

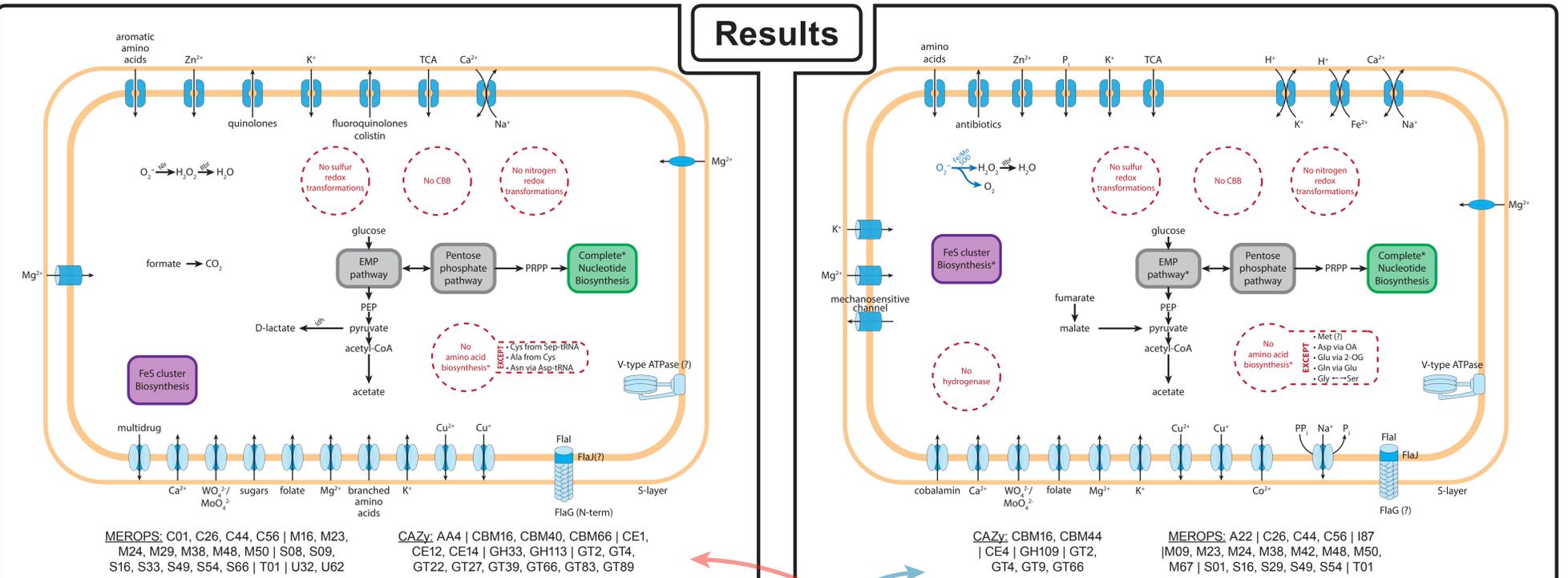
## Introduction / Motivation

- Mixed radiochemical (inc. <sup>239+240</sup>Pu and <sup>241</sup>Am) and organic/contaminated waste (steel drums, lab equipment, paper, etc.) was disposed of in shallow (3 m deep) trenches from 1960–1968.
- Previous research investigated functional and population dynamics during a rainfall event using shotgun metagenomics. This revealed a broad abundance of candidate and potentially undescribed taxa [1].

## Methods

- Assembly/Binning:** Megahit + CONCOCT + Anvi'o
- Phylogeny:** 44 concatenated ribosomal proteins (univ+arch). PMSF+C60+LG+F+I+G (IQ-TREE).
- Metabolic models:**
  - General: PathwayTools (MetaCyc) + KEGG + InterProScan + manual curation
  - Specific: dbCAN (CAZy) + MEROPS (proteases) + TCDB (transporters)

## Results



- No SOD, but neelaredoxin instead
- Formate dehydrogenase
- Facultative anaerobes or microaerophiles

Ca. 'Tiddalikarchaeum anstoanum' LFW-252\_1<sup>T</sup> CA00034

1 MAG

**LFWA-I**

75.3%C / 1.9%R  
5S, 16S, 23S<sup>P</sup>  
21/21 tRNA

- Facultative anaerobes

CA00041 Ca. 'Wianamattarchaeum fermentum' LFW-144\_1<sup>T</sup>

2 MAGs, 2 genera, 1 family

**LFWA-II**

85.8%C / 2.5%R  
5S, 16S, no 23S  
20/21 tRNA

**LFWA-III**

11 MAGs, 10 genera, 1 family

Ca. 'Gugararchaeum adminiculabundum' LFW-121\_3<sup>T</sup> CA00043

Ca. 'Wayembeharchaeum dharwalense' LFW-283\_2 [not shown] CA00042

- Facultative anaerobes or microaerophiles
- Biosynthesis of thiamine and riboflavin
- Complete *de novo* amino acid biosynthesis
- Partial TCA cycle

MEROPS: A08, A22 | C26, C44, C56, C82 | I39, I87 | M17, M20, M23, M24, M38, M48, M50, M79 | S08, S16, S33, S54 | T01, T05 | U32, U62  
CAZy: AA6 | CBM14 | CE4 | GH23, GH91, GH109 | GT2, GT4, GT19, GT66, GT83

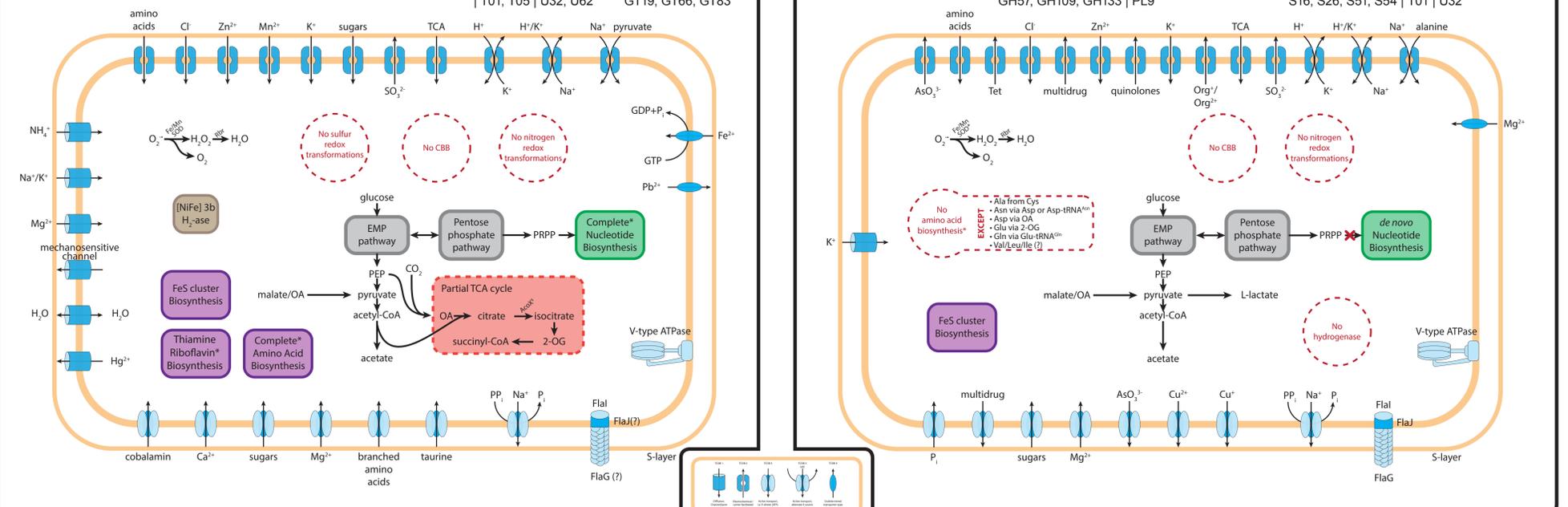
**LFWA-IV**

1 MAG

LFW-46<sup>T</sup>

- Facultative anaerobe
- High variety of antimicrobial resistance-related transporters, incl. organocations (e.g. quaternary amine detergents)
- Arsenite resistance
- Pectate lyase

CAZy: CBM16, CBM40 | CE3, CE4, CE7, CE14 | GH23, GH57, GH109, GH133 | PL9  
MEROPS: C26, C44, C45 | I51 | M14, M20, M24, M42, M48, M50 | S08, S09, S16, S26, S51, S54 | T01 | U32



## Contact

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## Conclusions

- 4 proposed new lineages in DPANN.
- All able to ferment, and facultative anaero/microaerophilic.
- LFWA-III: vitamin, aa and *de novo* nucleotide biosynthesis + partial TCA
- Rubredoxin + neelaredoxin or SOD

## References

[1] Vázquez-Campos, X. *et al.* 2017. *Appl Environ Microbiol*, 83(17), e00729. <https://doi.org/10.1128/AEM.00729-17>