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Contrasts between the cryoconite and ice-marginal bacterial communities of Svalbard glaciers

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Published in:
Polar Research

DOI:
[10.3402/polar.v32i0.19468](https://doi.org/10.3402/polar.v32i0.19468)

Publication date:
2013

Citation for published version (APA):

Edwards, A., Rassner, S. M., Anesio, A. M., Worgan, H. J., Irvine-Fynn, T. D., Williams, H. W., Sattler, B., & Griffith, G. W. (2013). Contrasts between the cryoconite and ice-marginal bacterial communities of Svalbard glaciers: Bacterial communities of Svalbard glaciers. *Polar Research*, 32, Article 19468.
<https://doi.org/10.3402/polar.v32i0.19468>

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Supplementary file for: Arwyn Edwards A., Rassner S.M., Anesio A.M., Worgan H.J., Irvine-Fynn T.D.L., Williams H.W., Sattler B. & Griffith G.W. 2013. Contrasts between the cryoconite and ice-marginal bacterial communities of Svalbard glaciers. *Polar Research* 32. Correspondence: Arwyn Edwards, Institute of Biological, Rural and Environmental Sciences, Cledwyn Building, Aberystwyth University, Aberystwyth SY23 3FG, UK. E-mail: aye@aber.ac.uk.

Supplementary methods

Terminal-restriction fragment (T-RF) relative abundances (in this instance normalized as ‰ integer values to satisfy formatting requirements) were plotted using the `radfit` function of the `Vegan` package (Oksanen et al. 2012) in the R 2.15.0 statistical environment (R Development Core Team 2012) to model broken-stick, pre-emption, log-normal, Zipf and Mandelbrot-Zipf relative abundance distributions against the T-RF relative abundance distributions. The fit of the zero-sum model to T-RF relative abundances was conducted using `TeTame` 2.1 (Jabot et al. 2008). The fit of all models to the T-RF data was evaluated using Akaike's Information Criterion (AIC; Akaike 1974) where the best-fitting model is accorded the lowest score. In the instance of zero-sum models, the output of `TeTame` 2.1 is provided as minimum log-likelihood values, requiring transformation to yield maximum likelihood values for calculation of AIC (Feinstein & Blackwood 2012).

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Supplementary Table S1. Akaike Information Criterion (AIC) values for model evaluation of terminal-restriction fragment (T-RF) abundance distribution; the lowest value, i.e., the best AIC score, for each sample is in boldface. The number of T-RFs per sample is indicated.

| Sample type | Sample | T-RFs | Null | Preemption | Lognormal | Zipf | Mandelbrot-Zipf | Zero-Sum |
|-------------|--------|-------|---------|------------|---------------|---------------|-----------------|----------|
| Cryoconite | MLC1 | 13 | 195.66 | 128.17 | 98.25 | 85.52 | 87.25 | 294.92 |
| Cryoconite | MLC2 | 12 | 243.44 | 173.50 | 151.33 | 108.02 | 109.45 | 259.35 |
| Cryoconite | MLC3 | 10 | 219.50 | 149.49 | 113.90 | 74.64 | 76.04 | 238.29 |
| Cryoconite | MLC4 | 32 | 262.23 | 254.57 | 200.83 | 197.05 | 194.28 | 414.95 |
| Cryoconite | MLC5 | 13 | 165.75 | 129.57 | 120.17 | 108.40 | 97.63 | 307.94 |
| Cryoconite | MLC6 | 21 | 248.51 | 165.82 | 136.74 | 123.48 | 121.24 | 293.07 |
| Cryoconite | MLC8 | 7 | 132.14 | 81.96 | 86.83 | 102.60 | 74.45 | 161.80 |
| Cryoconite | MLC9 | 15 | 200.14 | 185.40 | 137.31 | 103.64 | 105.64 | 336.73 |
| Cryoconite | MLC10 | 15 | 250.82 | 225.23 | 175.71 | 120.58 | 122.58 | 333.19 |
| Cryoconite | VBC1 | 30 | 344.25 | 305.08 | 219.69 | 168.41 | 170.41 | 445.20 |
| Cryoconite | VBC2 | 23 | 246.26 | 197.10 | 194.45 | 192.78 | 160.42 | 329.75 |
| Cryoconite | VBC3 | 18 | 180.53 | 128.35 | 136.93 | 147.56 | 114.15 | 424.30 |
| Cryoconite | VBC4 | 36 | 263.55 | 227.69 | 229.63 | 252.67 | 194.17 | 463.46 |
| Cryoconite | VBC5 | 20 | 229.97 | 203.15 | 158.32 | 128.54 | 127.17 | 295.16 |
| Cryoconite | VBC6 | 21 | 238.95 | 251.03 | 163.38 | 120.16 | 122.16 | 313.03 |
| Cryoconite | ABC1 | 10 | 208.02 | 141.49 | 105.01 | 82.77 | 84.77 | 242.13 |
| Cryoconite | ABC2 | 11 | 164.56 | 96.35 | 86.05 | 84.06 | 76.71 | 262.49 |
| Cryoconite | ABC3 | 9 | 250.72 | 102.13 | 79.79 | 94.65 | 74.28 | 211.47 |
| Cryoconite | ABC4 | 53 | 395.84 | 385.98 | 315.72 | 298.85 | 258.19 | 626.71 |
| Cryoconite | ABC5 | 62 | 387.31 | 377.56 | 312.62 | 304.04 | 286.30 | 745.65 |
| Cryoconite | ABC6 | 18 | 216.14 | 127.17 | 135.91 | 158.69 | 119.23 | 271.45 |
| Cryoconite | ABC7 | 18 | 213.36 | 122.74 | 131.74 | 154.08 | 114.25 | 272.73 |
| Cryoconite | ABC8 | 17 | 168.61 | 147.60 | 139.01 | 138.73 | 113.53 | 255.74 |
| Cryoconite | ABC9 | 18 | 212.21 | 188.12 | 170.53 | 152.78 | 132.30 | 284.67 |
| Cryoconite | ABC10 | 13 | 134.97 | 102.79 | 110.89 | 132.33 | 97.61 | 314.47 |
| Soil | ETT1 | 6 | 79.24 | 67.18 | 67.62 | 60.30 | 54.80 | 150.17 |
| Soil | ETT2 | 5 | 119.55 | 103.18 | 120.15 | 127.10 | 102.57 | 134.59 |
| Soil | ETT3 | 28 | 396.16 | 226.80 | 180.68 | 158.35 | 160.04 | 432.90 |
| Soil | ETT4 | 10 | 212.15 | 139.65 | 104.03 | 81.20 | 83.20 | 271.44 |
| Soil | ETT5 | 3 | 100.06 | 74.36 | 38.09 | 52.08 | 40.09 | 66.15 |
| Soil | SVT1 | 6 | 119.59 | 92.03 | 105.11 | 135.41 | 87.48 | 151.54 |
| Soil | SVT2 | 5 | 74.22 | 66.84 | 62.58 | 51.69 | 51.78 | 119.99 |
| Soil | SVT3 | 14 | 1535.77 | 833.76 | 400.86 | 258.06 | 260.06 | 350.28 |
| Soil | SVT5 | 5 | 177.17 | 112.53 | 77.03 | 58.90 | 60.90 | 121.35 |
| Soil | ABM1 | 44 | 322.99 | 333.76 | 243.52 | 217.44 | 215.92 | 567.43 |
| Soil | ABM2 | 34 | 352.99 | 202.30 | 189.93 | 202.36 | 179.64 | 464.71 |
| Soil | ABM3 | 26 | 359.46 | 190.64 | 153.04 | 153.22 | 153.64 | 395.26 |
| Soil | ABM5 | 25 | 381.16 | 192.10 | 153.36 | 142.51 | 142.69 | 406.87 |
| Soil | VBM1 | 19 | 344.29 | 359.70 | 223.87 | 130.58 | 132.58 | 487.26 |
| Soil | VBM2 | 23 | 328.71 | 206.56 | 160.29 | 131.53 | 133.53 | 384.74 |
| Soil | VBM3 | 19 | 322.53 | 233.11 | 175.22 | 121.64 | 123.64 | 491.59 |
| Soil | VBM4 | 22 | 318.18 | 147.92 | 140.20 | 166.27 | 130.35 | 358.95 |
| Soil | VBM5 | 24 | 355.58 | 165.84 | 143.87 | 163.29 | 142.89 | 380.19 |
| Soil | MLM1 | 46 | 354.27 | 336.15 | 260.65 | 231.96 | 231.35 | 626.24 |
| Soil | MLM2 | 23 | 268.24 | 194.70 | 159.24 | 137.75 | 137.09 | 362.78 |
| Soil | MLM3 | 23 | 404.26 | 216.72 | 165.52 | 139.94 | 141.94 | 361.88 |
| Soil | MLM4 | 20 | 340.95 | 261.47 | 188.97 | 133.23 | 135.23 | 509.50 |
| Soil | MLM5 | 11 | 183.94 | 129.00 | 107.72 | 86.85 | 87.32 | 294.08 |

Supplementary Table S2. Permutational analysis of variance (PERMANOVA) of bacterial 16S presence–absence terminal-restriction fragment length polymorphism (T-RFLP) profiles; $p(\text{perm})$ values of less than 0.01 have been highlighted in bold to indicate highly significant differences between sample groups upon pairwise PERMANOVA.

| | | Pairwise PERMANOVA $p(\text{perm})$ values) | | | | | | |
|------------|--|---|-----------------|-----------------|-----------------|-----------------|---------|---------|
| | | ML ^a | VB ^b | AB ^c | ET ^d | SV ^d | AB | VB |
| | | cryoconite | cryoconite | cryoconite | tundra | tundra | moraine | moraine |
| ML | | | | | | | | |
| cryoconite | | | | | | | | |
| VB | | | | | | | | |
| cryoconite | | 0.0008 | | | | | | |
| AB | | | | | | | | |
| cryoconite | | 0.0001 | 0.0102 | | | | | |
| ET | | | | | | | | |
| tundra | | 0.0005 | 0.0019 | 0.0008 | | | | |
| SV | | | | | | | | |
| tundra | | 0.0016 | 0.0051 | 0.0011 | 0.6423 | | | |
| AB | | | | | | | | |
| moraine | | 0.0016 | 0.0054 | 0.0009 | 0.0171 | 0.028 | | |
| VB | | | | | | | | |
| moraine | | 0.0003 | 0.0013 | 0.0006 | 0.0141 | 0.0251 | 0.0569 | |
| ML | | | | | | | | |
| moraine | | 0.0012 | 0.0024 | 0.0004 | 0.0312 | 0.0379 | 0.7912 | 0.235 |

^aMidtre Lovénbreen

^bVestre Brøggerbreen

^cAustre Brøggerbreen

^dSee Fig. 1 for location of the two tundra sites.