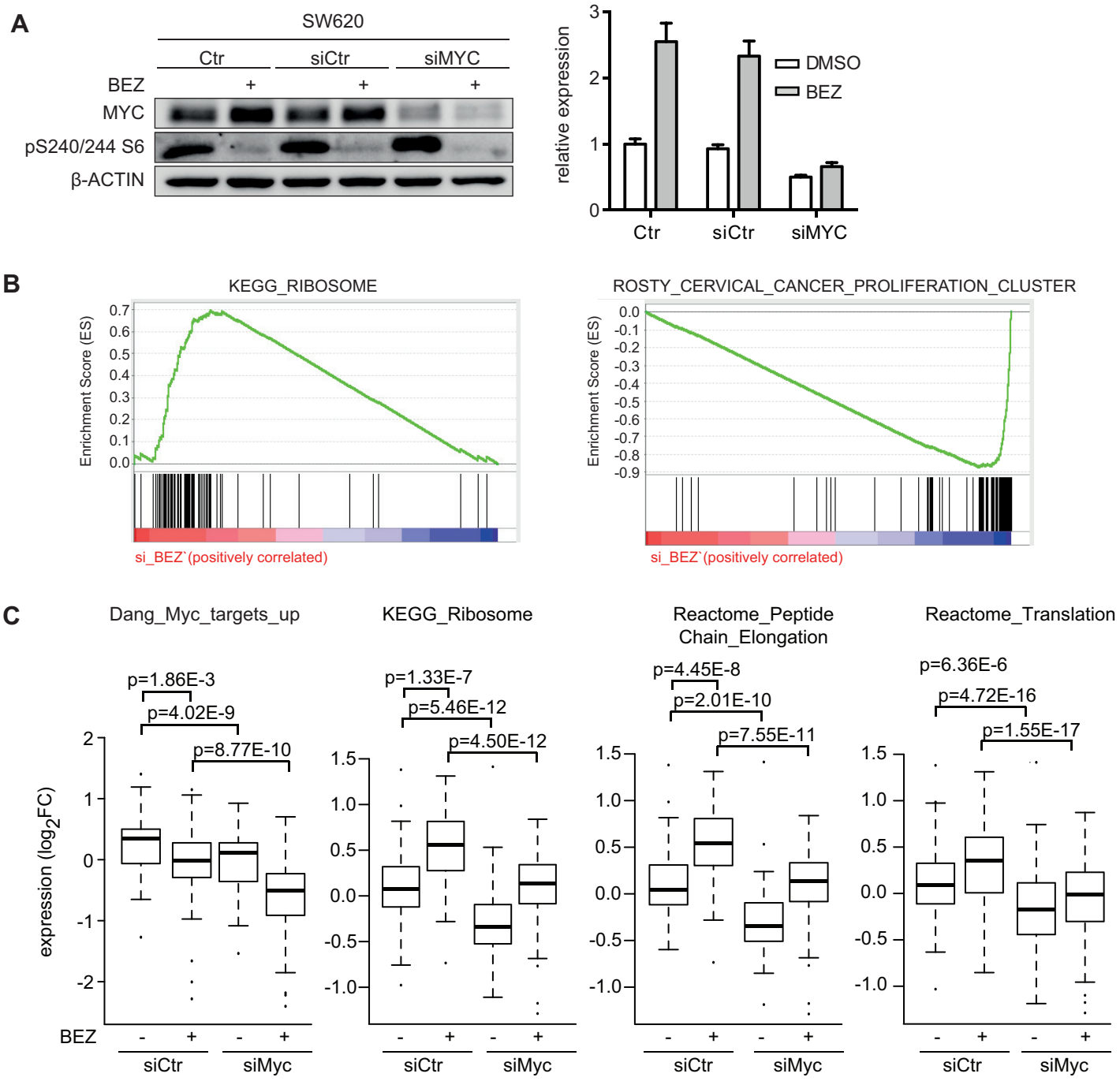


D

| Number | Gene set enriched | Nominal p-value | FDR q-value |
|---|---|---|---|
| 1 | KEGG_RIBOSOME | <0.001 | <0.001 |
| 2 | REACTOME_PEPTIDE_CHAIN_ELONGATION | <0.001 | <0.001 |
| 7 | REACTOME_TRANSLATION | <0.001 | <0.001 |

| Number | Gene set down | Nominal p-value | FDR q-value |
|---|---|---|---|
| 1 | KOBAYASHI_EGFR_SIGNALING_24HR_DN | <0.001 | <0.001 |
| 2 | ROSTY_CERVICAL_CANCER_PROLIFERATION_CLUSTER | <0.001 | <0.001 |
| 5 | GRAHAM_CML_DIVIDING_VS_NORMAL_QUIESCENT_UP | <0.001 | <0.001 |
| 6 | GRAHAM_NORMAL_QUIESCENT_VS_NORMAL_DIVIDING_DN | <0.001 | <0.001 |
| 16 | BENPORATH_PROLIFERATION | <0.001 | <0.001 |
| 20 | REACTOME_CELL_CYCLE_MITOTIC | <0.001 | <0.001 |