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Examining the Relationship Between Cu-ATSM Hypoxia Selectivity and Fatty Acid Synthase Expression in Human Prostate Cancer Cell Lines

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Abstract

Introduction—PET imaging with Cu-ATSM for delineating hypoxia has provided valuable clinical information, but investigations in animal models of prostate cancer have shown some inconsistencies. As a defense mechanism in prostate cancer cells, the fatty acid synthesis pathway harnesses its oxidizing power for improving the redox balance despite conditions of extreme hypoxia, potentially altering Cu-ATSM hypoxia-selectivity.

Methods—Human prostate tumor cultured cell lines (PC-3, 22Rv1, LNCaP, and LAPC-4), were treated with an FAS inhibitor (C75, 100 μ M) under anoxia. ⁶⁴Cu-ATSM uptake into these treated cells, and non-treated anoxic cells, was then examined. Fatty acid synthase (FAS) expression level in each cell line was subsequently quantified by ELISA. An additional study was performed in PC-3 cells to examine the relationship between the restoration of ⁶⁴Cu-ATSM hypoxia-selectivity and the concentration of C75 (100, 20, 4, or 0.8 μ M) administered to the cells.

Results—Inhibition of fatty acid synthesis with C75 resulted in a significant increase in ⁶⁴Cu-ATSM retention into prostate tumor cells *in vitro* under anoxia over 60 mins. Inhibition studies demonstrated higher uptake values of 20.9 ± 3.27 , 103.0 ± 32.6 , 144.2 ± 32.3 , and $200.1 \pm 79.3\%$ at 15 mins over control values for LAPC-4, PC-3, LNCaP, and 22Rv1 cells, respectively. A correlation was seen (R² = 0.911) with FAS expression plotted against % change in ⁶⁴Cu-ATSM uptake with C75 treatment.

Conclusions—Although Cu-ATSM has clinical relevance in the PET imaging of hypoxia in many tumor types, its translation to the imaging of prostate cancer may be limited by the over-expression of FAS associated with prostatic malignancies.

Keywords

Fatty Acid Synthase; Cu-ATSM; PET; prostate cancer

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Introduction

The onset of hypoxia in malignant tissues is fairly universal, though not homogeneous, and is associated and influenced by a myriad of complicated processes. It is thought that since tumor cells are metabolically more active and proliferative than normal cells, they drain the surrounding tissue and blood supply of oxygen faster than it is supplied [1-3]. Tumors with increased levels of hypoxia afford resistance to traditional radiation therapy, as well as reducing the effects of many chemotherapeutic agents [4-7]. Measuring tissue oxygenation is no trivial task. The most common method, the use of an Eppendorf O_2 probe to measure oxygen tension, is effective but also quite invasive. Although the values obtained through this method have been predictive of metastatic potential and patient response to treatment [8-10], the technical difficulties of this technique make it generally undesirable.

In recent years, investigations into alternative, non-invasive methods for measuring pO_2 have been pursued by numerous researchers [2]. The use of PET imaging in conjunction with radiolabeled molecules that undergo chemical changes in the presence or absence of oxygen has led to a few promising tracers. ¹⁸F-MISO (¹⁸F-fluormisonidazole) exploits a simple chemical cycle whereby it takes the place of oxygen as an electron acceptor and is reduced and trapped. In the presence of oxygen, the initially reduced compound returns to its original state by receiving an additional electron from oxygen in a futile cycle. Imaging with ¹⁸F-MISO has proved somewhat successful as a measure of oxygen levels, however the contrast between hypoxic and normoxic tissues is minimal (tumor to blood ratio > 1.2) [11-15]. ¹⁸F-FAZA (¹⁸F-fluoroazomycin arabinoside) has also been investigated as an alternative to ¹⁸F-MISO with faster background clearance while maintaining hypoxia selectivity [16, 17]. A recent pilot study showed promising results for clinical imaging in head and neck cancer patients [18].

Copper(II)-diacetyl-bis(N^4 -methylthiosemicarbazone), Cu-ATSM, labeled with a positron emitting isotope of copper (⁶⁰Cu, ⁶¹Cu, ⁶²Cu or ⁶⁴Cu) has been shown, *in vitro* and *in vivo*, to be selective for hypoxic tissue [19-21]. The clinical use of Cu-ATSM for delineating hypoxic human tumors using PET has provided valuable information in cervical, lung and rectal cancers [22-25]. Although a rapid >4-fold increase in retention of Cu-ATSM in hypoxic over normoxic cells was observed with overall retention at around 90% in hypoxic compared to 25% in normoxic cells [20, 26], these investigations did not include the use of prostate tumor cell lines. Burgmann et al. reported diminished hypoxic selectivity of Cu-ATSM *in vitro* in a rat prostate cancer cell line, R3327-AT [27]. In their analysis of 6 different cell lines ratios of hypoxic or anoxic uptake of ⁶⁴Cu-ATSM to normoxic uptake showed that the retention into the two prostate lines were consistently the lowest of the six lines examined. The same group also reported *in vivo* imaging studies comparing 64 Cu-ATSM and ¹⁸F-MISO in two animal models of cancer [28]. In the FaDu human squamous cell carcinoma tumor model, the early (\sim 1-4 h) and late (\sim 19 h) ⁶⁴Cu-ATSM images were similar and were in general accordance with ¹⁸F-MISO scans. However, their data did not support using Cu-ATSM as an indicator of hypoxia at early timepoints in the R3327-AT prostate model, with a negative correlation with ¹⁸F-MISO at 1 hour after administration. Therefore, although the use of Cu-ATSM has been validated in vitro and in vivo in multiple tumor models and clinically in various human cancers, there is a concern that its ability to delineate hypoxia in **prostate** tumors may be suspect.

Our recent work has demonstrated that there is extensive involvement of the fatty acid synthesis pathway in 1-¹¹C-acetate uptake in prostate tumors, leading to a possible marker for fatty acid synthase (FAS) expression in vivo by PET [29]. FAS is a multifunctional enzymatic protein that catalyzes fatty acid biosynthesis [30] and FAS levels are associated with tumor aggressiveness in late-stage prostatic adenocarcinomas as well as a prognostic

indicator for overall survival [31]. It has also been noted that the physiological significance of the fatty acid synthesis pathway in prostate cancers is in the harnessing of its oxidizing power for improving redox balance (i.e., lower NADH/NAD+ ratios) despite oxygen limiting (hypoxic) conditions [32]. Therefore, considering that the mechanism of Cu-ATSM hypoxia-selectivity is reliant on an hypoxia-induced intracellular chemical reduction of Cu(II) to Cu(I) but that the fatty acid synthesis pathway improves redox imbalance during hypoxia, we wanted to explore the relationships between the hypoxia-selectivity of Cu-ATSM and the expression levels of FAS in prostate tumor models.

Methods

All chemicals, unless otherwise stated, were purchased from Sigma-Aldrich Chemical Company, Inc (St. Louis, MO). All solutions were prepared using distilled, deionized water (>18 M Ω resistivity) by passing through a Milli-Q filtration system (Millipore Corp., Milford, MA). Radioactive samples were counted in a radioisotope calibrator (Capintec, Inc., Ramsey, NJ) for determination of mCi and an automated well scintillation Beckman 8000 gamma counter (Irvine, CA) for counts per minute. ⁶⁴Cu was produced at Washington University School of Medicine on a CS-15 biomedical cyclotron using previously published methods [33] and ⁶⁴Cu-ATSM was prepared as previously described [21]. Human prostate carcinoma tumor cell lines PC-3, LNCaP, and 22Rv1 were obtained from American Type Culture Collection (Manassas, VA) and LAPC-4 cells were a gift from Dr. Charles Sawyers at UCLA. Cells were maintained by serial passage in cell culture. LAPC-4 was shown in our laboratories to have relatively low FAS expression compared with the other prostate tumor cell lines and was therefore chosen as a 'negative' control.

In vitro ⁶⁴Cu-ATSM uptake

A comparison of ⁶⁴Cu-ATSM uptake under anoxic and normoxic conditions was made in three prostate cell lines – 22Rv1, LNCaP, and PC-3. The apparatus and methods used in the in vitro uptake study are based on those previously described in the literature [20]. Viability of the cells was measured with a hemocytometer utilizing trypan blue staining. Briefly, two three-neck flasks containing cells in suspension were immersed in a 37°C water bath by a rod connected to a stand rotating on an external orbital mixer. An anoxic (5% CO₂, 95% N₂) and normoxic (20% O₂, 5% CO₂, 95% N₂) gas mixture were humidified and brought to 37°C before being passed continuously through each of the flasks. 30-50 mL of cell suspension $(1 \times 10^6 \text{ cells/mL})$ was added to each flask and given 1 hour to equilibrate to the environment. After equilibration, 100 µCi ⁶⁴Cu-ATSM was added to each of the flasks in a small amount (~5 µL) of ethanol. At 1, 5, 15, 30, 45, and 60 minutes, 200 µL of cell suspension was removed via pipet and placed in a 1.5 mL Eppendorf tube which was immediately centrifuged for 30 s at 3000 rpm to pellet the cells. 180 μ L of the supernatant was then pipetted off and deposited in a separate tube. This was done in triplicate at each timepoint. All pellet and supernatant samples were counted on a gamma counter. Percent uptake was calculated as activity in the pellet (corrected for remaining supernatant) divided by the sum of the pellet and supernatant.

C75 treatment studies were performed in four prostate cell lines – 22Rv1, LNCaP, PC-3, and LAPC-4 following similar methods. In this case, both flasks were under anoxic conditions, while one flask received C75 for FAS inhibition. During the hour of equilibration, C75 was added to the cells (in 10 μ L or less of DMSO) so that the final concentration in the flask was 100 μ M, while the other flask received DMSO alone. At 1, 5, 15, 30, 45, and 60 minutes the same procedure of sample collection and analysis was performed.

FAS expression by ELISA

FAS expression of each tumor line was quantified using a FAS-detectTM ELISA kit (FASgen, Inc.) based on a two-site ELISA technique to quantitatively measure FAS in human serum. Cells were pelleted and lysed by addition of 1 mL of cell lysis buffer containing protease inhibitors and incubated on ice for 30 minutes. The cell debris was then repelleted and the final supernatant was analyzed. Results were compared to a standard curve of varying concentrations of FAS to determine the concentration in ng/mL. These values were normalized by protein concentration as determined by using a standard copper reduction/bicinchoninic acid assay (Pierce) with bovine serum albumin as the standard, resulting in a ratio with units of ng FAS/ μ g protein.

In vitro C75 dose response on ⁶⁴Cu-ATSM uptake

These *in vitro* studies were performed using the same apparatus and methods as previously described in the initial *in vitro* experiments. In this study however, various amounts of C75 were added to the cells (in 10 μ L or less of DMSO) so that the final concentration of C75 in each flask was 100, 20, 4, or 0.8 μ M.

Results

For each cell line, two three-neck flasks containing cells (35 mL) in suspension were placed under anoxic and normoxic conditions with or without pretreatment with C75 (100 μ M). Uptake of ⁶⁴Cu-ATSM was measured over time to determine if FAS inhibition (by C75) restored the cells' ability to trap Cu-ATSM under low oxygen conditions. Inhibition of FAS with C75 resulted in a dramatic increase in ⁶⁴Cu-ATSM retention into prostate tumor cells *in vitro* under anoxic conditions (Fig. 1). The treated cells demonstrated higher uptake values at 15 minutes of 20.93 ± 3.271, 103.0 ± 32.57, 144.2 ± 32.28, and 200.1 ± 79.27 % over control values for LAPC-4, PC-3, LNCaP, and 22Rv1 cell lines, respectively. These values would later be correlated to FAS expression of each cell line.

Ratios of anoxic to normoxic ⁶⁴Cu-ATSM uptake demonstrate the selectivity of the tracer in a particular model, and so these were also compared over the 60 minute experiment (Fig. 2, 15 min data shown). The value obtained for the retention of ⁶⁴Cu-ATSM into EMT-6 murine mammary carcinoma cells at 15 min is included for comparison [26]. The 15 minute time point was chosen since Cu-ATSM in these cells was shown to plateau at this point (Fig. 1). Similar kinetics were shown in Lewis et al., [26] and Dearling et al., 1998 [20]. Also given that this is entirely an *in vitro* study, the potential blood flow issues associated with *in vivo* studies was not felt to be a factor. It is also worth pointing out that in the human ⁶⁰Cu-ATSM studies uptake in tumors [22, 23] resulted in high contrast levels between hypoxic and normoxic tissues by as little as 10-15 minutes post injection, and yielded clinically relevant information about tumor oxygenation that was predictive of tumor behavior and response to therapy. In this current study, values in all three of the cell lines examined under normoxic conditions, were lower than the EMT-6 ratios suggesting that visualization of solid tumors, derived from these prostate cancer cell lines may be difficult due to normoxic background retention in the surrounding tissue.

Samples of each cell line were analyzed to determine if relative amounts of FAS were related to the magnitude of Cu-ATSM retention at 15 min post-incubation. Quantification of FAS expression, by ELISA, resulted in values of 0.00438 ± 0.00024 , 0.00832 ± 0.00038 , 0.01877 ± 0.00092 , and 0.02679 ± 0.00224 ng FAS/µg total protein for LAPC-4, PC-3, LNCaP, and 22Rv1, respectively. As expected, a correlation is seen (R² = 0.911) with FAS expression plotted against the change in hypoxic retention resulting from inhibition of FAS (Fig. 3).

The dose response effect of C75 on the restoration of hypoxic retention of Cu-ATSM was measured in PC-3 cells utilizing the same cell suspension apparatus and anoxic gas mixture. During a 1-hour equilibration period, the four flasks were treated with varying concentrations of C75 (100, 20, 4, or 0.8 μ M). Confirming the effect of FAS inhibition on ⁶⁴Cu-ATSM uptake, the experiment showed increasing retention directly related to C75 concentration with values of 0.0037 ± 0.0002, 0.0030 ± 0.0001, 0.0026 ± 0.0001, and 0.0024 ± 0.0001 % uptake/µg protein for 100, 20, 4, and 0.8 μ M respectively at 15 mins (Fig. 4).

Discussion

Low oxygen levels are a hallmark of malignant processes, leading to a cascade of oncogenic properties that support tumor growth and invasion [1-3, 34, 35]. Existing in this harsh environment, the cells go into survival mode and do not easily respond to DNA damaging chemotherapeutics or agents that promote apoptosis [6, 7]. In the absence of oxygen, the damaging radicals created by traditional radiation therapy recombine resulting in little damage to the target tissue [3-5]. Levels of tumor hypoxia have also been correlated to overall patient survival and tumor aggressiveness [1, 2, 36-38].

⁶⁴Cu-ATSM [19], which targets hypoxia as opposed to metabolism, has already been confirmed as a clinically important PET agent in the prediction of prognosis in several cancers. It has been shown to be a useful clinical PET imaging tracer that can distinguish responders to conventional therapies from non-responders in patients with lung [23], cervical [22], rectal [24] and head and neck (data not published) cancer. It has also been suggested as a potential tool for radiation oncologists to tailor the radiation dose based on hypoxic regions, giving the resistant regions an increased dose [39].

Recently, Yuan et al, showed that the regional ⁶⁴Cu-ATSM retention in two tumors types (R3230Ac and 9L) correlated closely with intrinsic markers for hypoxia, but in a third tumor type (FSA) a correlation was not observed [40]. The suggested reason for this low correlation between Cu-ATSM uptake and hypoxic distribution was the differing redox statuses of the tumor types. FSA tumors may have a lower-than-average redox potential with high concentrations of electron donors. If the speculated mechanism of Cu-ATSM uptake holds, then reduction, release and ultimate cellular trapping of Cu(I) would be caused by a decrease in the redox potential of the cell [19, 21, 27, 41, 42]. In the FSA tumor line, the lower-than-average redox potential caused reduction and trapping of ⁶⁴Cu-ATSM in both hypoxic and normoxic areas. It was also suggested that the higher perfusion levels surrounding the hypoxic regions of the FSA tumor could contribute to the higher level of retention of Cu-ATSM in normoxic FSA cells. In vitro and in vivo experiments have also revealed inconsistencies in the hypoxic selectivity of Cu-ATSM in prostate cancer models [27, 28]. Although a very limited number of other tumor cell lines such as FSA have shown inconsistencies, prostate tumors clearly stand out as the primary tumor type that Cu-ATSM hypoxia-selectivity is reduced to low or even undetectable levels [27, 28]. This current study was undertaken in human prostate tumor lines to better examine the reasons for the lack of hypoxia-selectivity of Cu-ATSM in prostate tumors.

Biologically, prostate tumors have some unique characteristics when compared to other malignancies. Prostate cancer is known to have low metabolism, and the most common PET tracer for detecting malignancies, ¹⁸F-FDG, is not effective in delineating it from surrounding tissue [43] and also goes to sites of inflammation. Fatty acid synthase (FAS) is a multi-functional enzymatic protein involved in many stages of fatty acid synthesis and has been found to be overexpressed in prostate carcinomas as well as other cancers [44-52]. It is an androgen-regulated enzyme that is overexpressed in the vast majority of prostate tumors

and its expression defines distinct molecular signatures in prostate cancer [53]. An increased level of FAS has been found to be indicative of aggressive and late-stage prostatic adenocarcinomas [31] as well as a prognostic indicator for overall survival [54].

As a defense mechanism in prostate cancer cells, the fatty acid synthesis pathway harnesses its oxidizing power for improving the redox balance (i.e., lower NADH/NAD⁺ ratios) despite conditions of extreme hypoxia [32]. This pathway is able to consume reducing equivalents (i.e., NADPH) as part of its normal processes. Under hypoxic conditions, anaerobic glycolysis creates excessive levels of lactate, which limits the respiratory chain and reduces its oxidizing power. The relationship between the fatty acid synthesis pathway and prostate tumors has been superbly reviewed by Hochachka et al., [32]. In regards to the hypoxia-selectivity of Cu-ATSM, it has been shown that Cu-ATSM enters all cells by passive diffusion, and under normoxic conditions freely exits the cell. Under hypoxic or anoxic conditions, the redox potential of the cell is altered such that the Cu(II) in the complex is reduced to Cu(I) and the complex falls apart [21, 27, 41, 42]. This retention mechanism is reliant on the reduction of the Cu(II) to Cu(I), and if cellular reducing equivalents such as NADPH are consumed by the overexpression of the fatty acid synthesis pathway this chemical reduction, even under conditions of hypoxia, is unlikely to take place.

C75 is a small molecule that binds to and inhibits mammalian FAS and inhibits fatty acid synthesis in human cancer cells. Researchers have shown that C75 inhibits FAS by 89-95% [55]. In this study, C75 was used to inhibit FAS activity in prostate tumor cell lines by blocking this enzymatic cycle's ability to offset the redox balance of a hypoxic cell, therefore, increasing retention of Cu-ATSM under hypoxic conditions. Typically a blocking study is designed to inhibit uptake of the radiotracer; in this case we were attempting to block the FAS so that it is unable to 'stabilize' the redox environment of the cell. Its inhibition should allow the anticipated retention of Cu-ATSM in hypoxic regions of the tumor. In vitro results in all cell lines demonstrate that FAS has a significant impact on the retention of ⁶⁴Cu-ATSM into human prostate tumor cells under oxygen-limited conditions. Inhibition of FAS with a single 1 hour treatment of C75 (50 μ M) resulted in increased retention of Cu-ATSM in all four tumor lines tested in vitro (Fig.1), but to a limited extent in the low-FAS-expressing LAPC-4. Results also showed that a linear change in uptake correlated directly to the cellular expression of FAS (Fig 3.). To demonstrate that restoration of retention is directly related to FAS inhibition, a dose response experiment was undertaken (Fig. 4).

Conclusion

Cu-ATSM has been a valuable tool as a marker of hypoxia, and it's possible that the nature of its mechanism of retention, though not completely understood, could lead us to answers for the incongruences in its selectivity. The uncertainty lies in the possibility that other physiological changes caused by malignant progression and hypoxia within the cell could also affect this redox balance regardless of oxygen concentration. If the suggested mechanism of Cu-ATSM uptake holds, then reduction, release and ultimate cellular trapping of Cu(I) would be caused by a decrease in the redox potential of the cell. And, although in most cases this is directly due to a change (reduction) in oxygen concentration within the cell, there are many other processes that could also alter the redox potential of this environment. In prostate tumors, a reduction of ⁶⁴Cu-ATSM hypoxia-selectivity is demonstrated in a manner that is related to FAS expression. However, if a reduction in Cu-ATSM hypoxia-selectivity is demonstrated in non-prostate tumor lines it cannot be assumed that FAS is involved and each line must be studied on a case-by-case basis. In conclusion, ⁶⁴Cu-ATSM is a very effective PET agent for clinically delineating many

hypoxic human malignancies, but, as with all radiopharmaceuticals, it is not a universal agent. Care should be taken in particular regard to the imaging of prostate tumors.

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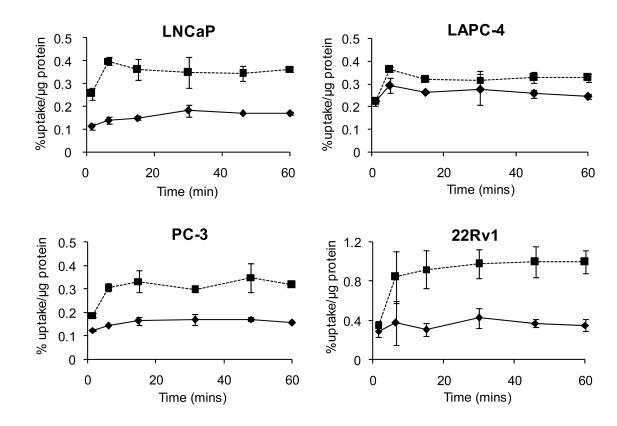


Figure 1.

Uptake of ⁶⁴Cu-ATSM into prostate tumor cells (in suspension) under anoxic conditions (5% $CO_2 + 95\% N_2$) treated with C75 (dashed) and compared to control (solid). (n = 3 at each timepoint)

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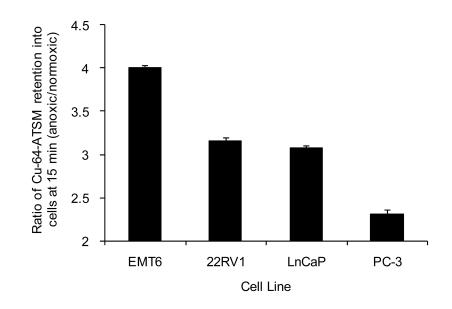


Figure 2.

Ratios of anoxic to normoxic ⁶⁴Cu-ATSM uptake in prostate cancer cells lines at 15 mins demonstrating differing selectivity of the tracer in different models. The value obtained for the retention of ⁶⁴Cu-ATSM into EMT-6 murine mammary carcinoma cells at 15 mins is included for comparison [26].

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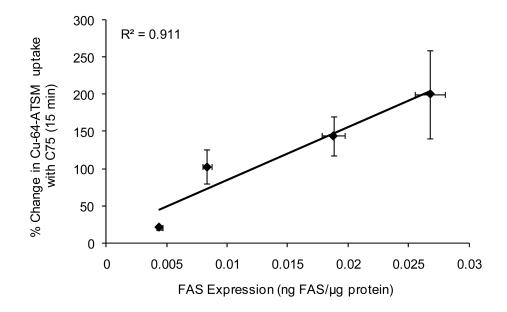


Figure 3.

Correlation of FAS expression as determined by ELISA to % change in ⁶⁴Cu-ATSM uptake between C75 treated (100 μ M) and control cells under anoxic conditions (5% CO₂, 95% N₂) at 15 min post-incubation. Points on the graph represent LAPC-4, PC-3, LNCaP, and 22Rv1 from left to right. (n = 3 at each timepoint)

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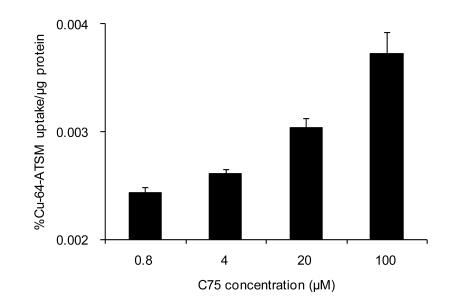


Figure 4.

Dose response of ⁶⁴Cu-ATSM uptake at 15 mins (under anoxic conditions) in PC-3 cells following FAS inhibition by C75. p = 0.0026 (100-20), p = 0.0002 (20-4).