

Supplementary data

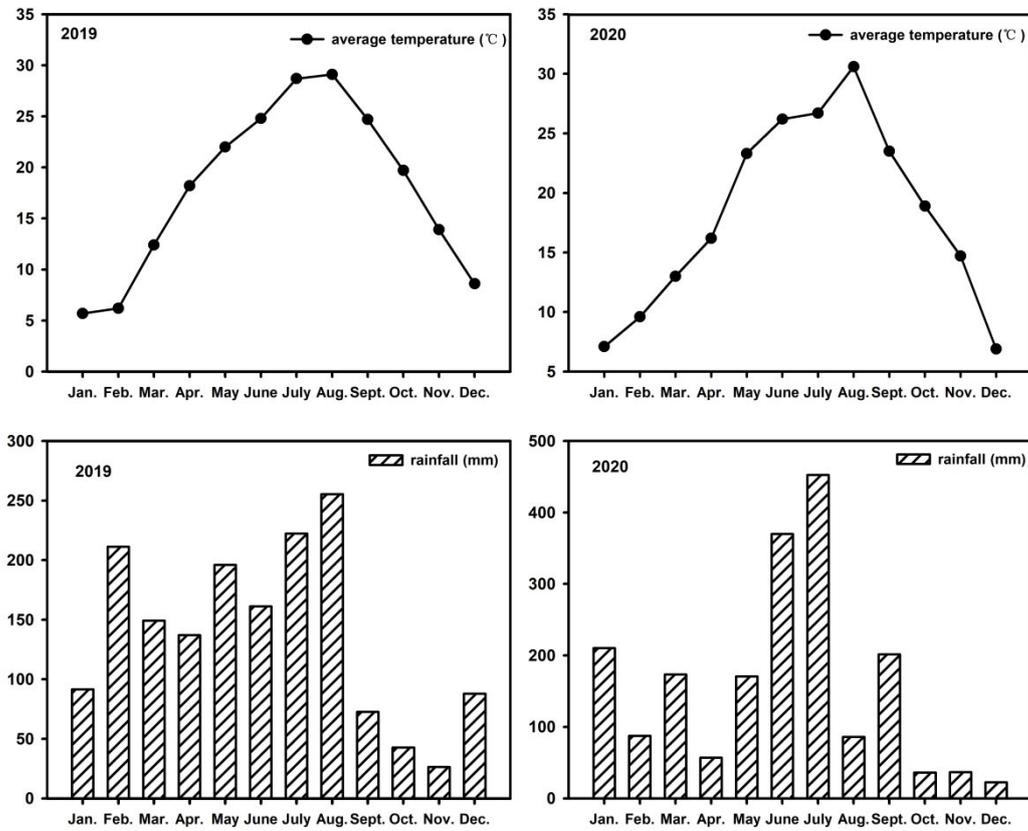


Figure S1. Weather data in the experimental location Hangzhou in 2019 and 2020.

Table S1. Heatmap data of soil bacterial community at genus level to be involved in greenhouse gas emission influenced by the mulching cultivation.

| Category | Microbe | CK-PI | BM-PI | CK-HD | BM-HD | |
|-------------------------|-----------------------|----------------|----------|----------|----------|----------|
| Methanogenic bacteria | Methanobacterium | 0.000938 | 0 | 0 | 0.000554 | |
| | Methanolinea | 0.000445 | 0 | 0 | 0.000554 | |
| Methanotrophic bacteria | Methylocaldum | 0.000513 | 0.00246 | 0.00164 | 0.005762 | |
| | Methylococcus | 0.026292 | 0.055058 | 0.067501 | 0.089393 | |
| | Methylomonas | 0.020758 | 0.073076 | 0.046221 | 0.082031 | |
| | Methyloparacoccus | 0.001918 | 0.025524 | 0.002123 | 0.008819 | |
| | Methylosarcina | 0.027123 | 0.032661 | 0.023958 | 0.049168 | |
| | Methyloversatilis | 0.056788 | 0.112903 | 0.161146 | 0.068164 | |
| | Methylobacter | 0.017916 | 0.026403 | 0.033377 | 0.034923 | |
| | Methylocystis | 0.340776 | 0.380144 | 0.240219 | 0.226578 | |
| | Methylogaea | 0.0015 | 0.00242 | 0.004418 | 0.005289 | |
| | Organic decomposition | Nocardia | 0.000515 | 0.002489 | 0.004926 | 0.006316 |
| Streptomyces | | 0.00687 | 0.009382 | 0.010495 | 0.018801 | |
| Syntrophorhabdus | | 0.37414 | 0.498595 | 0.499822 | 0.530512 | |
| Mycobacterium | | 0.062448 | 0.073567 | 0.096336 | 0.093036 | |
| Pseudomonas | | 0.00387 | 0.001023 | 0.037299 | 0.003136 | |
| Dehalogenimonas | | 0.164211 | 0.238048 | 0.22911 | 0.27139 | |
| Georgfuchsia | | 0.028758 | 0.046258 | 0.055211 | 0.069127 | |
| Nitrogen fixation | | Rhizomicrobium | 0.288196 | 0.329226 | 0.361877 | 0.268236 |
| | | Ktedonobacter | 0.009798 | 0.01238 | 0.03081 | 0.024152 |
| | | Sideroxydans | 0.163912 | 1.176299 | 1.779927 | 2.220718 |
| Nitrification | Nitrospira | 0.12786 | 0.096099 | 0.090731 | 0.138919 | |
| | Nitrospira | 0.301947 | 0.241455 | 0.234005 | 0.251962 | |
| Denitrification | Thiobacillus | 0.095359 | 0.188368 | 0.203669 | 0.157033 | |
| | Azospirillum | 0.005516 | 0.00294 | 0.003336 | 0.021849 | |
| | Bacillus | 0.00826 | 0.00895 | 0.012129 | 0.020852 | |
| | Pseudomonas | 0.00387 | 0.001023 | 0.037299 | 0.003136 | |
| | Ralstonia | 0.001454 | 0.000519 | 0.004366 | 0.001035 | |
| | Flavisolibacter | 0.069294 | 0.089169 | 0.084458 | 0.098238 | |
| | Acinetobacter | 0.001006 | 0.00048 | 0.01862 | 0.002597 | |
| | Chromobacterium | 0.000987 | 0.000438 | 0.003876 | 0.035229 | |
| | Flavobacterium | 0.005555 | 0.000956 | 0.001649 | 0.008311 | |
| | Hyphomicrobium | 0.017412 | 0.034536 | 0.076832 | 0.031958 | |
| Iron-sulfate reducer | Geobacter | 1.815637 | 1.720515 | 2.35395 | 2.583129 | |
| | Desulfatiglans | 0.071574 | 0.103849 | 0.101799 | 0.110119 | |
| | Desulfobacca | 0.024338 | 0.02957 | 0.053848 | 0.04088 | |
| | Desulfobulbus | 0.021712 | 0.017888 | 0.020348 | 0.035106 | |
| | Desulfocapsa | 0.005435 | 0.005488 | 0.01537 | 0.008951 | |
| | Desulfovibrio | 0.021691 | 0.034788 | 0.072577 | 0.078035 | |
| | Desulfuromonas | 0.000494 | 0.005401 | 0.010998 | 0.004704 | |

| | | | | |
|---------------------|----------|----------|----------|----------|
| Sulfuritalea | 0.004471 | 0.014022 | 0.020047 | 0.038184 |
| Thermodesulfovibrio | 4.665419 | 5.369562 | 6.284261 | 5.731618 |
