

Material and Methods: We studied the autophagic properties of human PC-3 prostate and MDA-MB-231 breast cancer models of acquired, stable resistance to metronomic cyclophosphamide therapy, compared to their parental counterparts. We also analyzed the anti-tumour effects of metronomic versus conventional cyclophosphamide \pm chloroquine (autophagy inhibitor) therapy on PC-3 xenografts. Furthermore, we compared the *in vivo* growth properties of paired autophagy-competent and autophagy-deficient (i.e., beclin1 haploinsufficient) baby mouse kidney epithelial cells treated with either metronomic or conventional cyclophosphamide.

Results: LC-3 Western blotting and acridine orange flow cytometry of parental PC-3 and MDA-MB-231 revealed strong autophagy induction under conditions of metabolic stress mimicking the microenvironment in tumours undergoing metronomic cyclophosphamide therapy (i.e., hypoxia, low pH and reduced nutrients). In contrast, the autophagic response was reduced in a number of metronomic cyclophosphamide resistant PC-3 and MDA-MB-231 variants. Chloroquine impaired the response of PC-3 xenografts to metronomic cyclophosphamide. Similarly, the impact of metronomic cyclophosphamide was reduced in autophagy-deficient versus autophagy-competent baby mouse kidney epithelial cell allografts. In contrast, both pharmacological and genetic autophagy deficiency enhanced the antitumour effects of conventional cyclophosphamide.

Conclusions: Our studies suggest that impaired autophagy contributes to resistance to metronomic cyclophosphamide chemotherapy, and possibly to other forms of antiangiogenic or chronic anticancer therapies. In other words, chronic metabolic stress associated with metronomic chemotherapy may favor cell death promoting autophagy effects. In contrast, cell survival promoting autophagy effects prevail during acute cellular stress due to conventional chemotherapy. Thus, the impact of autophagy modulators in clinical development may vary dramatically depending on the nature of concomitant anticancer therapies.

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POSTER

The Influence of the Combined Treatment With Vadimezan (ASA 404) and Taxol on the Growth of U251 Glioblastoma Xenografts

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Background: One of the most important biological characteristics of Glioblastoma multiforme (GBM) is high vascular density and targeting the vasculature in this tumour could be an attractive therapeutic strategy. Vadimezan (ASA 404, DMXAA) belongs to the class of small molecule vascular disrupting agents (VDA) that cause disruption of established tumour vessels and subsequent tumour hemorrhagic necrosis. Selective antivascular effects of ASA 404 are mediated by intratumoral induction of several cytokines including tumour necrosis factor- α (TNF- α), granulocyte-colony-stimulating factor (G-CSF), interleukin 6 (IL-6) and macrophage inflammatory protein 1 α (MIP-1 α). Preclinical studies have demonstrated that ASA 404 acts synergistically with Taxanes, which at lower concentrations inhibit angiogenesis. In this study, we investigated if treatment of mice bearing U251 human glioblastoma xenografts with ASA 404 and taxol may be synergistic. Therapy response was evaluated also by FDG-PET imaging.

Material and Methods: 1.5×10^6 U251 cells were inoculated s.c. into the right hind limb of NMRI-Foxn1^{nu} athymic female nude mice. Animals were randomly assigned in 4 groups (7–9 animals/group) for treatment: control, taxol, ASA 404 and ASA 404 plus taxol. The animals received either a single dose of taxol (10 mg/kg), ASA 404 (27.5 mg/kg), or taxol (10 mg/kg) plus ASA 404 (27.5 mg/kg) administered i.p.; ASA 404 was administered 24 hours after the treatment with taxol. 4 hours after treatment with ASA 404 (28 hours after treatment with taxol) FDG-PET scans were performed.

Results: The treatment with taxol did not affect the tumour growth in comparison to untreated controls. The treatment of animals with single dose ASA 404 alone or in combination with taxol caused a significant decrease in tumour volume. The combined treatment did not decrease the growth of the xenografts significantly more than ASA 404 alone. The final tumour weights were: control = 764 ± 168 mg, taxol = 651 ± 148 mg, ASA 404 = 283 ± 127 mg, ASA 404 + taxol = 180 ± 56 mg. FDG-PET imaging correlated with tumour response. SUV values were: control = 1.21 ± 0.39 , taxol = 1.14 ± 0.19 , ASA 404 = 0.36 ± 0.08 and ASA 404 + taxol = 0.50 ± 0.14 .

Conclusion: The treatment with ASA 404 alone or in combination with taxol showed antitumour effects in our glioblastoma model probably through destruction of blood vessels. The implications for the anticancer effect of this compound warrant further preclinical studies.

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POSTER

Overcoming the Acquired Resistance to Afatinib (BIBW2992) in HCC827, a Non-small Cell Lung Cancer Cell Line

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Understanding of the pharmacological responses to drug treatment in cancer cells is essential for discovery and development of novel anti-cancer therapies. In this study, drug resistant cell lines, HCC827-BR1 and HCC827-BR2, were developed by treatment of HCC827 cells with escalating concentration of afatinib (BIBW2992). The CC₅₀ of BIBW2992 in HCC827 ranges from 2 to 10 nM while the CC₅₀s of BIBW2992 in HCC827-BR1 and HCC827-BR2 are approximately 10 μ M. Gene expression analysis revealed that the epithelial-mesenchymal transition (EMT) may be involved in resistance to BIBW2992. The drug-resistant cells are more invasive as evaluated under *in vitro* assays. Results from this study have also identified that the drug-resistance cells are more sensitive to another kinase inhibitor; indicating that an oncogenic shift has occurred. When this drug is combined with BIBW2992 in treatment of HCC827 cells, much less colonies survived compared to cells treated by BIBW2992 alone. The clinical ramifications of these observations will be discussed.

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POSTER

Doxorubicin and Taxol Induce Apoptosis in Breast Cancer Cells by Activating Foxo3a Transcription Factor

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Background: Foxo3a is a member of the forkhead box O class of transcription factors (foxO), which are key regulators of apoptosis, cell cycle arrest and cell division. Foxo3a phosphorylation by kinases, such as Akt, leads to its nuclear exclusion, cytoplasmic accumulation and subsequent degradation. Foxo3a inactivation has been associated with tumorigenesis and poor survival in breast cancer, what turns this protein into a possible target for anticancer drugs. In this study, we aimed to investigate the role and regulation of Foxo3a in response to taxol and doxorubicin (doxo) treatment in breast cancer cells.

Materials and Methods: The human breast carcinoma cell lines MCF7 and MDA-MB-231 were exposed to clinically relevant concentrations of taxol and doxo and cytotoxicity was assessed by the MTT assay. Cell morphological changes were microscopically photographed. Western blot and Annexin V/PI by flow cytometry were used to detect caspases activation and apoptosis, respectively.

Results: After taxol exposure for 24h, there was an 85% cell viability inhibition, which was also observed after doxo treatment for 72h ($p < 0.01$), showing that both drugs display high toxicity against breast cancer cells. Morphological analysis of non-adherent cells revealed that the drugs induced 65% of cell death ($p < 0.05$), indicating that cytotoxicity was not resulted from cell proliferation inhibition. Taxol and doxo could effectively induce apoptosis, as detected by the Annexin-V/PI method and caspases-3, -7 and -9 activations. Western blot analysis showed that there was a 16 and 12-maximum fold increase in Foxo3a levels after treatment with taxol and doxo, respectively. However, the Real Time PCR analysis of mRNA Foxo3a expression in cells exposed to the drugs showed that Foxo3a levels were not increased, indicating that Foxo3a expression is not transcriptionally activated. This finding suggests that taxol and doxo may induce cellular mechanisms which prevent Foxo3a degradation. Data analysis was done using the t student test and a $p < 0.05$ was considered statistically significant.

Conclusion: The association between the increase in Foxo3a expression and cells sensitivity indicates that this transcription factor may act mediating taxol and doxo-induced apoptosis. These data point Foxo3a as a cellular target for anticancer drugs in breast cancer cells.

Acknowledgements: FINEP, Swissbridge Foundation and INCT para Controle do Câncer, CNPq 573806/2008-0; FAPERJ E26/170.026/2008.

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POSTER

Influence of Soluble CD40 Ligand on Colorectal Cancer Cells: a Flow Cytometric Study

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Background: The cell surface costimulatory molecule CD40 is a member of the Tumour Necrosis Factor Receptor family widely expressed on various