

SUPPLEMENTARY TABLES

Supplementary Table S1: List of genes overexpressed in cells treated with either TRAIL or C3/C2 or SN, or TRAIL+C3/C2 or TRAIL+SN.

See Supplementary File S1

Supplementary Table S2: KEGG pathway enrichment analysis of genes overexpressed in treated cells using FatiGO tool. List of genes are compared with genome and the significance of KEGG pathway enrichment was indicated as p-value. KEGG ID: KEGG Identifier. Hsa: homosapiens.

See Supplementary File S2