

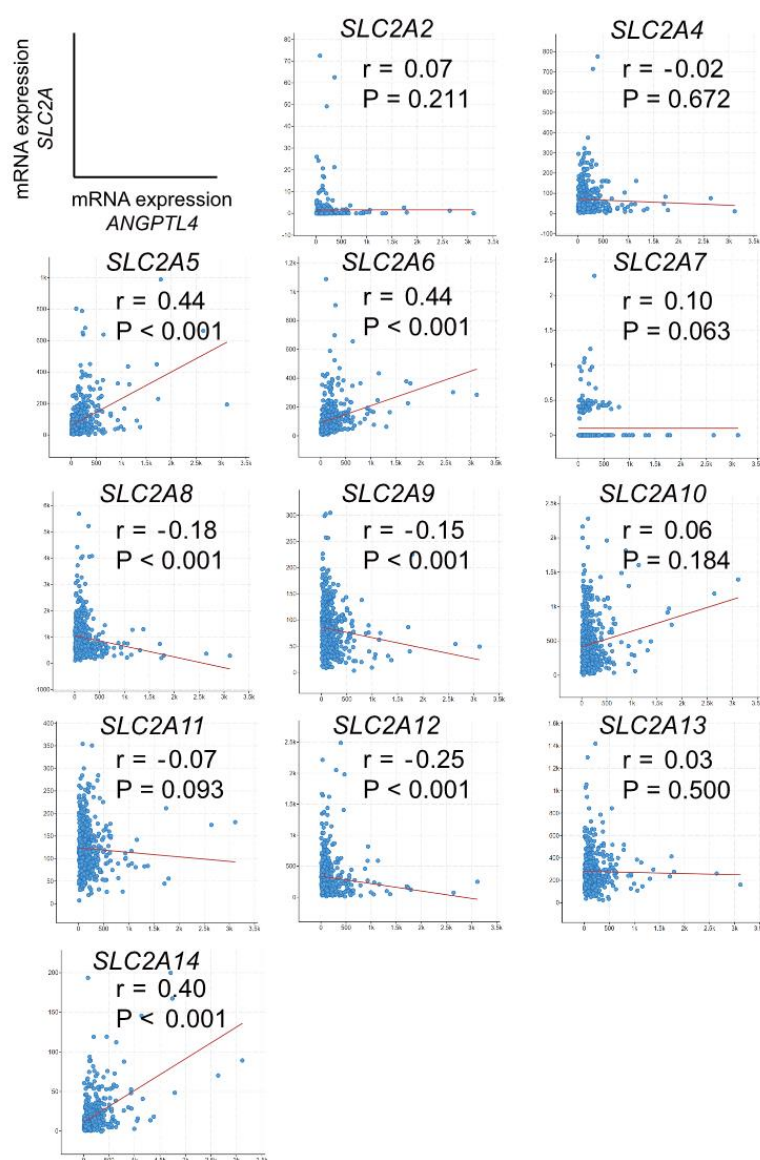


Fig. S2

Correlation of *ANGPTL4* and *SLC2A* mRNA expression levels in CRC.

The correlation between *ANGPTL4* and *SLC2A* mRNA expression levels in CRC was evaluated by using data from The Cancer Genome Atlas (TCGA) database. Coexpression analysis was performed, and the results were downloaded from the cBioPortal for Cancer Genomics (<http://www.cbioportal.org/>).