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Supplementary Materials

Environmental Risk Assessment of Living Modified Microorganisms (LMM) on the Indigenous Microbial Community

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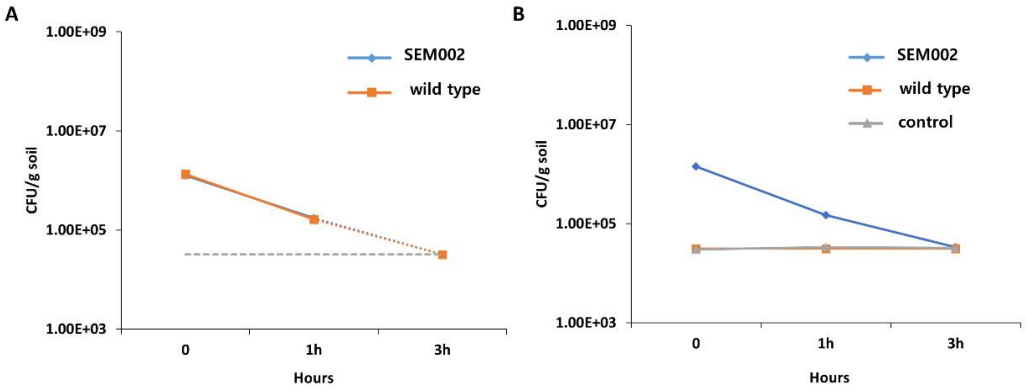

Figure S1. Changes in the bacterial counts of LMM and wild-type strains in LB medium and LPK medium after the release of LMM and wild-type strains into forest soil microcosm. A, LB medium; B, LPK medium. SEM002, number of SEM002 cells in the SEM002-inoculated soil; wild type, number of wild-type cells in the wild type-inoculated soil; control, antibiotic-resistant bacterial counts in the control group Dotted lines indicate that the inoculated strains were undetectable after 3 h of the experiment. Dotted lines parallel to the x-axis indicate the background level of indigenous microorganisms.

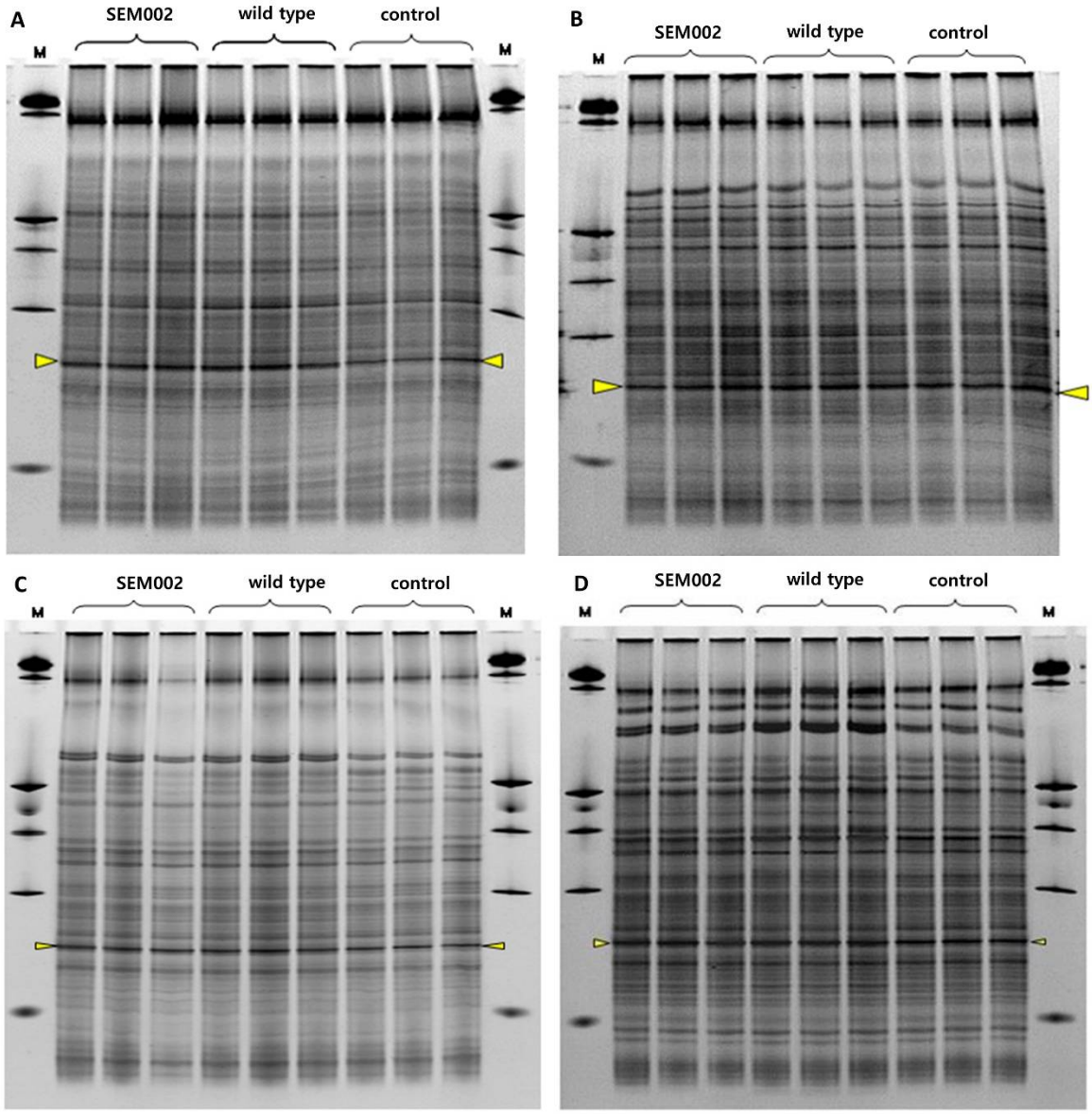

Figure S2. Analysis of structural changes of the bacterial community in the factory soil microcosm experiment using real-time PCR-DGGE. A, Day 15; B, Day 35; C, Day 55; D, Day 75. SEM002, LMM-inoculated soil; Wild type, ATCC 13032 strain-inoculated soil; Control, uninoculated soil. The arrows indicate DGGE bands of LMM and wild-type strains.

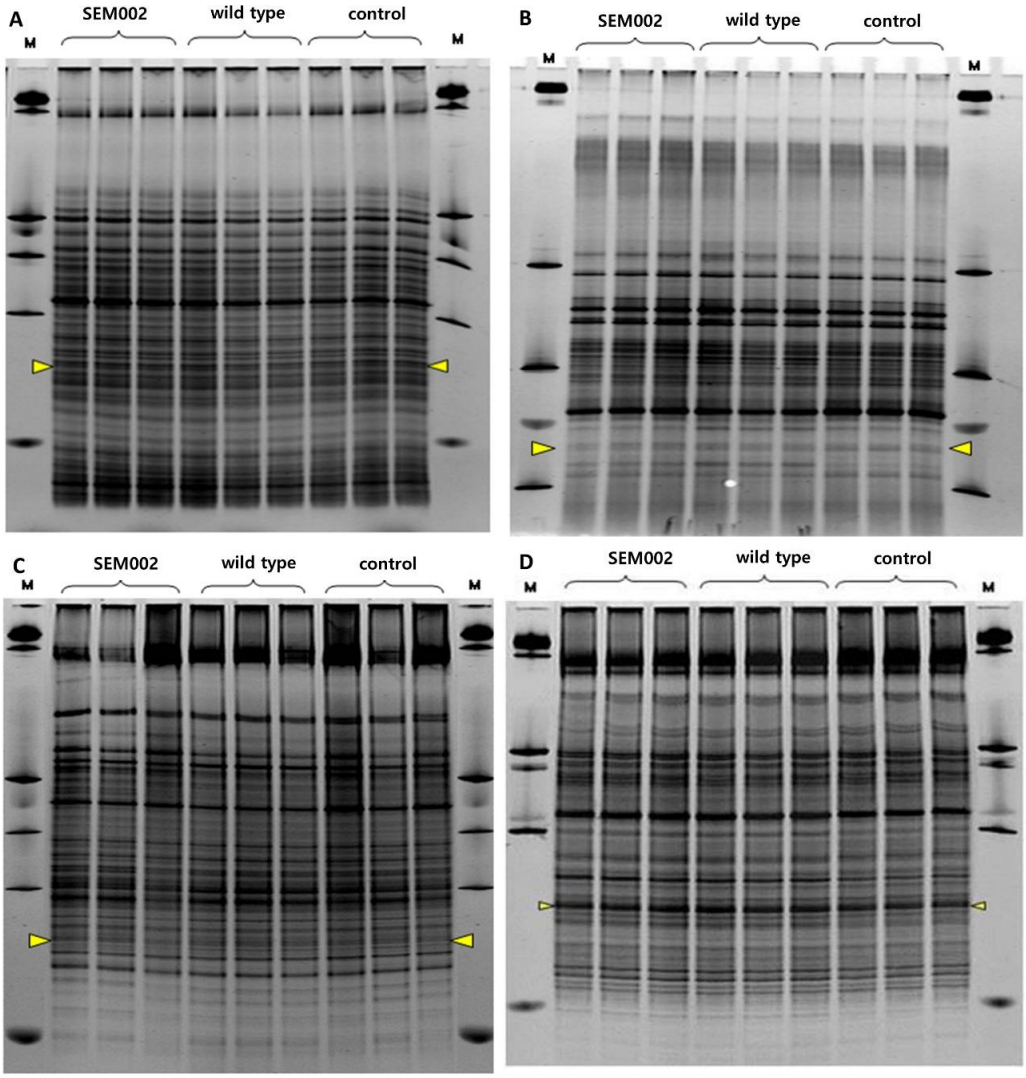

Figure S3. Analysis of structural changes of the bacterial community in the factory soil microcosm experiment using real-time PCR-DGGE. A, Day 13; B, Day 33; C, Day 54; D, Day 75. SEM002, LMM-inoculated soil; Wild type, ATCC 13032 strain-inoculated soil; Control, uninoculated soil. The arrows indicate DGGE bands of LMM and wild-type strains.

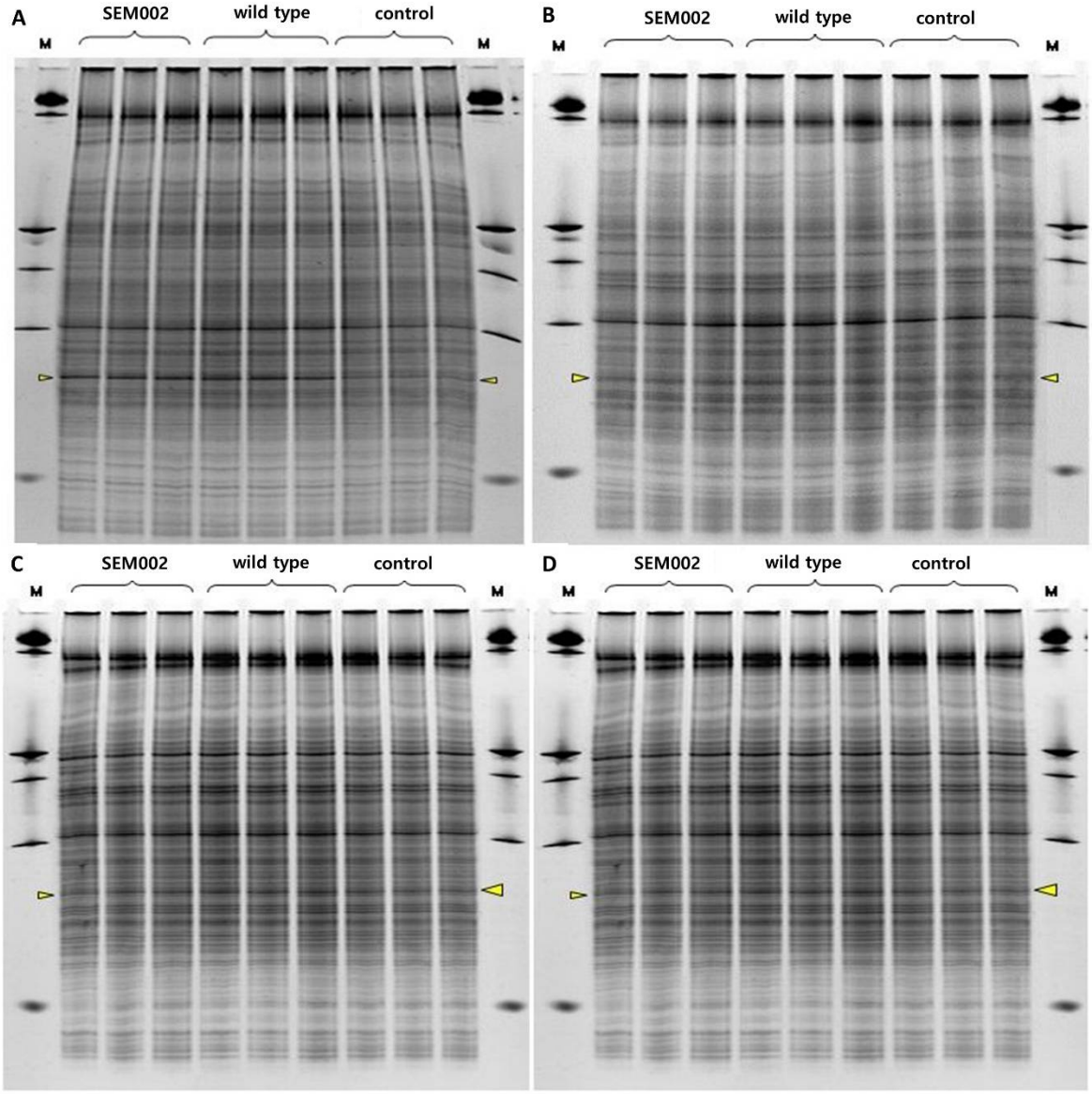

Figure S4. Analysis of structural changes of the bacterial community in the factory soil microcosm experiment using real-time PCR-DGGE. A, Day 15; B, Day 35; C, Day 55; D, Day 75. SEM002, LMM-inoculated soil; Wild type, ATCC 13032 strain-inoculated soil; Control, uninoculated soil. The arrows indicate DGGE bands of LMM and wild-type strains.