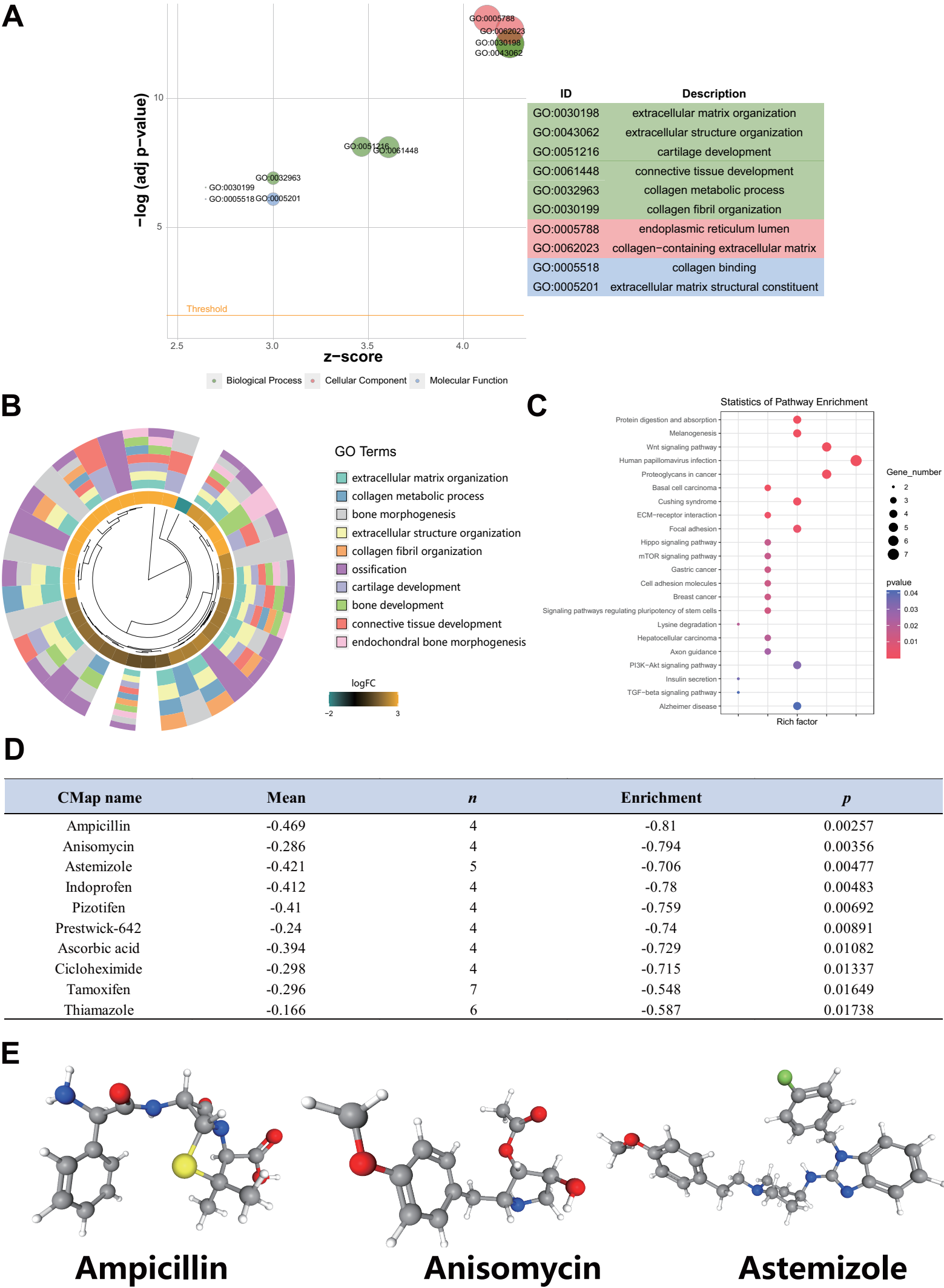




Supplementary Fig. 1 Function analysis of BML-specific DEGs
(A-B) Gene Ontology (GO) enrichment analysis. Y-axis, negative log-adjusted p-value; x-axis, z-score; Bubble area positively correlated with counts of gene numbers in indicated terms. Green, biological process (BP); pink, cellular component (CC); blue, molecular function (MF). Adjust P-value < 0.05 (orange cross-line) was considered significant.
(C) Results of Kyoto Encyclopedia of Genes and Genomes (KEGG) Pathway Enrichment Analysis.
(D) Results of CMap analysis, sorted by p-value from small to large order.
(E) The 3D chemical structure of the top three small molecule drugs for BMLs (from left to right respectively: Ampicillin, Anisomycin, Astemizole).