

Supporting information for:

Changes of microorganism composition in fresh and stored bee pollen from Southern Germany

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Tables S2-S5 with coding:

First: F = fresh; C = cold; RT = room temperature; W = warm

Second: Numbers 1-3 for replicates

Third: H = Hohenheim; F = Forbach; N = Nuertingen

Table S1 Detailed palynological analysis in all samples

| classification | Hohenheim 2018 | Forbach 2019 | Nuertingen 2019 |
|-----------------------------------|----------------|--------------|-----------------|
| <i>Aceraceae_Acer</i> | | 208 | |
| <i>Sapindaceae_Aesculus</i> | 2 | 49 | |
| <i>Apiaceae</i> | | | 4 |
| <i>Rosaceae_Aruncus dioicus</i> | | 2 | 209 |
| <i>Asteraceae_Bidens-T</i> | 2 | | |
| <i>Brassicaceae_Brassica-T</i> | 38 | | |
| <i>Asteraceae_Carduus</i> | 22 | | |
| <i>Cyperaceae_Carex</i> | | 8 | |
| <i>Fagaceae_Castanea sativa</i> | 4 | | |
| <i>Chenopodiaceae</i> | 2 | | |
| <i>Cornaceae_Cornus-T</i> | | 3 | |
| <i>Boraginaceae_Echium</i> | 8 | | |
| <i>Fabaceae</i> | | 34 | |
| <i>Rosaceae_Filipendula</i> | 52 | | 10 |
| <i>Rosaceae_Fragaria</i> | | | 2 |
| <i>Apiaceae_Heracleum</i> | | 12 | |
| <i>Cannabaceae_Humulus</i> | 2 | | |
| <i>Clusiaceae_Hypericum</i> | 1 | 4 | 3 |
| <i>Balsaminaceae_Impatiens</i> | 1 | | |
| <i>Lamiaceae_Lamium-T</i> | | 3 | 14 |
| <i>Oleaceae_Ligustrum</i> | | 1 | 78 |
| <i>Fabaceae_Melilotus</i> | 1 | 2 | |
| <i>Papaveraceae</i> | | 1 | 4 |
| <i>Boraginaceae_Phacelia</i> | 96 | | |
| <i>Pinaceae_Pinoideae</i> | | 31 | |
| <i>Plantaginaceae_Plantago</i> | 72 | 17 | 24 |
| <i>Poaceae</i> | 8 | | 2 |
| <i>Rosaceae_Potentilla</i> | | | 44 |
| <i>Ranunculaceae</i> | | 59 | |
| <i>Orobanchaceae_Rhinanthus</i> | | | 16 |
| <i>Anacardiaceae_Rhus typhina</i> | | | 2 |
| <i>Rosaceae_Rosa</i> | | | 41 |
| <i>Rosaceae_Rubus-T</i> | 3 | 6 | 1 |
| <i>Polygonaceae_Rumex</i> | 2 | | 3 |
| <i>Rosaceae_Sanguisorba minor</i> | | 16 | 1 |
| <i>Cyperaceae_Scirpus</i> | | 35 | |
| <i>Crassulaceae_Sedum</i> | 44 | | |
| <i>Brassicaceae_Sinapis-T</i> | 51 | | |
| <i>Asteraceae_Taraxacum-T</i> | 27 | | |
| <i>Malvaceae_Tilia</i> | 39 | | 39 |
| <i>Fabaceae_Trifolium repens</i> | 18 | 1 | 3 |
| <i>Adoxaceae_Viburnum</i> | | 8 | |
| <i>Poaceae_Zea mays</i> | 5 | | |

Table S2 Calculated relative abundance of all detected bacteria in the samples

| F1H | F2H | F3H | C1H | C2H | C3H | RT1H | RT2H | W1H | W2H | W3H | taxonomy |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.4335 | 0.6673 | 0.8104 | 0.5924 | 0.7750 | 0.6638 | 0.8669 | 0.9062 | 0.5248 | 0.0748 | 0.5476 | Bacteria;Firmicutes;Bacilli;Lactobacillales;Lactobacillaceae;Lactobacillus; |
| 0.1430 | 0.0173 | 0.0255 | 0.0070 | 0.0255 | 0.0864 | 0.0149 | 0.0067 | 0.0011 | 0 | 0.0174 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Pseudomonadaceae;Pseudomonas; |
| 0.0256 | 0.0505 | 0.0043 | 0.2206 | 0.0079 | 0.0834 | 0.0022 | 0.0033 | 0.0246 | 0.0200 | 0.2077 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Moraxellaceae;Acinetobacter; |
| 0.1204 | 0.0038 | 0.0084 | 0.0041 | 0.0025 | 0.0064 | 0.0976 | 0.0012 | 0 | 0 | 0.0321 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Rosenbergiella; |
| 0.0962 | 0.0874 | 0.0167 | 0.0070 | 0.0059 | 0.0787 | 0.0016 | 0.0014 | 0.0089 | 0 | 0.0056 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Erwiniaceae;Pantoea |
| 0.0124 | 0.0117 | 0.0053 | 0.0014 | 0.0032 | 0.0045 | 0.0001 | 0.0006 | 0.3230 | 0.7537 | 0.0683 | Bacteria;Actinobacteria;Actinobacteria;Actinomycetales;Micrococcaceae;Arthrobacter; |
| 0.0205 | 0.0085 | 0.0054 | 0.0106 | 0.0121 | 0.0163 | 0.0004 | 0.0095 | 0.0051 | 0 | 0.0137 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Erwinia; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Epilithonimonas; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Erwiniaceae;Phaseolibacter; |
| 0.0106 | 0.0030 | 0.0828 | 0.0004 | 0.0015 | 0.0026 | 0.0002 | 0.0007 | 0 | 0.0002 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Serratia; |
| 0.0538 | 0.0566 | 0.0148 | 0.0695 | 0.0862 | 0.0323 | 0.0024 | 0.0297 | 0.0212 | 0.0001 | 0.0249 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Arsenophonus; |
| 0.0118 | 0.0357 | 0.0068 | 0.0134 | 0.0211 | 0.0103 | 0.0086 | 0.0127 | 0.0080 | 0 | 0.0235 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodospirillales;Acetobacteraceae;Saccharibacter; |
| 0.0097 | 0.0130 | 0.0019 | 0.0236 | 0.0059 | 0.0034 | 0.0017 | 0.0096 | 0.0246 | 0.0215 | 0.0194 | Bacteria;Proteobacteria;Alphaproteobacteria;Sphingomonadales;Sphingomonadaceae;Sphingomonas; |
| 0.0005 | 0 | 0 | 0.0006 | 0.0002 | 0.0001 | 0.0001 | 0.0038 | 0 | 0 | 0 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Chryseobacterium; |
| 0.0063 | 0.0034 | 0.0012 | 0.0084 | 0.0027 | 0.0039 | 0.0007 | 0.0033 | 0.0159 | 0.0210 | 0.0132 | Bacteria;Proteobacteria;Betaproteobacteria;Burkholderiales;Oxalobacteraceae;Massilia; |
| 0.0004 | 0.0035 | 0.0006 | 0.0015 | 0.0097 | 0.0005 | 0.0000 | 0.0032 | 0 | 0 | 0.0011 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodospirillales;Acetobacteraceae;Gluconacetobacter; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Betaproteobacteria;Burkholderiales;Oxalobacteraceae;Duganella; |
| 0.0018 | 0.0016 | 0.0002 | 0.0018 | 0.0009 | 0.0001 | 0.0001 | 0.0005 | 0.0019 | 0.0101 | 0.0017 | Bacteria;Bacteroidetes;Sphingobacteriia;Sphingobacteriales;Sphingobacteriaceae;Pedobacter; |
| 0.0022 | 0.0085 | 0.0044 | 0.0081 | 0.0095 | 0.0043 | 0.0006 | 0.0013 | 0 | 0.0096 | 0.0064 | Bacteria;Proteobacteria;Gammaproteobacteria;Orbales;Orbaceae;Gilliamella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Cyanobacteria;Cyanobacteria;Synechococcales;Synechococcaceae;Cyanobium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.0114 | 0.0362 | 0 | Bacteria;Firmicutes;Clostridia;Clostridiales;Lachnospiraceae;Fusicatenibacter; |
| 0.0017 | 0.0085 | 0.0008 | 0.0056 | 0.0032 | 0.0006 | 0.0005 | 0.0020 | 0.0234 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Oceanospirillales;Halomonadaceae;Carnimonas; |
| 0.0003 | 0.0026 | 0.0002 | 0.0027 | 0.0046 | 0.0006 | 0.0001 | 0.0010 | 0.0060 | 0.0154 | 0.0079 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Escherichia/Shigella; |
| 0.0170 | 0.0081 | 0.0029 | 0.0014 | 0.0006 | 0.0006 | 0.0006 | 0.0004 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Citrobacter; |
| 0.0127 | 0 | 0.0004 | 0.0005 | 0.0006 | 0.0001 | 0.0000 | 0.0005 | 0 | 0.0139 | 0.0007 | Bacteria;Firmicutes;Bacilli;Lactobacillales;Streptococcaceae;Lactococcus; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhizobiales;Bradyrhizobiaceae;Bradyrhizobium; |

| | | | | | | | | | | | |
|--------|--------|--------|--------|--------|--------|--------|--------|---|--------|--------|---|
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.0182 | 0.0011 | Bacteria;Bacteroidetes;Bacteroidia;Bacteroidales;Bacteroidaceae;Bacteroides; |
| 0.0160 | 0.0004 | 0.0012 | 0.0002 | 0.0005 | 0.0002 | 0.0001 | 0.0003 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rickettsiales;Rickettsiaceae;Rickettsia; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodobacterales;Rhodobacteraceae;Actibacterium |
| 0.0012 | 0.0023 | 0.0019 | 0.0043 | 0.0039 | 0.0003 | 0.0000 | 0.0006 | 0 | 0 | 0.0010 | Bacteria;Proteobacteria;Betaproteobacteria;Neisseriales;Neisseriaceae;Snodgrassella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Pectobacterium |
| 0.0008 | 0.0027 | 0.0002 | 0.0036 | 0.0007 | 0.0001 | 0.0001 | 0.0003 | 0 | 0.0001 | 0.0035 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodospirillales;Acetobacteraceae;Neokomagataea; |
| 0.0011 | 0.0012 | 0.0013 | 0.0016 | 0.0033 | 0.0003 | 0.0002 | 0.0010 | 0 | 0 | 0.0015 | Bacteria;Proteobacteria;Gammaproteobacteria;Orbales;Orbaceae;Frischella; |
| 0.0004 | 0.0015 | 0.0024 | 0.0005 | 0.0050 | 0.0002 | 0.0001 | 0.0004 | 0 | 0 | 0.0005 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhizobiales;Bartonellaceae;Batronella; |
| 0 | 0.0003 | 0 | 0.0001 | 0.0075 | 0 | 0.0000 | 0.0000 | 0 | 0 | 0 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Apibacter; |
| 0.0002 | 0.0007 | 0.0000 | 0.0002 | 0.0002 | 0.0001 | 0.0000 | 0.0001 | 0 | 0.0052 | 0 | Bacteria;Firmicutes;Bacilli;Bacillales;Staphylococcaceae;Staphylococcus; |
| 0 | 0 | 0 | 0.0051 | 0 | 0.0001 | 0 | 0 | 0 | 0 | 0.0012 | Bacteria;Proteobacteria;Gammaproteobacteria;Oceanospirillales;Halomonadaceae;Halotalea; |
| 0 | 0 | 0 | 0.0039 | 0 | 0 | 0.0000 | 0.0000 | 0 | 0 | 0 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Flavobacterium; |

| F1F | F2F | F3F | C1F | C2F | C3F | RT1F | RT2F | RT3F | W1F | W2F | W3F | taxonomy |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.5824 | 0.8154 | 0.7050 | 0.7270 | 0.4291 | 0.6423 | 0.5668 | 0.5939 | 0.6394 | 0.9061 | 0.5689 | 0.8142 | Bacteria;Firmicutes;Bacilli;Lactobacillales;Lactobacillaceae;Lactobacillus; |
| 0.1086 | 0.0126 | 0.0356 | 0.0557 | 0.1557 | 0.1447 | 0.1173 | 0.1578 | 0.1004 | 0.0211 | 0.0641 | 0.0277 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Pseudomonadaceae;Pseudomonas; |
| 0.0233 | 0.0041 | 0.0137 | 0.0192 | 0.0062 | 0.0046 | 0.0398 | 0.0150 | 0.0285 | 0.0005 | 0.1711 | 0.0076 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Moraxellaceae;Acinetobacter; |
| 0.1586 | 0.0390 | 0.1071 | 0.0515 | 0.0477 | 0.0603 | 0.1209 | 0.0730 | 0.0275 | 0.0275 | 0.0572 | 0.0423 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Rosenbergiella; |
| 0.0139 | 0.0043 | 0.0072 | 0.0461 | 0.0092 | 0.0252 | 0.0072 | 0.0189 | 0.0017 | 0.0055 | 0.0016 | 0.0174 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Erwiniaceae;Pantoea |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Actinobacteria;Actinobacteria;Actinomycetales;Micrococcaceae;Arthrobacter; |
| 0.0080 | 0.0046 | 0.0436 | 0.0050 | 0.0101 | 0.0226 | 0.0202 | 0.0183 | 0.0062 | 0.0005 | 0.0276 | 0.0024 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Erwinia; |
| 0 | 0 | 0 | 0 | 0.0003 | 0.0003 | 0 | 0 | 0.0002 | 0 | 0 | 0.0003 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Epilithonimonas; |
| 0.0417 | 0.0623 | 0.0432 | 0.0398 | 0.2524 | 0.0529 | 0.0583 | 0.0469 | 0.0449 | 0.0033 | 0.0328 | 0.0484 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Erwiniaceae;Phaseolibacter; |
| 0.0242 | 0.0151 | 0.0104 | 0.0231 | 0.0359 | 0.0037 | 0.0101 | 0.0111 | 0.1169 | 0.0008 | 0.0126 | 0.0049 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Serratia; |
| 0 | 0.0006 | 0.0006 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.0004 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Arsenophonus; |
| 0.0116 | 0.0351 | 0.0259 | 0.0195 | 0.0096 | 0.0203 | 0.0081 | 0.0274 | 0.0212 | 0.0168 | 0.0126 | 0.0183 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodospirillales;Acetobacteraceae;Saccharibacter; |
| 0.0057 | 0.0007 | 0.0012 | 0.0021 | 0.0039 | 0.0057 | 0.0046 | 0.0124 | 0.0031 | 0.0025 | 0.0191 | 0.0058 | Bacteria;Proteobacteria;Alphaproteobacteria;Sphingomonadales;Sphingomonadaceae;Sphingomonas; |
| 0.0001 | 0 | 0 | 0 | 0 | 0 | 0 | 0.0007 | 0 | 0 | 0.0004 | 0.0003 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Chryseobacterium; |
| 0.0105 | 0.0041 | 0.0050 | 0.0067 | 0.0140 | 0.0129 | 0.0117 | 0.0176 | 0.0077 | 0.0146 | 0.0195 | 0.0058 | Bacteria;Proteobacteria;Betaproteobacteria;Burkholderiales;Oxalobacteraceae;Massilia; |
| 0 | 0.0011 | 0 | 0 | 0.0002 | 0 | 0 | 0 | 0 | 0.0005 | 0.0004 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodospirillales;Acetobacteraceae;Gluconacetobacter; |
| 0.0050 | 0.0004 | 0.0001 | 0.0011 | 0.0017 | 0.0017 | 0.0111 | 0.0065 | 0.0013 | 0 | 0.0032 | 0.0006 | Bacteria;Proteobacteria;Betaproteobacteria;Burkholderiales;Oxalobacteraceae;Duganella; |
| 0.0003 | 0.0002 | 0 | 0.0007 | 0 | 0.0026 | 0.0238 | 0.0007 | 0.0010 | 0.0003 | 0.0004 | 0 | Bacteria;Bacteroidetes;Sphingobacteriia;Sphingobacteriales;Sphingobacteriaceae;Pedobacter; |
| 0.0001 | 0.0002 | 0.0001 | 0 | 0.0226 | 0.0003 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Orbales;Orbaceae;Gilliamella; |
| 0.0060 | 0.0002 | 0.0013 | 0.0021 | 0 | 0 | 0 | 0 | 0 | 0 | 0.0077 | 0.0040 | Bacteria;Cyanobacteria;Cyanobacteria;Synechococcales;Synechococcaceae;Cyanobium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Firmicutes;Clostridia;Clostridiales;Lachnospiraceae;Fusicatenibacter; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Oceanospirillales;Halomonadaceae;Carnimonas; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Escherichia/Shigella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Citrobacter; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Firmicutes;Bacilli;Lactobacillales;Streptococcaceae;Lactococcus; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhizobiales;Bradyrhizobiaceae;Bradyrhizobium; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Bacteroidetes;Bacteroidia;Bacteroidales;Bacteroidaceae;Bacteroides; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rickettsiales;Rickettsiaceae;Rickettsia; |

| | | | | | | | | | | | | |
|---|---|---|--------|--------|---|---|---|---|---|--------|---|---|
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodobacterales;Rhodobacteraceae;Actibacterium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Betaproteobacteria;Neisseriales;Neisseriaceae;Snodgrassella; |
| 0 | 0 | 0 | 0.0004 | 0.0014 | 0 | 0 | 0 | 0 | 0 | 0.0004 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Pectobacterium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodospirillales;Acetobacteraceae;Neokomagataea; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Orbales;Orbaceae;Frischella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhizobiales;Bartonellaceae;Bartonella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Apiibacter; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Firmicutes;Bacilli;Bacillales;Staphylococcaceae;Staphylococcus; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Oceanospirillales;Halomonadaceae;Halotalea; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Flavobacterium; |

| F1N | F2N | F3N | C1N | C2N | C3N | RT1N | RT2N | RT3N | W1N | W2N | W3N | taxonomy |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.0479 | 0.0035 | 0.0143 | 0.0459 | 0.0570 | 0.0135 | 0.0015 | 0.0062 | 0.0524 | 0.0348 | 0.0273 | 0.0045 | Bacteria;Firmicutes;Bacilli;Lactobacillales;Lactobacillaceae;Lactobacillus; |
| 0.3984 | 0.1424 | 0.5023 | 0.3574 | 0.5189 | 0.3670 | 0.2967 | 0.2924 | 0.4553 | 0.1264 | 0.4771 | 0.2746 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Pseudomonadaceae;Pseudomonas; |
| 0.2276 | 0.2436 | 0.1581 | 0.1220 | 0.1436 | 0.1489 | 0.2432 | 0.1671 | 0.0531 | 0.3207 | 0.0647 | 0.3570 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Moraxellaceae;Acinetobacter; |
| 0.0617 | 0.3576 | 0.0128 | 0.0267 | 0.0121 | 0.0183 | 0.1409 | 0.0250 | 0.0128 | 0.0723 | 0.0066 | 0.0719 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Rosenbergiella; |
| 0.0806 | 0.0144 | 0.0951 | 0.0343 | 0.0410 | 0.0448 | 0.0661 | 0.0399 | 0.0838 | 0.1250 | 0.0755 | 0.0419 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Erwiniaceae;Pantoea |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Actinobacteria;Actinobacteria;Actinomycetales;Micrococcaceae;Arthrobacter; |
| 0.0369 | 0.0199 | 0.0373 | 0.0918 | 0.0236 | 0.0109 | 0.0425 | 0.0325 | 0.1743 | 0.2005 | 0.0360 | 0.0374 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Erwinia; |
| 0.0447 | 0.0756 | 0.0543 | 0.1232 | 0.0694 | 0.0446 | 0.0083 | 0.2012 | 0.0711 | 0.0182 | 0.1237 | 0.0746 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Epilithonimonas; |
| 0.0016 | 0 | 0 | 0.0029 | 0.0026 | 0 | 0.0001 | 0.0018 | 0 | 0.0016 | 0.0007 | 0.0009 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Erwiniaceae;Phaseolibacter; |
| 0.0152 | 0.0242 | 0.0126 | 0.0279 | 0.0644 | 0.0212 | 0.0250 | 0.0149 | 0.0092 | 0.0104 | 0.0273 | 0.0353 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Serratia; |
| 0.0010 | 0.0002 | 0 | 0.0174 | 0.0086 | 0.1032 | 0.0005 | 0.0010 | 0.0010 | 0.0088 | 0.0388 | 0.0003 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Arsenophonus; |
| 0.0131 | 0.0037 | 0.0049 | 0.0401 | 0.0132 | 0.0065 | 0.0041 | 0.0087 | 0.0074 | 0.0158 | 0.0119 | 0.0054 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodospirillales;Acetobacteraceae;Saccharibacter; |
| 0.0183 | 0.0067 | 0.0259 | 0.0540 | 0.0067 | 0.0102 | 0.0054 | 0.0238 | 0.0064 | 0.0075 | 0.0147 | 0.0150 | Bacteria;Proteobacteria;Alphaproteobacteria;Sphingomonadales;Sphingomonadaceae;Sphingomonas; |
| 0.0204 | 0.0680 | 0.0151 | 0.0267 | 0.0183 | 0.0216 | 0.0017 | 0.1237 | 0.0213 | 0.0080 | 0.0266 | 0.0249 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Chryseobacterium; |
| 0.0110 | 0.0088 | 0.0116 | 0.0139 | 0.0082 | 0.0220 | 0.0134 | 0.0050 | 0.0154 | 0.0147 | 0.0115 | 0.0174 | Bacteria;Proteobacteria;Betaproteobacteria;Burkholderiales;Oxalobacteraceae;Massilia; |
| 0.0031 | 0.0009 | 0.0012 | 0.0041 | 0.0060 | 0.0891 | 0.1298 | 0.0017 | 0.0173 | 0.0110 | 0.0021 | 0.0018 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodospirillales;Acetobacteraceae;Gluconacetobacter; |
| 0.0139 | 0.0098 | 0.0146 | 0.0023 | 0.0018 | 0.0698 | 0.0062 | 0.0189 | 0.0104 | 0.0118 | 0.0220 | 0.0279 | Bacteria;Proteobacteria;Betaproteobacteria;Burkholderiales;Oxalobacteraceae;Duganella; |
| 0.0044 | 0.0204 | 0.0086 | 0.0093 | 0.0044 | 0.0077 | 0.0017 | 0.0357 | 0.0087 | 0.0016 | 0.0091 | 0.0084 | Bacteria;Bacteroidetes;Sphingobacteriia;Sphingobacteriales;Sphingobacteriaceae;Pedobacter; |
| 0.0003 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Orbales;Orbaceae;Gilliamella; |
| 0 | 0.0002 | 0.0143 | 0 | 0.0001 | 0.0008 | 0 | 0.0003 | 0 | 0.0107 | 0 | 0.0009 | Bacteria;Cyanobacteria;Cyanobacteria;Synechococcales;Synechococcaceae;Cyanobium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Firmicutes;Clostridia;Clostridiales;Lachnospiraceae;Fusicatenibacter; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Oceanospirillales;Halomonadaceae;Carnimonas; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Escherichia/Shigella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Citrobacter; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Firmicutes;Bacilli;Lactobacillales;Streptococcaceae;Lactococcus; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.0245 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhizobiales;Bradyrhizobiaceae;Bradyrhizobium; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Bacteroidetes;Bacteroidia;Bacteroidales;Bacteroidaceae;Bacteroides; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rickettsiales;Rickettsiaceae;Rickettsia; |

| | | | | | | | | | | | | |
|---|---|--------|---|---|---|--------|---|---|---|---|---|---|
| 0 | 0 | 0.0168 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodobacterales;Rhodobacteraceae;Actibacterium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Betaproteobacteria;Neisseriales;Neisseriaceae;Snodgrassella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0.0129 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Pectobacterium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhodospirillales;Acetobacteraceae;Neokomagataea; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Orbales;Orbaceae;Frischella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Alphaproteobacteria;Rhizobiales;Bartonellaceae;Bartonella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Apidacter; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Firmicutes;Bacilli;Bacillales;Staphylococcaceae;Staphylococcus; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Proteobacteria;Gammaproteobacteria;Oceanospirillales;Halomonadaceae;Halotalea; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Bacteroidetes;Flavobacteriia;Flavobacteriales;Flavobacteriaceae;Flavobacterium; |

Table S3 Calculated relative abundance of all detected fungi in the samples

| F1H | F2H | F3H | C1H | C2H | C3H | RT1H | RT2H | RT3H | W1H | W2H | W3H | taxonomy |
|-----|-----|--------|-----|-----|-----|------|--------|--------|--------|--------|--------|--|
| 1 | 1 | 0.8186 | 1 | 0.5 | 1 | 1 | 0.4711 | 0.7432 | 0.1104 | 0.1276 | 0.5791 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Cladosporiaceae; Cladosporium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.3772 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Leotiomycetes; Erysiphales; Erysiphaceae; Podosphaera |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Mycosphaerella |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.1517 | 0 | 0.8525 | 0.6914 | 0.3141 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Saccharomycetaceae; Zygosaccharomyces |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Eurotiomycetes; Eurotiales; Aspergillaceae; Aspergillus |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Eurotiomycetes; Eurotiales; Aspergillaceae; Penicillium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Metschnikowiaceae; Metschnikowia |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Sordariomycetes; Hypocreales; Nectriaceae; Fusarium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Ramularia |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Taphrinomycetes; Taphrinales; Taphrinaceae; Taphrina |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Pleosporaceae; Alternaria |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Didymellaceae; Didymella |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Leotiomycetes; Thelebolales; Pseudoeurotiaceae; Bettisia |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Leotiomycetes; Helotiales; Sclerotiniaceae; Monilinia |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Dothideales; Aureobasidiaceae; Aureobasidium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Leotiomycetes; Helotiales; Helotiaceae; Tetracadium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Didymellaceae; Epicoccum |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Eurotiomycetes; Chaetothyriales; Trichomeriaceae; Trichomerium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Phaeosphaeriaceae; Pseudoophiobolus |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Leotiomycetes; Erysiphales; Erysiphaceae; Erysiphe |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Pleosporaceae; Pyrenophora |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Dothideomycetes; incertae sedis ; Monodictys |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Leotiomycetes; Erysiphales; Erysiphaceae; Blumeria |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Neodevriesiaceae; Neodevriesia |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Leptosphaeriaceae; Leptosphaeria |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Septoria |

| | | | | | | | | | | | | |
|---|---|--------|---|-----|---|---|---|--------|--------|--------|--------|--|
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Phaeosphaeriaceae; Neosetophoma |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Phaeothecaceae; Phaeotheca |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Debaryomycetaceae; Debaryomyces |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Melanommataceae; Alpinaria |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; incertae sedis; Periconia |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Sordariomycetes; Hypocreales; Bionectriaceae; Geosmithia |
| 0 | 0 | 0.1814 | 0 | 0.5 | 0 | 0 | 0 | 0.2568 | 0.0371 | 0.1809 | 0.1067 | Others |

| F1F | F2F | F3F | C1F | C2F | C3F | RT1F | RT2F | RT3F | W1F | W2F | W3F | taxonomy |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.5022 | 0.4722 | 0.3572 | 0.4118 | 0.3460 | 0.3924 | 0.4331 | 0.4817 | 0.3718 | 0.4458 | 0.4250 | 0.2067 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Cladosporiaceae; Cladosporium |
| 0.0131 | 0.0139 | 0.0096 | 0.0085 | 0.0162 | 0.0061 | 0.0096 | 0.0142 | 0.0168 | 0.0108 | 0.0022 | 0.0022 | Fungi; Ascomycota; Leotiomyces; Erysiphales; Erysiphaceae; Podosphaera |
| 0.0630 | 0.0859 | 0.0524 | 0.0693 | 0.0779 | 0.0540 | 0.0641 | 0.0635 | 0.0765 | 0.0520 | 0.0772 | 0.0332 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Mycosphaerella |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.0072 | 0.0019 | 0.0006 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Saccharomycetaceae; Zygosaccharomyces |
| 0.0198 | 0.0113 | 0.0111 | 0.0086 | 0.0097 | 0.0067 | 0.0063 | 0.0056 | 0.0092 | 0.2364 | 0.2488 | 0.2407 | Fungi; Ascomycota; Eurotiomycetes; Eurotiales; Aspergillaceae; Aspergillus |
| 0.0288 | 0.0657 | 0.0347 | 0.0379 | 0.0200 | 0.0407 | 0.0546 | 0.0176 | 0.0607 | 0.0539 | 0.0110 | 0.1054 | Fungi; Ascomycota; Eurotiomycetes; Eurotiales; Aspergillaceae; Penicillium |
| 0.0094 | 0.0237 | 0.0391 | 0.0182 | 0.0167 | 0.0071 | 0.0250 | 0.0077 | 0.0152 | 0.0068 | 0.0124 | 0.0200 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Metschnikowiaceae; Metschnikowia |
| 0.0376 | 0.0179 | 0.0511 | 0.0666 | 0.0592 | 0.0641 | 0.0658 | 0.1018 | 0.0863 | 0.0026 | 0.0301 | 0.0255 | Fungi; Ascomycota; Sordariomycetes; Hypocreales; Nectriaceae; Fusarium |
| 0.0199 | 0.0417 | 0.0349 | 0.0312 | 0.0365 | 0.0413 | 0.0251 | 0.0263 | 0.0360 | 0.0150 | 0.0099 | 0.0080 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Ramularia |
| 0.0466 | 0.0398 | 0.0435 | 0.0518 | 0.0526 | 0.0425 | 0.0379 | 0.0327 | 0.0403 | 0.0168 | 0.0151 | 0.0098 | Fungi; Ascomycota; Taphrinomycetes; Taphrinales; Taphrinaceae; Taphrina |
| 0.0129 | 0.0172 | 0.0658 | 0.0236 | 0.0259 | 0.0279 | 0.0179 | 0.0429 | 0.0155 | 0.0139 | 0.0148 | 0.0054 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Pleosporaceae; Alternaria |
| 0.0249 | 0.0231 | 0.0577 | 0.0264 | 0.0652 | 0.0171 | 0.0514 | 0.0334 | 0.0402 | 0.0293 | 0.0071 | 0.0032 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Didymellaceae; Didymella |
| 0.0002 | 0.0067 | 0.0001 | 0.0000 | 0.0003 | 0.0080 | 0.0135 | 0.0007 | 0.0052 | 0.0000 | 0.0069 | 0.2684 | Fungi; Ascomycota; Leotiomyces; Thelebolales; Pseudeurotiaceae; Bettisia |
| 0.0144 | 0.0123 | 0.0220 | 0.0627 | 0.0576 | 0.0306 | 0.0261 | 0.0167 | 0.0126 | 0.0170 | 0.0061 | 0.0080 | Fungi; Ascomycota; Leotiomyces; Helotiales; Sclerotiniaceae; Monilinia |
| 0.0561 | 0.0218 | 0.0159 | 0.0125 | 0.0231 | 0.0144 | 0.0154 | 0.0407 | 0.0142 | 0.0205 | 0.0356 | 0.0117 | Fungi; Ascomycota; Dothideomycetes; Dothideales; Aureobasidiaceae; Aureobasidium |
| 0.0255 | 0.0271 | 0.0298 | 0.0301 | 0.0266 | 0.0276 | 0.0344 | 0.0219 | 0.0354 | 0.0165 | 0.0291 | 0.0169 | Fungi; Ascomycota; Leotiomyces; Helotiales; Helotiaceae; Tetracladium |
| 0.0154 | 0.0127 | 0.0076 | 0.0117 | 0.0325 | 0.0049 | 0.0272 | 0.0021 | 0.0112 | 0.0013 | 0.0048 | 0.0031 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Didymellaceae; Epicoccum |
| 0.0218 | 0.0039 | 0.0169 | 0.0153 | 0.0142 | 0.0706 | 0.0167 | 0.0113 | 0.0178 | 0.0040 | 0.0037 | 0.0029 | Fungi; Ascomycota; Eurotiomycetes; Chaetothyriales; Trichomeriaceae; Trichomerium |
| 0.0014 | 0.0004 | 0.0016 | 0.0015 | 0.0017 | 0.0011 | 0.0019 | 0.0010 | 0.0016 | 0.0004 | 0.0024 | 0.0008 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Phaeosphaeriaceae; Pseudoophiobolus |
| 0.0090 | 0.0075 | 0.0067 | 0.0084 | 0.0083 | 0.0128 | 0.0053 | 0.0072 | 0.0135 | 0.0007 | 0.0021 | 0.0007 | Fungi; Ascomycota; Leotiomyces; Erysiphales; Erysiphaceae; Erysiphe |
| 0 | 0.0001 | 0 | 0 | 0 | 0 | 0 | 0.0022 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Pleosporaceae; Pyrenophora |
| 0.0007 | 0.0010 | 0.0017 | 0.0013 | 0.0008 | 0.0019 | 0.0002 | 0.0013 | 0.0003 | 0.0003 | 0.0026 | 0.0000 | Fungi; Ascomycota; Dothideomycetes; Dothideomycetes; incertae sedis ; Monodictys |
| 0.0005 | 0.0022 | 0.0016 | 0.0043 | 0.0094 | 0.0090 | 0.0065 | 0.0039 | 0.0168 | 0.0017 | 0.0018 | 0.0010 | Fungi; Ascomycota; Leotiomyces; Erysiphales; Erysiphaceae; Blumeria |
| 0.0045 | 0.0090 | 0.0219 | 0.0062 | 0.0091 | 0.0094 | 0.0064 | 0.0024 | 0.0136 | 0.0017 | 0.0117 | 0.0011 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Neodevriesiaceae; Neodevriesia |
| 0.0055 | 0.0026 | 0.0077 | 0.0019 | 0.0077 | 0.0036 | 0.0046 | 0.0043 | 0.0018 | 0.0067 | 0.0020 | 0.0056 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Leptosphaeriaceae; Leptosphaeria |
| 0.0035 | 0.0019 | 0.0104 | 0.0027 | 0.0012 | 0.0035 | 0.0013 | 0.0012 | 0.0014 | 0.0020 | 0.0011 | 0.0006 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Septoria |
| 0.0027 | 0.0023 | 0.0044 | 0.0021 | 0.0055 | 0.0026 | 0.0029 | 0.0026 | 0.0068 | 0.0108 | 0.0084 | 0.0012 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Phaeosphaeriaceae; Neosetophoma |
| 0.0020 | 0.0136 | 0.0042 | 0.0091 | 0.0039 | 0.0061 | 0.0021 | 0.0015 | 0.0032 | 0.0052 | 0.0063 | 0.0024 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Phaeothecaceae; Phaeotheca |

| | | | | | | | | | | | | |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.0010 | 0.0000 | 0.0000 | 0.0001 | 0.0002 | 0.0002 | 0.0001 | 0.0003 | 0.0000 | 0.0000 | 0.0003 | 0.0001 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Debaryomycetaceae; Debaryomyces |
| 0.0060 | 0.0026 | 0.0102 | 0.0005 | 0.0031 | 0.0263 | 0.0002 | 0.0034 | 0.0051 | 0.0004 | 0.0004 | 0.0005 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Melanommataceae; Alpinaria |
| 0.0035 | 0.0032 | 0.0010 | 0.0039 | 0.0022 | 0.0107 | 0.0029 | 0.0043 | 0.0052 | 0.0014 | 0.0006 | 0.0006 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; incertae sedis; Periconia |
| 0 | 0 | 0 | 0.0001 | 0 | 0 | 0 | 0.0001 | 0.0001 | 0 | 0 | 0 | Fungi; Ascomycota; Sordariomycetes; Hypocreales; Bionectriaceae; Geosmithia |
| 0.0480 | 0.0567 | 0.0791 | 0.0715 | 0.0669 | 0.0571 | 0.0416 | 0.0434 | 0.0657 | 0.0185 | 0.0184 | 0.0137 | Others |

| F1N | F2N | F3N | C1N | C2N | C3N | RT1N | RT2N | RT3N | W1N | W2N | W3N | taxonomy |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.2852 | 0.3448 | 0.1793 | 0.1398 | 0.2123 | 0.2065 | 0.2008 | 0.1668 | 0.1768 | 0.0903 | 0.1314 | 0.0748 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Cladosporiaceae; Cladosporium |
| 0.0683 | 0.1524 | 0.6562 | 0.3044 | 0.2928 | 0.1976 | 0.0351 | 0.1154 | 0.1335 | 0.1401 | 0.1702 | 0.1557 | Fungi; Ascomycota; Leotiomyces; Erysiphales; Erysiphaceae; Podosphaera |
| 0.1677 | 0.0924 | 0.0531 | 0.1946 | 0.1863 | 0.0950 | 0.2535 | 0.1904 | 0.1741 | 0.0744 | 0.0930 | 0.0879 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Mycosphaerella |
| 0.0002 | 0.0001 | 0.0001 | 0.0009 | 0.0005 | 0.0007 | 0.0095 | 0.0478 | 0.0193 | 0.5912 | 0.4577 | 0.4776 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Saccharomycetaceae; Zygosaccharomyces |
| 0.0021 | 0.0022 | 0.0005 | 0.0018 | 0.0020 | 0.0055 | 0.0007 | 0.0113 | 0.0042 | 0.0161 | 0.0268 | 0.0975 | Fungi; Ascomycota; Eurotiomycetes; Eurotiales; Aspergillaceae; Aspergillus |
| 0.0614 | 0.0108 | 0.0046 | 0.0118 | 0.0324 | 0.0257 | 0.0105 | 0.0113 | 0.0378 | 0.0010 | 0.0015 | 0.0018 | Fungi; Ascomycota; Eurotiomycetes; Eurotiales; Aspergillaceae; Penicillium |
| 0.0884 | 0.0768 | 0.0070 | 0.0133 | 0.0192 | 0.0407 | 0.0187 | 0.2069 | 0.0178 | 0.0218 | 0.0052 | 0.0104 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Metschnikowiaceae; Metschnikowia |
| 0.0011 | 0.0124 | 0.0005 | 0.0024 | 0.0031 | 0.0086 | 0.0061 | 0.0019 | 0.0027 | 0.0041 | 0.0033 | 0.0013 | Fungi; Ascomycota; Sordariomycetes; Hypocreales; Nectriaceae; Fusarium |
| 0.0455 | 0.0188 | 0.0159 | 0.0208 | 0.0277 | 0.0289 | 0.0404 | 0.0212 | 0.0342 | 0.0067 | 0.0077 | 0.0069 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Ramularia |
| 0.0296 | 0.0158 | 0.0084 | 0.0084 | 0.0077 | 0.0176 | 0.0159 | 0.0075 | 0.0116 | 0.0019 | 0.0213 | 0.0041 | Fungi; Ascomycota; Taphrinomycetes; Taphrinales; Taphrinaceae; Taphrina |
| 0.0565 | 0.0482 | 0.0046 | 0.0291 | 0.0322 | 0.0336 | 0.0290 | 0.0168 | 0.0274 | 0.0041 | 0.0081 | 0.0097 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Pleosporaceae; Alternaria |
| 0.0187 | 0.0197 | 0.0118 | 0.0045 | 0.0089 | 0.0298 | 0.0185 | 0.0056 | 0.0139 | 0.0043 | 0.0183 | 0.0063 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Didymellaceae; Didymella |
| 0.0004 | 0.0004 | 0.0002 | 0 | 0 | 0.0035 | 0.0050 | 0 | 0.1537 | 0.0001 | 0.0001 | 0.0000 | Fungi; Ascomycota; Leotiomyces; Thelebolales; Pseudeurotiaceae; Bettsia |
| 0.0209 | 0.0279 | 0.0095 | 0.0364 | 0.0320 | 0.0228 | 0.0141 | 0.0176 | 0.0382 | 0.0058 | 0.0047 | 0.0028 | Fungi; Ascomycota; Leotiomyces; Helotiales; Sclerotiniaceae; Monilinia |
| 0.0126 | 0.0122 | 0.0033 | 0.0054 | 0.0074 | 0.0224 | 0.0104 | 0.0047 | 0.0071 | 0.0021 | 0.0035 | 0.0022 | Fungi; Ascomycota; Dothideomycetes; Dothideales; Aureobasidiaceae; Aureobasidium |
| 0.0050 | 0.0019 | 0.0007 | 0.0032 | 0.0013 | 0.0033 | 0.0033 | 0.0025 | 0.0056 | 0.0019 | 0.0020 | 0.0025 | Fungi; Ascomycota; Leotiomyces; Helotiales; Helotiaceae; Tetracladium |
| 0.0083 | 0.0039 | 0.0026 | 0.0178 | 0.0050 | 0.0053 | 0.0312 | 0.0040 | 0.0385 | 0.0028 | 0.0044 | 0.0034 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Didymellaceae; Epicoccum |
| 0.0084 | 0.0054 | 0.0011 | 0.0009 | 0.0014 | 0.0145 | 0.0012 | 0.0008 | 0.0026 | 0.0004 | 0.0018 | 0.0007 | Fungi; Ascomycota; Eurotiomycetes; Chaetothyriales; Trichomeriaceae; Trichomerium |
| 0.0027 | 0.0075 | 0.0006 | 0.0079 | 0.0036 | 0.0050 | 0.1845 | 0.0032 | 0.0061 | 0.0017 | 0.0010 | 0.0016 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Phaeosphaeriaceae; Pseudoophiobolus |
| 0.0221 | 0.0098 | 0.0049 | 0.0080 | 0.0138 | 0.0122 | 0.0187 | 0.0060 | 0.0075 | 0.0011 | 0.0024 | 0.0028 | Fungi; Ascomycota; Leotiomyces; Erysiphales; Erysiphaceae; Erysiphe |
| 0 | 0.0084 | 0 | 0.1139 | 0.0192 | 0 | 0.0019 | 0.0220 | 0.0006 | 0 | 0.0007 | 0.0011 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Pleosporaceae; Pyrenophora |
| 0.0171 | 0.0041 | 0.0013 | 0.0088 | 0.0106 | 0.0830 | 0.0088 | 0.0092 | 0.0060 | 0.0010 | 0.0027 | 0.0046 | Fungi; Ascomycota; Dothideomycetes; Dothideomycetes; incertae sedis ; Monodictys |
| 0.0045 | 0.0282 | 0.0004 | 0.0054 | 0.0062 | 0.0036 | 0.0055 | 0.0061 | 0.0029 | 0.0017 | 0.0025 | 0.0236 | Fungi; Ascomycota; Leotiomyces; Erysiphales; Erysiphaceae; Blumeria |
| 0.0013 | 0.0022 | 0.0006 | 0.0003 | 0.0000 | 0.0021 | 0.0004 | 0.0004 | 0.0005 | 0.0006 | 0.0002 | 0.0002 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Neodevriesiaceae; Neodevriesia |
| 0.0075 | 0.0039 | 0.0008 | 0.0008 | 0.0017 | 0.0089 | 0.0016 | 0.0012 | 0.0021 | 0.0007 | 0.0007 | 0.0006 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Leptosphaeriaceae; Leptosphaeria |
| 0.0050 | 0.0050 | 0.0017 | 0.0024 | 0.0035 | 0.0093 | 0.0132 | 0.0033 | 0.0030 | 0.0017 | 0.0026 | 0.0009 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Septoria |
| 0.0015 | 0.0043 | 0.0008 | 0.0020 | 0.0005 | 0.0050 | 0.0026 | 0.0004 | 0.0061 | 0.0015 | 0.0019 | 0.0001 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Phaeosphaeriaceae; Neosetophoma |
| 0.0027 | 0.0030 | 0.0009 | 0.0026 | 0.0002 | 0.0037 | 0.0011 | 0.0004 | 0.0009 | 0.0005 | 0.0003 | 0.0003 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Phaeothecaceae; Phaeotheca |

| | | | | | | | | | | | | |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.0070 | 0.0082 | 0.0019 | 0.0113 | 0.0113 | 0.0074 | 0.0076 | 0.0097 | 0.0044 | 0.0005 | 0.0005 | 0.0023 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Debaryomycetaceae; Debaryomyces |
| 0 | 0 | 0.0000 | 0.0007 | 0 | 0 | 0 | 0.0003 | 0.0000 | 0.0001 | 0.0002 | 0 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; Melanommataceae; Alpinaria |
| 0.0008 | 0.0025 | 0.0001 | 0.0004 | 0.0016 | 0.0012 | 0.0006 | 0.0001 | 0.0043 | 0.0037 | 0.0006 | 0.0004 | Fungi; Ascomycota; Dothideomycetes; Pleosporales; incertae sedis; Periconia |
| 0.0002 | 0.0015 | 0.0001 | 0.0000 | 0.0011 | 0.0271 | 0.0001 | 0.0362 | 0.0027 | 0 | 0.0001 | 0 | Fungi; Ascomycota; Sordariomycetes; Hypocreales; Bionectriaceae; Geosmithia |
| 0.0475 | 0.0655 | 0.0264 | 0.0400 | 0.0543 | 0.0697 | 0.0496 | 0.0688 | 0.0536 | 0.0162 | 0.0227 | 0.0157 | Others |

Table S4 Calculated relative abundance of bacterial composition (> 10%) between different storage conditions

| F1H | F2H | F3H | C1H | C2H | C3H | RT1H | RT2H | W1H | W2H | W3H | taxonomy |
|--------|--------|--------|--------|--------|--------|--------|--------|---------|---------|---------|--|
| 0.4335 | 0.6673 | 0.8104 | 0.5924 | 0.7750 | 0.6638 | 0.8669 | 0.9062 | 0.52483 | 0.07480 | 0.54757 | Bacteria;Firmicutes;Bacilli;Lactobacillales;Lactobacillaceae;Lactobacillus; |
| 0.1430 | 0.0173 | 0.0255 | 0.0070 | 0.0255 | 0.0864 | 0.0149 | 0.0067 | 0.00106 | 0 | 0.01743 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Pseudomonadaceae;Pseudomonas; |
| 0.0256 | 0.0505 | 0.0043 | 0.2206 | 0.0079 | 0.0834 | 0.0022 | 0.0033 | 0.02460 | 0.02002 | 0.20771 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Moraxellaceae;Acinetobacter; |
| 0.1204 | 0.0038 | 0.0084 | 0.0041 | 0.0025 | 0.0064 | 0.0976 | 0.0012 | 0 | 0 | 0.03213 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Rosenbergiella; |
| 0.0124 | 0.0117 | 0.0053 | 0.0014 | 0.0032 | 0.0045 | 0.0001 | 0.0006 | 0.32299 | 0.75366 | 0.06831 | Bacteria;Actinobacteria;Actinobacteria;Actinomycetales;Micrococcaceae;Arthrobacter; |
| 0.2651 | 0.2494 | 0.1462 | 0.1746 | 0.1858 | 0.1554 | 0.0182 | 0.0821 | 0.12653 | 0.15152 | 0.12685 | Others |

| F1F | F2F | F3F | C1F | C2F | C3F | RT1F | RT2F | RT3F | W1F | W2F | W3F | taxonomy |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.5824 | 0.8154 | 0.7050 | 0.7270 | 0.4291 | 0.6423 | 0.5668 | 0.5939 | 0.6394 | 0.9061 | 0.5689 | 0.8142 | Bacteria;Firmicutes;Bacilli;Lactobacillales;Lactobacillaceae;Lactobacillus; |
| 0.1086 | 0.0126 | 0.0356 | 0.0557 | 0.1557 | 0.1447 | 0.1173 | 0.1578 | 0.1004 | 0.0211 | 0.0641 | 0.0277 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Pseudomonadaceae;Pseudomonas; |
| 0.0233 | 0.0041 | 0.0137 | 0.0192 | 0.0062 | 0.0046 | 0.0398 | 0.0150 | 0.0285 | 0.0005 | 0.1711 | 0.0076 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Moraxellaceae;Acinetobacter; |
| 0.1586 | 0.0390 | 0.1071 | 0.0515 | 0.0477 | 0.0603 | 0.1209 | 0.0730 | 0.0275 | 0.0275 | 0.0572 | 0.0423 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Rosenbergiella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Actinobacteria;Actinobacteria;Actinomycetales;Micrococcaceae;Arthrobacter; |
| 0.1271 | 0.1288 | 0.1387 | 0.1466 | 0.3613 | 0.1481 | 0.1551 | 0.1604 | 0.2042 | 0.0448 | 0.1387 | 0.1081 | Others |

| F1N | F2N | F3N | C1N | C2N | C3N | RT1N | RT2N | RT3N | W1N | W2N | W3N | taxonomy |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.0479 | 0.0035 | 0.0143 | 0.0459 | 0.0570 | 0.0135 | 0.0015 | 0.0062 | 0.0524 | 0.0348 | 0.0273 | 0.0045 | Bacteria;Firmicutes;Bacilli;Lactobacillales;Lactobacillaceae;Lactobacillus; |
| 0.3984 | 0.1424 | 0.5023 | 0.3574 | 0.5189 | 0.3670 | 0.2967 | 0.2924 | 0.4553 | 0.1264 | 0.4771 | 0.2746 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Pseudomonadaceae;Pseudomonas; |
| 0.2276 | 0.2436 | 0.1581 | 0.1220 | 0.1436 | 0.1489 | 0.2432 | 0.1671 | 0.0531 | 0.3207 | 0.0647 | 0.3570 | Bacteria;Proteobacteria;Gammaproteobacteria;Pseudomonadales;Moraxellaceae;Acinetobacter; |
| 0.0617 | 0.3576 | 0.0128 | 0.0267 | 0.0121 | 0.0183 | 0.1409 | 0.0250 | 0.0128 | 0.0723 | 0.0066 | 0.0719 | Bacteria;Proteobacteria;Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Rosenbergiella; |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Bacteria;Actinobacteria;Actinobacteria;Actinomycetales;Micrococcaceae;Arthrobacter; |
| 0.2645 | 0.2529 | 0.3124 | 0.4480 | 0.2684 | 0.4523 | 0.3177 | 0.5093 | 0.4263 | 0.4458 | 0.4243 | 0.2920 | Others |

Table S5 Calculated relative abundance of fungal composition (> 10%) between different storage conditions

| F1H | F2H | F3H | C1H | C2H | C3H | RT1H | RT2H | RT3H | W1H | W2H | W3H | taxonomy |
|-----|-----|--------|-----|-----|-----|------|--------|--------|--------|--------|--------|--|
| 1 | 1 | 0.8186 | 1 | 0.5 | 1 | 1 | 0.4711 | 0.7432 | 0.1104 | 0.1276 | 0.5791 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Cladosporiaceae; Cladosporium |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.3772 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Leotiomyces; Erysiphales; Erysiphaceae; Podosphaera |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Mycosphaerella |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.1517 | 0 | 0.8525 | 0.6914 | 0.3141 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Saccharomycetaceae; Zygosaccharomyces |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | Fungi; Ascomycota; Eurotiomycetes; Eurotiales; Aspergillaceae; Aspergillus |
| 0 | 0 | 0.1814 | 0 | 0.5 | 0 | 0 | 0 | 0.2568 | 0.0371 | 0.1809 | 0.1067 | Others |

| F1F | F2F | F3F | C1F | C2F | C3F | RT1F | RT2F | RT3F | W1F | W2F | W3F | taxonomy |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.5022 | 0.4722 | 0.3572 | 0.4118 | 0.3460 | 0.3924 | 0.4331 | 0.4817 | 0.3718 | 0.4458 | 0.4250 | 0.2067 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Cladosporiaceae; Cladosporium |
| 0.0131 | 0.0139 | 0.0096 | 0.0085 | 0.0162 | 0.0061 | 0.0096 | 0.0142 | 0.0168 | 0.0108 | 0.0022 | 0.0022 | Fungi; Ascomycota; Leotiomyces; Erysiphales; Erysiphaceae; Podosphaera |
| 0.0630 | 0.0859 | 0.0524 | 0.0693 | 0.0779 | 0.0540 | 0.0641 | 0.0635 | 0.0765 | 0.0520 | 0.0772 | 0.0332 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Mycosphaerella |
| 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0.0072 | 0.0019 | 0.0006 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Saccharomycetaceae; Zygosaccharomyces |
| 0.0198 | 0.0113 | 0.0111 | 0.0086 | 0.0097 | 0.0067 | 0.0063 | 0.0056 | 0.0092 | 0.2364 | 0.2488 | 0.2407 | Fungi; Ascomycota; Eurotiomycetes; Eurotiales; Aspergillaceae; Aspergillus |
| 0.4019 | 0.4166 | 0.5697 | 0.5018 | 0.5502 | 0.5409 | 0.4869 | 0.4351 | 0.5256 | 0.2477 | 0.2449 | 0.5166 | Others |

| F1N | F2N | F3N | C1N | C2N | C3N | RT1N | RT2N | RT3N | W1N | W2N | W3N | taxonomy |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--|
| 0.2852 | 0.3448 | 0.1793 | 0.1398 | 0.2123 | 0.2065 | 0.2008 | 0.1668 | 0.1768 | 0.0903 | 0.1314 | 0.0748 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Cladosporiaceae; Cladosporium |
| 0.0683 | 0.1524 | 0.6562 | 0.3044 | 0.2928 | 0.1976 | 0.0351 | 0.1154 | 0.1335 | 0.1401 | 0.1702 | 0.1557 | Fungi; Ascomycota; Leotiomyces; Erysiphales; Erysiphaceae; Podosphaera |
| 0.1677 | 0.0924 | 0.0531 | 0.1946 | 0.1863 | 0.0950 | 0.2535 | 0.1904 | 0.1741 | 0.0744 | 0.0930 | 0.0879 | Fungi; Ascomycota; Dothideomycetes; Capnodiales; Mycosphaerellaceae; Mycosphaerella |
| 0.0002 | 0.0001 | 0.0001 | 0.0009 | 0.0005 | 0.0007 | 0.0095 | 0.0478 | 0.0193 | 0.5912 | 0.4577 | 0.4776 | Fungi; Ascomycota; Saccharomycetes; Saccharomycetales; Saccharomycetaceae; Zygosaccharomyces |
| 0.0021 | 0.0022 | 0.0005 | 0.0018 | 0.0020 | 0.0055 | 0.0007 | 0.0113 | 0.0042 | 0.0161 | 0.0268 | 0.0975 | Fungi; Ascomycota; Eurotiomycetes; Eurotiales; Aspergillaceae; Aspergillus |
| 0.4766 | 0.4080 | 0.1109 | 0.3584 | 0.3061 | 0.4949 | 0.5004 | 0.4683 | 0.4921 | 0.0878 | 0.1209 | 0.1064 | Others |

Table S6 Results of statistical analysis 1 (GLM) for bacterial genera in Dataset 2 (additionally Dataset 1) (> 10%)

| GLM (Dataset 2-2019 (F and N)) | df1 | df2 | F.ratio | p.value |
|---------------------------------------|------------|------------|----------------|----------------|
| Lactobacillus | | | | |
| Location | 1 | Inf | 524.107 | <0.0001 |
| Storage condition | 3 | Inf | 2.792 | 0.0388 |
| Location:Storage condition | 3 | Inf | 3.014 | 0.0187 |
| | | | | |
| Pseudomonas | | | | |
| Location | 1 | Inf | 61.793 | <0.0001 |
| Storage condition | 3 | Inf | 2.303 | 0.0748 |
| Location:Storage condition | 3 | Inf | 0.338 | 0.798 |
| | | | | |
| Acinetobacter | | | | |
| Location | 1 | Inf | 43.311 | <0.0001 |
| Storage condition | 3 | Inf | 0.602 | 0.6138 |
| Location:Storage condition | 3 | Inf | 0.916 | 0.4319 |
| | | | | |
| Rosenbergiella | | | | |
| Location | 1 | Inf | 1.249 | 0.2637 |
| Storage condition | 3 | Inf | 0.954 | 0.4132 |
| Location:Storage condition | 3 | Inf | 0.132 | 0.941 |

| GLM (Dataset 1&2- 2018&2019 (H, F and N)) | df1 | df2 | F.ratio | p.value |
|--|------------|------------|----------------|----------------|
| Lactobacillus | | | | |
| Location | 2 | Inf | 260.289 | <0.0001 |
| Storage condition | 3 | Inf | 2.378 | 0.0677 |
| Location:Storage condition | 6 | Inf | 7.032 | <0.0001 |
| | | | | |
| Pseudomonas | | | | |
| Location | 2 | Inf | 62.422 | <0.0001 |
| Storage condition | 3 | Inf | 2.85 | 0.0359 |
| Location:Storage condition | 6 | Inf | 1.296 | 0.2551 |
| | | | | |
| Acinetobacter | | | | |
| Location | 2 | Inf | 18.475 | <0.0001 |
| Storage condition | 3 | Inf | 0.667 | 0.5724 |
| Location:Storage condition | 6 | Inf | 1.096 | 0.3616 |
| | | | | |
| Rosenbergiella | | | | |
| Location | 2 | Inf | 7.616 | 0.0005 |
| Storage condition | 3 | Inf | 1.266 | 0.2842 |
| Location:Storage condition | 6 | Inf | 0.214 | 0.9724 |
| | | | | |
| Arthrobacter | | | | |
| Location | 2 | Inf | 9.421 | 0.0001 |
| Storage condition | 3 | Inf | 5.425 | 0.001 |
| Location:Storage condition | 6 | Inf | 2.821 | 0.0096 |

Table S7 Results of statistical analysis 2 (Tukey Test) for bacterial genera in Dataset 2 (additionally Dataset 1) (> 10%)

| Tukey Test (Dataset 2-2019 (F and N)) | estimate | SE | df | z.ratio | p.value |
|--|----------|--------|-----|---------|---------|
| Lactobacillus | | | | | |
| cold-fresh | -0.0404 | 0.0396 | Inf | -1.02 | 0.7378 |
| cold-room temp | 0.0157 | 0.0407 | Inf | 0.386 | 0.9805 |
| cold-warm | -0.0815 | 0.038 | Inf | -0.2145 | 0.1391 |
| fresh-room temp | 0.0561 | 0.0387 | Inf | 1.45 | 0.4683 |
| fresh-warm | -0.0412 | 0.0358 | Inf | -1.15 | 0.6587 |
| room temp-warm | -0.0972 | 0.0371 | Inf | -2.623 | 0.0432 |

| Tukey Test (Dataset 1&2-2018&2019 (H, F and N)) | estimate | SE | df | z.ratio | p.value |
|--|----------|---------|-----|---------|---------|
| Lactobacillus | | | | | |
| F-H | 0.0381 | 0.0463 | Inf | 0.822 | 0.6892 |
| F-N | 0.6235 | 0.0356 | Inf | 17.516 | <0.0001 |
| H-N | 0.5854 | 0.0348 | Inf | 16.828 | <0.0001 |
| Pseudomonas | | | | | |
| cold-fresh | 0.0464 | 0.0309 | Inf | 1.505 | 0.4347 |
| cold-room temp | 0.0338 | 0.0321 | Inf | 0.71 | 0.893 |
| cold-warm | 0.0806 | 0.0294 | Inf | 2.745 | 0.0308 |
| fresh-room temp | -0.0236 | 0.0305 | Inf | -0.773 | 0.8667 |
| fresh-warm | 0.0342 | 0.0276 | Inf | 1.237 | 0.603 |
| room temp-warm | 0.0578 | 0.029 | Inf | 1.99 | 0.1917 |
| F-H | 0.0575 | 0.0172 | Inf | 3.339 | 0.0024 |
| F-N | -0.2525 | 0.0307 | Inf | -8.223 | <0.0001 |
| H-N | -0.3099 | 0.0281 | Inf | -11.047 | <0.0001 |
| Acinetobacter | | | | | |
| F-H | -0.0178 | 0.0143 | Inf | -1.246 | 0.436 |
| F-N | -0.01549 | 0.0255 | Inf | -6.078 | <0.0001 |
| H-N | -0.01371 | 0.0268 | Inf | -5.12 | <0.0001 |
| Rosenbergiella | | | | | |
| F-H | 0.0568 | 0.0165 | Inf | 3.445 | 0.0017 |
| F-N | 0.0212 | 0.0194 | Inf | 1.094 | 0.5175 |
| H-N | -0.0356 | 0.0142 | Inf | -2.513 | 0.0321 |
| Arthrobacter | | | | | |
| cold-fresh | -0.00407 | 0.00993 | Inf | -0.41 | 0.9767 |
| cold-room temp | 0.00351 | 0.00812 | Inf | 0.432 | 0.9729 |
| cold-warm | -0.10362 | 0.0269 | Inf | -3.853 | 0.0007 |
| fresh-room temp | 0.00758 | 0.00949 | Inf | 0.799 | 0.8548 |
| fresh-warm | -0.09955 | 0.02733 | Inf | -3.642 | 0.0015 |
| room temp-warm | -0.10713 | 0.02671 | Inf | -4.012 | 0.0004 |
| F-H | -0.0908 | 0.0211 | Inf | -4.302 | 0.0001 |
| F-N | 0 | 0.0056 | Inf | 0 | 1 |
| H-N | 0.0908 | 0.0211 | Inf | 4.302 | 0.0001 |

Table S8 Results of statistical analysis 1 (GLM) for fungal genera in Dataset 2 (> 10%)

| GLM (Dataset 2-2019 (F and N)) | df1 | df2 | Fratio | p.value |
|---------------------------------------|-----|-----|---------|---------|
| Cladosporium | | | | |
| Location | 1 | lnf | 95.146 | <0.0001 |
| Storage condition | 3 | lnf | 6.009 | 0.0004 |
| Location:Storage condition | 3 | lnf | 0.642 | 0.5879 |
| | | | | |
| Podoshaera | | | | |
| Location | 1 | lnf | 40.318 | <0.0001 |
| Storage condition | 3 | lnf | 2.218 | 0.0837 |
| Location:Storage condition | 3 | lnf | 2.389 | 0.0668 |
| | | | | |
| Mycosphaerella | | | | |
| Location | 1 | lnf | 45.613 | <0.0001 |
| Storage condition | 3 | lnf | 8.078 | <0.0001 |
| Location:Storage condition | 3 | lnf | 5.337 | 0.0011 |
| | | | | |
| Zygosaccharomyces | | | | |
| Location | 1 | lnf | 643.042 | <0.0001 |
| Storage condition | 3 | lnf | 221.964 | <0.0001 |
| Location:Storage condition | 3 | lnf | 217.176 | <0.0001 |
| | | | | |
| Aspergillus | | | | |
| Location | 1 | lnf | 139.089 | <0.0001 |
| Storage condition | 3 | lnf | 72.087 | <0.0001 |
| Location:Storage condition | 3 | lnf | 41.366 | <0.0001 |

Table S9 Results of statistical analysis 2 (Tukey Test) for fungal genera in Dataset 2 (> 10%)

| Tukey Test (Dataset 2-2019 (F and N)) | estimate | SE | df | zratio | p.value |
|--|------------|----------|-----|---------|---------|
| Cladosporium | | | | | |
| cold-fresh | -0.0688 | 0.0325 | Inf | -2.114 | 0.1486 |
| cold-room temp | -0.021 | 0.0316 | Inf | -0.664 | 0.9106 |
| cold-warm | 0.0595 | 0.0299 | Inf | 1.988 | 0.1926 |
| fresh-room temp | 0.0478 | 0.0327 | Inf | 1.463 | 0.46 |
| fresh-warm | 0.1283 | 0.0311 | Inf | 4.126 | 0.0002 |
| room temp-warm | 0.0808 | 0.0301 | Inf | 2.675 | 0.0375 |
| | | | | | |
| Mycosphaerella | | | | | |
| cold-fresh | 0.0288 | 0.0146 | Inf | 1.968 | 0.2 |
| cold-room temp | -0.0263 | 0.0163 | Inf | -1.621 | 0.3667 |
| cold-warm | 0.041 | 0.0142 | Inf | 2.893 | 0.0199 |
| fresh-room temp | -0.0552 | 0.0154 | Inf | -3.589 | 0.0019 |
| fresh-warm | 0.0122 | 0.0132 | Inf | 0.927 | 0.7903 |
| room temp-warm | 0.0674 | 0.0149 | Inf | 4.512 | <0.0001 |
| | | | | | |
| Zygosaccharomyces | | | | | |
| cold-fresh | 0.000471 | 0.000884 | Inf | 0.533 | 0.9511 |
| cold-room temp | -0.010296 | 0.002941 | Inf | -3.499 | 0.0026 |
| cold-warm | -0.254761 | 0.009996 | Inf | -25.485 | <0.0001 |
| fresh-room temp | -0.010767 | 0.002908 | Inf | -3.702 | 0.0012 |
| fresh-warm | -0.255232 | 0.009985 | Inf | -25.562 | <0.0001 |
| room temp-warm | -0.244466 | 0.010374 | Inf | -23.565 | <0.0001 |
| | | | | | |
| Aspergillus | | | | | |
| cold-fresh | -0.001901 | 0.0031 | Inf | -0.613 | 0.9279 |
| cold-room temp | 0.000427 | 0.00281 | Inf | 0.152 | 0.9987 |
| cold-warm | -0.132687 | 0.00916 | Inf | -14.488 | <0.0001 |
| fresh-room temp | 0.002328 | 0.00305 | Inf | 0.764 | 0.8708 |
| fresh-warm | -0.0130785 | 0.00925 | Inf | -14.161 | <0.0001 |
| room temp-warm | -0.0133114 | 0.00914 | Inf | -14.562 | <0.0001 |

Figure S1 Location area of sampling in Southern Germany, Baden-Wuerttemberg with different colors showing the different altitude regions. (Mapping with JMP® pro 15.0; basic data from the Environmental Information System (UIS) of the LUBW State Institute for the Environment Baden-Wuerttemberg and basic data © State Office for Geoinformation and Rural Development Baden-Wuerttemberg (www.lgl-bw.de))

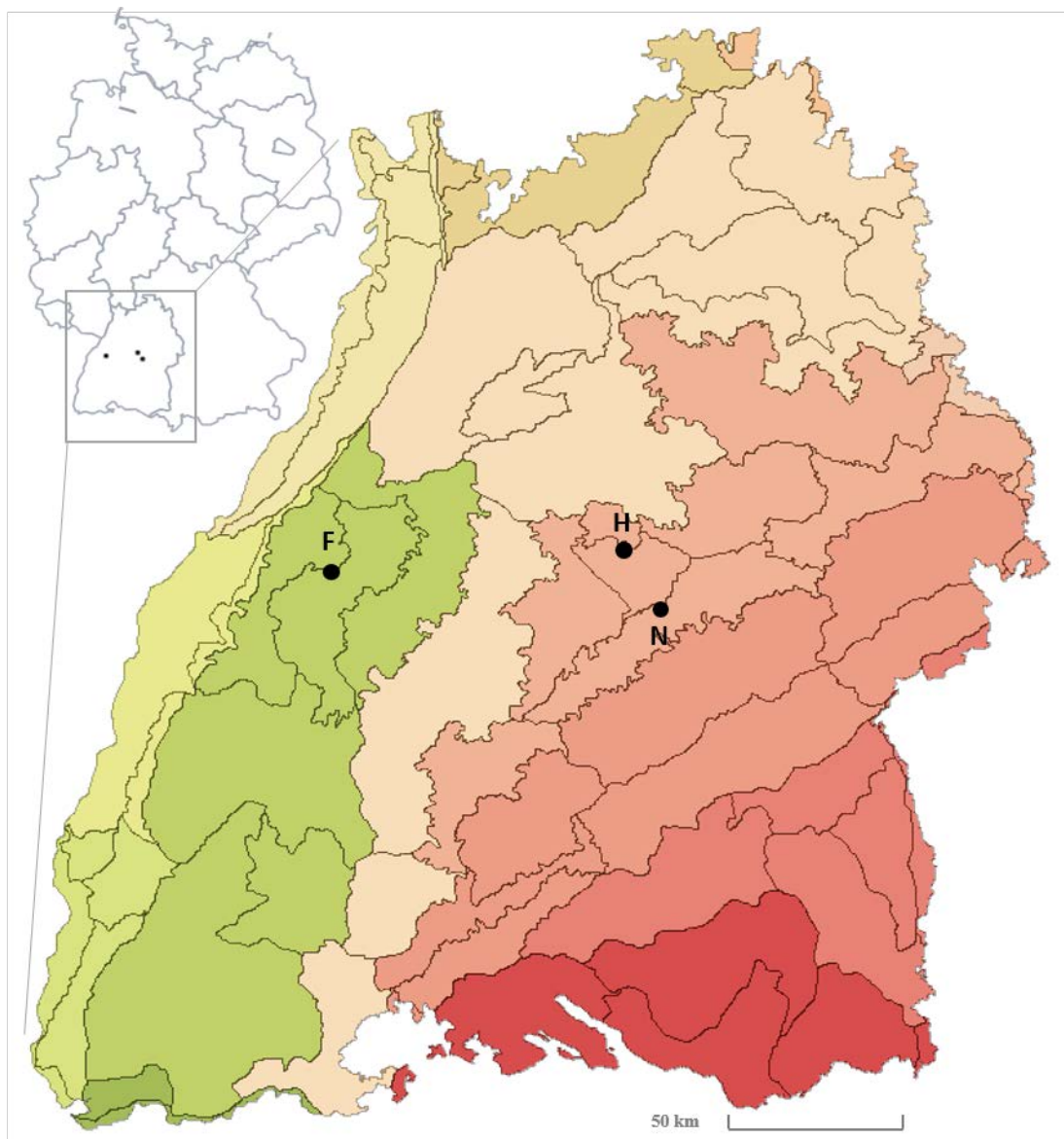


Figure S2 Storage cabinet with moisture meter and saturated sodium chloride solution (“warm” storage condition)



Figure S3 Stack bar chart, showing the composition of bacterial (a) and fungal (b) communities of Dataset 1 (H) and Dataset 2 (F and N) (filtered on minimum of 10% average) in fresh and stored bee pollen

