

## CD36 promotes chemotherapy resistance and stemness of cervical cancer cells through Hippo and ERK5 pathway activation

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| Abstract:                     | <p>Our previous work revealed that CD36 accelerates progression of cervical cancer. Here, we elucidate the effects of CD36 in regulating stemness and chemotherapy resistance of cervical cancer cells. We used bioinformatics analysis to assess the correlation between CD36 expression and chemotherapy drugs sensitivity in advanced cervical cancer. The effects of CD36 expression on proliferation of cervical cancer cells were analyzed using the CCK8 and colony formation assay. The stemness of cervical cancer cells were assessed by tumorsphere formation assay. Correlations between CD36 expression and stem cell markers were investigated by Western blot. The roles of CD36 on chemotherapy resistance of cervical cancer cells were evaluated by using colony formation assay, CCK8 assay and flow cytometry. Finally, the molecular mechanisms by which CD36 promotes stemness and chemotherapy resistance of cervical cancer cells were explored through bioinformatics analysis and Western blot. CD36 expression was positively correlated with cisplatin resistance in advanced cervical cancer. CD36 overexpression promoted the stemness and chemotherapy resistance of cervical cancer cells. Most importantly, we discovered that CD44, Hippo and MEK5/ERK5 pathways might be involved in CD36-mediated chemotherapy resistance and stemness of cervical cancer cells. CD36 may be an effective therapeutic target for improving the prognosis of cervical cancer.</p> |

**CD36 promotes chemotherapy resistance and stemness of cervical cancer cells  
through Hippo and ERK5 pathway activation**

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## Abstract

**Objective:** Our previous work revealed that CD36 accelerates progression of cervical cancer. Here, we elucidate the effects of CD36 in regulating stemness and chemotherapy resistance of cervical cancer cells.

**Methods:** We used bioinformatics analysis to assess the correlation between CD36 expression and chemotherapy drugs sensitivity in advanced cervical cancer. The effects of CD36 expression on proliferation of cervical cancer cells were analyzed using the CCK8 and colony formation assay. The stemness of cervical cancer cells were assessed by tumorsphere formation assay. Correlations between CD36 expression and stem cell markers were investigated by Western blot. Next, the roles of CD36 on chemotherapy resistance of cervical cancer cells were evaluated by using colony formation assay, CCK8 assay and flow cytometry. Finally, the molecular mechanisms by which CD36 promotes stemness and chemotherapy resistance of cervical cancer cells were explored through bioinformatics analysis and Western blot.

**Results:** CD36 expression was positively correlated with cisplatin resistance in advanced cervical cancer. CD36 overexpression promoted the stemness and chemotherapy resistance of cervical cancer cells. Most importantly, we discovered that CD44, Hippo and MEK5/ERK5 pathways might be involved in CD36-mediated chemotherapy resistance and stemness of cervical cancer cells.

**Conclusions:** CD36 may be an effective therapeutic target for improving the prognosis of cervical cancer.

**Keywords:** CD36; Stemness; Chemotherapy resistance; Cervical cancer

**Introduction**

Cervical cancer is the most common malignancy of the female genital tract [1]. Its progression involves infection with human papillomavirus, cervical intraepithelial neoplasia, local invasion, and metastasis [2-3]. Widespread use of cytological screening and biopsies has effectively reduced cervical cancer incidence. However, advanced cervical cancer continues to have a poor prognosis due to chemotherapy resistance [4]. Therefore, elucidating the mechanisms underlying chemotherapy resistance in cervical cancer is essential.

Cancer cells utilize fatty acids as an energy source to reprogram metabolism. CD36, a lipid transporter, is a membrane protein expressed on monocytes, epithelial cells, and endothelial cells [5-8]. In cancer cells, CD36 overexpression accelerates fatty acid uptake and lipid accumulation, promoting growth and metastasis [9-12]. Lipid accumulation also exerts immunosuppressive effects [13]. Owing to its multifaceted roles in cancer biology, CD36 has recently emerged as a promising therapeutic target in various cancers [14-18]. Notably, silencing CD36 inhibits clonogenicity and stemness marker expression in bladder cancer cells, highlighting its critical role in enhancing stemness properties [19]. However, current understanding of CD36's role in regulating stemness and chemotherapy resistance in cervical cancer cells remains limited.

Cancer stem cells (CSCs) are tumor-initiating cells that are capable of differentiating into multiple cell types and contributing to tumorigenicity and chemotherapy resistance in solid malignancies [20-22]. However, limited studies have examined the

relationship between cervical CSCs, chemotherapy resistance, and cancer progression. CD44, a cervical tumorigenesis stem cell marker, can enhance CSC self-renewal and chemotherapy resistance [23]. Yes-associated protein (YAP) promotes malignant transformation of cervical intraepithelial neoplasia by reinforcing CSC traits [24]. In this study, we explored the roles of CD36 in regulating stemness and chemotherapy resistance in cervical cancer cells. Further, we investigated the specific mechanisms through which CD36 facilitates these processes.

## Materials and methods

### Acquisition of Cervical Cancer Transcriptome Data

We downloaded transcriptome sequencing data (ribonucleic acid [RNA] sequencing) for cervical cancer from The Cancer Genome Atlas database (<https://www.cancer.gov/ccg/research/genome-sequencing/tcga>) using the Illumina HiSeq 2000 platform. The dataset included 62 advanced cervical cancer samples (41 stage 4 and 21 additional stage 4 cases). Details of the downloaded data are provided in Table S1.

### Half-maximal inhibitory concentration (IC<sub>50</sub>) correlation analysis

CD36 expression levels in cervical cancer specimens from The Cancer Genome Atlas database were analyzed in combination with data from the Genomics of Drug Sensitivity in Cancer database (<https://www.cancerrxgene.org/>) [25]. The R version 4.3.1 pRRophetic package (<https://github.com/paulgeeleher/pRRophetic>) was used to evaluate the sensitivity of advanced cervical cancer to various chemotherapeutic agents [26]. Using the downloaded drug data as a reference, the pRRophetic package

estimated the IC50 value of each sample based on whole-genome expression.

**Kyoto Encyclopedia of Genes and Genomes pathways analysis**

The correlation between CD36 gene expression and all of the other genes was calculated in 62 advanced cervical cancer samples using the Pearson correlation coefficient. Genes with correlation coefficients above 0.3 and P-values below 0.05 were retained. The Database for Annotation, Visualization and Integrated Discovery (DAVID) version 6.8 (<https://david.ncifcrf.gov/>) was then used to perform Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway enrichment analysis on the selected genes, applying a *P*-value threshold of less than 0.05 [27]. There were 10 KEGG pathways that met the significance criteria.

**Prognostic analysis**

For prognostic analysis, the R version 4.3.1 survival package (version 2.41-1; <https://cran.r-project.org/web/packages/finalfit/vignettes/survival.html>) was used to evaluate the correlation between CD36 expression and prognosis in advanced cervical cancer [28]. The effects of CD36 expression and clinical stage on patient prognosis were analyzed.

**Cell culture**

C33a and HeLa cells were obtained from the Type Collection Center (Wuhan, China) and cultured in Dulbecco's Modified Eagle Medium (Gibco, CA, USA). C33a cells were incubated with fluorescein isothiocyanate-labeled CD133 and sorted by flow cytometry. The CD133-positive C33a cell population was collected, expanded, and cultured *in vitro* as C33a-Cervical CSCs (C33a-CCSCs).

### Plasmid construction and transfection

The pIRES2-ZsGreen1-CD36 plasmid, designed to overexpress CD36, was constructed by cloning the CD36 coding sequence into the pIRES2-ZsGreen1 vector. C33a-CCSCs were transfected with the plasmid using Lipo<sup>TM</sup>6000 (Beyotime, Shanghai, China) and selected in medium containing 600 µg/ml G418 for three weeks. The empty pIRES2-ZsGreen1-NC plasmid served as a negative control (NC). CD36-overexpressing C33a and C33a-CCSCs cells were designated as “C33a/CD36” and “C33a-CCSCs/CD36,” respectively.

### Transfection with siRNA (small interfering RNA)

Human CD36 small interfering ribonucleic acid (siRNA) was purchased from Guangzhou Ribo (Guangzhou, China). Cells were seeded in six-well plates at a density of  $2 \times 10^5$  cells/well. HeLa cells were transfected with CD36 siRNA (HeLa/siRNA CD36) (sense: 5'-CCUGAUAGAAAUGAUCUUATT-3'; antisense: 5'-UAAGAUCAUUUCUAUCA GGTT-3') or NC siRNA (HeLa/NC-siRNA) (sense: 5'-UUCUCCGAACGUGUCACGUTT-3'; antisense: 5'-ACGUGACACGUUCGGAG AATT-3') using Lipo<sup>TM</sup>3000 (Thermo Fisher Scientific, Waltham, USA) at a final concentration of 100 nM.

### Flow cytometry analysis of the ratios of CD133 positive cervical cancer cells

C33a/CD36 and HeLa/CD36-siRNA cells were centrifuged at 1500 rpm for 5 min. The cells were resuspended in 200 µl of phosphate-buffered saline, followed by the addition of 5 µl of CD133 antibody to each flow tube. Samples were incubated at 4°C

for 30 min. After adding another 200 µl of phosphate-buffered saline, the proportion of CD133-positive cells was analyzed by flow cytometry.

**Tumorsphere formation assay**

C33a/CD36, C33a-CCSCs/CD36, or HeLa/siRNA CD36 cells were incubated in Dulbecco’s Modified Eagle Medium supplemented with 20 ng/ml basic fibroblast growth factor, 20 ng/ml epidermal growth factor, and 1×B27. Cells were seeded in low-attachment six-well plates at a density of  $2 \times 10^4$  cells/well and cultured for 10 days. Three fields were randomly selected from confocal images, and the number of stem cell spheres with diameters greater than 100 µm was counted for each group.

**Western blot**

After boiling for 8 min, total protein (80 µg) was loaded onto a 10% polyacrylamide gel and transferred to a polyvinylidene fluoride membrane. The membrane was incubated with the following primary antibodies: CD36 (1:1000; Ab252923, Abcam, Cambridge, UK), CD44 (1:3000; A19020, Abclonal, MA, USA), CD133 (1:1000; AF5120, Affinity, Cincinnati, USA), Nanog (1:5000; 67255-1-Ig, Sanying, Wuhan, China), YAP (1:1000; AF6328, Affinity, Cincinnati, USA), TAZ (1:300; A8202, Abclonal, MA, USA), Bcl-2 (1:3000; 26593-1-AP, Sanying, Wuhan, China), Bax (1:10000; 60627-1-Ig, Sanying, Wuhan, China), MEK5 (1:2000; DF7667, Affinity, Cincinnati, USA), ERK5 (1:1000; DF6835, Affinity, Cincinnati, USA), cleaved caspase-3 (1:1000; AF7022, Affinity, Cincinnati, USA), and GAPDH (1:1000; AB-P-R001, Xianzhi Biology, Hangzhou, China). Membranes were then incubated with horseradish peroxidase-conjugated secondary antibodies (1:10000; BA1051,

BA1054; Boster Biology, Wuhan, China). Proteins were visualized using a chemiluminescence kit (Pierce; Thermo Fisher Scientific, Inc., Massachusetts, USA).

### Colony formation assay

HeLa/siRNA CD36, HeLa/NC siRNA, and HeLa cells were seeded at 200 cells/well in six-well plates [29]. The cells were incubated at 5% CO<sub>2</sub>. After 14 days, colonies were stained with trypan blue, and the number of cell colonies was counted.

### CCK8 assay

C33a/CD36, C33a-CCSCs, HeLa/siRNA CD36 cells and the corresponding control groups were inserted into 96-well plates with  $5 \times 10^3$  cells per well and cultured 24h at 37°C. Add 10µl cck8 per well and culture at 37°C for 2h. The absorbance value OD450 of each well was determined by microplate reader. Cell survival rate % =  $\frac{[(\text{experimental hole absorbance}) - (\text{blank hole absorbance})]}{[(\text{control hole absorbance}) - (\text{blank hole absorbance})]} \times 100\%$

### TUNEL assay

C33a-CCSCs, C33a-CCSCs/CD36 and C33a-CCSCs/NC were seeded at a density of  $4 \times 10^5$  cells/well in six-well plates. After added 0.5 ml fixative, the cells were permeabilized by 0.2% Triton X-100. Removed the excess PBS, 100 µl of TUNEL equilibrium solution was added to each sample. A TUNEL reaction mixture containing TdT enzyme and fluorescein-dUTP was applied and incubated at 37 °C for 2h. Then diluted DAPI was added, and the samples were further incubated at room temperature for 5 min. The samples were observed under a fluorescence microscope.

### Immunofluorescence

After the tissue sections were deparaffinized, antigen repair was performed. After blocking non-specific binding with 5% diluted goat serum, sections were incubated with primary antibodies against CD44 (dilution, 1:100; A19020, Abclonal, MA, USA) and CD133(dilution, 1:100; AF5120, Affinity, Cincinnati, USA) at 4°C overnight, then incubated with secondary antibody (dilution, 1:100; BA1105, BOSTER, Wuhan, China). Nuclei were stained by 4',6- Diamidino-2-phenylindole (DAPI). After using anti-fluorescence quencher, the images were observed under a fluorescence microscope. Image-Pro Plus 6.0 software (Media Cybernetics, Maryland, USA) was used to quantitatively analyze the expressions of CD133 and CD44 in each group. The mean optical density value represents the protein expression level.

**Flow Cytometry analysis of apoptosis**

Cells were centrifuged at 1500rpm for 5 min. The ratios of apoptotic in cervical cancer cells were measured using Annexin V FITC/PI Apoptosis Detection Kit.

**Animal assay**

Four-week-old female BALB/C nude mice were purchased from SiBei Fu Animal Center (Beijing, China) . C33a/CD36 cells (2×10<sup>6</sup>/mL) were injected subcutaneously into the axillary of right forelimb of nude mice. Each group had six nude mice. In vivo imaging detection was performed with a small animal in vivo imager (IVIS Lumina III, Perkin Elmer). We euthanized the nude mice with anesthesia. The animal experiment protocol was carried out in accordance with the regulations of the Institutional Animal Ethical Care Committee of the Eighth Affiliated Hospital, Sun Yat-sen University (2024r005) .

## Statistical analysis

Data are presented as mean  $\pm$  standard error of the mean (SEM). The overall survival rate was calculated by plotting the survival curve with the Kaplan-Meier method. Statistical comparisons of data from the experiments on cultured cells or mice were performed using the two-tailed Student's *t* test. All of the statistical analyses were performed using Statistical Package for the Social Sciences version 17.0 (SPSS Inc., Chicago, USA). Differences were considered significant when the *p* value was less than 0.05.

## Results

### Correlation between CD36 expression level and resistance of chemotherapy drugs in advanced cervical cancer.

The IC<sub>50</sub> value reflects the sensitivity of chemotherapeutic drugs, and its correlation with CD36 expression was evaluated through bioinformatics analysis (**Table S2**). Notably, higher CD36 expression in advanced cervical cancer was associated with increased IC<sub>50</sub> values for cisplatin. Additionally, CD36 expression showed a positive correlation with cisplatin resistance and a negative correlation with vinorelbine resistance in advanced cervical cancer (**Figure 1A**).

### Analysis of downstream pathways regulated by CD36 in advanced cervical cancer

In 62 advanced cervical cancer samples, 648 genes were identified as significantly associated with CD36 expression (**Table S3**). KEGG pathway enrichment analysis

was conducted using DAVID version 6.8. There were 10 KEGG pathways that were identified based on a P-value threshold of  $< 0.05$  (**Figure 1B**). Among these, the Hippo and MEK5/ERK5 signaling pathways were associated with chemotherapy resistance and cancer cell stemness. However, no correlation was observed between CD36 expression and prognosis in patients with advanced cervical cancer (**Figure 1C–F**). CD36 expression was positively correlated with the expression of CD44, CD133, YAP, TAZ, MEK5, and ERK5 in cervical cancer cells (**Figure 1G–J**). These results suggest that CD36 may enhance stem cell marker expression and activate the Hippo and MEK5/ERK5 pathways *in vitro*.

**CD36 promotes stemness and tumor growth in vivo**

After six weeks, CD36 overexpression significantly promoted tumor growth in female BALB/c-nu mice compared with the C33a/Con group ( $1636.12 \pm 462.29 \text{ mm}^3$  versus  $732.64 \pm 329.89 \text{ mm}^3$ ; **Figure 2A–C**). Additionally, immunofluorescence assays showed that CD133 (mean optical density:  $0.81 \pm 0.04$  vs.  $0.31 \pm 0.01$ ; **Figure 2D**) and CD44 expression ( $0.95 \pm 0.05$  vs.  $0.48 \pm 0.03$ ; **Figure 2E**) were increased in the C33a/CD36 group compared with controls. These findings suggest that CD36 may promote cervical cancer cell growth by enhancing stemness *in vivo*.

**CD36 promoted the stemness of cervical cancer cells in vitro**

As shown in **Figure 3A–B**, the number of stem cell spheres with diameters greater than  $100 \text{ }\mu\text{m}$  was higher in the C33a/CD36 group ( $5.33 \pm 0.58$ ) than in the C33a/NC group ( $1.67 \pm 0.58$ ) and C33a group ( $2.33 \pm 0.57$ ). The Cell Counting Kit-8 assay revealed that CD36 overexpression significantly enhanced the proliferative capacity

of C33a cells (**Figure 3D**). In contrast, the number of spheres larger than 100  $\mu\text{m}$  decreased in the HeLa/siRNA CD36 group ( $2.33 \pm 0.58$ ) compared with the HeLa/NC-siRNA group ( $5.33 \pm 0.57$ ) and HeLa group ( $4.34 \pm 0.54$ ) (**Figure 3A, C**). Proliferation was also reduced in the HeLa/siRNA CD36 group relative to the HeLa/NC-siRNA and HeLa groups (**Figure 3E**). Flow cytometry analysis further demonstrated a higher percentage of CD133-positive cells in the C33a/CD36 group ( $8.35 \pm 0.30\%$ ) than in the C33a/NC group ( $3.45 \pm 0.21\%$ ) and C33a group ( $3.33 \pm 0.37\%$ ) (Figure 3F, H). Conversely, fewer CD133-positive cells were observed in the HeLa/siRNA CD36 group ( $2.44 \pm 0.33\%$ ) compared with the HeLa/NC-siRNA ( $5.07 \pm 0.11\%$ ) and HeLa groups ( $5.06 \pm 0.52\%$ ) (**Figure 3G and I**). These findings suggest that CD36 plays a critical role in promoting the stemness of cervical cancer cells *in vitro*.

### CD36 overexpression augmented stemness of cervical CSCs

We further sorted CD133-positive C33a cells using flow cytometry and established a CD36-overexpressing cervical cancer stem cell model (C33a-CCSCs/CD36) along with a corresponding control group (C33a-CCSCs/NC). Western blot analysis confirmed that CD36 expression was successfully upregulated in the C33a-CCSCs/CD36 group (**Figure 4A and D**). The number of stem cell spheres with diameters exceeding 100  $\mu\text{m}$  was significantly higher in the C33a-CCSCs/CD36 group ( $7.33 \pm 0.58$ ) than in the C33a-CCSCs/NC group ( $3.67 \pm 0.58$ ) and the C33a-CCSCs group ( $3.33 \pm 0.57$ ) (**Figure 4B**). Similarly, the proliferative capacity of the C33a-CCSCs/CD36 group was markedly elevated compared with both control

groups (**Figure 4C**). Immunofluorescence assays also showed increased CD44 expression in the C33a-CCSCs/CD36 group relative to the C33a-CCSCs/NC group (**Figure 4E**). These findings suggest that CD36 overexpression may enhance the stemness of cervical CSCs.

**Elevating CD36 expression potentiated the chemotherapy resistance of cervical CSCs through Hippo and MEK5/ERK5 pathways activation.**

To further assess whether CD36 expression influences chemotherapy resistance in cervical CSCs, C33a-CCSCs/CD36, C33a-CCSCs/NC, and C33a-CCSCs were treated with 20 µg/ml cisplatin for 24 h. As depicted in **Figure 5A**, the C33a-CCSCs/CD36 group exhibited higher proliferative capacity ( $130.98 \pm 3.15\%$ ) compared with the C33a-CCSCs/NC group ( $97.89 \pm 1.28\%$ ) and the C33a-CCSCs group ( $99.57 \pm 0.54\%$ ). Additionally, the C33a-CCSCs/CD36 group displayed the fewest apoptotic cells relative to the control groups (**Figure 5F**). The expression levels of YAP, TAZ, MEK5, and ERK5 were also elevated in the C33a-CCSCs/CD36 group compared with the controls (**Figure 5D-E**).

To further investigate whether the Hippo and MEK5/ERK5 pathways contribute to CD36-regulated chemotherapy resistance in cervical CSCs, C33a-CCSCs/CD36 cells were treated with Verteporfin (a YAP inhibitor) and XMD8-92 (an ERK5 inhibitor). As shown in **Figure 5B-C**, proliferative capacity decreased following treatment with Verteporfin ( $117.13 \pm 1.78\%$  vs.  $99.14 \pm 0.84\%$ ;  $99.18 \pm 0.47\%$ ) or XMD8-92 ( $120.99 \pm 2.32\%$  vs.  $99.37 \pm 0.74\%$ ;  $98.43 \pm 0.92\%$ ) compared with the C33a-CCSCs + cisplatin and C33a-CCSCs/NC+cisplatin groups. Additionally, the number of

apoptotic cells in the C33a-CCSCs/CD36 group increased relative to controls after treatment with either inhibitor (**Figure 5G, H**). These findings indicate that CD36 may induce chemotherapy resistance in cervical CSCs via activation of the Hippo and MEK5/ERK5 pathways.

### **Interfering CD36 expression alleviated the chemotherapy resistance of cervical cancer cells**

Conversely, we downregulated CD36 expression in HeLa cells and treated them with 20 µg/ml cisplatin for 24 hours. The colony formation assay revealed fewer colonies in the HeLa/siRNA CD36 group ( $106.67 \pm 14.57$ ) compared with the HeLa/NC siRNA group ( $161.33 \pm 8.08$ ) and HeLa group ( $148.33 \pm 10.02$ ) (**Figure 6A**). The proliferative capacity of the HeLa/siRNA CD36 group ( $75.10 \pm 3.12\%$ ) was significantly reduced relative to the HeLa group ( $94.15 \pm 1.18\%$ ) and HeLa/NC-siRNA group ( $97.48 \pm 2.57\%$ ) (**Figure 6B**). The HeLa/siRNA CD36 group also exhibited the highest number of apoptotic cells ( $35.28 \pm 1.15\%$  vs.  $21.27 \pm 0.88\%$ ;  $21.26 \pm 0.67\%$ ) (**Figure 6C, E**). Furthermore, Bcl-2 expression was decreased, Bax expression increased, and cleaved caspase-3 activity enhanced in the HeLa/siRNA CD36 group, consistent with CD36 downregulation (**Figures 6D, F**). These findings indicate that silencing CD36 inhibits proliferation and promotes apoptosis in HeLa cells following cisplatin treatment. Additionally, the expression of CD44, CD133, and Nanog was significantly reduced in the HeLa/siRNA CD36 group compared with controls (**Figure 6D, F**), suggesting that CD36 knockdown may alleviate chemotherapy resistance by suppressing cervical cancer cell stemness.

**Discussion**

Previous studies have shown that CD36 promotes the progression and metastasis of oral cancer by facilitating the uptake of exogenous fatty acids [11]. Our prior work also demonstrated that CD36 accelerates epithelial-mesenchymal transition and metastasis in cervical cancer [30]. Additionally, CD36-driven lipid uptake contributes to chemotherapy resistance. In acute myeloid leukemia, the CD36–fatty acid oxidation (FAO)–oxidative phosphorylation (OXPHOS) axis underlies resistance to chemotherapy [31]. In this study, we investigated the roles of CD36 in regulating stemness and chemotherapy resistance in cervical cancer cells.

In our study, we found that CD36 promoted the formation of larger stem cell spheres and enhanced the expression of stem cell markers. Additionally, both the proliferative capacity and sphere-forming ability increased following CD36 transfection in C33a-CCSCs. Similarly, Gyamfi et al. reported that CD36 knockdown reduced stemness marker expression in breast cancer cells [32]. Collectively, our results demonstrate that CD36 promotes the stemness of cervical cancer cells.

Chemotherapy resistance is a major factor limiting the clinical efficacy of cervical cancer treatment. Bioinformatics analysis revealed that CD36 expression was positively correlated with cisplatin resistance in advanced cervical cancer. Therefore, we investigated the role of CD36 in chemotherapy resistance. CD36 overexpression enhanced proliferation and reduced apoptosis in C33a-CCSCs treated with cisplatin. In contrast, CD36 silencing decreased colony formation and proliferation in HeLa cells while increasing apoptosis. These findings suggest that CD36 plays a key role in

promoting chemotherapy resistance in cervical cancer cells.

CD44 is a well-established cancer stem cell marker and a key regulator of epithelial–mesenchymal transition [33]. We found that CD36 enhances CD44 expression both *in vitro* and *in vivo*, thereby promoting the stemness of cervical cancer cells. The ERK5 and Hippo pathways are also recognized as important contributors to chemotherapy resistance and stemness in cancer cells [34–37]. YAP, a central effector of the Hippo pathway, promotes immune evasion in BRAF inhibitor-resistant melanoma by upregulating PD-L1 [38], while the MEK5–ERK5 –STAT3 axis accelerates self-renewal and tumorigenicity in glioma stem cells [39]. In our study, CD36 upregulated the expression of YAP, TAZ, MEK5, and ERK5, thereby activating both the Hippo and ERK5 pathways. Vittoria et al. reported that Hippo pathway activation is mediated through mitogen-activated protein kinase (MAPK) signaling [40], and CD36 has been shown to activate MAPK signaling [41]. Thus, we speculate that CD36 may activate the Hippo pathway via MAPK activation. Moreover, treatment with YAP and ERK5 inhibitors reduced proliferation and increased apoptosis in C33a-CCSCs/CD36 cells, further supporting the involvement of the Hippo and ERK5 pathways in CD36-mediated chemotherapy resistance.

## Conclusion

In summary, our findings demonstrate that CD36 promotes stemness and chemotherapy resistance in cervical cancer cells. Notably, CD44, Hippo, and MEK5/ERK5 pathways are involved in CD36-mediated resistance and contribute to enhanced stemness. Collectively, these results suggest that CD36 may serve as a

promising therapeutic target for improving the prognosis of cervical cancer.

**Declaration of conflicting interests**

The authors declare that they have no conflicts of interest.

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**Supplemental material**

The supplemental material for this article is available online.

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### Figure Legend

Figure 1. Correlation between CD36 expression and resistance of chemotherapy drugs and prognosis of cervical cancer and its regulated signaling pathways. A: CD36 expression had the most significant positive correlation with cisplatin resistance and the most significant negative correlation with vinorelbine in advanced cervical cancer by IC50 correlation analysis. Scale represents P value. B: KEGG analysis downstream pathways regulated by CD36 in advanced cervical cancer. C: Kaplan-Meier analysis showed the prognosis of stage 3 cervical cancer was significantly better than stage 4 cervical cancer. D: Kaplan-Meier analysis showed CD36 expression was not correlated with the prognosis of advanced cervical cancer. E: Kaplan-Meier analysis showed CD36 expression was not correlated with the prognosis of cervical cancer in stage 3. F: Kaplan-Meier analysis showed CD36 expression was not correlated with the prognosis of cervical cancer in stage 4. G, I: The expressions of CD44, CD133, Nanog, YAP, TAZ, MEK5 and ERK5 were increased after enhanced CD36 expression in C33a cells. H, J: The expressions of CD44, CD133, Nanog, YAP, TAZ, MEK5 and ERK5 in HeLa cells were reduced along with the down-regulation of CD36.

Figure 2. CD36 promotes stemness and tumor growth. A: CD36 over-expression (C33a/CD36 group) promoted tumor growth in vivo compared with the control group (C33a/Con group). B: The upper row: C33a /CD36 group; The bottom row: C33a/Con

group. C: Statistical plots of tumor volume in C33a/CD36 group and C33a/Con group (n=6). D: CD133 expression in C33a/CD36 group was significantly enhanced compared with the C33a/Con group ( $\times 400$ ). E: CD44 expression in C33a/CD36 group was significantly enhanced compared with the C33a/Con group ( $\times 400$ ).

Figure 3. CD36 promotes stemness of cervical cancer cells. A, B. The number of stem cell spheres with a diameter larger than 100 $\mu$ m in C33a/CD36 group was the highest. A-C: Interfering with CD36 expression in HeLa cells suppressed the number of stem cell spheres larger than 100 $\mu$ m (A: $\times 200$ ). D: The proliferative ability of C33a/CD36 group was increased compared with the C33a and C33a/NC group. E: Interfering with CD36 expression suppressed the proliferative ability of HeLa cells. F: The CD133-positive cells in the C33a/CD36 group were the highest compared with the C33a and C33a/NC group. G. The CD133-positive cells in the HeLa/siRNA CD36 group were the least compared with the HeLa and HeLa/NC siRNA group. H. Statistical chart of CD133-positive cell rate in C33a, C33a/NC and C33a/CD36 group. I. Statistical chart of CD133-positive cell rate in HeLa, HeLa/NC siRNA and HeLa/siRNA CD36 group. All experiments were performed in triplicate.  $*P<0.05$ .

Figure 4. Effect of CD36 overexpression on stemness of cervical cancer stem cells. A, D: Compared with C33a-CCSCs and C33a-CCSCs/NC, CD36 expression was significantly enhanced in the C33a-CCSCs/CD36 group. B: The number of stem cell spheres with a diameter larger than 100 $\mu$ m in C33a-CCSCs/CD36 group was the highest compared with the C33a-CCSCs and C33a-CCSCs/NC group (B: $\times 200$ ). C: The proliferative ability of C33a-CCSCs/CD36 group was increased compared with the C33a-CCSCs and C33a-CCSCs/NC group. E: CD44 expression was increased in C33a-CCSCs/CD36 group compared with C33a-CCSCs/NC group (E: $\times 400$ ).  $*P<0.05$ .

Figure 5. The effects of CD36 expression on chemotherapy resistance of cervical cancer stem cells. A: CD36 promotes proliferative ability of C33a-CCSCs. B,C: Verteporfin and XMD8-92 inhibited proliferation of C33a-CCSCs to some extent. D,E: CD36 increased the the expression levels of YAP, TAZ and ERK5. F: CD36 inhibited apoptosis of C33a-CCSCs. G: Verteporfin promotes apoptosis of C33a-CCSCs. H: XMD8-92 promotes apoptosis of C33a-CCSCs. All experiments were performed in triplicate.  $*P<0.05$ .

Figure 6. Effects of interfering CD36 expression on chemotherapy resistance of Hela cells. A: Interfering CD36 expression inhibited clone formation ability of HeLa cells after treated with cisplatin. B: Interfering CD36 expression inhibited proliferative ability of HeLa cells after treated with cisplatin. C, E: Interfering CD36 expression promoted apoptosis of HeLa cells after treated with cisplatin. D, F: Compared with the control groups, expression of CD44, CD133, Nanog, Bcl-2 were decreased and expression of Bax, Caspase-3 were increased in the HeLa/siRNA CD36 group. All experiments were performed in triplicate.  $*P<0.05$ .

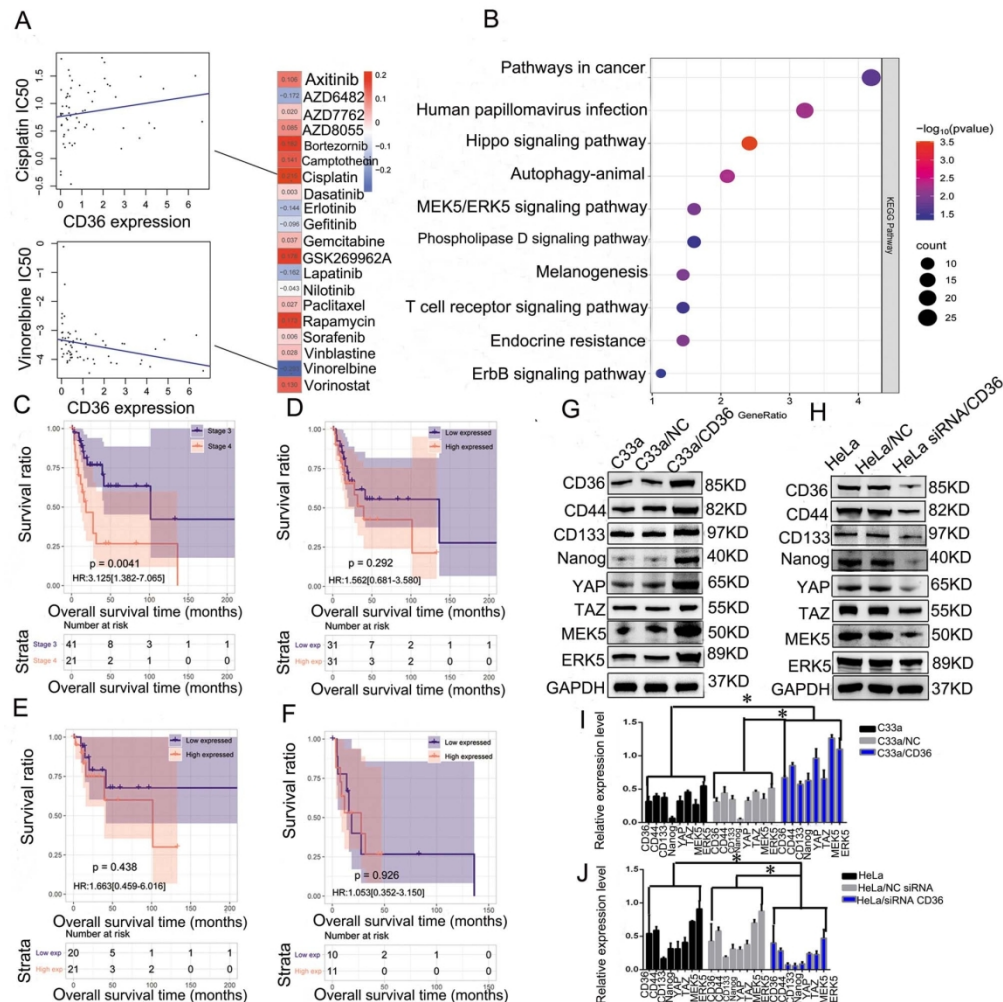


Figure 1. Correlation between CD36 expression and resistance of chemotherapy drugs and prognosis of cervical cancer and its regulated signaling pathways. A: CD36 expression had the most significant positive correlation with cisplatin resistance and the most significant negative correlation with vinorelbine in advanced cervical cancer by IC50 correlation analysis. Scale represents P value. B: KEGG analysis downstream pathways regulated by CD36 in advanced cervical cancer. C: Kaplan-Meier analysis showed the prognosis of stage 3 cervical cancer was significantly better than stage 4 cervical cancer. D: Kaplan-Meier analysis showed CD36 expression was not correlated with the prognosis of advanced cervical cancer. E: Kaplan-Meier analysis showed CD36 expression was not correlated with the prognosis of cervical cancer in stage 3. F: Kaplan-Meier analysis showed CD36 expression was not correlated with the prognosis of cervical cancer in stage 4. G, I: The expressions of CD44, CD133, Nanog, YAP, TAZ, MEK5 and ERK5 were increased after enhanced CD36 expression in C33a cells. H, J: The expressions of CD44, CD133, Nanog, YAP, TAZ, MEK5 and ERK5 in HeLa cells were reduced along with the down-regulation of CD36.

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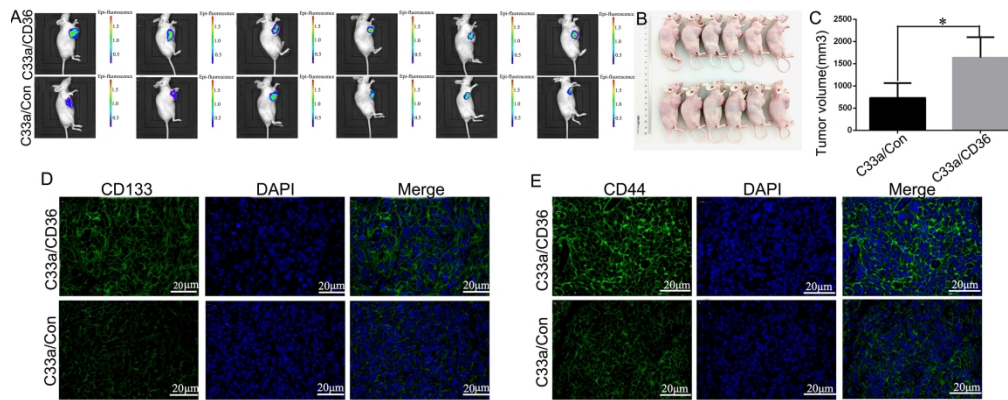


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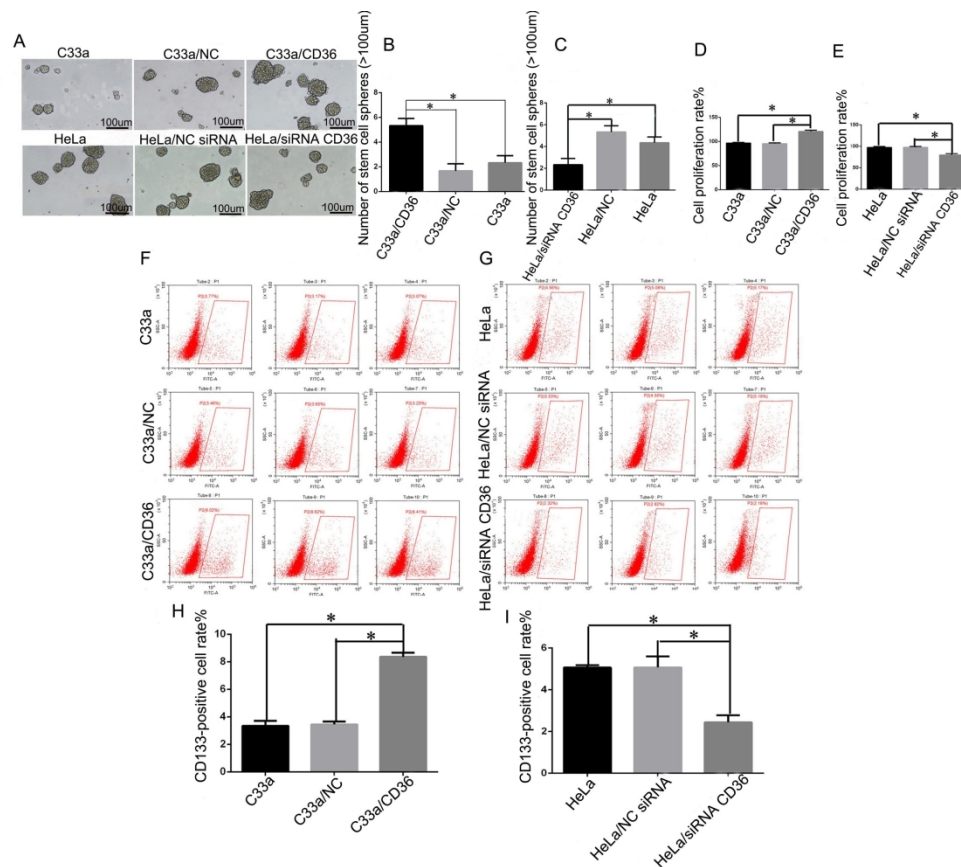


Figure 3. CD36 promotes stemness of cervical cancer cells. A, B. The number of stem cell spheres with a diameter larger than 100μm in C33a/CD36 group was the highest. A-C: Interfering with CD36 expression in HeLa cells suppressed the number of stem cell spheres larger than 100μm (A:×200). D: The proliferative ability of C33a/CD36 group was increased compared with the C33a and C33a/NC group. E: Interfering with CD36 expression suppressed the proliferative ability of HeLa cells. F: The CD133-positive cells in the C33a/CD36 group were the highest compared with the C33a and C33a/NC group. G: The CD133-positive cells in the HeLa/siRNA CD36 group were the least compared with the HeLa and HeLa/NC siRNA group. H. Statistical chart of CD133-positive cell rate in C33a, C33a/NC and C33a/CD36 group. I. Statistical chart of CD133-positive cell rate in HeLa, HeLa/NC siRNA and HeLa/siRNA CD36 group. All experiments were performed in triplicate. \*P<0.05.

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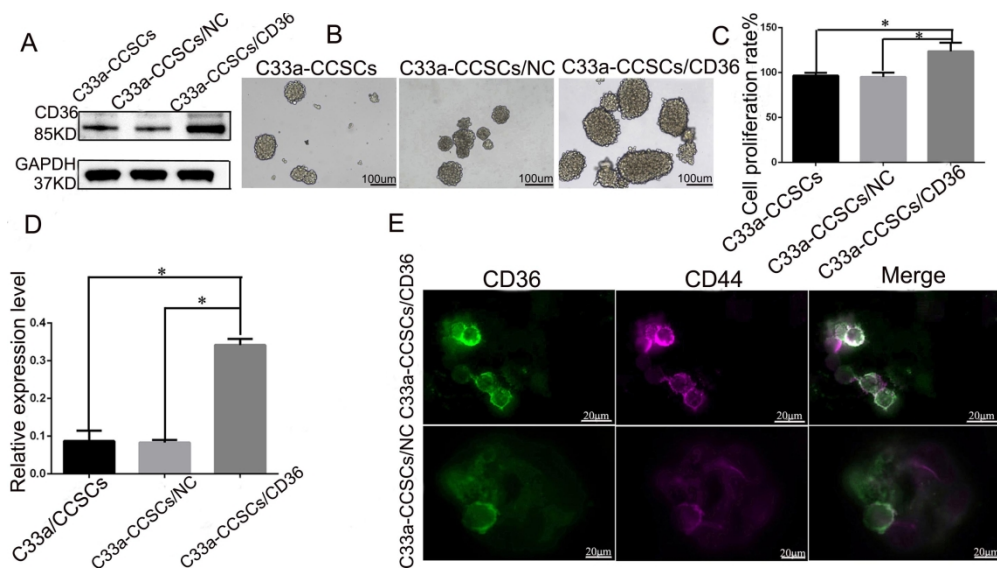


Figure 4. Effect of CD36 overexpression on stemness of cervical cancer stem cells. A, D: Compared with C33a-CCSCs and C33a-CCSCs/NC, CD36 expression was significantly enhanced in the C33a-CCSCs/CD36 group. B: The number of stem cell spheres with a diameter larger than 100µm in C33a-CCSCs/CD36 group was the highest compared with the C33a-CCSCs and C33a-CCSCs/NC group (B:×200). C: The proliferative ability of C33a-CCSCs/CD36 group was increased compared with the C33a-CCSCs and C33a-CCSCs/NC group. E: CD44 expression was increased in C33a-CCSCs/CD36 group compared with C33a-CCSCs/NC group (E:×400). \*P<0.05.

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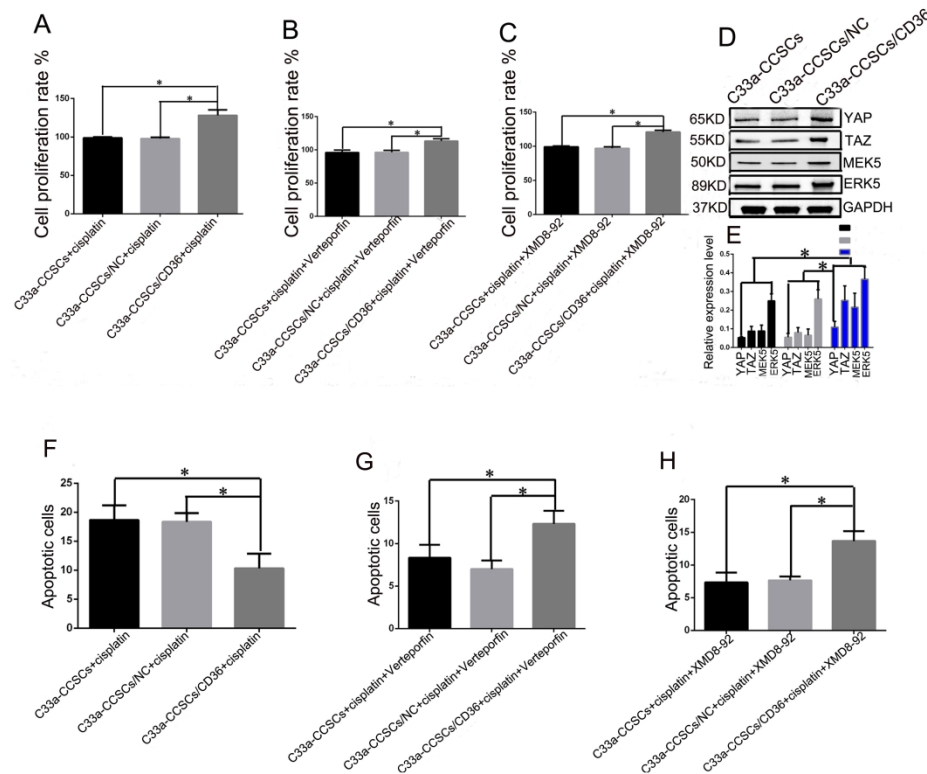


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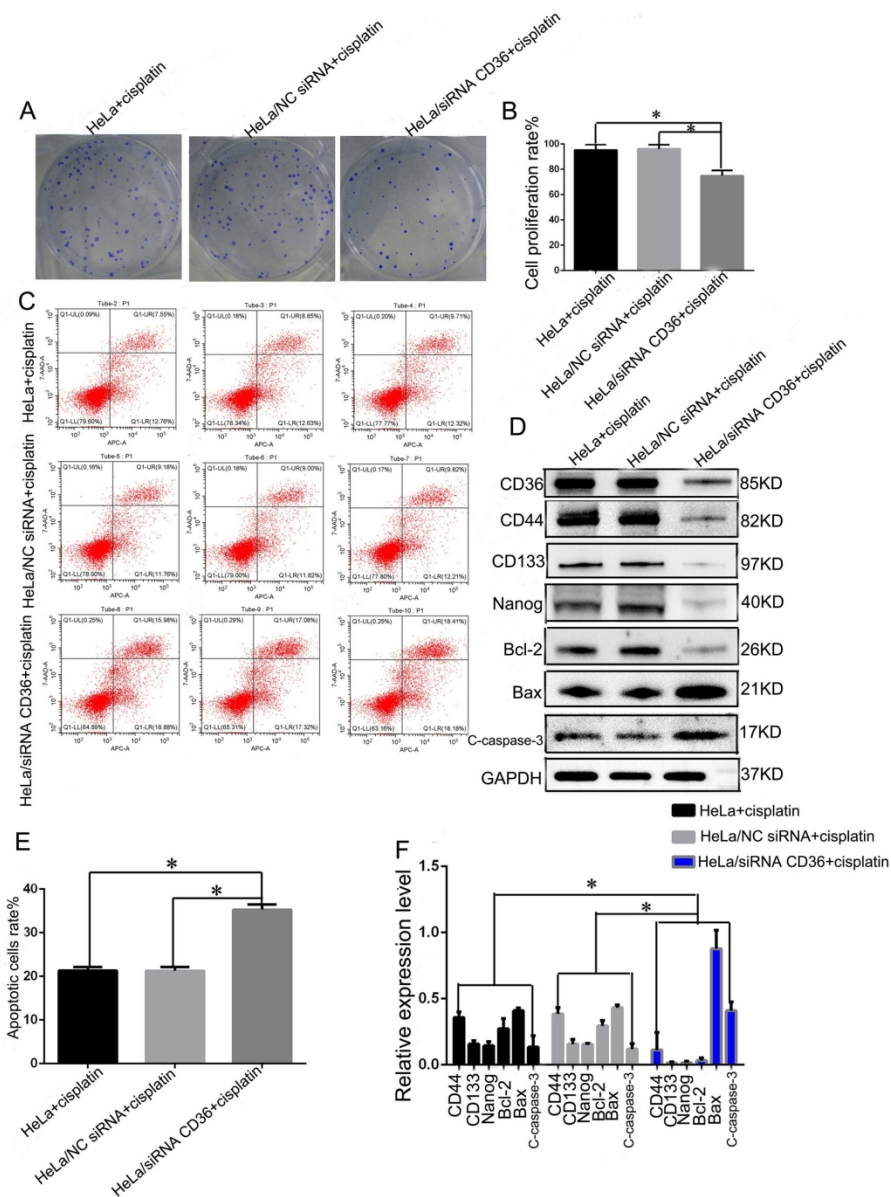


Figure 6. Effects of interfering CD36 expression on chemotherapy resistance of HeLa cells. A: Interfering CD36 expression inhibited clone formation ability of HeLa cells after treated with cisplatin. B: Interfering CD36 expression inhibited proliferative ability of HeLa cells after treated with cisplatin. C, E: Interfering CD36 expression promoted apoptosis of HeLa cells after treated with cisplatin. D, F: Compared with the control groups, expression of CD44, CD133, Nanog, Bcl-2 were decreased and expression of Bax, Caspase-3 were increased in the HeLa/siRNA CD36 group. All experiments were performed in triplicate.\*P<0.05.

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| gene          | time     | OS | Stage        | CD36     | status |
|---------------|----------|----|--------------|----------|--------|
| TCGA-JX-A3Q0- | 212.5    |    | 0 Stage III  | 0.106334 | 0      |
| TCGA-VS-A9UD- | 24.63333 |    | 0 Stage IIIA | 0.379796 | 0      |
| TCGA-VS-A950- | 40.7     |    | 0 Stage IIIA | 3.011863 | 1      |
| TCGA-LP-A5U2- | 0.3      |    | 0 Stage IIIB | 0.072316 | 0      |
| TCGA-C5-A1BQ- | 20.13333 |    | 1 Stage IIIB | 0.079746 | 0      |
| TCGA-FU-A40J- | 14.2     |    | 0 Stage IIIB | 0.11357  | 0      |
| TCGA-DR-A0ZM- | 59.7     |    | 0 Stage IIIB | 0.118669 | 0      |
| TCGA-DG-A2KJ- | 96.43333 |    | 0 Stage IIIB | 0.154352 | 0      |
| TCGA-FU-A23K- | 12.4     |    | 0 Stage IIIB | 0.200696 | 0      |
| TCGA-C5-A7X3- | 9.466667 |    | 1 Stage IIIB | 0.269161 | 0      |
| TCGA-BI-A0VR- | 50.16667 |    | 0 Stage IIIB | 0.340719 | 0      |
| TCGA-FU-A770- | 1.133333 |    | 0 Stage IIIB | 0.380776 | 0      |
| TCGA-EK-A2PI- | 19.53333 |    | 0 Stage IIIB | 0.397804 | 0      |
| TCGA-C5-A1MN- | 41.5     |    | 1 Stage IIIB | 0.427787 | 0      |
| TCGA-DG-A2KK- | 83.2     |    | 0 Stage IIIB | 0.457832 | 0      |
| TCGA-VS-A8EC- | 47.16667 |    | 0 Stage IIIB | 0.490488 | 0      |
| TCGA-EA-A6QX- | 24.33333 |    | 0 Stage IIIB | 0.498557 | 0      |
| TCGA-LP-A4AU- | 11.43333 |    | 0 Stage IIIB | 0.530072 | 0      |
| TCGA-C5-A1BI- | 37.06667 |    | 0 Stage IIIB | 0.55122  | 0      |
| TCGA-EK-A3GN- | 0.9      |    | 0 Stage IIIB | 0.55288  | 0      |
| TCGA-VS-A9UL- | 14.73333 |    | 1 Stage IIIB | 0.605414 | 1      |
| TCGA-MA-AA43- | 11.53333 |    | 0 Stage IIIB | 0.610311 | 1      |
| TCGA-ZX-AA5X- | 3.966667 |    | 0 Stage IIIB | 0.765881 | 1      |
| TCGA-C5-A7UH- | 132.9333 |    | 0 Stage IIIB | 0.779847 | 1      |
| TCGA-EA-A3QD- | 13.23333 |    | 0 Stage IIIB | 0.958895 | 1      |
| TCGA-C5-A1MH- | 39.53333 |    | 1 Stage IIIB | 1.105976 | 1      |
| TCGA-EK-A2GZ- | 12.76667 |    | 0 Stage IIIB | 1.20161  | 1      |
| TCGA-VS-A954- | 57.13333 |    | 0 Stage IIIB | 1.286538 | 1      |
| TCGA-FU-A3TQ- | 26.5     |    | 0 Stage IIIB | 1.364193 | 1      |
| TCGA-HM-A4S6- | 15.13333 |    | 0 Stage IIIB | 1.602762 | 1      |
| TCGA-DS-A3LQ- | 23.3     |    | 0 Stage IIIB | 1.664361 | 1      |
| TCGA-MA-AA3X- | 19.83333 |    | 0 Stage IIIB | 2.007529 | 1      |
| TCGA-UC-A7PG- | 12.33333 |    | 1 Stage IIIB | 2.123608 | 1      |
| TCGA-EA-A44S- | 12.3     |    | 0 Stage IIIB | 2.133504 | 1      |
| TCGA-C5-A1MK- | 2.466667 |    | 1 Stage IIIB | 2.269569 | 1      |
| TCGA-FU-A5XV- | 10.7     |    | 0 Stage IIIB | 2.381891 | 1      |
| TCGA-VS-A8EB- | 10.16667 |    | 1 Stage IIIB | 2.912379 | 1      |
| TCGA-EK-A2PL- | 0.433333 |    | 0 Stage IIIB | 3.820598 | 1      |
| TCGA-C5-A2LZ- | 101.5333 |    | 1 Stage IIIB | 4.217823 | 1      |
| TCGA-C5-A7CL- | 15.7     |    | 1 Stage IIIB | 4.889445 | 1      |
| TCGA-VS-A8EH- | 32.86667 |    | 0 Stage IIIB | 6.708641 | 1      |
| TCGA-VS-A953- | 15.9     |    | 1 Stage IVA  | 0.070301 | 0      |
| TCGA-VS-A9V4- | 4.4      |    | 1 Stage IVA  | 0.21431  | 0      |
| TCGA-C5-A7CK- | 136.2    |    | 1 Stage IVA  | 0.324765 | 0      |
| TCGA-VS-A8QC- | 11.66667 |    | 1 Stage IVA  | 0.463472 | 0      |
| TCGA-VS-A9UH- | 47.56667 |    | 0 Stage IVA  | 0.726318 | 1      |
| TCGA-VS-A9UV- | 3.466667 |    | 1 Stage IVA  | 0.823503 | 1      |
| TCGA-DS-A109- | 8.866667 |    | 1 Stage IVA  | 1.211253 | 1      |
| TCGA-VS-A8EK- | 27.63333 |    | 1 Stage IVA  | 2.647716 | 1      |
| TCGA-EA-A50E- | 7.566667 |    | 1 Stage IVA  | 4.472247 | 1      |
| TCGA-VS-A9V1- | 5.233333 |    | 1 Stage IVB  | 0.076624 | 0      |
| TCGA-IR-A3LI- | 83.1     |    | 0 Stage IVB  | 0.089749 | 0      |
| TCGA-EX-A449- | 14.9     |    | 0 Stage IVB  | 0.100071 | 0      |

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| 2  | TCGA-EK-A2RB- | 0.3      | 0 Stage IVB | 0.158049 | 0 |
| 3  | TCGA-C5-A7X5- | 13.8     | 1 Stage IVB | 0.238642 | 0 |
| 4  | TCGA-VS-A9UY- | 18.5     | 1 Stage IVB | 0.332576 | 0 |
| 5  | TCGA-VS-A9UM- | 27.63333 | 1 Stage IVB | 0.444087 | 0 |
| 6  | TCGA-VS-A9U6- | 44       | 0 Stage IVB | 0.485612 | 0 |
| 7  | TCGA-HM-A6W2- | 9.566667 | 0 Stage IVB | 0.935705 | 1 |
| 8  | TCGA-VS-A8QM- | 31.7     | 1 Stage IVB | 1.552829 | 1 |
| 9  | TCGA-JW-A5VH- | 3.333333 | 1 Stage IVB | 3.769822 | 1 |
| 10 | TCGA-VS-A9V3- | 18       | 0 Stage IVB | 6.390622 | 1 |
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For Peer Review

| ID               | CD36-exp | Axitinib | AZD6482  | AZD7762  | AZD8055  | Bortezomi | Camptothec |
|------------------|----------|----------|----------|----------|----------|-----------|------------|
| TCGA-BI-A0VR-01A | 0.340719 | 3.660418 | 3.420336 | -0.22302 | 0.347645 | -5.39689  | -4.39713   |
| TCGA-C5-A1BI-01B | 0.55122  | 3.821308 | 3.355618 | 0.432887 | 0.346588 | -5.32694  | -3.87563   |
| TCGA-C5-A1BQ-01C | 0.079746 | 3.443232 | 3.872432 | -0.41587 | 0.344161 | -4.75723  | -3.96468   |
| TCGA-C5-A1MH-01A | 1.105976 | 3.701603 | 3.575336 | -0.3089  | 0.310564 | -5.74141  | -4.30453   |
| TCGA-C5-A1MK-01A | 2.269569 | 3.77154  | 3.093486 | 0.474939 | 0.393432 | -4.97088  | -3.87943   |
| TCGA-C5-A1MN-01A | 0.427787 | 3.821797 | 3.650097 | 0.373932 | 0.457148 | -5.71754  | -4.25826   |
| TCGA-C5-A2LZ-01A | 4.217823 | 3.645571 | 3.404802 | -0.15555 | 0.355855 | -5.12616  | -4.35561   |
| TCGA-C5-A7CK-01A | 0.324765 | 3.44768  | 3.814793 | 0.326722 | 0.344459 | -5.53381  | -3.42399   |
| TCGA-C5-A7CL-01A | 4.889445 | 3.4843   | 3.770544 | 0.546695 | 0.400899 | -5.17453  | -3.63285   |
| TCGA-C5-A7UH-01A | 0.779847 | 3.381721 | 4.207259 | -0.45736 | 0.333283 | -4.56416  | -3.25684   |
| TCGA-C5-A7X3-01A | 0.269161 | 3.217006 | 4.354462 | -0.13998 | 0.328243 | -5.4701   | -4.61812   |
| TCGA-C5-A7X5-01A | 0.238642 | 3.603288 | 4.655408 | -0.55578 | 0.403608 | -5.17632  | -3.38106   |
| TCGA-DG-A2KJ-01A | 0.154352 | 3.515706 | 5.437954 | 0.370926 | 0.414918 | -6.45373  | -3.70228   |
| TCGA-DG-A2KK-01A | 0.457832 | 3.616131 | 3.660089 | -0.06385 | 0.375872 | -5.66756  | -3.5361    |
| TCGA-DR-A0ZM-01A | 0.118669 | 3.196214 | 3.903703 | -0.29216 | 0.351021 | -5.22454  | -4.36823   |
| TCGA-DS-A109-01A | 1.211253 | 3.575349 | 3.030317 | -0.28671 | 0.360602 | -6.07878  | -4.51177   |
| TCGA-DS-A3LQ-01A | 1.664361 | 3.883861 | 4.59546  | -0.10492 | 0.326878 | -5.76496  | -3.02048   |
| TCGA-EA-A3QD-01A | 0.958895 | 3.359368 | 3.995161 | 0.126281 | 0.235859 | -5.15652  | -4.7216    |
| TCGA-EA-A44S-01A | 2.133504 | 3.80991  | 3.331296 | 0.261129 | 0.343101 | -5.2033   | -3.88294   |
| TCGA-EA-A50E-01A | 4.472247 | 3.431899 | 3.693166 | -0.84592 | 0.328962 | -5.15719  | -3.66776   |
| TCGA-EA-A6QX-01A | 0.498557 | 4.077666 | 3.238327 | 0.422503 | 0.396389 | -5.97338  | -3.9408    |
| TCGA-EK-A2GZ-01A | 1.20161  | 3.789138 | 3.872351 | 0.173717 | 0.423928 | -5.25097  | -3.90783   |
| TCGA-EK-A2PI-01A | 0.397804 | 3.67011  | 3.399298 | 0.32117  | 0.373241 | -4.80234  | -3.6763    |
| TCGA-EK-A2PL-01A | 3.820598 | 3.708437 | 3.627253 | 0.39672  | 0.438634 | -5.02046  | -3.23183   |
| TCGA-EK-A2RB-01A | 0.158049 | 3.863114 | 3.830147 | 0.199523 | 0.417271 | -5.37024  | -4.24342   |
| TCGA-EK-A3GN-01A | 0.55288  | 3.332777 | 4.092873 | -0.37161 | 0.373236 | -5.50629  | -4.44259   |
| TCGA-EX-A449-01A | 0.100071 | 3.572283 | 3.572704 | 1.189197 | 0.39479  | -5.17276  | -2.88618   |
| TCGA-FU-A23K-01A | 0.200696 | 2.877506 | 4.767843 | -0.07508 | 0.257939 | -5.94106  | -5.38521   |
| TCGA-FU-A3TQ-01A | 1.364193 | 3.881185 | 3.315823 | 0.36409  | 0.399608 | -5.1765   | -3.14189   |
| TCGA-FU-A40J-01A | 0.11357  | 3.4372   | 3.799475 | 0.504944 | 0.366681 | -5.87639  | -3.95308   |
| TCGA-FU-A5XV-01A | 2.381891 | 3.836095 | 3.972429 | 0.64052  | 0.456533 | -5.62644  | -3.28136   |
| TCGA-FU-A770-01A | 0.380776 | 3.774985 | 4.147668 | 0.011399 | 0.429967 | -5.10582  | -3.30846   |
| TCGA-HM-A4S6-01A | 1.602762 | 3.33701  | 2.95623  | -0.29611 | 0.313672 | -5.017    | -4.53611   |
| TCGA-HM-A6W2-01A | 0.935705 | 2.074919 | 3.132999 | -0.15226 | 0.163985 | -5.07618  | -4.27599   |
| TCGA-IR-A3LI-01A | 0.089749 | 3.947109 | 4.246733 | 0.630191 | 0.492138 | -4.76356  | -2.73612   |
| TCGA-JW-A5VH-01A | 3.769822 | 2.735463 | 4.182868 | 0.267541 | 0.216361 | -4.72408  | -4.19689   |
| TCGA-JX-A3Q0-01A | 0.106334 | 3.691324 | 3.786072 | 0.027878 | 0.371796 | -6.02767  | -4.05645   |
| TCGA-LP-A4AU-01A | 0.530072 | 3.372171 | 3.818428 | 0.599816 | 0.365407 | -5.78797  | -3.2197    |
| TCGA-LP-A5U2-01A | 0.072316 | 2.711675 | 5.421288 | -0.443   | 0.196039 | -4.92803  | -5.47903   |
| TCGA-MA-AA3X-01A | 2.007529 | 3.867088 | 3.978834 | 0.28035  | 0.393076 | -4.30736  | -2.64029   |
| TCGA-MA-AA43-01A | 0.610311 | 2.947961 | 3.720534 | 0.284293 | 0.308717 | -5.86009  | -4.22463   |
| TCGA-UC-A7PG-01A | 2.123608 | 3.763666 | 3.363524 | 0.824915 | 0.40612  | -4.65957  | -2.75298   |
| TCGA-VS-A8EB-01A | 2.912379 | 2.908252 | 4.236173 | -0.71428 | 0.319418 | -5.23914  | -4.26459   |
| TCGA-VS-A8EC-01A | 0.490488 | 3.520431 | 4.173122 | 0.59125  | 0.278654 | -3.93276  | -4.01361   |
| TCGA-VS-A8EH-01A | 6.708641 | 3.712462 | 4.213105 | 0.437219 | 0.415806 | -4.76044  | -3.47403   |
| TCGA-VS-A8EK-01A | 2.647716 | 3.815696 | 3.535534 | 0.072239 | 0.364936 | -4.76463  | -3.04937   |
| TCGA-VS-A8QC-01A | 0.463472 | 3.704734 | 3.435021 | -0.09885 | 0.349535 | -5.6289   | -3.30529   |
| TCGA-VS-A8QM-01A | 1.552829 | 3.57686  | 2.879315 | 0.083879 | 0.295933 | -5.53267  | -3.8583    |
| TCGA-VS-A950-01A | 3.011863 | 3.724383 | 3.889678 | -0.29355 | 0.387764 | -4.75334  | -3.84142   |

|    |                  |          |          |          |          |          |          |          |
|----|------------------|----------|----------|----------|----------|----------|----------|----------|
| 1  |                  |          |          |          |          |          |          |          |
| 2  | TCGA-VS-A953-01A | 0.070301 | 3.033737 | 4.363747 | -0.32202 | 0.29878  | -5.0041  | -4.87205 |
| 3  | TCGA-VS-A954-01A | 1.286538 | 3.111671 | 3.793244 | -0.50902 | 0.296277 | -4.77603 | -3.91262 |
| 4  | TCGA-VS-A9U6-01A | 0.485612 | 3.302373 | 3.427697 | -0.2741  | 0.351269 | -4.23242 | -3.83392 |
| 5  | TCGA-VS-A9UD-01A | 0.379796 | 3.408373 | 4.742586 | 0.038005 | 0.303645 | -5.46668 | -4.63593 |
| 6  | TCGA-VS-A9UH-01A | 0.726318 | 3.247305 | 3.966814 | -1.06353 | 0.319375 | -6.26988 | -5.07129 |
| 7  | TCGA-VS-A9UL-01A | 0.605414 | 3.102712 | 3.57481  | 1.033875 | 0.310092 | -3.09785 | -3.89715 |
| 8  | TCGA-VS-A9UM-01A | 0.444087 | 3.415696 | 2.453643 | 0.241226 | 0.332617 | -5.14557 | -3.61167 |
| 10 | TCGA-VS-A9UV-01A | 0.823503 | 3.643072 | 3.569784 | 0.104707 | 0.348375 | -5.65504 | -4.08504 |
| 11 | TCGA-VS-A9UY-01A | 0.332576 | 3.527999 | 3.017786 | -0.05229 | 0.362303 | -5.73103 | -4.35238 |
| 12 | TCGA-VS-A9V1-01A | 0.076624 | 3.532575 | 3.893978 | 1.00086  | 0.435788 | -4.90999 | -3.19394 |
| 13 | TCGA-VS-A9V3-01A | 6.390622 | 3.598269 | 3.134113 | 0.524616 | 0.390481 | -5.17532 | -4.0378  |
| 14 | TCGA-VS-A9V4-01A | 0.21431  | 3.448576 | 5.621583 | 1.48837  | 0.488987 | -4.78114 | -2.76515 |
| 15 | TCGA-VS-A9V4-01A | 0.21431  | 3.448576 | 5.621583 | 1.48837  | 0.488987 | -4.78114 | -2.76515 |
| 16 | TCGA-ZX-AA5X-01A | 0.765881 | 3.386493 | 3.412547 | -0.73809 | 0.321007 | -5.26508 | -3.95934 |
| 17 |                  |          |          |          |          |          |          |          |
| 18 |                  |          |          |          |          |          |          |          |
| 19 |                  |          |          |          |          |          |          |          |
| 20 |                  |          |          |          |          |          |          |          |
| 21 |                  |          |          |          |          |          |          |          |
| 22 |                  |          |          |          |          |          |          |          |
| 23 |                  |          |          |          |          |          |          |          |
| 24 |                  |          |          |          |          |          |          |          |
| 25 |                  |          |          |          |          |          |          |          |
| 26 |                  |          |          |          |          |          |          |          |
| 27 |                  |          |          |          |          |          |          |          |
| 28 |                  |          |          |          |          |          |          |          |
| 29 |                  |          |          |          |          |          |          |          |
| 30 |                  |          |          |          |          |          |          |          |
| 31 |                  |          |          |          |          |          |          |          |
| 32 |                  |          |          |          |          |          |          |          |
| 33 |                  |          |          |          |          |          |          |          |
| 34 |                  |          |          |          |          |          |          |          |
| 35 |                  |          |          |          |          |          |          |          |
| 36 |                  |          |          |          |          |          |          |          |
| 37 |                  |          |          |          |          |          |          |          |
| 38 |                  |          |          |          |          |          |          |          |
| 39 |                  |          |          |          |          |          |          |          |
| 40 |                  |          |          |          |          |          |          |          |
| 41 |                  |          |          |          |          |          |          |          |
| 42 |                  |          |          |          |          |          |          |          |
| 43 |                  |          |          |          |          |          |          |          |
| 44 |                  |          |          |          |          |          |          |          |
| 45 |                  |          |          |          |          |          |          |          |
| 46 |                  |          |          |          |          |          |          |          |
| 47 |                  |          |          |          |          |          |          |          |
| 48 |                  |          |          |          |          |          |          |          |
| 49 |                  |          |          |          |          |          |          |          |
| 50 |                  |          |          |          |          |          |          |          |
| 51 |                  |          |          |          |          |          |          |          |
| 52 |                  |          |          |          |          |          |          |          |
| 53 |                  |          |          |          |          |          |          |          |
| 54 |                  |          |          |          |          |          |          |          |
| 55 |                  |          |          |          |          |          |          |          |
| 56 |                  |          |          |          |          |          |          |          |
| 57 |                  |          |          |          |          |          |          |          |
| 58 |                  |          |          |          |          |          |          |          |
| 59 |                  |          |          |          |          |          |          |          |
| 60 |                  |          |          |          |          |          |          |          |

For Peer Review

|    | Cisplatin | Dasatinib | Erlotinib | Gefitinib | Gemcitabine | GSK269962 | Lapatinib | Nilotinib | Paclitaxel |
|----|-----------|-----------|-----------|-----------|-------------|-----------|-----------|-----------|------------|
| 1  |           |           |           |           |             |           |           |           |            |
| 2  | 0.950916  | 2.16146   | 4.292524  | 1.942194  | -2.68046    | 3.885778  | 4.496173  | 4.073465  | -3.3324    |
| 3  | 1.280191  | 2.040422  | 4.09688   | 1.46317   | -2.33814    | 3.720014  | 4.127864  | 4.074138  | -3.22221   |
| 4  | 0.791731  | 2.270998  | 4.281362  | 1.04146   | -2.45544    | 4.086031  | 4.438828  | 3.936381  | -2.03606   |
| 5  | 0.862976  | 1.588231  | 4.478878  | 1.840501  | -2.73868    | 3.585858  | 4.423051  | 3.797859  | -3.46813   |
| 6  | 1.347367  | 1.684216  | 4.602803  | 1.706186  | -2.26619    | 3.537105  | 4.282921  | 4.20859   | -2.46862   |
| 7  | 0.517843  | -0.44383  | 3.942941  | 1.676915  | -2.60342    | 3.310771  | 3.595736  | 4.543347  | -3.28083   |
| 8  | 0.644857  | 2.195087  | 4.048214  | 1.507163  | -2.53056    | 4.183922  | 4.100727  | 3.757197  | -3.39065   |
| 9  | 1.107352  | 1.72778   | 4.242914  | 2.203004  | -2.40587    | 3.4335    | 4.539429  | 4.363176  | -2.96877   |
| 10 | 1.269807  | 2.211728  | 4.331621  | 1.797745  | -2.12939    | 4.103936  | 4.718706  | 4.143335  | -1.99005   |
| 11 | 0.815662  | 2.4743    | 4.41821   | 1.341986  | -2.40724    | 3.422286  | 4.587734  | 3.99647   | -3.19434   |
| 12 | 0.20158   | 1.608899  | 4.154716  | 1.971757  | -2.79699    | 3.2249    | 4.440855  | 4.511948  | -2.94607   |
| 13 | 0.435063  | 1.592724  | 4.263558  | 2.304     | -2.69002    | 3.683328  | 4.039312  | 4.201815  | -2.883     |
| 14 | 0.949452  | 0.684987  | 4.058886  | 0.919584  | -1.60783    | 3.292487  | 3.980774  | 4.395927  | -3.83971   |
| 15 | 0.824511  | 2.460132  | 4.442929  | 2.165901  | -2.2297     | 3.382978  | 4.449434  | 4.214166  | -2.64095   |
| 16 | 0.614431  | 0.809084  | 3.809476  | 0.500197  | -2.30752    | 3.73032   | 3.849072  | 3.847458  | -3.15874   |
| 17 | 0.875313  | 0.129579  | 4.284747  | 1.759617  | -2.55637    | 3.265818  | 4.111758  | 4.078554  | -3.01377   |
| 18 | 1.218169  | 1.442366  | 3.79738   | 2.329499  | -2.24124    | 3.75342   | 3.899908  | 4.346014  | -2.5397    |
| 19 | 1.328728  | 3.381758  | 4.43067   | 1.235639  | -2.7553     | 3.806557  | 5.048806  | 3.875422  | -2.56183   |
| 20 | 1.454748  | 3.06338   | 4.200706  | 1.115945  | -2.33515    | 4.100987  | 4.15903   | 4.158105  | -2.82483   |
| 21 | 0.551063  | 1.544337  | 4.34692   | 1.443783  | -2.58665    | 3.61115   | 4.232647  | 4.176752  | -2.75066   |
| 22 | 1.502916  | 1.866474  | 3.460722  | 1.347625  | -2.41153    | 4.012389  | 3.771293  | 4.064609  | -3.28292   |
| 23 | 0.571408  | 1.701636  | 4.204792  | 1.973101  | -2.32644    | 3.924281  | 3.778543  | 4.52083   | -2.55107   |
| 24 | 1.177973  | 2.133603  | 3.638119  | 1.773845  | -2.18676    | 3.594448  | 3.949456  | 4.342443  | -2.3834    |
| 25 | 1.488192  | 0.852337  | 3.885256  | 1.006714  | -2.25582    | 3.574798  | 3.353923  | 4.271532  | -3.58069   |
| 26 | 0.813896  | 2.462181  | 3.767896  | 1.161985  | -2.26427    | 3.65704   | 3.428604  | 4.422827  | -3.98214   |
| 27 | -0.46544  | 2.107433  | 3.975548  | 2.190334  | -2.61331    | 3.551901  | 4.329774  | 4.275381  | -3.41578   |
| 28 | 1.812286  | 4.136732  | 4.832369  | 3.726508  | -1.89225    | 3.873202  | 4.416812  | 4.587459  | -2.67936   |
| 29 | 0.264727  | 1.954422  | 4.766934  | 1.930096  | -2.51423    | 2.901969  | 4.994495  | 4.40953   | -3.79134   |
| 30 | 1.197446  | 1.781702  | 4.609894  | 1.793427  | -2.26321    | 3.759707  | 4.228811  | 4.048684  | -3.13732   |
| 31 | 0.613462  | 1.977146  | 3.949335  | 1.813302  | -2.57108    | 3.545913  | 4.438464  | 3.832837  | -2.89604   |
| 32 | 1.322097  | 0.603233  | 3.451542  | 1.263522  | -2.32832    | 4.08119   | 3.459017  | 4.226727  | -4.0684    |
| 33 | 0.704233  | 1.761162  | 4.563335  | 2.060188  | -2.31911    | 4.081772  | 4.226451  | 4.172772  | -2.67134   |
| 34 | 0.530375  | 1.923662  | 4.438524  | 1.672812  | -2.56336    | 3.848121  | 4.389686  | 3.712072  | -2.74779   |
| 35 | 0.66104   | 2.862238  | 4.773831  | 3.392513  | -2.56757    | 3.471464  | 4.802865  | 4.38787   | -1.37242   |
| 36 | 1.247729  | 2.335702  | 4.536737  | 2.339807  | -1.47514    | 4.032506  | 4.017111  | 4.704782  | -2.70531   |
| 37 | 0.370405  | 4.412203  | 5.185329  | 3.586953  | -2.63729    | 3.121692  | 5.359177  | 4.545868  | -1.76626   |
| 38 | 0.598716  | 1.421097  | 4.141602  | 2.088766  | -2.54957    | 3.869666  | 4.193122  | 3.9488    | -3.42172   |
| 39 | 0.778483  | 1.79602   | 4.293053  | 2.179184  | -2.30627    | 3.566302  | 4.259526  | 4.494897  | -3.02637   |
| 40 | -0.29877  | 3.601592  | 4.4615    | 1.88245   | -3.27667    | 3.533725  | 5.076574  | 3.68089   | -2.74268   |
| 41 | 1.830428  | 2.229908  | 5.007963  | 2.076224  | -2.07038    | 4.420872  | 4.686158  | 4.20891   | -2.55894   |
| 42 | 0.237193  | -0.39311  | 4.556381  | 3.017218  | -2.50775    | 2.196241  | 4.678821  | 4.364137  | -3.21134   |
| 43 | 1.746707  | 1.921272  | 4.382228  | 1.85471   | -1.9947     | 3.842081  | 4.273502  | 4.289822  | -2.42524   |
| 44 | 0.178984  | 1.256453  | 4.013303  | 1.974692  | -2.67835    | 2.801132  | 4.170863  | 4.327984  | -3.35409   |
| 45 | 1.131657  | 3.732407  | 4.142123  | 2.221708  | -2.38159    | 4.103348  | 4.55681   | 4.428819  | -2.5883    |
| 46 | 0.669996  | 2.040609  | 3.5454    | 1.842844  | -2.26414    | 4.022135  | 3.584562  | 4.254799  | -3.27743   |
| 47 | 1.117571  | 2.837008  | 4.237654  | 1.679256  | -2.52172    | 4.172164  | 4.240442  | 3.935036  | -2.41273   |
| 48 | 0.926228  | 1.73848   | 3.980862  | 1.799391  | -2.31361    | 3.102586  | 4.076423  | 4.223387  | -3.30432   |
| 49 | 1.035877  | 1.373136  | 4.057072  | 1.67835   | -2.48503    | 3.539969  | 4.274614  | 3.801111  | -3.27714   |
| 50 | 0.552994  | 2.46457   | 4.520832  | 2.141806  | -2.38556    | 3.623907  | 4.49742   | 4.645754  | -2.73316   |

|    |          |          |          |          |          |          |          |          |          |  |
|----|----------|----------|----------|----------|----------|----------|----------|----------|----------|--|
| 1  |          |          |          |          |          |          |          |          |          |  |
| 2  | -0.23502 | 1.788667 | 4.740385 | 1.392294 | -2.84267 | 3.260118 | 5.038803 | 4.227554 | -3.24122 |  |
| 3  | 0.614678 | 2.13734  | 4.703801 | 1.589829 | -2.39352 | 3.535996 | 4.706318 | 4.158191 | -3.12304 |  |
| 4  | 0.28043  | 2.452917 | 4.56252  | 2.197038 | -2.36584 | 3.91761  | 4.70694  | 3.900582 | -1.91587 |  |
| 5  | 1.130884 | 3.009235 | 4.458021 | 2.084026 | -2.51809 | 3.543144 | 4.897084 | 4.402465 | -3.46615 |  |
| 6  | 0.275101 | 0.916707 | 4.339056 | 1.502846 | -2.80574 | 2.948225 | 4.705291 | 4.09066  | -3.22035 |  |
| 7  | 1.418336 | 4.883698 | 5.457504 | 3.401156 | -2.01118 | 3.991336 | 5.481452 | 4.164013 | -1.13256 |  |
| 8  | 1.60786  | 1.750575 | 4.259087 | 1.926549 | -2.25013 | 3.799057 | 4.242302 | 3.73774  | -2.83269 |  |
| 9  | 0.737659 | 2.777494 | 4.382401 | 1.905761 | -2.42909 | 3.794484 | 4.324793 | 4.28861  | -2.93146 |  |
| 10 | 0.729908 | 1.123077 | 4.169695 | 1.577653 | -2.73642 | 4.112842 | 4.351852 | 3.847051 | -3.15063 |  |
| 11 | 0.817987 | 3.628466 | 4.91696  | 2.830861 | -1.95184 | 3.979    | 4.599951 | 4.616956 | -1.98749 |  |
| 12 | 1.504162 | 2.55454  | 4.162554 | 1.735807 | -2.32825 | 4.128369 | 4.41867  | 4.004019 | -2.45378 |  |
| 13 | 0.399013 | 4.452284 | 4.873152 | 2.939835 | -2.14496 | 4.620983 | 4.985564 | 4.367707 | 0.220104 |  |
| 14 | 0.553907 | 0.800853 | 4.237066 | 1.872171 | -2.59983 | 3.265563 | 4.524342 | 3.763652 | -2.64762 |  |
| 15 |          |          |          |          |          |          |          |          |          |  |
| 16 |          |          |          |          |          |          |          |          |          |  |
| 17 |          |          |          |          |          |          |          |          |          |  |
| 18 |          |          |          |          |          |          |          |          |          |  |
| 19 |          |          |          |          |          |          |          |          |          |  |
| 20 |          |          |          |          |          |          |          |          |          |  |
| 21 |          |          |          |          |          |          |          |          |          |  |
| 22 |          |          |          |          |          |          |          |          |          |  |
| 23 |          |          |          |          |          |          |          |          |          |  |
| 24 |          |          |          |          |          |          |          |          |          |  |
| 25 |          |          |          |          |          |          |          |          |          |  |
| 26 |          |          |          |          |          |          |          |          |          |  |
| 27 |          |          |          |          |          |          |          |          |          |  |
| 28 |          |          |          |          |          |          |          |          |          |  |
| 29 |          |          |          |          |          |          |          |          |          |  |
| 30 |          |          |          |          |          |          |          |          |          |  |
| 31 |          |          |          |          |          |          |          |          |          |  |
| 32 |          |          |          |          |          |          |          |          |          |  |
| 33 |          |          |          |          |          |          |          |          |          |  |
| 34 |          |          |          |          |          |          |          |          |          |  |
| 35 |          |          |          |          |          |          |          |          |          |  |
| 36 |          |          |          |          |          |          |          |          |          |  |
| 37 |          |          |          |          |          |          |          |          |          |  |
| 38 |          |          |          |          |          |          |          |          |          |  |
| 39 |          |          |          |          |          |          |          |          |          |  |
| 40 |          |          |          |          |          |          |          |          |          |  |
| 41 |          |          |          |          |          |          |          |          |          |  |
| 42 |          |          |          |          |          |          |          |          |          |  |
| 43 |          |          |          |          |          |          |          |          |          |  |
| 44 |          |          |          |          |          |          |          |          |          |  |
| 45 |          |          |          |          |          |          |          |          |          |  |
| 46 |          |          |          |          |          |          |          |          |          |  |
| 47 |          |          |          |          |          |          |          |          |          |  |
| 48 |          |          |          |          |          |          |          |          |          |  |
| 49 |          |          |          |          |          |          |          |          |          |  |
| 50 |          |          |          |          |          |          |          |          |          |  |
| 51 |          |          |          |          |          |          |          |          |          |  |
| 52 |          |          |          |          |          |          |          |          |          |  |
| 53 |          |          |          |          |          |          |          |          |          |  |
| 54 |          |          |          |          |          |          |          |          |          |  |
| 55 |          |          |          |          |          |          |          |          |          |  |
| 56 |          |          |          |          |          |          |          |          |          |  |
| 57 |          |          |          |          |          |          |          |          |          |  |
| 58 |          |          |          |          |          |          |          |          |          |  |
| 59 |          |          |          |          |          |          |          |          |          |  |
| 60 |          |          |          |          |          |          |          |          |          |  |

For Peer Review

|    | Rapamycin | Sorafenib | Vinblastine | Vinorelbine | Vorinostat |
|----|-----------|-----------|-------------|-------------|------------|
| 1  |           |           |             |             |            |
| 2  |           |           |             |             |            |
| 3  | -0.36763  | 3.926204  | -4.40076    | -4.05869    | 1.217697   |
| 4  | -0.31151  | 3.888592  | -3.63604    | -3.58695    | 1.165243   |
| 5  | 0.128674  | 3.837066  | -4.32029    | -3.08184    | 1.452139   |
| 6  | -0.50805  | 3.749559  | -4.58345    | -4.02613    | 1.073707   |
| 7  | 0.279735  | 3.978286  | -4.12423    | -3.89331    | 1.174451   |
| 8  | 0.368181  | 3.85032   | -4.04935    | -3.41349    | 1.529692   |
| 9  | -0.3731   | 3.828889  | -4.17399    | -4.25532    | 1.042122   |
| 10 | 0.105372  | 3.924886  | -3.96709    | -3.79634    | 1.467364   |
| 11 | 0.334165  | 3.899985  | -3.89481    | -3.31695    | 1.725247   |
| 12 | -0.43777  | 3.678772  | -3.86615    | -3.29145    | 1.100983   |
| 13 | -0.46275  | 3.790866  | -4.86329    | -3.48791    | 0.954243   |
| 14 | 0.553332  | 3.738785  | -3.76892    | -3.27325    | 1.633627   |
| 15 | -1.27594  | 3.6282    | -4.32662    | -0.11353    | 1.68845    |
| 16 | -0.32188  | 3.999454  | -3.55938    | -3.12191    | 1.268514   |
| 17 | 0.378836  | 3.885065  | -4.70798    | -3.25526    | 1.1478     |
| 18 | -0.33231  | 3.875805  | -4.8022     | -3.57573    | 1.452164   |
| 19 | 0.15774   | 3.878174  | -3.64028    | -3.13453    | 1.505887   |
| 20 | -0.2942   | 3.799399  | -4.58929    | -3.31525    | 0.736583   |
| 21 | -0.16274  | 3.938977  | -4.02878    | -3.50966    | 1.144405   |
| 22 | 0.339959  | 3.696684  | -4.29893    | -3.64355    | 1.129735   |
| 23 | -0.98141  | 3.797168  | -3.77963    | -3.51738    | 1.445567   |
| 24 | 0.319482  | 3.764885  | -3.75126    | -3.44114    | 1.821458   |
| 25 | -0.23965  | 4.004999  | -2.82277    | -3.71277    | 1.878713   |
| 26 | -0.5309   | 3.863681  | -4.05085    | -4.03054    | 1.578255   |
| 27 | -0.6203   | 3.784875  | -4.71775    | -3.81338    | 1.548686   |
| 28 | 0.132469  | 3.827771  | -4.33933    | -3.93456    | 1.159707   |
| 29 | -1.97817  | 3.819072  | -3.16193    | -2.76322    | 1.905009   |
| 30 | -1.06849  | 3.743708  | -5.44707    | -2.51057    | 0.634287   |
| 31 | -0.1235   | 3.990329  | -4.58515    | -4.47038    | 1.419217   |
| 32 | -0.55713  | 3.969997  | -2.65633    | -3.06668    | 1.197902   |
| 33 | 0.174358  | 3.723432  | -4.02699    | -4.11465    | 1.427325   |
| 34 | -0.18149  | 4.001736  | -2.93104    | -3.07576    | 1.524144   |
| 35 | -0.06503  | 3.85991   | -4.71869    | -4.15232    | 1.107298   |
| 36 | -0.35691  | 3.831306  | -4.22983    | -3.91501    | 0.666758   |
| 37 | -0.47534  | 4.011108  | -2.87439    | -2.42466    | 1.981495   |
| 38 | -0.44922  | 3.814991  | -3.93583    | -4.15772    | 0.654949   |
| 39 | -0.61335  | 3.895003  | -4.544      | -3.91173    | 1.394981   |
| 40 | -0.06237  | 3.786951  | -3.82227    | -3.82406    | 1.270271   |
| 41 | 0.58384   | 3.634564  | -5.36115    | -3.64019    | 0.004422   |
| 42 | 0.127203  | 3.929562  | -3.36072    | -3.36411    | 1.447872   |
| 43 | 0.089595  | 3.976097  | -4.34157    | -3.36803    | 1.00532    |
| 44 | 0.300597  | 4.023689  | -3.40491    | -3.59374    | 1.419987   |
| 45 | -0.56261  | 3.655679  | -3.74696    | -3.45268    | 1.16362    |
| 46 | -0.74413  | 3.911855  | -3.61467    | -3.94177    | 1.052804   |
| 47 | 0.153929  | 3.75463   | -4.01052    | -4.39556    | 1.880747   |
| 48 | 0.187387  | 3.918333  | -3.55395    | -4.14387    | 1.264746   |
| 49 | -0.4069   | 3.816594  | -4.18094    | -3.75712    | 1.159898   |
| 50 | -0.43469  | 3.907158  | -4.0395     | -3.99742    | 1.078783   |
| 51 | -0.32198  | 3.782265  | -3.9522     | -3.4632     | 1.521566   |

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|----|----------------------------------------------|
| 1  |                                              |
| 2  | -0.00975 3.917778 -4.33664 -3.35705 0.857542 |
| 3  | 0.129326 3.908113 -4.46493 -3.97244 0.817574 |
| 4  | -0.01661 3.791662 -3.73169 -3.84264 1.304176 |
| 5  | -1.51812 3.673048 -5.13845 -3.24524 1.347883 |
| 6  | -0.39385 3.757053 -5.18885 -3.02747 1.19654  |
| 7  | -0.14838 4.185658 -3.66177 -3.42134 1.277418 |
| 8  | -0.13575 3.838325 -4.19894 -4.02974 1.055105 |
| 9  | -0.13564 3.727249 -3.96745 -3.93668 1.185539 |
| 10 | 0.064978 3.933326 -4.02753 -4.11291 1.073728 |
| 11 | -0.60133 3.816778 -2.84908 -2.57004 1.644087 |
| 12 | -0.27338 4.045093 -3.41993 -3.16614 1.345514 |
| 13 | 0.962058 3.821037 -0.93194 -1.41253 0.937295 |
| 14 | 0.180313 3.904889 -4.17795 -3.72743 0.915782 |
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For Peer Review

| ID       | p        | Cor      |
|----------|----------|----------|
| FABP4    | 4.16E-07 | 0.591409 |
| NRG4     | 5.83E-07 | 0.585338 |
| SPINK9   | 7.28E-06 | 0.535555 |
| CLN8     | 1.38E-05 | 0.521561 |
| ARL5A    | 1.61E-05 | 0.518079 |
| ARRDC4   | 3.99E-05 | 0.496895 |
| CARTPT   | 4.22E-05 | 0.495513 |
| LCE4A    | 4.97E-05 | 0.49155  |
| SLC13A5  | 5.31E-05 | 0.489925 |
| FABP9    | 5.98E-05 | 0.486948 |
| LYSMD4   | 7.42E-05 | 0.481545 |
| OR13A1   | 8.42E-05 | 0.478334 |
| SRRM3    | 0.00011  | 0.471306 |
| CHMP4C   | 0.000114 | 0.47045  |
| YAP      | 0.000119 | 0.469322 |
| KRTAP4.4 | 0.000122 | 0.468766 |
| GCNT4    | 0.000126 | 0.467782 |
| C4orf17  | 0.000147 | 0.463735 |
| KRT10    | 0.000148 | 0.463557 |
| TMEM79   | 0.000152 | 0.462754 |
| GNAI1    | 0.000163 | 0.460891 |
| GAN      | 0.000166 | 0.460428 |
| C22orf23 | 0.000169 | 0.45993  |
| ACKR3    | 0.000176 | 0.458944 |
| ZMIZ1    | 0.000178 | 0.458537 |
| MEK5     | 0.000189 | 0.457009 |
| CHRNA3   | 0.000195 | 0.456124 |
| TMEM45A  | 0.000221 | 0.452625 |
| EPHA4    | 0.000224 | 0.452266 |
| RAB10    | 0.000227 | 0.45184  |
| USH1G    | 0.000251 | 0.449117 |
| SLC12A6  | 0.000257 | 0.448422 |
| PGLYRP4  | 0.000262 | 0.447904 |
| TAZ      | 0.000266 | 0.447494 |
| REL      | 0.000274 | 0.446642 |
| DUSP13   | 0.000279 | 0.446069 |
| MDM2     | 0.00028  | 0.446018 |
| FZD10    | 0.000313 | 0.442806 |
| PSAPL1   | 0.000325 | 0.441764 |
| BNIP1    | 0.000338 | 0.440647 |
| LENEP    | 0.000338 | 0.440615 |
| CD44     | 0.000342 | 0.440306 |
| ACVR1C   | 0.000348 | 0.439765 |
| RORA     | 0.000361 | 0.438756 |
| ERK5     | 0.000362 | 0.438632 |
| SEMA3C   | 0.000399 | 0.435854 |
| KCNK7    | 0.000403 | 0.435577 |
| TMEM155  | 0.000407 | 0.435224 |
| CST8     | 0.000408 | 0.435179 |
| STAU2    | 0.000412 | 0.43489  |
| SPTBN5   | 0.000416 | 0.434645 |
| ATAD2B   | 0.000424 | 0.43409  |
| FBXO22   | 0.000425 | 0.434004 |

|    |           |          |          |
|----|-----------|----------|----------|
| 1  |           |          |          |
| 2  | TNFRSF10I | 0.000427 | 0.43388  |
| 3  | CD133     | 0.000429 | 0.43375  |
| 4  | TGM5      | 0.000433 | 0.433455 |
| 5  | PIGF      | 0.000443 | 0.43278  |
| 6  | PGLYRP3   | 0.000446 | 0.432596 |
| 7  | RAB18     | 0.000448 | 0.432424 |
| 8  | KLHL20    | 0.000456 | 0.43189  |
| 9  | CEL       | 0.000469 | 0.431108 |
| 10 | NTRK3     | 0.000481 | 0.430357 |
| 11 | TMEM68    | 0.000489 | 0.429835 |
| 12 | C1orf61   | 0.000491 | 0.429738 |
| 13 | HIST2H2BE | 0.000532 | 0.427353 |
| 14 | SPANXN1   | 0.000547 | 0.426483 |
| 15 | LRRK1     | 0.000557 | 0.425972 |
| 16 | TMEM184A  | 0.000558 | 0.42591  |
| 17 | TMEM254   | 0.000561 | 0.425718 |
| 18 | PAQR7     | 0.000605 | 0.423446 |
| 19 | VPS37A    | 0.000607 | 0.423343 |
| 20 | IPMK      | 0.000616 | 0.422926 |
| 21 | LIPK      | 0.000627 | 0.422353 |
| 22 | PRSS42    | 0.000652 | 0.421195 |
| 23 | LYPD3     | 0.000676 | 0.420087 |
| 24 | UNC5B     | 0.000728 | 0.417797 |
| 25 | FAM84A    | 0.000734 | 0.417543 |
| 26 | CPNE3     | 0.000754 | 0.41674  |
| 27 | GPR1      | 0.000768 | 0.416142 |
| 28 | TPST2     | 0.000773 | 0.415939 |
| 29 | DLX5      | 0.000778 | 0.415738 |
| 30 | ARL8B     | 0.000803 | 0.414769 |
| 31 | RALGPS2   | 0.000814 | 0.414358 |
| 32 | RIC3      | 0.000844 | 0.413208 |
| 33 | RALGAPA1  | 0.000854 | 0.41284  |
| 34 | CHP2      | 0.000856 | 0.412777 |
| 35 | SCN3A     | 0.000869 | 0.412291 |
| 36 | SLC6A14   | 0.000878 | 0.411964 |
| 37 | CHRNA4    | 0.000881 | 0.411851 |
| 38 | YPEL5     | 0.000912 | 0.410769 |
| 39 | SECISBP2L | 0.000927 | 0.410267 |
| 40 | BICD2     | 0.000928 | 0.410246 |
| 41 | RXRA      | 0.000936 | 0.409972 |
| 42 | BAALC     | 0.00096  | 0.409149 |
| 43 | PIGN      | 0.000968 | 0.408891 |
| 44 | CYP4F3    | 0.000968 | 0.408882 |
| 45 | NCKAP1    | 0.000992 | 0.40813  |
| 46 | ERMP1     | 0.000994 | 0.408051 |
| 47 | FBXW2     | 0.001006 | 0.407686 |
| 48 | SNX16     | 0.001013 | 0.407441 |
| 49 | SKIL      | 0.001024 | 0.407109 |
| 50 | ZMAT3     | 0.001032 | 0.406859 |
| 51 | TMEM117   | 0.001039 | 0.406651 |
| 52 | PGM2      | 0.001081 | 0.405368 |
| 53 | CST9L     | 0.001099 | 0.404845 |
| 54 | ANKRD2    | 0.001105 | 0.404674 |
| 55 | ADAD2     | 0.001112 | 0.404478 |

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|----|----------|----------|----------|
| 1  |          |          |          |
| 2  | FZD6     | 0.001125 | 0.404105 |
| 3  | GRHL1    | 0.001136 | 0.403782 |
| 4  | RIMS3    | 0.001155 | 0.403251 |
| 5  | LCE2D    | 0.001203 | 0.40194  |
| 6  | NOTCH3   | 0.001213 | 0.40167  |
| 7  | INTU     | 0.001221 | 0.401441 |
| 8  | AADAC    | 0.001224 | 0.401362 |
| 9  | MYO1H    | 0.001226 | 0.401316 |
| 10 | SH3GLB2  | 0.001232 | 0.401153 |
| 11 | KLHL28   | 0.001233 | 0.401147 |
| 12 | CLIP1    | 0.001252 | 0.400643 |
| 13 | SERPINB3 | 0.00127  | 0.400182 |
| 14 | FAM219B  | 0.001297 | 0.399482 |
| 15 | MXD1     | 0.0013   | 0.39942  |
| 16 | LRRC57   | 0.001323 | 0.398833 |
| 17 | SLC6A11  | 0.001328 | 0.398711 |
| 18 | KLHL42   | 0.001335 | 0.398555 |
| 19 | MPP7     | 0.001339 | 0.398454 |
| 20 | RC3H1    | 0.001361 | 0.397906 |
| 21 | NEMF     | 0.001378 | 0.397511 |
| 22 | TBC1D2B  | 0.00141  | 0.396756 |
| 23 | ABCA5    | 0.001446 | 0.395925 |
| 24 | ALAD     | 0.001446 | 0.395925 |
| 25 | BTG1     | 0.001463 | 0.395529 |
| 26 | HOXD4    | 0.00147  | 0.395387 |
| 27 | FRRS1    | 0.001471 | 0.395361 |
| 28 | AMFR     | 0.001484 | 0.395063 |
| 29 | TANC1    | 0.001488 | 0.394987 |
| 30 | FLG2     | 0.00152  | 0.394272 |
| 31 | LCE2C    | 0.001536 | 0.393931 |
| 32 | SLC4A9   | 0.001556 | 0.39349  |
| 33 | DCST1    | 0.001565 | 0.393308 |
| 34 | ABI1     | 0.00159  | 0.392784 |
| 35 | AGL      | 0.001614 | 0.392278 |
| 36 | SLC37A2  | 0.001619 | 0.392171 |
| 37 | KIFAP3   | 0.001638 | 0.391779 |
| 38 | CCR3     | 0.00165  | 0.391545 |
| 39 | TXNDC8   | 0.001664 | 0.391261 |
| 40 | DSG4     | 0.001664 | 0.39126  |
| 41 | IVNS1ABP | 0.001672 | 0.391089 |
| 42 | FAM206A  | 0.001672 | 0.391089 |
| 43 | CNDP1    | 0.001687 | 0.390789 |
| 44 | GNAL     | 0.00173  | 0.389951 |
| 45 | TP53INP1 | 0.00174  | 0.389753 |
| 46 | IPPK     | 0.001742 | 0.389722 |
| 47 | NR6A1    | 0.001742 | 0.389719 |
| 48 | SYCN     | 0.00175  | 0.389571 |
| 49 | CALCRL   | 0.001754 | 0.389487 |
| 50 | PUS10    | 0.001764 | 0.389304 |
| 51 | KIAA1841 | 0.001768 | 0.389218 |
| 52 | CDK5R1   | 0.001799 | 0.38864  |
| 53 | CREG1    | 0.001803 | 0.388555 |
| 54 | GPR155   | 0.001811 | 0.388416 |
| 55 | ESRP2    | 0.001824 | 0.38817  |

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|----|----------|----------|----------|
| 1  |          |          |          |
| 2  | CHST2    | 0.001825 | 0.388147 |
| 3  | C18orf25 | 0.001836 | 0.387943 |
| 4  | STEAP4   | 0.001857 | 0.387557 |
| 5  | USP8     | 0.001858 | 0.387538 |
| 6  | MAST4    | 0.001862 | 0.387477 |
| 7  | LCE3E    | 0.001896 | 0.386867 |
| 8  | RNF111   | 0.001949 | 0.385918 |
| 9  | LRRC28   | 0.001998 | 0.385072 |
| 10 | PHLDB3   | 0.002018 | 0.384739 |
| 12 | FAM214A  | 0.00203  | 0.384534 |
| 13 | AKTIP    | 0.002032 | 0.384502 |
| 14 | HS6ST3   | 0.002048 | 0.384232 |
| 15 | FAM214B  | 0.00205  | 0.384194 |
| 16 | NPL      | 0.002053 | 0.384158 |
| 17 | MAP2K1   | 0.002098 | 0.383416 |
| 18 | ZNF106   | 0.002106 | 0.383278 |
| 19 | RHCG     | 0.002121 | 0.383039 |
| 20 | DNAJB4   | 0.002125 | 0.382974 |
| 22 | DPP8     | 0.002128 | 0.382922 |
| 23 | ABHD2    | 0.002154 | 0.382499 |
| 24 | UBE2V2   | 0.00218  | 0.382088 |
| 25 | CMIP     | 0.002186 | 0.381996 |
| 26 | HDAC4    | 0.002196 | 0.381835 |
| 27 | ATP9B    | 0.002202 | 0.381738 |
| 28 | RNF217   | 0.002206 | 0.38169  |
| 29 | PTPDC1   | 0.002217 | 0.38151  |
| 30 | BNIP2    | 0.002243 | 0.381104 |
| 31 | ATP6V1A  | 0.002277 | 0.380595 |
| 32 | KCTD11   | 0.002281 | 0.380522 |
| 33 | LPCAT3   | 0.002291 | 0.380378 |
| 34 | PRKACB   | 0.002341 | 0.379631 |
| 35 | YTHDF3   | 0.002348 | 0.379526 |
| 36 | ZNF770   | 0.002392 | 0.378877 |
| 37 | ZBTB1    | 0.002406 | 0.378674 |
| 38 | RALGDS   | 0.002429 | 0.378343 |
| 39 | SYPL1    | 0.002434 | 0.378279 |
| 40 | MALT1    | 0.002447 | 0.378087 |
| 41 | CCDC144A | 0.002478 | 0.377646 |
| 42 | ATG14    | 0.002519 | 0.37707  |
| 43 | EPHB3    | 0.002547 | 0.376696 |
| 44 | HOXD1    | 0.002568 | 0.376402 |
| 45 | C9orf85  | 0.00258  | 0.376238 |
| 46 | CELA2B   | 0.002598 | 0.376001 |
| 47 | SLC23A2  | 0.002624 | 0.375645 |
| 48 | MAP3K2   | 0.002649 | 0.37531  |
| 49 | NAB1     | 0.002661 | 0.375158 |
| 50 | PTBP3    | 0.002704 | 0.374593 |
| 51 | CAPNS2   | 0.002713 | 0.374471 |
| 52 | SAMD8    | 0.002759 | 0.373885 |
| 53 | WWP1     | 0.002767 | 0.37377  |
| 54 | ARIH1    | 0.002796 | 0.373412 |
| 55 | OR1Q1    | 0.0028   | 0.373353 |
| 56 | COBLL1   | 0.002827 | 0.37302  |
| 57 | NIPAL2   | 0.002843 | 0.372819 |

1 KRTAP5.11 0.002875 0.372422  
2 HYAL1 0.002892 0.372217  
3 ZNF117 0.002935 0.371685  
4 RAB14 0.002947 0.371539  
5 ABCA1 0.003002 0.370883  
6 C15orf41 0.003047 0.370357  
7 STXBP5 0.003073 0.370051  
8 MOB4 0.003099 0.369743  
9 EVPL 0.003107 0.36966  
10 CA6 0.003128 0.36941  
11 RASAL2 0.00317 0.368936  
12 NANOS1 0.003178 0.368847  
13 AXDND1 0.003213 0.368451  
14 PTGS1 0.003232 0.368246  
15 ARL15 0.00327 0.367822  
16 RNASE10 0.003302 0.367476  
17 ZNF236 0.003303 0.367458  
18 PCMTD1 0.003326 0.367212  
19 HBP1 0.00333 0.367167  
20 SLC22A25 0.003338 0.367083  
21 ART1 0.003343 0.367021  
22 BLOC1S2 0.003385 0.36657  
23 HOOK3 0.003453 0.365861  
24 LRRC8A 0.00352 0.365162  
25 UNKL 0.003534 0.365013  
26 RBP3 0.003597 0.364372  
27 CLU 0.00364 0.363942  
28 KLHL36 0.003642 0.363918  
29 ZNF780A 0.003652 0.363821  
30 COL24A1 0.003688 0.363465  
31 SYK 0.003724 0.363113  
32 ZBTB6 0.003761 0.362747  
33 OR1L3 0.00377 0.36266  
34 TMEM87A 0.003812 0.362252  
35 DAAM1 0.003817 0.362204  
36 MEGF9 0.00386 0.361796  
37 SCAPER 0.003889 0.361517  
38 IMPA1 0.003913 0.361291  
39 PPP1R13L 0.003995 0.360534  
40 PAPOLG 0.004085 0.359708  
41 TSC1 0.00409 0.359665  
42 SLTM 0.004146 0.359165  
43 RETSAT 0.004183 0.358835  
44 KCTD21 0.004184 0.35883  
45 PPP2R2A 0.004192 0.358755  
46 CES4A 0.004235 0.358373  
47 BBS10 0.004236 0.358368  
48 TOM1L2 0.004242 0.358319  
49 ZNF658 0.004262 0.358143  
50 HOXD8 0.004265 0.358119  
51 FAM160B2 0.004273 0.358043  
52 SNTB2 0.004301 0.357802  
53 SLC35G1 0.004502 0.356101  
54 SERPINB7 0.00451 0.356041

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|----|----------|----------|----------|
| 1  |          |          |          |
| 2  | PAWR     | 0.00454  | 0.355793 |
| 3  | P2RY1    | 0.00455  | 0.355706 |
| 4  | CACNA2D1 | 0.004561 | 0.355614 |
| 5  | MEX3B    | 0.004564 | 0.355592 |
| 6  | LONP2    | 0.004575 | 0.355501 |
| 7  | NGRN     | 0.00458  | 0.355463 |
| 8  | ZNF385A  | 0.004617 | 0.355159 |
| 9  | ERICH1   | 0.004619 | 0.355148 |
| 10 | UBQLN1   | 0.004684 | 0.354623 |
| 11 | CDC42SE1 | 0.004789 | 0.353788 |
| 12 | SLC39A2  | 0.004799 | 0.353714 |
| 13 | PYGL     | 0.004824 | 0.353512 |
| 14 | MUCL1    | 0.004933 | 0.352674 |
| 15 | IGIP     | 0.004946 | 0.352573 |
| 16 | HSPA4L   | 0.004968 | 0.352404 |
| 17 | PDE2A    | 0.004979 | 0.352322 |
| 18 | PEX13    | 0.005014 | 0.352055 |
| 19 | YTHDC2   | 0.005017 | 0.352037 |
| 20 | XKRX     | 0.005019 | 0.35202  |
| 21 | RIOK3    | 0.005038 | 0.351876 |
| 22 | NT5C1B   | 0.005211 | 0.350598 |
| 23 | HECTD2   | 0.005217 | 0.350557 |
| 24 | GANC     | 0.005231 | 0.350449 |
| 25 | ANKRD9   | 0.005244 | 0.35036  |
| 26 | SP6      | 0.00525  | 0.350312 |
| 27 | ADAM23   | 0.005252 | 0.3503   |
| 28 | DSP      | 0.005291 | 0.350016 |
| 29 | PITPNM2  | 0.005303 | 0.34993  |
| 30 | GOLGA1   | 0.005349 | 0.349603 |
| 31 | NOD2     | 0.005368 | 0.349471 |
| 32 | CNOT7    | 0.005387 | 0.349336 |
| 33 | ZNF280D  | 0.005427 | 0.349049 |
| 34 | CEACAM19 | 0.00544  | 0.348957 |
| 35 | MTA3     | 0.005458 | 0.348832 |
| 36 | PSMD5    | 0.005468 | 0.348766 |
| 37 | MTSS1L   | 0.005543 | 0.34824  |
| 38 | LY6G6C   | 0.005563 | 0.348105 |
| 39 | OSTF1    | 0.005585 | 0.347951 |
| 40 | REEP4    | 0.005636 | 0.347602 |
| 41 | CTSB     | 0.005652 | 0.347499 |
| 42 | MTMR3    | 0.005661 | 0.347433 |
| 43 | MBLAC2   | 0.005705 | 0.34714  |
| 44 | NPC1     | 0.005751 | 0.346829 |
| 45 | DENND1B  | 0.005791 | 0.346564 |
| 46 | KRAS     | 0.005847 | 0.346196 |
| 47 | ESRP1    | 0.005939 | 0.345591 |
| 48 | EPG5     | 0.005954 | 0.345493 |
| 49 | RBM11    | 0.00596  | 0.345458 |
| 50 | PLD1     | 0.005966 | 0.345414 |
| 51 | PON2     | 0.005976 | 0.345349 |
| 52 | GAPVD1   | 0.005985 | 0.345295 |
| 53 | CCL26    | 0.005999 | 0.345201 |
| 54 | NIPA1    | 0.006063 | 0.34479  |
| 55 | UACA     | 0.006106 | 0.344519 |

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| 1  |           |          |          |
| 2  | GM2A      | 0.006138 | 0.34432  |
| 3  | RIT1      | 0.006173 | 0.344095 |
| 4  | ATP6V1G1  | 0.006221 | 0.343799 |
| 5  | PTPN12    | 0.006232 | 0.343729 |
| 6  | MFAP1     | 0.006233 | 0.343722 |
| 7  | CYB561D1  | 0.006238 | 0.343688 |
| 8  | ASB7      | 0.006269 | 0.3435   |
| 9  | CEP350    | 0.006282 | 0.343416 |
| 10 | SPINK5    | 0.006293 | 0.343351 |
| 11 | KIFC2     | 0.00632  | 0.343185 |
| 12 | ERV3.1    | 0.006358 | 0.342952 |
| 13 | ASPH      | 0.006361 | 0.342933 |
| 14 | ZNF655    | 0.006408 | 0.342645 |
| 15 | TMEM40    | 0.006467 | 0.342289 |
| 16 | TAS2R20   | 0.006518 | 0.341981 |
| 17 | EPN2      | 0.006521 | 0.341962 |
| 18 | CHMP1B    | 0.006538 | 0.341861 |
| 19 | NRBP1     | 0.006589 | 0.341556 |
| 20 | PCSK5     | 0.006592 | 0.341541 |
| 21 | SCD       | 0.006626 | 0.341338 |
| 22 | DUOXA1    | 0.006669 | 0.341089 |
| 23 | F11R      | 0.006689 | 0.340967 |
| 24 | ZSCAN29   | 0.006733 | 0.340711 |
| 25 | CD46      | 0.006765 | 0.340528 |
| 26 | TMOD3     | 0.006771 | 0.34049  |
| 27 | TPPP      | 0.006772 | 0.340485 |
| 28 | ANKRD46   | 0.006788 | 0.340395 |
| 29 | ABHD17B   | 0.006788 | 0.340394 |
| 30 | TM7SF3    | 0.006799 | 0.340329 |
| 31 | TAF7      | 0.006856 | 0.340003 |
| 32 | CYP3A7    | 0.006885 | 0.339835 |
| 33 | LCE1D     | 0.006906 | 0.339717 |
| 34 | PDILT     | 0.006948 | 0.339477 |
| 35 | SNX33     | 0.006954 | 0.339447 |
| 36 | ZNF264    | 0.007032 | 0.339006 |
| 37 | BRMS1L    | 0.007033 | 0.338999 |
| 38 | CDK17     | 0.00706  | 0.338846 |
| 39 | ASPA      | 0.007131 | 0.338455 |
| 40 | SF3B1     | 0.007145 | 0.338378 |
| 41 | CLIP4     | 0.007148 | 0.338361 |
| 42 | OSBP2     | 0.007165 | 0.338266 |
| 43 | FBX034    | 0.007225 | 0.337937 |
| 44 | GSN       | 0.007264 | 0.337725 |
| 45 | PRPF39    | 0.007386 | 0.337066 |
| 46 | NOTCH1    | 0.007397 | 0.337003 |
| 47 | CDK8      | 0.007399 | 0.336997 |
| 48 | LRFN2     | 0.007422 | 0.336872 |
| 49 | PLEKHA1   | 0.007438 | 0.336785 |
| 50 | CD58      | 0.00747  | 0.336615 |
| 51 | CDK9      | 0.007505 | 0.336428 |
| 52 | ARRDC3    | 0.007518 | 0.33636  |
| 53 | PCMTD2    | 0.007519 | 0.336355 |
| 54 | FEM1B     | 0.007522 | 0.336341 |
| 55 | TMPRSS11L | 0.007567 | 0.336103 |

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| 1  |           |          |          |
| 2  | NPHP3     | 0.007589 | 0.33599  |
| 3  | INPP4A    | 0.00761  | 0.335877 |
| 4  | HSPA2     | 0.007652 | 0.335658 |
| 5  | NOX5      | 0.007663 | 0.335601 |
| 6  | WNT11     | 0.007712 | 0.335349 |
| 7  | MAP7      | 0.007778 | 0.335006 |
| 8  | RXR1      | 0.007819 | 0.334798 |
| 9  | ADCY7     | 0.007824 | 0.334773 |
| 10 | SLURP1    | 0.007843 | 0.334675 |
| 11 | SOX7      | 0.007871 | 0.334533 |
| 12 | ATP6V1H   | 0.007912 | 0.334326 |
| 13 | FUT11     | 0.007958 | 0.334095 |
| 14 | MIB1      | 0.00797  | 0.334035 |
| 15 | CST6      | 0.008026 | 0.333751 |
| 16 | KRT74     | 0.008081 | 0.333479 |
| 17 | DHX32     | 0.008098 | 0.333397 |
| 18 | SFT2D2    | 0.008215 | 0.33282  |
| 19 | NAPG      | 0.008222 | 0.332784 |
| 20 | GDPGP1    | 0.008226 | 0.332767 |
| 21 | RAET1G    | 0.008237 | 0.33271  |
| 22 | GPR157    | 0.008325 | 0.332286 |
| 23 | LOR       | 0.008349 | 0.332171 |
| 24 | ACVR1     | 0.00844  | 0.331734 |
| 25 | ALDH1A3   | 0.008453 | 0.331668 |
| 26 | RCOR1     | 0.008479 | 0.331547 |
| 27 | CMTR2     | 0.008496 | 0.331466 |
| 28 | DSTYK     | 0.008511 | 0.331395 |
| 29 | BLOC1S6   | 0.008547 | 0.331222 |
| 30 | ALDH3B2   | 0.008616 | 0.330899 |
| 31 | HIST1H2BC | 0.008622 | 0.330869 |
| 32 | STX17     | 0.008648 | 0.33075  |
| 33 | WDR37     | 0.008672 | 0.330637 |
| 34 | SERPINB4  | 0.008679 | 0.330602 |
| 35 | DHCR24    | 0.008698 | 0.330513 |
| 36 | RGSL1     | 0.00878  | 0.330134 |
| 37 | SPINK6    | 0.008793 | 0.330074 |
| 38 | UNC119B   | 0.0088   | 0.330044 |
| 39 | TLK1      | 0.008804 | 0.330023 |
| 40 | GJB6      | 0.008841 | 0.329856 |
| 41 | TNK2      | 0.008846 | 0.329834 |
| 42 | WEE2      | 0.008855 | 0.32979  |
| 43 | NEBL      | 0.00888  | 0.329678 |
| 44 | NCCRP1    | 0.008896 | 0.329602 |
| 45 | CHD9      | 0.008904 | 0.329566 |
| 46 | CDX4      | 0.008931 | 0.329443 |
| 47 | SLC35G3   | 0.008974 | 0.32925  |
| 48 | BTBD11    | 0.009107 | 0.328649 |
| 49 | ARNT      | 0.009149 | 0.328462 |
| 50 | RAB3GAP2  | 0.009155 | 0.328436 |
| 51 | PTHLH     | 0.009178 | 0.328333 |
| 52 | ZNF624    | 0.0092   | 0.328238 |
| 53 | GALR2     | 0.009296 | 0.327811 |
| 54 | CLPX      | 0.0093   | 0.327794 |
| 55 | PPP3R1    | 0.00932  | 0.327706 |

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| 1  |           |          |          |
| 2  | LEPROT    | 0.009326 | 0.327679 |
| 3  | RBL2      | 0.009338 | 0.32763  |
| 4  | ABO       | 0.00936  | 0.327531 |
| 5  | EEA1      | 0.009381 | 0.327442 |
| 6  | MAPKAP1   | 0.009418 | 0.32728  |
| 7  | ADAMTS17  | 0.00943  | 0.327228 |
| 8  | FAM129B   | 0.009436 | 0.327202 |
| 9  | CDKN2B    | 0.009455 | 0.32712  |
| 10 | CCDC105   | 0.009479 | 0.327016 |
| 11 | ATP11B    | 0.009506 | 0.326901 |
| 12 | PUM2      | 0.009566 | 0.326641 |
| 13 | TGFA      | 0.009588 | 0.326549 |
| 14 | OR10A3    | 0.009747 | 0.325873 |
| 15 | FASTKD1   | 0.009859 | 0.325404 |
| 16 | LACC1     | 0.009897 | 0.325246 |
| 17 | DCST2     | 0.009922 | 0.325139 |
| 18 | DYNC1LI2  | 0.009926 | 0.325123 |
| 19 | DNAJA4    | 0.009971 | 0.32494  |
| 20 | CASP14    | 0.009972 | 0.324935 |
| 21 | VPS13A    | 0.009996 | 0.324835 |
| 22 | RBBP6     | 0.010109 | 0.324371 |
| 23 | MME       | 0.010155 | 0.324182 |
| 24 | GALNT1    | 0.010172 | 0.324114 |
| 25 | TRAPPC8   | 0.010202 | 0.323995 |
| 26 | BCL2L2    | 0.010215 | 0.323942 |
| 27 | CHAC2     | 0.010258 | 0.323766 |
| 28 | LNK2      | 0.010307 | 0.323571 |
| 29 | FAM43A    | 0.010367 | 0.32333  |
| 30 | BTAF1     | 0.010401 | 0.323194 |
| 31 | KATNBL1   | 0.010483 | 0.322868 |
| 32 | EFCAB14   | 0.010484 | 0.322865 |
| 33 | SERPINB8  | 0.010497 | 0.322812 |
| 34 | SLC35D1   | 0.010544 | 0.322628 |
| 35 | TMEM132B  | 0.010553 | 0.32259  |
| 36 | PHF2      | 0.010563 | 0.322551 |
| 37 | BRWD1     | 0.010586 | 0.322462 |
| 38 | CNEP1R1   | 0.010683 | 0.322083 |
| 39 | MORC3     | 0.0107   | 0.322019 |
| 40 | DMTF1     | 0.010705 | 0.321996 |
| 41 | PRR23B    | 0.010714 | 0.321963 |
| 42 | KRTAP19.3 | 0.010806 | 0.321609 |
| 43 | HDHD2     | 0.010812 | 0.321585 |
| 44 | WNT5A     | 0.010827 | 0.321527 |
| 45 | NCOA4     | 0.010908 | 0.321217 |
| 46 | PRRC2C    | 0.01093  | 0.32113  |
| 47 | LGALSL    | 0.010935 | 0.321112 |
| 48 | MFSD6     | 0.011008 | 0.320834 |
| 49 | ALOXE3    | 0.011038 | 0.320722 |
| 50 | MTSS1     | 0.011048 | 0.320682 |
| 51 | TMEM245   | 0.011057 | 0.32065  |
| 52 | CPEB4     | 0.011066 | 0.320616 |
| 53 | SCRN3     | 0.011156 | 0.320275 |
| 54 | MAP4K5    | 0.011159 | 0.320267 |
| 55 | GFOD1     | 0.011166 | 0.320241 |

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| 1  |          |          |          |
| 2  | ETV3     | 0.011171 | 0.320222 |
| 3  | UNC45A   | 0.011287 | 0.319787 |
| 4  | NIN      | 0.011332 | 0.31962  |
| 5  | NDEL1    | 0.011378 | 0.319452 |
| 6  | MAP3K9   | 0.011398 | 0.31938  |
| 7  | SLC44A1  | 0.011398 | 0.319379 |
| 8  | ADSSL1   | 0.0114   | 0.319372 |
| 9  | AP4E1    | 0.011524 | 0.318917 |
| 10 | PPP1R1C  | 0.011538 | 0.318866 |
| 11 | RRM2B    | 0.011562 | 0.318779 |
| 12 | IL20RA   | 0.011572 | 0.318741 |
| 13 | ARAP2    | 0.011595 | 0.318659 |
| 14 | GDAP2    | 0.011606 | 0.318617 |
| 15 | PAFAH1B1 | 0.011621 | 0.318566 |
| 16 | ZBTB37   | 0.011629 | 0.318535 |
| 17 | PPP2CB   | 0.011658 | 0.31843  |
| 18 | C9orf3   | 0.011749 | 0.318104 |
| 19 | MREG     | 0.011786 | 0.31797  |
| 20 | PLXNB3   | 0.011833 | 0.317802 |
| 21 | CYFIP1   | 0.011955 | 0.317371 |
| 22 | SVIL     | 0.011988 | 0.317254 |
| 23 | MEF2A    | 0.012023 | 0.317132 |
| 24 | METTL9   | 0.012044 | 0.317056 |
| 25 | RNF20    | 0.012076 | 0.316946 |
| 26 | PRDM1    | 0.012101 | 0.316859 |
| 27 | FAM110A  | 0.012135 | 0.316738 |
| 28 | ANKRD29  | 0.012147 | 0.316699 |
| 29 | MYH14    | 0.01218  | 0.316584 |
| 30 | BCL2L10  | 0.012192 | 0.316542 |
| 31 | GOLGA2   | 0.012227 | 0.316419 |
| 32 | SPPL2A   | 0.012284 | 0.316224 |
| 33 | NMB      | 0.012303 | 0.316158 |
| 34 | ECM1     | 0.012311 | 0.316129 |
| 35 | FKTN     | 0.012329 | 0.316068 |
| 36 | MYO1B    | 0.012344 | 0.316016 |
| 37 | GALR3    | 0.012356 | 0.315976 |
| 38 | SOST     | 0.012412 | 0.315784 |
| 39 | WDR59    | 0.012494 | 0.315504 |
| 40 | IGFL3    | 0.012526 | 0.315396 |
| 41 | DMTN     | 0.012622 | 0.315069 |
| 42 | SOS1     | 0.01263  | 0.315045 |
| 43 | CTRC     | 0.012707 | 0.314786 |
| 44 | PIM1     | 0.012769 | 0.314578 |
| 45 | BARX1    | 0.01278  | 0.314541 |
| 46 | SPTBN2   | 0.012893 | 0.314165 |
| 47 | ARHGEF4  | 0.012997 | 0.313821 |
| 48 | ABHD4    | 0.013057 | 0.313627 |
| 49 | AANAT    | 0.013061 | 0.313612 |
| 50 | ANKRD26  | 0.01313  | 0.313388 |
| 51 | EMP2     | 0.013135 | 0.313373 |
| 52 | SRPK2    | 0.013169 | 0.31326  |
| 53 | LIPI     | 0.013173 | 0.313247 |
| 54 | ITGAV    | 0.013184 | 0.313212 |
| 55 | PXT1     | 0.013265 | 0.31295  |

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| 1  |          |          |          |
| 2  | RDH16    | 0.013281 | 0.312897 |
| 3  | TVP23B   | 0.013303 | 0.312827 |
| 4  | GSDMC    | 0.013332 | 0.312733 |
| 5  | RAPGEFL1 | 0.013335 | 0.312725 |
| 6  | SCYL3    | 0.013347 | 0.312685 |
| 7  | LPAR5    | 0.013392 | 0.312542 |
| 8  | DUSP11   | 0.013462 | 0.312318 |
| 9  | ARID4A   | 0.013464 | 0.312312 |
| 10 | CDC42BPG | 0.013526 | 0.312116 |
| 11 | ZNF532   | 0.013604 | 0.311868 |
| 12 | BMP7     | 0.013661 | 0.311687 |
| 13 | SBN01    | 0.013739 | 0.311444 |
| 14 | SDHC     | 0.013743 | 0.311431 |
| 15 | CLCA2    | 0.01378  | 0.311316 |
| 16 | TTC21B   | 0.013799 | 0.311256 |
| 17 | THEM5    | 0.013803 | 0.311242 |
| 18 | CHD7     | 0.013811 | 0.311217 |
| 19 | AADACL2  | 0.013813 | 0.311212 |
| 20 | SOCS4    | 0.013832 | 0.311151 |
| 21 | CADM4    | 0.013859 | 0.31107  |
| 22 | TREML4   | 0.013877 | 0.311013 |
| 23 | KLC3     | 0.013906 | 0.310921 |
| 24 | GABRE    | 0.013919 | 0.310881 |
| 25 | DUOX1    | 0.013949 | 0.31079  |
| 26 | HAUS2    | 0.013963 | 0.310745 |
| 27 | UBE2G1   | 0.013972 | 0.310719 |
| 28 | STAM     | 0.01406  | 0.310447 |
| 29 | UBE2D1   | 0.014152 | 0.310167 |
| 30 | STRN     | 0.014195 | 0.310033 |
| 31 | NCOA3    | 0.014205 | 0.310003 |
| 32 | RTF1     | 0.014219 | 0.309962 |
| 33 | SREBF1   | 0.014265 | 0.309822 |
| 34 | MPP5     | 0.014391 | 0.309441 |
| 35 | HOXC13   | 0.014467 | 0.309212 |
| 36 | NACC2    | 0.014477 | 0.309185 |
| 37 | SCRIB    | 0.014518 | 0.30906  |
| 38 | RASGRP1  | 0.014539 | 0.308997 |
| 39 | BAZ2B    | 0.014548 | 0.30897  |
| 40 | NDST1    | 0.014575 | 0.30889  |
| 41 | ZKSCAN1  | 0.014622 | 0.308752 |
| 42 | BEST2    | 0.014656 | 0.30865  |
| 43 | PNLIPRP3 | 0.014694 | 0.308539 |
| 44 | SMAD7    | 0.014717 | 0.308468 |
| 45 | AMMECR1L | 0.014735 | 0.308417 |
| 46 | RSP03    | 0.014882 | 0.307985 |
| 47 | KCTD9    | 0.01489  | 0.307962 |
| 48 | HERC4    | 0.014986 | 0.307683 |
| 49 | ABCA10   | 0.014992 | 0.307666 |
| 50 | ATP1B3   | 0.015027 | 0.307563 |
| 51 | GAB1     | 0.015067 | 0.307447 |
| 52 | ATP8B3   | 0.015089 | 0.307384 |
| 53 | PURA     | 0.015185 | 0.307107 |
| 54 | GK5      | 0.015222 | 0.307    |
| 55 | FLG      | 0.015312 | 0.306744 |

|    |          |          |          |
|----|----------|----------|----------|
| 1  |          |          |          |
| 2  | OXR1     | 0.015348 | 0.306641 |
| 3  | RGS6     | 0.015435 | 0.306393 |
| 4  | ESCO1    | 0.015457 | 0.306331 |
| 5  | AP1G1    | 0.015463 | 0.306313 |
| 6  | ERCC6    | 0.015469 | 0.306298 |
| 7  | CARD6    | 0.015471 | 0.306291 |
| 8  | PAQR6    | 0.015541 | 0.306094 |
| 9  | IGSF3    | 0.015614 | 0.305888 |
| 10 | EPAS1    | 0.015656 | 0.30577  |
| 11 | IER5     | 0.015735 | 0.30555  |
| 12 | ZMYND11  | 0.015772 | 0.305447 |
| 13 | FAM177A1 | 0.01588  | 0.305147 |
| 14 | PEX11A   | 0.015911 | 0.305063 |
| 15 | ZNF319   | 0.015989 | 0.304848 |
| 16 | S100A7   | 0.01603  | 0.304733 |
| 17 | SPRR2B   | 0.01604  | 0.304706 |
| 18 | VCPIP1   | 0.016145 | 0.304419 |
| 19 | LCE3D    | 0.016163 | 0.30437  |
| 20 | PAPLN    | 0.016166 | 0.304363 |
| 21 | HS3ST6   | 0.016261 | 0.304104 |
| 22 | BLZF1    | 0.016287 | 0.304033 |
| 23 | NCK1     | 0.016306 | 0.303983 |
| 24 | STX16    | 0.016309 | 0.303975 |
| 25 | GAS1     | 0.016402 | 0.303725 |
| 26 | SRPX2    | 0.016427 | 0.303657 |
| 27 | UBE2Q2   | 0.016477 | 0.303523 |
| 28 | GTF3C3   | 0.016479 | 0.303518 |
| 29 | SCYL2    | 0.016487 | 0.303495 |
| 30 | ZNF710   | 0.016538 | 0.303359 |
| 31 | PITPNM3  | 0.016581 | 0.303244 |
| 32 | RNASEL   | 0.016645 | 0.303075 |
| 33 | UBXN2A   | 0.016726 | 0.30286  |
| 34 | TNKS     | 0.016739 | 0.302825 |
| 35 | ZNF654   | 0.016754 | 0.302785 |
| 36 | BMP10    | 0.016802 | 0.302658 |
| 37 | TNFAIP2  | 0.016819 | 0.302614 |
| 38 | ZDHHC21  | 0.016834 | 0.302574 |
| 39 | NRBF2    | 0.016934 | 0.302313 |
| 40 | PI4K2A   | 0.016988 | 0.30217  |
| 41 | PMS1     | 0.017038 | 0.302042 |
| 42 | ZBTB43   | 0.017066 | 0.301968 |
| 43 | A2ML1    | 0.017125 | 0.301816 |
| 44 | RHOV     | 0.01719  | 0.301646 |
| 45 | LCE1B    | 0.017197 | 0.301628 |
| 46 | ASH1L    | 0.017239 | 0.30152  |
| 47 | CSNK1G1  | 0.01725  | 0.301491 |
| 48 | OR2G2    | 0.017301 | 0.301362 |
| 49 | TMC01    | 0.017337 | 0.301267 |
| 50 | MLLT10   | 0.017348 | 0.301239 |
| 51 | CRCP     | 0.017439 | 0.301006 |
| 52 | KIF16B   | 0.017454 | 0.300968 |
| 53 | OR52E2   | 0.017471 | 0.300927 |
| 54 | GTF2IRD1 | 0.017626 | 0.300531 |
| 55 | HERC3    | 0.017649 | 0.300473 |

ARPP19 0.017828 0.300024

For Peer Review