

Original Paper

Na⁺-ATPase, (Na⁺ + K⁺)-ATPase and NaPi-II Crosstalk in Ovarian Cancer Cells: Potential Role in Proliferation and Survival

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Key Words

SLC34A2 • NaPi-IIb • Na⁺-ATPase • (Na⁺ + K⁺)-ATPase • Epithelial ovarian cancer

Abstract

Background/Aims: Epithelial ovarian cancer (EOC) is the second leading cause of death among malignant neoplasms of the female reproductive system. In the U.S., approximately 20,890 new EOC cases and 12,730 EOC-related deaths are projected for 2025. Currently, there is no sufficiently accurate screening test for the early detection of EOC, nor have any specific or sensitive EOC markers proven effective in identifying the disease at its initial and potentially curable stages. **Methods:** EOC cell lines (A2780, ACRP, ES-2) and the non-tumoral IOSE cell line were analyzed using functional assays, including ATPase activity and phosphate uptake measurements, western blotting, and assays for proliferation, apoptosis, and motility. Pharmacological inhibitors were employed to assess the involvement of the PI3K/Akt/mTOR and MEK/ERK signaling pathways in the regulation of these transporters. **Results:** NaPi-IIb and Na⁺-ATPase hyperactivity were associated with increased proliferation, whereas only Na⁺-ATPase supported the survival and motility of EOC cells. In contrast, (Na⁺ + K⁺)-ATPase activity was not significantly implicated in EOC progression. Importantly, the PI3K/Akt/mTOR and MEK/ERK pathways modulated these effects in a lineage-dependent manner. **Conclusion:** The functional cooperation between NaPi-IIb and Na⁺-ATPase, regulated by major oncogenic pathways, contributes to proliferation, survival, and motility in EOC. These findings reveal a mechanism underlying ovarian cancer progression and highlight NaPi-IIb and Na⁺-ATPase as promising therapeutic targets for the development of more effective and selective treatments.

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Introduction

Epithelial ovarian cancer (EOC) is the second most common cause of death among neoplasms of the female reproductive system [1]. In the U.S., an estimated 20,890 new EOC cases and 12,730 EOC-related deaths are expected in 2025 [2]. Currently, there is no sufficiently accurate screening test for the early detection of EOC, nor have any specific or sensitive EOC markers proven effective in identifying the disease at its initial and potentially curable stages. Consequently, approximately 70% of EOC cases are diagnosed at advanced stages, when metastatic tumor cells have acquired a drug-resistant phenotype, and the 5-year survival rate drops to about 25% (compared with 80% for localized disease) [3]. SLC34A2, which encodes NaPi-IIb, has been shown to be overexpressed across all EOC histological subtypes compared with normal ovarian surface epithelium; therefore, NaPi-IIb has emerged as a promising candidate biomarker for EOC [4, 5].

Despite substantial evidence supporting the clinical significance of targeting NaPi-IIb to combat EOC (for example, using the MX35 antibody) [5, 6, 7, 8, 9, 10], its regulation and function in ovarian cells remain unclear. One proposed role of NaPi-IIb in the ovary is the maintenance of folate levels essential for oocyte maturation, a theory supported by the expression of NaPi-IIb in the epithelial cells lining the fallopian tubes [11]. NaPi-IIb is a secondary active transporter that utilizes the sodium electrochemical gradient to import phosphate from the extracellular space into the cell cytoplasm. In relation to the establishment and maintenance of this sodium electrochemical gradient in epithelial cells by primary active transporters, the classical ouabain-sensitive (Na⁺+K⁺)-ATPase and a second Na⁺-ATPase—insensitive to ouabain, sensitive to furosemide, and previously cloned in enterocytes—are critical. Notably, the expression of Na⁺-ATPase in ovarian cells has not yet been investigated [12].

This study aimed to investigate the expression of NaPi-IIb and its correlation with the two sodium pumps in EOC cells. Additionally, the potential roles of these transporters in EOC were examined. Herein, we present novel findings indicating that Na⁺-ATPase and NaPi-IIb, but not (Na⁺+K⁺)-ATPase, are modulated during EOC progression. Furthermore, Na⁺-ATPase appears to coordinate with NaPi-II to enhance cellular proliferation in EOC. In contrast, only Na⁺-ATPase promotes the survival and motility of EOC cells.

Materials and Methods

Reagents

ATP (sodium salt), ouabain, furosemide, phosphonoformic acid (PFA), sodium chloride, potassium chloride, magnesium chloride, EDTA, EGTA, Hepes, Tris, sodium deoxycholate and sodium orthovanadate were purchased from Sigma-Aldrich. Sodium pyrophosphate was purchased from Reagen. SDS was purchased from GE Healthcare. Sucrose was from Merck. Wortmannin was purchased from Calbiochem. U0126, polyclonal phospho-PKB (Ser-473), polyclonal PKB, polyclonal phospho-ERK (Thr-202/Tyr-204), polyclonal ERK, polyclonal GAPDH and polyclonal β -actin antibodies were from Cell Signaling Technology. Anti-NPT-2b was from Santa Cruz Biotechnology. ³²Pi was obtained from the Brazilian Institute of Energetic and Nuclear Research (São Paulo, Brazil). γ -³²Pi-ATP was synthesized according to the procedures described by Maia et al. [13]. All other reagents were of the highest purity available.

Cell culture

Human EOC cells A2780, ACRP and ES-2 were kindly provided by Dr. Patrice J. Morin, NIH, USA, and maintained in RPMI 1640 medium (Sigma-Aldrich) supplemented with 10% fetal bovine serum (FBS, Gibco/Invitrogen) and stabilized solution of penicillin (100 un/mL), streptomycin (100 μ g/mL) and amphotericin B (250 μ g/mL) (Gibco/Invitrogen). Immortalized cell line IOSE was kindly provided by Dr. Nelly Auersperg of the Canadian Ovarian Tissue Bank (University of British Columbia, Vancouver, Canada), and cultured in 1:1 mixture of medium 199 and MCDB 105 (Sigma-Aldrich), supplemented with 10% FBS, L-Glutamine (5 mM), and stabilized solution of penicillin (100 un/ml), streptomycin (100 μ g/ml) and amphotericin B

(250 ug/ml). Cells were grown at 37°C in a humidified atmosphere of 95% air and 5% CO₂. Cells were preincubated with different compounds indicated in the in serum-depleted medium. A2780 is an ovarian carcinoma cell line sensitive to cisplatin, established from tumor tissue from an untreated patient. ACRP is a cell line originated from the intermittent treatment of A2780 with cisplatin, and it was determined that the cisplatin IC₅₀ for A2780 and ACRP was 0.75 uM and 3 uM, respectively [14]. ES-2 is a clear cell ovarian carcinoma (CCOC) that exhibits low to moderate resistance to various chemotherapeutic agents such as doxorubicin, cisplatin, carmustine and etoposide. IOSE is an immortalized cell line with epithelial ovarian cell characteristics without tumorigenic function in Nu/nu mice [15].

Antibodies

All primary antibodies were rabbit IgG and purchased from Cell Signaling Technology®. Primary antibodies were used at a 1:1000 dilution and incubated overnight. The phosphorylated secondary antibodies were applied at a 1:4000 dilution, while the total ones were used at a 1:5000 dilution. All secondary antibodies were incubated for 1 hour.

Measurement of (Na⁺+K⁺)-ATPase and Na⁺-ATPase activities

ATPase activity was measured according to the method described by Grubmeyer and Penefsky [16]. Briefly, treated cells were washed in PBS, harvested, and cleared by centrifugation for 5 min at 600 x g, and incubated for 30 min in lysis buffer (1 mM EGTA, 0.1% sodium deoxycholate, 20 mM Hepes (pH 7.0), 250 mM sucrose). The composition of the standard assay medium was 10 mM MgCl₂, 8.3 mM ATP (specific activity 90mCi/mmol [γ -³²P] ATP), 33 mM HEPES (pH 7.0), 120 mM NaCl, and 50 mM KCl. The reaction was started by the addition of cell lysate. After 10 min, the reaction was stopped with charcoal activated with 0.1 N HCl. The ³²Pi released to the medium was measured in liquid scintillation counter (Packard Tri-Carb 2100 TR). (Na⁺+K⁺)-ATPase activity was determined by its sensitivity to 1 mM ouabain and expressed in nanomoles of Pi per milligram of protein per minute. Na⁺-ATPase activity was evaluated in the presence of 1 mM ouabain and in the absence of KCl added to the medium and determined by its sensitivity to 2 mM furosemide and expressed in nanomoles of Pi per milligram of protein per minute. Protein quantification was determined by the Folin phenol method, using BSA as standard.

Measurement of sodium-dependent phosphate uptake in EOC cells

Sodium-dependent phosphate uptake was measured by the uptake of radiolabeled phosphate into EOC cell monolayers. Cells were pretreated with the indicated conditions in serum-free RPMI medium. Measurement of phosphate uptake was initiated by the addition of transport medium containing NaCl 140 mM, KCl 5 mM, MgCl₂ 1 mM, CaCl₂ 1.5 mM, HEPES 15 mM (pH 7.4), Na₂HPO₄/NaH₂PO₄.H₂O 0.1 mM (pH 7.4) and ³²Pi (1 mCi/mmmole), which allowed the analysis of the total phosphate transport. Another transport buffer containing choline chloride instead of NaCl was used to measure Na⁺-independent Pi uptake. Uptake was stopped after 10 min by aspiration of the transport medium and washing the cells with ice-cold PBS. Cell monolayers were solubilized by the addition of 1% SDS and the radioactivity was counted in liquid scintillation counter (Packard Tri-Carb 2100 TR). Absolute values of phosphate uptake were calculated. Na⁺-dependent Pi transport was calculated by subtraction of the Na⁺-independent component from the total Pi uptake in the presence of Na⁺ and expressed in nanomoles of Pi per milligram of protein per minute. Protein quantification was determined by the Folin phenol method, using BSA as standard.

Western Blot

The cells were washed with PBS 1x, harvested, and incubated for 40 min in lysis buffer (25 mM Tris (pH 7.5), 50 mM NaCl, 1 mM EDTA, 1% Triton X-100, 0.5% sodium deoxycholate, 10 mM SDS, 10 mM NaF, 5 mM sodium vanadate, 5 mM sodium pyrophosphate, protease inhibitor mixture (Sigma-Aldrich) and cleared by centrifugation for 10 min at 15,000 x g. The supernatant was retained, and the protein concentrations were determined by the Folin phenol method, using BSA as standard. Proteins were resolved in SDS-polyacrylamide gels and transferred to polyvinylidene fluoride membranes (Millipore), according to the manufacturer's instructions. After antibody labeling, detection was acquired with ECL-prime (GE Amersham Biosciences).

Phagokinetic Track Motility Assay

The phagokinetic track motility assay is based on the ability of a moving cell to clear gold particles from its path to create a measurable track on a colloidal gold-coated plate [17]. Briefly, 24-well plates were coated with the colloidal gold particles. Cells were trypsinized and resuspended in culture medium at a density of 2×10^3 cells/ml, and 0.5 milliliter of the cell suspension, treated or not, were added to each gold particle-coated well. After incubation at 37°C in 5% CO₂ for 16 h, the tracks created by single migrating cells were visible and photographs under 10X magnification were taken using a Leica inverted phase-contrast microscope (Leica DM IL LED, Leica Microsystems) equipped with Quick Imaging system. Cell motility was evaluated by measuring the areas free of the gold particles using the ImageJ software (version 1.6.0). At least 20 cells per condition were analyzed in three independent experiments.

CFSE assay

Cell proliferation was analyzed using a carboxyfluorescein succinimidyl ester (CFSE) kit (CellTrace™ CFSE Cell Proliferation Kit, Invitrogen) according to the manufacturer's instructions. Briefly, cells were stained with CFSE and seeded in 12-wells plates for 24 hours. Following, cells from some wells were harvested with trypsin-EDTA (0.25 mg/ml trypsin and 1 mg/ml EDTA), centrifuged at 2000 x g for 5 minutes, resuspended in culture medium and analyzed by flow cytometry (BD FACScan) to evaluate the binding intensity with CFSE to the cells at the beginning of each described treatment. The remaining cells were treated with furosemide (2.0 mM) or vehicle (0.4% NaOH) or PFA (1.0 mM) or only with culture medium and cultured for 24 hours. After 24 hours of treatment, cells were harvested with trypsin-EDTA, centrifuged, and resuspended in culture medium, and then analyzed on a flow cytometer. The results were analyzed by FlowJo software (version 7.6.2) and expressed as a percentage of divided cells. The experiments were performed in triplicate.

Apoptosis assay

For determination of cell death by apoptosis, 2×10^5 cells were seeded in 12-well culture plates. After 48 hours, cells were treated with furosemide (2.0 mM) or vehicle (0.4% NaOH) or PFA (1.0 mM) or with culture medium alone and incubated at 37°C under 5% CO₂ for 24 hours. The supernatant was collected (for evaluation of the cells in suspension) and the adherent cells were gently collected with PBS-EDTA (0.5 mM). Samples were then centrifuged, resuspended in binding buffer (Annexin V-FITC Kit, BD, USA), incubated for 15 minutes in the dark at room temperature with Annexin V-FITC, then incubated with propidium iodide and analyzed immediately by flow cytometry. Data were analyzed using the FlowJo software (version 7.6.2). The experiments were run in triplicate.

Statistical Analysis

All statistical analyses were performed using GraphPad Prism 9 (GraphPad Software, San Diego, CA, USA). Results are presented as the mean \pm SEM. For comparisons between two groups, an unpaired Student's t-test was performed. Comparisons among three or more groups were conducted using one-way ANOVA followed by Bonferroni post hoc test, when applicable. Statistical significance was defined as $p < 0.05$.

Results

Expression profile of NaPi-IIb in EOC cells

Our data proved that all tested ovarian cancer cells express NaPi-IIb in higher magnitude than immortalized non-tumoral normal epithelial ovarian cells (IOSE) cells. Amongst the EOC cells, A2780 and ACRP express the transporter two-fold more than ES-2 ($p < 0.05$) (Fig. 1). Altogether, our results suggest that serous ovarian carcinoma (SOC) overexpresses NaPi-IIb in a higher magnitude when compared to CCOC. Even though these are both low prognostic diseases, mainly due to late diagnosis and chemoresistance, it is worthwhile to mention that the later occur much more frequently than the later.

Sodium dependent phosphate uptake in EOC cells

Motivated by the fact that EOC cells overexpress NaPi-IIb compared to the normal counterpart IOSE cells, we argued whereas there was transporter activity of sodium dependent phosphate uptake sensitive to phosphonoformate acid (PFA), a selective inhibitor of NaPi-II. Sodium dependent phosphate uptake sensitive to PFA increased in a dose dependent manner in all lineages studied, as ACRP=A2780>ES-2>IOSE (Fig. 2) ($p<0.05$). These findings corroborate to our previous data concerning the expression of the protein NaPi-IIb (Fig. 1). The kinetic parameters of the sodium dependent phosphate transporter were calculated following the equation $v=V_{max} [S]/K_{0.5}+[S]$ and are summarized in Table 1. V_{max} was higher in ACRP and in A2780, (0.3514 ± 0.0264 and 0.2701 ± 0.0153 nmol Pi x mg⁻¹ x min⁻¹, respectively) than in ES-2 (0.2035 ± 0.0225 nmol Pi x mg⁻¹ x min⁻¹) ($p<0.05$). On the other hand, IOSE had higher V_{max} than EOC cells (0.0812 ± 0.0220 nmol Pi x mg⁻¹ x min⁻¹) ($p<0, 005$). The differences observed regarding NaPi-II kinetic parameters in ovarian cells lead us to investigate the regulation and the function of NaPi-II in EOC.

ATPase and Na⁺ATPase

NaPi-IIb is a secondary active transporter, thus depends on a primary active transporter, such as (Na⁺+K⁺)-ATPase, to function properly. As shown in Fig. 3, preliminary results (N=2) suggest that (Na⁺+K⁺)-ATPase activity may partially modulate NaPi-II activity in the presence of sodium [140 mM] or choline [140 mM] in EOC cells. Although statistical analysis could not be performed due to the limited number of replicates, an approximately 50% decrease in NaPi-II activity was observed following 30 min treatment of ovarian cells with ouabain, supporting a potential functional link between these transporters that warrants further investigation.

We further questioned the proportional contribution of (Na⁺+K⁺)-ATPase and Na⁺-ATPase on ovarian malignant transformation, and got interest results: i) the ratio of the activities of (Na⁺+K⁺)-ATPase and Na⁺-ATPase was 3.81-, 1.92-, 2.14-, and 1.03-fold in IOSE, ES-2, ACRP, and A2780, respectively (Figure 4) ($p<0.05$); ii) the activity of (Na⁺+K⁺)-ATPase did not influence ovarian carcinogenesis, despite the fact that its activity was sustained in

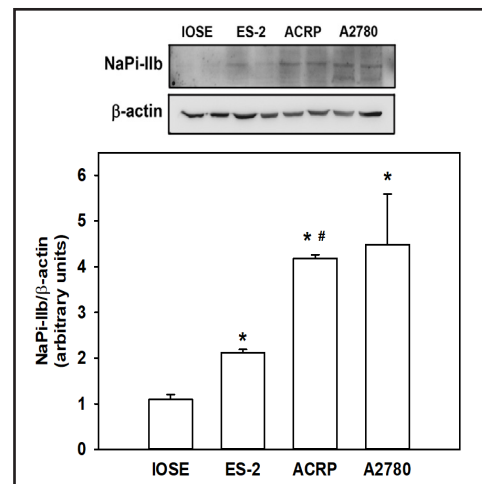


Fig. 1. NaPi-IIb expression was analyzed in EOC cells: A2780 and ACRP, both SOC; ES-2, CCOC; IOSE, immortalized non-tumoral normal epithelial ovarian cells. N=3; * $p<0.05$ compared to IOSE. # $p<0.05$ compared to ES-2.

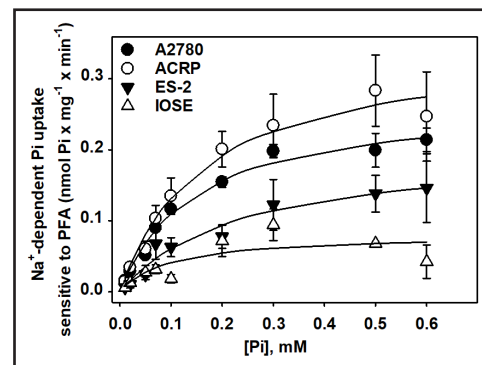


Fig. 2. Sodium dependent phosphate uptake, sensitive to PFA, in ovarian cell lines. The experiment was conducted in a dose dependent manner, with phosphate varying from 0.01 to 0.6 mM ³²Pi (1 mCi/mmol Pi) was used as a tracer, and the protein activity expressed in nmol Pi x min⁻¹ x mg⁻¹. N=4.

Lineage	V_{max} (nmol Pi x mg ⁻¹ x min ⁻¹)	$K_{0.5}$ (mM Pi)
IOSE	0,0812 ± 0,0220	0,0981 ± 0,0823
ES-2	0,2035 ± 0,0225 *	0,2354 ± 0,0684
ACRP	0,3514 ± 0,0264 *#	0,1677 ± 0,0328
A2780	0,2701 ± 0,0153 *#§	0,147 ± 0,0227

Table 1. Kinetic parameters of the sodium dependent phosphate uptake, sensitive to PFA, in ovarian cells. * $p<0,05$ compared to IOSE; # $p<0,05$ in comparison with ES-2; § $p<0,05$ in relation to ACRP

Fig. 3. Sodium dependent phosphate uptake in EOC cells: A2780, ACRP, and ES-2 cells were treated with ouabain 1 mM, and NaPi-II was evaluated in a time-course fashion in these cells. ³²Pi (1 mCi/mmol Pi) was used as a reaction tracer. Results were expressed as percentage of the sodium phosphate caption. N=2.

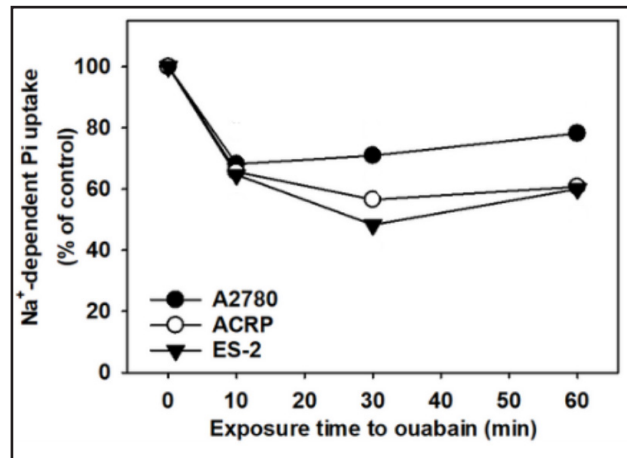
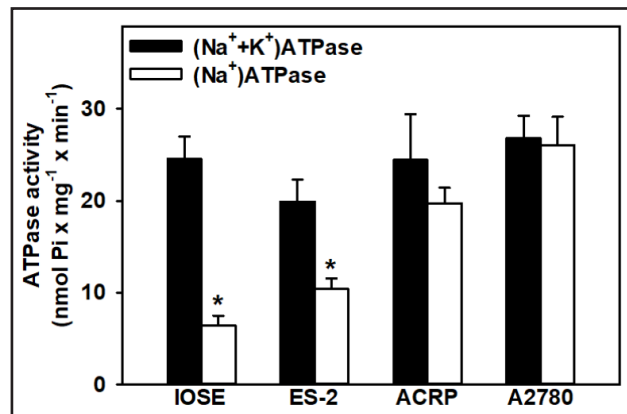


Fig. 4. Correlation between Na⁺-ATPase and (Na⁺+K⁺)-ATPase activities in ovarian lineages (IOSE, ES-2, ACRP and A2780). N=4. *p<0.05 in comparison with the (Na⁺+K⁺)-ATPase activity within the same lineage.



aggressive tumors, therefore it might contribute to the phenomenon; iii) Na⁺-ATPase was over-activated in EOC cells vs. IOSE cells; iv) whereas (Na⁺+K⁺)-ATPase was the same in EOC vs. IOSE cells, Na⁺-ATPase and NaPi-II profiles were similar in the ovary. One possibility is that Na⁺-ATPase might provide the fine sodium gradient that is probable required for the higher activity of NaPi-II in EOC, especially in SOC (the most frequent and highly aggressive form of the disease) (Fig. 5). We then postulated that Na⁺-ATPase would modulate NaPi-II sensitive to PFA. We observed that furosemide inhibited the sodium dependent phosphate uptake in a dose dependent manner, the maximum effect being at 2.0 mM. Under this experimental condition, NaPi-II decreased from 0.264 ± 0.027 without furosemide to 0.135 ± 0.029 nmol Pi x mg⁻¹ x min⁻¹ in the presence of furosemide in A2780; from 0.244 ± 0.027 (in the absence of furosemide) to 0.121 ± 0.006 nmol Pi x mg⁻¹ x min⁻¹ (in the presence of furosemide) in ACRP; from 0.169 ± 0.007 (without furosemide) to 0.091 ± 0.007 nmol Pi x mg⁻¹ x min⁻¹ (with furosemide) (Fig. 5) (p<0.001 for all cases). Taking in conjunction, our results strongly support the correlation between the activity of NaPi-II and Na⁺-ATPase in ovarian cells.

Na⁺-ATPase activity insensitive to ouabain and sensitive to furosemide, and of NaPi-II sensitive to PFA modulate ovarian cells motility and survival

Cell motility, measured as arbitrary unit, was reduced in 31% (p<0.001), 40% (p<0.0001), 50% (p<0.0001) and 35% (p<0.0005) in IOSE, ES-2, ACRP and A2780 by furosemide (2.0 mM), respectively. In turn, motility was decreased in IOSE by PFA (1.0 mM) in 17% (p<0.05) (Fig. 6). Our findings suggest that NaPi-II is unlikely to regulate the motility of EOC cells, being its effect restricted to IOSE. On the other hand, Na⁺-ATPase contributes, at least partially, to EOC cells' motility (Fig. 7), which is a key phenomenon for cancer cells' metastasis. Next, we argued whether NaPi-II, sensitive to PFA and Na⁺-ATPase, sensitive to furosemide

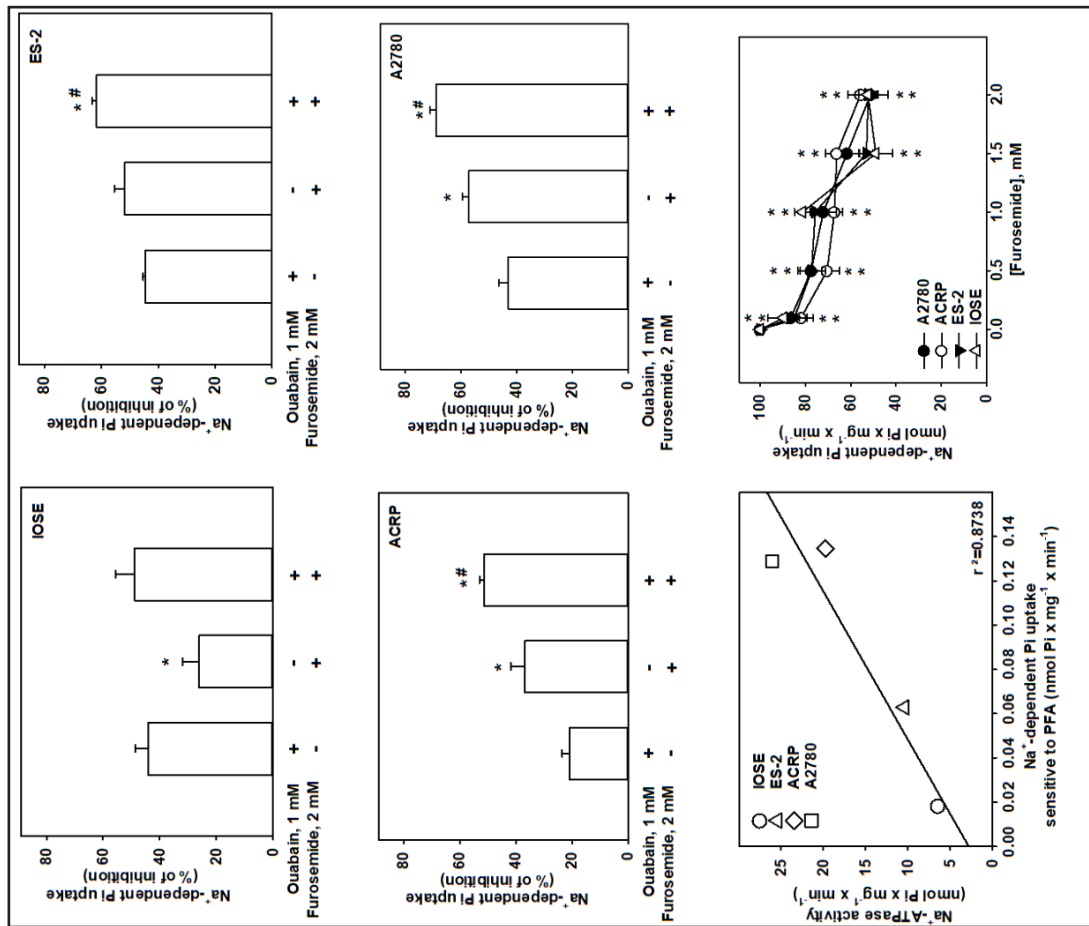
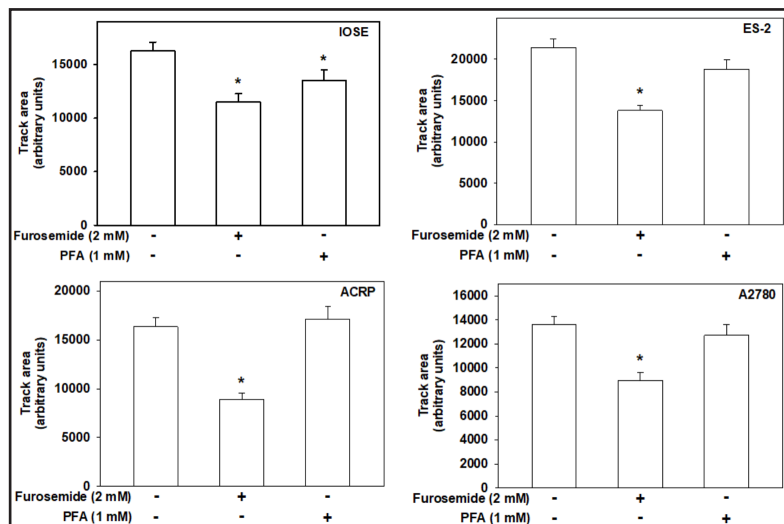


Fig. 5. Modulation of sodium dependent phosphate uptake in ovarian cells modulated by (Na⁺+K⁺)-ATPase and Na⁺-ATPase. (A-D) Ouabain 1.0 mM and Furosemide 2.0 mM were added as shown in the Fig. ³²Pi (1 mCi/mmol Pi) was used as a tracer. Enzyme activity was expressed as percentage in relation to control. N=6. *p< 0,01 in comparison with cells treated with ouabain or furosemide#p<0,05. (E) Fig. shows the correlation between Na⁺-ATPase and the sodium dependent phosphate uptake, sensitive to PFA, in ovarian cells. N=4. $r^2=0.8738$. (F) The Fig. demonstrates the positive effect of Na⁺-ATPase on NaPi-II sensitive to PFA. ³²Pi (1 mCi/mmol Pi) was used as a tracer. Data were expressed as a percentage of control. N=6. *p< 0.05 in comparison with non-treated cells.

Fig. 6. Colloidal gold assay was used to analyze ovarian cells' motility. Cells were plated into wells covered with a thin gold layer, then treated with furosemide (2.0 mM), vehicle (NaOH 0.4%) or PFA (1.0 mM) for 16 hours. While the furosemide-treated cells decreased the motility of all ovarian cells evaluated, PFA treatment comprised the same effect exclusively in IOSE. *p<0.05 in comparison with non-treated cells.



and insensitive to ouabain, would play a role in ovarian cell proliferation. Therefore, we conducted the carboxyfluorescein succinimidyl ester incorporation assay (CFSE). Cells were labelled with CFSE, and then treated with furosemide (2.0 mM), vehicle (NaOH 0.4%) or PFA (1.0 mM). Proliferative cells were detected by flow cytometry. Furosemide 2.0 mM reduced the ovarian cellular proliferation by 17% ($p < 0.05$), 45% ($p < 0.001$), 28% ($p < 0.01$) and 46% ($p < 0.001$) in IOSE, ES-2, ACRP and A2780, respectively. On the other hand, PFA reduced ovarian cell proliferation in 38% ($p < 0.05$), 40% ($p < 0.001$), 55% ($p < 0.01$) and 62% ($p < 0.001$) in IOSE, ES-2, ACRP and A2780, respectively (Fig. 8). Of note, the proliferation index of IOSE was significantly lower than the EOC cells. Therefore, our results suggest that whereas NaPi-II modulates the proliferative index of normal and cancerous ovarian cells, Na⁺ATPase induces the proliferation of cancerous cells exclusively. In conjunction, our data strongly suggest that whereas NaPi-II regulates the proliferation of normal and cancerous ovarian cells, Na⁺ATPase likely increases the proliferative index of EOC cells more prominently than normal ovarian cells.

Evaluation of the role of both transporters activity in ovarian cells survival was assessed by the annexin V/Pi analysis assay using flow cytometry. Our results show that furosemide (2.0 mM) increased the percentage of apoptotic cells, 2.4-1.8- ($p < 0.0001$); 2.0- ($p < 0.0001$) and 2.0-fold ($p < 0.0001$), compared to control, in IOSE (Fig. 9), ES-2 (Fig. 10), ACRP (Fig. 11) e A2780 (Fig. 12), respectively. On the other side, PFA increased the percentage of apoptotic cells exclusively in ES-2 (1.6-fold; $p < 0.0001$), as shown in Fig. 10.

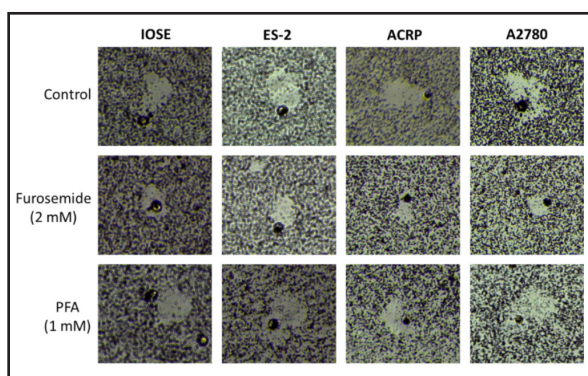


Fig. 7. Role of NaPi-II and Na⁺-ATPase in ovarian cells' motility. The colloidal gold motility assay was used to evaluate ovarian cells' motility. Cells were photographed, and the area displaced by the cells was quantified by the Image J Software. N = 3.

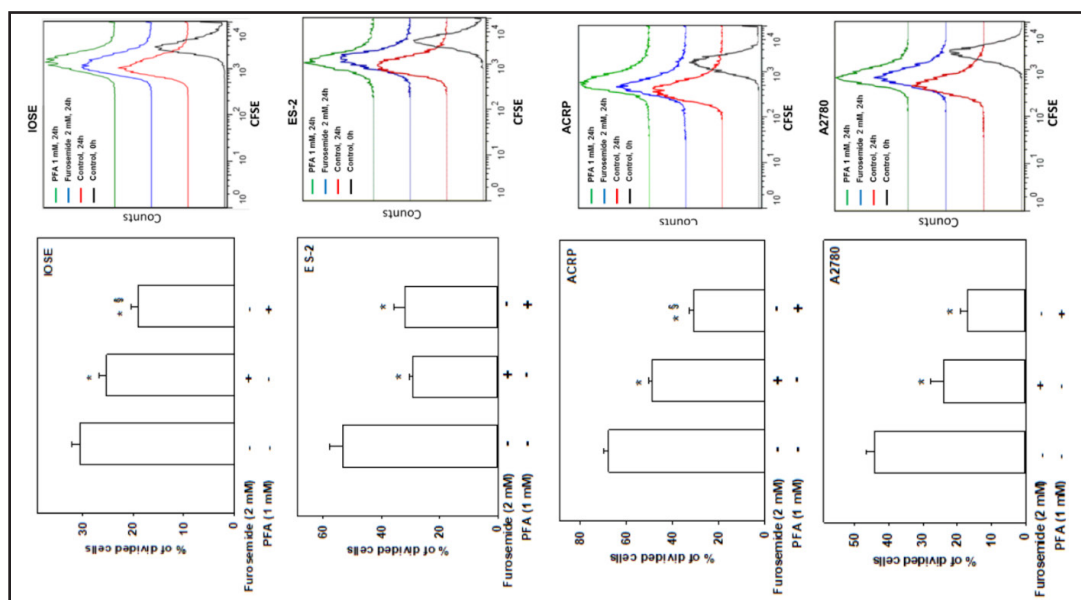


Fig. 8. Effect of NaPi-II and Na⁺-ATPase on ovarian cell proliferation. Cell proliferation was assessed using the carboxyfluorescein succinimidyl ester (CFSE) incorporation assay and analyzed by flow cytometry. The bar graphs represent the mean \pm SD of three independent replicates (N=3), while the line graph illustrates a representative CFSE experiment. Cells were treated with furosemide (2.0 mM) or PFA (1.0 mM) to evaluate the roles of Na⁺-ATPase and NaPi-II in ovarian cells, respectively. * $p < 0.01$ compared to non-treated cells; $\$p < 0.05$ compared to cells treated with furosemide.

Fig. 9. Evaluation of the role of the NaPi-II cotransporter and Na⁺-ATPase in ovarian epithelial cells' death. Non-cancerous ovarian IOSE cells were seeded and, after 48 hours, treated for 24 hours with furosemide (2,0 mM) or PFA (1,0 mM). Following treatment, cells were stained with annexin V-FITC and propidium iodide (PI) to assess the percentage of apoptotic (annexin V-positive/PI-negative) and necrotic (annexin V-positive/PI-positive) cells. N = 6. *p<0.01 compared to untreated cells.

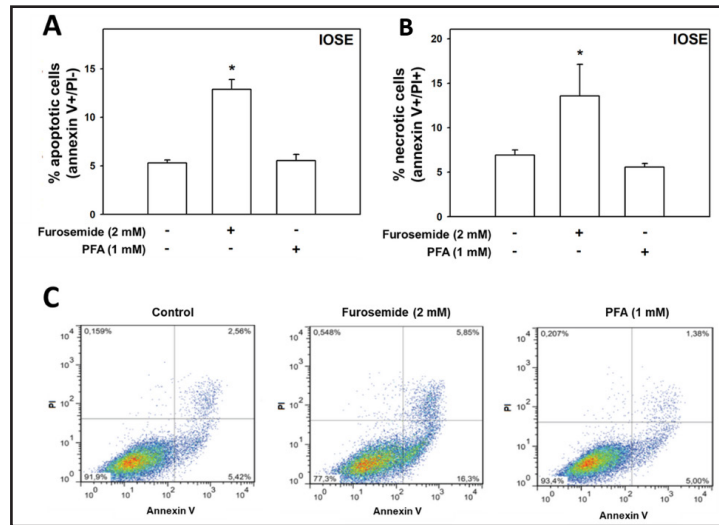


Fig. 10. Evaluation of the role of the NaPi-II cotransporter and Na⁺-ATPase in ovarian epithelial cells' death. EOC ES-2 cells were seeded and, after 48 hours, treated for 24 hours with furosemide (2,0 mM) or PFA (1,0 mM). Following treatment, cells were stained with annexin V-FITC and propidium iodide (PI) to assess the percentage of apoptotic (annexin V-positive/PI-negative) and necrotic (annexin V-positive/PI-positive) cells. N = 6. *p<0.01 compared to untreated cells.

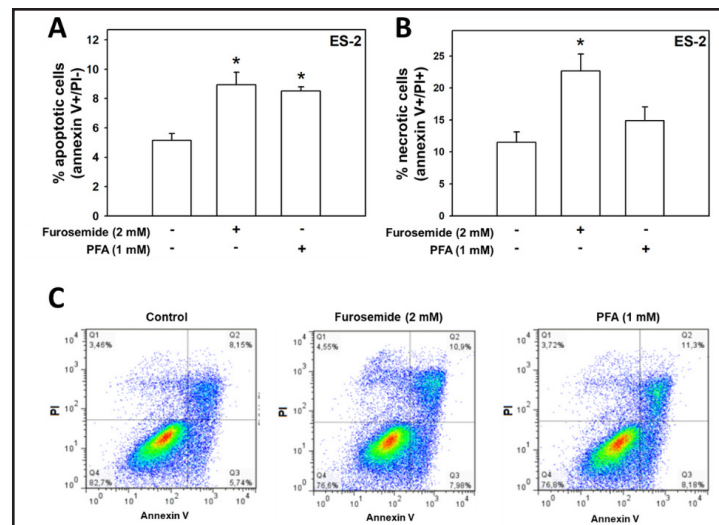


Fig. 11. Evaluation of the role of the NaPi-II cotransporter and Na⁺-ATPase in ovarian epithelial cells' death. EOC ACRP cells were seeded and, after 48 hours, treated for 24 hours with furosemide (2,0 mM) or PFA (1,0 mM). Following treatment, cells were stained with annexin V-FITC and propidium iodide (PI) to assess the percentage of apoptotic (annexin V-positive/PI-negative) and necrotic (annexin V-positive/PI-positive) cells. N = 6. *p<0.01 compared to untreated cells.

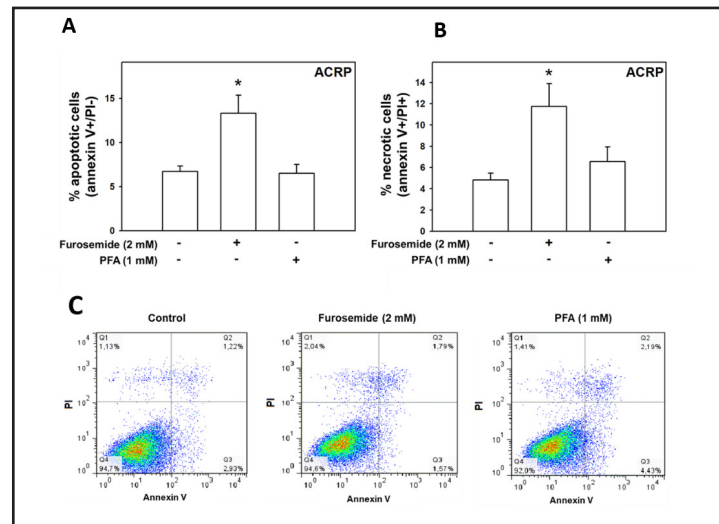
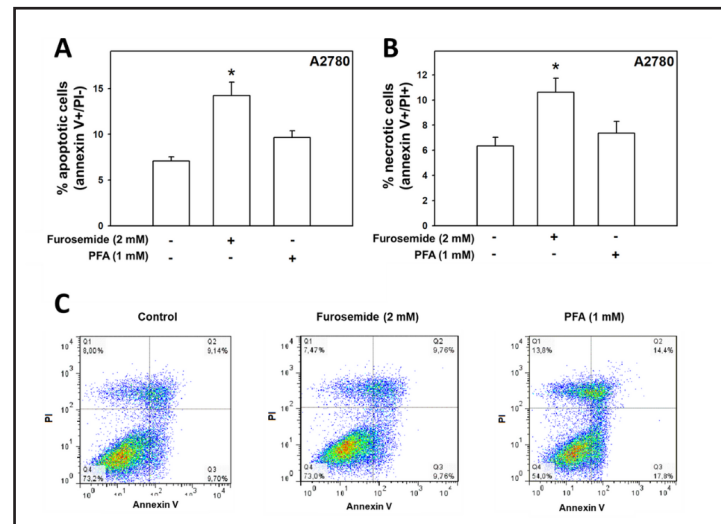


Fig. 12. Evaluation of the role of the NaPi-II cotransporter and Na⁺-ATPase in ovarian epithelial cells' death. EOC A2780 cells were seeded and, after 48 hours, treated for 24 hours with furosemide (2,0 mM) or PFA (1,0 mM). Following treatment, cells were stained with annexin V-FITC and propidium iodide (PI) to assess the percentage of apoptotic (annexin V-positive/PI-negative) and necrotic (annexin V-positive/PI-positive) cells. N = 6. *p<0.01 compared to untreated cells.



MEK/ERK and PI3K/AKT/mTOR signalling pathways regulate NaPi-II in ovarian cells

Considering the key role of MAPK (MEK/ERK) and PI3K/AKT/mTOR signalling pathways in cancer, we decided to investigate their status in ovarian cells, as well as their role in NaPi-II and Na⁺-ATPase activity modulation. Protein expression was evaluated by Western blot, and data are compiled in Fig. 13. MAPK pathway was differentially expressed in EOC cells compared to IOSE cells. Indeed, the activation of MAPK signalling pathway was more intense in EOC vs. IOSE cells. Amongst EOC cells, however, the observation was ACRP>A2780>ES-2. On the other hand, PI3K pathway was activated in ACRP and A2780 cells, which is known to have the protein constitutively activated, when compared with IOSE and ES-2, as previously observed by Hua et al, 2008 and 2009 [18, 19].

Considering the expression profile of the MAPK and the PI3K pathways in ovarian cells, we sought to investigate their effect on NaPi-II activity in the same cell lines studied. To do so, cells were cultured in the presence or in the absence of ERK inhibitor, U0126 (1.0 μM) or PI3K inhibitor, wortmannin (10⁻⁷M). We concluded that in all ovarian cells, NaPi-II activity was stimulated by the PI3K/AKT/mTOR pathway as the co-transporter activity decreased to 43% (p<0, 01), 27% (p<0, 01), 67% (p<0, 001) and 40% (p<0, 01) in IOSE, ES-2, ACRP e A2780, respectively, in the presence of wortmannin.

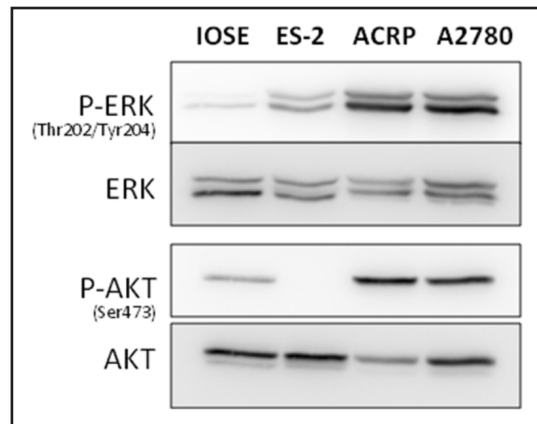


Fig. 13. Profiles of MAPK and PI3K pathways in EOC and IOSE cells by Western blot. N = 3.

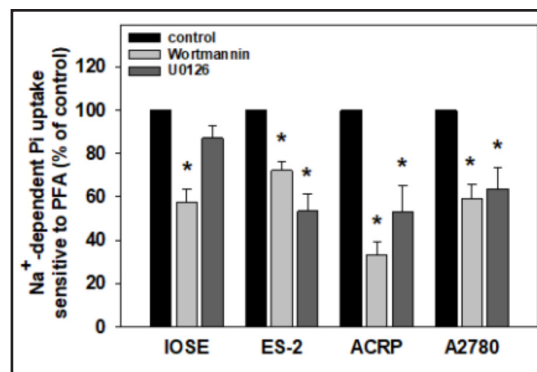


Fig. 14. Modulation of NaPi-II activity by MAPK and PI3K pathways in ovarian cells. Cells were treated for 2 hours with wortmannin (10⁻⁷ M) or U0126 (10 μM). Sodium-dependent phosphate uptake sensitive to PFA activity was measured using ³²Pi (1 mCi/mmol Pi) as a tracer and was expressed as percentage of control. N=3. * p<0.05 compared to the control.

On the other hand, the MAPK pathway is unlikely to modulate NaPi-II activity in all ovarian cells. In fact, U0126 reduced the activity of NaPi-IIb in 46% ($p < 0, 05$), 47% ($p < 0, 05$) and 36% ($p < 0, 05$) in ES-2, ACRP and A2780, respectively. Nonetheless, no significant effect was observed in IOSE cells. These results are shown in Fig. 14.

Discussion

Recent evidence highlights that phosphate metabolism is not merely a secondary biochemical process in malignancies but rather serves as a central regulatory axis essential for sustaining oncogenic growth and metabolic plasticity [20]. Elevated intracellular phosphate levels provide the necessary substrate for ATP synthesis, nucleotide production, and activation of key signalling kinases, including Akt, ERK, and mTOR, thereby supporting increased cell proliferation, biosynthetic activity, and metabolic reprogramming in cancer cells [21]. Moreover, dysregulated phosphate homeostasis has been linked to alterations in tumor immunity, extracellular matrix remodeling, and epithelial–mesenchymal transition (EMT), further underscoring phosphate as a multifaceted determinant of tumor progression and microenvironmental adaptation [22, 23].

Within this broader metabolic context, the NaPi family of sodium-dependent phosphate transporters (SLC34 and SLC20) has emerged as a key regulatory hub in cancer-associated phosphate homeostasis [24]. Among these, NaPi-IIb (SLC34A2) is notable for its pronounced overexpression in various epithelial malignancies, including ovarian, lung, colorectal, and gastric cancers, where it imparts proliferative and survival advantages to tumor cells [24]. In addition to mediating phosphate uptake, NaPi-IIb has attracted growing interest as both a diagnostic and therapeutic target. Several antibody–drug conjugates (ADCs) and monoclonal antibodies targeting NaPi-IIb are currently under investigation, showing promising preclinical efficacy and tumor selectivity with minimal toxicity in non-malignant tissues [25]. Conversely, the phosphate exporter XPR1 (SLC53A1) plays an equally important yet complementary role, regulating intracellular phosphate efflux and influencing tumor aggressiveness and metabolic balance [26]. This reciprocal regulation between NaPi-mediated import and XPR1-mediated export suggests that cancer cells dynamically adjust phosphate fluxes to meet anabolic demands while avoiding cytotoxic phosphate overload. Such a mechanism implies that distinct tumor types may utilize differential phosphate-handling strategies, determined by their metabolic context and signaling architecture [26].

Our present findings are consistent with this model, revealing a functional coupling between NaPi-IIb and the PI3K/Akt and MEK/ERK signaling cascades. These pathways are well established as key regulators of transporter expression and function, integrating nutrient sensing, growth factor responses, and metabolic control [27]. Consistent with previous reports in other tumor systems, our data indicate that NaPi-IIb activation is lineage-dependent and mechanistically linked to oncogenic kinase activity, further reinforcing its role as an adaptive metabolic element within EOC cells [4].

From a translational perspective, these results support the notion that targeting phosphate metabolism represents a promising yet underexplored therapeutic approach in EOC [26]. The tissue-restricted expression profile of NaPi-IIb, combined with its functional interdependence with PI3K and MEK signaling, underscores its potential for tumor-selective intervention [28, 29]. Furthermore, combination strategies utilizing phosphate transporter blockade in conjunction with PI3K or MEK inhibitors may yield synergistic antitumor effects by simultaneously disrupting the metabolic and signaling dependencies that sustain tumor viability and progression.

In this study, we investigated the role of the phosphate co-transporter NaPi-IIb (encoded by the SLC34A2 gene) and the Na⁺-ATPase enzyme in EOC cell lines, as well as their relationship with signaling pathways implicated in tumorigenesis, including PI3K/Akt/mTOR and MEK/ERK. The functional modulation of these proteins was assessed in key cellular processes, such as proliferation, motility, phosphate uptake, and cell viability.

Tumor cell lines A2780, ACRP (cisplatin-resistant), and ES-2 (CCOC) were utilized alongside the non-tumorigenic IOSE line to compare tumor and non-tumor expression and functional patterns of the proteins of interest.

Our findings confirm that NaPi-IIb is markedly overexpressed in EOC cell lines compared to the non-tumorigenic IOSE line [30, 31]. Notably, the study by Nurgalieva et al. (2021) further indicates that NaPi-IIb may serve as a potential biomarker in ovarian cancer patients. Its upregulation is functionally associated with a significant increase in phosphate (Pi) uptake, which was effectively inhibited by PFA. The enhanced Pi uptake observed in tumor cells was more pronounced and demonstrated a dose-dependent response to PFA, indicating increased transporter activity in the malignant context. These results strongly support a pivotal role for NaPi-IIb in sustaining the proliferative capacity of EOC cells. In agreement with our observations, Nagayoshi and colleagues (2022) reported that another phosphate transporter, XPR1 (encoded by the SLC53A1 gene), also contributes to the proliferation of CCOC, underscoring the broader relevance of phosphate homeostasis in ovarian cancer biology [32]. Supporting our hypothesis, pharmacological inhibition with PFA significantly reduced ($p < 0.05$) the proliferative index in tumor cell lines, while a milder effect was observed in IOSE cells. This differential sensitivity reinforces the notion that tumor cells are more dependent on elevated Pi influx to support metabolic activity and cell cycle progression than normal cells [29].

Conversely, the effects of PFA on cell motility were limited to IOSE cells, with no notable impact on tumor cells. This finding suggests that, in the tumor context, NaPi-IIb primarily supports proliferation rather than migration, whereas in non-tumorigenic cells, even subtle changes in phosphate transport can influence cellular motility. These results indicate that NaPi-IIb-mediated regulation of motility is not a significant mechanism driving EOC progression. In contrast, Russo and colleagues (2018) demonstrated that, in MDA-MB-231 cells—a widely used *in vitro* model of triple-negative breast cancer—blockade of phosphate transport with PFA significantly impaired migratory capacity. These findings underscore the importance of phosphate uptake in maintaining pro-tumorigenic features such as motility in this breast cancer model [33].

The functional activity of Na⁺-ATPase was also assessed, revealing that this enzyme is significantly hyperactivated in tumor cell lines compared to IOSE cells. These findings suggest a reprogramming of ion transport mechanisms associated with tumorigenesis, consistent with emerging evidence that ion transporters may play non-canonical roles in cancer progression [34]. Notably, in lung carcinoma, the (Na⁺+K⁺)-ATPase $\alpha 1$ subunit has been shown to cooperate with endogenous ligands to actively reshape the immune microenvironment and promote immune escape [34]. In this context, our data further support the concept that dysregulation of ion transport systems—such as Na⁺-ATPase, the Na⁺/Pi-II cotransporter, and other ATPases—may represent a hallmark of oncogenic reprogramming, with potential implications for tumor cell adaptation, survival, and resistance to therapy.

Na⁺-ATPase activity is insensitive to the classical inhibitor ouabain but responsive to furosemide, suggesting the involvement of atypical ATPases or alternative regulatory mechanisms in ovarian malignant transformation. Supporting our findings, previous studies have shown that furosemide inhibits proliferation of poorly differentiated gastric cancer cells via the Na⁺/K⁺/2Cl⁻ cotransporter, inducing G₀/G₁ cell-cycle arrest and highlighting its ability to impair tumor growth through modulation of sodium transport [35]. Notably, as furosemide treatment significantly reduced sodium-dependent phosphate uptake, cell proliferation, and motility in tumor cells without inducing immediate cell death—thereby excluding nonspecific cytotoxic effects—these results further reinforce the functional relevance of this enzyme in carcinogenesis [36].

Numerous studies have also linked (Na⁺+K⁺)-ATPase to cancer progression: ouabain targeting the $\alpha 3$ isoform of the enzyme inhibits proliferation and induces apoptosis in renal and lung carcinoma models [37], while blockade of (Na⁺+K⁺)-ATPase by cardiac glycosides diminishes proliferation, migration, invasion, and macropinocytosis in breast and colorectal cancer cells [38]. Additionally, acute inhibition disrupts the Na⁺-dependent glycolytic flux

in breast cancer, further connecting Na⁺-ATPase function to the metabolic reprogramming characteristic of malignancy [39].

Furthermore, Na⁺-ATPase activity surpassed that of (Na⁺+K⁺)-ATPase in the analyzed cell lines, exerting a stronger influence on proliferation and migration than NaPi-II itself. This suggests that Na⁺-ATPase may play a central role in supporting tumor cell survival and aggressiveness. Taken together, these complementary findings across different sodium transporters and cancer models underscore the critical role of atypical Na⁺ pumps in tumor biology, paving the way for novel therapeutic strategies targeting sodium homeostasis in cancer.

The relationship between NaPi-IIb and (Na⁺+K⁺)-ATPase was also investigated. Although the latter partially modulates the activity of the co-transporter, its activity levels did not correlate with ovarian carcinogenesis, suggesting that its functional contribution, while present, is secondary. In contrast, Na⁺-ATPase significantly modulated both motility and survival of tumor cells, thereby emerging as a more promising potential therapeutic target.

Regarding intracellular signaling, the data indicate that both the PI3K/Akt/mTOR and MEK/ERK pathways contribute to the functional regulation of NaPi-IIb, exhibiting lineage-dependent variations. Basal phosphorylation of Akt at Ser473 was elevated in A2780 and ACRP cells, and inhibition with wortmannin significantly reduced phosphate uptake in these lines, suggesting that the PI3K/Akt pathway is functionally coupled to NaPi-IIb activation. In contrast, this effect was attenuated in ES-2 cells. This pattern suggests that alternative pathways may compensate for phosphate transport regulation in this lineage, or that NaPi-IIb plays a less prominent functional role in ES-2 cells.

Consistent with our observations, other solute carriers—such as the bicarbonate transporter SLC4A7—have also been shown to promote tumor progression by engaging the PI3K/Akt/mTOR signaling cascade, as reported in head and neck squamous cell carcinoma (HNSCC), where SLC4A7 activation enhanced epithelial-to-mesenchymal transition and metastatic potential [40]. Similarly, the involvement of the MEK/ERK axis in modulating ion transport and cellular responses is supported by studies on trigeminal neuralgia, in which modulation of the MAPK pathway altered sodium channel activity and neuronal sensitivity [41]. Collectively, these findings reinforce the concept that ion transporters, beyond their canonical roles in homeostasis, may participate in oncogenic signaling and phenotypic reprogramming via lineage- and context-specific kinase pathways.

Regarding the MEK/ERK pathway, inhibition with U0126 markedly reduced phosphate uptake in cell lines exhibiting high ERK1/2 phosphorylation (A2780 and ACRP), but not in IOSE cells, which displayed low basal pathway activation. These findings suggest that, similar to PI3K/Akt, the MEK/ERK pathway regulates NaPi-IIb in a lineage-specific manner, underscoring the importance of detailed molecular characterization in the design of targeted therapeutic strategies. Consistently, in non-small cell lung cancer (NSCLC), downregulation of SLC34A2 (NaPi-IIb) significantly suppressed cell growth, migration, and invasion, an effect associated with reduced activation of both PI3K/Akt and Ras/Raf/MEK signaling pathways [27].

To assess the role of phosphate in the homeostasis of non-tumorigenic lung cells, Chang and colleagues (2006) reported that high concentrations of phosphate promote increased cell viability and significantly upregulate the expression of phosphate transporters in these cells. Furthermore, they found that elevated phosphate levels stimulate the phosphorylation of Akt and other kinases, including MEK, ERK, and Raf-1. Conversely, treatment of lung cells with PFA—a known phosphate transporter inhibitor—resulted in inhibition of Akt phosphorylation and subsequent reduction in cell growth, suggesting a role for phosphate in Akt-mediated cell survival [37]. Additionally, Jin and colleagues (2009) demonstrated that overexpression of NaPi-IIb led to phosphate-mediated activation of Akt and promoted increased carcinogenesis and proliferation of non-small cell lung carcinoma (NSCLC) cells [42]. This parallel regulation suggests that the functional coupling of NaPi-IIb to MEK/ERK and other pathways is conserved across tumor types, further reinforcing its potential as a context-dependent therapeutic target.

Although our findings strongly support a mechanistic link between NaPi-IIb expression, phosphate uptake, and activation of oncogenic kinases, the directionality of this relationship warrants further investigation. It remains possible that the enhanced proliferative and signaling responses observed following NaPi-IIb upregulation are secondary to increased intracellular phosphate availability, which fuels ATP and nucleotide synthesis and thereby sustains kinase activation [21]. Conversely, transporter-independent functions of NaPi-IIb cannot be excluded. Previous studies have suggested that certain membrane transporters, including members of the SLC34 and SLC20 families, may act as molecular scaffolds or modulators of intracellular signaling independent of their transport activity [24].

Conclusion

Whether NaPi-IIb directly participates in signaling complex formation or indirectly modulates downstream cascades through phosphate flux remains to be determined. Future experiments employing transport-deficient NaPi-IIb mutants or controlled phosphate supplementation and deprivation will be essential to disentangle these possibilities and clarify the precise causal relationship between NaPi-IIb activity and oncogenic signaling in EOC. Nonetheless, the results of this study indicate that hyperactivity of NaPi-IIb and Na⁺-ATPase is associated with proliferation, survival, and motility of EOC cells. The functional modulation of these proteins by pathways such as PI3K/Akt/mTOR and MEK/ERK, in a lineage-dependent manner, highlights promising molecular targets for the development of more effective and selective therapeutic approaches for EOC treatment.

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Statement of Ethics

The authors have no ethical conflicts to disclose.

Disclosure Statement

The authors have no conflicts of interest to declare.

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