

# Investigation of New Habitats for Mercury Methylation

- case studies at rice paddy and landfill leachate environments



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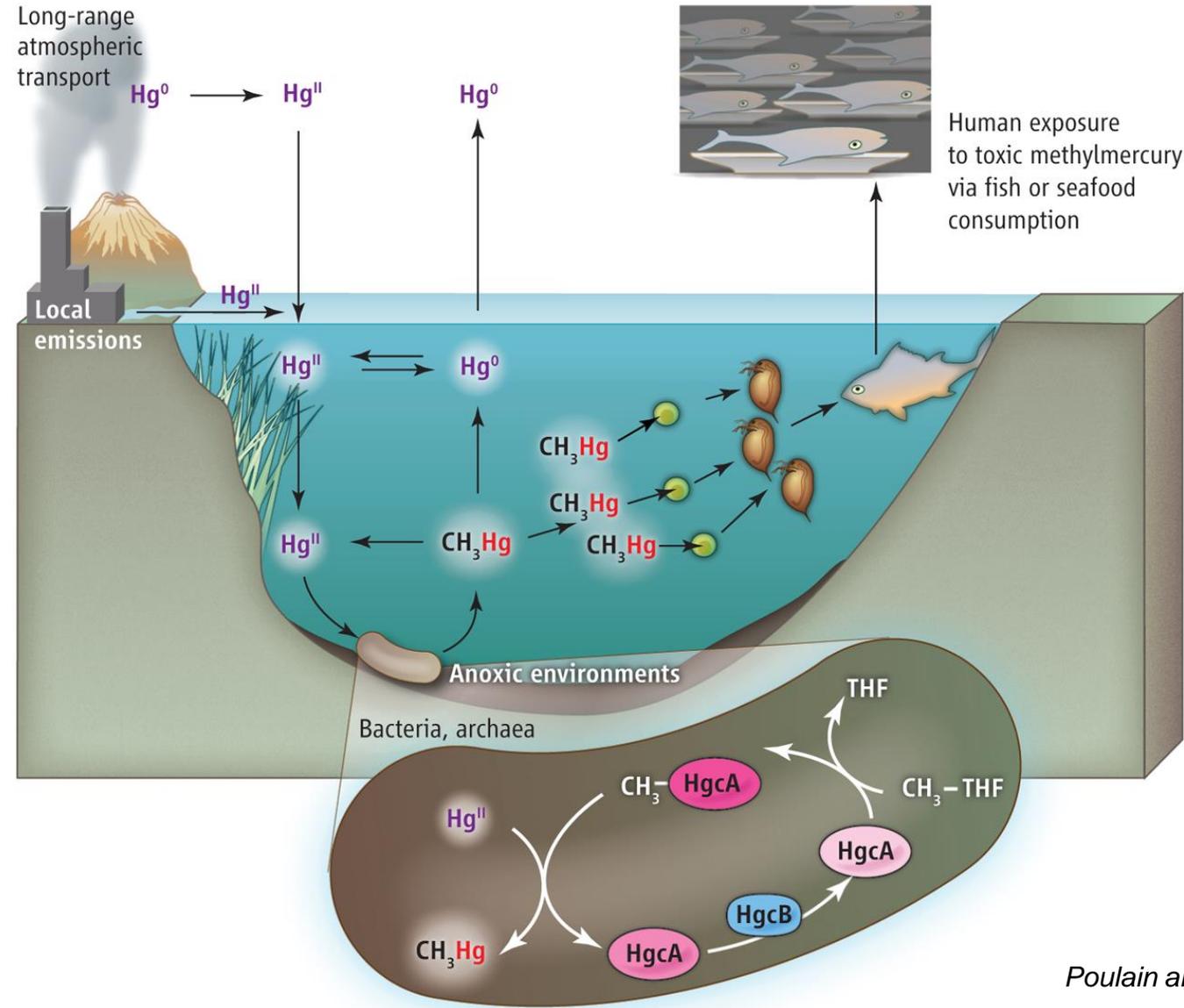


行政院環境保護署  
Environmental Protection Administration  
Executive Yuan, R.O.C (Taiwan)

科技部 Ministry of Science and Technology



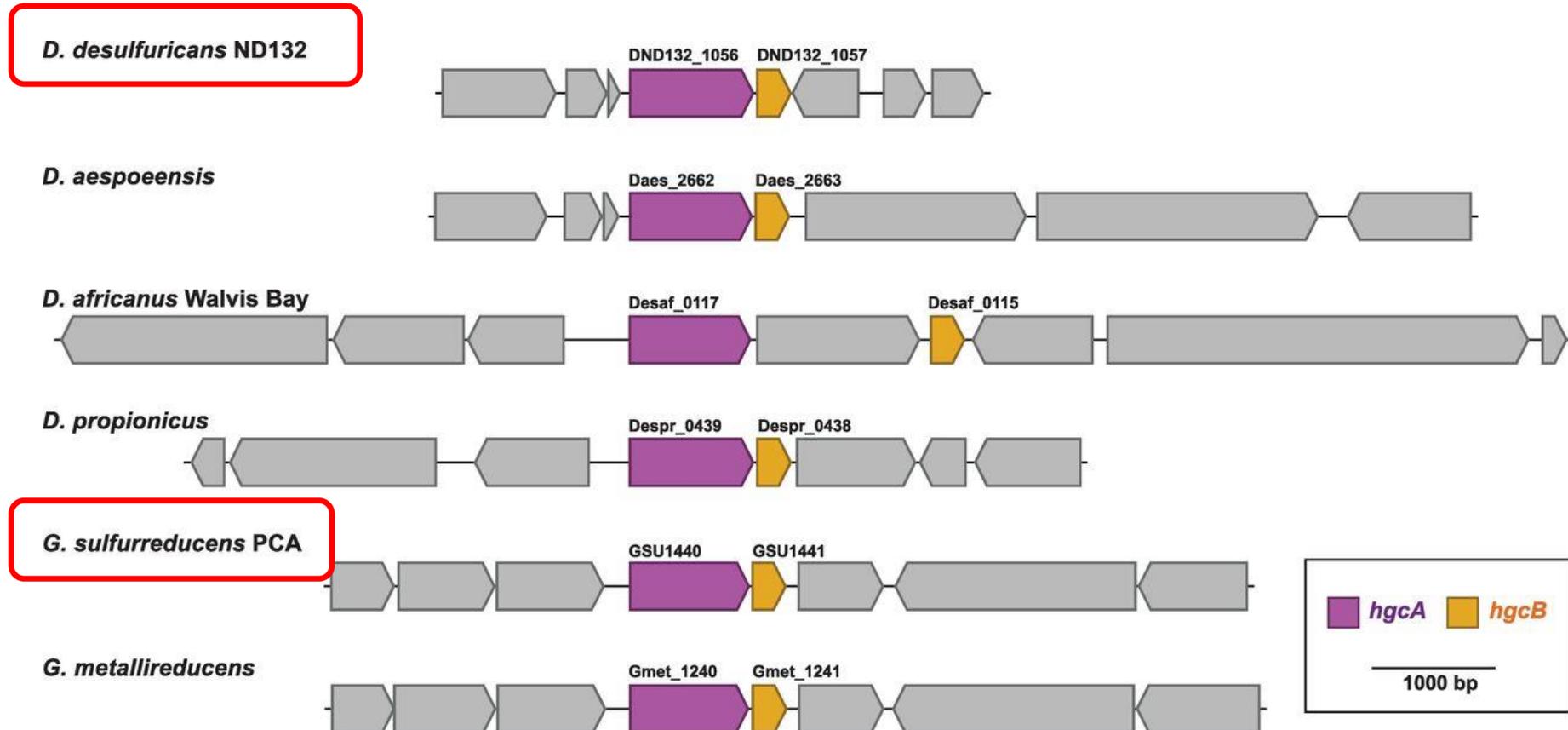
To date, much of what we learn about Hg biogeochemistry and bioaccumulation pathways have come from the studies mostly with the lake ecosystem.



Poulain and Barkay (2013) Science

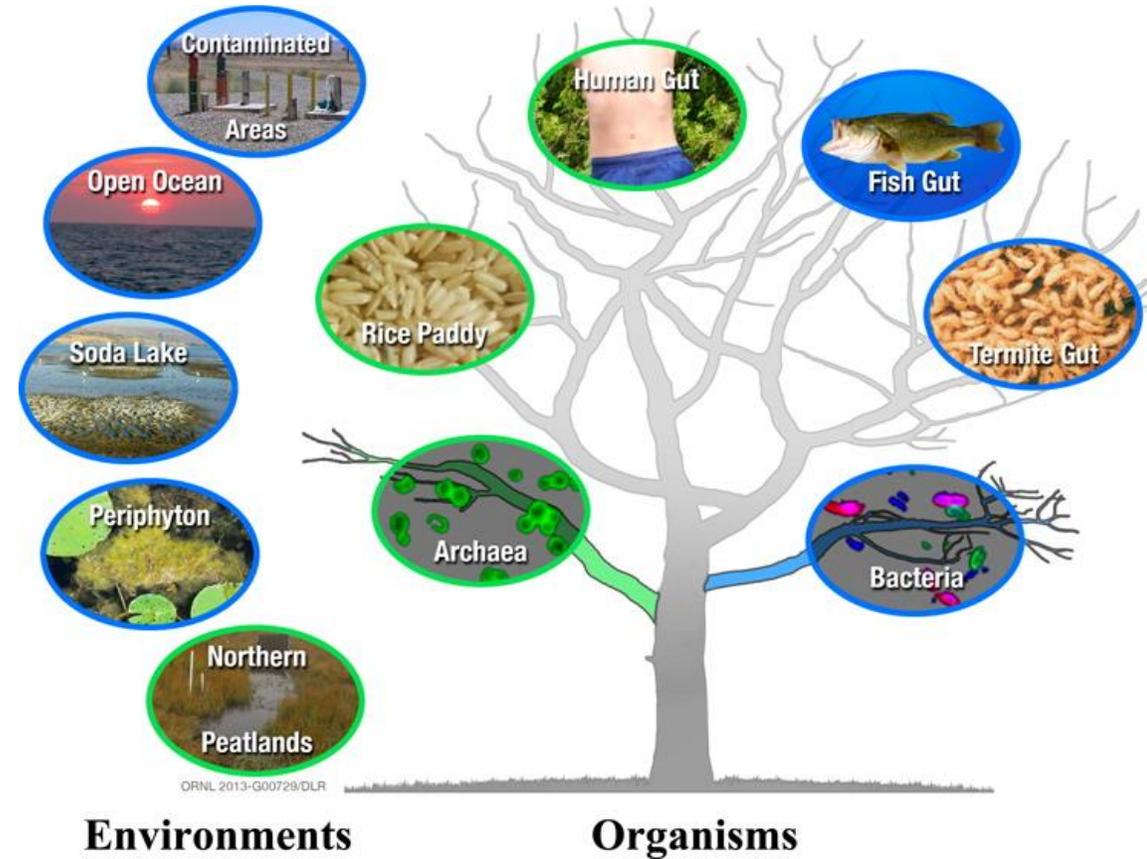
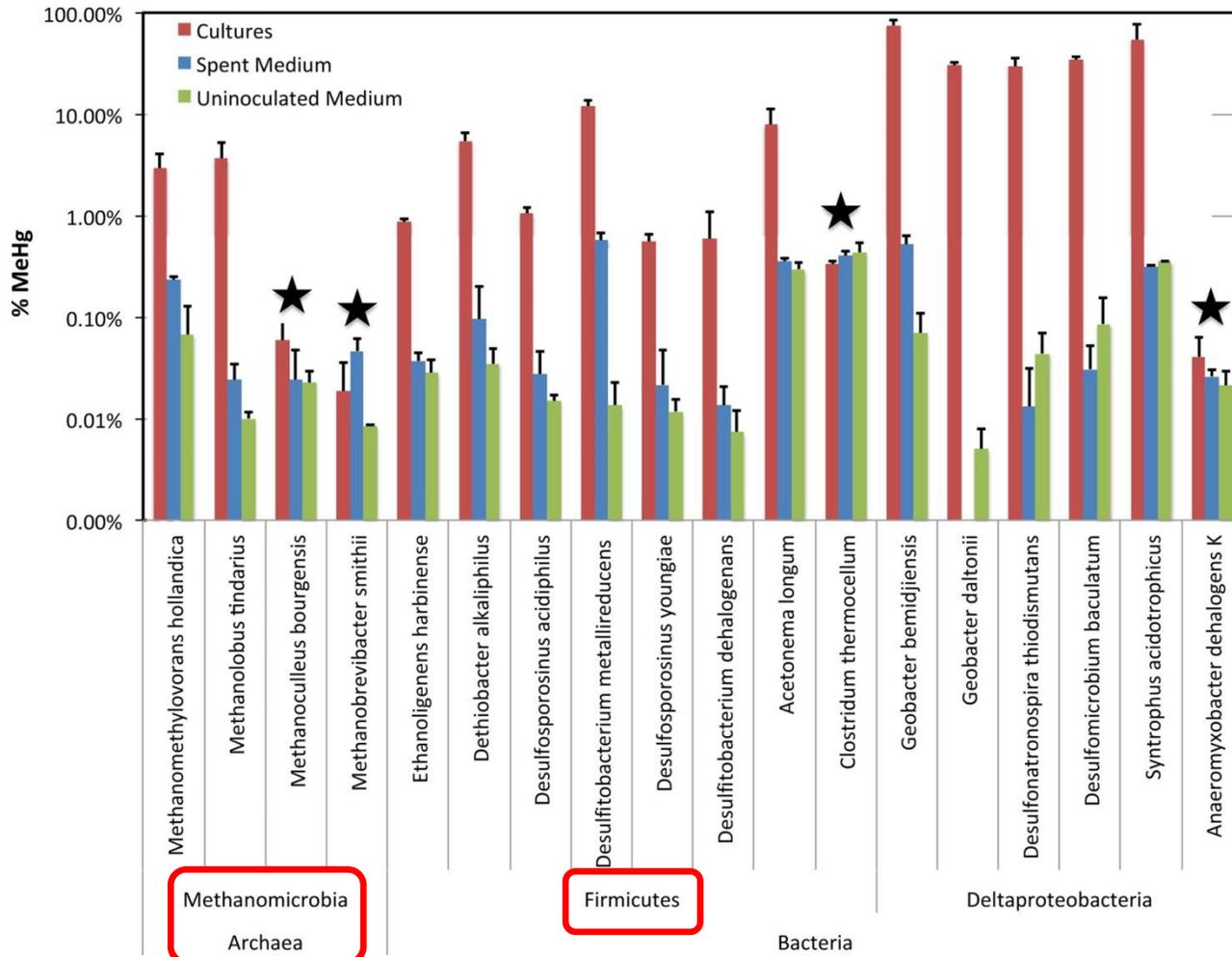
# Putative mercury methylation gene cluster and genomic context

HgcA: a putative methyltransferase corrinoid protein; HgcB: a putative [4Fe-4S] ferredoxin



Parks et al. (2013) Science

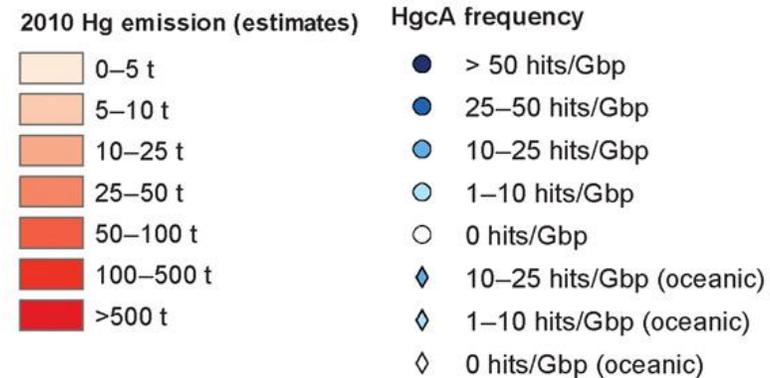
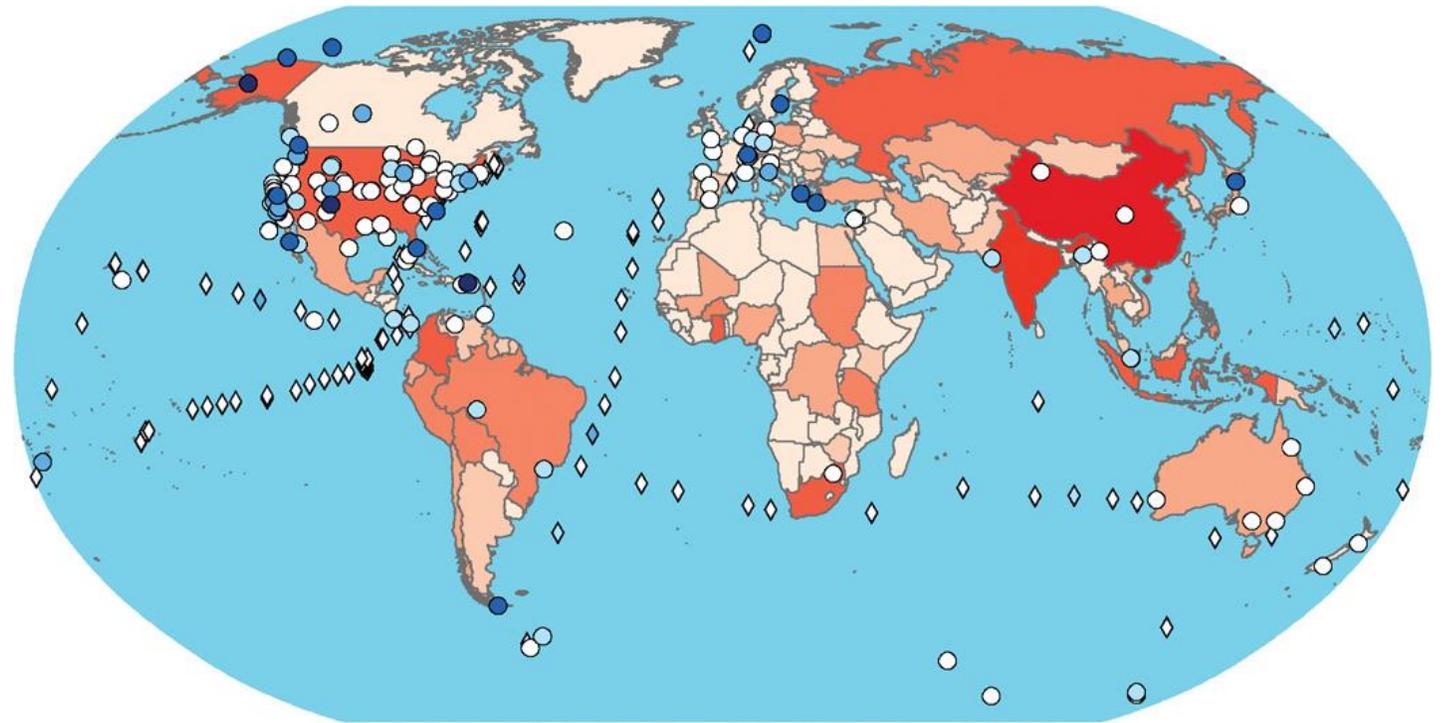
# The Hg-methylating gene cluster, *hgcAB*, is a reliable bio-marker that would be helpful in the development of monitoring and management strategies.



Gilmour et al. (2013) ES&T

The UN Environment Programme recently identified two pressing global issues with regard to mercury pollution (2013):

- (1) establishing the link among deposition, methylation, and uptake by living organisms;
- (2) characterizing methylation and demethylation and how these reactions are affected by climate change.



# **Hg methylation potential in the rice paddy system**

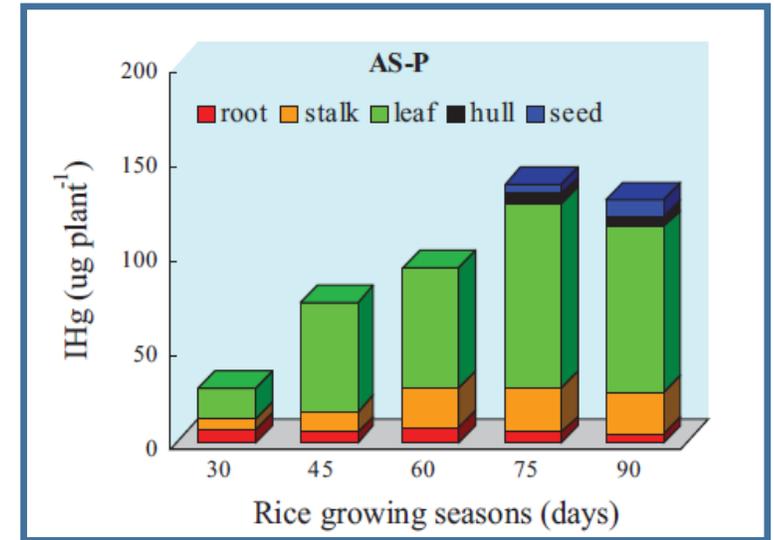
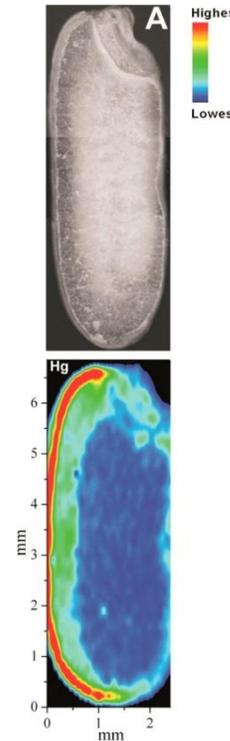
**Hg bioaccumulation in the terrestrial food web has been considered negligible. However, in inland mining areas of China:**

- Hair Hg levels significantly correlate with rice MeHg intake
- Other crops have 10-100 fold lower MeHg in the edible portion
- Rice seeds have the highest capacity to accumulate MeHg
- Accumulation pathways of IHg and MeHg in rice are different

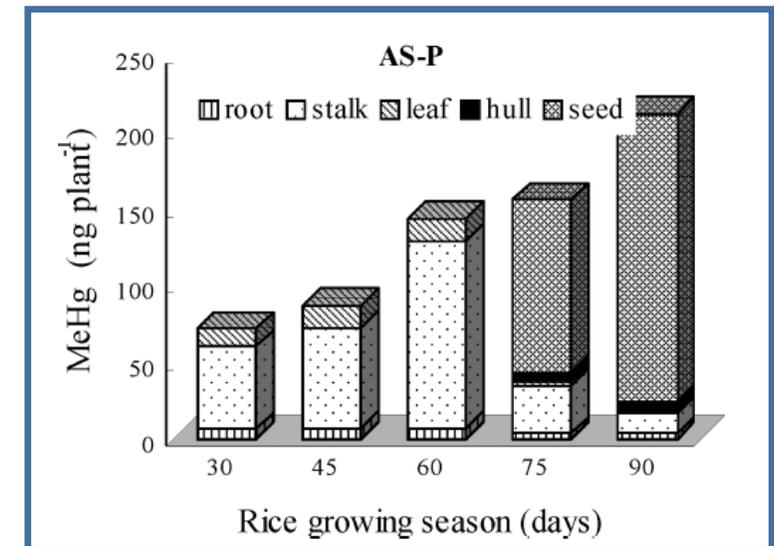
**Table 1. Concentrations of IHg and MeHg in Fractions of Rice Grain from Different Sampling Sites (Mean ± S.D,  $\mu\text{g kg}^{-1}$ )**

sampling sites	fractions	Hg concentration		
		IHg	MeHg	MeHg/THg(%)
Control Site (n = 10)	hull	6.2 ± 0.51	0.26 ± 0.05	4.1 ± 0.73
	bran	24 ± 1.4	3.0 ± 0.48	11 ± 1.8
	white rice	2.7 ± 0.25	2.4 ± 0.16	48 ± 3.3

Meng et al.(2014) ES&T 2014



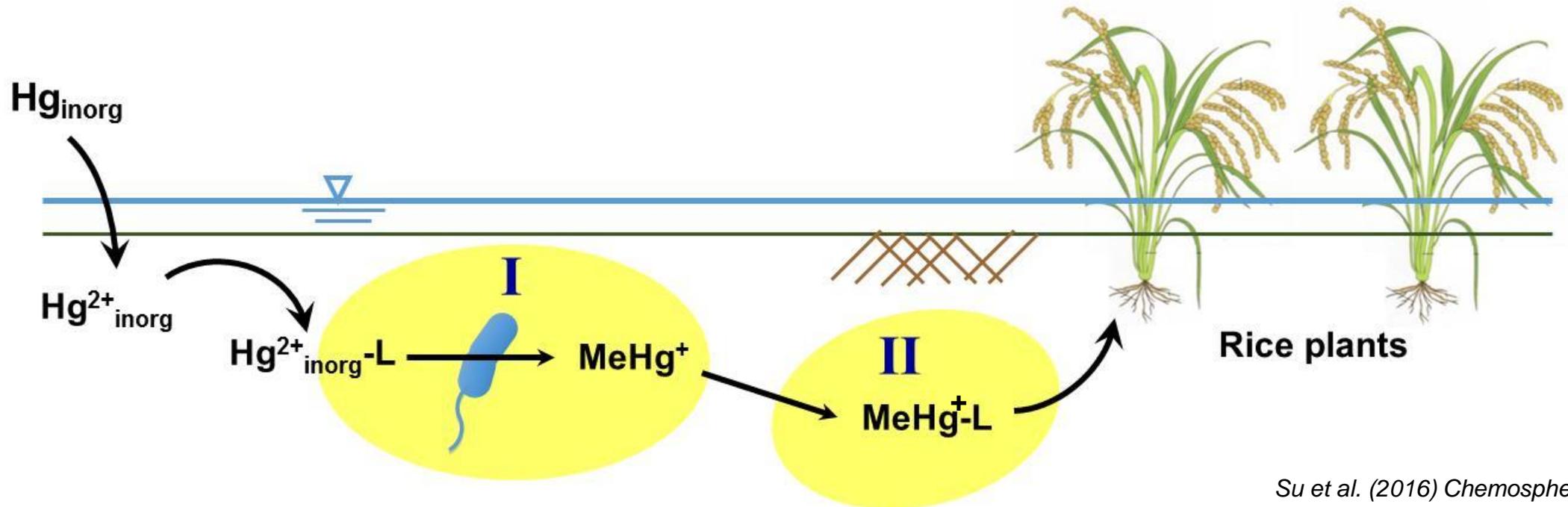
Meng et al. (2012) Environ. Toxicol. Chem.



Meng et al.(2011) ES&T 2011

Given that we do not yet have a complete picture of the mechanisms that underlie the formation, uptake and accumulation of MeHg in the paddy ecosystem, in this study we sought to examine and explore:

- why rice paddies are conducive for Hg methylation?
- what are the major biogeochemical factors involved in this process?
- who are the primary *in situ* Hg-methylators in the paddy rhizosphere?
- what is the role of porewater coordination chemistry in MeHg uptake by rice roots?



Approaches: to answer these questions, we conducted field campaigns over a rice growing season at

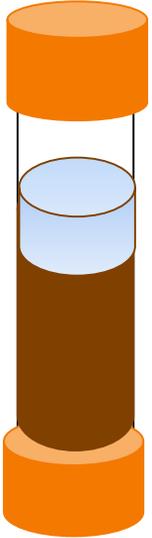
### The Beitou Municipal Solid Waste Incinerator (2013)



### The Taichung Coal-Fired Power Station (2014)

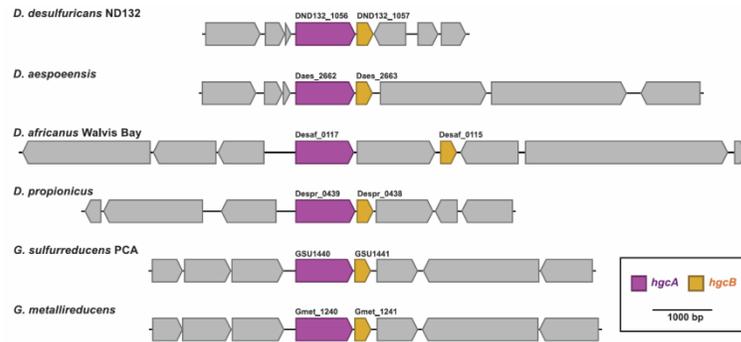


# Approaches: we carefully processed the field samples using strict anaerobic techniques in the lab



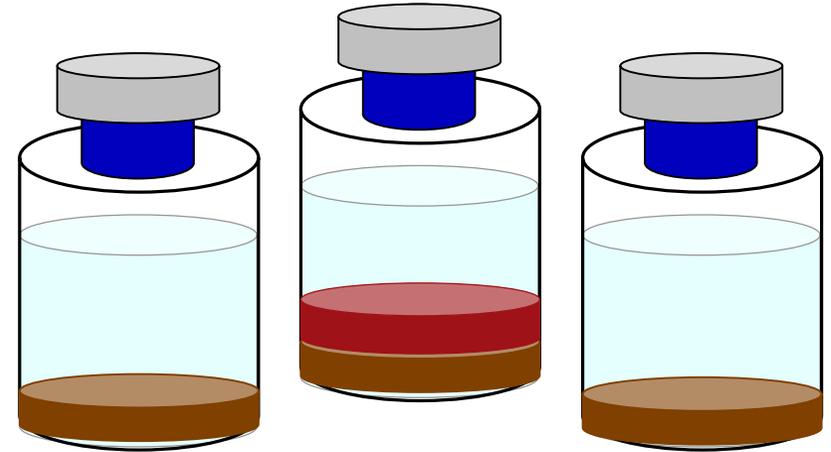
Took sediment cores while maintaining the redox status

Extracted **porewater** and quantified the concentrations of total Hg, MeHg, and ancillary geochemical parameters (iron, sulfur, organic carbon, pH...)



*Parks et al. (2013) Science*

Other **soil samples**: in addition to the exactly the same geochemical parameters, we also designed the primers targeting the *hgcA* gene



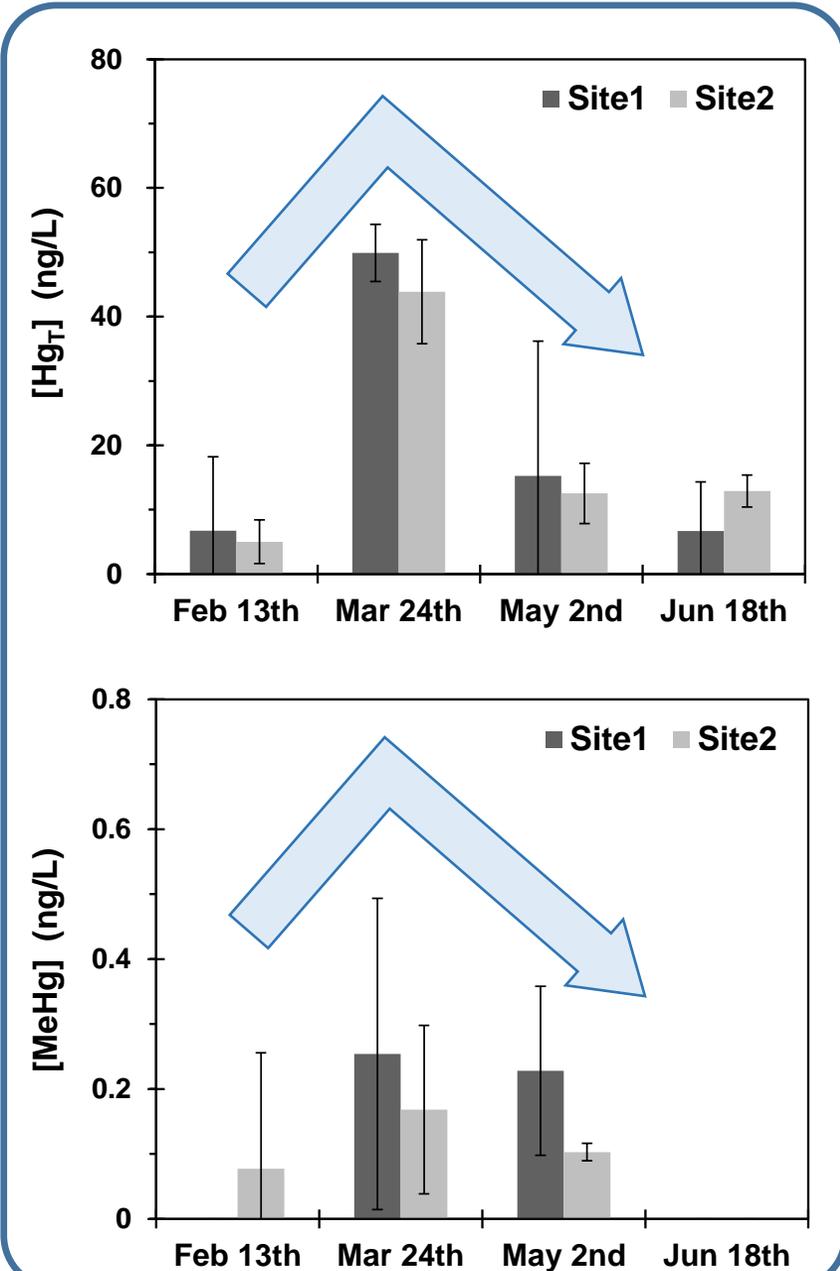
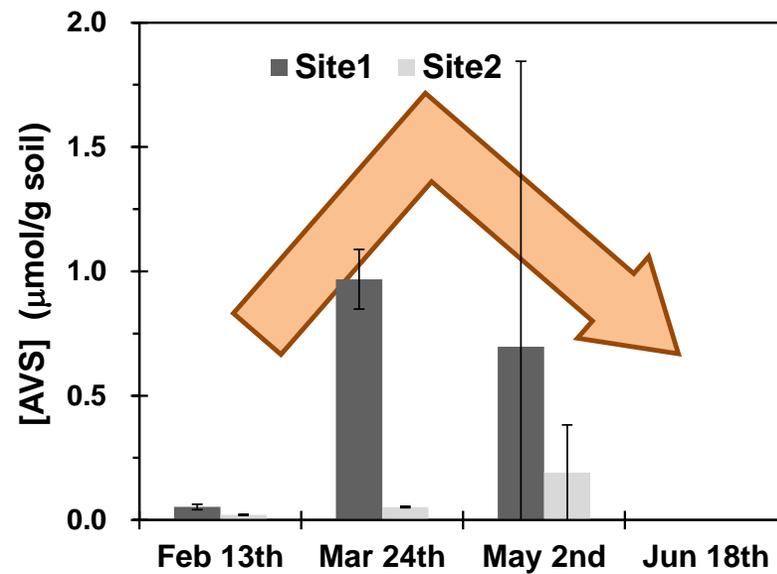
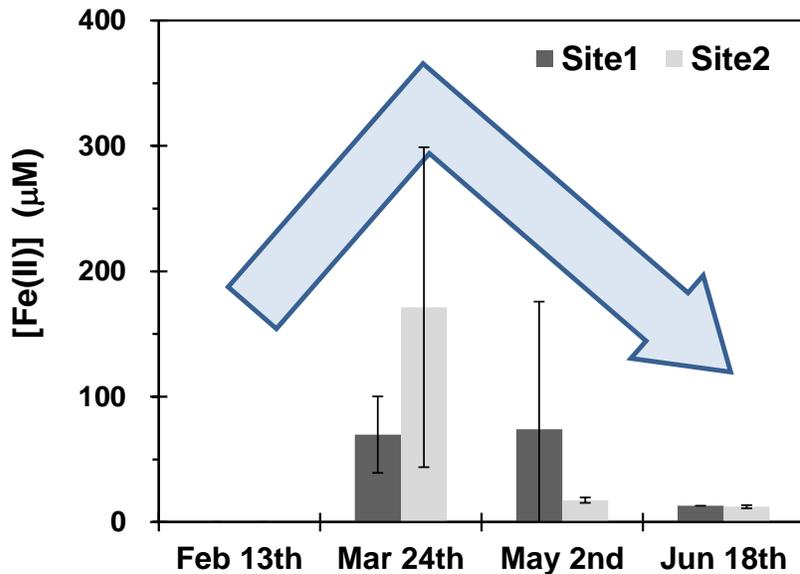
Set up incubation microcosm tests to

- Stimulate/inhibit SRB:  $\text{SO}_4^{2-}$ /molybdate
- Stimulate FeRB: ferrihydrite
- Stimulate/inhibit MPA:  $\text{H}_2 + \text{CO}_2$ /BESA

We also conducted hydroponic experiments by cultivating rice in a defined nutrient solution amended with fixed MeHg and varying levels of MeHg-binding ligands

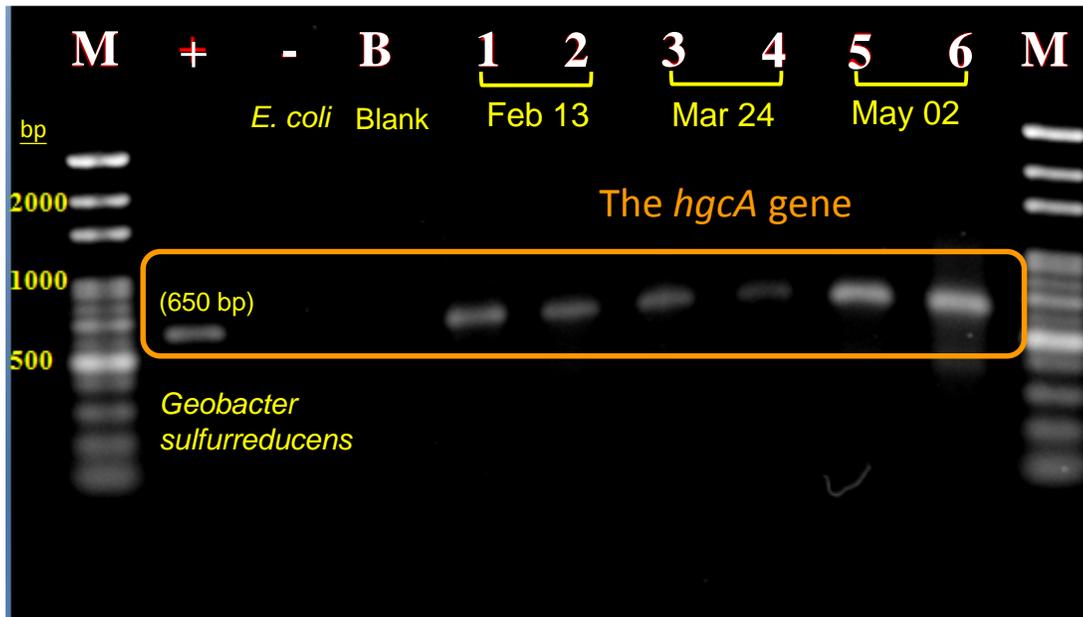
## We observed Hg cycling associated with rhizosphere biogeochemical dynamics in the paddy:

- Total Hg and MeHg levels in paddy soil and rice grains did not exceed the control standards for farmland soil and edible rice in Taiwan.
- *In situ* bioavailability of inorganic Hg and activity of Hg-methylating microbes in the rhizosphere increased from the early-season and peaked at the mid-season, presumably due to the anoxia created under flooded conditions and root exudation of organic compounds.



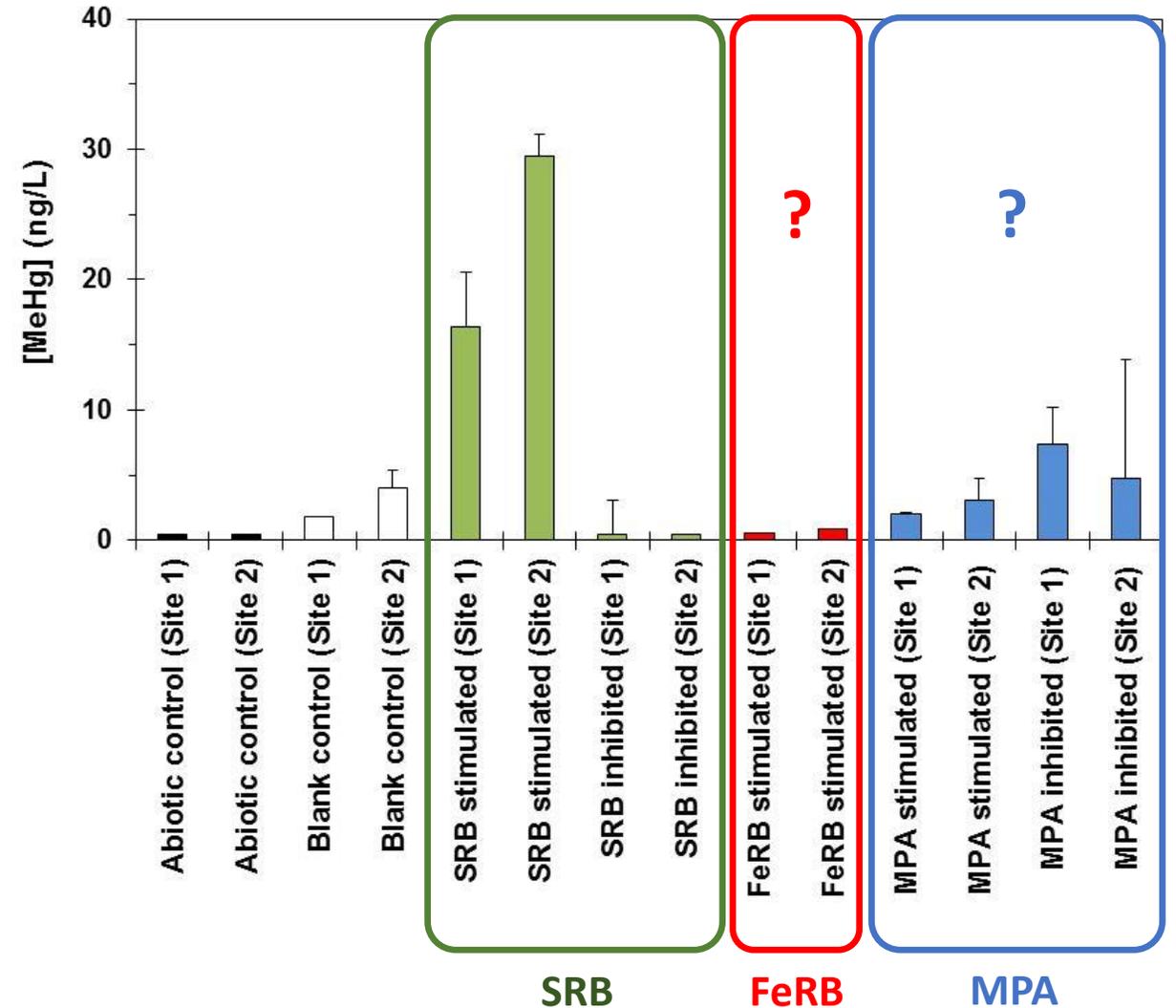
In addition, we identified **the potential primary *in situ* Hg-methylators in the paddy rhizosphere:**

- The presence of Hg-methylators was also confirmed by the detection of the bacterial Hg-methylating gene, *hgcA*, in all root soils.
- Microcosm incubation tests revealed that sulfate reducers might have been the primary Hg-methylating guild at our study sites.



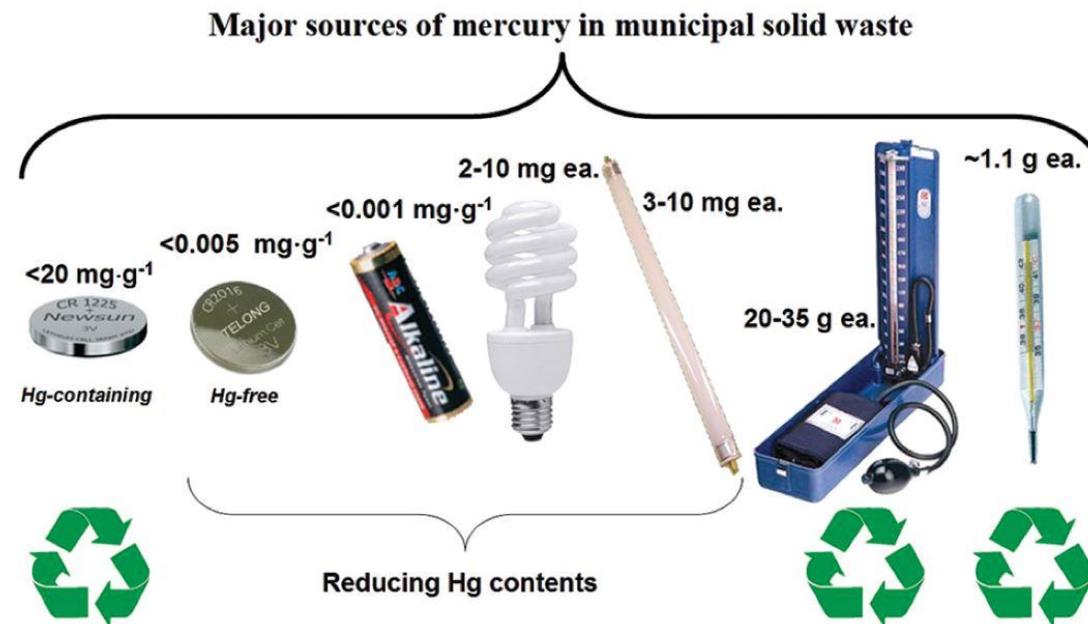
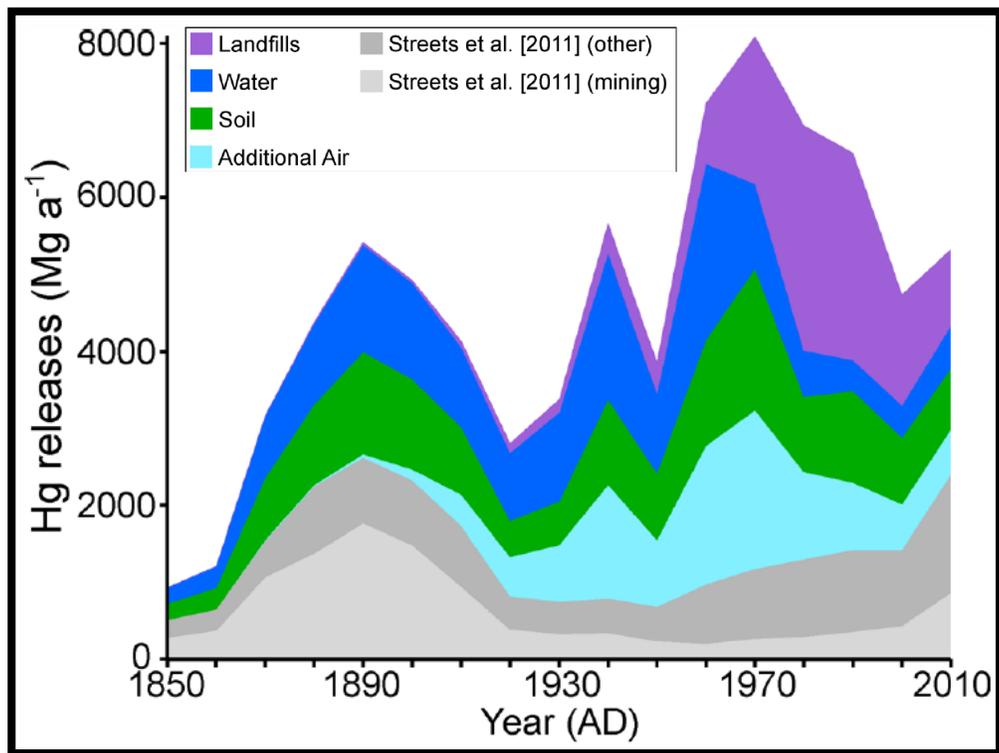
Su et al.(2016) Chemosphere

- **Should there be a more significant role for FeRB and MPA to play in MeHg production in paddies?**



# **Hg methylation potential in the landfill system**

# Hg-containing products may be disposed of at landfills

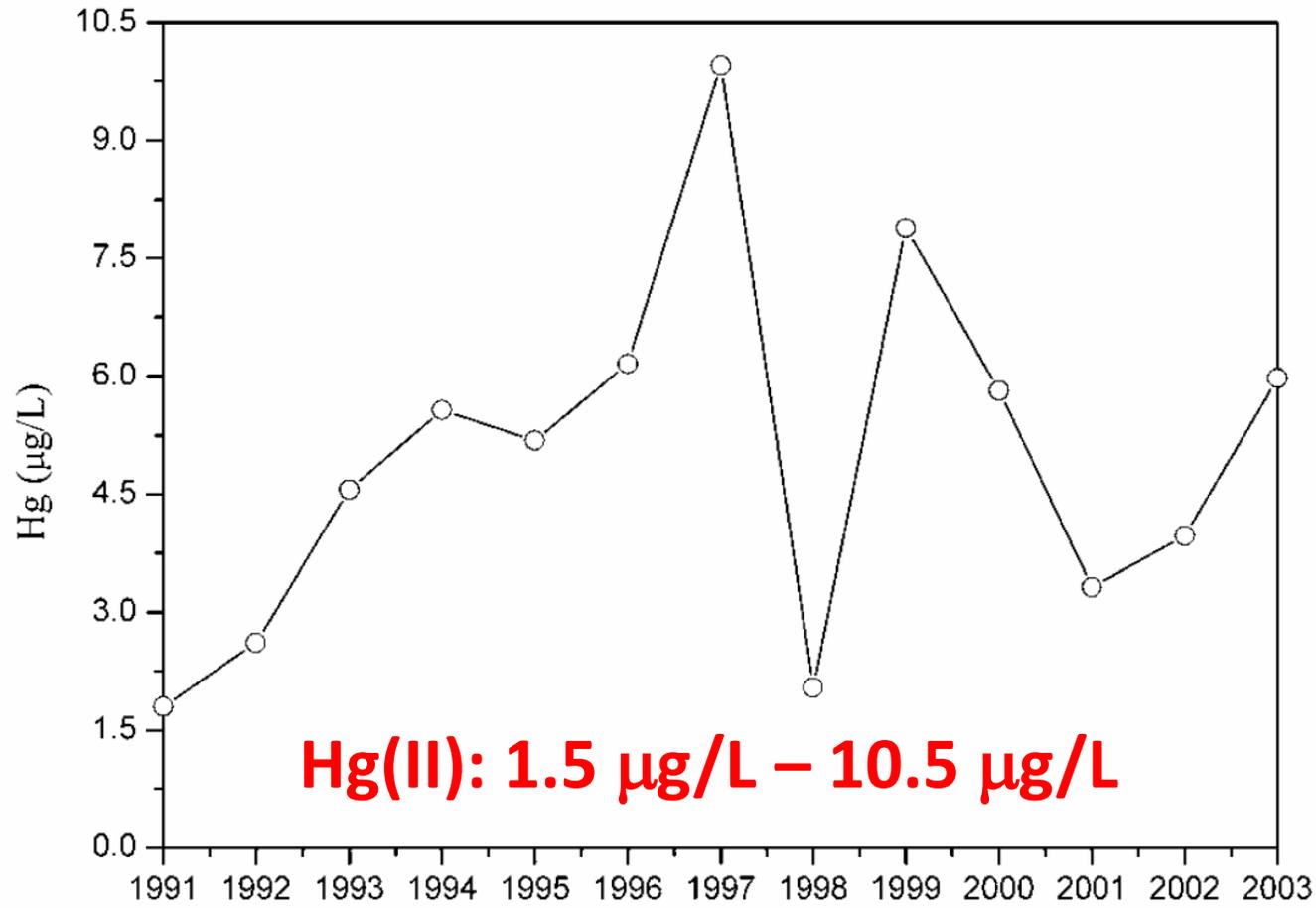


Cheng et al. (2012) ES&T

reservoirs	Hg (Gg)
Mining (air)	215
air, water, soil	310
Landfill	230
<b>total</b>	<b>755</b>

# Levels of mercury detected in landfill leachates

Environmental level ~ 1-500 ng/L



**Fig. 1** The mercury concentration of landfill leachate (the discharge standard of Hg in landfill leachate is 0.001 mg l<sup>-1</sup>).

Xiaoli et al. (2011) J. Environ. Monit.

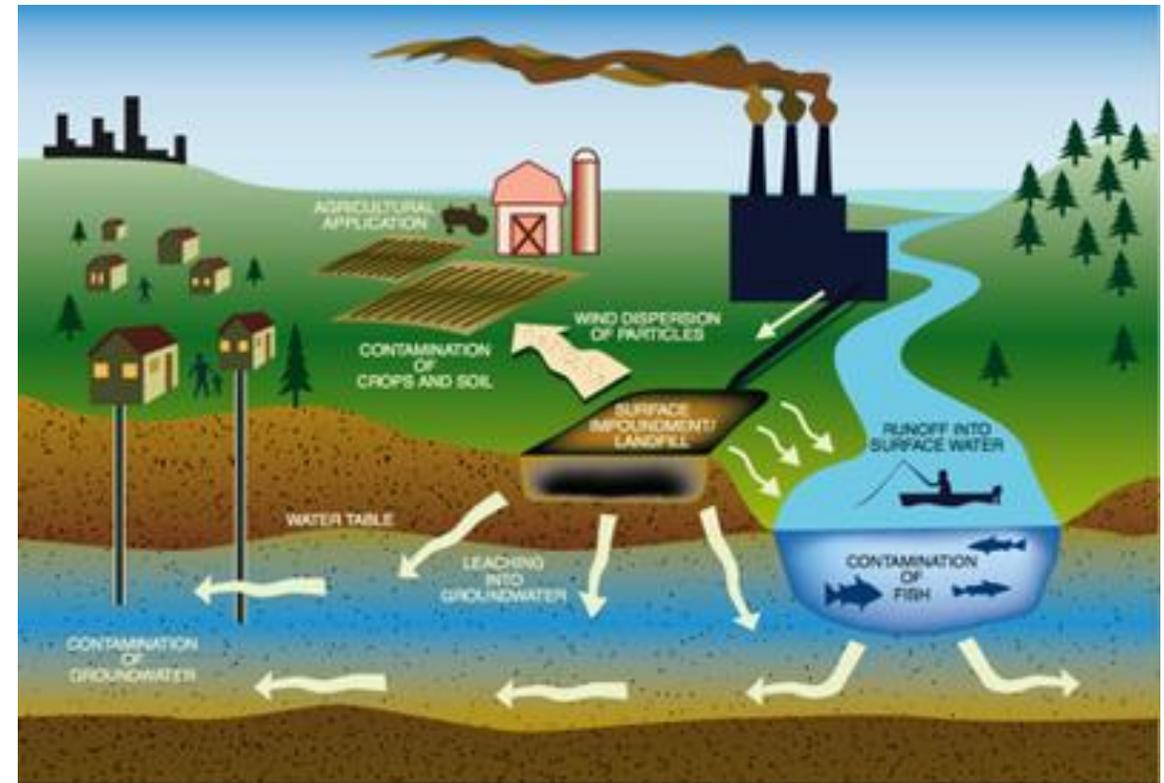
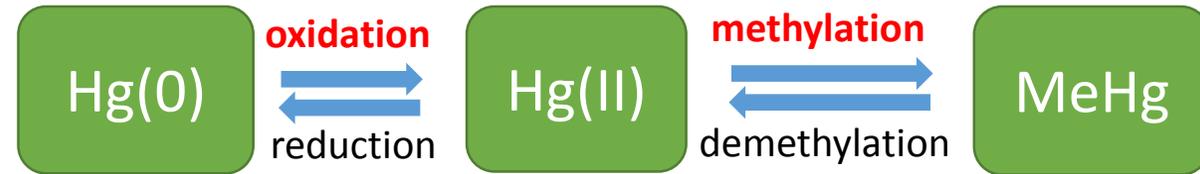
Parameter	Range
pH	4.5–9
Specific conductivity (µS cm <sup>-1</sup> )	2500–35 000
Total solids	2000–60 000
<i>Organic matter</i>	
Total organic carbon (TOC)	30–29 000
Biological oxygen demand (BOD <sub>5</sub> )	20–57 000
Chemical oxygen demand (COD)	140–152 000
BOD <sub>5</sub> /COD (ratio)	0.02–0.80
Organic nitrogen	14–2500
<i>Inorganic macrocomponents</i>	
Total phosphorous	0.1–23
Chloride	150–4500
Sulphate	8–7750
Hydrogencarbonate	610–7320
Sodium	70–7700
Potassium	50–3700
Ammonium-N	50–2200
Calcium	10–7200
Magnesium	30–15 000

**Hg(II): 50 ng/L - 160 µg/L**

<i>Inorganic trace elements</i>	
Arsenic	0.01–1
Cadmium	0.0001–0.4
Chromium	0.02–1.5
Cobalt	0.005–1.5
Copper	0.005–10
Lead	0.001–5
<b>Mercury</b>	<b>0.00005–0.16</b>
Nickel	0.015–13
Zinc	0.03–1000

# Mercury Transformations in Landfill Sites

- Recent available data prompt a need to re-examine the landfill environment as a potential hot-spot for MeHg production, given that **the mechanisms underlying Hg transformations in this system have not been studied.**
- To approach and address this issue, the following factors/processes have to be determined:
  - existence of the *hgcAB* gene pair
  - existence of a  $\text{Hg(II)}_{(\text{aq})}$  pool
  - bioavailability of  $\text{Hg(II)}_{(\text{aq})}$
  - net MeHg production
- Our hypothesis: the extent of MeHg formation in this system is still a function of both the activity of Hg-methylating bacteria and  $\text{Hg(II)}$ -bioavailability.



Source : [catawbariverkeeper.org](http://catawbariverkeeper.org)

Site C



# Geochemical conditions

Site A



Site B



Site C



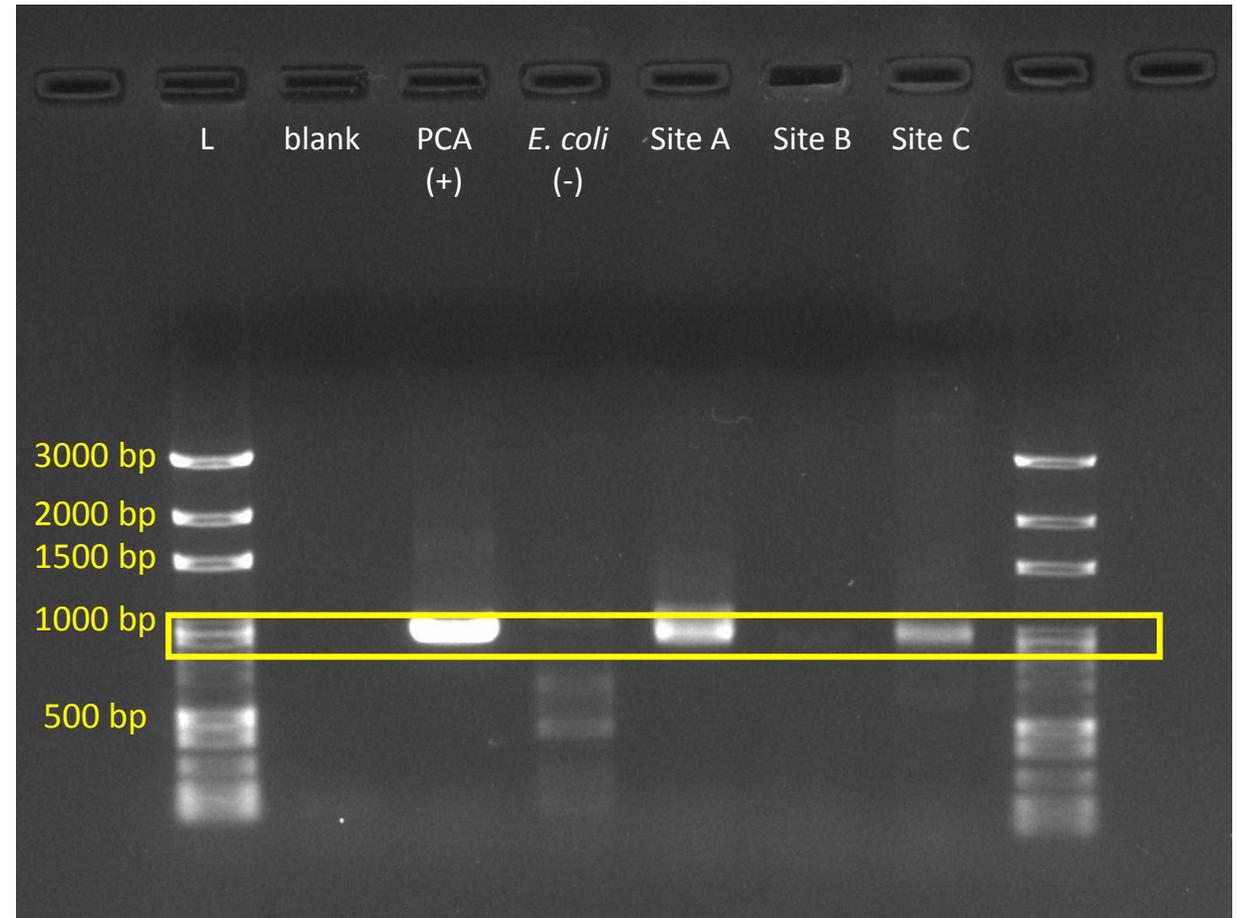
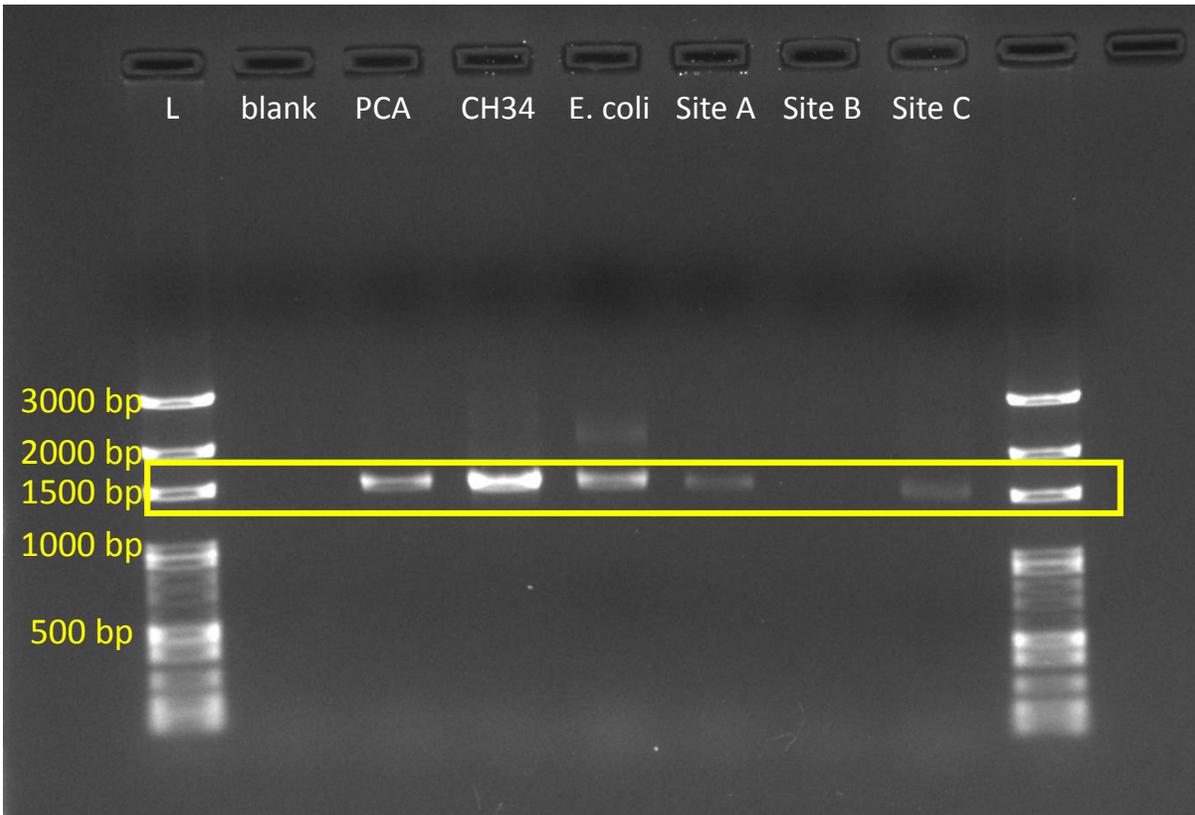
	Site A	Site B	Site C
Temp °C	25.6	24.7	20.8 - 27.3
pH	7.32	7.62	6.96 - 7.06
EC μS/cm	5652 - 7612	3054 - 4132	8634 - 14789
DO mg/L	0.07 - 5.18	0.10 - 0.23	0.16 - 0.73
ORP mV	205 - 220	207 - 219	23.4 - 221
COD mg/L as COD	357	147 - 247	453 - 988
TOC mg/L as C	81-116	35 - 154	109 - 153
Sulfate mg/L as SO <sub>4</sub> <sup>2-</sup>	2.08	< 2.00	< 2.00
Sulfide μg/L as S <sup>2-</sup>	36	20 - 141	54 - 65
Nitrate mg/L as NO <sub>3</sub> <sup>-</sup> -N	0.39	7.07	6.80
Nitrite mg/L as NO <sub>2</sub> <sup>-</sup> -N	0.10	1.74	5.35
TFe mM	< 10	< 10	< 10

# THg & MeHg



		Site A	Site B	Site C
total Hg	pM	68.7 (9.70 - 249)	82.7 (8.72 - 259)	245.8 (39.1 - 483)
dissolved Hg (% in THg)	pM	12.5 (6.00 - 20.1) 18.2%	14.4 (5.29 - 54.4) 17.5%	92.4 (88.3 - 96.7) 23.31%
non-purgeable Hg (% in THg)	pM	18.0 (12.0 - 26.2) 96.6%	80.0 (75.3 - 86.4) 69.0%	266.6 (168 - 336) 70.4%
MeHg (% in THg)	pM	1.07 ± 0.102 1.56%	1.07 ± 0.11 1.30%	0.53 ± 0.11 0.22%

# Detection of the *hgcAB* gene cluster



Blank: DDW as sample

PCA: *Geobacter sulfurreducens* PCA

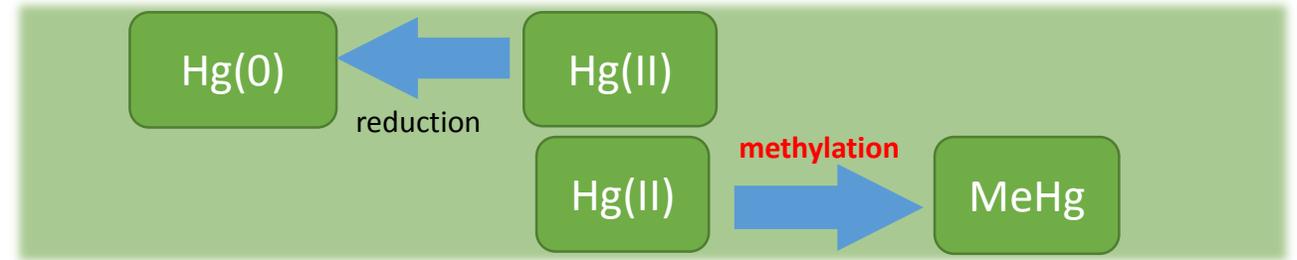
CH34: *Cupriavidus metallidurans* CH34

*E. coli*: *Escherichia coli*

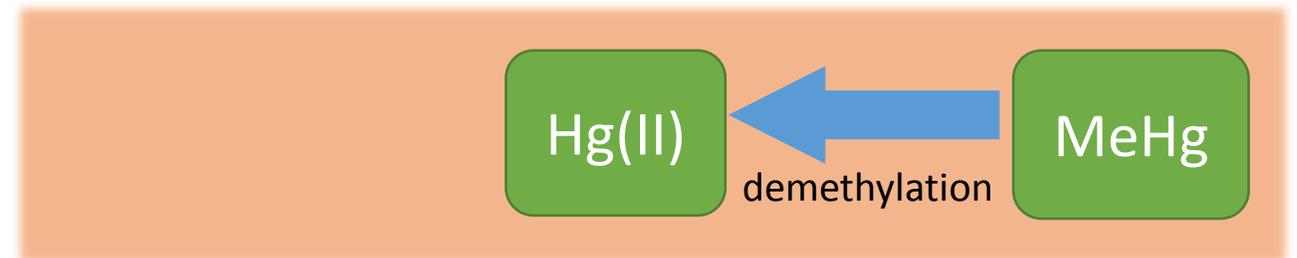
# Microcosm tests



in the dark under anaerobic conditions  
at static, room temperature

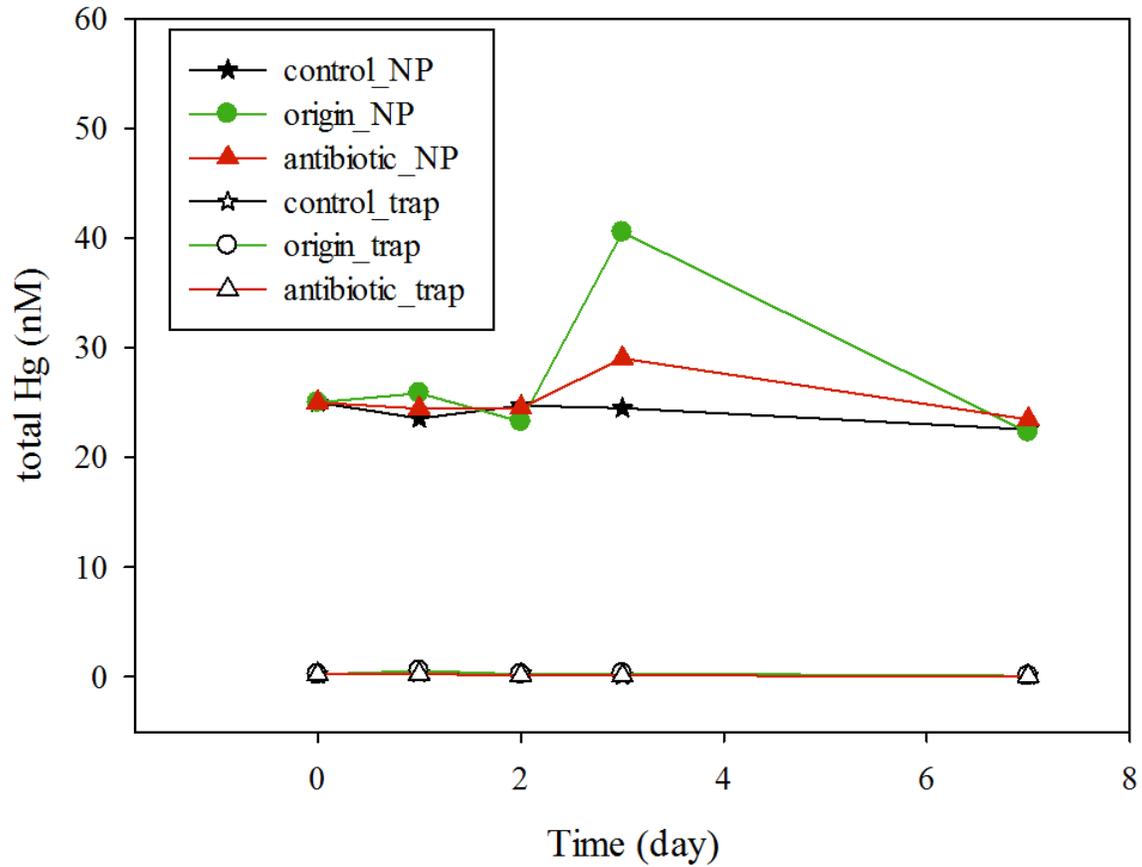


- Hg(II) reduction
- Hg(II) methylation (Potential)
- Hg(II) methylation (Energy)
- Hg(II) methylation (Bioavailability): PCA

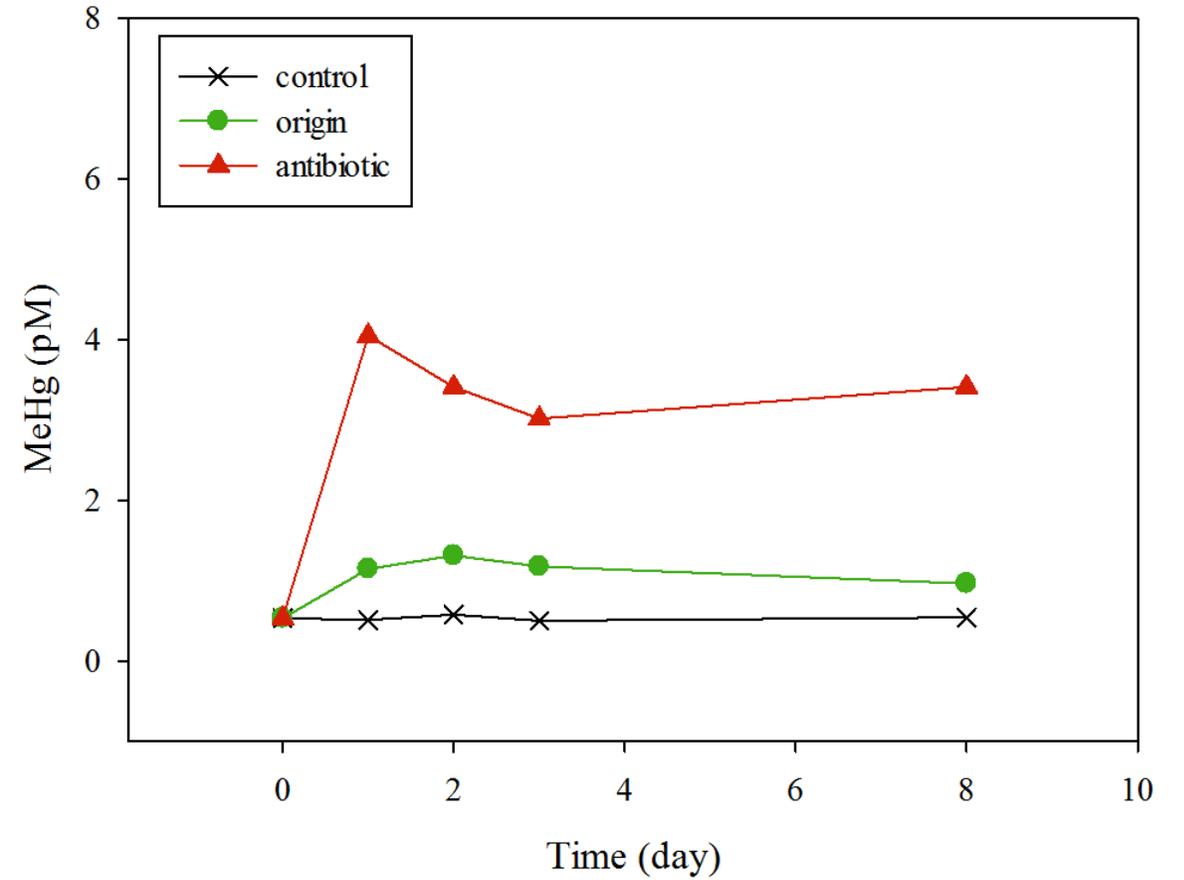


- MeHg demethylation

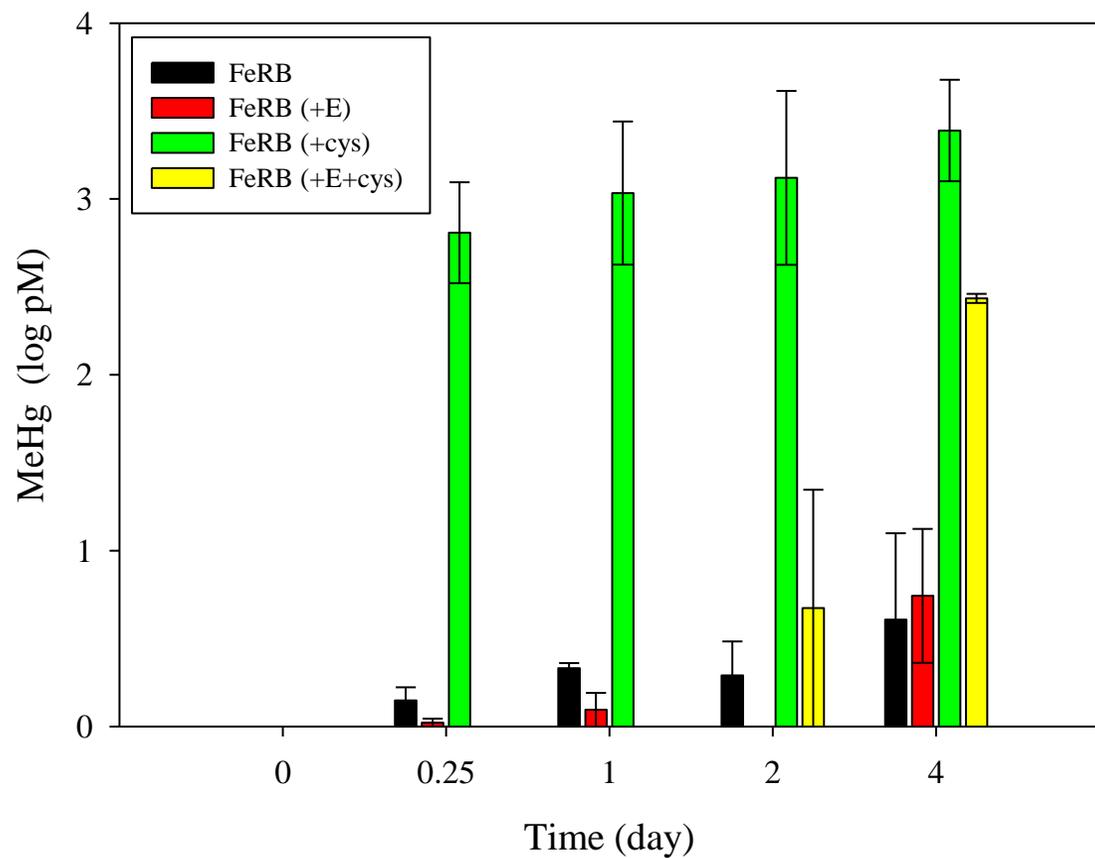
### Hg(II) reduction is not significant



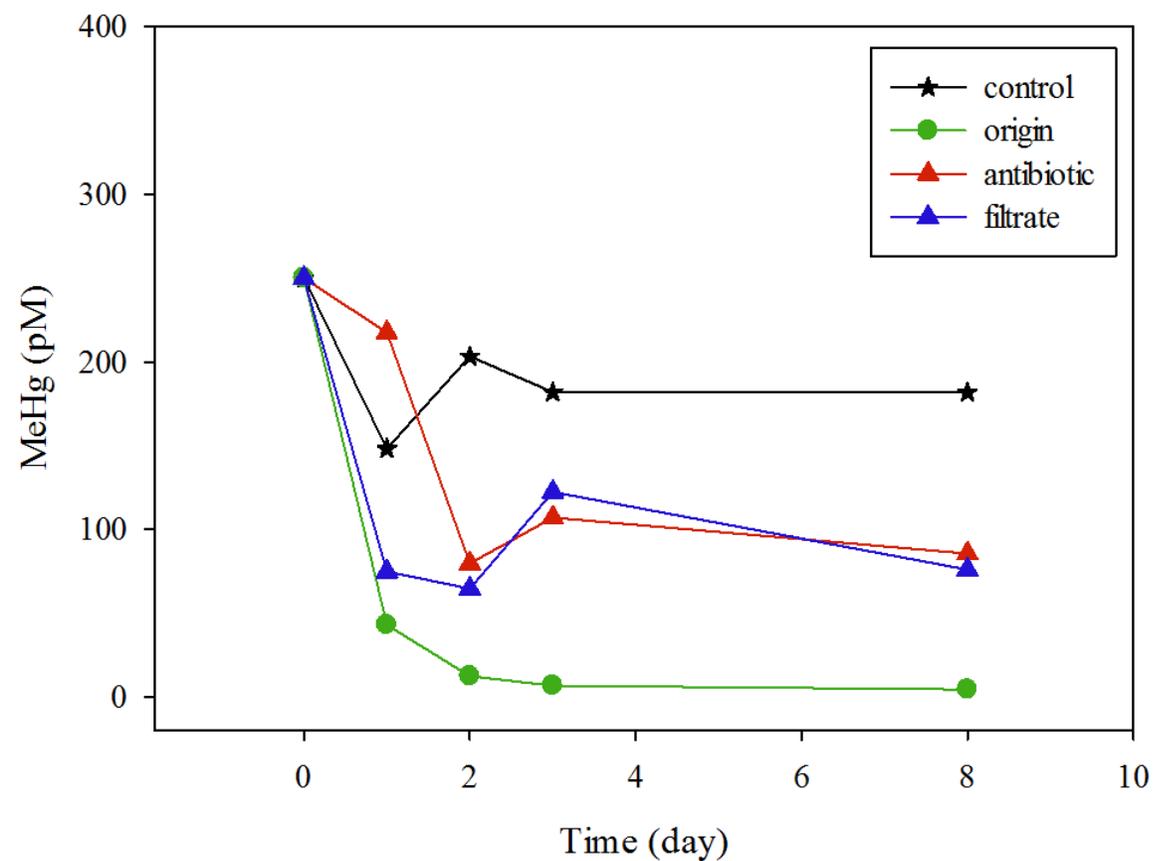
### Abiotic Hg methylation is dominant



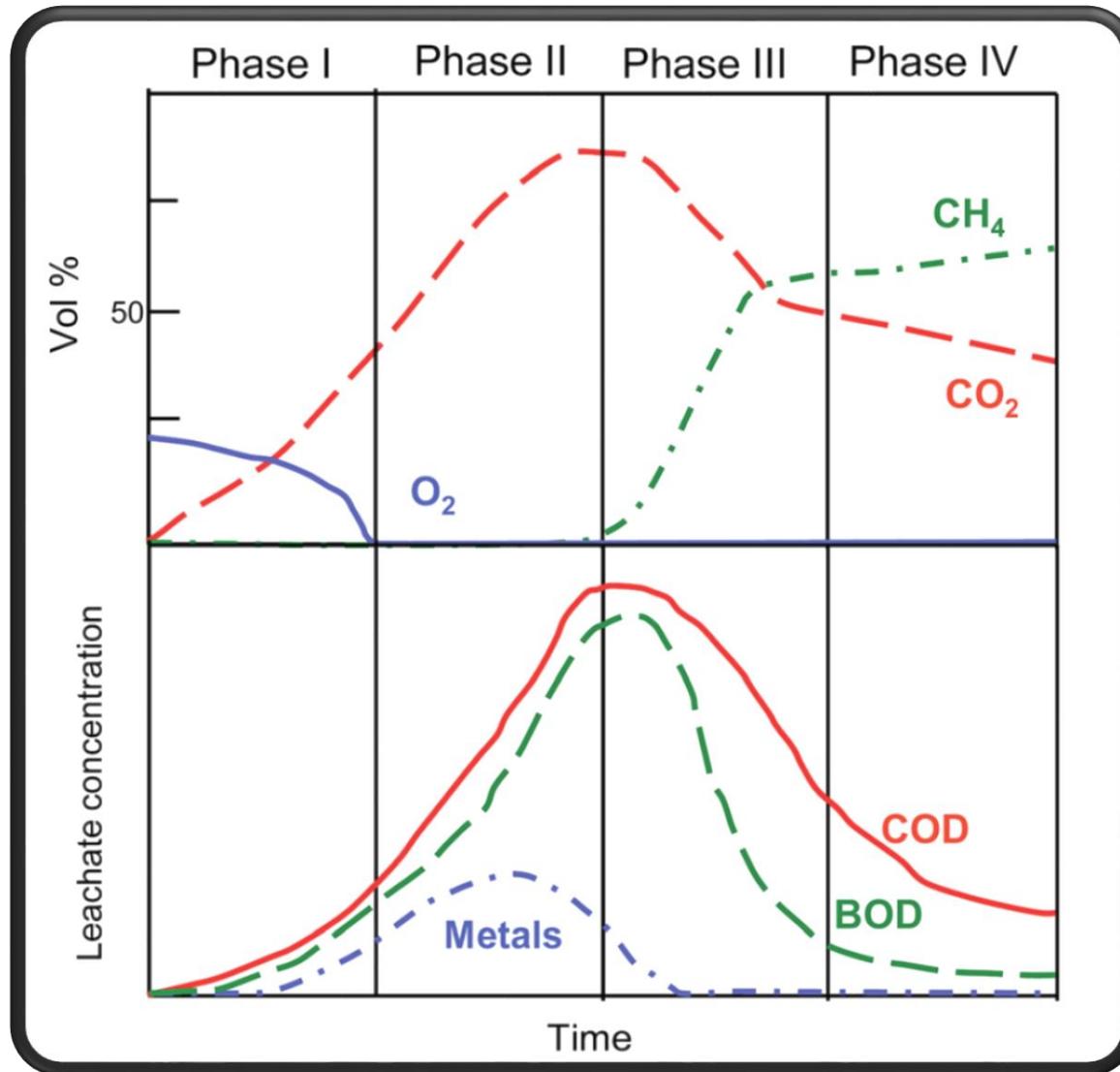
## Dissolved Hg(II) is available to microbes



## Significantly indigenous MeHg degradation



# Landfill maturation process



# Summary

- Paddy rhizosphere has higher Hg methylation potential during the rice mid-growing season.
- The role of methanogens in MeHg production in the paddy ecosystem deserves further investigations.
- Hg methylation is not significant in the landfill leachate of the Phase IV and later maturation processes.
- A solid understanding of the mechanisms that underpin the important environmental processes may eventually lead to developments of better ecological assessments and more sound remedial actions.

# Thank you for your attention!

Special thanks to  
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**Yen-Bin Su**  
(宿彦彬)



**Wei-Chun Chang**  
(張惟竣)



**Chih-Kuen Hsu**  
(徐志昆)