

The background image shows a forest with many thin, light-colored tree trunks. A stream flows through the center of the forest, with water that is a distinct reddish-brown color, likely due to mineral content or erosion. The ground is covered with fallen leaves and some green plants.

The SURRI project: Developing new biorecovery approaches at legacy uranium and polymetallic mining sites

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Bioremediation/biorecovery challenges

1. Extreme pH and Redox Conditions
2. High Toxicity of Heavy Metals and Metalloids (As, Hg, Cd, Pb, Zn, Cu)
3. Low Nutrient Availability – C, N, P
4. Poor Oxygen Penetration and Mass Transfer
5. Maintaining Functional Microbial Communities
6. Scaling Up and Field Application

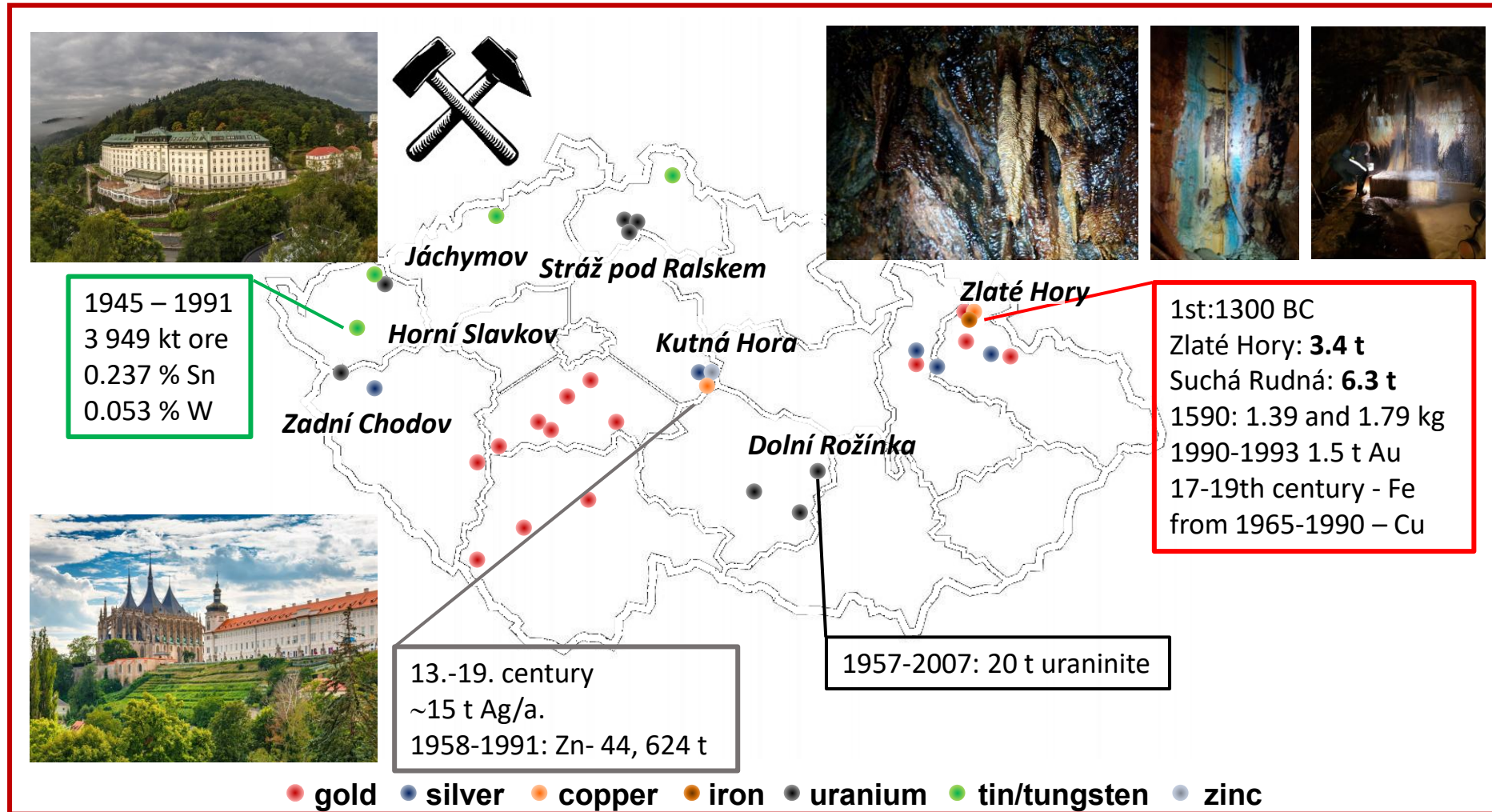
Element (limit)	Mine water (mg/L)	Tailings (mg/kg)	Tailings (mg/kg)	Sediment (mg/kg)
Al	26.5	9,658	5,024	1,252
As	18.7	63.1	147	<0.01
Ca	80.1	8,688	11,388	637
Cd	NA	8.8	33.9	<0.01
Co	0.26	15.7	32	<0.01
Cr	NA	21.6	9.9	<0.01
Cu	18.7	662	2,669	<0.01
Fe	34.4	39,646	66,703	454,464
K	0.94	683	507	<0.01
Mg	33.8	8,771	5,694	<0.01
Mn	7.56	844	765	446
Na	3.74	NA	NA	<0.01
Ni	NA	25.2	NA	<0.01
Pb	NA	549	2,584	806
V	<0.01	14.4	NA	<0.01
Zn	2.9	1,689	7,273	880

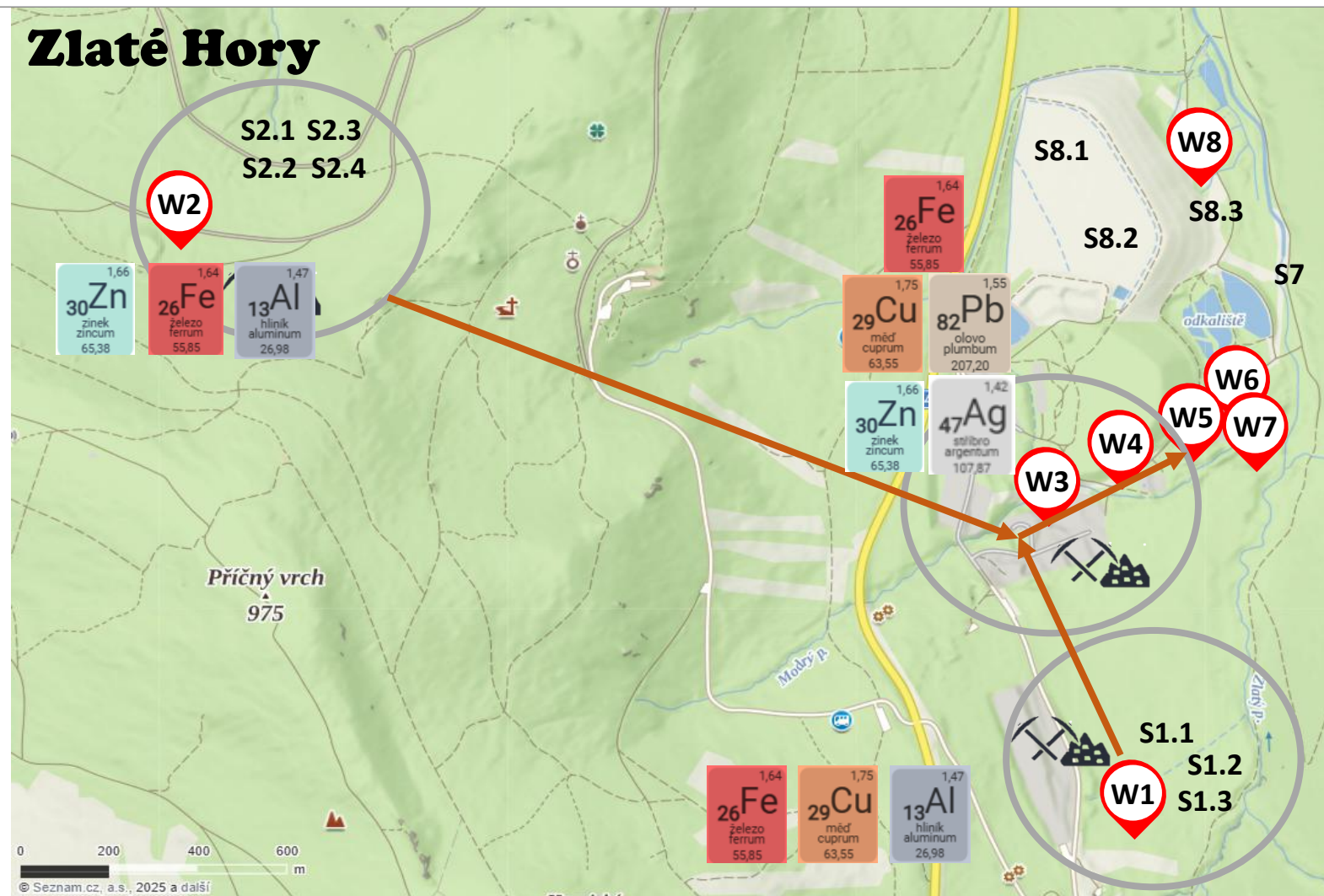
AIMs

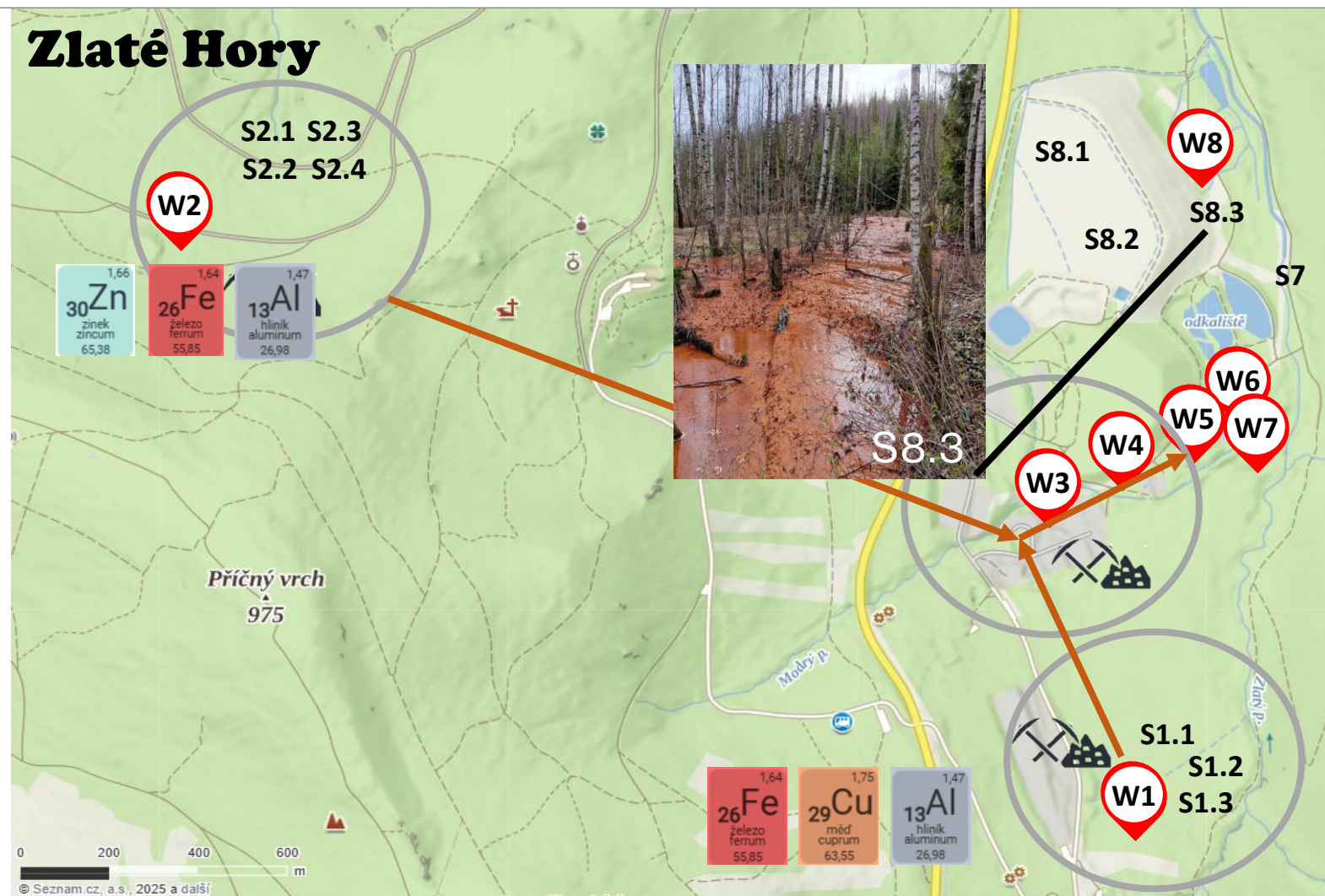
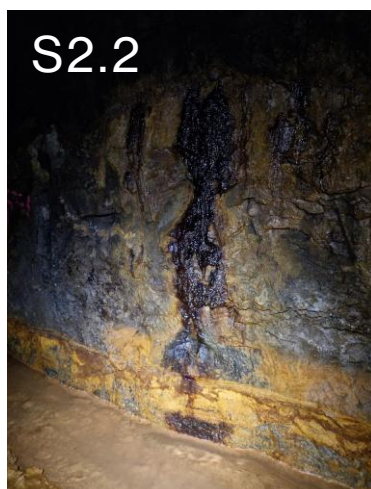
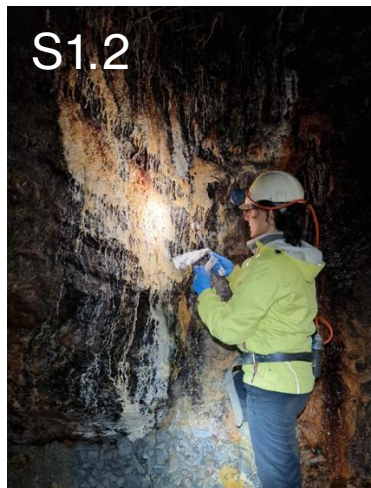
- Describe the effect of mine water treatment on microbial diversity
- Characterize microbial isolates in terms of applicability for bioremediation and biorecovery
- Develop novel strategy combining chemical and bio-based approach and determine its efficiency

Tools

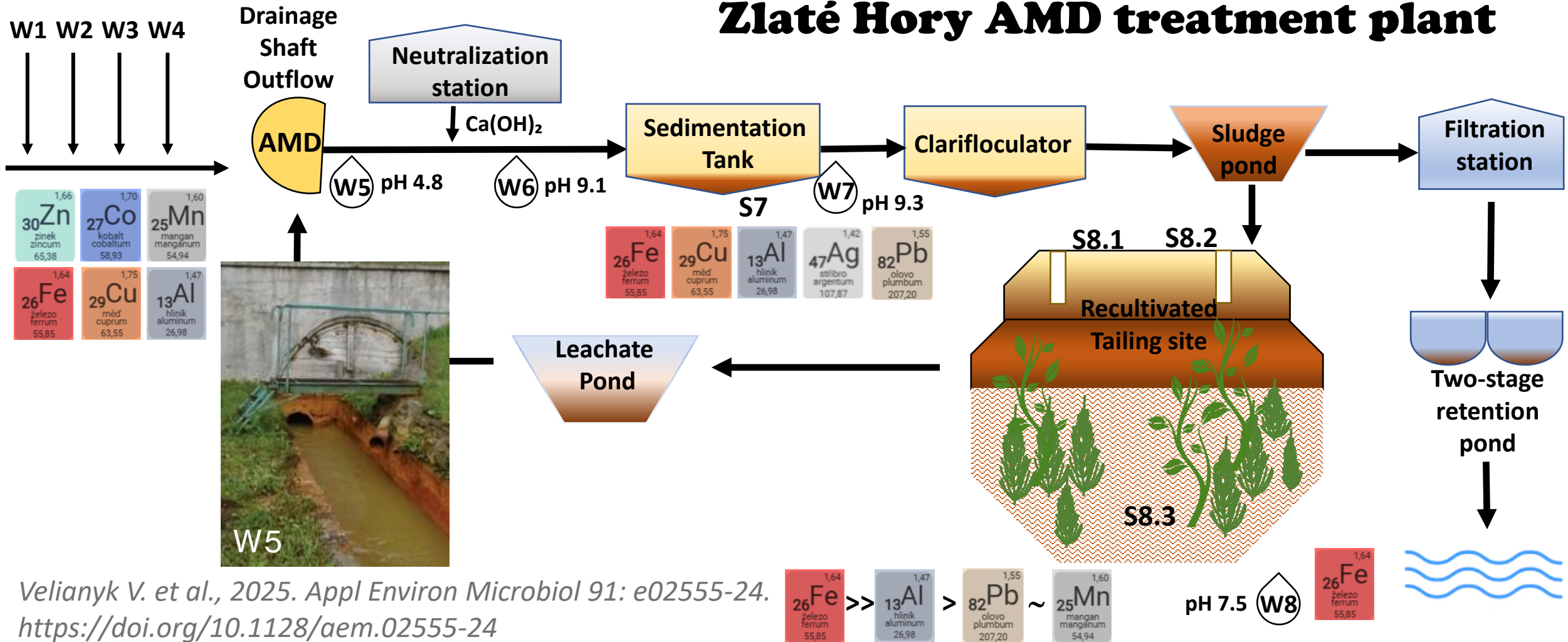
- Alpha/beta diversity, relative abundance changes, correlation of taxa with physico-chemical parameters
- Isolate, identify (sequencing), growth kinetics, MIC/tolerance tests, biosorption and bioaccumulation tests
- Recovery yield of target metals, microbial monitoring (amplicon sequencing and viability)

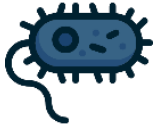




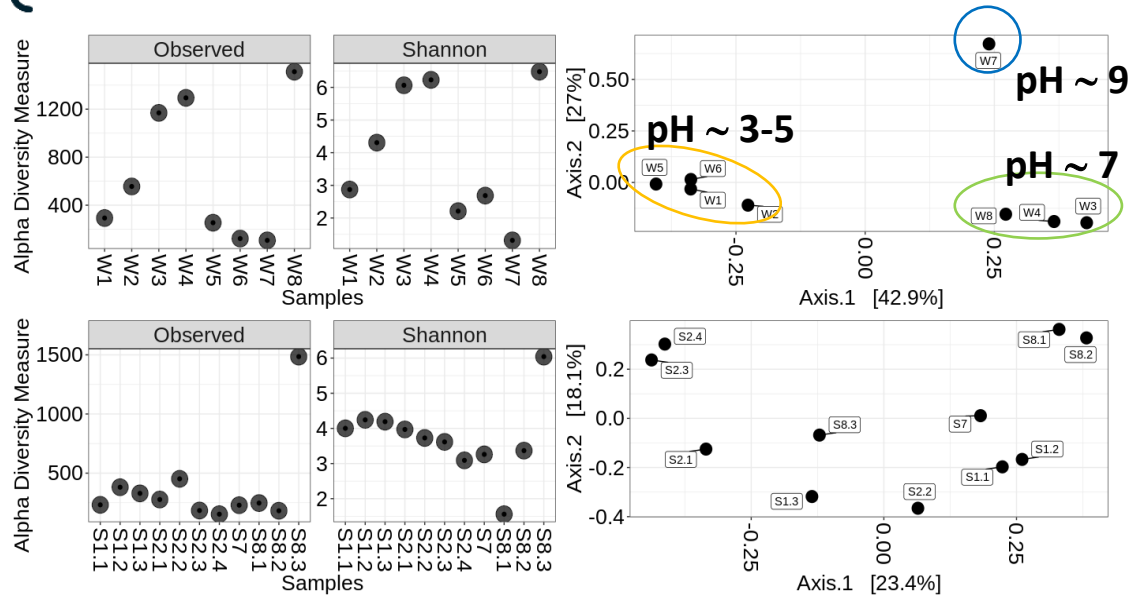


Zlaté Hory AMD treatment plant

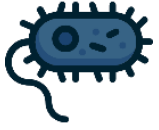




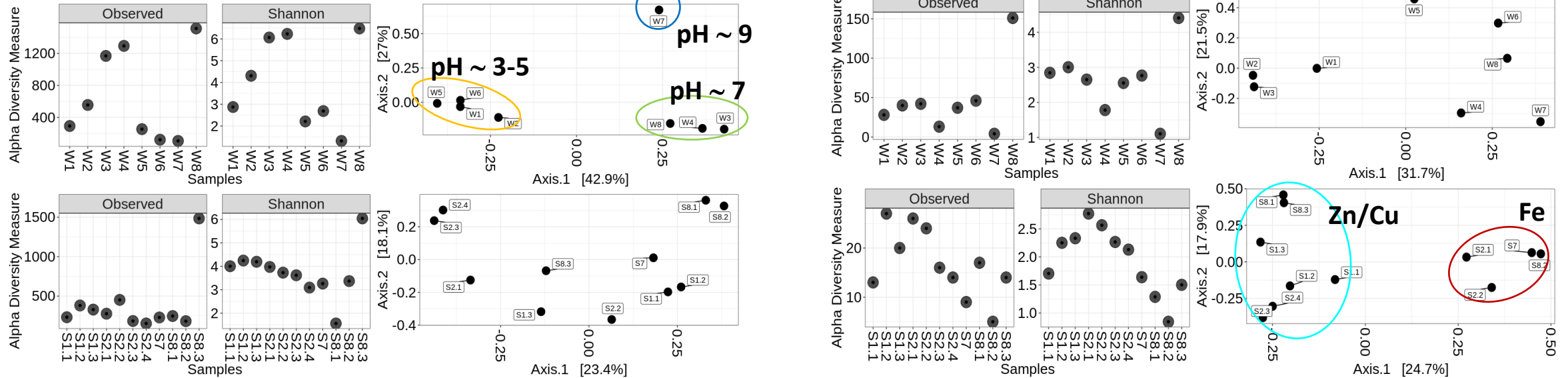
Microbial diversity



- Generally low microbial diversity was observed in all samples.
- High bacterial diversity was found in more neutral waters and sediment.
- The lowest microbial diversity was detected in sedimentation tank.
- PCoA showed clusters correlating with pH

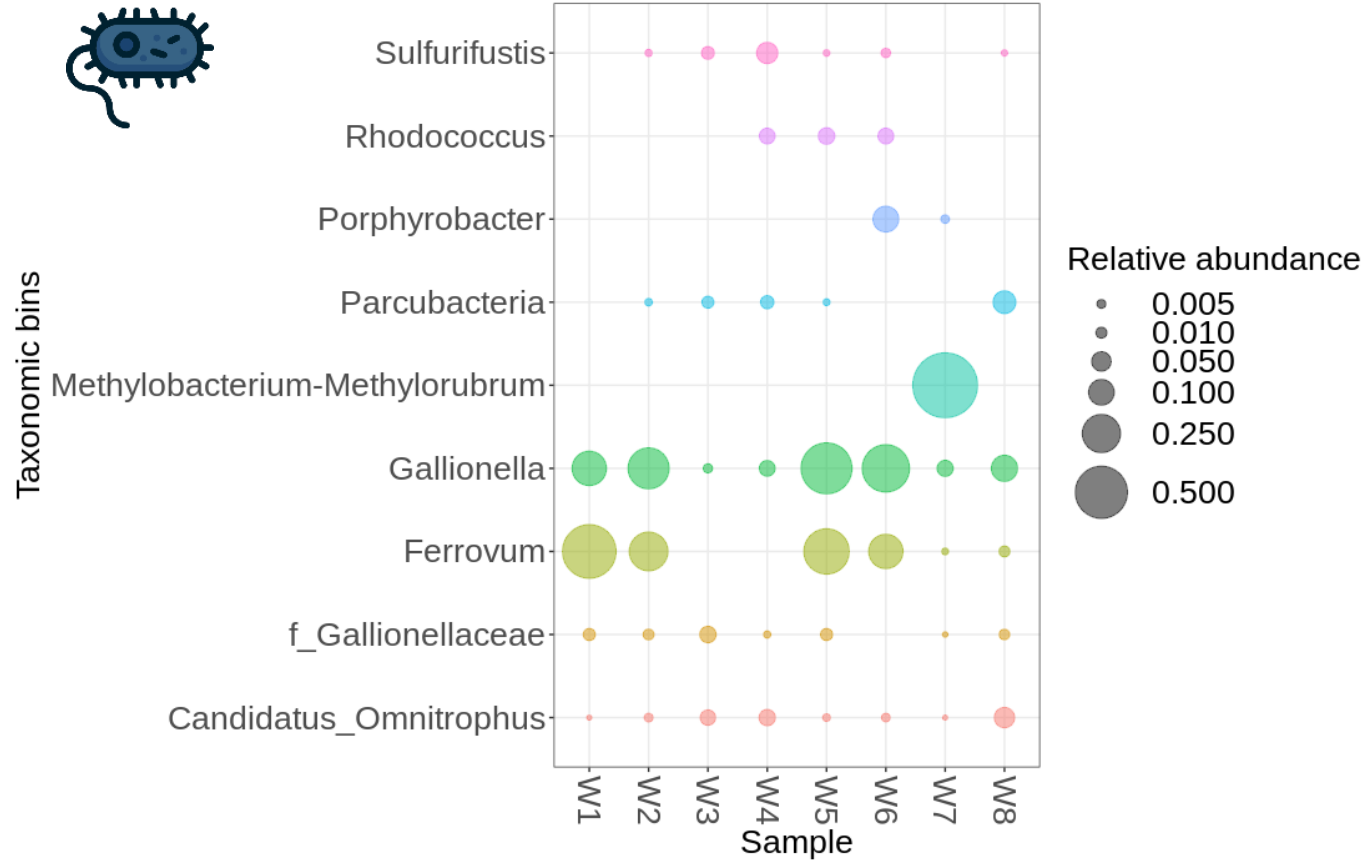


Microbial diversity

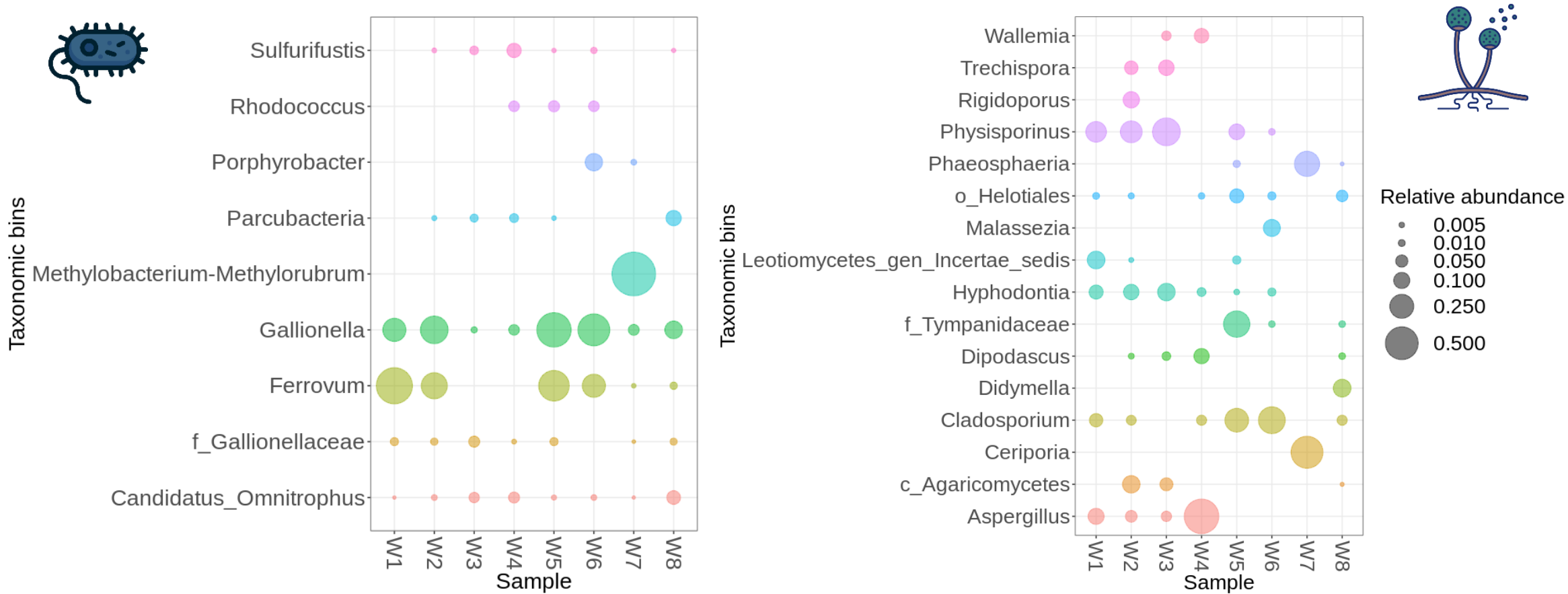


- Generally low microbial diversity was observed in all samples.
- High bacterial diversity was found in more neutral waters and sediment.
- The lowest microbial diversity was detected in sedimentation tank.
- PCoA showed clusters correlating with pH of waters or reflecting geochemical properties of sinters

Microbial community composition

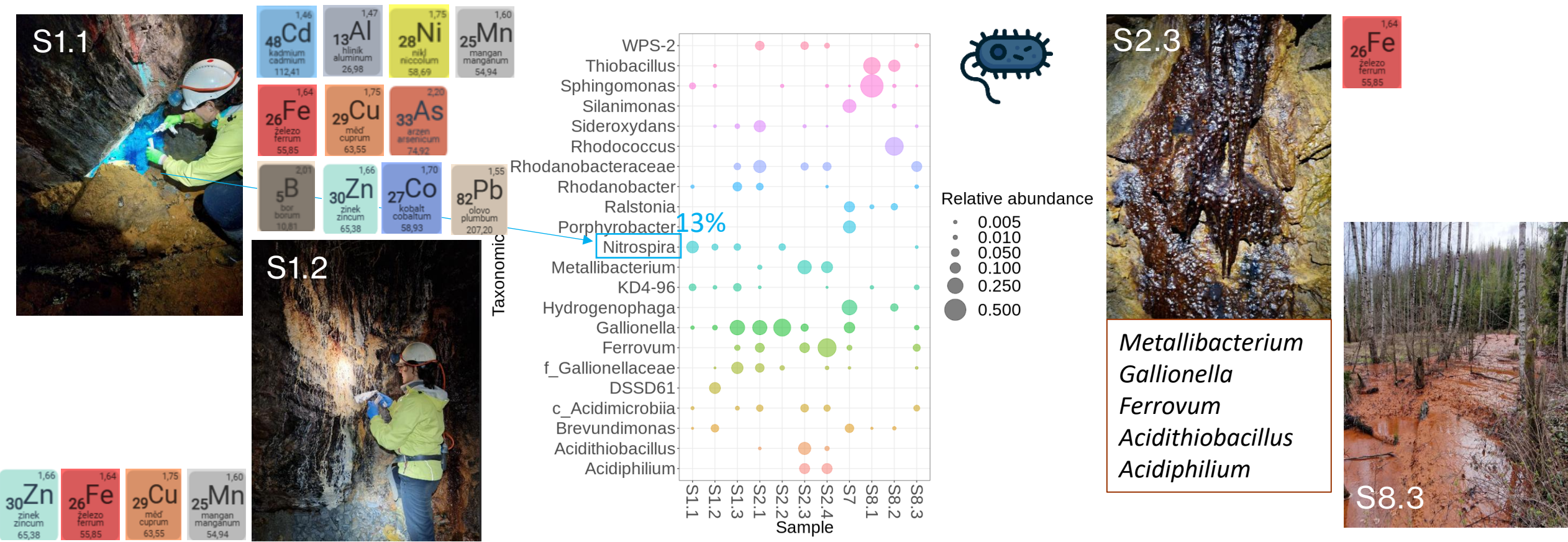


Microbial community composition



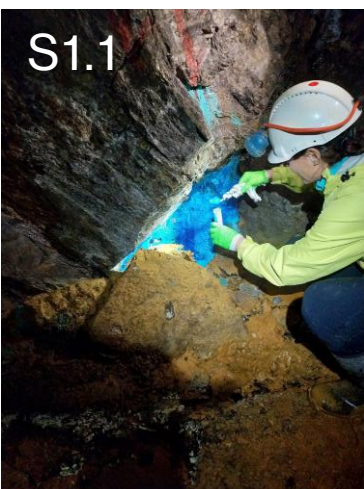
Microbial community composition

Solid samples: Bacterial community composition is dependent on geochemical characteristic of samples

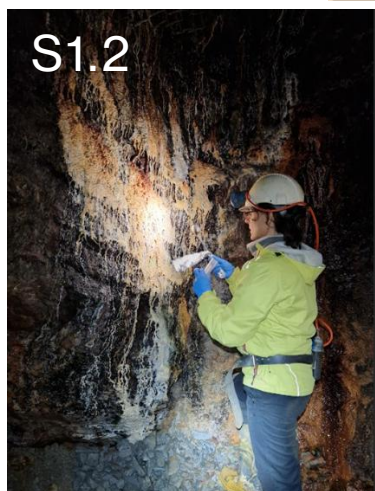


Microbial community composition

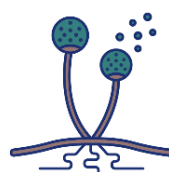
Solid samples: Bacterial community composition is dependent on geochemical characteristic of samples



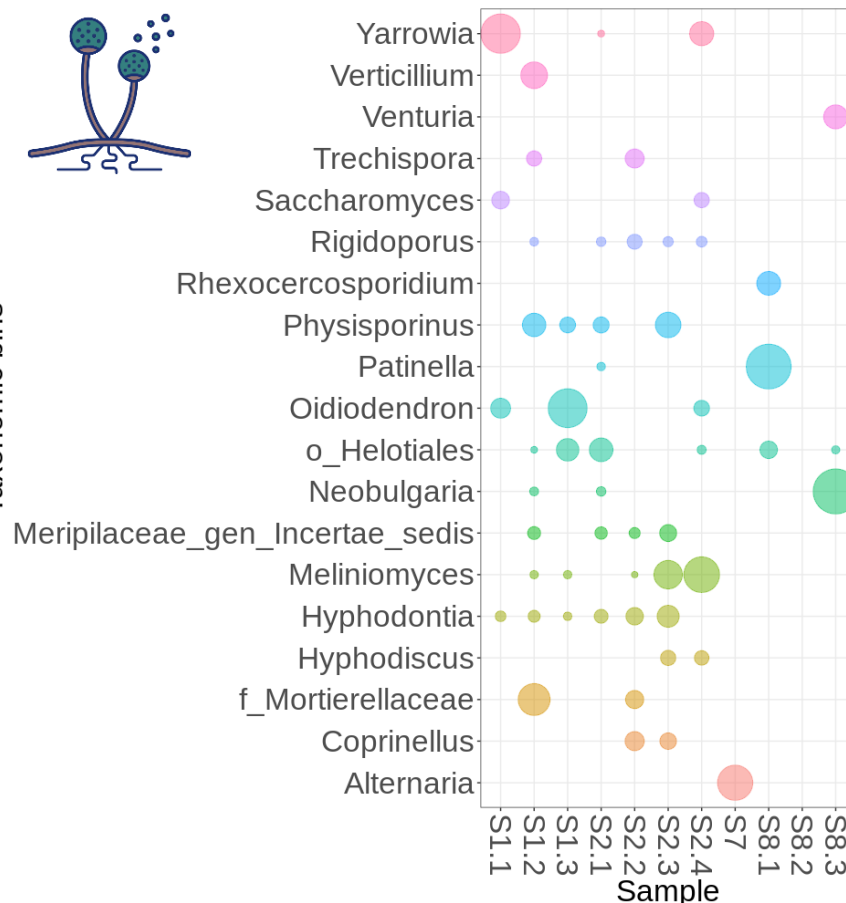
48Cd kadmium 112,41	13Al hliník 26,98	28Ni nikl 58,69	25Mn mangan 54,94
26Fe železo 55,85	29Cu měď 63,55	33As arzen 74,92	
5B bor 10,81	30Zn zinek 65,38	27Co kobalt 58,93	82Pb olovo 207,20



30Zn zinek 65,38	26Fe železo 55,85	29Cu měď 63,55	25Mn mangan 54,94
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Taxonomic bins

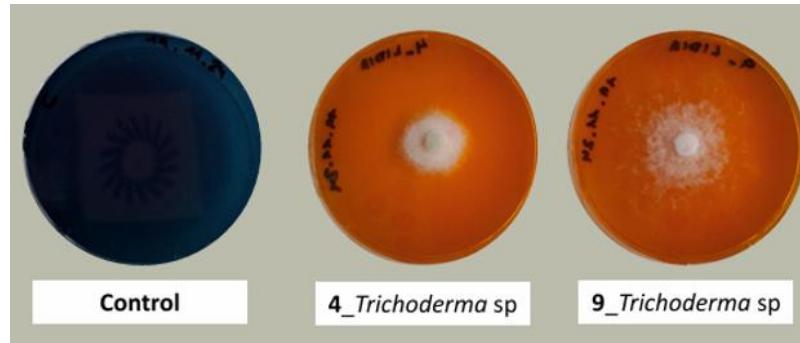


26Fe železo 55,85

Coprinellus
Hyphodiscus
Hyphodontia
Meliniomyces
Physisporinus



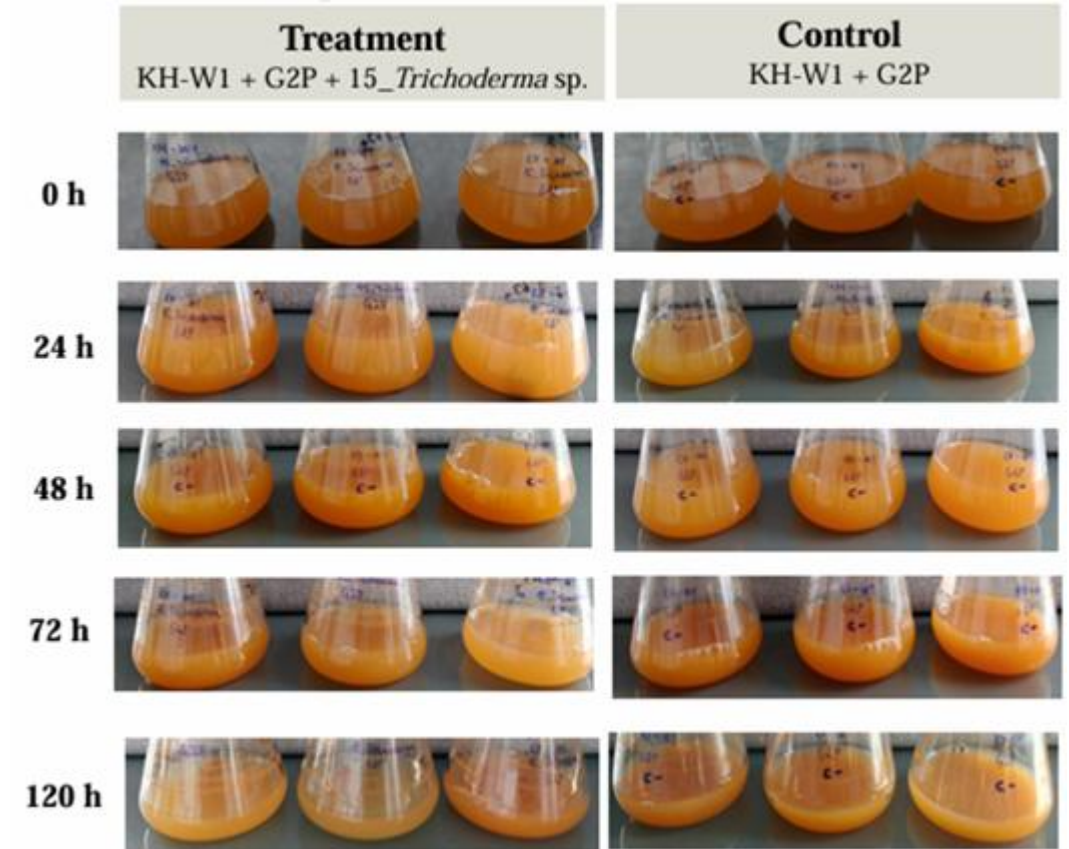
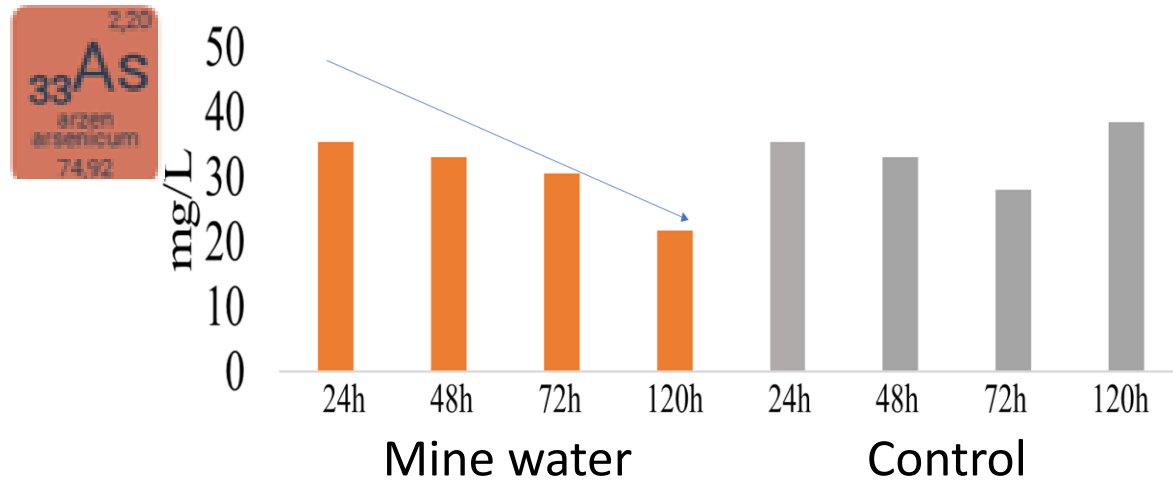
Removal of arsenic from mining water using *Trichoderma* sp.



Bromocresol green
assay →
organic acid
production
Initial pH: 6.5

S8.3

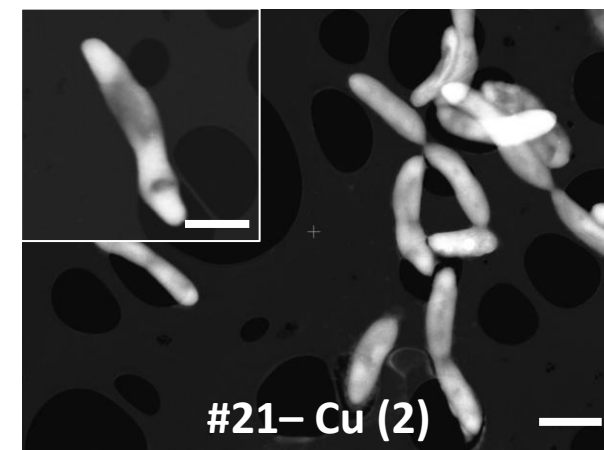
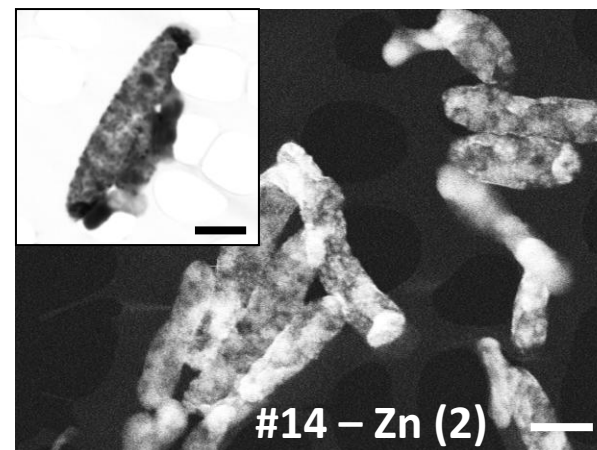
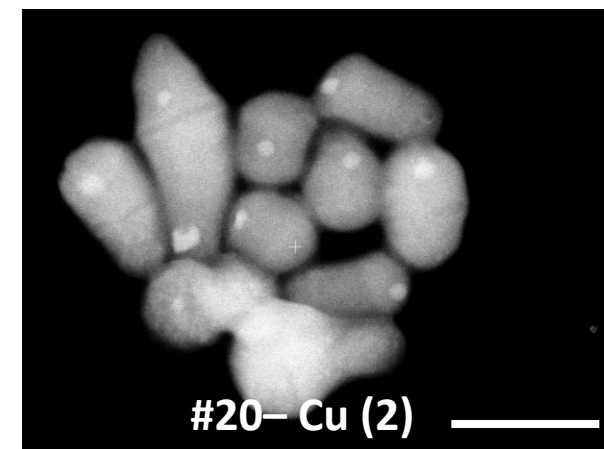
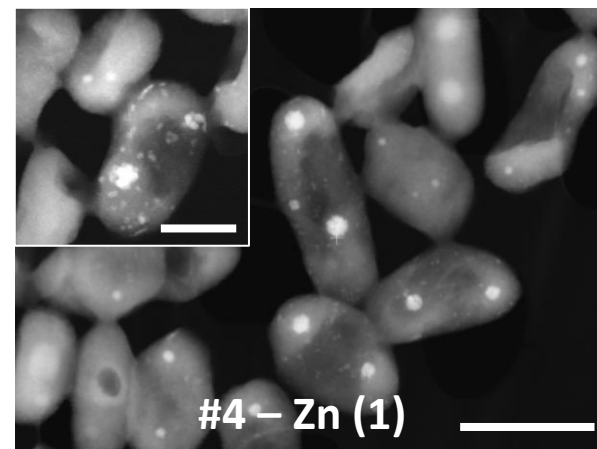
W1

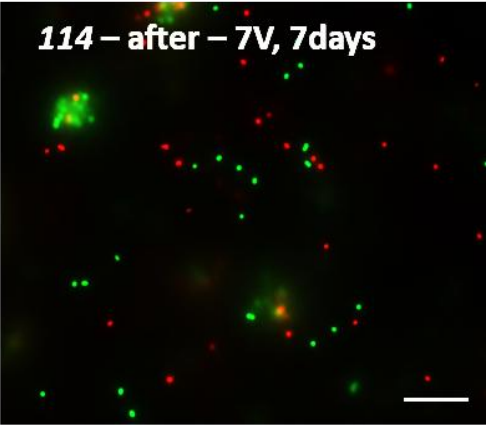
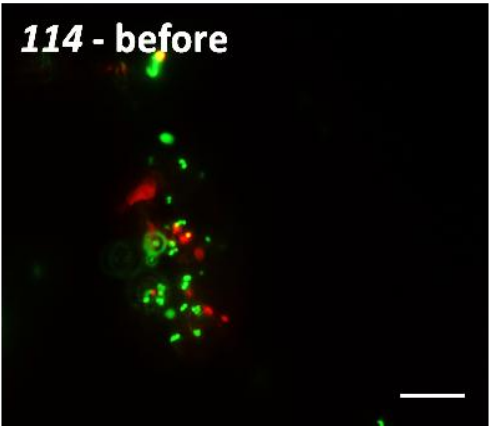


Bioaccumulation and Biosorption

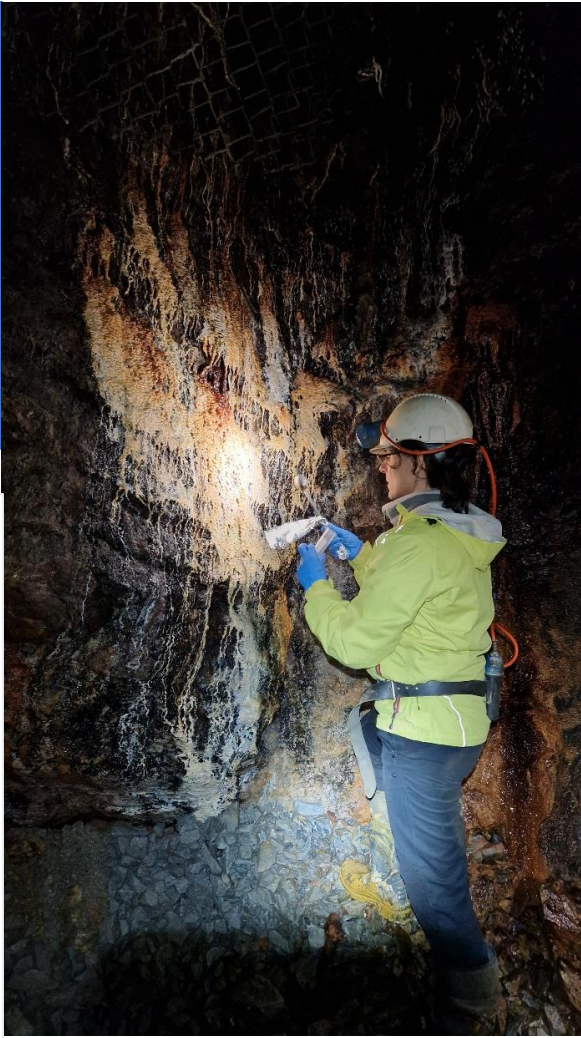


Culture #	Strain	% identity	Cu	Co	Zn	Pb	Se
4	<i>Variovorax</i> sp. SIB_Cd_R3	99.67	>4	n.d.	8	8	>8
14	<i>Microbaterium</i> sp. HaHa633	99.87	>4	>4	8	>8	>8
20	<i>Arthrobacter</i> sp. MB118	98.95	4	2	8	8	>8
21	<i>Pseudomonas</i> <i>mandelii</i>	100	>4	>4	8	8	16

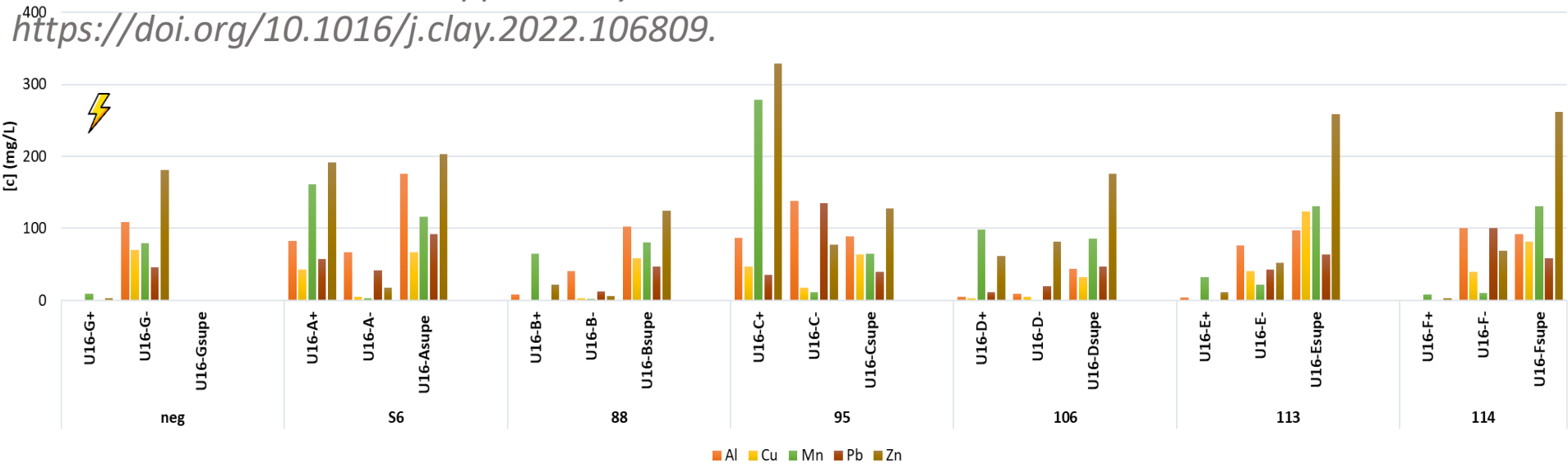




Bacteria	% Identity
<i>Variovorax</i> sp. SIB_Cd_R3	99.7
<i>Variovorax boronicumulans</i>	100
<i>Variovorax</i> sp. +	NA
<i>Variovorax boronicumulans</i>	100
<i>Microbacterium profundus</i>	99.8
<i>Microbaterium</i> sp. HaHa633	99.9



Hlavackova V. et al., 2023: *Applied Clay Science*, 232
<https://doi.org/10.1016/j.clay.2022.106809>.

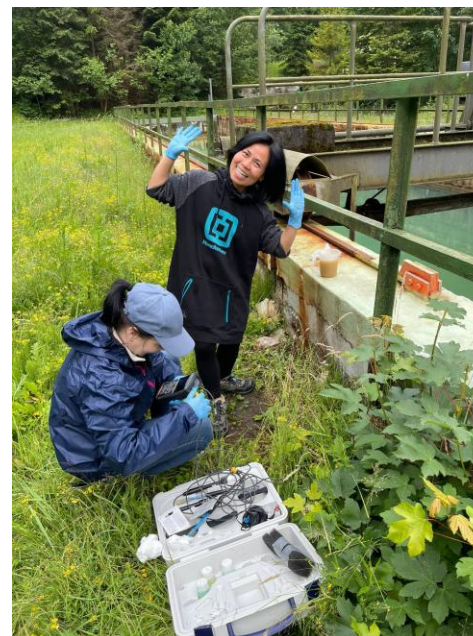


Conclusion

1. The microbial environment in postmining sites is highly selective and fragile.
2. Both microorganisms capable of leaching and bio-precipitation are already on site
3. Indigenous microbiome can be used for combined bio-electrochemical approaches
4. The comprehensive evaluation provides a framework for optimizing biorecovery technologies.



Thank you for your attention



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P. Kejzlar - SEM