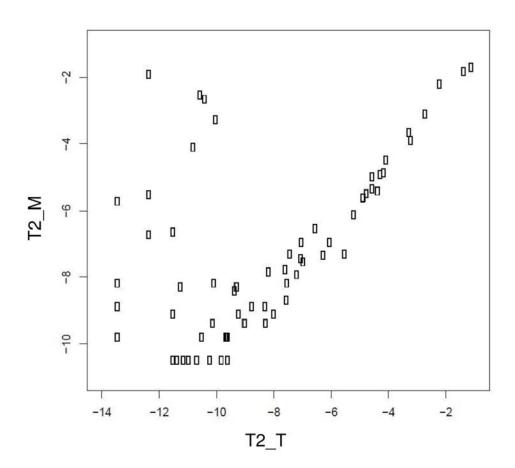
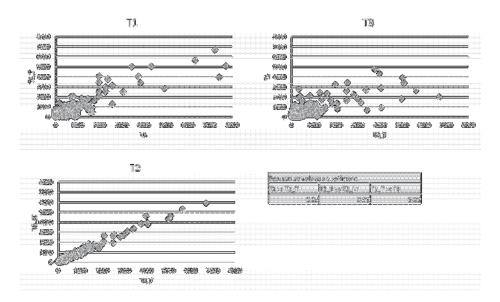
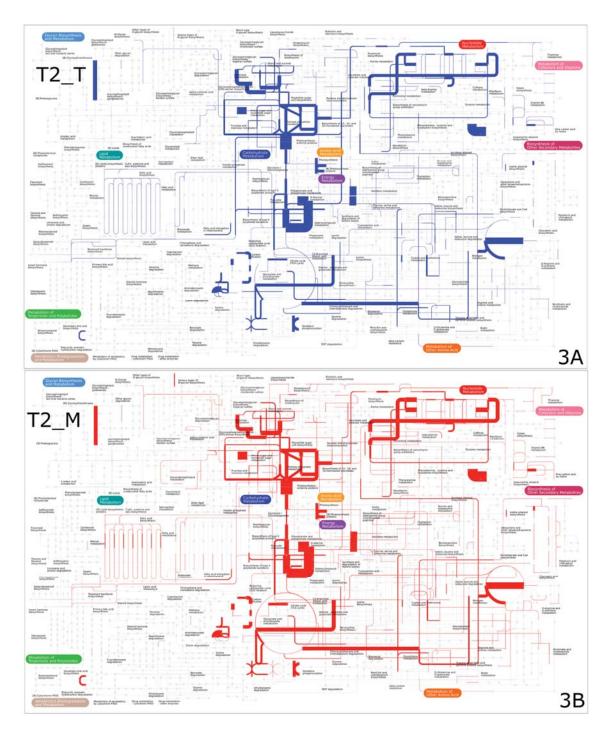
Table and Figure Legends

Supplementary Figure 1. Log relative abundances of common genera between 2D_1 and 2D_2.



Supplementary Figure 2. Correlations of the expression pattern between the samples.





- K01176 amyA, malS; alpha-amylase;
- KO1182 oligo-1,6-glucosidase;
- K01186 NEU1; sialidase-1;
- KO1187 malZ; alpha-glucosidase;
- K01190 lacZ; beta-galactosidase;
- K01191 alpha-mannosidase;
- KO1192 MANBA, manB; beta-mannosidase;
- K01193 sacA; beta-fructofuranosidase;
- K01197 hya; hyaluronoglucosaminidase;
- KO1198 xynB; xylan 1,4-beta-xylosidase;
- K01200 pullulanase;
- KO1205 NAGLU; alpha-N-acetylglucosaminidase;
- K01206 FUCA; alpha-L-fucosidase;
- KO1207 nagZ; beta-N-acetylhexosaminidase;
- K01209 abfA; alpha-N-arabinofuranosidase;
- K01210 glucan 1,3-beta-glucosidase;
- KO1215 dexB; glucan 1,6-alpha-glucosidase;
- K01220 lacG; 6-phospho-beta-galactosidase;
- KO1223 bglA; 6-phospho-beta-glucosidase;
- KO1227 mannosyl-glycoprotein endo-beta-N-acetylglucosaminidase;
- K01232 glvA; maltose-6'-phosphate glucosidase;
- K01234 nplT; neopullulanase;
- K01239 iunH; purine nucleosidase;
- K01240 URH1; uridine nucleosidase;
- K01243 mtnN, mtn, pfs; adenosylhomocysteine nucleosidase;
- KO1246 tag; DNA-3-methyladenine glycosylase I;
- KO1250 rihA; pyrimidine-specific ribonucleoside hydrolase;
- KO1811 yicl; alpha-D-xyloside xylohydrolase;
- KO2438 treX, glgX; glycogen operon protein;
- K03332 fruA; fructan beta-fructosidase;
- KO3575 mutY; A/G-specific adenine glycosylase;
- K03648 UNG, UDG; uracil-DNA glycosylase;
- K03660 OGG1; N-glycosylase/DNA lyase;
- K05349 bglX; beta-glucosidase;
- K05520 pfpl; protease l;
- K05546 GANAB; alpha 1,3-glucosidase;
- KO6113 abnA B; arabinan endo-1,5-alpha-L-arabinosidase;
- K07407 galA, rafA; alpha-galactosidase;
- K08309 slt; soluble lytic murein transglycosylase;
- K12111 ebgA; evolved beta-galactosidase subunit alpha;
- K12308 bgaB, lacA; beta-galactosidase;
- K12309 GLB1, ELNR1; beta-galactosidase;
- K12373 HEXA_B; hexosaminidase;
- K15532 yteR, yesR; unsaturated rhamnogalacturonyl hydrolase;

- NCOAT, MGEA5; protein O-GlcNAcase / histone
- K15719 acetyltransferase
- K15921 xynD; arabinoxylan arabinofuranohydrolase;
- K15923 AXY8, FUC95A, afcA; alpha-L-fucosidase 2
- 411 **Supplementary Figure 4.** Overall expression of mRNA functions from LGG on the visualization tool
- 412 iPATH2.0.

