

WILLIAMS, ASHLEY S., M.S. Detecting a Microbial Response in Sediment of the Dan River Following a Coal Ash Spill. (2019)
Directed by Dr. Parke Rublee. 112 pp.

Coal ash is the residual material of coal combustion for electricity generation. It contains heavy metals and other pollutants and is generally deposited into reservoir ponds for storage, although it may spill or leach into nearby water and possibly disrupt aquatic communities. In February of 2014, a coal ash spill occurred in the Dan River in Eden, NC. Coal ash contains constituents that may stimulate the growth of mercury methylating bacteria. This study aimed to determine if a microbial response is detectable 1.5 years following the spill using qPCR. We tested three primers targeting mercury methylation. We detected an elevated signal 0.5 km downstream from the spill site relative to some other upstream and downstream locations. However, the highest abundance of amplified targets was observed in the furthest upstream site. We also undertook a survey of bacteria present in a coal ash sample from the coal ash pond that was the source of the spill, located at the retired Dan River Steam Station in Eden, NC. SSU rDNA extracted from 31 isolated organisms was sequenced and the organisms identified to genus level. The community was predominantly composed of *Bacillus* and *Arthrobacter* spp. 14 Isolates were grown in 50% nutrient broth amended with heavy metals commonly found in coal ash waste (As, Cd, Cr, Hg, Pb, Se, Zn) at environmentally relevant concentrations to characterize their metal tolerance.

Overall, 1) we could not confirm a spike in our mercury markers, and 2) coal ash isolates exhibited metal tolerance.

DETECTING A MICROBIAL RESPONSE IN SEDIMENT OF THE DAN RIVER
FOLLOWING A COAL ASH SPILL

by

Ashley S. Williams

A Thesis Submitted to
the Faculty of The Graduate School at
The University of North Carolina at Greensboro
in Partial Fulfillment
of the Requirements for the Degree
Master of Science

Greensboro
2019

Approved by

Committee Chair

To my kids and my family.

APPROVAL PAGE

This thesis written by Ashley S. Williams has been approved by the following committee of the Faculty of The Graduate School at The University of North Carolina at Greensboro.

Committee Chair _____
Parke Rublee

Committee Members _____
Anne Hershey

Martin Tsui

Date of Acceptance by Committee

Date of Final Oral Examination

ACKNOWLEDGMENTS

I sincerely appreciate the continuous encouragement and guidance from my advisor and mentor Dr. Parke Rublee, and from my committee members, Drs. Anne Hershey and Martin Tsui. I thank Brian Williams of the Dan River Basin Association for his help and guidance on the river. I also thank Kimber Corson, Matthew Monteverde, and Peija Ku for their assistance during field sampling, as well as Jacob Cleary and our lab mates for their assistance in the lab, including many late night assay reads. A very special thanks to Louisa Liberman for her editorial support and, along with John Rice, Jennifer Petite, and Leah Blasiak for their copious amounts of encouragement and support. Duke Energy provided access to their boat launch at the spill site. This work was funded by a North Carolina Water Resources Research Institute grant and the UNCG Department of Biology.

TABLE OF CONTENTS

	Page
LIST OF TABLES.....	vi
LIST OF FIGURES.....	viii
CHAPTER	
I. INTRODUCTION	1
II. RESPONSE OF MERCURY METHYLATING BACTERIA TO THE COAL ASH SPILL IN THE DAN RIVER	4
Introduction	4
Methods.....	7
Results	12
Discussion	19
III. IDENTIFICATION AND CHARACTERIZATION OF HEAVY METAL TOLERANCE OF BACTERIA ISOLATED FROM COAL ASH	21
Introduction	21
Methods.....	24
Results	27
Discussion	35
IV. CONCLUSIONS.....	37
REFERENCES	38
APPENDIX A. SUPPLEMENTAL TABLES	44

LIST OF TABLES

	Page
Table 2.1. Sampling Locations.	9
Table 2.2. Sediment Sampling Schemes.	10
Table 2.3. Primers Used in this Study.	11
Table 2.4. Comparisons of Each Sampling Location to the Location Directly Downstream of the Spill	16
Table 2.5. Comparisons of Each Sampling Location to the Furthest Upstream Location	16
Table 2.6. Comparisons of Each Sampling Location to the Leaching Site Location	17
Table 3.1. Final Concentrations of Metal in Broth (nM)	25
Table 3.2. P-values of Kruskal-Wallis Test of Carrying Capacity.	34
Table 3.3. P-values of Kruskal-Wallis Test of Growth Rate	35
Table A.1. DR 13 Gompertz Model Parameters.	44
Table A.2. DR 24 Gompertz Model Parameters.	49
Table A.3. DR 26 Gompertz Model Parameters.	53
Table A.4. DR 3.2 Gompertz Model Parameters.	58
Table A.5. DR 5 Gompertz Model Parameters.	63
Table A.6. DR 52 Gompertz Model Parameters.	68
Table A.7. DR 60 Gompertz Model Parameters.	73
Table A.8. DR 76 Gompertz Model Parameters.	78
Table A.9. DR p51 Gompertz Model Parameters.	83
Table A.10. DR p72 Gompertz Model Parameters.	88
Table A.11. DR p74 Gompertz Model Parameters.	93
Table A.12. NO 14 Gompertz Model Parameters.	98

Table A.13. NO 17 Gompertz Model Parameters.....	103
Table A.14. NO 22 Gompertz Model Parameters.....	108

LIST OF FIGURES

	Page
Figure 2.1. Google Earth Image of Dan River and Sampling Locations.....	8
Figure 2.2. <i>hgcA</i> Abundance Normalized to Organic Matter at Each Sampling Location.....	13
Figure 2.3. FeRB Abundance Normalized to Organic Matter at Each Sampling Location.....	14
Figure 2.4. SRB Abundance Normalized to Organic Matter at Each Sampling Location.....	15
Figure 2.5. Target DNA Concentration at Each Depth	18
Figure 2.6. River Channel Abundance at Sampling Sites Upstream and Downstream Normalized to Organic Matter	19
Figure 3.1. Phylogenetic Analysis of Isolates Cultured from Coal Ash Amended Media.....	28
Figure 3.2. Growth of Isolate DR13.....	31
Figure 3.3. Growth of Isolate DR3.2	32
Figure 3.4. Growth of Isolate DR52.....	33

CHAPTER I

INTRODUCTION

Coal is the second most used fuel for electricity generation in the United States. In 2017, approximately 30% of all electricity production was fueled by coal combustion (Energy Information Administration, 2018). Although the percentage has fallen in recent years due to retirement of coal-fired plants and increases in natural gas and other energy sources, coal remains a main fuel for electricity. Coal combustion results in the production of the waste material, coal ash.

Coal ash is composed of fly ash, bottom ash, and flue gas desulfurized gypsum. Fly ash is a fine powdery substance, comprised primarily of silica that is exhausted through the smokestack. It is produced during the combustion of finely ground coal and most is captured from the exhaust process using electrostatics and scrubber systems. Bottom ash is formed during the combustion of pulverized coal in boilers. It ranges in size from fine sand to fine gravel and is grey to black in color. Bottom ash is too large to be carried up the exhaust system and is collected in an ash hopper. Flue gas desulfurized gypsum is not a direct product of coal combustion, but a product of the scrubber system to remove SO_2 emissions from exhaust (Kisku *et al.*, 2018; Messinger and Silman, 2016).

Physical and chemical properties of coal ash are determined by the geographical location where the raw coal was mined, the type of boiler, and the operating conditions of the power plant (Jayaranjan *et al.*, 2014). Fly ash is composed mainly of oxides such as SiO_2 , Al_2O_3 , Fe_2O_3 , TiO_2 , and CaO . Most natural elements can be found in coal ash, and trace elements include, As, Cd, Cr, Hg, Pb,

Se, and Zn (Greely Jr. *et al.*, 2014; Jayaranjan *et al.*, 2014; Shaheen *et al.*, 2014). Coal bottom ash consists of silicate, carbonate, aluminate, ferrous materials, and several heavy metals and metalloids. Like fly ash, the chemical composition of the bottom ash is dependent on the source of the raw coal, boiler type, and the refinement process of the raw coal (Jayaranjan *et al.*, 2014).

Once produced and collected, coal ash, in many cases, is mixed with water to form a slurry and stored wet in settling ponds. These ponds are constructed either lined or unlined; open to the atmosphere or capped. The coal ash storage pond located at the Dan River Steam Station, was an open, unlined two-pond system where, the coal ash was pumped into one pond where it settled out of the water column, then pumped to a second pond for further settling before liquid effluent was discharged into the river (Messinger and Silman, 2016). In the US, of the approximately 120 Mt of coal ash is produced annually, 54% is disposed of in landfills or settling ponds (American Coal Ash Association, 2012). Possible catastrophic impoundment failures and chronic leaching from unlined impoundments allow the mobilization of coal ash including their associated heavy metals into the environment where these metals may enter the food web directly or indirectly through microbially-mediated transformations (Cabral *et al.*, 2016; Deonaraine *et al.*, 2013; Hershey *et al.*, 2016; Otter *et al.*, 2012).

On February 2, 2014, two storm water drainage pipes located under a coal ash impoundment pond at the Duke Energy Dan River Steam Station near Eden, NC collapsed, releasing approximately 28,000 cubic yards of coal ash and about 27 million gallons of untreated ash wastewater into the Dan River (Lemly, 2015). Following the spill, water and sediment was sampled from the river and Kerr Reservoir downstream of the spill to assess water quality and human health concerns. Test results showed no

constituents to be at levels exceeding safe limits in the water column (Hesterberg *et al.*, 2014; US EPA, 2014). Duke Energy dredged ash deposits at two locations along the river, but likely over 90% of the ash remains buried in river sediments or has been deposited into Kerr Lake (NC DEQ, 2014). While the test results were encouraging for immediate water quality, the long-term concern is the effect of mobilization of coal ash constituents into the riverine food webs.

CHAPTER II

RESPONSE OF MERCURY METHYLATING BACTERIA TO THE COAL ASH SPILL IN THE DAN RIVER

Introduction

The February 2014 coal ash spill mobilized coal ash into the Dan River. One constituent of particular concern during leaching and/or impoundment failure is mercury (Schwartz *et al.*, 2016). Inorganic mercury often passes through an organism, but the bioavailable form of mercury, methylmercury (MeHg) is a known neurotoxin and potential endocrine disruptor and has a high affinity for sulfhydryl groups in proteins (Boyd and Barkay, 2012). This may destabilize proteins and lead to decreased enzymatic activity and reduced overall fitness of organisms (Driscoll *et al.*, 2013; Ehrlich and Newman, 2008).

MeHg is produced in anaerobic conditions predominately by sulfate reducing bacteria (SRB), iron reducing bacteria (FeRB), and methanogens (Liu *et al.*, 2014, Schwartz *et al.*, 2016). Coal ash may provide the necessary substrates, such as sulfate and/or iron, to stimulate the microbial methylation of Hg (Deonarine *et al.*, 2013, Schwartz *et al.*, 2016). Microorganisms have developed various mechanisms to mitigate effects of high concentrations of heavy metal toxins, such as Hg. These include reduction of the metal to a less toxic form, metal complexation, efflux pumps via an energy-dependent membrane transporter, and extracellular sequestration (Binkley and Simpson, 2003; Poulain and Barkay, 2013). In submerged anoxic sediments under certain conditions, inorganic mercury (Hg^{2+}) can be converted into MeHg through microbial metabolism (Dash and Das, 2014; Hershey *et al.*, 2016;

Schaefer *et al.*, 2011, Schwartz *et al.*, 2016). If Hg is methylated, it is bioavailable where, if ingested, could bioaccumulate and biomagnify in the river food webs, posing a health risk to local residents who consume fish. (Dash and Das, 2014; Otter *et al.*, 2012; Rowe, 2014).

The total available amount of MeHg within an ecosystem is controlled by multiple microbial and abiotic processes that reduce availability of Hg^{2+} or degradation of MeHg. Hg^{2+} can be volatilized as Hg^0 through photoreduction or by bacteria with the *merA* gene (Boyd and Barkay, 2012). Additionally, MeHg can be demethylated into Hg^{2+} by sunlight (Tsui *et al.*, 2013) or microbes with the *merB* gene (Bizily *et al.*, 1999).

Two genes are required for methylation of Hg, *hgcA* and *hgcB*. As Hg^{2+} enters the cell, a methylated-HgcA protein transfers a $-\text{CH}_3$ group to Hg^{2+} within the cytosol. HgcB protein is then required to recycle the methylated-HgcA protein (Poulain and Barkay, 2013). The *hgcAB* sequence is conserved across multiple genera and therefore could be utilized as a molecular biomarker for suspected contaminated sites with real-time quantitative PCR (Christensen *et al.*, 2016; Dash and Das, 2014; Lima de Silva *et al.*, 2012). Liu et al. (2014) found that the *hgcA* abundance and the concentration of MeHg in rice paddy soil in China near a mercury mining area is positively correlated (Liu *et al.*, 2014). This finding suggests that microbes may be contributing to the MeHg in the sampled soils. They also found high genetic diversity within the microbial community and that environmental factors such as total Hg, SO_4 , NH_4 , and organic matter influenced the community structure. After phylogenetic analysis, the representative taxa in the community consisted of *Deltaproteobacteria*, *Firmicutes*, *Chloroflexi*, *Euryarchaeota*, and two novel taxa (Liu *et al.*, 2014).

In 2008, a dike failure at the Tennessee Valley Authority Kingston Fossil Plant coal ash pond in Harriman, Tennessee, released an estimated 5.4 million cubic yards of ash into the surrounding community and rivers (Ruhl *et al.*, 2010). The release ruptured a natural gas line, disrupted power and transportation, destroyed three homes, and resulted in the evacuation of nearby neighborhoods. The impoundment pond has since been rebuilt and reinforced to resist natural disasters including earthquakes (TVA, 2011). In sediment samples collected downstream following the spill, total mercury concentrations were three to four times greater than sediments upstream of the spill. MeHg was also slightly higher than upstream (Deonarine *et al.*, 2013).

The coal ash spill into the Dan River similarly mobilized heavy metals into the environment (NC DEQ, 2014). The extent of long-term effects of potential introduction methylated mercury into the food web of the river is unknown. Mercury, along with other coal ash constituents, such as sulfur and iron, may stimulate mercury-methylating microorganisms in anaerobic sediments (Schwartz *et al.*, 2016). The goal of this study was to characterize the response of key microbial community constituents, specifically, *hgcA* abundance as a result of the Dan River coal ash spill.

Objective and Hypothesis

Determine the spatial distribution of mercury-methylating taxa as a result of the coal ash spill using qPCR. I hypothesize that there will be increased abundance of the in the SSU rDNA of mercury methylating taxonomic groups and the *hgcA* gene downstream of spill site due to stimulation by coal ash constituents present in the sediment.

Methods

Study Sites and Sediment Collection

The Dan River is a 344 km river that rises in Patrick Co. Virginia and crosses into North Carolina in Stokes County. It flows across the border between NC and VA several times before flowing into the Kerr Reservoir on the Roanoke River which then flows to the Atlantic Ocean at the Albemarle Sound in North Carolina. This study encompasses sites up to 3.6 km upstream of the spill site in Eden, NC and 64.7 km downstream to Milton, NC, sampled in July 2015, about 17 months following the spill.

To characterize the extent of the coal ash spill impact on the microbial community, samples were collected at three upstream reference sites, one site parallel to the ash ponds but upstream of the spill (leaching site), and five downstream sites including near two sites that were dredged for remediation, one at Town Creek, near the spill site and one near Abreu-Grogan Park, Danville, Va., and potential depositional sites that were not dredged near Danville and downstream (Figure 2.1, Table 2.1).

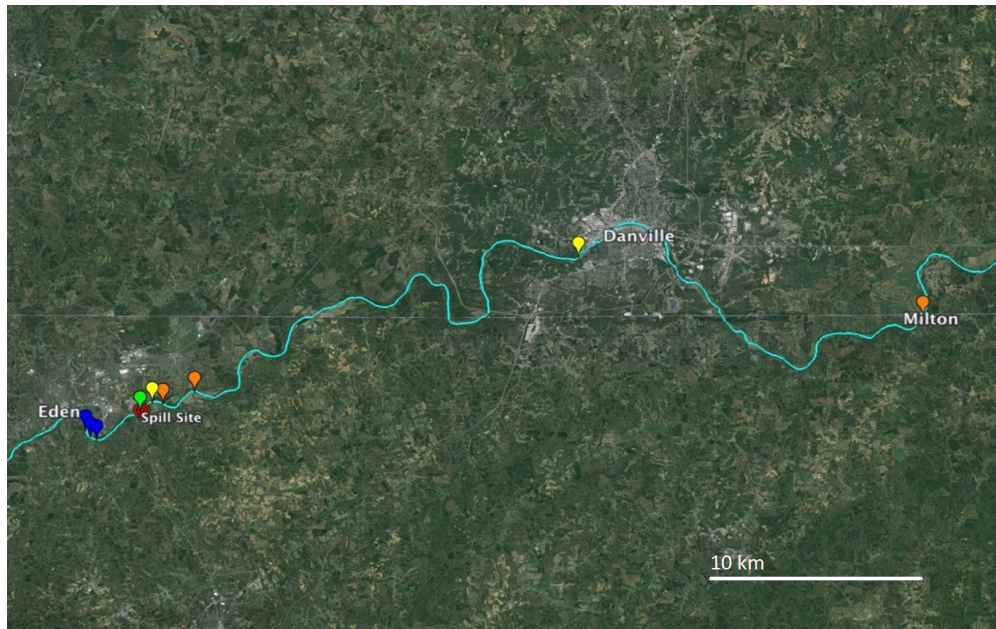


Figure 2.1. Google Earth Image of Dan River and Sampling Locations. Google Earth Image of Dan River and Sampling Locations. Markers indicate sampling sites: Blue, upstream of spill site; Green, leaching site; Yellow, downstream dredged locations; Orange, downstream not dredged.

Each site was accessed by boat, where sediment from the riverbank and channel was collected. Riverbank sediment cores were collected in triplicate using a piston-style coring device. Channel samples were collected using a small dredge. Sediment cores were sectioned by depth and individual segments homogenized according to one of three sampling schemes to reduce the total number of samples to be assayed (Table 2.2). 0.25 cm³ samples were preserved in CTAB (cetyltrimethylammonium bromide) for DNA extraction. Channel sediment was homogenized then 0.25 cm³ was preserved in CTAB. This field study was constrained by access with few boat ramps, and two dams. Frequent high water levels also provided a logistical obstacle for repeated field sampling. Additionally, the leaching site was accessed after permission from Duke Energy using their onsite boat ramp.

Table 2.1. Sampling Locations..

Site ID	Distance from Spill (km)	Longitude and Latitude	Sampling Scheme	Shore Samples (n)	Channel Samples (n)
U-03	3.6	N 36 28.605, W 079 45.018	B	6	1
U-02	2.7	N 36 28.261, W 079 44.598	C	4	1
U-01	1.6	N 36 28.574, W 079 43.989	A	2	1
L-01	0.0	N 36 29.188, W 079 43.025	B	6	0
D-01	0.3	N 36 29.471, W 079 42.722	B	6	1
D-02	1.0	N 36 29.895, W 079 40.836	C	4	1
D-03	4.3	N 36 34.716, W 079 29.596	C	5	1
D-04	36.6	N 36 34.514, W 079 27.024	B	6	1
D-05	38.0	N 36 34.448, W 079 26.211	A	3	1
D-06	64.7	N 36 32.279, W 079 13.038	A	2	1

Table 2.2. Sediment Sampling Schemes..

Scheme	Description	Total (n)
A	Triplicate cores segmented at 8 cm and 16 cm, segments pooled	2
B	Triplicate cores segmented at 8 cm and 16 cm, each core sampled	6
C	Triplicate sediment cores pooled at 0-4 cm, 4-8 cm, 8-12 cm, 12-16 cm	4

DNA Extraction and qPCR

A CTAB extraction of DNA was performed using standard protocol for each sample (Stewart and Via, 1993). The DNA extracted was quantified and subsequently diluted to a standard concentration of 5 ng/ μ L in TE (Tris-EDTA) buffer pH 8.0 and stored at 4°C. Extracted DNA quantity and purity were determined from 2 μ L subsamples of each extraction using Thermo Scientific Nanodrop Spectrophotometer based on the 260/280 nm wavelength ratio.

An Applied Biosystems StepOne real-time PCR System was utilized to detect, amplify, and quantify target DNA and representative taxa to meet the objective. Primers were chosen from literature targeting a general metabolic category. These include generic primers to the *hgcA* gene, a sulfate reducing gene, *dsr*, and the 16S rDNA of iron reducing bacteria. (Geets *et al.* (2006); Schaefer *et al.* (2014); Wagner *et al.* (1998); Daly *et al.* (2000), Table 2.3). Each reaction contained the following: 10 μ L of Power Sybr® Green PCR master mix, 1 μ L of forward primer (10 μ M), 1 μ L of reverse primer (10 μ M), 8 μ L of sterile deionized water, and 1 μ L of extracted DNA (5 ng/ μ L). Three negative control reactions, samples repeated in triplicate, and positive standard controls serially diluted in triplicate, were ran in each 48 well plate. Extracted DNA from a pure culture of *Desulfovibrio africanus* (ATCC 19997), an isolate known to contain the *hgcA* gene was used as a standard for the *hgcA* and SRB

targets (Christensen *et al.*, 2016). Genomic DNA extracted from *Geobacter metallireducens* (ATCC 53774) served as the positive control and standard for FeRB primers (Christensen *et al.*, 2016). The real-time qPCR run method consisted of a holding stage for 5 minutes at 95°C, followed by 40 cycles of 15 seconds at 95°C, 1 minute at 57°C, one minute at 72°C, and a 15-second data collection at 80°C, followed by a melt curve analysis beginning at 95°C, dropping to 60°C and increasing at 0.3°C every second until it reaches 95°C. The relative abundance of targets was computed by the StepOne software using the standard curve. The melt curve was examined to ensure amplicon specificity.

The target abundance was normalized to the volume of extracted sediment as well as to organic matter. The amount of organic matter present in each sample was determined by ash free dry mass (AFDM). Each sample was dried at 60°C for 48 hours, then combusted in a muffle furnace at 500°C for 2 hours. Samples were weighed before and after combustion to calculate the AFDM.

Table 2.3. Primers Used in this Study.

Target	Primer Name	Primer Sequence 5'-3'	Reference
<i>hgcA</i>	hgcA4R	CGCATYTCCTTYTYBACNCC	Liu <i>et al.</i> , 2014
	hgcA4F	GGNRTYAAYRTCTGGTGYGC	
FeRB	Geo-R	TACCCGCTACACCTAGT	Medihala <i>et al.</i> , 2012
	Geo-F	AGGAAGCAACGGCTAACTCC	
SRB	DSV230R	GRGYCYGCGTTYCATTAGC	Daly <i>et al.</i> , 2000
	DSV838F	SYCCGRCAYCTAGYRTYCATC	

Statistical Analysis

Statistical analyses included nonparametric tests for significance to determine the overall effect of sampling location on the abundance of target per sample. To determine whether abundance of the targets vary in the shore sediment based on the

distance from the spill site, the data was first tested for normality with the Shapiro-Wilk test of normality. The data did not follow a Gaussian distribution, therefore, the data were transformed ($\text{LN}(x + 1)$), but still failed the test for normality. Samples were binned according to location irrespective of depth due to limited replication of depths, and evaluated with nonparametric tests. For each target, a Kruskal-Wallis test was performed to assess whether a difference in target abundance exists between sites. Where a significant difference was observed, a Wilcoxon Rank Sum test was performed to determine differences between sampling locations of interest. We hypothesized an increase in DNA abundance with proximity to the spill site, therefore, pairwise comparisons were assessed between Site ID D-01 and all others, as well as L-01 to all others. These tests were then repeated with the data normalized to the mass of sediment and to the amount of organic matter present in each sample.

Due to the lack of replication of river channel samples, the samples collected from the river channel bottom were binned into upstream and downstream groups. A Wilcoxon Rank Sum test was used to determine significance. All analyses were conducted using R (R Core Team, 2019).

Results

Shore Sediment Samples

The Kruskal-Wallis test revealed a significant difference between the abundance of amplified DNA and sampling locations for each of the three qPCR targets assessed (Kruskal-Wallis p-value < 0.05). Each of the targets amplified from the D-01 sampling site, the first sampling site downstream (0.3 km) from the spill site, was detected at a higher abundance compared to at least one of the sites downstream and upstream (Figures 2.2 - 2.4, Table 2.4). However, D-01 was not the

location with the highest overall abundance of these targets. The sampling location U-03, the farthest upstream location, 3.6 km upstream of the spill, and 0.5 km downstream from the Smith River confluence, showed the highest values for each of the targets tested (Figures 2.2 - 2.4, Table 2.5).

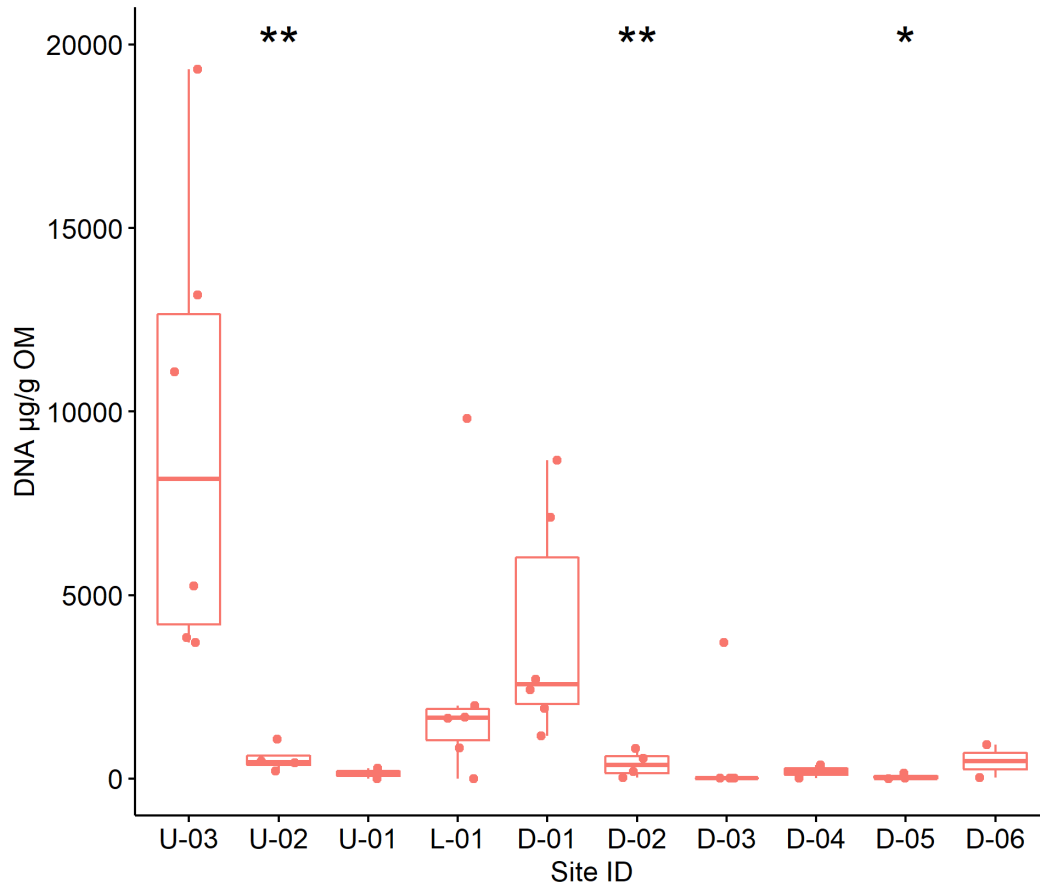


Figure 2.2. *hgcA* Abundance Normalized to Organic Matter at Each Sampling Location. *hgcA* Abundance Normalized to Organic Matter at Each Sampling Location. Asterisks signify difference between the reference, D-01 using the Wilcoxon Rank Sum test. * : $p \leq 0.05$, ** : $p \leq 0.01$

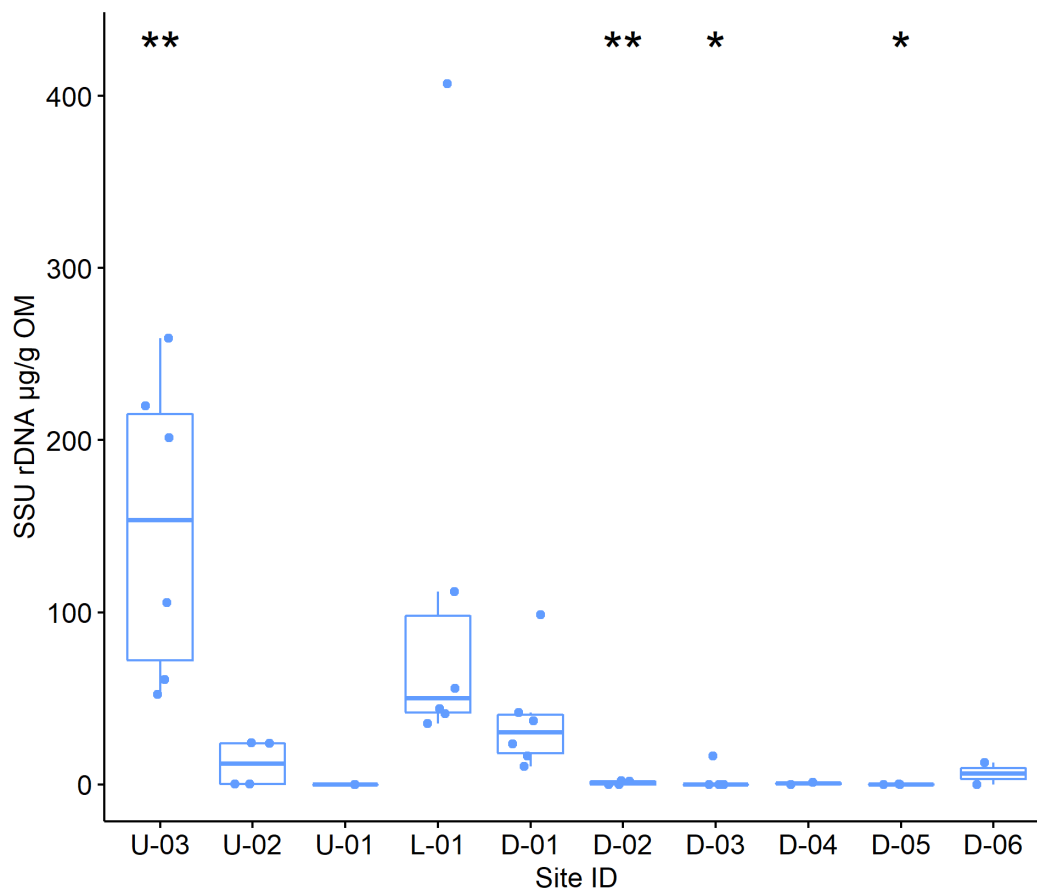


Figure 2.3. FeRB Abundance Normalized to Organic Matter at Each Sampling Location. FeRB Abundance Normalized to Organic Matter at Each Sampling Location. Asterisks signify difference between the reference, D-01 using the Wilcoxon Rank Sum test. * : $p \leq 0.05$, ** : $p \leq 0.01$

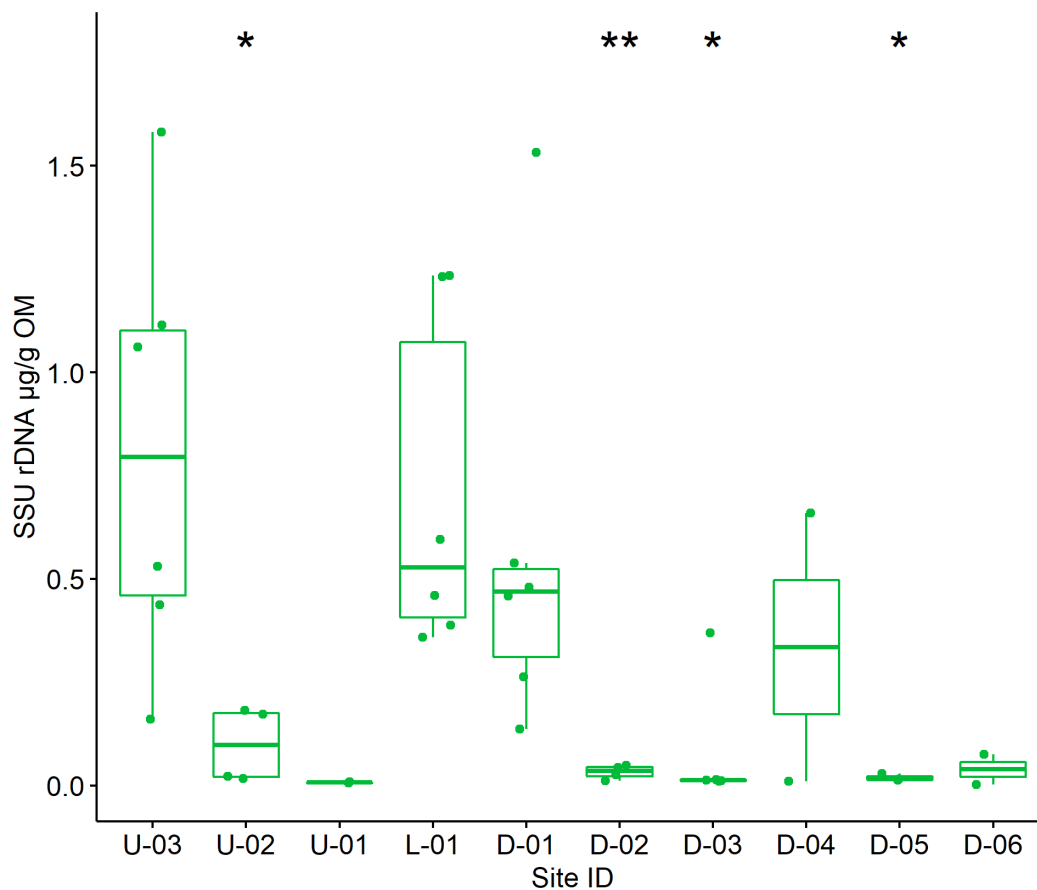


Figure 2.4. SRB Abundance Normalized to Organic Matter at Each Sampling Location. SRB Abundance Normalized to Organic Matter at Each Sampling Location. Asterisks signify difference between the reference, D-01 using the Wilcoxon Rank Sum test. * : $p \leq 0.05$, ** : $p \leq 0.01$

Table 2.4. Comparisons of Each Sampling Location to the Location Directly Downstream of the Spill. Comparisons of Each Sampling Location to the Location Directly Downstream of the Spill. Wilcoxon Rank Sum Test, ns: not significant, * : $p \leq 0.05$, ** : $p \leq 0.01$

Location	<i>hgcA</i>	FeRB	SRB
U-03	ns	**	ns
U-02	**	ns	*
U-01	ns	ns	ns
L-01	ns	ns	ns
D-02	**	**	**
D-03	ns	*	*
D-04	ns	ns	ns
D-05	*	*	*
D-06	ns	ns	ns

Table 2.5. Comparisons of Each Sampling Location to the Furthest Upstream Location. Comparisons of Each Sampling Location to the Furthest Upstream Location. Wilcoxon Rank Sum Test, ns: not significant, * : $p \leq 0.05$, ** : $p \leq 0.01$

Location	<i>hgcA</i>	FeRB	SRB
U-03	ns	**	ns
U-02	**	ns	*
U-01	ns	ns	ns
L-01	ns	ns	ns
D-02	**	**	**
D-03	ns	*	*
D-04	ns	ns	ns
D-05	*	*	*
D-06	ns	ns	ns

Additionally, the leaching site, L-01 is not significantly different from the D-01 location. When comparing the leaching site to all others, targets abundance is slightly higher than most of the other sampling locations. This pattern is similar to the D-01 comparisons, but overall, less pairs are significantly different (Table 2.6). The same pattern was observed between the data normalized to the mass of sediment in each sample as well as the amount of organic matter present.

Table 2.6. Comparisons of Each Sampling Location to the Leaching Site Location. Comparisons of Each Sampling Location to the Leaching Site Location. Wilcoxon Rank Sum Test, ns: not significant, * : $p \leq 0.05$, ** : $p \leq 0.01$

Location	<i>hgcA</i>	FeRB	SRB
U-03	ns	**	ns
U-02	**	ns	*
U-01	ns	ns	ns
L-01	ns	ns	ns
D-02	**	**	**
D-03	ns	*	*
D-04	ns	ns	ns
D-05	*	*	*
D-06	ns	ns	ns

Within this study, replication of core depth was limited, therefore, the interaction between the depth of samples was not evaluated. However, a scatterplot of the data did not show any clear patterns indicating a depth interaction, but the effect is inconclusive (Figure 2.5).

River Channel Samples

The samples collected in the river channel were limited by one sample at each site. To assess significance, samples were binned according to upstream and downstream. For each of the targets tested, there is no significant difference between the groups (Figure 2.6).

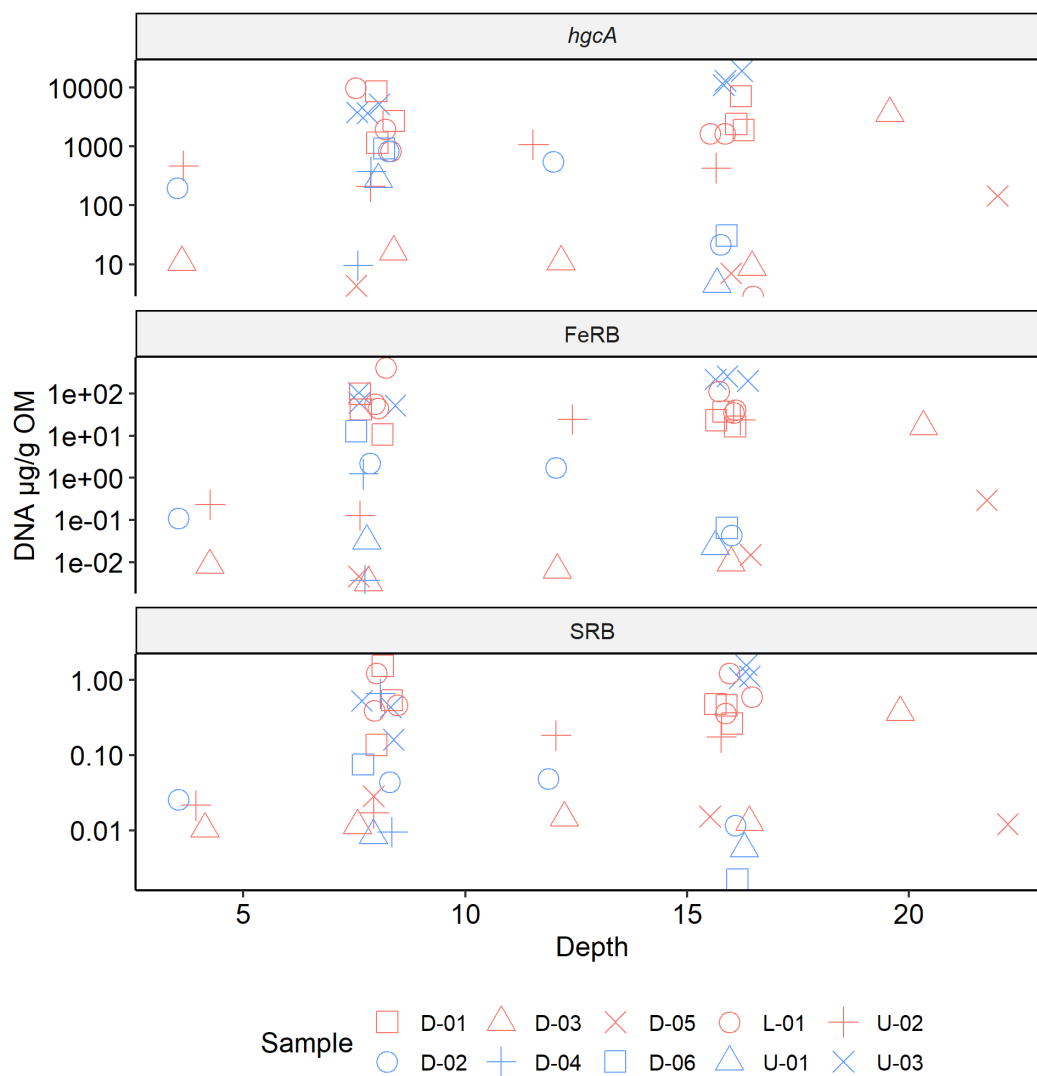


Figure 2.5. Target DNA Concentration at Each Depth. Target DNA Concentration at Each Depth.

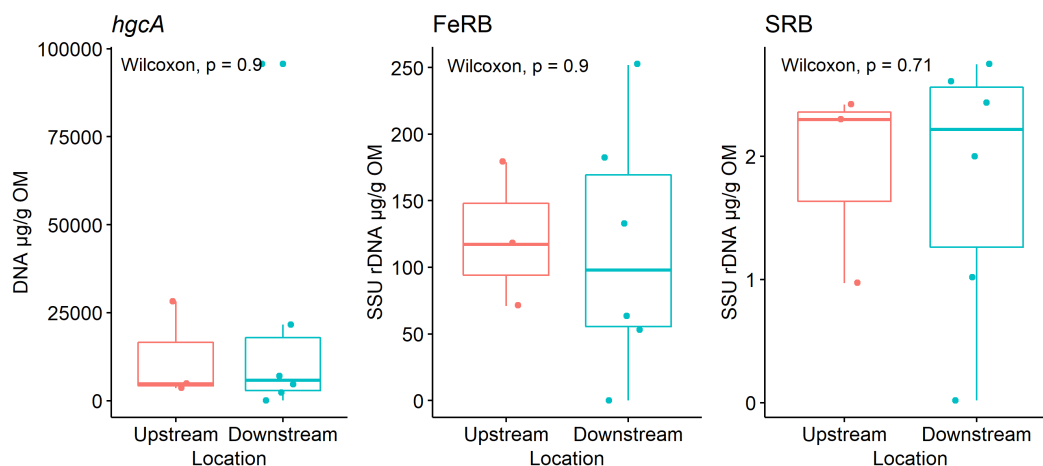


Figure 2.6. River Channel Abundance at Sampling Sites Upstream and Downstream Normalized to Organic Matter. River Channel Abundance at Sampling Sites Upstream and Downstream Normalized to Organic Matter.

Discussion

All primers tested exhibited similar patterns of significance and abundance per location. The abundance of targets detected at the sampling location 0.3 km downstream of the spill site, D-01, is higher than some locations upstream and downstream. However, the highest abundance of these three targets was found at the furthest upstream site of all locations. Also, the leaching site, L-01, exhibited higher target signal over some locations. This may signify that there could be stimulation of microbial growth, but assessment of DNA abundance does not inform methylation activity. Mercury methylation is dependent on environmental factors, where substrates such as inorganic mercury, sulfate, and organic matter must be unbound and freely available for uptake under anoxic conditions.

The results from this microbial study are inconclusive. We attempted to better define the mercury methylating potential of the microbial community using primers from (Christensen *et al.*, 2016), but experienced technical difficulties and

were not able to successfully use them. Hg contamination within the sediment may still be possible, and our methods of sampling sediment may not have been robust to the variable and “hotspot” nature of the coal ash deposition and dispersal into the river sediments. It is plausible that 1.5 years after the spill event, the coal ash is buried in the sediment, the substrates may not be available for uptake, or the coal ash has been transported further downstream than our furthest downstream location.

CHAPTER III

IDENTIFICATION AND CHARACTERIZATION OF HEAVY METAL TOLERANCE OF BACTERIA ISOLATED FROM COAL ASH

Introduction

Coal ash is a waste product generated from coal-fired power plants. These power plants are located near water sources to meet water demands during the electricity generation process. The ash and other waste products are discharged as a slurry into a settling pond, after the combustion process (Connors, 2015). To prevent overflow the pond is maintained by pumping surface water into the nearby waters, usually a river. In North Carolina, 14 such coal ash ponds are maintained by Duke Energy. Unlined and uncapped coal ash waste ponds therefore have the potential for leaching contaminants into the groundwater or spilling into nearby waterways possibly polluting drinking water and/or disrupting aquatic ecological communities (Otter *et al.*, 2012; Ruhl *et al.*, 2009).

On February 2, 2014, a coal ash pond located at the retired Dan River Steam Station near Eden, NC expelled approximately 39,000 tons of coal ash slurry into the Dan River of North Carolina due to an underground pipe collapse (NC DEQ, 2014). The ash was dispersed along the river bottom between the spill site and the Kerr Reservoir, approximately 120 km downstream (Hesterberg *et al.*, 2014). Dredging operations conducted by Duke Energy at Town Creek and an area upstream from the Schoolfield Dam in Danville, VA, removed approximately 2,500 tons of coal ash-laden sediment, but the remainder was deposited in the river bottom (Lemly, 2015).

Physical and chemical properties of coal ash are determined by the geographic location where the raw coal was mined, the type of boiler, and the operating conditions of the power plant (Jayaranjan *et al.*, 2014). Most natural elements can be found in coal ash including trace elements such as arsenic, cadmium, chromium, mercury, lead, selenium, and zinc (Greely Jr. *et al.*, 2014; Jayaranjan *et al.*, 2014). Coal ash is wholly comprised of multiple waste products with differing chemical properties including fly ash, bottom ash, and byproducts of pollution mitigation processes, such as scrubber systems. Fly ash is composed mainly of oxides such as, SiO_2 , Al_2O_3 , Fe_2O_3 . Bottom ash consists of silicate, carbonate, aluminate, ferrous materials and high concentrations of several heavy metals and metalloids (Kisku *et al.*, 2018).

Heavy metals are characterized by a density greater than 5 g/cm^3 , mostly transition elements, and play an important role as trace elements in biochemical reactions. These heavy metals, due to an incompletely filled d orbital, allow the cations to form complex compounds with the potential to be redox reactive. Heavy metal ions form unspecific complex compounds in living cells, leading to toxic effects. For example, Hg^{2+} , Cd^{2+} , and Ag^+ form strong toxic complexes that are not conducive to any physiological function. Trace metals such as Zn^{2+} , Ni^{2+} , and Cu^{2+} are required for some biological functions but they are toxic at high concentrations Hesterberg *et al.* (2014). Therefore, the intracellular concentration of heavy metals must be controlled, and organisms have adapted heavy metal resistance strategies (Binkley and Simpson, 2003; Poulain and Barkay, 2013).

Through the process of coal combustion, the coal ash is rendered sterile. Therefore, inoculation of coal ash in waste ponds occurs by natural processes including atmospheric deposition through rainfall and windblown particulates.

Opportunistic microorganisms may be adapted, or may be able to adapt, to the high concentration of heavy metals through resistance or metal detoxification.

Perturbations such as coal ash spills result in altered environments in receiving waters, which microbes may adapt to and subsequently mobilize coal ash constituents into food webs. Knowledge of the microbial community structure and distribution is important to estimate the extent of biological mobility of these pollutants. Further, organisms which exhibit metal tolerance may warrant further investigation into their potential for bioremediation (Keshri *et al.*, 2014; Naik and Dubey, 2013; Pepi *et al.*, 2011; Raja and Omine, 2012).

The objective of this study was to determine if a microbial community is viable and present in coal ash ponds, and if present, identify the taxa and assess the metal tolerance of isolated organisms. Klubek *et al.*, 1992 analyzed the microbial community of coal ash ponds, but this study did not identified groups of metabolic capabilities. Roychowdhury *et al.* 2018, isolated 10 bacteria representing three genera from wheathered fly ash pond samples. Raja and Omine 2013, identified boron tolerant isolates from a fly ash dumping site, and Stepanauskas *et al.* 2005, evaluated the metal tolerance of microbial communities of intake and discharge of coal ash ponds. In a preliminary PCR analysis, a small amount of DNA was present in a coal ash sample from the Dan River Steam Station coal ash pond, but we were not able to amplify SSU rDNA from bacteria with universal prokaryotic primers, possibly due to contaminants present in the sample or low concentrations of bacterial DNA. Therefore, in this study, we increased the abundance of microbes and their DNA by culturing and isolating bacteria from samples of coal ash, to ensure adequate amounts of DNA for analyses.

Objectives and Hypotheses

Objective 1: To identify isolated bacteria using qPCR amplification and rDNA sequencing. I seek to identify these taxa using qPCR amplification and rDNA sequencing to compare to the GenBank database.

I hypothesize novel organisms may be discovered which may have unique metal tolerance capabilities that may be useful for bioremediation.

Objective 2: To determine the metal tolerance of bacteria isolated from coal ash collected from the Dan River impoundment site.

I hypothesize that microbial growth of bacterial isolates from pure coal ash collected at the Dan River Steam Station impoundment pond will be tolerant to heavy metals.

Methods

Pure Culture Isolation from Coal Ash

Samples of coal ash provided by our collaborator, Brian Williams of the Dan River Basin Association, were taken directly from a coal ash retention pond at the retired Dan River Steam Station near Eden, NC. To culture organisms, aliquots (0.5 g) of coal ash from the pond were added to six 50 mL conical tubes. 40 mL of filter sterilized (0.22 μ m pore) Dan River water was added to three tubes and filter sterilized Dan River water supplemented with 10% nutrient broth to the other three tubes. Additionally, filter sterilized Dan River water alone was evaluated in triplicate to serve as a sterility control. Tubes were incubated at ambient room temperature for 48 hours without agitation. 1 mL aliquots were then transferred from each culture and spread on 50% nutrient agar plates for isolation. Morphologically unique colonies

identified on spread plates were isolated and pure cultures maintained on 50% nutrient agar slants for heavy metal tolerance experimentation.

Heavy Metal Tolerance Characterization

Optical density (OD) at 580 nm, of liquid broth culture was assessed as a proxy for growth of organisms, since growth is related to the increase in turbidity of a bacterial culture. Metals tested include, arsenic, cadmium, chromium, mercury, lead, selenium and zinc. Stock concentrations of Na_2HAsO_4 , CdCl_2 , CrCl_3 , HgCl_2 , PbCl_2 , Na_2SeO_3 , and ZnCl_2 were serially diluted to span several orders of magnitude greater and less than natural environmental concentrations reported in the literature for each metal, resulting in five concentration levels per metal (Table 3.1).

Table 3.1. Final Concentrations of Metal in Broth (nM). Final Concentrations of Metal in Fermentation Broth (nM).

Factor	As	Cd	Cr	Hg	Pb	Se	Zn
0	0	0	0	0	0	0	0
1	0.000135	0.008896	0.001923	0.004985	0.04826	0.1266	0.0153
2	0.00135	0.08896	0.01923	0.04985	0.4826	1.266	0.153
3	0.0135	0.8896	0.1923	0.4985	4.826	12.66	1.53
4	0.135	8.896	1.923	4.985	48.26	126.6	15.3
5	1.35	88.96	19.23	49.85	482.6	1266	153

Metal stocks were prepared by dissolving each metal salt in sterile reverse osmosis deionized (RO/DI) water and subsequently diluted. To ensure a viable culture inoculum, sufficient volume of cells, and standardization for experimentation, each unique isolate was incubated in a 50% nutrient broth culture in a shaking water bath at 25°C overnight until an optical density of at least 0.7 at 580 nm was observed. For each metal-isolate experiment, 100 μL of 0.7 OD~580 nm~ inoculum was added to 2.4 mL of 50% nutrient broth amended with 100 μL each concentration of metal in

triplicate. For the control, 100 μ L of RO/DI water was substituted for the metal spike in triplicate. Growth patterns were measured by absorbance at 580 nm, and recorded at ten time points, approximately 0, 1, 2, 4, 8, 16, 24, 30, 36, and 48 hours post inoculation.

Isolate Identification and Phylogenetic Analysis

Isolates were grown in 50% nutrient broth then pelleted by centrifugation. The pellet was resuspended in CTAB buffer. DNA was extracted and purified using the CTAB method (Stewart and Via, 1993), then amplified with universal 16S primers (Bruce *et al.*, 1992; Edwards *et al.*, 1989). PCR products were then purified and commercially sequenced. Sequence chromatogram data was evaluated and edited manually to optimize the sequence. The FASTA files of sequences were imported into Molecular Evolutionary Genetics Analysis version 7.0 (MEGA7) (Kumar *et al.*, 2016). Sequences were aligned using MUSCLE and manual adjustments (Edgar, 2004). The resulting aligned sequences were then subjected to phylogenetic tree analysis using MEGA7. The maximum likelihood tree was computed using MEGA7 using the Jukes-Cantor model (Jukes and Cantor, 1969). The bootstrap consensus tree was inferred from 1000 replicates. Initial trees with a greater log likelihood value were calculated by applying Neighbor-Join and BioNJ algorithms to a matrix of pairwise distances estimated using the Maximum Composite Likelihood approach. To identify the genus of each isolate, the sequences were compared to the NCBI GenBank database with BLASTn (Nucleotide Basic Local Alignment Search Tool). Genus calls were made with similarities greater than 97% (Clark *et al.*, 2016).

Statistical Analyses

To characterize the heavy metal tolerance of isolates in liquid broth culture, the carrying capacity and growth rate of each isolate-metal experiment was modeled

with a Gompertz sigmoidal function (Zwietering *et al.*, 1990). Significant differences of growth rate and carrying capacity within each experiment were determined using the Kruskal-Wallis one-way analysis of variance test (p-value < 0.05). Experiments were excluded from analysis where no growth was observed in the control. Isolates were deemed metal tolerant at the concentrations tested if no significant difference between carrying capacity or growth rate was observed between concentrations.

Results

Phylogenetic Analysis

A total of 31 microorganisms were isolated from the coal ash sample and sequenced. Sequences were aligned and a phylogenetic tree was constructed (Figure 3.1). Sequences were predominantly identified as *Bacillus* or *Arthrobacter* spp. One isolate, DR 76, was not identified in GenBank with the BLASTn search.

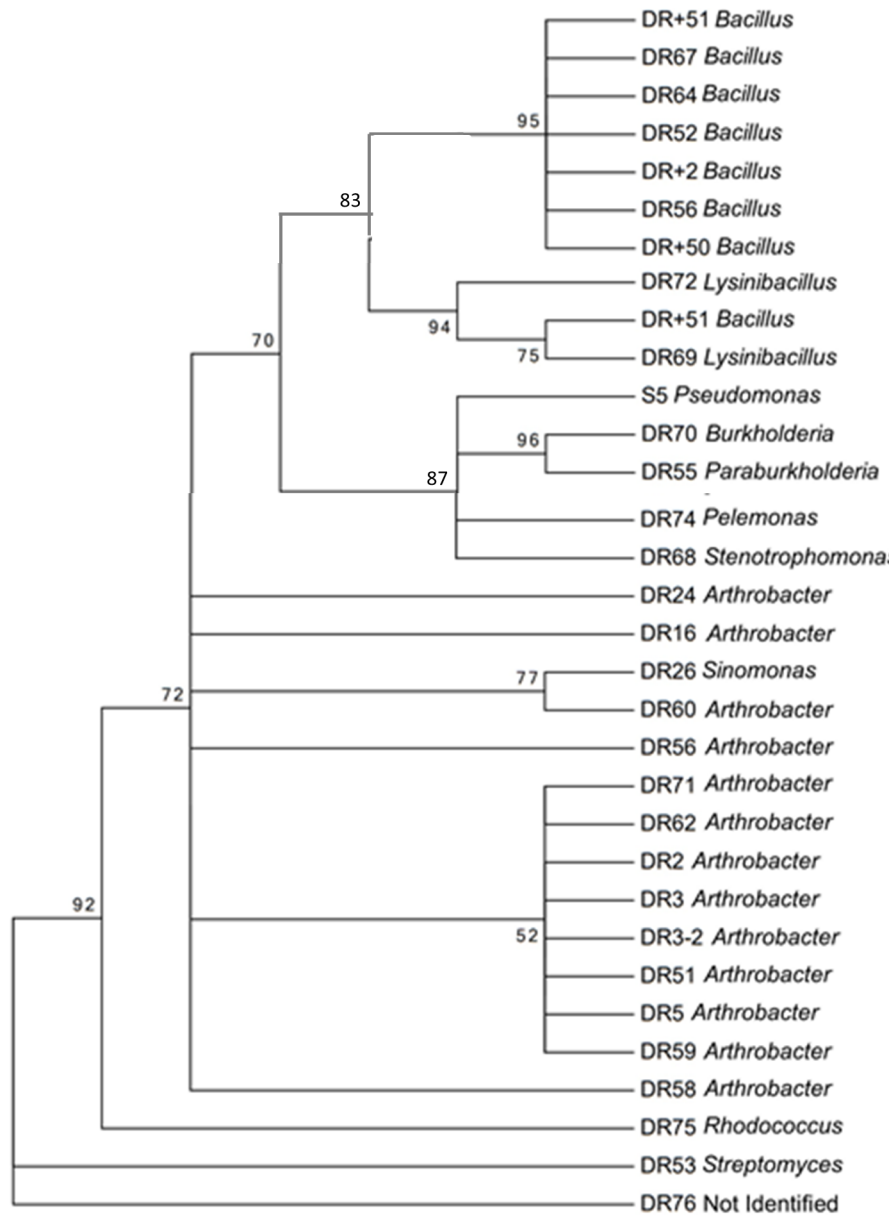


Figure 3.1. Phylogenetic Analysis of Isolates Cultured from Coal Ash Amended Media.
Phylogenetic Analysis of Isolates Cultured from Coal Ash Amended Media.

Metal Tolerance Assays

We tested metal tolerance of 14 isolates. The isolates were challenged against seven metals, resulting in a total of 94 trials. Isolates generally grew well. Figures 3.2 - 3.4 are exemplary of typical growth responses. Figure 3.2 is representative of similar growth response among differing metal concentrations. Carrying capacity is significantly different between metal concentrations of Cd and Zn, and growth rates are different with As and Zn. Figure 3.3 showed a varied response where carrying capacity is significantly different between metal concentrations of Cd and Se, but no difference is observed when testing maximal growth rates. Figure 3.4, Se is an example of no growth. When evaluating the growth of DR 52 challenged by the other metals, the carrying capacity is significantly different between metal concentrations of Cd, and growth rates are different with As, Cd, and Hg (Kruskall-Wallis test $p\text{-value} < 0.05$).

When evaluating carrying capacity, 70 trials were not different from the control, that is, they were metal tolerant (Kruskall-Wallis test $p\text{-value} > 0.05$, Table 3.2). Evaluation of maximal growth rate produced 71 isolate-metal trials that were tolerant (Table 3.3). Metal intolerance (i.e. reduced growth rate or lower carrying capacity) was found in 64% of the isolates tested with Cd and 72% are susceptible to Se. The other metals tested (As, Cr, Hg, Pb, and Zn) resulted in much less susceptibility, ranging from 1-3 susceptible isolates.

High concentrations of Cd had an effect on most of the isolates tested. Of the five isolates that were tolerant of Cd, one is *Arthrobacter*, although not all *Arthrobacter* were tolerant of Cd. *Bacillus* spp. are genomically diverse, ubiquitous endospore-formers are highly resistant to stressors (Earl *et al.*, 2008). Two *Bacillus*

spp. were subjected to metal tolerance assays. The *Bacillus* spp. DR52 and DRp51 exhibited the same metal tolerance characterization; tolerant to all except Cd and Se.

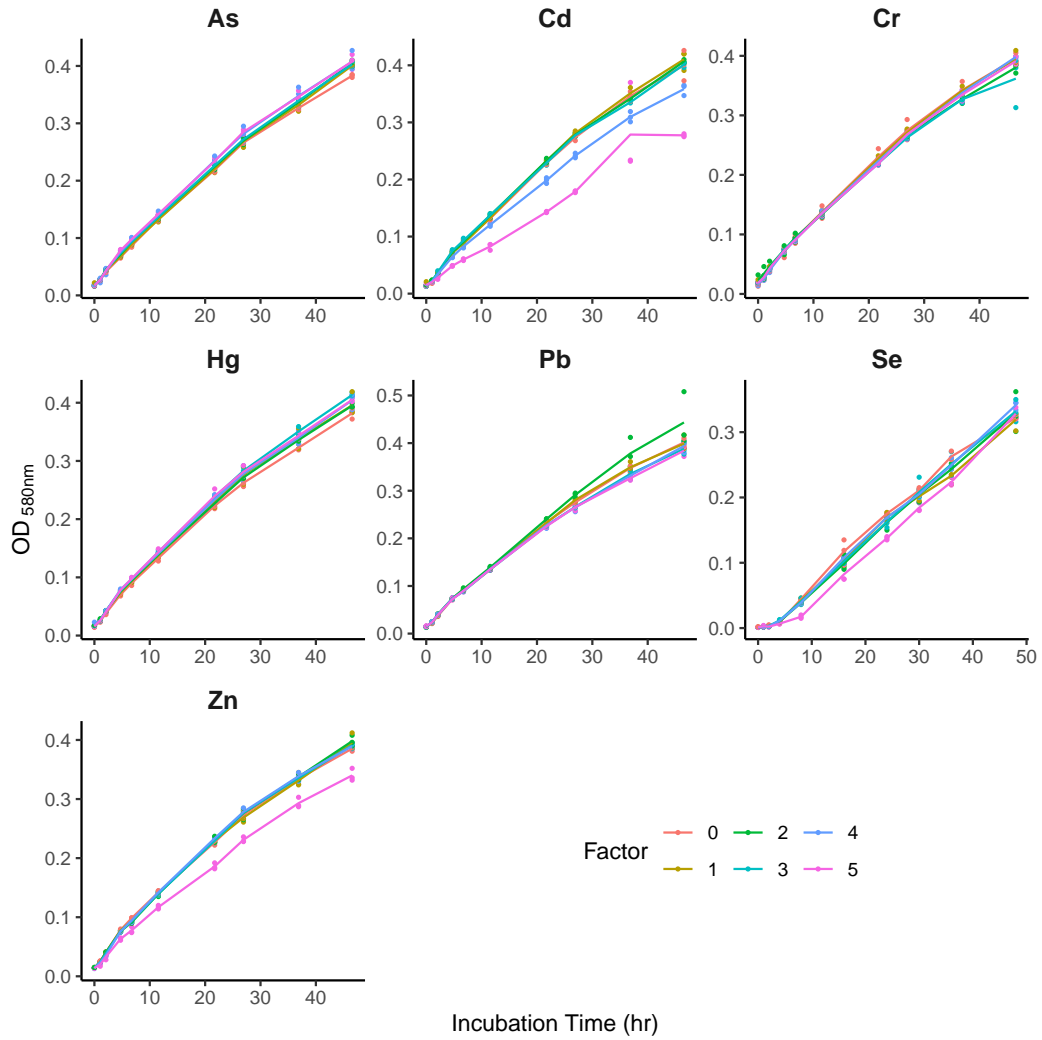


Figure 3.2. Growth of Isolate DR13. Growth of Isolate DR13. Lines connect means of three replicates, colors represent concentration levels (Table 3.1). Carrying capacity is significantly different between metal concentrations of Cd and Zn. Growth rates are different with As and Zn (Kruskal-Wallis test p-value < 0.05)

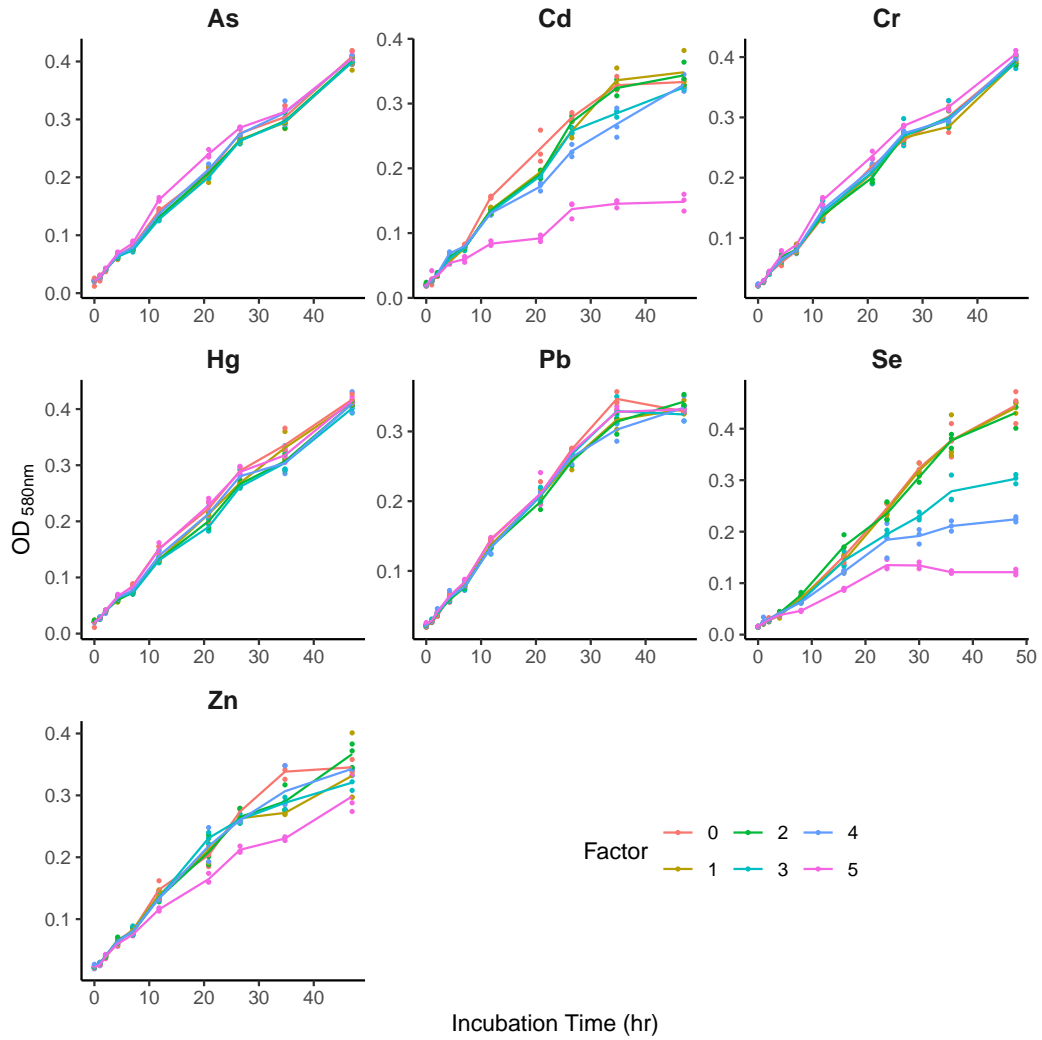


Figure 3.3. Growth of Isolate DR3.2. Growth of Isolate DR3.2. Lines connect means of three replicates, colors represent concentration levels (Table 3.1). Carrying capacity is significantly different between metal concentrations of Cd and Se. No difference is observed when testing maximal growth rates (Kruskal-Wallis test p-value < 0.05)

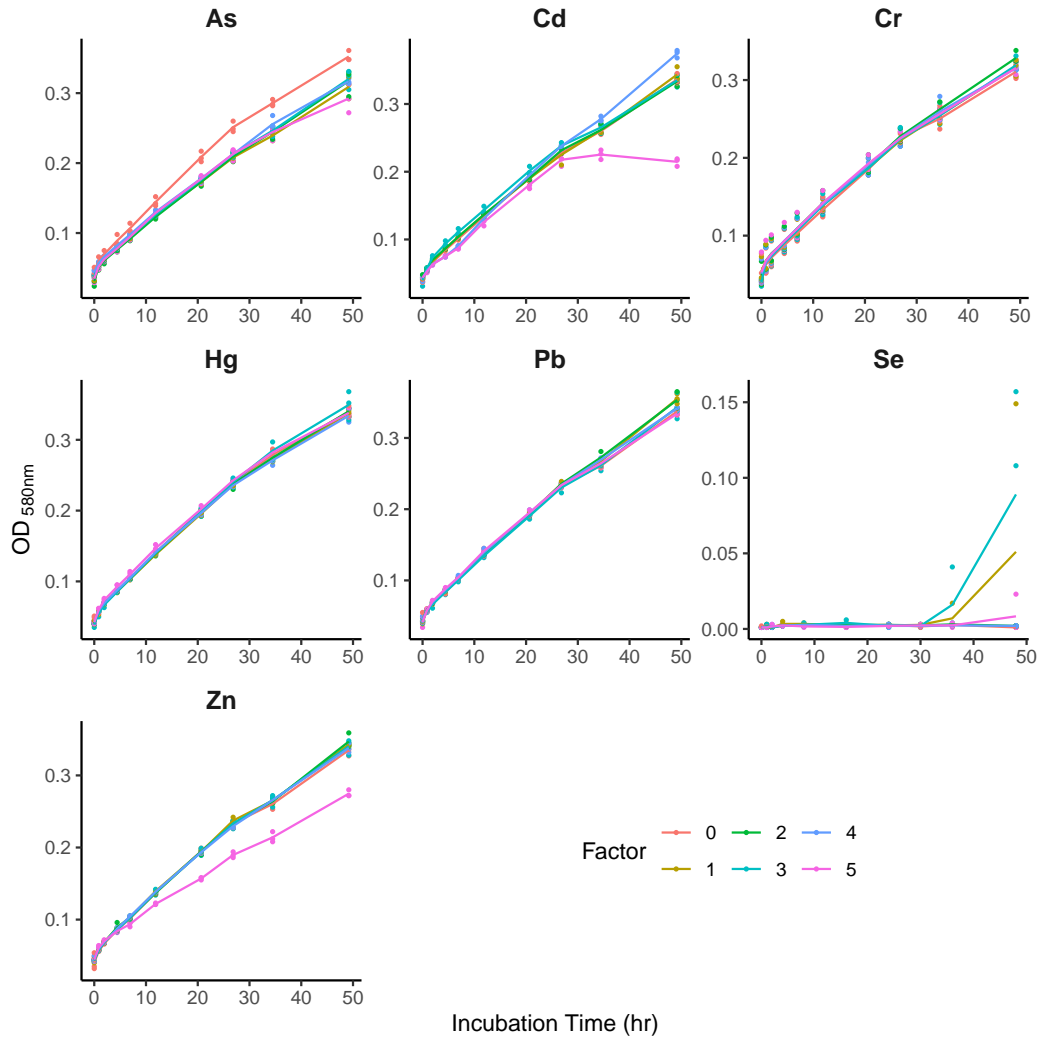


Figure 3.4. Growth of Isolate DR52. Growth of Isolate DR52. Lines connect means of three replicates, colors represent concentration levels (Table 3.1). Se was excluded from analysis. Carrying capacity is significantly different between metal concentrations of Cd. Growth rates are different with As, Cd, and Hg. (Kruskall-Wallis test p -value < 0.05)

Table 3.2. P-values of Kruskal-Wallis Test of Carrying Capacity. P-values of Kruskal-Wallis Test of Carrying Capacity. Bold typeface signifies p-value <0.05. NA: Not Analyzed (Data listed in Appendix)

Genus	Isolate	As	Cd	Cr	Hg	Pb	Se	Zn
Not Sequenced	DR13	0.138	0.046	0.148	0.086	0.281	0.083	0.025
<i>Arthrobacter</i>	DR24	0.141	0.025	0.554	0.504	0.163	NA	0.022
<i>Sinomonas</i>	DR26	0.482	0.02	0.612	0.547	0.134	0.011	0.067
<i>Arthrobacter</i>	DR3.2	0.367	0.027	0.637	0.091	0.669	0.011	0.344
<i>Arthrobacter</i>	DR5	0.171	0.203	0.961	0.494	0.075	0.074	0.16
<i>Bacillus</i>	DR52	0.364	0.009	0.691	0.425	0.066	NA	0.081
<i>Arthrobacter</i>	DR60	0.038	0.04	0.097	0.135	0.074	NA	0.164
Unknown Genus	DR76	0.125	0.336	0.517	NA	0.18	0.014	0.02
<i>Bacillus</i>	DRp51	0.116	0.014	0.54	0.713	0.787	0.008	0.193
<i>Lysinibacillus</i>	DRp72	0.173	0.018	0.096	0.057	0.43	NA	0.104
<i>Peletonas</i>	DRp74	0.385	0.105	0.015	0.016	0.131	0.032	0.669
Not Sequenced	NO14	0.243	0.031	0.297	0.97	0.822	0.03	0.741
Not Sequenced	NO17	0.051	0.067	0.22	0.329	0.035	0.054	0.063
Not Sequenced	NO22	NA	NA	NA	NA	NA	0.014	NA

Table 3.3. P-values of Kruskal-Wallis Test of Growth Rate. P-values of Kruskal-Wallis Test of Growth Rate. Bold typeface signifies p-value <0.05. NA: Not Analyzed (Data listed in Appendix)

Genus	Isolate	As	Cd	Cr	Hg	Pb	Se	Zn
Not Sequenced	DR13	0.042	0.177	0.445	0.825	0.251	0.114	0.021
<i>Arthrobacter</i>	DR24	0.284	0.026	0.073	0.183	0.637	NA	0.02
<i>Sinomonas</i>	DR26	0.306	0.128	0.149	0.383	0.114	0.009	0.334
<i>Arthrobacter</i>	DR3.2	0.206	0.068	0.171	0.074	0.213	0.055	0.083
<i>Arthrobacter</i>	DR5	0.419	0.036	0.192	0.637	0.56	0.012	0.511
<i>Bacillus</i>	DR52	0.028	0.007	0.889	0.041	0.063	NA	0.078
<i>Arthrobacter</i>	DR60	0.028	0.071	0.054	0.082	0.141	NA	0.03
Unknown Genus	DR76	0.218	0.451	0.534	NA	0.399	0.039	0.028
<i>Bacillus</i>	DRp51	0.154	0.138	0.025	0.439	0.975	0.136	0.279
<i>Lysinibacillus</i>	DRp72	0.263	0.035	0.255	0.158	0.385	NA	0.173
<i>Pelemonas</i>	DRp74	0.669	0.261	0.048	0.021	0.056	0.141	0.488
Not Sequenced	NO14	0.324	0.086	0.713	0.9	0.801	0.014	0.648
Not Sequenced	NO17	0.174	0.107	0.68	0.825	0.112	0.013	0.286
Not Sequenced	NO22	NA	NA	NA	NA	NA	0.05	NA

Discussion

We isolated cultures of aerobic bacteria from samples of coal ash. The coal ash isolates were predominantly identified as *Arthrobacter* and *Bacillus* spp. *Arthrobacter* spp. are ubiquitous and have been found in common soils as well as extreme environments. Several studies have isolated bacteria from coal ash.

Roychowdhury *et al.* 2018 found 10 isolates represented by *Bacillus*, *Micrococcus*, *Kytococcus*, and *Staphylococcus* genera. In our study we isolated 31 bacteria including 8 *Bacillus* spp. but none of the other genera. Raja and Omine 2013, using 16S rRNA sequencing and phylogenetic analysis, identified 4 genera from a fly ash dumping site, including *Bacillus* and *Lysinibacillus*, which we also found. They also found *Microbacterium* and *Ralstonia* spp. Pangayao *et al.* 2018, isolated *Pseudomonas* spp. from multiple coal ash ponds, which we did not find. The *Arthrobacter* sp. that constituted the majority of our isolates (16 of 31), are spore-formers that are able to survive long periods of time under stressful conditions

such as nutrient deficiency, temperature shifts, and toxic chemicals. They are metabolically diverse, have been utilized to biodegrade environmental pollutants, and are highly resistant to heavy metals (Mongodin *et al.*, 2006). The predominance of *Arthrobacter* sp. among our coal ash isolates may be due to the selective pressure of the coal ash or the isolation methods.

One isolate could not be identified by a BLASTn search of the GenBank database, and may be an interesting organism to for further testing and characterization. Since we only studied organisms which were easily grown and maintained within the lab under room temperature conditions, likely only a very small portion of the organisms that were present in coal ash were cultivated. More novel organisms may be discoverable using alternate culturing methods.

Our isolates were generally metal tolerant as in previous studies. For example, Stepanauskas *et al.* 2005 documented metal tolerance in bacteria from ash settling basins by flow cytometric analyses. Roychowdhury *et al.* 2018 found As tolerance in isolates from wheathered pond ash samples. Raja and Omine *et al.* 2013, found boron tolerant microbes from a fly ash dumping site in Japan. All of these studies along with ours, demonstrate metal tolerance, and our assay appears useful as a screen to find interesting isolates for further study.

CHAPTER IV

CONCLUSIONS

We found no significant evidence of increased mercury methylation potential from the coal ash spill into the Dan River. We observed similar patterns of abundance of each qPCR target among the sample locations, indicating a similar response by multiple taxonomic groups. Unexpectedly, the highest abundance among all targets was observed at a site that seems unlikely to have been contaminated with coal ash due to its distance upstream from the spill site.

The coal ash isolates were generally tolerant to the concentrations of heavy metals tested, except for cadmium and selenium. However, we did not measure the ion concentrations directly to determine if the metal was unbound and available for uptake. Alternate isolation protocols to capture a more diverse group of isolates may better improve our understanding of coal ash bacteria.

Due to the dynamic nature of river systems, assaying for polluting spills or other perturbations is difficult. A study with improved sampling methods may better elucidate risk of mercury-methylation and mobilization into the food webs.

REFERENCES

- American Coal Ash Association (2012). 2012 Coal Combustion Product (CCP) Production and Use Survey Report. Tech. rep., ACAA.
- Barkay, T., Miller, S. M., & Summers, A. O. (2003). Bacterial mercury resistance from atoms to ecosystems. *FEMS Microbiology Reviews*, *27*(2-3), 355–384.
- Bizily, S. P., Rugh, C. L., Summers, A. O., & Meagher, R. B. (1999). Phytoremediation of methylmercury pollution: merB expression in *Arabidopsis thaliana* confers resistance to organomercurials. *PNAS*, *96*(12), 6808–6813.
- Boyd, E., & Barkay, T. (2012). The mercury resistance operon: From an origin in a geothermal environment to an efficient detoxification machine. *Frontiers in Microbiology*, *3*, 349.
- Bruce, K. D., Hiorns, W. D., Hobman, J. L., Osborn, A. M., Strike, P., & Ritchie, D. A. (1992). Amplification of DNA from native populations of soil bacteria by using the polymerase chain reaction. *Applied and Environmental Microbiology*, *58*(10), 3413–3416.
- Cabral, L., Yu, R.-Q., Crane, S., Giovanella, P., Barkay, T., & Camargo, F. A. O. (2016). Methylmercury degradation by *Pseudomonas putida* V1. *Ecotoxicology and Environmental Safety*, *130*, 37–42.
- Christensen, G. A., Wymore, A. M., King, A. J., Podar, M., Hurt, R. A., Santillan, E. U., Soren, A., Brandt, C. C., Brown, S. D., Palumbo, A. V., Wall, J. D., Gilmour, C. C., & Elias, D. A. (2016). Development and validation of broad-range qualitative and clade-specific quantitative molecular probes for assessing mercury methylation in the environment. *Applied and Environmental Microbiology*, *82*(19), 6068–6078.
- Clark, K., Karsch-Mizrachi, I., Lipman, D. J., Ostell, J., & Sayers, E. W. (2016). GenBank. *Nucleic Acids Research*, *44* (Database issue), D67–D72.
- Connors, E. (2015). Coal-ash management by U.S. electric utilities: Overview and recent developments. *Utilities Policy*, *34*, 30–33.
- Daly, K., Sharp, R. J., & McCarthy, A. J. (2000). Development of oligonucleotide probes and PCR primers for detecting phylogenetic subgroups of sulfate-reducing bacteria. *Microbiology*, *146*(7), 1693–1705.

- Dash, H. R., & Das, S. (2014). Bioremediation Potential of Mercury by *Bacillus* Species Isolated from Marine Environment and Wastes of Steel Industry. *Bioremediation Journal*, 18(3), 204–212.
- Deonaraine, A., Bartov, G., Johnson, T. M., Ruhl, L., Vengosh, A., & Hsu-Kim, H. (2013). Environmental Impacts of the Tennessee Valley Authority Kingston Coal Ash Spill. 2. Effect of Coal Ash on Methylmercury in Historically Contaminated River Sediments. *Environmental Science & Technology*, 47(4), 2100–2108.
- Driscoll, C. T., Mason, R. P., Chan, H. M., Jacob, D. J., & Pirrone, N. (2013). Mercury as a Global Pollutant: Sources, Pathways, and Effects. *Environmental Science & Technology*, 47(10), 4967–4983.
- Earl, A. M., Losick, R., & Kolter, R. (2008). Ecology and genomics of *Bacillus subtilis*. *Trends in Microbiology*, 16(6), 269–275.
- Edgar, R. C. (2004). MUSCLE: Multiple sequence alignment with high accuracy and high throughput. *Nucleic Acids Research*, 32(5), 1792–1797.
- Edwards, U., Rogall, T., Blöcker, H., Emde, M., & Böttger, E. C. (1989). Isolation and direct complete nucleotide determination of entire genes. Characterization of a gene coding for 16S ribosomal RNA. *Nucleic Acids Research*, 17(19), 7843–7853.
- Ehrlich, H. L., & Newman, D. K. (2008). *Geomicrobiology, Fifth Edition*. Boca Raton: CRC Press.
- Energy Information Administration (2018). Table 1.1. Net Generation by Energy Source: Total (All Sectors), 2008-September 2018. In *Electric Power Monthly (September 2018)*.
- Geets, J., Borremans, B., Diels, L., Springael, D., Vangronsveld, J., van der Lelie, D., & Vanbroekhoven, K. (2006). DsrB gene-based DGGE for community and diversity surveys of sulfate-reducing bacteria. *Journal of Microbiological Methods*, 66(2), 194–205.
- Greely Jr., M. S., Elmore, L. R., McCracken, M. K., & Sherrard, R. M. (2014). Effects of Sediment Containing Coal Ash from the Kingston Ash Release on Embryo-Larval Development in the Fathead Minnow, *Pimephales promelas* (Rafinesque, 1820). *Bulletin of Environmental Contamination and Toxicology*, 92(2), 154–159.
- Hershey, A., Rublee, P., & Tsui, M. (2016). Heavy metal analysis, gene proxies, and stable isotope tracers of coal ash contamination in the dan river food web. Tech. Rep. 463, Water Resources Research Institute of The University of North Carolina.

- Hesterberg, D., Polizzotto, M., & Crozier, C. (2014). Assessment of trace-element impacts on agricultural use of the water from the Dan River following the Eden coal ash spill. Tech. rep., N.C. State University.
- Hughes, M. N., & Poole, R. K. (1991). Metal speciation and microbial growth-the hard (and soft) facts. *Journal of General Microbiology*, 137, 725–734.
- Jayaranjan, M. L. D., van Hullebusch, E. D., & Annachhatre, A. P. (2014). Reuse options for coal fired power plant bottom ash and fly ash. *Reviews in Environmental Science and Biotechnology*, 13(4), 467–486.
- Jukes, T., & Cantor, C. (1969). Evolution of protein molecules. In M. HN (Ed.) *Mammalian Protein Metabolism*, (pp. 21–132). New York: Academic Press.
- Keshri, J., Mankazana, B. B. J., & Momba, M. N. B. (2014). Profile of bacterial communities in South African mine-water samples using Illumina next-generation sequencing platform. *Applied Microbiology and Biotechnology*, 99(7), 3233–3242.
- Kisku, G. C., Kumar, V., Sahu, P., Kumar, P., & Kumar, N. (2018). Characterization of coal fly ash and use of plants growing in ash pond for phytoremediation of metals from contaminated agricultural land. *International Journal of Phytoremediation*, 20(4), 330–337.
- Klubek, B., Carison, C. L., Oliver, J., & Adriano, D. C. (1992). Characterization of microbial abundance and activity from three coal ash basins. *Soil Biology and Biochemistry*, 24(11), 1119–1125.
- Kumar, S., Stecher, G., & Tamura, K. (2016). MEGA7: Molecular Evolutionary Genetics Analysis Version 7.0 for Bigger Datasets. *Molecular Biology and Evolution*, 33(7), 1870–1874.
- Lemly, A. (2015). Damage cost of the Dan River coal ash spill. *Environmental Pollution*, 197, 55–61.
- Lima de Silva, A. A., de Carvalho, M. A. R., de Souza, S. A. L., Dias, P. M. T., da Silva Filho, R. G., de Meirelles Saramago, C. S., de Melo Bento, C. A., & Hofer, E. (2012). Heavy metal tolerance (Cr, Ag AND Hg) in bacteria isolated from sewage. *Brazilian Journal of Microbiology*, 43(4), 1620–1631.
- Liu, Y.-R., Yu, R.-Q., Zheng, Y.-M., & He, J.-Z. (2014). Analysis of the Microbial Community Structure by Monitoring an Hg Methylation Gene (hgcA) in Paddy Soils along an Hg Gradient. *Applied and Environmental Microbiology*, 80(9), 2874–2879.

- Mongodin, E. F., Shapir, N., Daugherty, S. C., DeBoy, R. T., Emerson, J. B., Shvartzbeyn, A., Radune, D., Vamathevan, J., Riggs, F., Grinberg, V., Khouri, H., Wackett, L. P., Nelson, K. E., & Sadowsky, M. J. (2006). Secrets of Soil Survival Revealed by the Genome Sequence of *Arthrobacter aurescens* TC1. *PLOS Genetics*, 2(12), e214.
- Naik, M. M., & Dubey, S. K. (2013). Lead resistant bacteria: Lead resistance mechanisms, their applications in lead bioremediation and biomonitoring. *Ecotoxicology and Environmental Safety*, 98, 1–7.
- NC DEQ (2014). Rapid response timeline of events. Tech. rep., NC DEQ.
- Otter, R. R., Bailey, F. C., Fortner, A. M., & Adams, S. M. (2012). Trophic status and metal bioaccumulation differences in multiple fish species exposed to coal ash-associated metals. *Ecotoxicology and Environmental Safety*, 85, 30–36.
- Pangayao, D., Gallardo, S., Promentilla, M., & Van Hullebusch, E. (2018). Bioleaching of trace metals from coal ash using local isolate from coal ash ponds. In *MATEC Web of Conferences*, vol. 156.
- Parks, J., Johs, A., Podar, M., Bridou, R., Hurt Jr., R., Smith, S., Tomanicek, S., Qian, Y., Brown, S., Brandt, C., Palumbo, A., Smith, J., Wall, J., Elias, D., & Liang, L. (2013). The genetic basis for bacterial mercury methylation. *Science*, 339(6125), 1332–1335.
- Pepi, M., Gaggi, C., Bernardini, E., Focardi, S., Lobianco, A., Ruta, M., Nicolardi, V., Volterrani, M., Gasperini, S., Trinchera, G., Renzi, P., Gabellini, M., & Focardi, S. E. (2011). Mercury-resistant bacterial strains *Pseudomonas* and *Psychrobacter* spp. isolated from sediments of Orbetello Lagoon (Italy) and their possible use in bioremediation processes. *International Biodeterioration & Biodegradation*, 65(1), 85–91.
- Poulain, A. J., & Barkay, T. (2013). Cracking the Mercury Methylation Code. *Science*, 339(6125), 1280–1281.
- R Core Team (2019). R: A language and environment for statistical computing.
- Raja, C. E., & Omine, K. (2012). Characterization of boron tolerant bacteria isolated from a fly ash dumping site for bacterial boron remediation. *Environmental Geochemistry and Health*, 35(4), 431–438.
- Ramamoorthy, S., & Kushner, D. J. (1975). Binding of mercuric and other heavy metal ions by microbial growth media. *Microbial Ecology*, 2(2), 162–176.

- Rowe, C. L. (2014). Bioaccumulation and effects of metals and trace elements from aquatic disposal of coal combustion residues: Recent advances and recommendations for further study. *Science of The Total Environment*, 485, 490–496.
- Roychowdhury, R., Roy, M., Rakshit, A., Sarkar, S., & Mukherjee, P. (2018). Arsenic Bioremediation by Indigenous Heavy Metal Resistant Bacteria of Fly Ash Pond. *Bulletin of Environmental Contamination and Toxicology*, 101(4), 527–535.
- Ruhl, L., Vengosh, A., Dwyer, G. S., Hsu-Kim, H., & Deonaraine, A. (2010). Environmental impacts of the coal ash spill in Kingston, Tennessee: An 18-month survey. *Environmental Science & Technology*, 44(24), 9272–9278.
- Ruhl, L., Vengosh, A., Dwyer, G. S., Hsu-Kim, H., Deonaraine, A., Bergin, M., & Kravchenko, J. (2009). Survey of the potential environmental and health impacts in the immediate aftermath of the coal ash spill in Kingston, Tennessee. *Environmental Science & Technology*, 43(16), 6326–6333.
- Schaefer, J. K., Kronberg, R.-M., Morel, F. M. M., & Skyllberg, U. (2014). Detection of a key Hg methylation gene, *hgcA*, in wetland soils. *Environmental Microbiology Reports*, 6(5), 441–447.
- Schaefer, J. K., Rocks, S. S., Zheng, W., Liang, L., Gu, B., & Morel, F. M. M. (2011). Active transport, substrate specificity, and methylation of Hg(II) in anaerobic bacteria. *PNAS*, 108(21), 8714–8719.
- Schwartz, G. E., Redfern, L. K., Ikuma, K., Gunsch, C. K., Ruhl, L. S., Vengosh, A., & Hsu-Kim, H. (2016a). Impacts of coal ash on methylmercury production and the methylating microbial community in anaerobic sediment slurries. *Environmental Science: Processes & Impacts*, 18(11), 1427–1439.
- Schwartz, G. E., Redfern, L. K., Ikuma, K., Gunsch, C. K., Ruhl, L. S., Vengosh, A., & Hsu-Kim, H. (2016b). Impacts of coal ash on methylmercury production and the methylating microbial community in anaerobic sediment slurries. *Environmental Science: Processes & Impacts*, 18(11), 1427–1439.
- Shaheen, S. M., Hooda, P. S., & Tsadilas, C. D. (2014). Opportunities and challenges in the use of coal fly ash for soil improvements – A review. *Journal of Environmental Management*, 145, 249–267.
- Stepanauskas, R., Glenn, T. C., Jagoe, C. H., Tuckfield, R. C., Lindell, A. H., & McArthur, J. (2005). Elevated Microbial Tolerance to Metals and Antibiotics in Metal-Contaminated Industrial Environments. *Environmental Science & Technology*, 39(10), 3671–3678.

- Stewart, C. N., & Via, L. E. (1993). A rapid CTAB DNA isolation technique useful for RAPD fingerprinting and other PCR applications. *BioTechniques*, 14(5), 748–750.
- Tsui, M. T. K., Blum, J. D., Finlay, J. C., Balogh, S. J., Kwon, S. Y., & Nollet, Y. H. (2013). Photodegradation of methylmercury in stream ecosystems. *Limnology and Oceanography*, 58(1), 13–22.
- TVA (2011). Fact Sheet Kingston Ash Recovery Project: December 7, 2011. Tech. rep., TVA.
- US EPA (2014). EPA’s Response to the Duke Energy Coal Ash Spill in Eden, NC. <http://www.epa.gov/region4/duke-energy/>. EPA’s Response to the Duke Energy Coal Ash Spill in Eden, NC.
- Wagner, M., Roger, A. J., Flax, J. L., Brusseau, G. A., & Stahl, D. A. (1998). Phylogeny of dissimilatory sulfite reductases supports an early origin of sulfate respiration. *Journal of Bacteriology*, 180(11), 2975–2982.
- Zwietering, M. H., Jongenburger, I., Rombouts, F. M., & van ’t Riet, K. (1990). Modeling of the Bacterial Growth Curve. *Applied and Environmental Microbiology*, 56(6), 1875–1881.

APPENDIX A
SUPPLEMENTAL TABLES

Table A.1. DR 13 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.031	0.064	0.431	0.997
As	0	2	0.032	0.065	0.427	0.994
As	0	3	0.030	0.064	0.428	0.996
As	1	1	0.034	0.060	0.460	0.996
As	1	2	0.034	0.055	0.477	0.993
As	1	3	0.032	0.061	0.459	0.996
As	2	1	0.035	0.059	0.471	0.994
As	2	2	0.034	0.055	0.483	0.994
As	2	3	0.034	0.061	0.457	0.995
As	3	1	0.034	0.062	0.453	0.994
As	3	2	0.034	0.059	0.473	0.995
As	3	3	0.034	0.060	0.458	0.994
As	4	1	0.036	0.062	0.482	0.995
As	4	2	0.035	0.060	0.449	0.994
As	4	3	0.030	0.069	0.439	0.995
As	5	1	0.035	0.063	0.471	0.994
As	5	2	0.035	0.063	0.459	0.994
As	5	3	0.034	0.066	0.434	0.994
Cd	0	1	0.027	0.072	0.406	0.995
Cd	0	2	0.029	0.066	0.447	0.993
Cd	0	3	0.030	0.060	0.496	0.993
Cd	1	1	0.028	0.069	0.431	0.997
Cd	1	2	0.029	0.064	0.478	0.996
Cd	1	3	0.031	0.061	0.481	0.994

Table A.1. DR 13 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	2	1	0.031	0.065	0.450	0.993
Cd	2	2	0.030	0.065	0.450	0.995
Cd	2	3	0.030	0.066	0.452	0.994
Cd	3	1	0.031	0.066	0.442	0.991
Cd	3	2	0.030	0.067	0.435	0.994
Cd	3	3	0.033	0.062	0.453	0.992
Cd	4	1	0.027	0.063	0.395	0.994
Cd	4	2	0.028	0.061	0.421	0.994
Cd	4	3	0.030	0.060	0.422	0.993
Cd	5	1	0.017	0.062	0.385	0.905
Cd	5	2	0.023	0.050	0.364	0.997
Cd	5	3	0.024	0.053	0.343	0.997
Cr	0	1	0.028	0.063	0.445	0.995
Cr	0	2	0.033	0.063	0.439	0.994
Cr	0	3	0.034	0.070	0.440	0.995
Cr	1	1	0.035	0.060	0.471	0.997
Cr	1	2	0.033	0.064	0.434	0.996
Cr	1	3	0.033	0.063	0.439	0.996
Cr	2	1	0.032	0.062	0.438	0.997
Cr	2	2	0.044	0.059	0.443	0.998
Cr	2	3	0.033	0.065	0.411	0.994
Cr	3	1	0.033	0.062	0.433	0.995
Cr	3	2	0.035	0.061	0.446	0.994
Cr	3	3	0.029	0.085	0.337	0.993
Cr	4	1	0.032	0.059	0.461	0.993
Cr	4	2	0.034	0.062	0.452	0.994
Cr	4	3	0.032	0.062	0.452	0.996

Table A.1. DR 13 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cr	5	1	0.033	0.060	0.444	0.994
Cr	5	2	0.034	0.061	0.454	0.995
Cr	5	3	0.034	0.062	0.446	0.996
Hg	0	1	0.030	0.066	0.430	0.994
Hg	0	2	0.033	0.066	0.407	0.993
Hg	0	3	0.030	0.063	0.429	0.994
Hg	1	1	0.032	0.068	0.423	0.995
Hg	1	2	0.034	0.062	0.430	0.993
Hg	1	3	0.032	0.066	0.465	0.996
Hg	2	1	0.034	0.064	0.438	0.995
Hg	2	2	0.033	0.064	0.438	0.994
Hg	2	3	0.035	0.061	0.451	0.993
Hg	3	1	0.033	0.065	0.467	0.995
Hg	3	2	0.035	0.063	0.458	0.993
Hg	3	3	0.033	0.064	0.459	0.994
Hg	4	1	0.034	0.067	0.455	0.994
Hg	4	2	0.036	0.060	0.465	0.992
Hg	4	3	0.034	0.064	0.432	0.993
Hg	5	1	0.032	0.072	0.434	0.994
Hg	5	2	0.034	0.064	0.448	0.993
Hg	5	3	0.034	0.064	0.447	0.993
Pb	0	1	0.030	0.067	0.435	0.995
Pb	0	2	0.030	0.064	0.463	0.995
Pb	0	3	0.029	0.067	0.443	0.995
Pb	1	1	0.029	0.071	0.423	0.996
Pb	1	2	0.030	0.069	0.430	0.995
Pb	1	3	0.032	0.063	0.474	0.995

Table A.1. DR 13 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	2	1	0.030	0.067	0.466	0.995
Pb	2	2	0.030	0.068	0.445	0.996
Pb	2	3	0.024	0.072	0.500	0.980
Pb	3	1	0.030	0.067	0.424	0.995
Pb	3	2	0.032	0.060	0.463	0.994
Pb	3	3	0.030	0.069	0.411	0.994
Pb	4	1	0.032	0.063	0.448	0.993
Pb	4	2	0.031	0.064	0.425	0.993
Pb	4	3	0.032	0.063	0.449	0.994
Pb	5	1	0.030	0.068	0.407	0.994
Pb	5	2	0.031	0.065	0.428	0.993
Pb	5	3	0.032	0.066	0.432	0.994
Se	0	1	0.007	0.070	0.360	0.995
Se	0	2	0.008	0.072	0.351	0.989
Se	0	3	0.006	0.069	0.372	0.995
Se	1	1	0.008	0.062	0.391	0.993
Se	1	2	0.007	0.067	0.341	0.992
Se	1	3	0.008	0.064	0.377	0.989
Se	2	1	0.006	0.076	0.328	0.996
Se	2	2	0.008	0.055	0.424	0.995
Se	2	3	0.008	0.055	0.476	0.995
Se	3	1	0.006	0.066	0.411	0.998
Se	3	2	0.008	0.060	0.383	0.995
Se	3	3	0.009	0.056	0.421	0.994
Se	4	1	0.007	0.062	0.413	0.995
Se	4	2	0.009	0.055	0.440	0.991
Se	4	3	0.009	0.059	0.407	0.991

Table A.1. DR 13 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Se	5	1	0.007	0.049	0.497	0.991
Se	5	2	0.005	0.055	0.433	0.996
Se	5	3	0.004	0.056	0.438	0.996
Zn	0	1	0.032	0.071	0.416	0.993
Zn	0	2	0.033	0.066	0.427	0.992
Zn	0	3	0.034	0.066	0.420	0.992
Zn	1	1	0.032	0.066	0.421	0.993
Zn	1	2	0.034	0.063	0.461	0.992
Zn	1	3	0.034	0.064	0.429	0.990
Zn	2	1	0.032	0.064	0.435	0.994
Zn	2	2	0.032	0.066	0.447	0.994
Zn	2	3	0.032	0.068	0.432	0.994
Zn	3	1	0.031	0.067	0.430	0.994
Zn	3	2	0.032	0.068	0.431	0.993
Zn	3	3	0.031	0.069	0.419	0.994
Zn	4	1	0.029	0.071	0.426	0.995
Zn	4	2	0.031	0.071	0.412	0.993
Zn	4	3	0.031	0.071	0.419	0.993
Zn	5	1	0.030	0.059	0.410	0.993
Zn	5	2	0.028	0.060	0.389	0.993
Zn	5	3	0.025	0.064	0.377	0.994

Table A.2. DR 24 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.032	0.103	0.312	0.985
As	0	2	0.030	0.092	0.269	0.989
As	0	3	0.030	0.098	0.334	0.988
As	1	1	0.028	0.101	0.272	0.984
As	1	2	0.032	0.087	0.286	0.989
As	1	3	0.032	0.087	0.283	0.990
As	2	1	0.033	0.087	0.263	0.944
As	2	2	0.030	0.090	0.276	0.990
As	2	3	0.031	0.091	0.280	0.989
As	3	1	0.028	0.100	0.268	0.989
As	3	2	0.031	0.091	0.259	0.952
As	3	3	0.030	0.090	0.284	0.989
As	4	1	0.030	0.097	0.281	0.992
As	4	2	0.034	0.084	0.311	0.986
As	4	3	0.031	0.092	0.288	0.989
As	5	1	0.028	0.102	0.288	0.989
As	5	2	0.029	0.092	0.286	0.985
As	5	3	0.030	0.097	0.287	0.988
Cd	0	1	0.032	0.096	0.277	0.995
Cd	0	2	0.030	0.093	0.282	0.990
Cd	0	3	0.030	0.094	0.269	0.983
Cd	1	1	0.031	0.089	0.286	0.988
Cd	1	2	0.030	0.088	0.286	0.987
Cd	1	3	0.029	0.093	0.274	0.989
Cd	2	1	0.031	0.092	0.286	0.988
Cd	2	2	0.030	0.091	0.279	0.988
Cd	2	3	0.029	0.092	0.284	0.990

Table A.2. DR 24 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.029	0.094	0.288	0.990
Cd	3	2	0.030	0.096	0.271	0.990
Cd	3	3	0.030	0.088	0.266	0.987
Cd	4	1	0.030	0.096	0.255	0.985
Cd	4	2	0.028	0.104	0.217	0.984
Cd	4	3	0.028	0.100	0.222	0.990
Cd	5	1	0.016	0.175	0.151	0.991
Cd	5	2	0.018	0.152	0.163	0.997
Cd	5	3	0.018	0.160	0.159	0.999
Cr	0	1	0.029	0.095	0.275	0.989
Cr	0	2	0.030	0.092	0.278	0.987
Cr	0	3	0.032	0.097	0.288	0.988
Cr	1	1	0.030	0.096	0.271	0.989
Cr	1	2	0.033	0.086	0.305	0.989
Cr	1	3	0.032	0.095	0.256	0.948
Cr	2	1	0.031	0.091	0.281	0.989
Cr	2	2	0.030	0.095	0.276	0.989
Cr	2	3	0.032	0.094	0.292	0.987
Cr	3	1	0.030	0.096	0.274	0.989
Cr	3	2	0.034	0.080	0.325	0.981
Cr	3	3	0.035	0.083	0.306	0.986
Cr	4	1	0.031	0.100	0.251	0.946
Cr	4	2	0.031	0.094	0.281	0.989
Cr	4	3	0.030	0.096	0.276	0.989
Cr	5	1	0.029	0.101	0.277	0.987
Cr	5	2	0.031	0.105	0.273	0.984
Cr	5	3	0.028	0.105	0.278	0.989

Table A.2. DR 24 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.027	0.093	0.263	0.991
Hg	0	2	0.031	0.088	0.283	0.990
Hg	0	3	0.032	0.087	0.293	0.987
Hg	1	1	0.030	0.097	0.271	0.992
Hg	1	2	0.030	0.090	0.282	0.989
Hg	1	3	0.031	0.087	0.291	0.987
Hg	2	1	0.028	0.100	0.274	0.990
Hg	2	2	0.029	0.099	0.276	0.988
Hg	2	3	0.030	0.094	0.281	0.988
Hg	3	1	0.029	0.096	0.286	0.989
Hg	3	2	0.029	0.093	0.277	0.991
Hg	3	3	0.031	0.094	0.287	0.990
Hg	4	1	0.030	0.095	0.280	0.990
Hg	4	2	0.030	0.095	0.275	0.988
Hg	4	3	0.030	0.090	0.279	0.986
Hg	5	1	0.030	0.095	0.279	0.987
Hg	5	2	0.033	0.084	0.303	0.983
Hg	5	3	0.033	0.078	0.330	0.987
Pb	0	1	0.032	0.088	0.281	0.985
Pb	0	2	0.031	0.096	0.330	0.989
Pb	0	3	0.031	0.082	0.344	0.991
Pb	1	1	0.031	0.083	0.281	0.990
Pb	1	2	0.032	0.082	0.310	0.989
Pb	1	3	0.031	0.084	0.297	0.990
Pb	2	1	0.036	0.078	0.388	0.988
Pb	2	2	0.034	0.090	0.343	0.991
Pb	2	3	0.032	0.082	0.307	0.991

Table A.2. DR 24 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.029	0.087	0.305	0.991
Pb	3	2	0.032	0.082	0.312	0.990
Pb	3	3	0.031	0.089	0.283	0.991
Pb	4	1	0.031	0.087	0.292	0.985
Pb	4	2	0.030	0.090	0.284	0.988
Pb	4	3	0.031	0.091	0.287	0.984
Pb	5	1	0.032	0.080	0.302	0.983
Pb	5	2	0.031	0.087	0.277	0.986
Pb	5	3	0.029	0.092	0.266	0.987
Zn	0	1	0.029	0.089	0.276	0.985
Zn	0	2	0.029	0.097	0.273	0.989
Zn	0	3	0.029	0.090	0.264	0.984
Zn	1	1	0.033	0.084	0.306	0.986
Zn	1	2	0.031	0.085	0.276	0.985
Zn	1	3	0.031	0.093	0.287	0.986
Zn	2	1	0.031	0.082	0.322	0.985
Zn	2	2	0.034	0.075	0.332	0.983
Zn	2	3	0.034	0.080	0.315	0.985
Zn	3	1	0.032	0.083	0.293	0.986
Zn	3	2	0.036	0.075	0.340	0.986
Zn	3	3	0.032	0.082	0.308	0.989
Zn	4	1	0.028	0.098	0.275	0.988
Zn	4	2	0.030	0.089	0.324	0.990
Zn	4	3	0.031	0.085	0.319	0.988
Zn	5	1	0.026	0.095	0.212	0.990
Zn	5	2	0.026	0.100	0.208	0.991
Zn	5	3	0.025	0.096	0.215	0.993

Table A.3. DR 26 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.013	0.037	0.011	0.047
As	0	2	0.016	0.056	0.357	0.999
As	0	3	0.013	0.059	0.334	1.000
As	1	1	0.016	0.073	0.279	0.992
As	1	2	0.017	0.061	0.330	0.998
As	1	3	0.017	0.064	0.333	0.996
As	2	1	0.016	0.008	0.500	0.721
As	2	2	0.014	0.063	0.334	0.997
As	2	3	0.014	0.060	0.353	0.996
As	3	1	0.018	0.063	0.332	0.996
As	3	2	0.010	0.000	0.500	0.024
As	3	3	0.017	0.070	0.293	0.995
As	4	1	0.016	0.014	0.500	0.871
As	4	2	0.016	0.064	0.313	0.999
As	4	3	0.016	0.064	0.344	0.998
As	5	1	0.016	0.243	0.023	0.580
As	5	2	0.016	0.061	0.350	0.998
As	5	3	0.017	0.064	0.334	0.998
Cd	0	1	0.018	0.057	0.271	0.990
Cd	0	2	0.018	0.067	0.243	0.992
Cd	0	3	0.018	0.071	0.234	0.993
Cd	1	1	0.018	0.066	0.255	0.991
Cd	1	2	0.019	0.064	0.252	0.992
Cd	1	3	0.022	0.051	0.368	0.989
Cd	2	1	0.018	0.068	0.244	0.991
Cd	2	2	0.019	0.069	0.244	0.991
Cd	2	3	0.021	0.064	0.246	0.988

Table A.3. DR 26 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.020	0.056	0.295	0.984
Cd	3	2	0.018	0.063	0.244	0.989
Cd	3	3	0.019	0.062	0.264	0.987
Cd	4	1	0.018	0.080	0.161	0.986
Cd	4	2	0.021	0.059	0.215	0.963
Cd	4	3	0.017	0.085	0.157	0.980
Cd	5	1	0.014	0.077	0.076	0.976
Cd	5	2	0.016	0.071	0.086	0.980
Cd	5	3	0.015	0.069	0.077	0.988
Cr	0	1	0.014	0.086	0.011	0.100
Cr	0	2	0.017	0.039	0.500	0.995
Cr	0	3	0.017	0.066	0.299	0.997
Cr	1	1	0.033	0.038	0.454	0.908
Cr	1	2	0.014	0.011	0.500	0.763
Cr	1	3	0.012	0.542	0.012	0.010
Cr	2	1	0.015	0.000	0.000	0.025
Cr	2	2	0.015	0.001	0.000	0.157
Cr	2	3	0.010	0.001	0.500	0.085
Cr	3	1	0.011	0.001	0.500	0.079
Cr	3	2	0.012	-0.001	0.500	0.178
Cr	3	3	0.011	0.001	0.500	0.096
Cr	4	1	0.014	0.600	0.011	0.346
Cr	4	2	0.013	-0.003	0.500	0.290
Cr	4	3	0.012	0.000	0.482	0.001
Cr	5	1	0.011	0.002	0.500	0.134
Cr	5	2	0.011	0.000	0.499	0.003
Cr	5	3	0.013	0.000	0.500	0.016

Table A.3. DR 26 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.009	0.003	0.500	0.305
Hg	0	2	0.016	0.075	0.331	0.993
Hg	0	3	0.018	0.051	0.460	0.994
Hg	1	1	0.004	0.116	0.188	0.894
Hg	1	2	0.017	0.067	0.346	0.999
Hg	1	3	0.017	0.061	0.361	0.998
Hg	2	1	0.013	0.030	0.500	0.985
Hg	2	2	0.014	0.501	0.017	0.495
Hg	2	3	0.016	0.011	0.500	0.797
Hg	3	1	0.018	0.055	0.407	0.992
Hg	3	2	0.017	0.061	0.357	0.998
Hg	3	3	0.016	0.055	0.371	0.999
Hg	4	1	0.017	0.068	0.319	0.996
Hg	4	2	0.015	0.069	0.316	0.994
Hg	4	3	0.016	0.061	0.343	0.998
Hg	5	1	0.013	0.004	0.500	0.771
Hg	5	2	0.010	0.067	0.288	0.999
Hg	5	3	0.011	0.010	0.500	0.895
Pb	0	1	0.018	0.073	0.244	0.989
Pb	0	2	0.019	0.071	0.235	0.989
Pb	0	3	0.019	0.068	0.239	0.992
Pb	1	1	0.019	0.066	0.266	0.991
Pb	1	2	0.020	0.054	0.316	0.987
Pb	1	3	0.018	0.075	0.240	0.991
Pb	2	1	0.020	0.054	0.312	0.987
Pb	2	2	0.021	0.049	0.369	0.986
Pb	2	3	0.018	0.077	0.221	0.989

Table A.3. DR 26 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.021	0.049	0.352	0.987
Pb	3	2	0.020	0.058	0.285	0.986
Pb	3	3	0.023	0.046	0.384	0.981
Pb	4	1	0.018	0.071	0.234	0.989
Pb	4	2	0.019	0.071	0.239	0.991
Pb	4	3	0.020	0.067	0.253	0.988
Pb	5	1	0.018	0.076	0.238	0.991
Pb	5	2	0.018	0.074	0.235	0.990
Pb	5	3	0.018	0.076	0.238	0.973
Se	0	1	0.000	0.086	0.304	0.998
Se	0	2	0.000	0.073	0.316	0.998
Se	0	3	0.000	0.081	0.318	0.998
Se	1	1	0.000	0.076	0.327	0.997
Se	1	2	0.000	0.080	0.297	0.998
Se	1	3	0.000	0.073	0.313	0.997
Se	2	1	0.000	0.071	0.283	0.998
Se	2	2	0.000	0.080	0.296	0.998
Se	2	3	0.000	0.066	0.338	0.996
Se	3	1	0.000	0.122	0.119	0.963
Se	3	2	0.000	0.118	0.140	0.974
Se	3	3	0.000	0.123	0.118	0.969
Se	4	1	0.000	0.132	0.036	0.959
Se	4	2	0.000	0.128	0.038	0.969
Se	4	3	0.000	0.126	0.039	0.965
Se	5	1	0.000	0.041	0.500	0.973
Se	5	2	0.000	0.019	0.500	0.983
Se	5	3	0.000	0.042	0.500	0.961

Table A.3. DR 26 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.014	-0.001	0.500	0.009
Zn	0	2	0.019	0.068	0.240	0.991
Zn	0	3	0.019	0.069	0.238	0.991
Zn	1	1	0.019	0.070	0.241	0.991
Zn	1	2	0.019	0.076	0.221	0.994
Zn	1	3	0.019	0.075	0.234	0.989
Zn	2	1	0.020	0.068	0.248	0.988
Zn	2	2	0.018	0.076	0.230	0.985
Zn	2	3	0.020	0.071	0.241	0.992
Zn	3	1	0.019	0.072	0.264	0.993
Zn	3	2	0.020	0.069	0.241	0.989
Zn	3	3	0.022	0.069	0.285	0.990
Zn	4	1	0.021	0.071	0.273	0.990
Zn	4	2	0.019	0.078	0.222	0.989
Zn	4	3	0.019	0.072	0.240	0.992
Zn	5	1	0.019	0.065	0.158	0.985
Zn	5	2	0.016	0.079	0.141	0.988
Zn	5	3	0.018	0.061	0.179	0.985

Table A.4. DR 3.2 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.028	0.065	0.457	0.995
As	0	2	0.036	0.053	0.500	0.986
As	0	3	0.032	0.061	0.443	0.985
As	1	1	0.030	0.066	0.422	0.995
As	1	2	0.033	0.052	0.500	0.990
As	1	3	0.033	0.052	0.500	0.987
As	2	1	0.030	0.060	0.457	0.996
As	2	2	0.035	0.052	0.500	0.991
As	2	3	0.032	0.057	0.462	0.988
As	3	1	0.033	0.053	0.485	0.992
As	3	2	0.032	0.053	0.487	0.992
As	3	3	0.032	0.057	0.466	0.994
As	4	1	0.033	0.055	0.488	0.991
As	4	2	0.034	0.056	0.471	0.992
As	4	3	0.029	0.068	0.441	0.996
As	5	1	0.034	0.065	0.443	0.981
As	5	2	0.030	0.080	0.403	0.992
As	5	3	0.034	0.069	0.428	0.991
Cd	0	1	0.024	0.089	0.352	0.994
Cd	0	2	0.020	0.106	0.352	0.997
Cd	0	3	0.022	0.096	0.342	0.995
Cd	1	1	0.024	0.077	0.363	0.990
Cd	1	2	0.024	0.075	0.384	0.980
Cd	1	3	0.028	0.066	0.432	0.995
Cd	2	1	0.025	0.074	0.376	0.993
Cd	2	2	0.027	0.078	0.361	0.986
Cd	2	3	0.027	0.071	0.408	0.993

Table A.4. DR 3.2 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.025	0.076	0.351	0.997
Cd	3	2	0.029	0.074	0.347	0.993
Cd	3	3	0.027	0.078	0.344	0.993
Cd	4	1	0.030	0.067	0.346	0.990
Cd	4	2	0.037	0.045	0.449	0.982
Cd	4	3	0.032	0.063	0.370	0.989
Cd	5	1	0.030	0.062	0.176	0.949
Cd	5	2	0.027	0.095	0.141	0.969
Cd	5	3	0.026	0.080	0.162	0.958
Cr	0	1	0.031	0.058	0.472	0.996
Cr	0	2	0.029	0.064	0.438	0.996
Cr	0	3	0.033	0.064	0.414	0.982
Cr	1	1	0.034	0.052	0.481	0.988
Cr	1	2	0.032	0.059	0.440	0.989
Cr	1	3	0.034	0.064	0.419	0.985
Cr	2	1	0.033	0.054	0.466	0.990
Cr	2	2	0.034	0.059	0.461	0.993
Cr	2	3	0.034	0.058	0.441	0.986
Cr	3	1	0.032	0.053	0.469	0.991
Cr	3	2	0.032	0.061	0.426	0.991
Cr	3	3	0.032	0.073	0.427	0.993
Cr	4	1	0.034	0.061	0.441	0.986
Cr	4	2	0.033	0.065	0.427	0.987
Cr	4	3	0.033	0.059	0.454	0.989
Cr	5	1	0.035	0.067	0.438	0.987
Cr	5	2	0.033	0.068	0.433	0.990
Cr	5	3	0.034	0.074	0.416	0.991

Table A.4. DR 3.2 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.029	0.071	0.459	0.995
Hg	0	2	0.032	0.066	0.465	0.992
Hg	0	3	0.031	0.059	0.471	0.992
Hg	1	1	0.029	0.060	0.489	0.997
Hg	1	2	0.030	0.058	0.492	0.997
Hg	1	3	0.032	0.057	0.479	0.993
Hg	2	1	0.032	0.058	0.485	0.995
Hg	2	2	0.031	0.053	0.500	0.995
Hg	2	3	0.031	0.052	0.500	0.990
Hg	3	1	0.032	0.055	0.476	0.990
Hg	3	2	0.032	0.052	0.500	0.988
Hg	3	3	0.029	0.056	0.482	0.995
Hg	4	1	0.032	0.066	0.422	0.986
Hg	4	2	0.034	0.051	0.497	0.986
Hg	4	3	0.028	0.063	0.490	0.995
Hg	5	1	0.032	0.068	0.448	0.987
Hg	5	2	0.033	0.067	0.440	0.987
Hg	5	3	0.031	0.061	0.474	0.995
Pb	0	1	0.026	0.084	0.360	0.984
Pb	0	2	0.022	0.086	0.366	0.981
Pb	0	3	0.023	0.092	0.357	0.994
Pb	1	1	0.026	0.083	0.345	0.998
Pb	1	2	0.025	0.080	0.375	0.997
Pb	1	3	0.028	0.077	0.355	0.992
Pb	2	1	0.026	0.072	0.393	0.996
Pb	2	2	0.028	0.070	0.380	0.993
Pb	2	3	0.026	0.079	0.356	0.998

Table A.4. DR 3.2 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.023	0.080	0.361	0.995
Pb	3	2	0.024	0.086	0.351	0.998
Pb	3	3	0.026	0.085	0.354	0.978
Pb	4	1	0.028	0.067	0.386	0.997
Pb	4	2	0.022	0.092	0.336	0.997
Pb	4	3	0.030	0.079	0.358	0.996
Pb	5	1	0.029	0.079	0.360	0.991
Pb	5	2	0.028	0.077	0.365	0.989
Pb	5	3	0.025	0.094	0.354	0.997
Se	0	1	0.012	0.070	0.500	0.998
Se	0	2	0.018	0.069	0.460	0.999
Se	0	3	0.015	0.073	0.500	0.993
Se	1	1	0.017	0.067	0.493	0.997
Se	1	2	0.008	0.077	0.500	0.988
Se	1	3	0.019	0.063	0.500	0.998
Se	2	1	0.020	0.079	0.428	0.997
Se	2	2	0.016	0.068	0.500	0.993
Se	2	3	0.018	0.064	0.500	0.994
Se	3	1	0.019	0.072	0.320	1.000
Se	3	2	0.022	0.068	0.335	0.989
Se	3	3	0.017	0.074	0.345	0.990
Se	4	1	0.021	0.074	0.234	0.997
Se	4	2	0.015	0.103	0.236	0.983
Se	4	3	0.019	0.091	0.235	0.994
Se	5	1	0.013	0.126	0.133	0.944
Se	5	2	0.015	0.121	0.128	0.971
Se	5	3	0.017	0.113	0.137	0.970

Table A.4. DR 3.2 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.028	0.071	0.389	0.985
Zn	0	2	0.028	0.081	0.366	0.992
Zn	0	3	0.027	0.083	0.384	0.992
Zn	1	1	0.025	0.087	0.312	0.995
Zn	1	2	0.024	0.101	0.301	0.996
Zn	1	3	0.034	0.061	0.430	0.972
Zn	2	1	0.031	0.064	0.382	0.992
Zn	2	2	0.031	0.058	0.436	0.987
Zn	2	3	0.029	0.078	0.389	0.998
Zn	3	1	0.023	0.097	0.313	0.997
Zn	3	2	0.024	0.093	0.328	0.996
Zn	3	3	0.028	0.087	0.341	0.998
Zn	4	1	0.028	0.072	0.367	0.994
Zn	4	2	0.025	0.077	0.385	0.991
Zn	4	3	0.029	0.086	0.345	0.992
Zn	5	1	0.031	0.066	0.308	0.992
Zn	5	2	0.033	0.046	0.434	0.986
Zn	5	3	0.031	0.071	0.291	0.990

Table A.5. DR 5 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.016	0.071	0.416	0.993
As	0	2	0.016	0.078	0.368	0.996
As	0	3	0.020	0.072	0.415	0.992
As	1	1	0.018	0.069	0.379	0.990
As	1	2	0.017	0.073	0.355	0.993
As	1	3	0.020	0.056	0.456	0.992
As	2	1	0.016	0.082	0.358	0.989
As	2	2	0.018	0.068	0.379	0.992
As	2	3	0.017	0.074	0.353	0.992
As	3	1	0.016	0.071	0.370	0.994
As	3	2	0.018	0.065	0.380	0.991
As	3	3	0.018	0.068	0.377	0.993
As	4	1	0.018	0.069	0.411	0.992
As	4	2	0.017	0.068	0.375	0.992
As	4	3	0.019	0.068	0.373	0.992
As	5	1	0.020	0.079	0.421	0.989
As	5	2	0.022	0.065	0.419	0.990
As	5	3	0.020	0.069	0.456	0.992
Cd	0	1	0.019	0.069	0.448	0.991
Cd	0	2	0.012	0.092	0.341	0.978
Cd	0	3	0.016	0.079	0.360	0.994
Cd	1	1	0.018	0.069	0.472	0.992
Cd	1	2	0.017	0.070	0.366	0.994
Cd	1	3	0.014	0.090	0.309	0.991
Cd	2	1	0.016	0.074	0.402	0.994
Cd	2	2	0.016	0.075	0.354	0.993
Cd	2	3	0.016	0.077	0.340	0.992

Table A.5. DR 5 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.020	0.060	0.464	0.990
Cd	3	2	0.016	0.068	0.366	0.993
Cd	3	3	0.016	0.077	0.333	0.992
Cd	4	1	0.018	0.045	0.395	0.990
Cd	4	2	0.016	0.051	0.373	0.990
Cd	4	3	0.016	0.064	0.340	0.990
Cd	5	1	0.014	0.060	0.256	0.990
Cd	5	2	0.013	0.055	0.254	0.991
Cd	5	3	0.013	0.054	0.252	0.993
Cr	0	1	0.018	0.070	0.412	0.994
Cr	0	2	0.024	0.062	0.473	0.989
Cr	0	3	0.022	0.068	0.464	0.990
Cr	1	1	0.019	0.074	0.365	0.992
Cr	1	2	0.023	0.057	0.500	0.988
Cr	1	3	0.024	0.066	0.476	0.989
Cr	2	1	0.019	0.073	0.397	0.993
Cr	2	2	0.021	0.065	0.406	0.991
Cr	2	3	0.021	0.067	0.491	0.996
Cr	3	1	0.021	0.080	0.364	0.987
Cr	3	2	0.021	0.067	0.454	0.991
Cr	3	3	0.024	0.065	0.491	0.990
Cr	4	1	0.020	0.074	0.401	0.992
Cr	4	2	0.025	0.056	0.464	0.988
Cr	4	3	0.022	0.077	0.447	0.990
Cr	5	1	0.019	0.085	0.414	0.995
Cr	5	2	0.019	0.082	0.479	0.995
Cr	5	3	0.018	0.090	0.454	0.997

Table A.5. DR 5 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.013	0.085	0.424	0.997
Hg	0	2	0.011	0.085	0.416	0.997
Hg	0	3	0.016	0.070	0.408	0.994
Hg	1	1	0.017	0.075	0.417	0.994
Hg	1	2	0.016	0.069	0.409	0.992
Hg	1	3	0.017	0.068	0.359	0.992
Hg	2	1	0.014	0.089	0.427	0.997
Hg	2	2	0.018	0.064	0.422	0.991
Hg	2	3	0.017	0.063	0.434	0.991
Hg	3	1	0.015	0.090	0.434	0.997
Hg	3	2	0.018	0.064	0.429	0.993
Hg	3	3	0.019	0.061	0.405	0.989
Hg	4	1	0.014	0.093	0.437	0.997
Hg	4	2	0.014	0.089	0.383	0.990
Hg	4	3	0.018	0.066	0.381	0.993
Hg	5	1	0.013	0.096	0.429	0.997
Hg	5	2	0.018	0.073	0.411	0.992
Hg	5	3	0.018	0.065	0.416	0.994
Pb	0	1	0.018	0.068	0.408	0.993
Pb	0	2	0.019	0.063	0.434	0.992
Pb	0	3	0.018	0.077	0.438	0.992
Pb	1	1	0.014	0.079	0.345	0.993
Pb	1	2	0.016	0.072	0.359	0.992
Pb	1	3	0.018	0.071	0.388	0.994
Pb	2	1	0.015	0.081	0.341	0.993
Pb	2	2	0.015	0.074	0.349	0.994
Pb	2	3	0.018	0.070	0.384	0.992

Table A.5. DR 5 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.016	0.077	0.364	0.993
Pb	3	2	0.017	0.074	0.379	0.992
Pb	3	3	0.017	0.076	0.358	0.991
Pb	4	1	0.015	0.078	0.347	0.994
Pb	4	2	0.017	0.074	0.363	0.993
Pb	4	3	0.017	0.074	0.359	0.994
Pb	5	1	0.021	0.062	0.447	0.989
Pb	5	2	0.017	0.074	0.377	0.993
Pb	5	3	0.018	0.073	0.458	0.995
Se	0	1	0.017	0.070	0.500	0.994
Se	0	2	0.017	0.071	0.494	0.998
Se	0	3	0.014	0.075	0.467	0.987
Se	1	1	0.015	0.074	0.456	0.992
Se	1	2	0.019	0.067	0.469	0.998
Se	1	3	0.015	0.070	0.500	0.997
Se	2	1	0.016	0.069	0.500	0.998
Se	2	2	0.018	0.064	0.500	0.998
Se	2	3	0.017	0.068	0.499	0.996
Se	3	1	0.015	0.076	0.450	0.995
Se	3	2	0.018	0.071	0.447	0.996
Se	3	3	0.015	0.075	0.442	0.997
Se	4	1	0.018	0.066	0.433	0.998
Se	4	2	0.022	0.057	0.467	0.994
Se	4	3	0.022	0.056	0.469	0.997
Se	5	1	0.020	0.037	0.489	0.998
Se	5	2	0.021	0.045	0.371	0.982
Se	5	3	0.021	0.038	0.475	0.993

Table A.5. DR 5 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.020	0.055	0.500	0.989
Zn	0	2	0.021	0.067	0.467	0.986
Zn	0	3	0.012	0.112	0.376	0.981
Zn	1	1	0.010	0.079	0.459	0.978
Zn	1	2	0.016	0.082	0.366	0.996
Zn	1	3	0.013	0.097	0.425	0.998
Zn	2	1	0.018	0.069	0.395	0.994
Zn	2	2	0.018	0.067	0.391	0.994
Zn	2	3	0.015	0.091	0.425	0.997
Zn	3	1	0.018	0.074	0.395	0.994
Zn	3	2	0.018	0.066	0.392	0.993
Zn	3	3	0.017	0.084	0.458	0.996
Zn	4	1	0.020	0.073	0.385	0.991
Zn	4	2	0.021	0.056	0.464	0.992
Zn	4	3	0.019	0.078	0.390	0.991
Zn	5	1	0.012	0.084	0.275	0.995
Zn	5	2	0.014	0.077	0.261	0.994
Zn	5	3	0.015	0.087	0.261	0.993

Table A.6. DR 52 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.050	0.063	0.384	0.993
As	0	2	0.050	0.055	0.393	0.997
As	0	3	0.050	0.058	0.387	0.998
As	1	1	0.050	0.049	0.340	0.996
As	1	2	0.048	0.050	0.364	0.993
As	1	3	0.050	0.043	0.410	0.994
As	2	1	0.050	0.045	0.407	0.994
As	2	2	0.046	0.045	0.408	0.994
As	2	3	0.046	0.050	0.344	0.990
As	3	1	0.050	0.048	0.360	0.995
As	3	2	0.050	0.046	0.400	0.996
As	3	3	0.050	0.045	0.408	0.993
As	4	1	0.050	0.047	0.396	0.995
As	4	2	0.050	0.051	0.362	0.996
As	4	3	0.050	0.051	0.363	0.993
As	5	1	0.050	0.051	0.362	0.995
As	5	2	0.046	0.054	0.329	0.992
As	5	3	0.050	0.068	0.292	0.996
Cd	0	1	0.050	0.049	0.391	0.993
Cd	0	2	0.050	0.052	0.379	0.992
Cd	0	3	0.050	0.052	0.397	0.995
Cd	1	1	0.050	0.047	0.401	0.989
Cd	1	2	0.050	0.051	0.412	0.992
Cd	1	3	0.050	0.049	0.396	0.993
Cd	2	1	0.050	0.056	0.359	0.994
Cd	2	2	0.050	0.052	0.378	0.994
Cd	2	3	0.050	0.053	0.390	0.993

Table A.6. DR 52 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.050	0.065	0.349	0.988
Cd	3	2	0.050	0.060	0.348	0.991
Cd	3	3	0.050	0.056	0.380	0.988
Cd	4	1	0.050	0.045	0.466	0.997
Cd	4	2	0.050	0.043	0.497	0.998
Cd	4	3	0.050	0.042	0.496	0.996
Cd	5	1	0.042	0.097	0.237	0.983
Cd	5	2	0.042	0.090	0.237	0.986
Cd	5	3	0.043	0.096	0.223	0.989
Cr	0	1	0.050	0.051	0.346	0.997
Cr	0	2	0.050	0.069	0.331	0.960
Cr	0	3	0.049	0.056	0.341	0.996
Cr	1	1	0.050	0.073	0.299	0.938
Cr	1	2	0.050	0.053	0.363	0.997
Cr	1	3	0.050	0.054	0.369	0.995
Cr	2	1	0.050	0.071	0.337	0.952
Cr	2	2	0.050	0.053	0.368	0.994
Cr	2	3	0.050	0.049	0.379	0.995
Cr	3	1	0.050	0.069	0.335	0.962
Cr	3	2	0.050	0.049	0.367	0.994
Cr	3	3	0.050	0.054	0.351	0.994
Cr	4	1	0.050	0.071	0.311	0.947
Cr	4	2	0.050	0.051	0.362	0.995
Cr	4	3	0.050	0.063	0.345	0.998
Cr	5	1	0.050	0.080	0.311	0.929
Cr	5	2	0.050	0.054	0.358	0.996
Cr	5	3	0.050	0.057	0.337	0.996

Table A.6. DR 52 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.050	0.057	0.372	0.997
Hg	0	2	0.050	0.056	0.390	0.997
Hg	0	3	0.050	0.057	0.371	0.997
Hg	1	1	0.050	0.056	0.372	0.996
Hg	1	2	0.050	0.054	0.384	0.996
Hg	1	3	0.050	0.056	0.382	0.996
Hg	2	1	0.050	0.056	0.384	0.995
Hg	2	2	0.050	0.056	0.386	0.996
Hg	2	3	0.050	0.054	0.377	0.996
Hg	3	1	0.050	0.056	0.367	0.996
Hg	3	2	0.050	0.051	0.432	0.996
Hg	3	3	0.050	0.054	0.401	0.996
Hg	4	1	0.050	0.059	0.353	0.994
Hg	4	2	0.050	0.057	0.368	0.995
Hg	4	3	0.050	0.056	0.386	0.996
Hg	5	1	0.050	0.063	0.354	0.992
Hg	5	2	0.050	0.062	0.369	0.992
Hg	5	3	0.050	0.058	0.375	0.996
Pb	0	1	0.050	0.050	0.396	0.994
Pb	0	2	0.050	0.054	0.372	0.992
Pb	0	3	0.050	0.055	0.378	0.995
Pb	1	1	0.050	0.051	0.410	0.992
Pb	1	2	0.050	0.046	0.443	0.993
Pb	1	3	0.050	0.050	0.405	0.992
Pb	2	1	0.050	0.047	0.445	0.996
Pb	2	2	0.050	0.050	0.414	0.993
Pb	2	3	0.050	0.052	0.394	0.994

Table A.6. DR 52 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.050	0.045	0.451	0.990
Pb	3	2	0.050	0.050	0.380	0.995
Pb	3	3	0.050	0.053	0.385	0.997
Pb	4	1	0.050	0.057	0.372	0.995
Pb	4	2	0.050	0.051	0.398	0.995
Pb	4	3	0.050	0.052	0.395	0.994
Pb	5	1	0.050	0.057	0.366	0.993
Pb	5	2	0.050	0.054	0.374	0.988
Pb	5	3	0.050	0.058	0.367	0.995
Se	0	1	0.001	0.425	0.002	0.295
Se	0	2	0.000	0.531	0.002	0.403
Se	0	3	0.002	0.177	0.002	0.019
Se	1	1	0.000	0.600	0.002	0.363
Se	1	2	0.000	0.600	0.003	0.704
Se	1	3	0.000	0.047	0.500	0.922
Se	2	1	0.000	0.600	0.003	0.796
Se	2	2	0.001	0.600	0.002	0.379
Se	2	3	0.000	0.600	0.002	0.742
Se	3	1	0.000	0.600	0.002	0.417
Se	3	2	0.000	0.049	0.500	0.962
Se	3	3	0.000	0.043	0.500	0.899
Se	4	1	0.001	0.600	0.003	0.575
Se	4	2	0.000	0.595	0.002	0.801
Se	4	3	0.000	0.600	0.002	0.493
Se	5	1	0.000	0.600	0.002	0.503
Se	5	2	0.001	0.117	0.002	0.261
Se	5	3	0.000	0.030	0.500	0.902

Table A.6. DR 52 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.050	0.052	0.392	0.992
Zn	0	2	0.050	0.054	0.365	0.990
Zn	0	3	0.050	0.057	0.371	0.996
Zn	1	1	0.050	0.052	0.389	0.993
Zn	1	2	0.050	0.053	0.390	0.992
Zn	1	3	0.050	0.055	0.384	0.994
Zn	2	1	0.050	0.050	0.417	0.989
Zn	2	2	0.050	0.050	0.396	0.994
Zn	2	3	0.050	0.053	0.388	0.993
Zn	3	1	0.050	0.052	0.373	0.994
Zn	3	2	0.050	0.052	0.396	0.995
Zn	3	3	0.050	0.054	0.392	0.993
Zn	4	1	0.050	0.055	0.372	0.994
Zn	4	2	0.050	0.050	0.401	0.994
Zn	4	3	0.050	0.055	0.376	0.994
Zn	5	1	0.050	0.049	0.312	0.990
Zn	5	2	0.050	0.050	0.307	0.984
Zn	5	3	0.050	0.048	0.326	0.988

Table A.7. DR 60 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.036	0.049	0.558	0.999
As	0	2	0.037	0.047	0.565	0.999
As	0	3	0.037	0.048	0.557	0.999
As	1	1	0.036	0.049	0.558	0.999
As	1	2	0.037	0.049	0.543	1.000
As	1	3	0.033	0.050	0.557	0.999
As	2	1	0.036	0.044	0.600	0.998
As	2	2	0.036	0.048	0.567	0.998
As	2	3	0.036	0.048	0.548	0.999
As	3	1	0.036	0.044	0.586	0.999
As	3	2	0.037	0.042	0.595	0.999
As	3	3	0.035	0.042	0.621	0.999
As	4	1	0.036	0.046	0.579	0.999
As	4	2	0.035	0.046	0.573	0.999
As	4	3	0.037	0.039	0.673	0.998
As	5	1	0.037	0.044	0.608	0.999
As	5	2	0.038	0.043	0.602	0.999
As	5	3	0.039	0.043	0.620	0.999
Cd	0	1	0.033	0.057	0.536	1.000
Cd	0	2	0.033	0.062	0.493	0.999
Cd	0	3	0.033	0.059	0.501	0.999
Cd	1	1	0.035	0.057	0.511	1.000
Cd	1	2	0.036	0.055	0.507	0.999
Cd	1	3	0.036	0.057	0.506	0.999
Cd	2	1	0.035	0.059	0.498	0.999
Cd	2	2	0.035	0.058	0.498	1.000
Cd	2	3	0.035	0.057	0.518	0.999

Table A.7. DR 60 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.036	0.059	0.502	0.999
Cd	3	2	0.036	0.056	0.502	0.999
Cd	3	3	0.034	0.060	0.484	1.000
Cd	4	1	0.035	0.053	0.539	0.999
Cd	4	2	0.043	0.043	0.568	0.941
Cd	4	3	0.036	0.057	0.513	0.999
Cd	5	1	0.032	0.046	0.576	0.999
Cd	5	2	0.037	0.044	0.591	0.999
Cd	5	3	0.036	0.047	0.561	1.000
Cr	0	1	0.037	0.046	0.573	0.999
Cr	0	2	0.035	0.049	0.556	0.999
Cr	0	3	0.036	0.048	0.555	0.999
Cr	1	1	0.035	0.049	0.541	0.999
Cr	1	2	0.038	0.046	0.584	0.998
Cr	1	3	0.038	0.046	0.584	0.999
Cr	2	1	0.033	0.048	0.554	0.999
Cr	2	2	0.036	0.047	0.567	0.999
Cr	2	3	0.038	0.045	0.576	0.999
Cr	3	1	0.037	0.044	0.574	0.999
Cr	3	2	0.036	0.044	0.580	0.999
Cr	3	3	0.035	0.043	0.611	0.999
Cr	4	1	0.035	0.042	0.589	0.999
Cr	4	2	0.035	0.044	0.581	0.998
Cr	4	3	0.036	0.043	0.600	0.999
Cr	5	1	0.031	0.047	0.576	0.996
Cr	5	2	0.038	0.042	0.631	0.999
Cr	5	3	0.037	0.045	0.594	0.998

Table A.7. DR 60 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.040	0.040	0.649	0.994
Hg	0	2	0.040	0.045	0.601	0.999
Hg	0	3	0.037	0.044	0.610	0.997
Hg	1	1	0.037	0.046	0.598	0.999
Hg	1	2	0.037	0.046	0.587	0.999
Hg	1	3	0.035	0.048	0.558	0.999
Hg	2	1	0.038	0.045	0.605	0.999
Hg	2	2	0.034	0.049	0.557	0.999
Hg	2	3	0.034	0.050	0.546	0.999
Hg	3	1	0.035	0.049	0.566	0.999
Hg	3	2	0.036	0.049	0.554	0.999
Hg	3	3	0.036	0.050	0.554	1.000
Hg	4	1	0.036	0.047	0.592	0.999
Hg	4	2	0.035	0.051	0.553	0.999
Hg	4	3	0.034	0.049	0.550	1.000
Hg	5	1	0.030	0.050	0.570	0.999
Hg	5	2	0.032	0.049	0.561	0.999
Hg	5	3	0.035	0.048	0.556	0.999
Pb	0	1	0.034	0.053	0.532	0.999
Pb	0	2	0.033	0.055	0.515	1.000
Pb	0	3	0.036	0.052	0.546	0.999
Pb	1	1	0.034	0.052	0.540	0.999
Pb	1	2	0.035	0.053	0.533	1.000
Pb	1	3	0.034	0.056	0.515	1.000
Pb	2	1	0.035	0.055	0.540	0.999
Pb	2	2	0.036	0.056	0.519	0.999
Pb	2	3	0.034	0.056	0.519	0.999

Table A.7. DR 60 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.036	0.055	0.523	0.999
Pb	3	2	0.034	0.058	0.511	0.999
Pb	3	3	0.035	0.056	0.514	1.000
Pb	4	1	0.034	0.055	0.515	1.000
Pb	4	2	0.033	0.064	0.440	0.999
Pb	4	3	0.035	0.053	0.529	0.999
Pb	5	1	0.037	0.051	0.569	0.999
Pb	5	2	0.035	0.055	0.547	0.999
Pb	5	3	0.037	0.050	0.574	0.999
Se	0	1	0.001	0.008	1.000	0.525
Se	0	2	0.000	0.300	0.003	0.651
Se	0	3	0.002	0.300	0.002	0.003
Se	1	1	0.001	0.002	1.000	0.106
Se	1	2	0.000	0.300	0.002	0.558
Se	1	3	0.000	0.300	0.002	0.436
Se	2	1	0.001	0.010	0.037	0.577
Se	2	2	0.001	0.300	0.002	0.177
Se	2	3	0.002	0.300	0.003	0.043
Se	3	1	0.001	0.300	0.003	0.726
Se	3	2	0.000	0.300	0.002	0.490
Se	3	3	0.000	0.041	1.000	0.977
Se	4	1	0.001	0.300	0.003	0.558
Se	4	2	0.001	0.300	0.003	0.514
Se	4	3	0.001	0.300	0.003	0.532
Se	5	1	0.001	0.300	0.003	0.452
Se	5	2	0.001	0.300	0.003	0.306
Se	5	3	0.001	0.300	0.003	0.421

Table A.7. DR 60 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.036	0.048	0.558	1.000
Zn	0	2	0.035	0.048	0.572	0.994
Zn	0	3	0.036	0.052	0.525	1.000
Zn	1	1	0.035	0.053	0.514	0.999
Zn	1	2	0.036	0.051	0.552	1.000
Zn	1	3	0.036	0.050	0.552	0.999
Zn	2	1	0.036	0.050	0.541	1.000
Zn	2	2	0.036	0.048	0.579	0.999
Zn	2	3	0.036	0.049	0.555	1.000
Zn	3	1	0.037	0.051	0.537	0.999
Zn	3	2	0.035	0.052	0.521	1.000
Zn	3	3	0.035	0.053	0.528	0.999
Zn	4	1	0.037	0.049	0.556	0.999
Zn	4	2	0.035	0.050	0.556	0.998
Zn	4	3	0.038	0.049	0.567	1.000
Zn	5	1	0.033	0.040	0.574	0.999
Zn	5	2	0.033	0.042	0.550	0.998
Zn	5	3	0.036	0.033	0.744	0.998

Table A.8. DR 76 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.002	0.060	0.059	0.963
As	0	2	0.003	0.059	0.059	0.996
As	0	3	0.004	0.025	0.356	0.961
As	1	1	0.007	0.034	1.000	0.984
As	1	2	0.006	0.043	0.699	0.990
As	1	3	0.001	0.081	0.354	0.997
As	2	1	0.000	0.046	0.683	0.999
As	2	2	0.000	0.086	0.282	0.998
As	2	3	0.000	0.052	0.374	0.996
As	3	1	0.001	0.080	0.333	0.997
As	3	2	0.001	0.073	0.415	0.996
As	3	3	0.002	0.088	0.259	0.997
As	4	1	0.001	0.037	1.000	0.999
As	4	2	0.002	0.029	0.827	0.997
As	4	3	0.001	0.031	1.000	0.995
As	5	1	0.000	0.101	0.217	0.992
As	5	2	0.002	0.057	0.235	0.997
As	5	3	0.002	0.032	1.000	0.997
Cd	0	1	0.002	0.021	1.000	0.958
Cd	0	2	0.000	0.082	0.280	0.997
Cd	0	3	0.000	0.038	1.000	0.836
Cd	1	1	0.003	0.044	0.157	0.997
Cd	1	2	0.003	0.029	0.248	0.968
Cd	1	3	0.003	0.044	0.532	0.997
Cd	2	1	0.003	0.028	0.364	0.982
Cd	2	2	0.002	0.060	0.365	1.000
Cd	2	3	0.000	0.133	0.064	0.923

Table A.8. DR 76 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.002	0.038	0.698	0.996
Cd	3	2	0.000	0.094	0.368	0.994
Cd	3	3	0.006	0.045	0.361	0.955
Cd	4	1	0.000	0.090	0.256	0.986
Cd	4	2	0.000	0.136	0.063	0.924
Cd	4	3	0.003	0.028	0.476	0.994
Cd	5	1	0.000	0.086	0.283	0.995
Cd	5	2	0.000	0.090	0.255	0.988
Cd	5	3	0.000	0.086	0.270	0.995
Cr	0	1	0.000	0.046	1.000	0.999
Cr	0	2	0.004	0.019	0.580	0.970
Cr	0	3	0.002	0.039	0.286	0.997
Cr	1	1	0.004	0.033	0.167	0.990
Cr	1	2	0.001	0.081	0.053	0.961
Cr	1	3	0.004	0.022	0.444	0.985
Cr	2	1	0.007	0.043	0.801	0.992
Cr	2	2	0.001	0.086	0.345	0.997
Cr	2	3	0.001	0.080	0.405	0.995
Cr	3	1	0.000	0.094	0.254	0.995
Cr	3	2	0.000	0.084	0.291	0.986
Cr	3	3	0.002	0.033	1.000	1.000
Cr	4	1	0.004	0.052	0.072	0.936
Cr	4	2	0.002	0.056	0.064	0.981
Cr	4	3	0.005	0.031	1.000	0.990
Cr	5	1	0.000	0.090	0.234	0.993
Cr	5	2	0.003	0.038	0.111	0.977
Cr	5	3	0.000	0.029	1.000	0.922

Table A.8. DR 76 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.004	0.035	0.123	0.973
Hg	0	2	0.003	0.021	1.000	0.961
Hg	0	3	0.003	0.027	0.709	0.999
Hg	1	1	0.000	0.083	0.285	0.998
Hg	1	2	0.002	0.072	0.082	0.990
Hg	1	3	0.004	0.031	0.178	0.957
Hg	2	1	0.004	0.035	0.146	0.973
Hg	2	2	0.003	0.027	0.307	0.990
Hg	2	3	0.002	0.075	0.062	0.959
Hg	3	1	0.002	0.077	0.053	0.985
Hg	3	2	0.001	0.112	0.052	0.988
Hg	3	3	0.001	0.098	0.059	0.907
Hg	4	1	0.001	0.092	0.050	0.991
Hg	4	2	0.003	0.074	0.055	0.991
Hg	4	3	0.002	0.076	0.053	0.975
Hg	5	1	0.001	0.024	1.000	0.986
Hg	5	2	0.002	0.061	0.045	0.958
Hg	5	3	0.004	0.023	0.266	0.988
Pb	0	1	0.000	0.079	0.311	0.986
Pb	0	2	0.000	0.053	0.625	0.999
Pb	0	3	0.004	0.018	1.000	0.957
Pb	1	1	0.000	0.090	0.260	0.991
Pb	1	2	0.000	0.058	0.493	0.999
Pb	1	3	0.000	0.075	0.379	0.999
Pb	2	1	0.000	0.070	0.411	0.999
Pb	2	2	0.003	0.072	0.245	0.984
Pb	2	3	0.003	0.075	0.321	0.995

Table A.8. DR 76 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.000	0.057	0.407	0.995
Pb	3	2	0.000	0.091	0.246	0.985
Pb	3	3	0.000	0.087	0.223	0.981
Pb	4	1	0.000	0.095	0.251	0.985
Pb	4	2	0.000	0.094	0.216	0.988
Pb	4	3	0.000	0.077	0.214	0.987
Pb	5	1	0.000	0.047	0.702	0.998
Pb	5	2	0.000	0.073	0.378	0.999
Pb	5	3	0.000	0.097	0.240	0.990
Se	0	1	0.009	0.105	0.107	0.943
Se	0	2	0.016	0.055	0.122	0.973
Se	0	3	0.010	0.062	0.118	0.965
Se	1	1	0.011	0.087	0.107	0.993
Se	1	2	0.012	0.088	0.114	0.978
Se	1	3	0.013	0.079	0.112	0.985
Se	2	1	0.012	0.067	0.148	0.985
Se	2	2	0.012	0.067	0.122	0.992
Se	2	3	0.013	0.060	0.134	0.983
Se	3	1	0.014	0.062	0.123	0.986
Se	3	2	0.012	0.072	0.107	0.990
Se	3	3	0.014	0.070	0.107	0.985
Se	4	1	0.013	0.070	0.090	0.973
Se	4	2	0.013	0.073	0.085	0.975
Se	4	3	0.012	0.081	0.076	0.979
Se	5	1	0.010	0.196	0.042	0.973
Se	5	2	0.009	0.201	0.039	0.970
Se	5	3	0.011	0.161	0.047	0.983

Table A.8. DR 76 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.000	0.094	0.303	0.994
Zn	0	2	0.000	0.065	0.409	0.998
Zn	0	3	0.000	0.063	0.429	0.999
Zn	1	1	0.000	0.092	0.262	0.985
Zn	1	2	0.000	0.100	0.234	0.984
Zn	1	3	0.000	0.103	0.270	0.992
Zn	2	1	0.000	0.096	0.247	0.990
Zn	2	2	0.001	0.077	0.280	0.988
Zn	2	3	0.000	0.090	0.331	0.995
Zn	3	1	0.000	0.059	0.573	0.999
Zn	3	2	0.000	0.085	0.332	0.999
Zn	3	3	0.001	0.056	0.438	0.999
Zn	4	1	0.003	0.036	1.000	1.000
Zn	4	2	0.002	0.038	1.000	0.998
Zn	4	3	0.000	0.061	0.483	1.000
Zn	5	1	0.000	0.071	0.399	0.997
Zn	5	2	0.000	0.062	0.486	0.998
Zn	5	3	0.001	0.042	1.000	0.998

Table A.9. DR p51 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.049	0.033	0.479	0.993
As	0	2	0.041	0.045	0.396	0.996
As	0	3	0.040	0.051	0.370	0.997
As	1	1	0.039	0.056	0.329	0.998
As	1	2	0.042	0.042	0.403	0.993
As	1	3	0.050	0.020	1.000	0.959
As	2	1	0.038	0.061	0.316	0.995
As	2	2	0.040	0.054	0.335	0.994
As	2	3	0.043	0.050	0.358	0.994
As	3	1	0.042	0.060	0.331	0.989
As	3	2	0.041	0.061	0.309	0.992
As	3	3	0.040	0.055	0.344	0.995
As	4	1	0.048	0.055	0.351	0.982
As	4	2	0.045	0.055	0.351	0.986
As	4	3	0.045	0.053	0.367	0.990
As	5	1	0.044	0.065	0.342	0.993
As	5	2	0.045	0.055	0.362	0.987
As	5	3	0.046	0.052	0.382	0.987
Cd	0	1	0.042	0.066	0.355	0.988
Cd	0	2	0.039	0.072	0.325	0.987
Cd	0	3	0.035	0.072	0.319	0.991
Cd	1	1	0.040	0.065	0.356	0.992
Cd	1	2	0.040	0.058	0.345	0.972
Cd	1	3	0.044	0.065	0.313	0.986
Cd	2	1	0.031	0.077	0.321	0.989
Cd	2	2	0.044	0.061	0.328	0.990
Cd	2	3	0.042	0.064	0.330	0.989

Table A.9. DR p51 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.035	0.071	0.318	0.990
Cd	3	2	0.029	0.079	0.294	0.987
Cd	3	3	0.037	0.063	0.310	0.995
Cd	4	1	0.032	0.076	0.288	0.993
Cd	4	2	0.038	0.071	0.291	0.995
Cd	4	3	0.035	0.076	0.282	0.992
Cd	5	1	0.028	0.099	0.191	0.997
Cd	5	2	0.032	0.074	0.180	0.990
Cd	5	3	0.024	0.081	0.179	0.994
Cr	0	1	0.040	0.052	0.361	0.994
Cr	0	2	0.041	0.050	0.362	0.993
Cr	0	3	0.043	0.050	0.416	0.995
Cr	1	1	0.039	0.055	0.349	0.996
Cr	1	2	0.040	0.051	0.332	0.989
Cr	1	3	0.039	0.053	0.386	0.987
Cr	2	1	0.038	0.053	0.358	0.991
Cr	2	2	0.042	0.055	0.363	0.994
Cr	2	3	0.041	0.057	0.382	0.995
Cr	3	1	0.038	0.066	0.292	0.997
Cr	3	2	0.040	0.058	0.365	0.993
Cr	3	3	0.039	0.063	0.356	0.992
Cr	4	1	0.039	0.059	0.361	0.983
Cr	4	2	0.043	0.057	0.368	0.990
Cr	4	3	0.047	0.055	0.382	0.986
Cr	5	1	0.040	0.060	0.363	0.984
Cr	5	2	0.042	0.054	0.405	0.988
Cr	5	3	0.042	0.065	0.366	0.986

Table A.9. DR p51 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.043	0.052	0.350	0.992
Hg	0	2	0.044	0.040	0.422	0.993
Hg	0	3	0.040	0.056	0.336	0.990
Hg	1	1	0.041	0.055	0.376	0.993
Hg	1	2	0.041	0.050	0.392	0.996
Hg	1	3	0.039	0.053	0.358	0.996
Hg	2	1	0.043	0.059	0.364	0.992
Hg	2	2	0.041	0.058	0.347	0.994
Hg	2	3	0.040	0.051	0.361	0.996
Hg	3	1	0.043	0.064	0.344	0.991
Hg	3	2	0.039	0.068	0.327	0.993
Hg	3	3	0.044	0.049	0.383	0.994
Hg	4	1	0.044	0.066	0.358	0.991
Hg	4	2	0.040	0.063	0.348	0.995
Hg	4	3	0.042	0.050	0.362	0.993
Hg	5	1	0.044	0.056	0.378	0.988
Hg	5	2	0.042	0.057	0.364	0.993
Hg	5	3	0.042	0.056	0.349	0.995
Pb	0	1	0.038	0.059	0.333	0.984
Pb	0	2	0.035	0.068	0.302	0.985
Pb	0	3	0.032	0.076	0.310	0.987
Pb	1	1	0.028	0.068	0.311	0.988
Pb	1	2	0.046	0.061	0.326	0.994
Pb	1	3	0.042	0.082	0.276	0.996
Pb	2	1	0.040	0.053	0.331	0.984
Pb	2	2	0.030	0.071	0.292	0.988
Pb	2	3	0.039	0.076	0.285	0.995

Table A.9. DR p51 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.028	0.058	0.311	0.980
Pb	3	2	0.027	0.070	0.297	0.983
Pb	3	3	0.029	0.092	0.273	0.993
Pb	4	1	0.035	0.064	0.303	0.982
Pb	4	2	0.020	0.081	0.245	0.989
Pb	4	3	0.034	0.071	0.302	0.985
Pb	5	1	0.040	0.060	0.327	0.982
Pb	5	2	0.029	0.066	0.308	0.980
Pb	5	3	0.025	0.078	0.276	0.976
Se	0	1	0.020	0.058	0.446	0.998
Se	0	2	0.020	0.066	0.430	0.997
Se	0	3	0.021	0.064	0.465	0.997
Se	1	1	0.022	0.058	0.429	0.997
Se	1	2	0.021	0.065	0.406	0.998
Se	1	3	0.023	0.068	0.425	0.995
Se	2	1	0.023	0.059	0.406	0.997
Se	2	2	0.017	0.076	0.384	0.997
Se	2	3	0.019	0.071	0.421	0.997
Se	3	1	0.024	0.052	0.403	0.996
Se	3	2	0.021	0.067	0.383	0.996
Se	3	3	0.023	0.071	0.408	0.998
Se	4	1	0.022	0.059	0.341	0.997
Se	4	2	0.023	0.060	0.348	0.998
Se	4	3	0.027	0.059	0.356	0.998
Se	5	1	0.021	0.078	0.215	0.996
Se	5	2	0.022	0.073	0.226	0.996
Se	5	3	0.023	0.078	0.254	0.995

Table A.9. DR p51 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.034	0.067	0.318	0.986
Zn	0	2	0.050	0.066	0.348	0.982
Zn	0	3	0.038	0.072	0.325	0.988
Zn	1	1	0.042	0.061	0.301	0.985
Zn	1	2	0.050	0.048	0.324	0.978
Zn	1	3	0.039	0.072	0.331	0.987
Zn	2	1	0.045	0.049	0.331	0.986
Zn	2	2	0.042	0.052	0.354	0.985
Zn	2	3	0.042	0.069	0.334	0.993
Zn	3	1	0.048	0.047	0.351	0.981
Zn	3	2	0.040	0.053	0.351	0.986
Zn	3	3	0.043	0.064	0.345	0.988
Zn	4	1	0.043	0.055	0.340	0.989
Zn	4	2	0.045	0.045	0.398	0.992
Zn	4	3	0.043	0.061	0.345	0.990
Zn	5	1	0.046	0.053	0.325	0.974
Zn	5	2	0.048	0.044	0.381	0.984
Zn	5	3	0.048	0.059	0.356	0.985

Table A.10. DR p72 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.033	0.057	0.395	0.989
As	0	2	0.038	0.057	0.382	0.987
As	0	3	0.032	0.057	0.375	0.985
As	1	1	0.034	0.061	0.376	0.987
As	1	2	0.034	0.059	0.364	0.987
As	1	3	0.040	0.046	0.416	0.976
As	2	1	0.037	0.051	0.403	0.962
As	2	2	0.032	0.062	0.357	0.986
As	2	3	0.031	0.064	0.359	0.984
As	3	1	0.029	0.066	0.369	0.993
As	3	2	0.032	0.061	0.370	0.986
As	3	3	0.029	0.069	0.355	0.991
As	4	1	0.034	0.058	0.391	0.983
As	4	2	0.037	0.042	0.478	0.983
As	4	3	0.034	0.060	0.385	0.991
As	5	1	0.035	0.056	0.402	0.984
As	5	2	0.036	0.054	0.422	0.989
As	5	3	0.035	0.064	0.391	0.985
Cd	0	1	0.031	0.069	0.354	0.987
Cd	0	2	0.035	0.069	0.349	0.981
Cd	0	3	0.031	0.064	0.361	0.989
Cd	1	1	0.031	0.065	0.396	0.992
Cd	1	2	0.037	0.046	0.445	0.978
Cd	1	3	0.025	0.086	0.309	0.995
Cd	2	1	0.030	0.064	0.371	0.989
Cd	2	2	0.030	0.065	0.359	0.990
Cd	2	3	0.030	0.068	0.350	0.987

Table A.10. DR p72 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.031	0.070	0.340	0.990
Cd	3	2	0.032	0.068	0.333	0.990
Cd	3	3	0.034	0.064	0.347	0.986
Cd	4	1	0.024	0.096	0.253	0.990
Cd	4	2	0.028	0.087	0.253	0.984
Cd	4	3	0.026	0.092	0.248	0.988
Cd	5	1	0.017	0.146	0.173	0.980
Cd	5	2	0.019	0.147	0.173	0.980
Cd	5	3	0.025	0.139	0.187	0.979
Cr	0	1	0.028	0.071	0.352	0.991
Cr	0	2	0.029	0.067	0.365	0.989
Cr	0	3	0.029	0.067	0.371	0.989
Cr	1	1	0.033	0.057	0.360	0.968
Cr	1	2	0.029	0.070	0.340	0.989
Cr	1	3	0.034	0.063	0.386	0.989
Cr	2	1	0.030	0.073	0.343	0.987
Cr	2	2	0.032	0.060	0.398	0.986
Cr	2	3	0.033	0.059	0.394	0.988
Cr	3	1	0.030	0.070	0.358	0.990
Cr	3	2	0.031	0.064	0.364	0.992
Cr	3	3	0.033	0.059	0.397	0.987
Cr	4	1	0.031	0.064	0.377	0.988
Cr	4	2	0.036	0.056	0.406	0.989
Cr	4	3	0.033	0.060	0.400	0.982
Cr	5	1	0.037	0.057	0.398	0.982
Cr	5	2	0.034	0.055	0.400	0.986
Cr	5	3	0.034	0.061	0.406	0.986

Table A.10. DR p72 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.034	0.061	0.368	0.975
Hg	0	2	0.035	0.051	0.424	0.985
Hg	0	3	0.032	0.063	0.348	0.984
Hg	1	1	0.035	0.055	0.416	0.985
Hg	1	2	0.036	0.059	0.384	0.991
Hg	1	3	0.036	0.052	0.405	0.985
Hg	2	1	0.034	0.061	0.402	0.988
Hg	2	2	0.031	0.066	0.348	0.988
Hg	2	3	0.033	0.053	0.405	0.980
Hg	3	1	0.032	0.059	0.397	0.990
Hg	3	2	0.033	0.055	0.409	0.990
Hg	3	3	0.033	0.056	0.409	0.990
Hg	4	1	0.050	0.043	0.456	0.984
Hg	4	2	0.037	0.051	0.417	0.985
Hg	4	3	0.036	0.048	0.462	0.986
Hg	5	1	0.033	0.058	0.418	0.988
Hg	5	2	0.034	0.055	0.427	0.985
Hg	5	3	0.036	0.051	0.426	0.986
Pb	0	1	0.031	0.067	0.368	0.990
Pb	0	2	0.034	0.048	0.431	0.982
Pb	0	3	0.033	0.059	0.365	0.984
Pb	1	1	0.032	0.061	0.377	0.989
Pb	1	2	0.031	0.060	0.367	0.987
Pb	1	3	0.026	0.081	0.315	0.993
Pb	2	1	0.033	0.056	0.383	0.983
Pb	2	2	0.029	0.063	0.368	0.988
Pb	2	3	0.032	0.061	0.368	0.991

Table A.10. DR p72 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.038	0.054	0.396	0.981
Pb	3	2	0.030	0.067	0.359	0.989
Pb	3	3	0.033	0.058	0.346	0.983
Pb	4	1	0.030	0.068	0.372	0.988
Pb	4	2	0.030	0.067	0.368	0.991
Pb	4	3	0.033	0.059	0.371	0.984
Pb	5	1	0.032	0.065	0.367	0.989
Pb	5	2	0.033	0.067	0.350	0.988
Pb	5	3	0.031	0.065	0.362	0.986
Se	0	1	0.015	0.051	0.139	0.969
Se	0	2	0.016	0.047	0.140	0.980
Se	0	3	0.015	0.063	0.107	0.989
Se	1	1	0.018	0.025	0.246	0.969
Se	1	2	0.017	0.052	0.109	0.978
Se	1	3	0.014	0.063	0.098	0.990
Se	2	1	0.014	0.061	0.105	0.989
Se	2	2	0.014	0.058	0.113	0.994
Se	2	3	0.011	0.068	0.128	0.988
Se	3	1	0.016	0.061	0.089	0.980
Se	3	2	0.013	0.065	0.085	0.963
Se	3	3	0.013	0.067	0.086	0.995
Se	4	1	0.014	0.075	0.064	0.975
Se	4	2	0.015	0.049	0.073	0.977
Se	4	3	0.013	0.101	0.058	0.982
Se	5	1	0.012	0.222	0.040	0.939
Se	5	2	0.014	0.106	0.043	0.920
Se	5	3	0.015	0.139	0.040	0.923

Table A.10. DR p72 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.033	0.055	0.399	0.986
Zn	0	2	0.026	0.078	0.350	0.992
Zn	0	3	0.033	0.060	0.377	0.985
Zn	1	1	0.034	0.061	0.351	0.987
Zn	1	2	0.033	0.057	0.379	0.984
Zn	1	3	0.033	0.064	0.353	0.984
Zn	2	1	0.031	0.069	0.360	0.988
Zn	2	2	0.031	0.067	0.359	0.990
Zn	2	3	0.032	0.060	0.399	0.990
Zn	3	1	0.034	0.057	0.376	0.989
Zn	3	2	0.034	0.057	0.392	0.988
Zn	3	3	0.033	0.059	0.391	0.990
Zn	4	1	0.033	0.061	0.372	0.988
Zn	4	2	0.035	0.058	0.386	0.985
Zn	4	3	0.033	0.063	0.389	0.991
Zn	5	1	0.030	0.075	0.306	0.989
Zn	5	2	0.031	0.070	0.316	0.987
Zn	5	3	0.034	0.068	0.330	0.985

Table A.11. DR p74 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.015	0.133	0.086	0.988
As	0	2	0.018	0.072	0.116	0.969
As	0	3	0.019	0.148	0.117	0.953
As	1	1	0.019	0.156	0.082	0.980
As	1	2	0.017	0.123	0.085	0.978
As	1	3	0.018	0.058	0.140	0.979
As	2	1	0.018	0.117	0.097	0.966
As	2	2	0.017	0.124	0.086	0.972
As	2	3	0.022	0.101	0.132	0.982
As	3	1	0.016	0.112	0.088	0.969
As	3	2	0.020	0.124	0.089	0.973
As	3	3	0.020	0.062	0.167	0.985
As	4	1	0.018	0.083	0.126	0.981
As	4	2	0.018	0.089	0.124	0.986
As	4	3	0.022	0.081	0.138	0.986
As	5	1	0.024	0.082	0.146	0.976
As	5	2	0.020	0.111	0.126	0.991
As	5	3	0.024	0.100	0.135	0.970
Cd	0	1	0.017	0.176	0.084	0.958
Cd	0	2	0.017	0.160	0.075	0.983
Cd	0	3	0.019	0.164	0.076	0.971
Cd	1	1	0.016	0.167	0.074	0.975
Cd	1	2	0.015	0.196	0.080	0.912
Cd	1	3	0.018	0.081	0.091	0.966
Cd	2	1	0.013	0.230	0.078	0.979
Cd	2	2	0.014	0.179	0.082	0.970
Cd	2	3	0.020	0.068	0.128	0.961

Table A.11. DR p74 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.015	0.146	0.081	0.977
Cd	3	2	0.015	0.143	0.082	0.986
Cd	3	3	0.021	0.055	0.173	0.991
Cd	4	1	0.016	0.099	0.107	0.971
Cd	4	2	0.021	0.059	0.132	0.965
Cd	4	3	0.018	0.068	0.165	0.995
Cd	5	1	0.019	0.114	0.121	0.992
Cd	5	2	0.020	0.096	0.134	0.981
Cd	5	3	0.021	0.086	0.138	0.990
Cr	0	1	0.016	0.118	0.133	0.989
Cr	0	2	0.020	0.081	0.149	0.985
Cr	0	3	0.022	0.086	0.146	0.989
Cr	1	1	0.021	0.107	0.140	0.988
Cr	1	2	0.018	0.142	0.102	0.950
Cr	1	3	0.020	0.095	0.122	0.980
Cr	2	1	0.016	0.108	0.131	0.990
Cr	2	2	0.013	0.102	0.119	0.993
Cr	2	3	0.025	0.077	0.147	0.984
Cr	3	1	0.021	0.115	0.110	0.979
Cr	3	2	0.022	0.090	0.120	0.969
Cr	3	3	0.020	0.100	0.120	0.980
Cr	4	1	0.024	0.078	0.157	0.978
Cr	4	2	0.021	0.082	0.159	0.989
Cr	4	3	0.019	0.074	0.166	0.990
Cr	5	1	0.030	0.061	0.224	0.987
Cr	5	2	0.028	0.064	0.214	0.988
Cr	5	3	0.030	0.067	0.198	0.992

Table A.11. DR p74 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.019	0.115	0.137	0.984
Hg	0	2	0.018	0.132	0.131	0.976
Hg	0	3	0.020	0.107	0.144	0.979
Hg	1	1	0.015	0.098	0.143	0.991
Hg	1	2	0.017	0.090	0.139	0.986
Hg	1	3	0.024	0.058	0.182	0.977
Hg	2	1	0.023	0.092	0.147	0.980
Hg	2	2	0.023	0.074	0.161	0.992
Hg	2	3	0.024	0.080	0.154	0.973
Hg	3	1	0.022	0.097	0.143	0.984
Hg	3	2	0.021	0.074	0.156	0.986
Hg	3	3	0.021	0.073	0.159	0.979
Hg	4	1	0.011	0.445	0.028	0.768
Hg	4	2	0.014	0.435	0.039	0.758
Hg	4	3	0.019	0.315	0.034	0.830
Hg	5	1	0.016	0.000	0.000	0.183
Hg	5	2	0.017	0.078	0.014	0.157
Hg	5	3	0.018	0.001	0.000	0.234
Pb	0	1	0.021	0.115	0.139	0.991
Pb	0	2	0.025	0.083	0.147	0.984
Pb	0	3	0.021	0.115	0.130	0.974
Pb	1	1	0.027	0.069	0.161	0.965
Pb	1	2	0.026	0.080	0.142	0.985
Pb	1	3	0.023	0.096	0.148	0.986
Pb	2	1	0.022	0.081	0.141	0.977
Pb	2	2	0.025	0.068	0.169	0.990
Pb	2	3	0.024	0.085	0.156	0.981

Table A.11. DR p74 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.020	0.079	0.142	0.983
Pb	3	2	0.021	0.068	0.153	0.986
Pb	3	3	0.024	0.078	0.158	0.982
Pb	4	1	0.018	0.091	0.140	0.991
Pb	4	2	0.017	0.096	0.135	0.977
Pb	4	3	0.019	0.062	0.194	0.987
Pb	5	1	0.018	0.141	0.138	0.981
Pb	5	2	0.009	0.151	0.130	0.988
Pb	5	3	0.016	0.124	0.135	0.984
Se	0	1	0.018	0.114	0.142	0.980
Se	0	2	0.017	0.119	0.134	0.974
Se	0	3	0.026	0.097	0.150	0.959
Se	1	1	0.023	0.109	0.135	0.986
Se	1	2	0.024	0.090	0.146	0.968
Se	1	3	0.021	0.096	0.146	0.971
Se	2	1	0.023	0.069	0.180	0.986
Se	2	2	0.023	0.072	0.168	0.988
Se	2	3	0.021	0.063	0.192	0.986
Se	3	1	0.014	0.114	0.136	0.986
Se	3	2	0.022	0.064	0.204	0.993
Se	3	3	0.022	0.064	0.181	0.985
Se	4	1	0.017	0.133	0.112	0.992
Se	4	2	0.018	0.115	0.116	0.995
Se	4	3	0.023	0.087	0.141	0.984
Se	5	1	0.025	0.100	0.128	0.992
Se	5	2	0.025	0.100	0.123	0.995
Se	5	3	0.027	0.092	0.112	0.990

Table A.11. DR p74 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.017	0.137	0.109	0.965
Zn	0	2	0.013	0.141	0.111	0.897
Zn	0	3	0.018	0.087	0.141	0.973
Zn	1	1	0.018	0.141	0.097	0.989
Zn	1	2	0.022	0.094	0.103	0.953
Zn	1	3	0.019	0.064	0.169	0.994
Zn	2	1	0.016	0.147	0.088	0.966
Zn	2	2	0.022	0.088	0.105	0.948
Zn	2	3	0.024	0.066	0.141	0.958
Zn	3	1	0.021	0.092	0.126	0.971
Zn	3	2	0.024	0.059	0.156	0.984
Zn	3	3	0.021	0.083	0.131	0.983
Zn	4	1	0.023	0.078	0.128	0.975
Zn	4	2	0.019	0.109	0.121	0.988
Zn	4	3	0.023	0.095	0.131	0.983
Zn	5	1	0.020	0.118	0.112	0.968
Zn	5	2	0.023	0.112	0.121	0.975
Zn	5	3	0.025	0.101	0.130	0.975

Table A.12. NO 14 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.007	0.063	0.308	0.994
As	0	2	0.006	0.069	0.369	0.998
As	0	3	0.006	0.067	0.335	0.996
As	1	1	0.010	0.034	0.950	0.985
As	1	2	0.009	0.035	1.000	0.990
As	1	3	0.011	0.034	1.000	0.968
As	2	1	0.008	0.050	0.524	0.996
As	2	2	0.006	0.070	0.301	0.995
As	2	3	0.007	0.061	0.358	0.992
As	3	1	0.004	0.082	0.270	0.999
As	3	2	0.010	0.030	1.000	0.977
As	3	3	0.006	0.049	0.496	0.998
As	4	1	0.010	0.033	1.000	0.958
As	4	2	0.006	0.066	0.347	0.998
As	4	3	0.006	0.071	0.283	0.995
As	5	1	0.010	0.039	0.885	0.985
As	5	2	0.003	0.093	0.261	0.994
As	5	3	0.001	0.136	0.205	0.995
Cd	0	1	0.007	0.054	0.308	0.982
Cd	0	2	0.000	0.264	0.082	0.941
Cd	0	3	0.008	0.058	0.288	0.984
Cd	1	1	0.011	0.086	0.191	0.998
Cd	1	2	0.006	0.096	0.162	0.986
Cd	1	3	0.011	0.046	0.333	0.986
Cd	2	1	0.007	0.066	0.271	0.990
Cd	2	2	0.003	0.095	0.216	0.909
Cd	2	3	0.007	0.060	0.304	0.990

Table A.12. NO 14 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.009	0.046	0.418	0.984
Cd	3	2	0.007	0.048	0.477	0.981
Cd	3	3	0.005	0.045	0.484	0.992
Cd	4	1	0.009	0.054	0.401	0.983
Cd	4	2	0.010	0.047	0.423	0.982
Cd	4	3	0.009	0.038	0.513	0.985
Cd	5	1	0.006	0.071	0.173	0.979
Cd	5	2	0.006	0.056	0.206	0.954
Cd	5	3	0.007	0.053	0.187	0.974
Cr	0	1	0.004	0.085	0.248	0.966
Cr	0	2	0.010	0.041	0.575	0.985
Cr	0	3	0.006	0.078	0.293	0.991
Cr	1	1	0.012	0.048	0.439	0.996
Cr	1	2	0.004	0.086	0.275	0.994
Cr	1	3	0.005	0.091	0.283	0.998
Cr	2	1	0.007	0.079	0.266	0.994
Cr	2	2	0.004	0.087	0.238	0.999
Cr	2	3	0.004	0.081	0.291	1.000
Cr	3	1	0.007	0.072	0.300	0.993
Cr	3	2	0.006	0.070	0.317	0.999
Cr	3	3	0.003	0.090	0.270	0.997
Cr	4	1	0.004	0.080	0.311	0.999
Cr	4	2	0.005	0.087	0.290	0.990
Cr	4	3	0.005	0.080	0.277	0.996
Cr	5	1	0.007	0.060	0.409	0.996
Cr	5	2	0.007	0.064	0.342	0.988
Cr	5	3	0.004	0.084	0.324	0.995

Table A.12. NO 14 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.008	0.053	0.510	0.996
Hg	0	2	0.010	0.036	1.000	0.997
Hg	0	3	0.008	0.070	0.327	0.989
Hg	1	1	0.009	0.048	0.502	0.990
Hg	1	2	0.007	0.053	0.478	0.994
Hg	1	3	0.005	0.066	0.354	0.998
Hg	2	1	0.009	0.038	0.790	0.991
Hg	2	2	0.008	0.051	0.472	0.999
Hg	2	3	0.003	0.081	0.285	0.989
Hg	3	1	0.005	0.063	0.397	0.993
Hg	3	2	0.004	0.065	0.425	0.996
Hg	3	3	0.006	0.067	0.360	0.996
Hg	4	1	0.005	0.082	0.278	0.990
Hg	4	2	0.007	0.036	1.000	0.978
Hg	4	3	0.004	0.073	0.344	1.000
Hg	5	1	0.006	0.064	0.367	0.995
Hg	5	2	0.007	0.038	0.897	0.996
Hg	5	3	0.004	0.072	0.326	0.999
Pb	0	1	0.003	0.124	0.145	0.993
Pb	0	2	0.007	0.057	0.341	0.992
Pb	0	3	0.007	0.027	1.000	0.950
Pb	1	1	0.009	0.039	0.521	0.985
Pb	1	2	0.007	0.027	1.000	0.967
Pb	1	3	0.005	0.095	0.171	0.981
Pb	2	1	0.009	0.058	0.399	0.985
Pb	2	2	0.011	0.038	0.558	0.983
Pb	2	3	0.007	0.068	0.257	0.992

Table A.12. NO 14 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.009	0.032	1.000	0.980
Pb	3	2	0.009	0.040	0.481	0.966
Pb	3	3	0.009	0.033	1.000	0.988
Pb	4	1	0.011	0.029	1.000	0.977
Pb	4	2	0.007	0.065	0.263	0.990
Pb	4	3	0.011	0.029	0.999	0.963
Pb	5	1	0.005	0.062	0.465	0.965
Pb	5	2	0.009	0.034	1.000	0.965
Pb	5	3	0.008	0.058	0.373	0.990
Se	0	1	0.001	0.074	0.114	0.986
Se	0	2	0.001	0.067	0.148	0.993
Se	0	3	0.006	0.068	0.347	0.995
Se	1	1	0.001	0.065	0.146	0.988
Se	1	2	0.001	0.054	0.183	0.991
Se	1	3	0.000	0.085	0.200	0.998
Se	2	1	0.000	0.078	0.146	0.989
Se	2	2	0.000	0.086	0.141	0.992
Se	2	3	0.000	0.096	0.186	0.998
Se	3	1	0.000	0.112	0.124	0.990
Se	3	2	0.001	0.085	0.145	0.994
Se	3	3	0.001	0.076	0.191	0.994
Se	4	1	0.000	0.165	0.079	0.981
Se	4	2	0.000	0.180	0.079	0.989
Se	4	3	0.000	0.173	0.090	0.989
Se	5	1	0.000	0.158	0.047	0.996
Se	5	2	0.000	0.158	0.052	0.995
Se	5	3	0.000	0.170	0.053	0.994

Table A.12. NO 14 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.010	0.031	1.000	0.966
Zn	0	2	0.010	0.053	0.358	0.983
Zn	0	3	0.010	0.033	1.000	0.977
Zn	1	1	0.009	0.035	1.000	0.971
Zn	1	2	0.009	0.049	0.440	0.977
Zn	1	3	0.010	0.034	1.000	0.972
Zn	2	1	0.009	0.048	0.540	0.983
Zn	2	2	0.011	0.038	0.829	0.984
Zn	2	3	0.010	0.034	1.000	0.975
Zn	3	1	0.009	0.051	0.480	0.989
Zn	3	2	0.009	0.044	0.680	0.992
Zn	3	3	0.010	0.047	0.556	0.986
Zn	4	1	0.010	0.043	0.604	0.983
Zn	4	2	0.011	0.031	1.000	0.974
Zn	4	3	0.010	0.036	1.000	0.980
Zn	5	1	0.009	0.032	1.000	0.966
Zn	5	2	0.006	0.045	0.685	0.997
Zn	5	3	0.007	0.046	0.592	0.995

Table A.13. NO 17 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.017	0.078	0.389	0.999
As	0	2	0.017	0.078	0.379	0.998
As	0	3	0.015	0.085	0.353	0.999
As	1	1	0.019	0.068	0.423	0.996
As	1	2	0.022	0.054	0.548	0.992
As	1	3	0.016	0.077	0.393	0.998
As	2	1	0.015	0.078	0.387	0.998
As	2	2	0.016	0.080	0.381	0.998
As	2	3	0.015	0.081	0.393	0.998
As	3	1	0.016	0.077	0.395	0.998
As	3	2	0.016	0.077	0.398	0.998
As	3	3	0.015	0.082	0.388	0.998
As	4	1	0.016	0.077	0.415	0.997
As	4	2	0.015	0.081	0.384	0.998
As	4	3	0.016	0.075	0.424	0.996
As	5	1	0.014	0.080	0.399	0.996
As	5	2	0.016	0.068	0.446	0.996
As	5	3	0.015	0.073	0.434	0.998
Cd	0	1	0.017	0.058	0.482	0.999
Cd	0	2	0.016	0.069	0.421	0.997
Cd	0	3	0.019	0.060	0.468	0.999
Cd	1	1	0.023	0.044	0.677	0.994
Cd	1	2	0.033	0.056	0.473	0.996
Cd	1	3	0.019	0.063	0.443	0.999
Cd	2	1	0.020	0.058	0.449	0.994
Cd	2	2	0.018	0.068	0.418	0.999
Cd	2	3	0.017	0.070	0.411	0.999

Table A.13. NO 17 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.019	0.062	0.450	0.994
Cd	3	2	0.020	0.068	0.410	0.999
Cd	3	3	0.019	0.068	0.415	0.999
Cd	4	1	0.020	0.066	0.425	0.999
Cd	4	2	0.018	0.068	0.409	0.999
Cd	4	3	0.019	0.053	0.492	0.998
Cd	5	1	0.020	0.049	0.383	0.998
Cd	5	2	0.019	0.052	0.345	0.999
Cd	5	3	0.021	0.045	0.379	0.997
Cr	0	1	0.019	0.058	0.492	0.996
Cr	0	2	0.017	0.079	0.405	0.998
Cr	0	3	0.016	0.081	0.418	0.998
Cr	1	1	0.012	0.099	0.336	0.997
Cr	1	2	0.016	0.080	0.402	0.998
Cr	1	3	0.015	0.084	0.398	0.999
Cr	2	1	0.012	0.075	0.407	0.997
Cr	2	2	0.013	0.083	0.389	0.996
Cr	2	3	0.015	0.087	0.391	0.999
Cr	3	1	0.015	0.082	0.409	0.998
Cr	3	2	0.013	0.087	0.387	0.998
Cr	3	3	0.016	0.078	0.447	NA
Cr	4	1	0.016	0.077	0.402	0.997
Cr	4	2	0.014	0.085	0.394	0.998
Cr	4	3	0.013	0.087	0.431	0.998
Cr	5	1	0.015	0.084	0.411	0.997
Cr	5	2	0.016	0.075	0.440	0.995
Cr	5	3	0.013	0.089	0.404	0.997

Table A.13. NO 17 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.017	0.075	0.404	0.998
Hg	0	2	0.019	0.073	0.406	0.998
Hg	0	3	0.017	0.075	0.386	0.999
Hg	1	1	0.018	0.075	0.405	0.999
Hg	1	2	0.018	0.069	0.431	0.997
Hg	1	3	0.018	0.075	0.396	0.996
Hg	2	1	0.021	0.065	0.399	0.946
Hg	2	2	0.018	0.075	0.367	0.974
Hg	2	3	0.016	0.077	0.390	0.999
Hg	3	1	0.019	0.060	0.464	0.994
Hg	3	2	0.017	0.073	0.404	0.998
Hg	3	3	0.015	0.076	0.392	0.998
Hg	4	1	0.013	0.088	0.389	0.998
Hg	4	2	0.017	0.070	0.462	0.998
Hg	4	3	0.016	0.073	0.442	0.998
Hg	5	1	0.016	0.078	0.415	0.998
Hg	5	2	0.017	0.072	0.417	0.998
Hg	5	3	0.014	0.080	0.410	0.999
Pb	0	1	0.020	0.054	0.467	0.999
Pb	0	2	0.020	0.056	0.471	0.998
Pb	0	3	0.019	0.060	0.462	0.998
Pb	1	1	0.019	0.055	0.475	0.996
Pb	1	2	0.020	0.058	0.466	0.998
Pb	1	3	0.018	0.059	0.463	0.999
Pb	2	1	0.019	0.064	0.436	0.999
Pb	2	2	0.019	0.063	0.439	0.998
Pb	2	3	0.018	0.064	0.446	0.999

Table A.13. NO 17 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.022	0.062	0.401	0.973
Pb	3	2	0.021	0.058	0.438	0.996
Pb	3	3	0.020	0.064	0.425	0.999
Pb	4	1	0.022	0.054	0.463	0.993
Pb	4	2	0.019	0.069	0.395	0.999
Pb	4	3	0.014	0.078	0.435	0.995
Pb	5	1	0.016	0.068	0.414	0.999
Pb	5	2	0.018	0.065	0.406	0.999
Pb	5	3	0.017	0.067	0.416	0.998
Se	0	1	0.006	0.110	0.231	0.993
Se	0	2	0.008	0.082	0.286	0.991
Se	0	3	0.006	0.077	0.285	0.994
Se	1	1	0.007	0.059	0.137	0.961
Se	1	2	0.008	0.034	0.212	0.943
Se	1	3	0.009	0.042	0.142	0.902
Se	2	1	0.009	0.038	0.184	0.938
Se	2	2	0.009	0.031	0.251	0.936
Se	2	3	0.006	0.063	0.102	0.943
Se	3	1	0.006	0.068	0.102	0.947
Se	3	2	0.006	0.070	0.103	0.955
Se	3	3	0.007	0.064	0.111	0.937
Se	4	1	0.002	0.172	0.067	0.940
Se	4	2	0.001	0.205	0.063	0.974
Se	4	3	0.004	0.107	0.070	0.942
Se	5	1	0.003	0.034	0.112	0.965
Se	5	2	0.003	0.059	0.050	0.936
Se	5	3	0.007	0.059	0.258	0.988

Table A.13. NO 17 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.018	0.065	0.422	0.999
Zn	0	2	0.017	0.065	0.435	0.997
Zn	0	3	0.021	0.050	0.565	0.994
Zn	1	1	0.017	0.066	0.408	0.999
Zn	1	2	0.021	0.057	0.451	0.988
Zn	1	3	0.019	0.071	0.413	0.999
Zn	2	1	0.021	0.056	0.547	0.998
Zn	2	2	0.019	0.061	0.504	0.998
Zn	2	3	0.019	0.066	0.485	0.999
Zn	3	1	0.020	0.059	0.456	0.998
Zn	3	2	0.018	0.077	0.382	0.999
Zn	3	3	0.018	0.074	0.404	0.998
Zn	4	1	0.018	0.067	0.411	0.997
Zn	4	2	0.017	0.070	0.414	0.998
Zn	4	3	0.015	0.076	0.392	0.997
Zn	5	1	0.020	0.057	0.415	0.990
Zn	5	2	0.016	0.068	0.379	0.999
Zn	5	3	0.016	0.076	0.376	0.998

Table A.14. NO 22 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity

Metal	factor	Replicate	y0	mumax	K	r2
As	0	1	0.000	0.063	0.026	0.983
As	0	2	0.001	0.028	0.142	0.994
As	0	3	0.000	0.066	0.043	0.992
As	1	1	0.001	0.016	1.000	0.994
As	1	2	0.001	0.026	0.137	0.988
As	1	3	0.000	0.039	0.141	0.999
As	2	1	0.001	0.015	1.000	0.987
As	2	2	0.000	0.105	0.028	0.974
As	2	3	0.000	0.118	0.027	0.999
As	3	1	0.001	0.018	0.820	0.975
As	3	2	0.001	0.017	1.000	0.980
As	3	3	0.000	0.088	0.037	0.983
As	4	1	0.001	0.018	1.000	0.974
As	4	2	0.001	0.018	1.000	0.988
As	4	3	0.001	0.016	1.000	0.977
As	5	1	0.000	0.082	0.051	0.999
As	5	2	0.001	0.019	1.000	0.998
As	5	3	0.000	0.038	1.000	0.998
Cd	0	1	0.000	0.044	0.215	0.991
Cd	0	2	0.000	0.023	1.000	0.995
Cd	0	3	0.000	0.025	1.000	0.996
Cd	1	1	0.001	0.022	0.743	0.998
Cd	1	2	0.000	0.086	0.081	0.990
Cd	1	3	0.001	0.020	1.000	0.992
Cd	2	1	0.001	0.020	1.000	0.996
Cd	2	2	0.000	0.076	0.041	0.998
Cd	2	3	0.000	0.080	0.052	0.998

Table A.14. NO 22 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Cd	3	1	0.002	0.053	0.005	0.513
Cd	3	2	0.000	0.083	0.086	0.986
Cd	3	3	0.002	0.009	0.556	0.694
Cd	4	1	0.001	0.252	0.001	0.037
Cd	4	2	0.002	0.300	0.001	0.010
Cd	4	3	0.001	0.029	0.114	0.968
Cd	5	1	0.002	0.300	0.001	0.266
Cd	5	2	0.002	0.300	0.001	0.125
Cd	5	3	0.002	0.002	0.000	0.253
Cr	0	1	0.000	0.089	0.051	0.965
Cr	0	2	0.001	0.021	1.000	0.997
Cr	0	3	0.000	0.084	0.084	0.979
Cr	1	1	0.000	0.059	0.081	0.986
Cr	1	2	0.000	0.020	1.000	0.994
Cr	1	3	0.000	0.023	1.000	0.994
Cr	2	1	0.002	0.014	1.000	0.954
Cr	2	2	0.000	0.033	0.142	0.988
Cr	2	3	0.000	0.062	0.106	0.993
Cr	3	1	0.000	0.059	0.063	0.999
Cr	3	2	0.000	0.026	1.000	0.995
Cr	3	3	0.000	0.021	1.000	0.992
Cr	4	1	0.000	0.070	0.052	0.996
Cr	4	2	0.002	0.015	1.000	0.566
Cr	4	3	0.000	0.148	0.052	0.854
Cr	5	1	0.001	0.018	1.000	0.997
Cr	5	2	0.000	0.078	0.097	0.991
Cr	5	3	0.000	0.088	0.076	0.987

Table A.14. NO 22 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Hg	0	1	0.000	0.144	0.018	0.816
Hg	0	2	0.000	0.138	0.026	0.963
Hg	0	3	0.000	0.189	0.021	0.991
Hg	1	1	0.001	0.037	0.111	0.992
Hg	1	2	0.001	0.014	1.000	0.976
Hg	1	3	0.000	0.028	1.000	0.992
Hg	2	1	0.000	0.020	1.000	0.984
Hg	2	2	0.001	0.017	1.000	0.984
Hg	2	3	0.000	0.043	0.110	0.989
Hg	3	1	0.000	0.117	0.042	0.926
Hg	3	2	0.000	0.058	0.071	0.988
Hg	3	3	0.000	0.115	0.044	0.969
Hg	4	1	0.000	0.037	0.121	0.993
Hg	4	2	0.000	0.040	1.000	0.998
Hg	4	3	0.000	0.020	1.000	0.994
Hg	5	1	0.000	0.070	0.058	0.991
Hg	5	2	0.001	0.015	1.000	0.940
Hg	5	3	0.000	0.104	0.032	0.987
Pb	0	1	0.001	0.018	1.000	0.995
Pb	0	2	0.000	0.020	1.000	0.995
Pb	0	3	0.000	0.048	0.111	0.993
Pb	1	1	0.000	0.032	1.000	0.997
Pb	1	2	0.000	0.036	1.000	0.997
Pb	1	3	0.000	0.037	1.000	0.997
Pb	2	1	0.000	0.061	0.068	0.998
Pb	2	2	0.000	0.022	1.000	0.996
Pb	2	3	0.000	0.089	0.193	0.999

Table A.14. NO 22 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Pb	3	1	0.000	0.021	1.000	0.996
Pb	3	2	0.001	0.019	1.000	0.983
Pb	3	3	0.000	0.058	0.520	0.999
Pb	4	1	0.000	0.060	0.069	0.959
Pb	4	2	0.000	0.146	0.051	0.939
Pb	4	3	0.000	0.044	1.000	0.999
Pb	5	1	0.000	0.019	1.000	0.991
Pb	5	2	0.000	0.029	1.000	0.998
Pb	5	3	0.001	0.038	0.115	0.991
Se	0	1	0.003	0.087	0.306	0.998
Se	0	2	0.005	0.068	0.368	0.992
Se	0	3	0.011	0.077	0.355	0.993
Se	1	1	0.000	0.018	1.000	0.990
Se	1	2	0.001	0.030	0.219	0.995
Se	1	3	0.000	0.080	0.148	0.997
Se	2	1	0.001	0.067	0.160	0.990
Se	2	2	0.000	0.097	0.158	0.956
Se	2	3	0.001	0.060	0.196	0.991
Se	3	1	0.000	0.097	0.147	0.991
Se	3	2	0.001	0.081	0.161	0.994
Se	3	3	0.001	0.053	0.207	0.990
Se	4	1	0.000	0.091	0.096	0.993
Se	4	2	0.001	0.085	0.094	0.988
Se	4	3	0.000	0.099	0.099	0.989
Se	5	1	0.000	0.105	0.037	0.978
Se	5	2	0.000	0.102	0.043	0.996
Se	5	3	0.000	0.103	0.050	0.997

Table A.14. NO 22 Gompertz Model Parameters. Estimations of maximal growth rates and carrying capacities. factor = Metal Concentration Factor (Table 3.1), mumax = Maximal Growth Rate, K = Carrying Capacity (*continued*)

Metal	factor	Replicate	y0	mumax	K	r2
Zn	0	1	0.001	0.025	0.438	0.988
Zn	0	2	0.000	0.020	1.000	0.981
Zn	0	3	0.001	0.018	1.000	0.994
Zn	1	1	0.000	0.061	0.077	0.990
Zn	1	2	0.001	0.019	1.000	0.987
Zn	1	3	0.000	0.022	1.000	0.995
Zn	2	1	0.001	0.067	0.264	0.992
Zn	2	2	0.002	0.062	0.295	0.994
Zn	2	3	0.002	0.054	0.462	0.996
Zn	3	1	0.001	0.018	1.000	0.991
Zn	3	2	0.000	0.036	1.000	0.997
Zn	3	3	0.000	0.069	0.057	0.997
Zn	4	1	0.000	0.080	0.082	0.985
Zn	4	2	0.001	0.015	1.000	0.986
Zn	4	3	0.001	0.025	0.208	0.989
Zn	5	1	0.001	0.033	0.050	0.953
Zn	5	2	0.001	0.048	0.029	0.694
Zn	5	3	0.000	0.016	1.000	0.966