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Examination of Hypoxanthine Guanine Phosphoribosyltransferase as a biomarker for colorectal cancer patients

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ABSTRACT
The aim of this study is to investigate these enzymes as possible biomarkers in two colorectal cancer cell lines: HT29, SW620, and Colo205. With 1,168,929 individuals currently diagnosed with colorectal cancer in the United States, there remains a need to find biomarkers to improve diagnosis and expand treatment options for patients. Due to their role in proliferation and cell cycle regulation, we hypothesized an increase in salvage pathway enzyme (APRT, DCK, and HPRT) expression and possible presentation within colon cancer cells. Enzyme surface localization was assessed utilizing confocal microscopy, flow cytometry, and scanning electron microscopy. General protein expression was evaluated utilizing immunohistochemistry and Western blot analysis. While we found no statistically significant presence of either APRT or DCK on the membranes of SW620, Colo205, and HT29 cells, but found significant expression of HPRT on the surface of HT29, SW480, and SW620 cells. The average population fluorescence increased by 28%, 58%, and 40% in HT29, SW620, and SW480 cells, respectively, when compared to isotype controls. Confocal microscopy images revealed direct overlap between SW620 cells stained with a membrane dye and anti-HPRT antibody, indicating co-localization on the plasma membrane. In addition, cells treated with gold labelled HPRT antibody experienced significant changes in gold weight percentage on both SW620 and HT29 cells when compared to isotype controls. When evaluating expression within normal tissue, there was insignificant levels of HPRT binding. These data collectively suggest that HPRT may be a possible biomarker target for the identification and treatment of colorectal cancer.

Introduction
Colorectal Cancer (CRC) is one of the leading causes of death in the United States. Every year 49,700 individuals die as a result of CRC while an estimated 1,168,929 are currently diagnosed with the disease. CRC is one of the most common cancers in the western world as 1 in 21 men and 1 in 23 women are predicted to develop the disease. These markers are valuable when determining treatment options for individuals with a unique blend of cancer characteristics. Recent research using cultured cancer cells have identified cancer biomarkers such as the 1,25-Dihydroxyvitamin D3 receptor that may act as a marker for colon carcinoma cell differentiation and analysis. Receptors such as Vitamin D receptor are upregulated on colon cancer cell surfaces and can serve as a target for tumor reduction and elimination. Additionally, markers such as CD133 and CD44 have also been identified for the elimination of cancer stem cell. While a number of tumor antigens have been identified, additional markers will aid in better understanding colorectal cancer disease progression and could lead to additional treatment options.

In the search to further characterize colorectal cancer cells, we decided to evaluate the salvage pathway enzymes Hypoxanthine Guanine Phosphoribosyltransferase (HPRT), Adenine Phosphoribosyltransferase (APRT) and Deoxycytidine Kinase (DCK) as a possible upregulated target. Salvage pathway enzymes act as recycling agents, reusing the components of old nucleotides to skip energetically expensive steps in the formation of nucleotide base. The salvage pathway is the chosen method of nucleotide synthesis for a majority of the cell cycle in humans as 90% of free purines are recycle. Responsible for the salvage of adenine in the cell cycle, APRT is found constitutively expressed in a majority of mammalian cell. DCK is primarily involved in the phosphorylation of deoxycytidine in the production of cytosin. HPRT functions by transferring phosphoribose from phosphoribosyl pyrophosphate (PRPP) to hypoxanthine or guanine bases in the purine biosynthesis of inosine and guanin.

A deficiency of HPRT results in a spectrum of diseases that directly correspond with the availability of the protein. Individuals with a complete lack of functional HPRT will develop Lesh-Nyhan syndrome, while individuals with a partial deficiency will develop gout-like symptoms characteristic of Kelley-Seegmiller syndrom. Because the gene is located on the X chromosome, it is an X-linked recessive condition that predominantly affects males of diseased families. Evaluation of the hprt gene has become a common biomarker.
for mutational assessment, and over 500 mutations in the gene have been described.\textsuperscript{17}

Having functional salvage pathway enzymes is important in the survival and functionality of mammalian cells. Salvage enzymes, such as HPRT, are known as common housekeeping genes, and are integral in several daily cellular functions regulating cell proliferation and cell cycle progression.\textsuperscript{16,18} We evaluated these enzymes because of their intimate role in the production of nucleotides necessary to maintain rapid cell proliferation. Additionally, these enzymes maintain responsibility for synthesizing GTP and ATP which provide the critical energy source for several cellular processes that are found upregulated within malignant cell.\textsuperscript{19–21}

The aim of this study was to evaluate HPRT, APRT, and DCK as potential biomarkers for CRC. We assessed the expression of the proteins on the surface of four CRC cell lines in addition to evaluation within tumor tissue and normal tissue to determine the clinical relevance of the protein expression. Results from these experiments could provide an additional marker for the characterization of colorectal cancer.

**Results**

**Flow cytometry reveals an overall increase in fluorescence when colon cancer cell lines were exposed to HPRT antibody, but not when treated with DCK or APRT antibodies**

Flow cytometry revealed no significant presence of APRT (p-value = 0.93) or DCK (p-value = 0.243) on the surface of SW620 cells (Figure 1). There was also no statistically significant presence of both enzymes (APRT, p-value = 0.39; DCK, p-value = 0.57) on the surface of HT29 cells or Colo205 cells (APRT, p-value = 0.75; DCK, p-value = 0.96). We did find that SW480, SW620, and HT29 cells had statistically significant HPRT expression on the surface of the cells. The average fluorescence of the cell population increased by 27.73% in HT29 (p-value = 0.013), 39.6% in SW480 (p-value = 0.0095), and 58.85% in SW620 cells (p-value = 0.0079) when compared to isotype controls (Figure 2). This indicates a strong presence of HPRT on the surface of the cells. Figure 2 shows insignificant isotypic binding, lower than 3% average of the total population (p-value = 0.374) in SW620 cells, 7% (p-value = 0.11) in HT29 cells, and 3.94% (p-value = 0.058) in SW480 cells. While these cell lines showed positive HPRT surface localization, Colo205 cells showed no significant increase in the surface presence of HPRT (p-value = 0.99). All cells were gated to exclude dead cells and cell doublet populations.

**HPRT is strongly associated with the plasma membrane of SW620 cells**

To ensure antibody binding was towards surface HPRT and not cytoplasmic HPRT, confocal microscopy was performed to visualize protein localization on SW620 cells (Figure 3). In all controls we observed a minimal FITC signal, indicating insignificant antibody binding, with the exception of samples treated with anti-HPRT. SW620 cells treated with anti-HPRT FITC antibody had a noteworthy association with the plasma membrane. These images reveal a direct overlap between the plasma membrane and antibodies targeting HPRT. In addition, the FITC channel reveals a distinguishable external presence of HPRT as fluorescent antibody binding is only seen on the periphery of the cells.

Western Blot analysis shows there is a significant presence of HPRT within SW620 cancer cells. Along with a clear presence of the protein, this analysis also confirmed HPRT as a membrane associated protein, as it is found in the biotinylated fraction of the cells (Figure 4).

**Scanning electron microscopy reveals a random distribution of the protein across the surface of HT29 and SW620 cells**

To evaluate whether surface HPRT binding was distributed across the membrane randomly, we utilized scanning electron microscopy to physically visualize the position of the enzyme on the surface of the cells. As pictured in Figures 5 and 6, both cell lines show an increase in gold particle binding when exposed to anti-HPRT antibodies. This same increase in expression is not seen with isotype controls and further implicates HPRT on the surface of the cells. The protein appears to be randomly presented across the plasma membrane with no clear pattern of expression. EDAX analysis (Figure 6) for each sample shows an increase in the gold percentage when cells were treated with anti-HPRT antibodies. A significant increase in sample elemental gold percentages is seen as SW620 cells (p-value of 8.14x10\textsuperscript{-6}) and HT29 cells (p-value of 1.74x10\textsuperscript{-4}) were treated with gold labelled HPRT antibodies. This analysis provides a further confirmation that HPRT is present on the surface of SW620 and HT29 colon cancer cells.

**Within normal colon samples from patients, there is insignificant levels of HPRT binding**

To evaluate whether HPRT would be useful as a biomarker target for CRC patients, we measured the proteins levels on the surface of normal patient tissue. Flow cytometry revealed insignificant HPRT presence within normal colon tissue from healthy patients when compared to isotype controls (p value = 0.998) and unstained controls (p value = 0.996). When compared to a CD44 control, HPRT levels were minimal and shared similar binding to that of the isotype control, indicating its presence to be negligible in normal tissue (Figure 7).

**HPRT expression within malignant tissue demonstrates the variable nature of HPRT upregulation**

In order to evaluate HPRT expression levels within malignant tissue, 94 patient samples were stained (Table 1). While the overall average staining intensity of all malignant samples (p-value = 0.545) was insignificant when compared to normal patients, in 59% of the malignant tissue stained for HPRT, there was a substantial increase in the protein expression,
Figure 1. Analysis of APRT and DCK expression on SW620 and HT29 colon cancer cells. A, Cells treated with APRT antibodies experienced no shift in the fluorescent population and had similar fluorescent signatures to isotype controls. Statistical analysis of APRT and DCK binding reveal insignificant levels of either protein on the surface of SW620 cells. APRT antibodies were mouse and were compared against mouse isotype controls, and DCK antibodies were rabbit and were compared against rabbit isotype controls for statistical analysis. Insignificant shifts in the fluorescent intensity of the cells was observed when treated with both B, APRT and DCK antibodies. Statistical analysis of APRT and DCK binding in HT29 cells showed insignificant levels of the proteins on the surface. C, Insignificant surface binding was also observed in Colo205 cells as well with no shifts in the fluorescent population upon treatment with either APRT or DCK antibodies.
while 41% of the patient samples showed insignificant increases in HPRT expression (Figure 8). This elevation was statistically significant from normal tissue ($p < 0.001$) and demonstrates unique HPRT production within a cohort of the patients (Figure 9). This expression was also significant when compared to the GAPDH control, which was utilized to assess housekeeping levels of protein expression.

**Analysis within malignant colon samples confirms the variable nature of HPRT surface localization within patients**

Three malignant biopsy samples were obtained from patients with colorectal cancer. Of the three samples obtained, two of them had no HPRT surface localization, while one of the samples had elevated surface HPRT (Figure 10). This confirms the variation found within the tissues and the cell lines evaluated as the observed HPRT over-expression and subsequent surface presentation was not found within all the patients. Patients without surface localization had expression similar to the Colo205 cancer cell line tested, while patients with surface localization had HPRT expression similar to levels observed in SW620, HT29, and SW480 cell lines. As one of the patients had a significant level of HPRT on the surface of their tumor cells, it demonstrates there is potential for the protein to be targeted within those individuals who experience an up-regulation of the enzyme within their tumors.

**Discussion**

HPRT is a common housekeeping gene critical to the successful production and regulation of nucleotides within the cell cycle. Our results show a significant presence of HPRT on the surface of HT29, SW620, and SW480 colorectal cancer cell lines. These results show a different role of HPRT within a malignant environment that has not been reported. Currently, HPRT is understood to be expressed at a relatively constant level within tumor cells as it is commonly used as an endogenous control for several molecular technique. Our results question the current view of HPRT within colon cancer tissue as it has shown to possess unique characteristics within cancer cell lines in addition to within malignant colon tissue. HPRT expression appears to be very similar to the expression of other biomarkers for colorectal cancer, such as the Vitamin D3 receptor which also shows presence on the surface of colon cancer cells and serves as a marker for cell differentiation and growth. This marker is currently used to reduce tumors and it is likely that HPRT could serve this same purpose within colon cancer tissue as it seems to be related to cell proliferation, and appears to be absent on the surface of healthy tissue.
There is other evidence of salvage enzymes serving as diagnostic and prognostic biomarkers to diagnose and monitor cancer development in patients. Thymidine Kinase 1 (TK1), another salvage pathway enzyme, serves as a serum biomarker for cancer recurrence and has shown to have potential as a therapeutic biomarker as well. HPRT may also be used in a similar setting to aid in diagnosing cancers as it appears to only be upregulated in cancer tissue. Unlike TK1, it does not appear to be stage dependent, which would be useful as an early diagnostic companion tool to detect early stage cancer. While pathologists analyze patient biopsy tissue, HPRT could also be evaluated to help in the initial diagnosis.

While HPRT is present on SW620, SW480 and HT29 cells, the relative abundance of the protein is not equal between the cell lines. SW620, the highest expressing cell line, has upwards of 25% more protein on the surface when compared to the lowest expressing HT29 cells. SW620 cells are derived from a metastatic lymph node and are aggressive, fast growing cancer cells. In contrast, HT29 cells are derived from a colorectal adenocarcinoma originating in the mucus glands in the colon and rectum and are consequently less aggressive. Our results potentially indicate that HPRT surface expression may be more abundant in aggressive, rapidly proliferating cells, but this needs to be further explored.

Figure 3. Plasma membrane co-localization of HPRT in SW620 cells. SW620 cells were dyed with both a FITC dye and a Rhodamine Red membrane dye to label antibody treatments and the plasma membrane respectively. Utilizing unstained cells, IgG treated cells, and NF-kB treated cells as controls, plasma membrane associations were evaluated to determine whether any of the treatments significantly bound to the membrane of SW620 cells. A, Each sample was analyzed and imaged by a 488nm laser to illuminate FITC positive cells. These images show the binding of the respective antigen treatment. B, Samples were also imaged in a 594nm laser to show rhodamine positive cells. This dye binds to the plasma membrane of all cells. C, The two images obtained from column A and B were merged to show associations between treated antibodies and the plasma membrane of cells. These results show a clear overlap between cells treated with anti-HPRT antibody and those treated with the membrane dye.

Figure 4. Western analysis of HPRT expression in both cytosolic and membrane fractions. Surface proteins were biotinylated and isolated for analysis. SW620 cell extract, membrane fractions, and cytosolic fractions were probed for HPRT along with a non-biotinylated control. This data shows that there is a very significant presence of HPRT within SW620 cytosol in addition to a clear presence on the surface of the cells.
We hypothesize the observed surface expression of HPRT in these cell lines may point to a regulatory element of HPRT expression that has lost function in cancer cells within certain patient. The HPRT gene has several regulatory transcription factors that control its expression (P53, NF-κB, FOXL1, etc.) which may be altered in SW620, SW480 and HT29 cells due to mutation. Loss of hprt gene control may increase levels of the protein in the cell and subsequently result in the export of the protein to the extracellular matrix where it may transiently reside on the surface of the cell. Further investigation into the mechanism by which this cytosolic protein is transported to the plasma membrane of these cancer cells needs to be evaluated to elucidate how HPRT is able to localize to the surface, and if it provides any functional advantage to the cancer cell. Furthermore, HPRT has shown to have a unique expression profile within a cohort of patients as determined by IHC staining. The over-expression of HPRT within these patients also points to a loss of HPRT regulation and may aid in determining which patients may experience this unique surface expression HPRT. While we were able to evaluate a few malignant cell lines and tumors for HPRT surface presentation, further testing with more patient samples will need to be done to determine how prevalent HPRT surface expression is and which patients could benefit from potentially targeting the protein. As colo205 cells do not show significant HPRT surface localization, further investigation into the mechanism

**Figure 5.** Scanning Electron Microscopy Images and resulting EDAX in HT29 and SW620 cells. Cells were labelled with gold towards their respective antibody treatment. The size scale is shown in each image and represents a 2µm distance. Sa-A & Sb-A, Images were obtained using a Back Scatter Electron (BSE) detector. This detector is specialized to image heavy metals within samples, and highlights enhanced gold within the sample. Any distinguishable large particles of gold represent a bound antibody. Sa-B & Sb-B, Images were also obtained with a Gaseous Side Electron (GSE) detector, which showed cell morphology to ensure correct cell structure and integrity. Sa-C & Sb-C, EDAX analysis of each sample showed the gold elemental peaks for all the elements present within the sample. Silicon is the highest represented element because cells were mounted on silicon cover slips for analysis. The gold elemental peak is indicated with a gold arrow. Images obtained from this analysis show the exact location of the HPRT bound to the surface of the cell, and show no patterns indicating a random distribution of the protein across the surface of the cell.
of surface presentation needs to be conducted to determine the reason some cells express HPRT on the surface while others do not.

HPRT could be used as a valuable marker for studies evaluating biomarker targetin. Testing against this antigen could provide researchers with significant advantages when evaluating

Figure 5. (Continued).

Figure 6. Gold percentage of SW620 and HT29 cells. The gold elemental composition of each sample is denoted on the Y-axis. The increase in the gold percentage when cells were exposed to HPRT and CD44 shows a quantifiable increase in the gold present on the outside of the cell. A, Gold elemental percentages in SW620 cells exposed to HPRT antibody had a gold weight of approximately 11.2%, which is statistically significant to the IgG controls used for background binding (p value < 0.0001). These data indicate a statistically significant presence of HPRT on the surface of SW620 cells. B, Gold elemental percentages in HT29 cells. Gold weight was approximately 10.4% with a p value < 0.0001.
therapy efficiency and may lead to a new biomarker target for the treatment of a subset of colorectal cancer patients who experience an upregulation and surface presentation of the protein.

Conclusions

As a surface biomarker that is not present in normal cells, HPRT could be used as a valuable target for immunotherapies. Patients who experience an elevation in HPRT within their tumors may use the protein as a means to reduce tumor burden by targeting HPRT+ cells.

Materials and methods

Chemicals

Anti-HPRT monoclonal antibody (Thermo Fischer Scientific) was aliquoted and stored at −20°C. Mouse-FITC and Rabbit-FITC antibodies (Sigma Aldrich) were stored at 4°C and were used in dark conditions. Bovine Serum Albumin (Sigma Aldrich) and Sodium thiosulfate (Macron Fine Chemicals) were dissolved in PBS at a 1% concentration and stored at 4°C. 50% Glutaraldehyde stock (Electron Microscopy Sciences) was stored at −20°C and workable solutions were diluted to 0.25% in PBS and stored at 4°C. Glycine (Thermo Fischer Scientific) was diluted to 0.2mM in PBS and stored at 4°C. NF-kB polyclonal antibody (Bioss Antibodies), DCK polyclonal antibody (rabbit: Abnova) and monoclonal antibody (mouse: Santa Cruz, Dallas TX) and APRT polyclonal antibodies (mouse: One World Labs, San Diego, Cal; rabbit: Abnova) were stored at −20°C. CD44 monoclonal antibody (One World Lab) was stored at −20°C. Propidium Iodide (Sigma Aldrich Inc.) was stored at 4°C and aliquoted for use. Fc Block was purchased from Biolegend and stored at 4°C. An APC-Conjugation Kit (Abcam) was stored at −20°C.

Cell culture conditions

The human colon carcinoma cell lines SW620, SW480, Colo205 and HT29 were obtained from the American Type Cell Culture Collection.
Culture Collection. HT29 and Colo205 cells were grown in RPMI 1640 medium supplemented with 10% fetal bovine serum and 2mM L-Glutamine (all from Hyclone). SW620 and SW480 cells were grown in DMEM medium supplemented with 10% fetal bovine serum and 4mM L-Glutamine (all from Hyclone). The cell media was replaced every 48 hours and cells were trypsinized and the cell population was reduced by half once 90% confluence was obtained. Cells were treated with Acutase (Stem Cell Technology) when utilized for flow cytometry and when plated for all other applications. Cell viability was evaluated using a trypan blue staining, and cells were utilized for all applications when the viability was greater than 98%. All cells were grown at 37°C and 5% CO₂. Cell lines were authenticated in May of 2016 by the University of Arizona Genetics Core.

**Flow cytometry**

The surface presence of HPRT, APRT, and DCK in cultured cells was evaluated measuring the fluorescence intensity of antibodies against each of the salvage pathway enzymes. All samples were analyzed on a Blue/Red Attune (Applied Biosystems), and 25,000–50,000 events were recorded per sample. Briefly, 3-5x10⁵ cells were incubated with 200µL of

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**Figure 8. Evaluation of HPRT expression within patient tissue.** All tissues were stained with a monoclonal anti-HPRT antibody. The resulting converted grayscale image is pictured in the top left corner of each image while the grayscale plot is below. A, Tissue from a 79-year-old female patient with stage IIB colon adenocarcinoma and B, tissue from a 48-year-old female patient with stage IV colon adenocarcinoma. These malignant tissues are significantly darker stained than normal colon tissue. C, Normal colon tissue from a 36-year-old male patient and D, tissue from a 31-year-old male patient. These tissues show an upregulation of HPRT within malignant cells.
PBS containing 1µg of primary antibody treatment for 15 minutes on ice. Cells were then labelled with FITC-conjugated secondary (mouse or rabbit) antibody for 15 minutes on ice. Isotypic IgG and unstained cells served as negative controls to ensure correct cell gating. The forward/side-scatter plots were used to gate out cell doublets, dead cells, and cell debris. Using unstained and isotype controls as guides, the positive population was determined by the overall shift in the fluorescent intensity. Each cell line was independently analyzed four times and the data was plotted using FlowJo Software (FlowJo Enterprise). Cells were washed appropriately after each step of the protocol.

**Patient tissue dissociation and analysis**

Healthy and malignant colon tissue samples were obtained from the Utah Valley Regional Medical Center. Tissue samples were minced using sterile scalpels into pieces ranging from 2-3mm in length and were suspended in Hanks media (5% FBS). Minced tissue was then placed in a solution of Collagenase IV on a shaker for 1–4 hours depending on the fat percentage of the tissue. Once dissociated the solution was washed through a 100 micron filter, to produce a single cell suspension. These cells were then washed and treated with an Fc block to minimize non-specific antibody binding. Following blocking, cells were treated with anti-CD45 FITC, anti-HPRT APC and PI to aid in the selection of correct cell population.
populations for analysis. Cells were gated on CD45- and PI-

cell populations to avoid analysis of lymphocytes resident in the
tissue and dead cells.

**Biotinylation and western blot analysis**

Cells were analysed for surface presence along with general expres-
sion within the cell utilizing the Pierce Cell Surface Protein
Isolation Kit (Thermo Scientific). Briefly, 3 flasks of SW620 cells
were grown to 95% confluency, washed, and treated with a kit-
provided biotin solution. Following rocking on a shaker for
30 minutes at 4°C, the cells were treated with a quenching solution.
Cells were detached from the flask via cell scraping and transferred
to a 50mL conical vial for washing. Then, cells were treated with a
lysis solution and incubated for 30 minutes at 4°C. Cell lysis was
added to a neutravidin gel and incubated for 60 minutes at room
temperature. This solution was then passed through a filter and
proteins bound to biotin are trapped within the column. The

neutravidin gel was washed 4 times and the flow through was
collected and labelled “cytosolic fraction”. The biotin labelled
protein was then eluted from the column utilizing a 50mM DTT
solution and labelled “membrane fraction”.

Both membrane fractions and cytosolic fractions along with
cell extract from SW620s were evaluated for protein expression
utilizing standard Western Blotting techniques described in Sewda
et al. with slight modifications.38 Briefly, each sample was boiled
for 5 minutes prior to running on a 12% polyacrylamide gel under
reducing conditions. Gels were then transferred to a nitrocellulose
membrane (Biorad Laboratories), blocked, and treated with an
anti-HPRT monoclonal antibody overnight at 4°C. Following
primary antibody treatment, membranes were washed and treated
with a rabbit fluorescent secondary antibody (Licor) for 1 hour.
Membranes were then imaged on a Licor Odyssey CLx. SW620
cells were utilized for this analysis because their expression of
HPRT on the membrane is significantly higher than that of
HT29 cells.

**Confocal microscopy**

Fluorescently-stained cells were examined under an epifluorescence
microscope (Olympus, Tokyo, Japan) equipped with a laser con-
focal system (Bio-Rad Laboratories, Hercules, CA) containing a
15mW Krypton/Argon laser. Image processing was carried out
with Laser Sharp Computer Software (Bio Rad Laboratories). After
treatment with acutase, cells were plated at a concentration of
400,000 cells/mL on glass coverslips. Following one to two days
of growth, cells were incubated in 500µL of PBS containing 2.5µg
of anti-HPRT antibody for 15 minutes on a shaker at 4°C. Cells
were then labelled with 2.5µg of FITC-conjugated secondary anti-
body for 15 minutes at 4°C. Then, cells were incubated at 37°C for
10 minutes with a 1:1000 dilution of a Cell Mask Deep Red plasma
membrane dye (Fisher Scientific).

**Electron microscopy**

Following acutase treatment, cells were plated at a concentration of
400,000 cells/mL on glass coverslips. After one to two days of
growth, cells were placed in 6 well plates and washed with PBS
three times and with 1% PBS-BSA for 5 minutes at 4°C followed by
a sodium azide wash. Cells were then incubated with 2.5µg or 5µg
of primary antibody conjugated to Biotin for 15 minutes on a
shaker at 4°C. After primary incubation, cells were washed with
1% PBS-BSA followed by two washes with PBS. Then, cells were
washed with 1% PBS-BSA and 1% PBS-sodium thiosulfate for
5 minutes on a shaker at 4°C. Cells were incubated with 2.5µg of
Streptavidin-gold conjugate (Nanoprobes, Yaphank, NY) for
15 minutes on a shaker at 4°C. This is followed by a 1% PBS-
BSA wash and three PBS washes. Cells were then fixed with a
0.25% Glutaraldehyde solution diluted in PBS for 5 minutes. The
reaction was then extinguished by adding a 0.2mM PBS-Glycine
Solution and incubating for 10 minutes until the solution turned
a slight yellow color. Cells were then washed three times with
ddH2O. Solutions A and B from the Nanoprobes gold enhance-
ment kit (Nanoprobes Inc.) were incubated together for 5 minutes.
Solutions C and D were then added, vortexed, and 40µL of the gold
enhancement were added to each sample and incubated for 5 min-
utes. Each sample was put through a series of dehydrations with
70%, 80%, 90%, and 100% ethanol. Gold labelled samples were
examined under a Phillips XL-30 ESEM using a 15kV electron
stream under low vacuum conditions at 1 Torr. A Gaseous Side
Electron (GSE) detector was utilized to image the cell morphology
topography. A Back Scatter Electron (BSE) detector was
used to visualize gold particles on the cell surface. Once images
for the cells were obtained, the elemental composition of the cells
was evaluated using energy dispersive spectroscopy (EDAX) and
X-rays. EDAX analysis will provide a k-ratio, a Z value, an A value,
and an F value. The k-ratio represents the element’s peak height
compared to a sample of the pure element collected under the
same conditions. The Z value represents a correction in the atomic
number taking backscattered electron yield of the pure element
and the sample. The A value represents a compensation for X-rays
generated in the sample that are cannot emit energy. The F value
represents a correction for the generation of X-rays. We used these
EDAX output values to normalize our samples gold weight per-
centages using the following equation:

\[
\text{Normalized Weight Percentage} = \frac{K \times \text{ratio}}{Z \times A \times F}
\]

**Immunohistochemistry**

Colorectal Adenocarcinoma tissue arrays were obtained from
BioMax. These tissues contain various stages of cancer along
with corresponding benign and normal tissue from 100 different
patients. HPRT levels were assessed utilizing standard immuno-

histochemistry staining. Tissues were rehydrated in a series of
ethanol washes before treatment with a DIVA (Biocare Medical)
solution to retrieve antigen. Tissues were then incubated with a
Background Sniper (Biocare Medical) solution to reduce non-

specific antibody binding. Following blocking, a primary antibody
is added to the tissue at a concentration of 1:100 to 1:200 and
incubated overnight at 4°C. Following primary staining, tissues
were washed and then treated with secondary antibody conjugated
to a HRP polymer (Biocare Medical) and incubated for an hour.
Following washing, a DAB Peroxidase solution was incubated
with the tissues. Areas of antibody binding converted the colorless
substrate to a brown product, effectively highlighting the target
protein. Tissues were treated with hematoxylin (Biocare Medical)
to stain the nucleus of the cells. Along with HPRT treatment, a universal negative antibody (Biocare Medical) was used as a negative control, and a GAPDH antibody was used as a positive control.

Tissues were quantified utilizing ImageJ software. All images were evaluated using the IHC toolbox ImageJ plugin. The DAB option is chosen and the tissue image is then removed of all other staining except for DAB. Following this analysis, the image is then converted to a grayscale and a threshold is applied in order to eliminate areas of no staining inherit within the tissue image. Once the threshold is applied the average gray value of the tissue is collected. The same threshold is applied to all tissue samples in order to ensure consistency.

**Statistical analysis**

ANOVA statistical analysis with the multiple comparison method were used to determine the differential surface expression of the various treatments for flow cytometry data on all cell lines. In addition, two-way ANOVA tests were performed to compare the mean expression of HPRT between SW620 and HT29 cells. EDAX data was analyzed using an ANOVA with the multiple comparison method in addition to unpaired t tests to determine significance between samples. All statistical analysis was evaluated using GraphPad Prism 7 software. Differences were considered significant when the p value was < 0.05.

**Abbreviations**

HPRT Hypoxanthine Guanine Phosphoribosyltransferase
APRT Adenine Phosphoribosyltransferase
DCK Deoxycytidine Kinase
GSE Gaseous Side Electron
BSE Back Scatter Electron
EDAX Energy Dispersive Spectroscopy

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**Disclosure of Potential Conflicts of Interest**

The authors declare that they have no competing interests

**Ethics approval and consent to participate**

All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

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