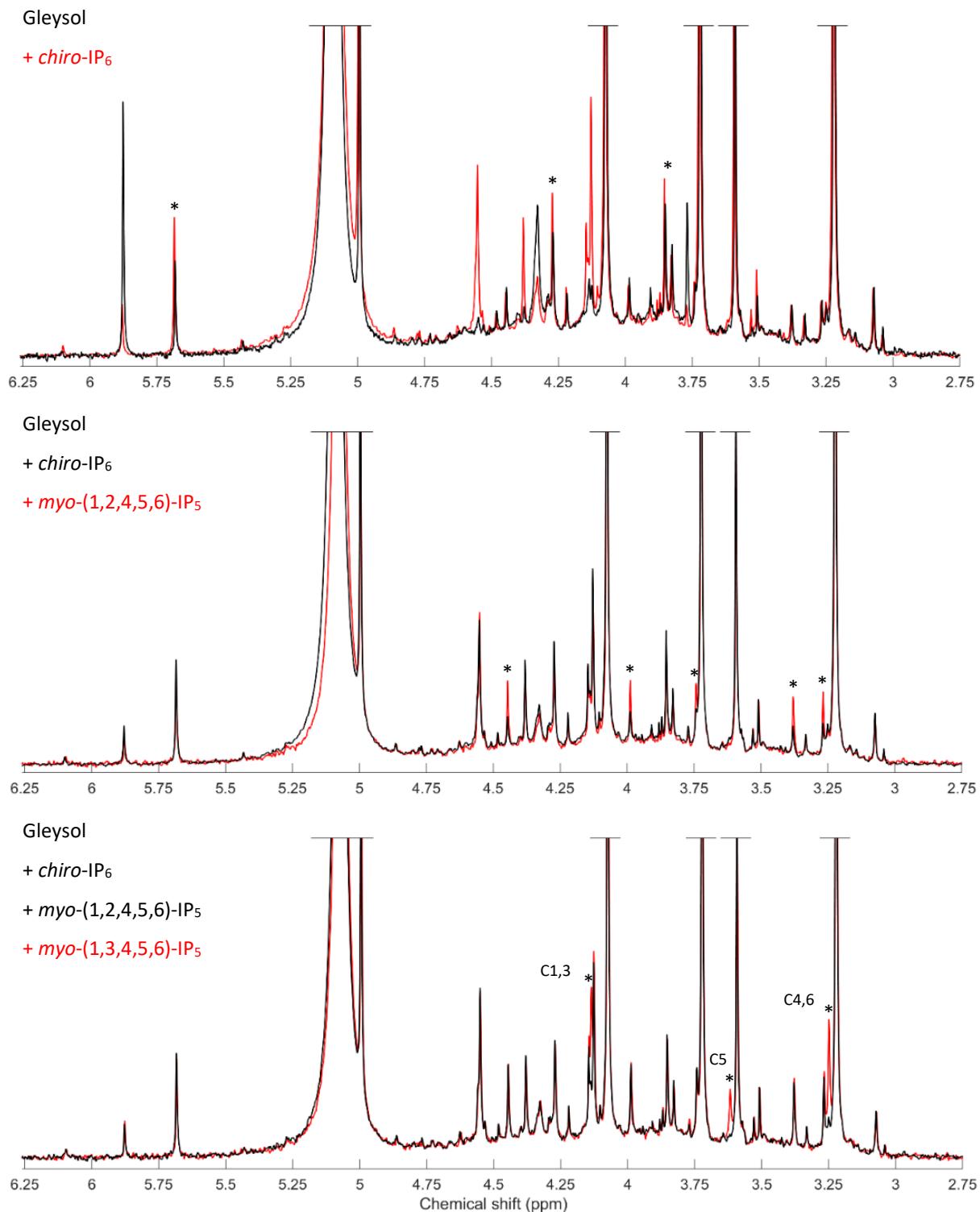
**Figure SI2. Solution  $^{31}\text{P}$  NMR spectra of phosphomonoester region of hypobromite oxidised 0.25 M NaOH + 0.05 M EDTA Vertisol extract. Spiked spectrum with indicated standard in red. Peaks assigned to standard marked with \*.**



**Figure SI3.** Solution <sup>31</sup>P NMR spectra of phosphomonoester region of hypobromite oxidised 0.25 M NaOH + 0.05 M EDTA Cambisol extract. Spiked spectrum with indicated standard in red. Peaks assigned to standard marked with \*.

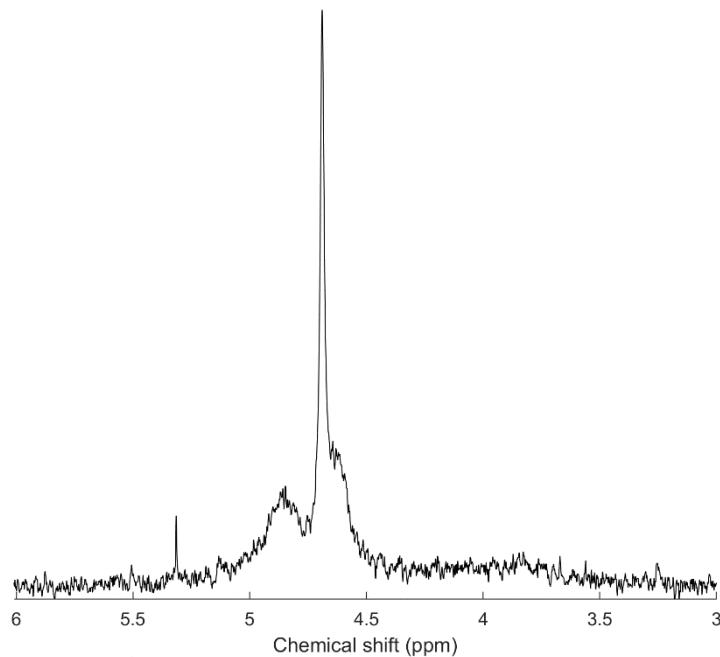


**Figure SI4.** Solution <sup>31</sup>P NMR spectra of phosphomonoester region of hypobromite oxidised 0.25 M NaOH + 0.05 M EDTA Cambisol extract. Spiked spectrum with indicated standard in red. Peaks assigned to 4-equatorial/2-axial conformation marked with ‡, peaks assigned to 2-equatorial/4-axial conformation marked with +.

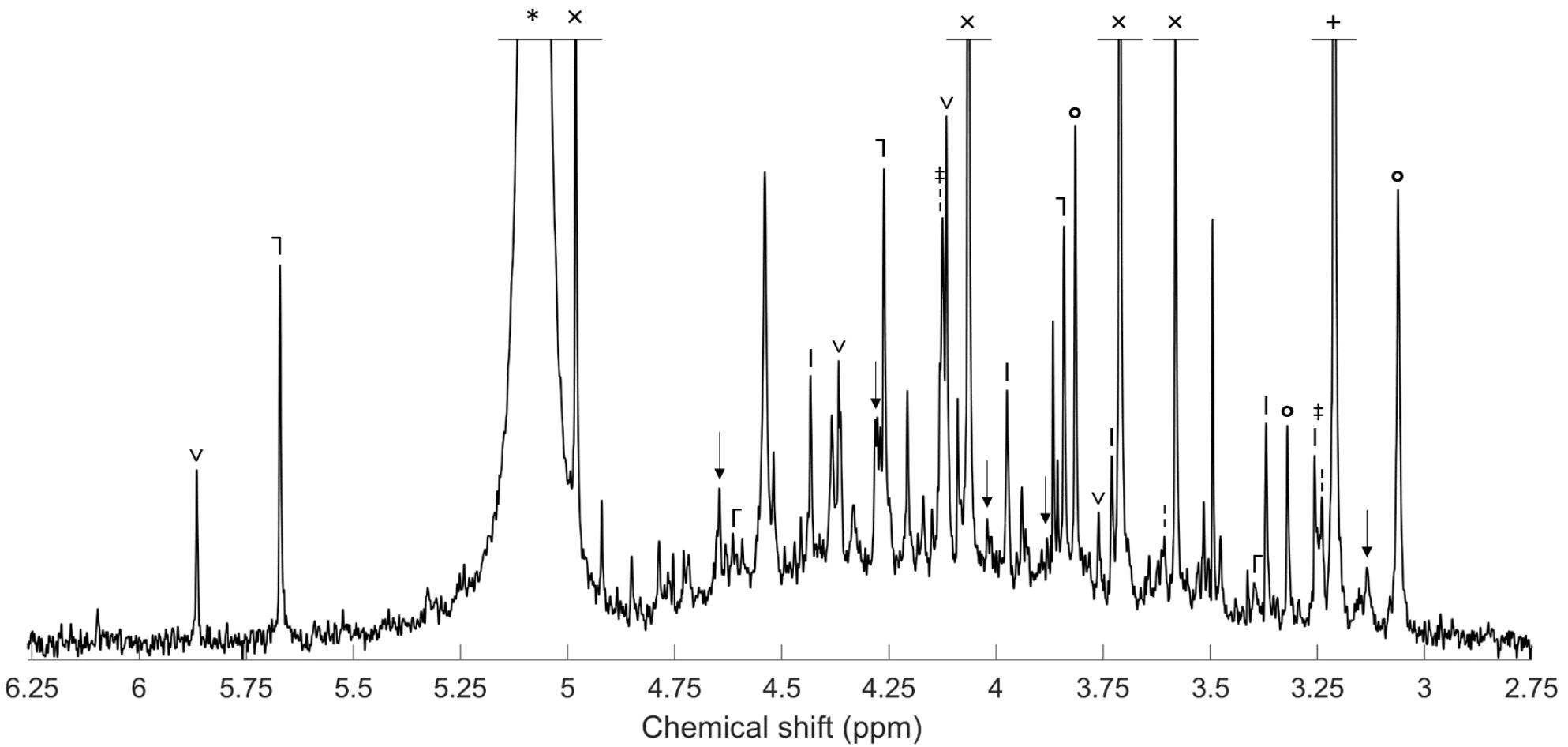


**Figure S15.** Solution  $^{31}\text{P}$  NMR spectra of phosphomonoester region of hypobromite oxidised 0.25 M NaOH + 0.05 M EDTA Gleysol extract. Spiked spectrum with indicated standard in red. Peaks assigned to standard marked with \*. For *myo*-(1,3,4,5,6)-IP<sub>5</sub>, the respective phosphorylated carbon nuclei of the inositol have been marked based on the  $^{31}\text{P}$  NMR spectrum prediction of the program Mnova 11.0.4 (©Mestrelab Research).

**Transverse relaxation time of an orthophosphate solution.** The analysis of a 0.25 M NaOH + 0.05 M EDTA solution containing 910 mg KH<sub>2</sub>PO<sub>4</sub>/L resulted in a single orthophosphate peak in the NMR spectrum ( $\delta$  5.09 ppm) with a linewidth at peak half height of 0.56 Hz. Transverse relaxation experiments were carried out (similar to that previously described) on the solution, which resulted in a T<sub>2</sub> time of 203 ms for orthophosphate.



**Figure SI7.** Solution  $^{31}\text{P}$  NMR spectrum of phosphomonoester region of purchased *myo*-(1,2,3,4,6)-IP<sub>5</sub> standard dissolved in 0.25 M NaOH + 0.05 M EDTA.



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Figure SI8. Solution  $^{31}\text{P}$  NMR spectra of phosphomonoester region of hypobromite oxidised 0.25 M NaOH + 0.05 M EDTA Cambisol extract. All identified peaks are marked: orthophosphate (\*), *myo*-IP<sub>6</sub> (x), *scyllo*-IP<sub>6</sub> (+), *neo*-IP<sub>6</sub> (v), *chiro*-IP<sub>6</sub> (↑), *myo*-(1,2,4,5,6)-IP<sub>5</sub> (l), *myo*-(1,3,4,5,6)-IP<sub>5</sub> (↑), *scyllo*-IP<sub>5</sub> (°), *neo*-IP<sub>5</sub> (↓), *chiro*-IP<sub>5</sub> (Γ), *scyllo*-(1,2,3,4)-IP<sub>4</sub> (‡). The chemical shifts in ppm of all identified peaks are listed in Table 5.