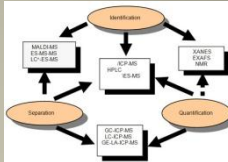


The Mechanisms of Arsenic bioremediation from water by the Green Microalgae *Chlorella vulgaris*



Middlesex University

In collaboration with



TESLA
Trace element
Speciation
Laboratory
Aberdeen

Leonardo Pantoja, Diane Purchase, Huw Jones, Jörg Feldmann and Hemda Garelick

OUTLINE



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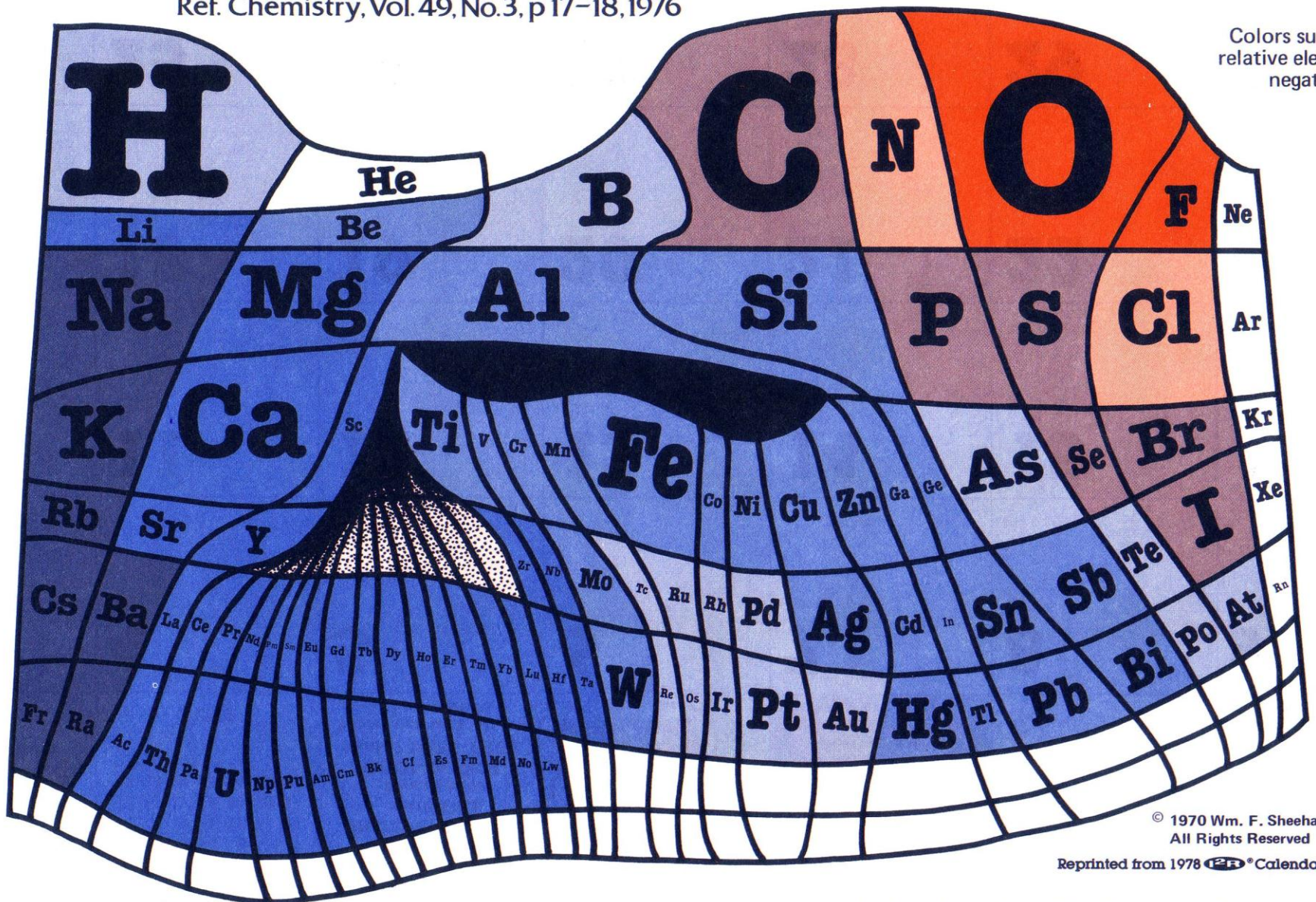
- ▶ Toxicity of arsenic to *C. vulgaris*
 - ASTM E1218 and flow cytometry
- ▶ Focused sonication for extraction for As-phytochelatin complexes
- ▶ Formation of As-GS/PC complexes as detoxification mechanism
 - Exposure to As(III) (Sodium arsenite)
 - Exposure to DMA(V) (Dimethylarsinic acid)
 - Exposure to As(V) (Sodium arsenate)
- ▶ Total As-GS/PC formation
- ▶ Transport of As-GS/PC to vacuoles
- ▶ Conclusions

The Elements According to Relative Abundance

A Periodic Chart by Prof. Wm. F. Sheehan, University of Santa Clara, CA 95053

Ref. Chemistry, Vol. 49, No. 3, p 17-18, 1976

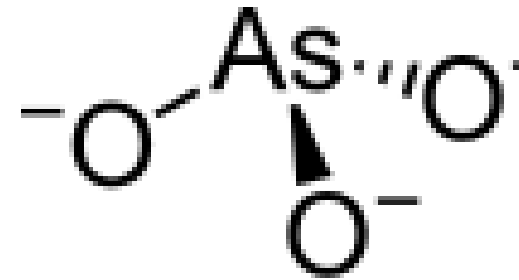
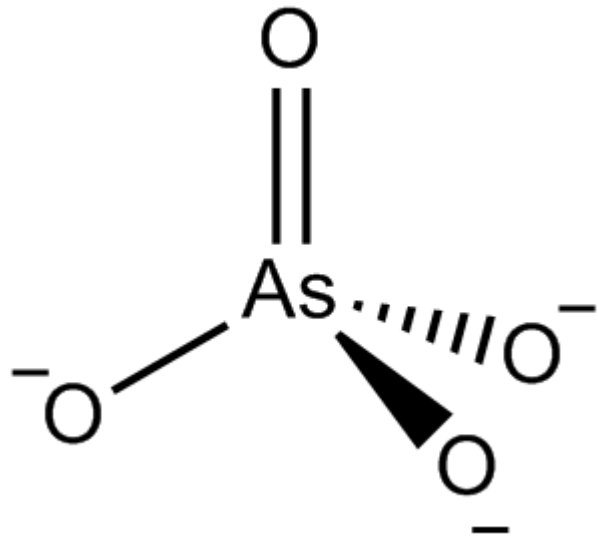
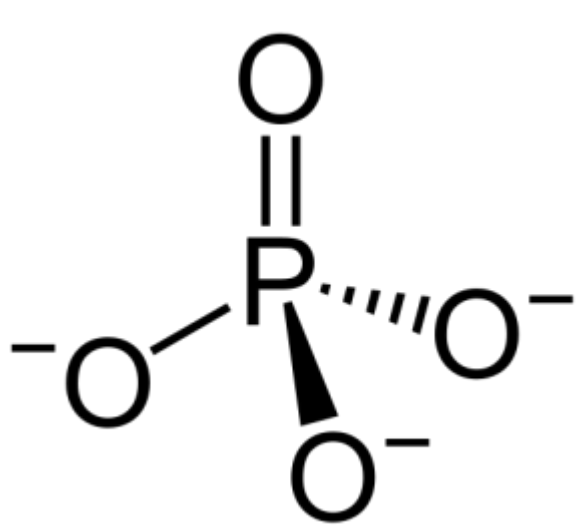
Colors suggest relative electro-negativity



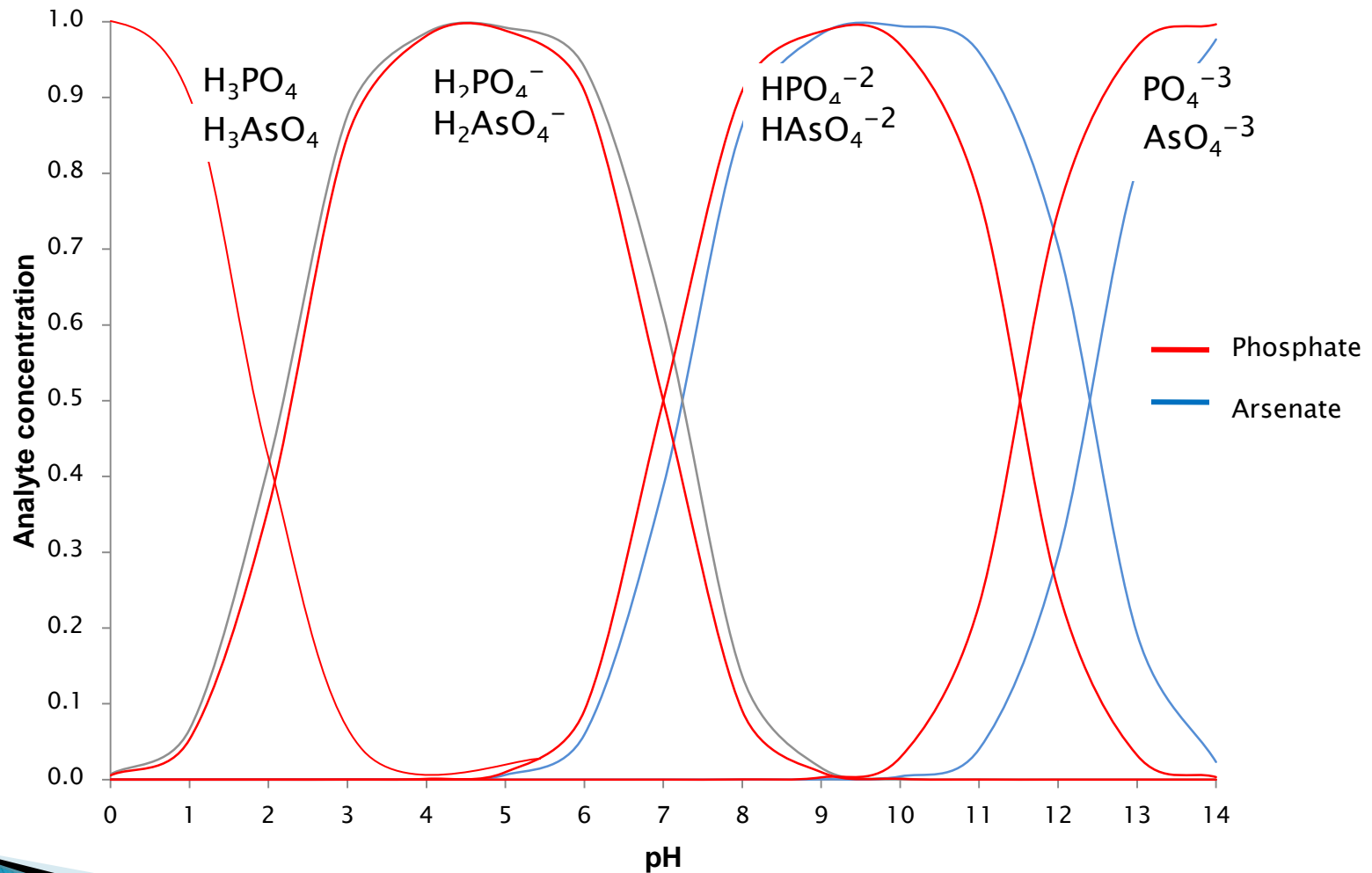
© 1970 Wm. F. Sheehan
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Reprinted from 1978  Calendar.

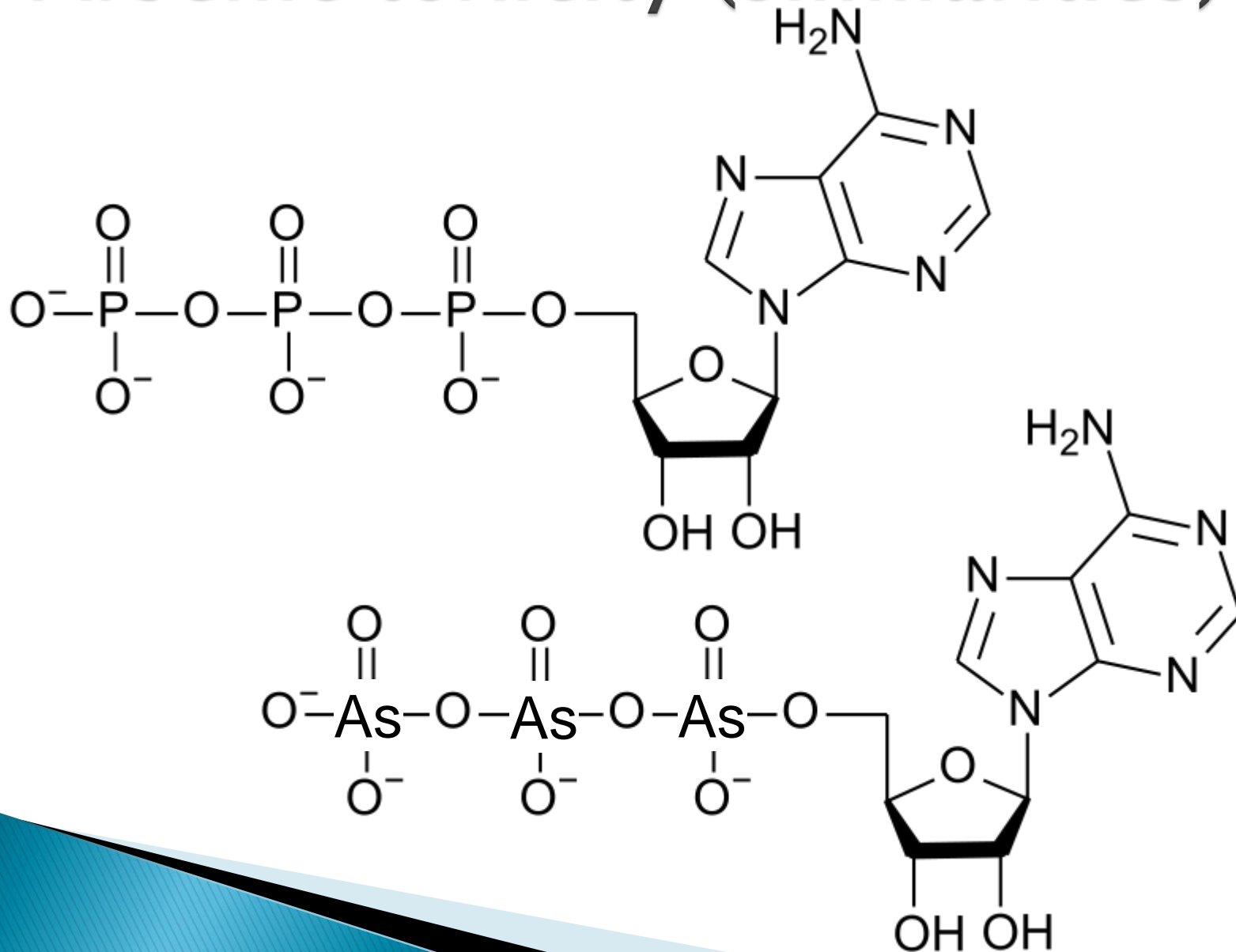
Arsenic toxicity (similarities)



Arsenic toxicity (similarities)



Arsenic toxicity (similarities)



TOXICITY OF ARSENIC TO *C. VULGARIS*



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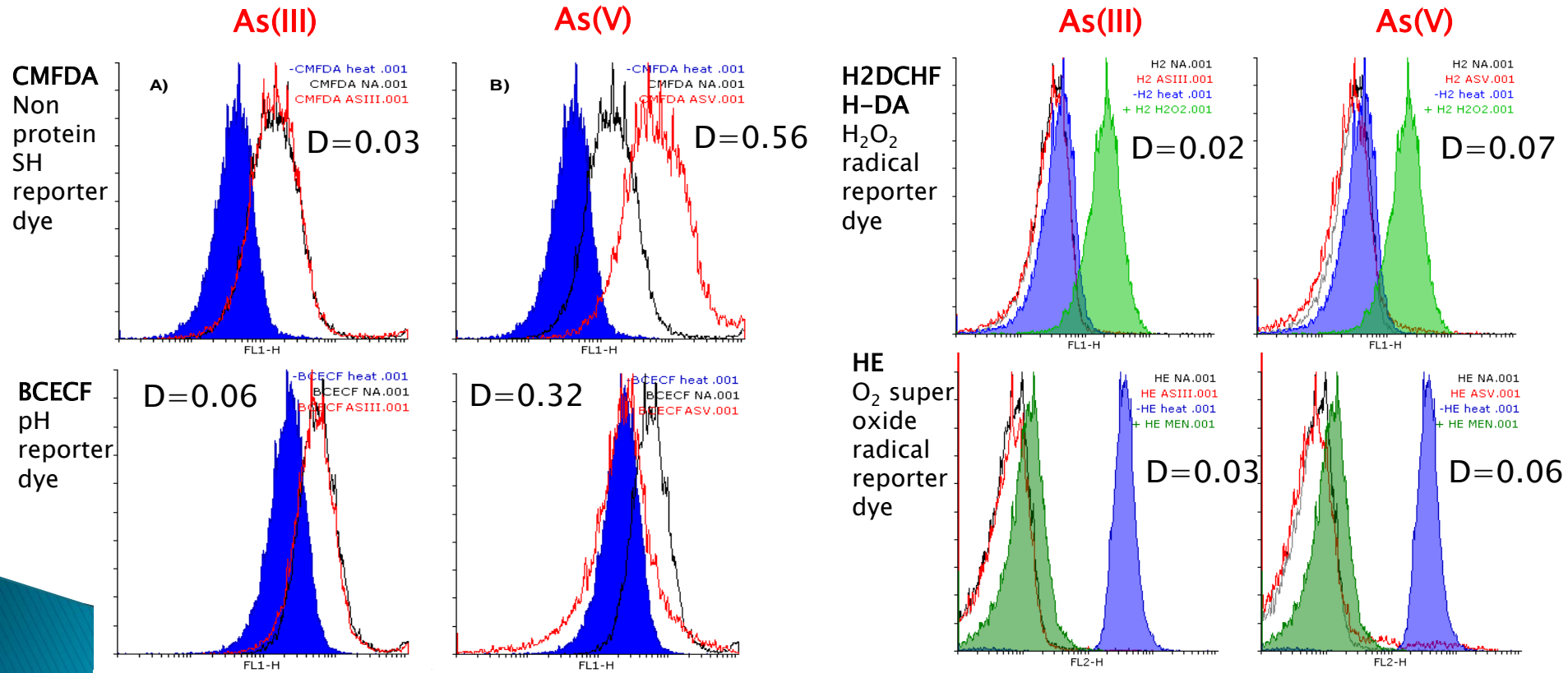
- ▶ Toxicity of As(III) (72 h @ 0.3mg L⁻¹ phosphate)

IC₅₀=64.23 mg L⁻¹

- ▶ Toxicity of As(V) (72 h @ 0.3mg L⁻¹ phosphate)

IC₅₀=1.07 mg L⁻¹

Chlorophyll as surrogate for cell health

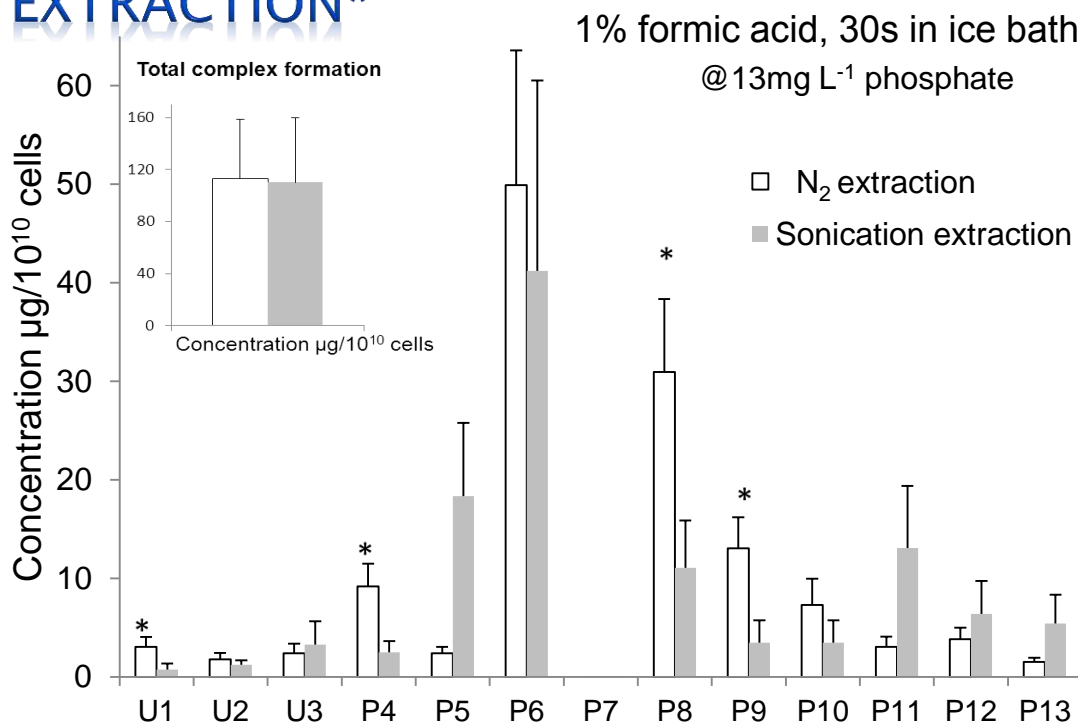


n=20,000 cells, Strong D>0.3, Moderate 0.2< D <= 0.3, Weak 0.15< D <=0.20, Negligible D <= 0.15
Kolmogorov-Smirnov statistics

ENHANCED DETERMINATION OF ARSENIC-PHYTOCHELATIN COMPLEXES IN *C. VULGARIS* USING FOCUSED SONICATION EXTRACTION*



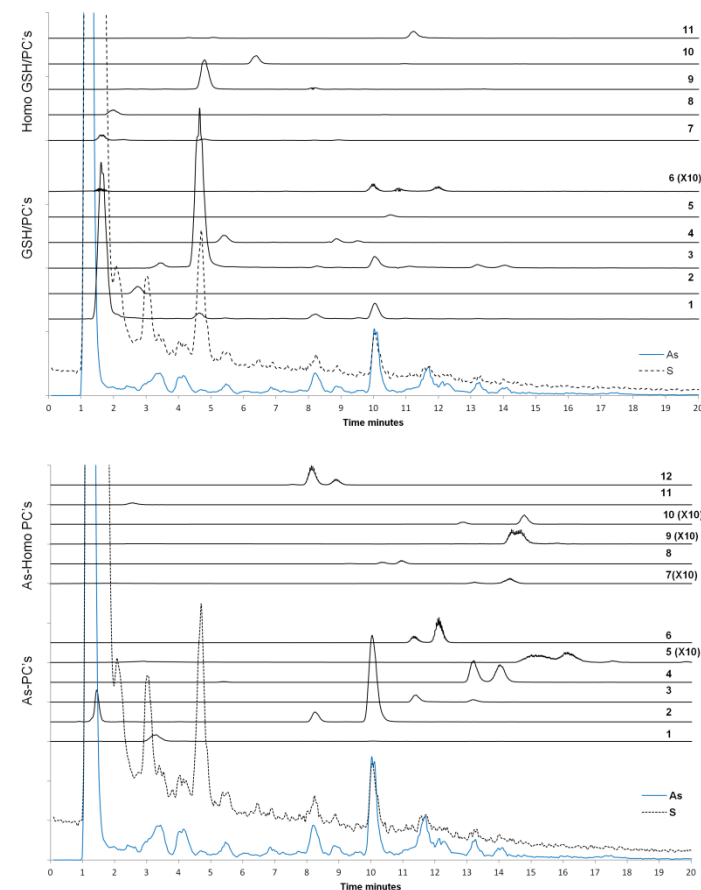
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Sample	$\mu\text{g As g}^{-1}$	% Recovery
Total extraction <i>C. vulgaris</i>	83.2 SE = 2.04, n = 6	
Sonication <i>C. vulgaris</i>	59.2 SE = 1.14, n = 6	71.10%
Quality control	$\mu\text{g As L}^{-1}$	
Total extraction Kelp	23.5 SE = 0.60, n = 9	
Sonication Kelp	22.2 SE = 0.28, n = 11	94.70%
SRM 2669	48.1 %RSD = 2.16	
Certified value	50.7 \pm 6.3 (95% CI)	

SE = Standard error, n = Number of samples, % RSD = Relative standard deviation

Online HPLC-ICPMS/ESMS



21 peptides

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Anal. Methods, 2014,6, 791-797

EXPOSURE TO As(III)*

Cells exposed to 50mg L⁻¹ for 48h



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Unbound peptides

Molecule	Formula	Monoisotopic mass (M+H ⁺ or M+2H ⁺)	Experimental mass	Difference ppm
GSH/PC				
GSH	C ₁₀ H ₁₇ N ₃ O ₆ S	308.0916	308.0912	-1.34
GSSG	C ₂₀ H ₃₂ N ₆ O ₁₂ S ₂	613.1598	613.1598	0
Reduced PC ₂	C ₁₈ H ₂₉ N ₅ O ₁₀ S ₂	540.1434	540.1433	-0.16
Oxidised PC ₂	C ₁₈ H ₂₇ N ₅ O ₁₀ S ₂	538.1278	538.1289	2.04
Reduced PC ₃	C ₂₆ H ₄₁ N ₇ O ₁₄ S ₃	772.1952	772.1952	-0.01
Oxidised PC ₃	C ₂₆ H ₃₉ N ₇ O ₁₄ S ₃	770.1795	770.1795	0
Reduced PC ₄	C ₃₄ H ₅₃ N ₉ O ₁₈ S ₄	1004.247	Not found	
Ala GSH/PC				
γ-(Glu-Cys)-Ala	C ₁₁ H ₁₉ N ₃ O ₆ S	322.1073	322.1072	-0.21
γ-(Glu-Cys) ₂ -Ala	C ₁₉ H ₃₁ N ₅ O ₁₀ S ₂	554.1591	554.1578	-2.2
desGly GSH/PC				
γ-(Glu-Cys)	C ₈ H ₁₄ N ₂ O ₅ S	251.0702	251.0706	1.55
γ-(Glu-Cys) ₂	C ₁₆ H ₂₆ N ₄ O ₉ S ₂	483.1219	483.1217	-0.58
γ-(Glu-Cys) ₃	C ₂₄ H ₃₈ N ₆ O ₁₃ S ₃	715.1737	715.1747	1.41

Glutathione/PC homologues (terminal amino acid Gly substituted by Ala, Ser, Gln, Glu or is absent)

Arsenic bound peptides

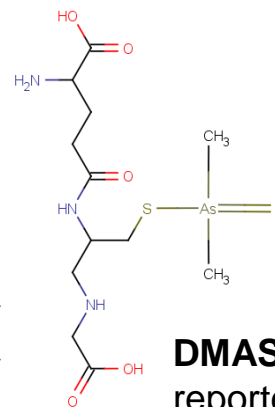
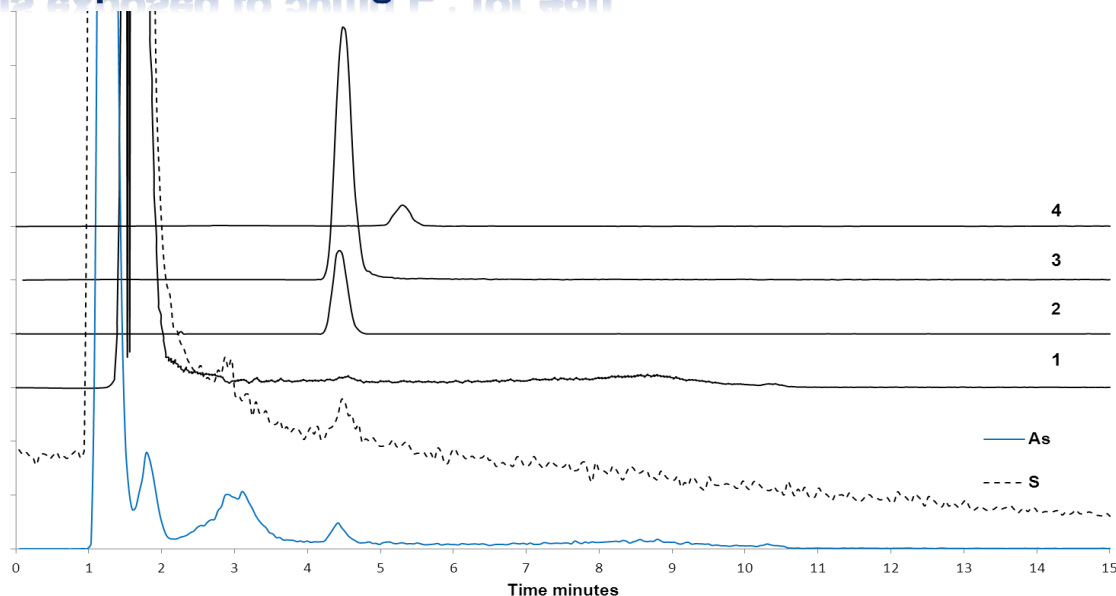
Molecule	Formula	Monoisotopic mass (M+H ⁺ or M+2H ⁺)	Experimental mass	Difference ppm
GSH/PC				
As(III)-PC ₂	C ₁₈ H ₂₈ N ₅ O ₁₁ S ₂ As	630.0443	630.0437	-0.88
As(III)-PC ₃	C ₂₆ H ₃₈ N ₇ O ₁₄ S ₃ As	844.0933	844.0931	-0.19
GS-As(III)-PC ₂	C ₂₈ H ₄₃ N ₈ O ₁₆ S ₃ As	460.0666	460.0663	-0.06
		919.1253	919.1247	-0.69
As(III)-(PC ₂) ₂	C ₃₆ H ₅₄ N ₁₀ O ₂₀ S ₄ As	576.0925	576.0923	-0.37
As(III)-PC ₄	C ₃₄ H ₅₀ N ₉ O ₁₈ S ₄ As	1076.1451	1076.1455	0.37
MMA(III)-PC ₂	C ₁₉ H ₃₀ N ₅ O ₁₀ S ₂ As	628.0728	628.0729	0.12
Ala GSH/PC				
As(III)-γ-(Glu-Cys) ₃ -Ala	C ₂₇ H ₄₀ N ₇ O ₁₄ S ₃ As	858.109	858.1082	-0.87
GS-As(III)-γ-(Glu-Cys) ₂ -Ala	C ₂₉ H ₄₅ N ₈ O ₁₆ S ₃ As	467.0744	467.0744	0.09
As(III)-γ-((Glu-Cys) ₂) ₂ -Ala	C ₃₇ H ₅₇ N ₁₀ O ₂₀ S ₄ As	583.1003	583.101	1.19
MMA(III)-γ-(Glu-Cys) ₂ -Ala	C ₂₀ H ₃₂ N ₅ O ₁₀ S ₂ As	642.0885	642.0889	0.66
desGly GSH/PC				
As(III)-γ-(Glu-Cys) ₂	C ₁₆ H ₂₅ N ₄ O ₁₀ S ₂ As	573.0306	573.0318	2.11
GS-As(III)-γ-(Glu-Cys) ₂	C ₂₆ H ₄₀ N ₇ O ₁₅ S ₃ As	431.5558	431.5558	-0.17

Newly reported peptides

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Anal. Methods, 2014,6, 791-797

EXPOSURE TO DMA*

Cells exposed to 50mg L⁻¹ for 48h



DMAS^V-GS has only been reported once in *Brassica Oleracea* plants:

Cabbage, broccoli, cauliflower, kale, Brussels sprouts, collard greens, savoy, among other

1 GSH (308), 2 **DMAS^V-GS (444)**, 3 Reduced PC₂ (540) and 4 Oxidised PC₂ (538), As and S signals from HR-ICP-MS

Molecule	Formula (M)	Monoisotopic mass (M+H ⁺)	Experimental mass	Difference ppm
GSH	C ₁₀ H ₁₇ N ₃ O ₆ S	308.0916	308.0918	0.65
Red PC ₂	C ₁₈ H ₂₉ N ₅ O ₁₀ S ₂	540.1434	540.1433	-0.16
Ox PC ₂	C ₁₈ H ₂₈ N ₅ O ₁₀ S ₂	538.1278	538.1288	1.86
DMAS ^V -GS	C ₁₂ H ₂₃ N ₃ O ₆ S ₂ As	444.0244	444.0247	0.56

@13mg L⁻¹
phosphate

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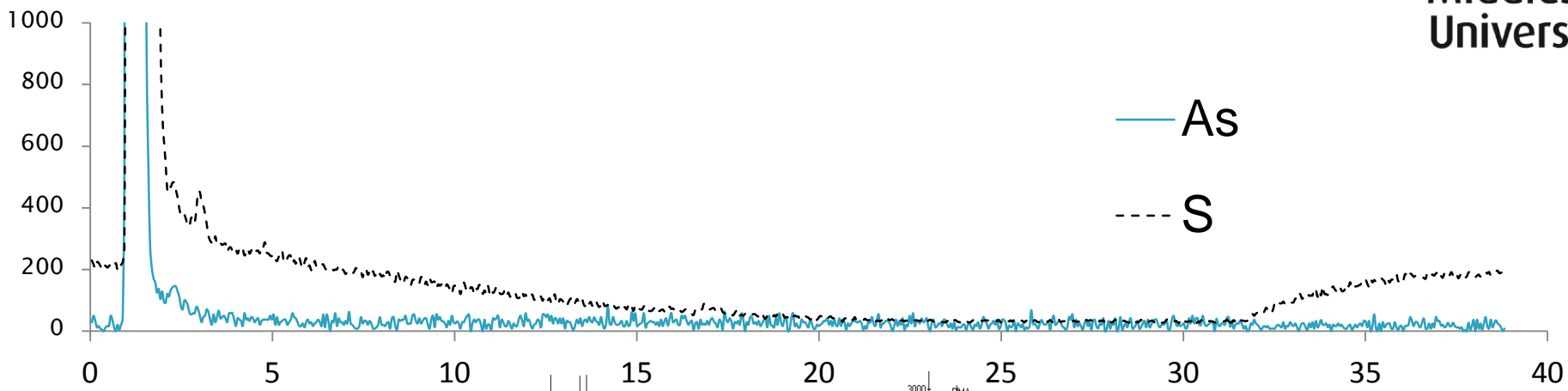
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EXPOSURE TO As(V)*

Cells exposed to 50mg L⁻¹ for 48h

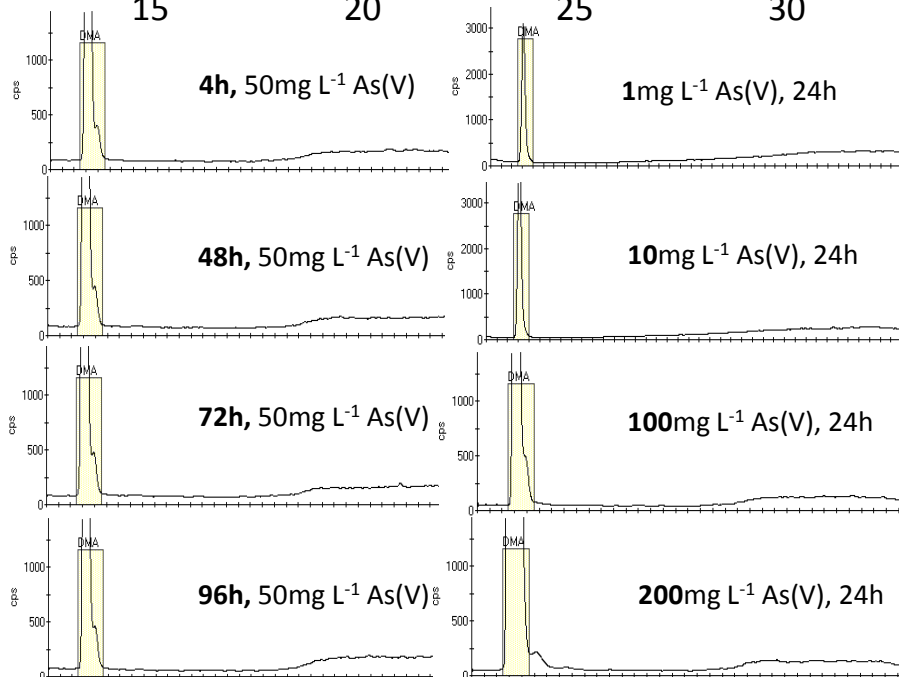


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@13mg L⁻¹ phosphate

At this level of phosphate cells are not under stress

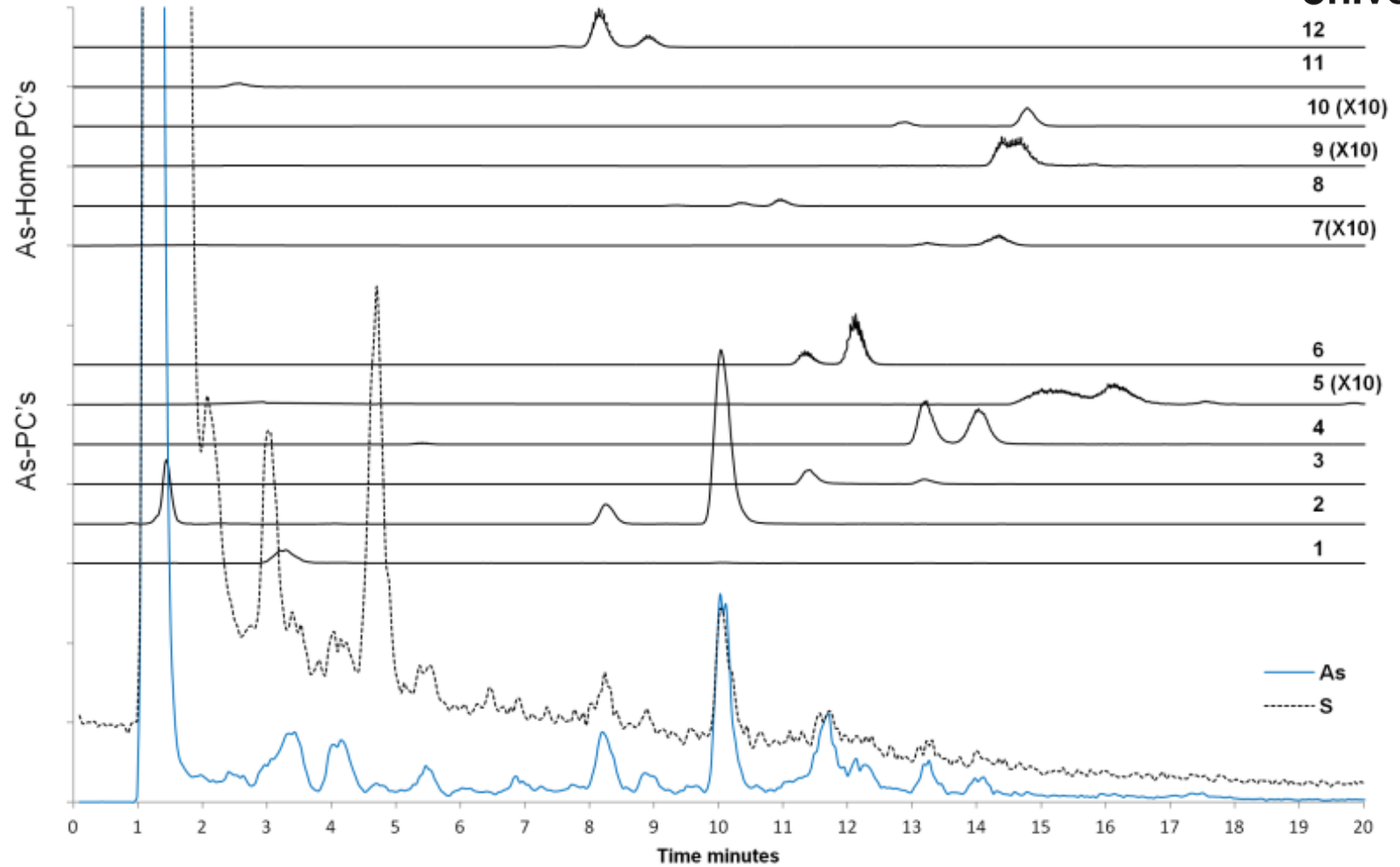


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EXPOSURE TO As(III)*



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@13mg L⁻¹
phosphate

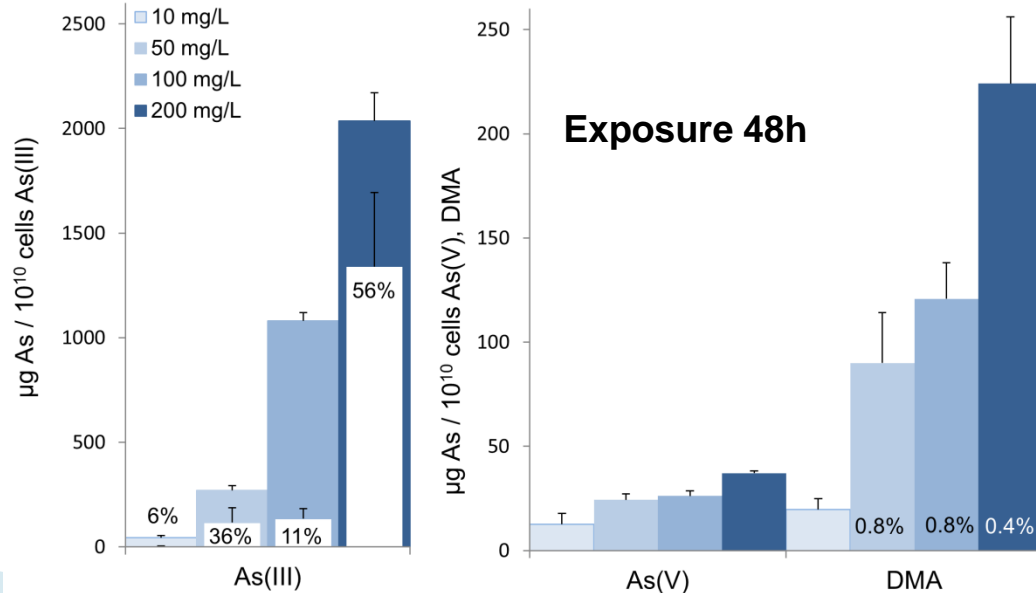
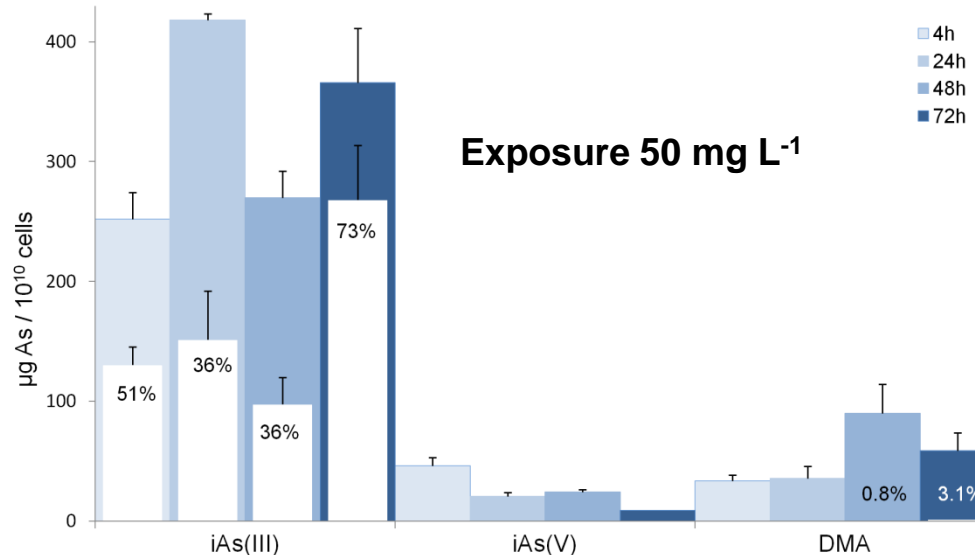
TOTAL As-GS/PC FORMATION



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@13mg L⁻¹ phosphate

- Coloured bars - total amount of arsenic
- White bars- amount of As-GS/PC (percentage of total arsenic)



Vertical bars denote + 1 standard error, n = 3

TRANSPORT OF As-GS/PC TO VACUOLES



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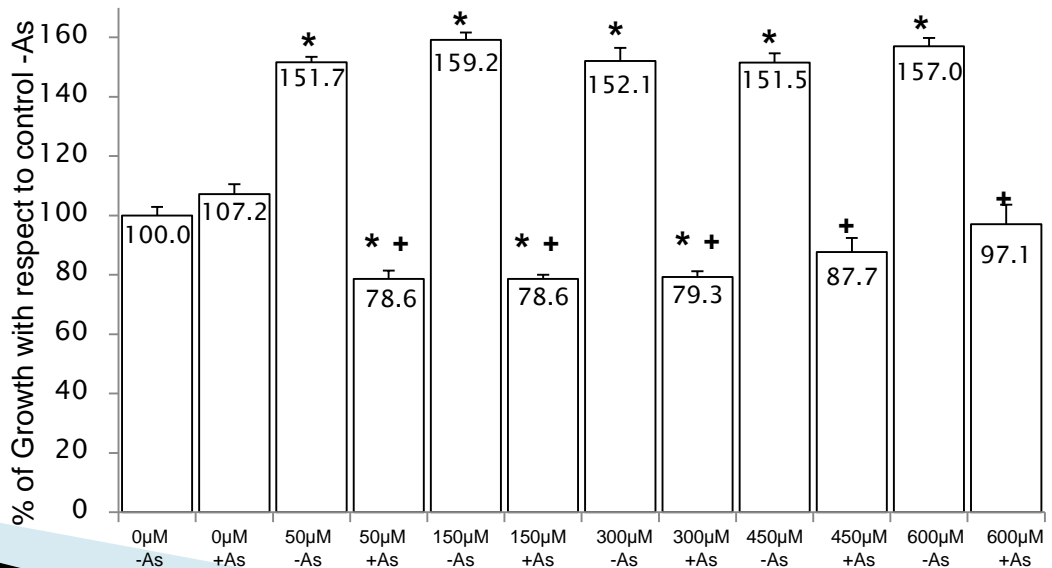
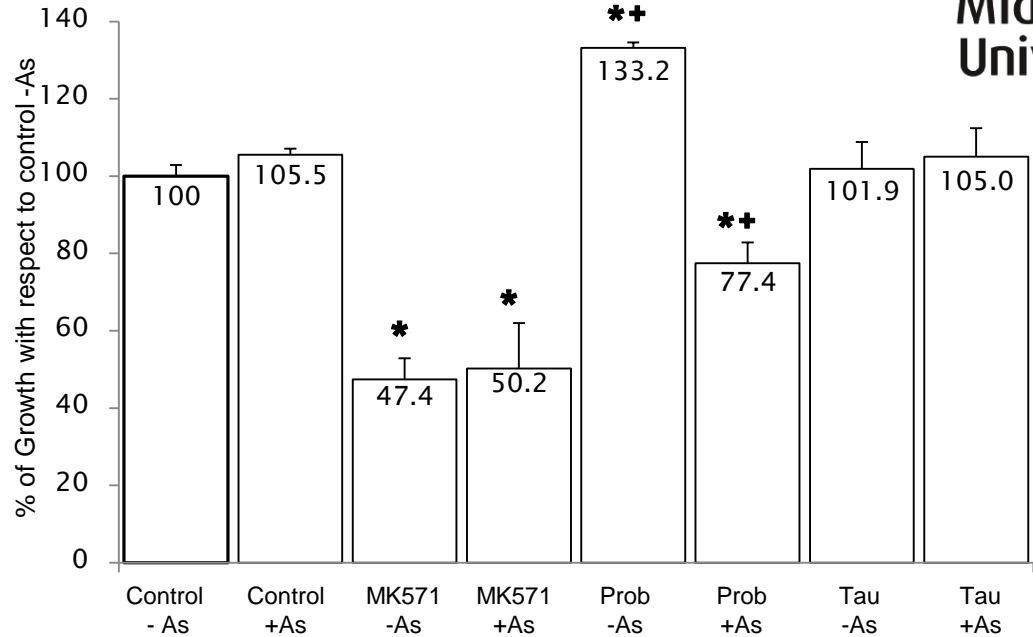
Toxicity (72 h) to As(III) (50 mg L⁻¹)
@13mg L⁻¹ phosphate

Test the presence of ABCC1 and ABCC2 inhibitors:

- MK571 (25 μM)
- Probenecid (Prob, 500 μM)
- Sodium taurocholate (Tau, 50 μM).

ABC = ATP-binding cassette transporter
MRP = Multidrug resistance-associated protein

* (P < 0.05) with respect to control
+ (P < 0.05) with respect to absence/presence of arsenic
Vertical bars + 1 standard error
Control n=6, experiments n = 3

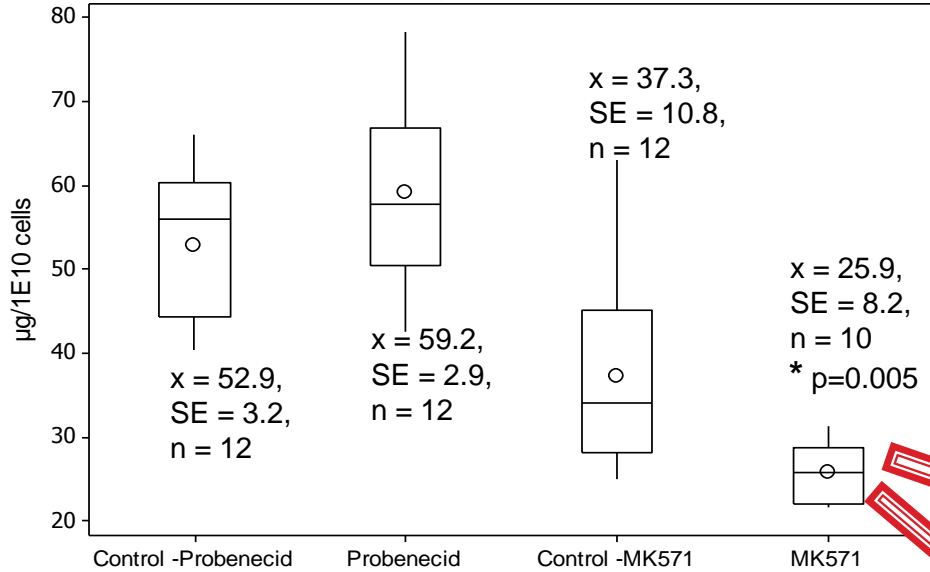


TRANSPORT OF As-GS/PC TO VACUOLES

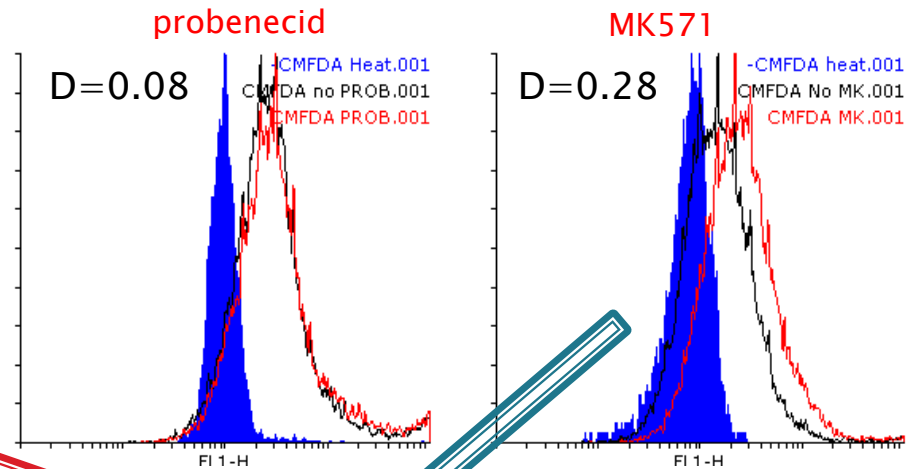


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Cells exposed to 150 mg L⁻¹ of As(III) for 24 h

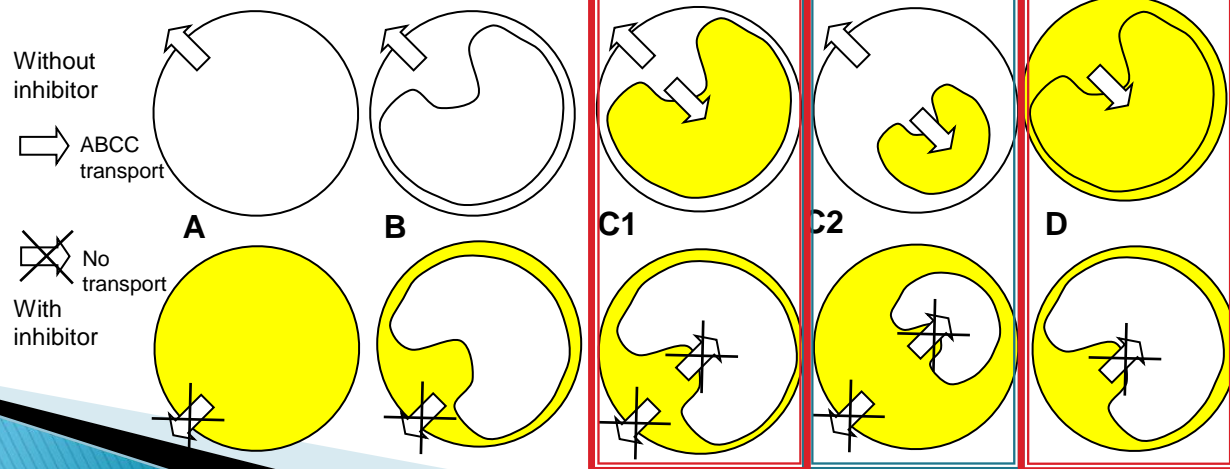


CMFDA substrate for ABCC transport



D > 0.3, Moderate 0.2 < D <= 0.3, Weak 0.15 < D <= 0.20, Negligible D <= 0.15

Treatment time 60 min, n=20,000 cells



CONCLUSIONS



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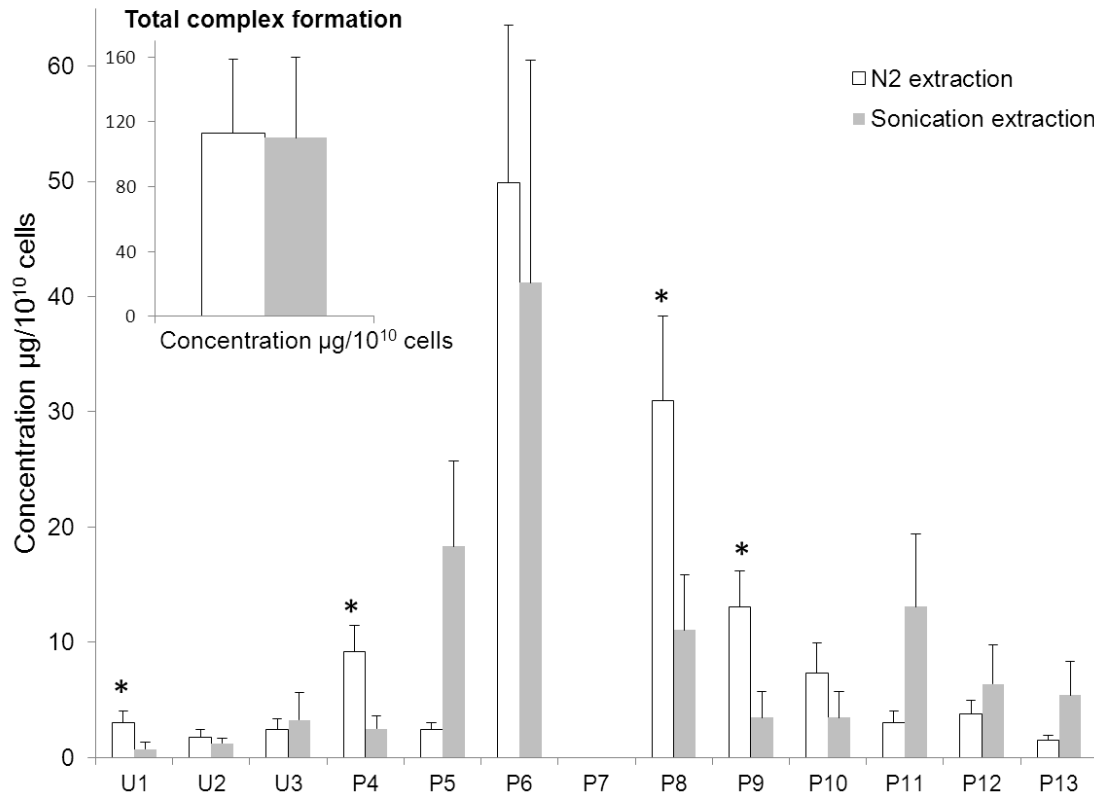
- As(V) is more toxic than As(III) to *C. vulgaris* cells at the same concentration of phosphate
- As(III) triggers the formation of **As-GS/PC** molecules
- As(V) does not trigger the formation of **As-GS/PC** molecules when cells are not under stress
- DMA triggers the formation of **DMAS^V-GS** but it is unlikely that this is part of a detoxification mechanism
- ABCC1 and ABCC2 are involved in **As-GS/PC** transport to acidic vacuoles in *C. vulgaris*



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Thank you for listening.

Any questions?



U1-3

Unknowns

P4

GS-As(III)-PC₂/GS-As(III)- γ -(Glu-Cys)₂,

P5

As(III)- γ -(Glu-Cys)₂

P6

GS-As(III)-PC₂

P7

GS-As(III)- γ -(Glu-Cys)₂-Ala

P8

As(III)-PC₃/MMA(III)-PC₂

P9

MMA(III)-PC₂

P10

As(III)-PC₃/As(III)-(PC₂)₂

P11

As(III)-(PC₂)₂/As(III)- γ -(Glu-Cys)₃-Ala/As(III)- γ -((Glu-Cys)₂)₂-Ala/

MMA(III)- γ -(Glu-Cys)₂-Ala,

P12

As(III)-PC₄

P13

As(III)-PC₄