

Supplementary Figure S1. Doxycycline-induced reconstitution of TGFBR2 expression and downstream signaling. Western blot analysis demonstrated dox-induced reconstituted expression of TGFBR2 and phosphorylation of SMAD2 (pSMAD2) in protein lysates of HCT116-TGFBR2 cells. Total SMAD2 protein was used as a loading control. Protein sizes are indicated. .


Supplementary Figure S2 Incorporation of heavy-labeled amino acids in EVs isolated from TGFBR2proficient HCT116-TGFBR2 cells. Incorporation efficiency of heavy-labeled amino acids was calculated based on 144 identified peptides.


Supplementary Figure S3. Distribution of quantified protein ratios. Histograms are shown for $\log _{2}-$ transformed protein ratios (dT/pT) from each biological replicate of EVs (A) and cells (B).

Supplementary Table S1 TGFBR2-regulated proteins in cells.

| Gene names | Protein names | Protein ratio ${ }^{1}$ (dT/pT) | Peptides (unique) | Coverage (\%) | Identification score |
| :---: | :---: | :---: | :---: | :---: | :---: |
| Upregulated cellular proteins by TGFBR2 deficiency |  |  |  |  |  |
| NUMA1 | Nuclear mitotic apparatus protein 1 | 2.23 | 12 (12) | 8.3 | 36.38 |
| CAD | CAD protein | 1.91 | 20 (20) | 13.1 | 85.112 |
| GLUL | Glutamine synthetase | 1.82 | 7 (7) | 25.5 | 29.43 |
| UBE2Z | Ubiquitin-conjugating enzyme E2 Z | 1.77 | 4 (4) | 13.3 | 13.23 |
| TUBB4A | Tubulin beta-4A chain | 1.71 | 17 (17) | 6.3 | 26.76 |
| EDC4 | Enhancer of mRNAdecapping protein 4 | 1.56 | 7 (7) | 9.0 | 27.22 |
| RANBP2 | E3 SUMO-protein ligase RanBP2 | 1.51 | 6 (6) | 3.4 | 20.66 |


| KRT18 | Keratin, type I <br> cytoskeletal 18 | 0.61 | $30(19)$ | 77 | 323.31 |
| :---: | :---: | :---: | :---: | :---: | :---: |
| VDAC3 | Voltage-dependent <br> anion-selective channel <br> protein 3 | 0.63 | $9(9)$ | 41.7 | 63.09 |
| EMG1 | Ribosomal RNA small <br> subunit <br> methyltransferase NEP1 | 0.64 | $4(4)$ | 22.1 | 15.02 |
| KRT8 | Keratin, type II <br> cytoskeletal 8 <br> Voltage-dependent <br> Vnion-selective channel <br> protein 2 | 0.66 | $39(31)$ | 64.2 | 323.31 |
| HNAC2 | 0.65 | $13(13)$ | 59.5 | 174.22 |  |
|  | Hematological and <br> neurological expressed 1 | 0.67 | $3(3)$ | 54.5 | 18.21 |

[^0]
[^0]:    ${ }^{1}$ Mean ratios calculated from three biological replicates.

