

Legend

Supplementary Figure 1. Bioinformatic analysis of differentially expressed proteins in HCT 116 cells versus NCM460 cells obtained by LFQ LC-MS/MS. STRING v11.0 was used to analyze the differentially expressed 901 genes listed in Supplementary Table 1 (899 were recognized by STRING). The proteins have significantly more interactions (7610 edges) than the expected 5050 ($p < 1 \times 10^{-16}$). The analysis indicated that the proteins are at least partially biologically related. Details are given in the text.