**Title:** Metformin Increases Sensitivity of Pancreatic Cancer Cells to Gemcitabine by Reducing $CD133^+$ Cell Populations and Suppressing ERK/P70S6K Signaling

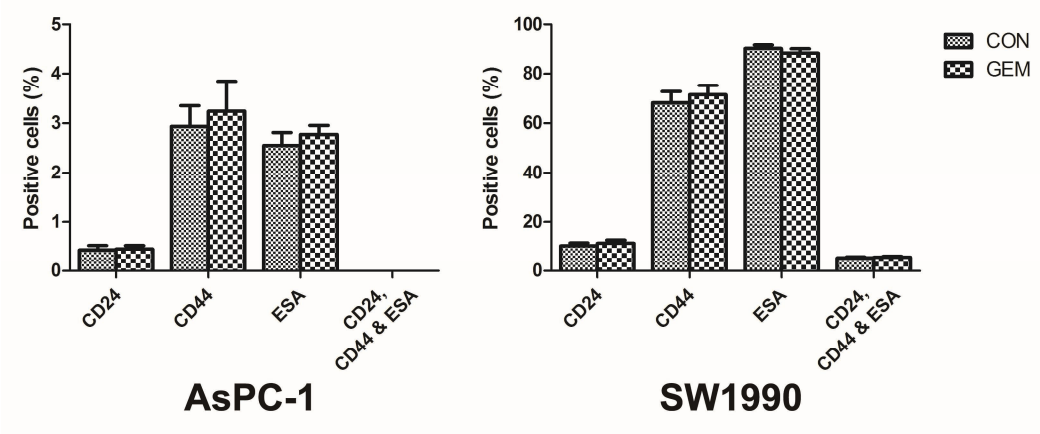
**Authors:**

Xinqun Chai[1], Hongpeng Chu[1], Xuan Yang[1], Yuanpu Meng[1], Pengfei Shi[2], Shanmiao Gou[3].

[1] Department of Hepatobiliary Surgery, Union Hospital, Tongji Medical College, Huazhong University of Science and Technology.

[2] Department of Breast and Thyroid Surgery, Central Hospital of Wuhan.

[3] Pancreatic Disease Institute, Union Hospital, Tongji Medical College, Huazhong University of Science and Technology.




**Supplementary Figure S1. Gemcitabine didn't change the proportion of $CD24^+CD44^+ESA^+$ cells in pancreatic cancer cells.** AsPC-1 and SW1990 pancreatic cancer cells were treated with 300 nM gemcitabine for 48 h, and the proportion of $CD24^+$, $CD44^+$, $ESA^+$ and $CD24^+CD44^+ESA^+$ cells was determined by flow cytometry. Gemcitabine did not affect $CD24^+$, $CD44^+$, $ESA^+$ or $CD24^+CD44^+ESA^+$ cells ($CD24^+CD44^+ESA^+$ cells are not detectable among AsPC-1 cells). CON, control; GEM, gemcitabine.

Supplementary Table S1. Effect of gemcitabine on $CD133^+$ and $CD133^-$ pancreatic cancer cells

| | Control | | | Gemcitabine (300 nM, 48 hours) | | | | |
|---|---|---|---|---|---|---|---|---|
| | **A**: Cell numbers (relative to control, mean ± standard deviation, %) | **B**: $CD133^+$ cells (mean ± standard deviation, %) | **C**: $CD133^-$ cells (mean ± standard deviation, %) | **D**: Cell numbers (relative to control, mean ± standard deviation, %) | **E**: $CD133^+$ cells (mean ± standard deviation, %) | **F**: $CD133^-$ cells (mean ± standard deviation, %) | **G:** Proportion of survived $CD133^+$ cells ((D*E)/(A*B), %) | **H:** Proportion of survived $CD133^-$ cells ((D*F)/(A*C), %) |
| AsPC-1 | 100 ± 2.83 | 5.18 ± 0.94 | 94.82 ± 0.94 | 40.73 ± 2.51 | 11.00 ± 2.34 | 89.00 ± 2.34 | **86.41** | **38.23** |
| SW1990 | 100 ± 2.51 | 4.38 ± 1.08 | 95.62 ± 1.08 | 31.33 ± 1.93 | 8.40 ± 1.70 | 91.60 ± 1.70 | **60.14** | **30.02** |

Supplementary Table S2. Correlation between 193 proteins and histological grade of pancreatic cancer using reverse phase protein array (RPPA)

| Protein | CC | 95% CI (Lower) | 95% CI (Upper) | P | t | df |
|---|---|---|---|---|---|---|
| beta | 0.0995 | -0.132 | 0.3207 | 0.3991 | 0.8482 | 72 |
| epsilon | -0.026 | -0.253 | 0.2037 | 0.826 | -0.2207 | 72 |
| zeta | -0.0996 | -0.3208 | 0.1319 | 0.3986 | -0.8492 | 72 |
| 4E-BP1 | -0.1277 | -0.3461 | 0.1039 | 0.2783 | -1.0923 | 72 |
| 4E-BP1_pS65 | -0.0936 | -0.3153 | 0.1379 | 0.4278 | -0.7975 | 72 |
| 4E-BP1_pT37_T46 | -0.0963 | -0.3178 | 0.1352 | 0.4143 | -0.8211 | 72 |
| 4E-BP1_pT70 | 0.0694 | -0.1616 | 0.2933 | 0.5567 | 0.5906 | 72 |
| 53BP1 | 0.0697 | -0.1614 | 0.2935 | 0.5553 | 0.5926 | 72 |
| A-Raf_pS299 | -0.1267 | -0.3452 | 0.1048 | 0.2819 | -1.0842 | 72 |
| ACC_pS79 | 0.1251 | -0.1064 | 0.3438 | 0.2881 | 1.0701 | 72 |
| ACC1 | -0.0121 | -0.24 | 0.217 | 0.9184 | -0.1028 | 72 |
| Acetyl-a-Tubulin-Lys40 | 0.0495 | -0.181 | 0.2749 | 0.6752 | 0.4207 | 72 |
| ACVRL1 | -0.0371 | -0.2634 | 0.193 | 0.7534 | -0.3153 | 72 |
| ADAR1 | 0.0071 | -0.2217 | 0.2352 | 0.952 | 0.0604 | 72 |
| Akt | 0.1417 | -0.0897 | 0.3586 | 0.2284 | 1.2149 | 72 |
| Akt_pS473 | 0.0888 | -0.1426 | 0.311 | 0.4518 | 0.7564 | 72 |
| Akt_pT308 | 0.0161 | -0.2132 | 0.2437 | 0.8918 | 0.1365 | 72 |
| alpha-Catenin | -0.1482 | -0.3643 | 0.0832 | 0.2077 | -1.2712 | 72 |
| AMPK_alpha | -0.0453 | -0.271 | 0.1851 | 0.7015 | -0.3848 | 72 |
| AMPK_pT172 | 0.2365 | 0.0085 | 0.4412 | 0.0425 | 2.0653 | 72 |
| Annexin-1 | 0.0695 | -0.1616 | 0.2933 | 0.5564 | 0.5909 | 72 |
| Annexin_VII | -0.2238 | -0.4303 | 0.0049 | 0.0552 | -1.9489 | 72 |
| AR | 0.0851 | -0.1462 | 0.3076 | 0.471 | 0.7247 | 72 |
| ARHI | -0.1072 | -0.3277 | 0.1243 | 0.3631 | -0.9152 | 72 |
| ASNS | -0.0378 | -0.264 | 0.1923 | 0.749 | -0.3212 | 72 |
| ATM | 0.1152 | -0.1164 | 0.3348 | 0.3286 | 0.9837 | 72 |
| B-Raf | 0.1457 | -0.0856 | 0.3621 | 0.2155 | 1.2497 | 72 |
| Bad_pS112 | -0.0224 | -0.2496 | 0.2071 | 0.8496 | -0.1903 | 72 |
| Bak | 0.1118 | -0.1197 | 0.3318 | 0.3428 | 0.9549 | 72 |
| Bap1c-4 | 0.0735 | -0.1576 | 0.297 | 0.5337 | 0.6254 | 72 |
| Bax | 0.04 | -0.1903 | 0.266 | 0.7354 | 0.3393 | 72 |
| Bcl-2 | -0.0054 | -0.2336 | 0.2234 | 0.9635 | -0.0459 | 72 |
| Bcl-xL | -0.0473 | -0.2729 | 0.1831 | 0.6888 | -0.4021 | 72 |
| Beclin | 0.0184 | -0.211 | 0.2459 | 0.8763 | 0.1562 | 72 |
| beta-Catenin | 0.0541 | -0.1766 | 0.2791 | 0.6473 | 0.4594 | 72 |
| Bid | 0.1336 | -0.0979 | 0.3514 | 0.2563 | 1.1442 | 72 |
| Bim | 0.125 | -0.1066 | 0.3437 | 0.2887 | 1.0689 | 72 |
| BRCA2 | 0.0867 | -0.1446 | 0.3091 | 0.4625 | 0.7388 | 72 |

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|---|---|---|---|---|---|---|
| c-Jun_pS73 | -0.1051 | -0.3258 | 0.1264 | 0.3729 | -0.8967 | 72 |
| c-Kit | -0.0837 | -0.3064 | 0.1476 | 0.4782 | -0.7129 | 72 |
| c-Met_pY1235 | -0.0802 | -0.3032 | 0.1511 | 0.4969 | -0.6828 | 72 |
| c-Myc | -0.2441 | -0.4477 | -0.0166 | 0.0361 | -2.1362 | 72 |
| C-Raf | 0.1563 | -0.0749 | 0.3715 | 0.1836 | 1.3428 | 72 |
| C-Raf_pS338 | -0.0187 | -0.2462 | 0.2107 | 0.8742 | -0.1589 | 72 |
| Caspase-7_cleavedD198 | 0.1411 | -0.0903 | 0.358 | 0.2305 | 1.2092 | 72 |
| Caveolin-1 | 0.0583 | -0.1725 | 0.283 | 0.6219 | 0.4953 | 72 |
| CD20 | 0.1229 | -0.1086 | 0.3418 | 0.2968 | 1.051 | 72 |
| CD31 | -0.0708 | -0.2946 | 0.1602 | 0.5486 | -0.6027 | 72 |
| CD49b | -0.0736 | -0.2971 | 0.1575 | 0.533 | -0.6265 | 72 |
| CDK1 | -0.0896 | -0.3117 | 0.1418 | 0.4476 | -0.7635 | 72 |
| Chk1 | 0.186 | -0.0444 | 0.3976 | 0.1126 | 1.6062 | 72 |
| Chk1_pS345 | -0.0871 | -0.3095 | 0.1442 | 0.4603 | -0.7423 | 72 |
| Chk2 | -0.137 | -0.3544 | 0.0945 | 0.2445 | -1.1734 | 72 |
| Chk2_pT68 | -0.0824 | -0.3052 | 0.1489 | 0.485 | -0.7019 | 72 |
| cIAP | 0.0614 | -0.1695 | 0.2859 | 0.6034 | 0.5219 | 72 |
| Claudin-7 | -0.1579 | -0.373 | 0.0732 | 0.1789 | -1.3573 | 72 |
| Collagen_VI | 0.0278 | -0.2019 | 0.2547 | 0.8139 | 0.2363 | 72 |
| Cyclin_B1 | 0.1054 | -0.1262 | 0.326 | 0.3716 | 0.8991 | 72 |
| Cyclin_D1 | 0.124 | -0.1075 | 0.3428 | 0.2923 | 1.0608 | 72 |
| Cyclin_E1 | 0.0576 | -0.1731 | 0.2824 | 0.6257 | 0.4899 | 72 |
| Cyclin_E2 | 0.0843 | -0.147 | 0.3069 | 0.4749 | 0.7182 | 72 |
| DJ-1 | -0.2208 | -0.4277 | 0.0081 | 0.0587 | -1.9208 | 72 |
| Dvl3 | 0.0329 | -0.1971 | 0.2594 | 0.7808 | 0.2793 | 72 |
| E-Cadherin | -0.0309 | -0.2576 | 0.199 | 0.7935 | -0.2627 | 72 |
| eEF2 | -0.0332 | -0.2597 | 0.1968 | 0.7791 | -0.2815 | 72 |
| eEF2K | -0.0382 | -0.2643 | 0.192 | 0.7469 | -0.324 | 72 |
| EGFR | 0.1643 | -0.0667 | 0.3786 | 0.1619 | 1.4134 | 72 |
| EGFR_pY1068 | -0.0328 | -0.2593 | 0.1972 | 0.7815 | -0.2784 | 72 |
| EGFR_pY1173 | 0.0618 | -0.1691 | 0.2863 | 0.6008 | 0.5256 | 72 |
| eIF4E | -0.2292 | -0.4349 | -8.00E-04 | 0.0495 | -1.9982 | 72 |
| eIF4G | 0.0714 | -0.1597 | 0.2951 | 0.5454 | 0.6076 | 72 |
| ER-alpha | 0.1886 | -0.0417 | 0.3999 | 0.1076 | 1.6294 | 72 |
| ER-alpha_pS118 | 0.0934 | -0.138 | 0.3152 | 0.4285 | 0.7963 | 72 |
| ERCC1 | 0.1383 | -0.0931 | 0.3556 | 0.24 | 1.1849 | 72 |
| ERK2 | 0.0327 | -0.1973 | 0.2592 | 0.7823 | 0.2774 | 72 |
| ETS-1 | 0.1862 | -0.0442 | 0.3977 | 0.1123 | 1.6077 | 72 |
| FASN | 0.0642 | -0.1667 | 0.2885 | 0.5867 | 0.546 | 72 |
| Fibronectin | 0.1393 | -0.0921 | 0.3565 | 0.2365 | 1.1936 | 72 |
| FoxM1 | 0.0224 | -0.2071 | 0.2497 | 0.8495 | 0.1904 | 72 |
| FOXO3a | -0.1608 | -0.3755 | 0.0703 | 0.1711 | -1.3825 | 72 |
| FOXO3a_pS318_S321 | -0.1412 | -0.3582 | 0.0902 | 0.23 | -1.2106 | 72 |
| G6PD | -0.0234 | -0.2505 | 0.2062 | 0.8434 | -0.1982 | 72 |

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|---|---|---|---|---|---|---|
| Gab2 | 0.0842 | -0.1471 | 0.3068 | 0.4758 | 0.7168 | 72 |
| GAPDH | 0.0393 | -0.1909 | 0.2654 | 0.7394 | 0.334 | 72 |
| GATA3 | 0.1152 | -0.1164 | 0.3348 | 0.3286 | 0.9837 | 72 |
| GSK3-alpha-beta | -0.0489 | -0.2743 | 0.1816 | 0.6791 | -0.4153 | 72 |
| GSK3-alpha-beta_pS21_S9 | -0.0018 | -0.2302 | 0.2268 | 0.9876 | -0.0156 | 72 |
| GSK3_pS9 | 0.0912 | -0.1402 | 0.3132 | 0.4397 | 0.7771 | 72 |
| HER2 | 0.0602 | -0.1706 | 0.2848 | 0.6102 | 0.5121 | 72 |
| HER2_pY1248 | 0.0261 | -0.2037 | 0.253 | 0.8256 | 0.2212 | 72 |
| HER3 | 0.0759 | -0.1553 | 0.2992 | 0.5204 | 0.6459 | 72 |
| HER3_pY1289 | 0.0299 | -0.1999 | 0.2567 | 0.8002 | 0.254 | 72 |
| Heregulin | 0.1615 | -0.0695 | 0.3761 | 0.1692 | 1.3888 | 72 |
| HSP70 | 0.1496 | -0.0817 | 0.3656 | 0.2033 | 1.284 | 72 |
| IGFBP2 | -0.1047 | -0.3254 | 0.1268 | 0.3746 | -0.8935 | 72 |
| INPP4B | 0.0343 | -0.1957 | 0.2607 | 0.7718 | 0.2911 | 72 |
| IRS1 | 0.1536 | -0.0776 | 0.3691 | 0.1914 | 1.3188 | 72 |
| JNK_pT183_pY185 | -0.0196 | -0.247 | 0.2098 | 0.868 | -0.1668 | 72 |
| JNK2 | 8.00E-04 | -0.2277 | 0.2293 | 0.9944 | 0.007 | 72 |
| Ku80 | 0.0887 | -0.1427 | 0.3109 | 0.4522 | 0.7559 | 72 |
| Lck | 0.0926 | -0.1388 | 0.3145 | 0.4326 | 0.7892 | 72 |
| LKB1 | -0.019 | -0.2464 | 0.2105 | 0.8726 | -0.1609 | 72 |
| MAPK_pT202_Y204 | 0.047 | -0.1835 | 0.2726 | 0.6908 | 0.3994 | 72 |
| MEK1 | -0.0286 | -0.2554 | 0.2012 | 0.8088 | -0.2429 | 72 |
| MEK1_pS217_S221 | 0.0043 | -0.2244 | 0.2326 | 0.9711 | 0.0364 | 72 |
| MIG-6 | -2.00E-04 | -0.2287 | 0.2283 | 0.9984 | -0.002 | 72 |
| Mre11 | 0.048 | -0.1825 | 0.2735 | 0.6844 | 0.4081 | 72 |
| MSH2 | 0.1119 | -0.1197 | 0.3319 | 0.3426 | 0.9553 | 72 |
| MSH6 | 0.1509 | -0.0803 | 0.3668 | 0.1992 | 1.2956 | 72 |
| mTOR | 0.0309 | -0.199 | 0.2576 | 0.794 | 0.2621 | 72 |
| mTOR_pS2448 | 0.0625 | -0.1684 | 0.2869 | 0.5968 | 0.5313 | 72 |
| MYH11 | 0.0183 | -0.2111 | 0.2457 | 0.8772 | 0.155 | 72 |
| Myosin-IIa-pS1943 | -0.0107 | -0.2387 | 0.2183 | 0.9276 | -0.0911 | 72 |
| N-Cadherin | 0.1432 | -0.0882 | 0.3599 | 0.2235 | 1.2278 | 72 |
| N-Ras | 0.1115 | -0.1201 | 0.3315 | 0.3443 | 0.952 | 72 |
| NDRG1_pT346 | 0.0382 | -0.192 | 0.2644 | 0.7468 | 0.3241 | 72 |
| NF-kB-p65_pS536 | 0.1331 | -0.0983 | 0.351 | 0.2581 | 1.1399 | 72 |
| NF2 | 0.0088 | -0.2201 | 0.2368 | 0.9407 | 0.0747 | 72 |
| Notch1 | 0.0599 | -0.1709 | 0.2845 | 0.612 | 0.5094 | 72 |
| P-Cadherin | 0.12 | -0.1116 | 0.3392 | 0.3085 | 1.0256 | 72 |
| p21 | 0.088 | -0.1434 | 0.3102 | 0.4561 | 0.7493 | 72 |
| p27 | -0.0203 | -0.2477 | 0.2092 | 0.8637 | -0.1723 | 72 |
| p27_pT157 | -0.0291 | -0.2559 | 0.2007 | 0.8055 | -0.2472 | 72 |
| p27_pT198 | -0.0561 | -0.281 | 0.1747 | 0.6352 | -0.4765 | 72 |
| p38 | 0.0706 | -0.1605 | 0.2943 | 0.5502 | 0.6003 | 72 |
| p38_pT180_Y182 | -0.0437 | -0.2696 | 0.1866 | 0.7113 | -0.3716 | 72 |

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|---|---|---|---|---|---|---|
| p53 | -0.0641 | -0.2884 | 0.1669 | 0.5876 | -0.5448 | 72 |
| p62-LCK-ligand | 0.1677 | -0.0632 | 0.3816 | 0.1532 | 1.4434 | 72 |
| p70S6K | 0.0966 | -0.1348 | 0.3181 | 0.4128 | 0.8238 | 72 |
| p70S6K_pT389 | 0.2292 | 7.00E-04 | 0.4349 | 0.0495 | 1.998 | 72 |
| p90RSK | 0.0554 | -0.1753 | 0.2803 | 0.6394 | 0.4706 | 72 |
| p90RSK_pT359_S363 | 0.0765 | -0.1547 | 0.2998 | 0.517 | 0.6513 | 72 |
| PAI-1 | 0.1024 | -0.1291 | 0.3233 | 0.3853 | 0.8734 | 72 |
| Paxillin | 0.1432 | -0.0882 | 0.3599 | 0.2235 | 1.2278 | 72 |
| PCNA | -0.0594 | -0.2841 | 0.1714 | 0.6151 | -0.505 | 72 |
| PDCD4 | -0.1865 | -0.398 | 0.0439 | 0.1116 | -1.6106 | 72 |
| PDK1 | 0.0211 | -0.2084 | 0.2484 | 0.8586 | 0.1788 | 72 |
| PDK1_pS241 | -0.0512 | -0.2765 | 0.1794 | 0.6646 | -0.4353 | 72 |
| PEA15 | -0.133 | -0.3508 | 0.0985 | 0.2587 | -1.1385 | 72 |
| PEA15_pS116 | 0.1255 | -0.106 | 0.3441 | 0.2867 | 1.0734 | 72 |
| PI3K-p110-alpha | -0.0105 | -0.2384 | 0.2185 | 0.9292 | -0.0891 | 72 |
| PI3K-p85 | 0.0506 | -0.18 | 0.2759 | 0.6685 | 0.43 | 72 |
| PKC-alpha | -0.1195 | -0.3388 | 0.1121 | 0.3105 | -1.0213 | 72 |
| PKC-alpha_pS657 | -0.0436 | -0.2694 | 0.1868 | 0.7123 | -0.3702 | 72 |
| PKC-delta_pS664 | 0.0077 | -0.2212 | 0.2358 | 0.948 | 0.0654 | 72 |
| PKC-pan_BetaII_pS660 | 0.0062 | -0.2226 | 0.2343 | 0.9583 | 0.0524 | 72 |
| PR | -0.0534 | -0.2785 | 0.1772 | 0.6512 | -0.454 | 72 |
| PRAS40_pT246 | 0.0379 | -0.1923 | 0.2641 | 0.7486 | 0.3217 | 72 |
| PRDX1 | -0.1545 | -0.37 | 0.0767 | 0.1886 | -1.3273 | 72 |
| PREX1 | -0.0277 | -0.2546 | 0.2021 | 0.815 | -0.2349 | 72 |
| PTEN | 0.0412 | -0.189 | 0.2672 | 0.7272 | 0.3502 | 72 |
| Rab11 | 0.1243 | -0.1073 | 0.343 | 0.2915 | 1.0627 | 72 |
| Rab25 | -0.1367 | -0.3541 | 0.0947 | 0.2455 | -1.171 | 72 |
| Rad50 | 0.0385 | -0.1917 | 0.2646 | 0.7448 | 0.3267 | 72 |
| Rad51 | 0.0999 | -0.1316 | 0.3211 | 0.3971 | 0.8519 | 72 |
| Raptor | -0.0169 | -0.2444 | 0.2124 | 0.8865 | -0.1433 | 72 |
| Rb-M-QC | -0.1813 | -0.3935 | 0.0492 | 0.1221 | -1.5643 | 72 |
| Rb_pS807_S811 | -0.0207 | -0.248 | 0.2088 | 0.861 | -0.1758 | 72 |
| RBM15 | 0.102 | -0.1295 | 0.323 | 0.3872 | 0.87 | 72 |
| Rictor | 0.1098 | -0.1218 | 0.33 | 0.3518 | 0.9372 | 72 |
| Rictor_pT1135 | 0.0851 | -0.1462 | 0.3076 | 0.4709 | 0.7248 | 72 |
| S6 | -0.064 | -0.2883 | 0.167 | 0.5882 | -0.5439 | 72 |
| S6_pS235_S236 | 7.00E-04 | -0.2279 | 0.2291 | 0.9956 | 0.0056 | 72 |
| S6_pS240_S244 | 0.0354 | -0.1947 | 0.2617 | 0.7649 | 0.3002 | 72 |
| SCD1 | -0.209 | -0.4176 | 0.0204 | 0.0739 | -1.8137 | 72 |
| SF2 | -0.0265 | -0.2534 | 0.2032 | 0.8228 | -0.2248 | 72 |
| Shc_pY317 | -0.0874 | -0.3097 | 0.1439 | 0.4589 | -0.7447 | 72 |
| Smad1 | 0.0645 | -0.1664 | 0.2888 | 0.5848 | 0.5488 | 72 |
| Smad3 | 0.1242 | -0.1074 | 0.3429 | 0.2919 | 1.0619 | 72 |
| Smad4 | 0.042 | -0.1883 | 0.2679 | 0.7226 | 0.3564 | 72 |

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|---|---|---|---|---|---|---|
| Src | -0.1168 | -0.3363 | 0.1148 | 0.3219 | -0.9975 | 72 |
| Src_pY416 | -0.0894 | -0.3115 | 0.142 | 0.4487 | -0.7617 | 72 |
| Src_pY527 | -0.0362 | -0.2625 | 0.1939 | 0.7595 | -0.3073 | 72 |
| STAT3_pY705 | 0.0584 | -0.1724 | 0.2832 | 0.6209 | 0.4967 | 72 |
| STAT5-alpha | 0.1285 | -0.103 | 0.3468 | 0.2752 | 1.0996 | 72 |
| Stathmin | 0.037 | -0.1931 | 0.2633 | 0.7544 | 0.314 | 72 |
| Syk | -0.0153 | -0.2429 | 0.214 | 0.8973 | -0.1295 | 72 |
| TAZ | 0.1077 | -0.1239 | 0.3281 | 0.3612 | 0.9189 | 72 |
| TIGAR | -9.00E-04 | -0.2293 | 0.2277 | 0.9942 | -0.0073 | 72 |
| Transglutaminase | 0.126 | -0.1056 | 0.3446 | 0.2849 | 1.0775 | 72 |
| TFRC | -0.0169 | -0.2445 | 0.2124 | 0.886 | -0.1438 | 72 |
| TSC1 | 0.0496 | -0.181 | 0.275 | 0.6748 | 0.4213 | 72 |
| TTF1 | 0.1042 | -0.1274 | 0.3249 | 0.3772 | 0.8886 | 72 |
| Tuberin | 0.1071 | -0.1244 | 0.3276 | 0.3635 | 0.9145 | 72 |
| Tuberin_pT1462 | -0.0174 | -0.2449 | 0.212 | 0.8833 | -0.1473 | 72 |
| VEGFR2 | 0.0618 | -0.1691 | 0.2863 | 0.6008 | 0.5256 | 72 |
| VHL | 0.0606 | -0.1703 | 0.2851 | 0.6082 | 0.5149 | 72 |
| XBP1 | 0.0905 | -0.1409 | 0.3125 | 0.4433 | 0.7709 | 72 |
| XRCC1 | -0.1005 | -0.3216 | 0.131 | 0.3943 | -0.8571 | 72 |
| YAP | 1.00E-04 | -0.2284 | 0.2286 | 0.9994 | 8.00E-04 | 72 |
| YAP_pS127 | -0.0602 | -0.2848 | 0.1707 | 0.6106 | -0.5115 | 72 |
| YB-1 | -0.1665 | -0.3806 | 0.0644 | 0.1561 | -1.4331 | 72 |
| YB-1_pS102 | -0.0415 | -0.2675 | 0.1888 | 0.7256 | -0.3524 | 72 |

CC, correlation coefficient; CI, confidence interval; df, degree of freedom.