Serotonin activates glycolysis and mitochondria biogenesis in human breast cancer cells through activation of the Jak1/STAT3/ERK1/2 and adenylate cyclase/PKA, respectively

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BACKGROUND: Although produced by several types of tumours, the role of serotonin on cancer biology is yet to be understood.

METHODS: The effects of serotonin (5-HT) on human breast cancer cells proliferation, signalling pathways and metabolic profile were evaluated by cytometry, western blotting, qPCR, enzymology and confocal microscopy.

RESULTS: Our results revealed that incubation of MCF-7 cells with 10 µM 5-HT increased cell growth rate by 28%, an effect that was prevented by the 5-HTR2A/C antagonist, ketanserin. Conversely, increasing concentrations of 5-HT promoted glucose consumption and lactate production by MCF-7 cells. We also showed that increased glucose metabolism is provoked by the upregulation of pyruvate kinase M2 (PKM2) isoform through 5-HTR2A/C-triggered activation of Jak1/STAT3 and ERK1/2 subcellular pathways. However, we noticed a decrease in the rate of produced lactate per consumed glucose as a function of the hormone concentration, suggesting a disruption of the Warburg effect. The latter effect is due to 5-HTR2A/C-dependent mitochondrial biogenesis and metabolism, which is triggered by adenylyl cyclase/PKA, enhancing the oxidation of lactate within these cells.

CONCLUSIONS: We showed that serotonin, through 5-HTR2A/C, interferes with breast cancer cells proliferation and metabolism by triggering two distinct signalling pathways: Jak1/STAT3 that boosts glycolysis through upregulation of PKM2, and adenylyl cyclase/PKA that enhances mitochondrial biogenesis.

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cancer cell survival necessitates shifting to oxidative metabolism, the latter that is correlated to epithelial-to-mesenchymal transition, cancer invasion and aggressiveness.\textsuperscript{19,20}

We have previously demonstrated that 5-HT triggers glycolysis in diverse tissues by activating glycolytic enzymes, which involves 5-HT\textsubscript{2A} receptor activation.\textsuperscript{9–11} Since breast tumour cells produce and secrete 5-HT, we hypothesised that the hormone might have an autocrine action on these cells by modulating their energy metabolism. In the current work, we used the human breast cancer cell line MCF-7 to evaluate 5-HT effects on cancer metabolism. We compared the results with other cell lines, the non-tumorigenic MCF10A and the potentially metastatic MDA-mb-231. We showed that 5-HT modulates breast cancer biology through interfering with mitochondria biogenesis and shifting the fate of glycolysis products to oxidative metabolism, thus altering aerobic glycolysis and mitochondrial metabolism.

**METHODS**

Materials

Human breast cancer cell lineages MCF-7 and MDA-mb-231 were, respectively, obtained from the Cell Bank of Rio de Janeiro (www.bcrj.org.br Duque de Caxias, RJ, Brazil) and gifted by Dr. Mitzi Brentani (USP, São Paulo, Brazil), and were grown and maintained at 37 °C (5% CO\textsubscript{2} atmosphere) in Dulbecco’s Modified Eagle’s Medium (DMEM) with 25 mM glucose supplemented with 10% (vol/vol) heat-inactivated foetal bovine serum (FBS) and 5 mM L-glutamine (Invitrogen, São Paulo, SP, Brazil). The non-tumorigenic human breast cell line MCF10A was kindly supplied by Dr. M. Brentani (USP, São Paulo, Brazil), and was grown and maintained at 37 °C (5% CO\textsubscript{2} atmosphere) in DMEM/F12 medium with 25 mM glucose supplemented with 10% (vol/vol) FBS, 0.02 mg/ml EGF, 5 mg/ml insulin, 1.25 mg/ml hydrocortisone, 0.1 mg/ml cholera toxin and 5 mM L-glutamine (Invitrogen, São Paulo, SP, Brazil). All the chemicals used were of the highest quality commercially available.

Cell proliferation rate, apoptosis and viability

To evaluate the cell proliferation rate, cells were seeded in 24-well plates (4 × 10\textsuperscript{4} cells/well) and incubated for 3 h to adhere. After this short incubation, cells were treated with 5-HT (Sigma Chemicals Co., St. Louis, MO, USA) and/or ketanserin (Sigma Chemicals Co., St. Louis, MO, USA) and/or spiperone (Sigma Chemicals Co., St. Louis, MO, USA), accordingly, and then incubated for 18 h until reaching almost 50% confluency. Then, cells were treated with PE Annexin V reagent (1:20 dilution; BD Pharmingen, BD Biosciences, Franklin Lakes, NJ, USA) and 5 µM 7-AAD (BD Pharmingen, BD Biosciences, Franklin Lakes, NJ, USA), and the proliferation rate, apoptosis and cell viability were evaluated by counting the number of cells negative for both markers (viable), positive for Annexin V (apoptotic) and positive for 7-AAD staining (non-viable) by using a cell-sorting system (Muse\textsuperscript{®} Cell Analyzer, Merck Millipore, Billerica, MA, USA).

Glucose consumption and lactate production

To evaluate glucose consumption and lactate production, cells were seeded in 96-well plates (8 × 10\textsuperscript{3} cells/well) and grown to 70% confluency. Then, the medium was replaced by a fresh one containing 2-(N-(7-nitrobenz-2-oxa-1,3-diazol-4-yl)amino)-2-deoxyglucose (2-NBDG; 1 mM/mol glucose) to trace glucose consumption or not containing the tracer for lactate production. After 24-h incubation, the medium was removed to evaluate the amount of produced lactate, and cells were harvested and resuspended in phosphate-buffered saline (PBS; 137 mM NaCl, 2.7 mM KCl, 10 mM Na\textsubscript{2}HPO\textsubscript{4} and 1.8 mM KH\textsubscript{2}PO\textsubscript{4}, pH 7.4) containing 0.1% bovine serum albumin (BSA, heat-shock fraction, fatty acid free, protease free, Sigma-Aldrich, St. Louis, MO, USA). Suspended cells were evaluated for glucose content by using a cell cytometer (Muse Cell Analyzer, Merck Millipore, Billerica, MA, USA). A calibration curve was performed to determine the amount of glucose taken up by the cells. For determination of lactate production, the amount of lactate in the culture medium was evaluated by using a commercial kit (Labtest Diagnóstica S/A, Lagoa Santa, MG, Brazil). The lactate production/glucose consumption ratio was obtained by simply the product of these two measurements.

Mitochondrial activity

Mitochondrial activity was assessed by the mitochondrial reductive potential via incubating the cells in the presence of 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT, Sigma Chemicals Co., St. Louis, MO, USA) and evaluating the formation of formazan products. Briefly, cells were seeded in 96-well plates (8 × 10\textsuperscript{3} cells/well) and grown to 70% confluency; then cells were treated with different concentrations of 5-HT for 24 h. After this period, the medium was removed and a solution of 5 mg/ml MTT was added followed by 3-h incubation at 37 °C. Then, the medium was removed, and 100 µl of DMSO was added to dissolve the formazan crystals formed, which was evaluated by measuring the absorbance at 560 nm subtracted by the absorbance at 670 nm.

Enzymatic activities

The enzymatic activities were evaluated after the treatment for 24 h of cells seeded in 96-well plates and grown to 70% confluency. Following treatment, the medium was removed, and cells were lysed with 100 µl of 10 mM phosphate buffer (pH 7.4) by pipetting up and down several times. The cell-free homogenates were used for evaluating the enzymes’ activities. Hexokinase activity was assessed in a reaction medium containing 50 mM Tris-HCl (pH 7.4), 5 mM Mg\textsubscript{2+}, 1 mM glucose, 1 mM ATP, 150 mM KCl, 0.2 mM NADP\textsuperscript{+} and 0.2 mM/L glucose-6-phosphate dehydrogenase. Phosphofructokinase activity was assessed in a reaction medium containing 50 mM Tris-HCl (pH 7.4), 1 mM Mg\textsubscript{2+}, 1 mM (NH\textsubscript{4})\textsubscript{2}SO\textsubscript{4}, 1 mM fructose-6-phosphate, 0.1 mM ATP, 150 mM KCl, 0.2 mM NADH, 0.2 µM/ml aldolase, 0.4 mM/ml triosephosphate isomerase and 0.2 mM/ml α-glycerophosphate dehydrogenase. Pyruvate kinase activity was assessed in a reaction medium containing 50 mM Tris-HCl (pH 7.4), 1 mM Mg\textsubscript{2+}, 1 mM phosphoenolpyruvate, 1 mM ADP, 150 mM KCl, 0.2 mM NADH and 2 mM/l lactate dehydrogenase. Lactate dehydrogenase activity: 50 mM Tris-HCl (pH 7.4), 1 mM Mg\textsubscript{2+}, 1 mM pyruvate and 0.2 mM NADH. Glucose-6-phosphate dehydrogenase activity was assessed in a reaction medium containing 50 mM Tris-HCl (pH 7.4), 5 mM Mg\textsubscript{2+}, 1 mM glucose-6-phosphate, 150 mM KCl and 0.2 mM NADP\textsuperscript{+}. Reactions were started by adding 10 µl of the cell-free homogenates, and NAD(P) reduction/oxidation was followed spectrophotometrically at 340 nm. The slopes of the curves were used to determine the enzymatic activities. Succinate dehydrogenase was assessed in a reaction medium containing 50 mM Tris-HCl (pH 7.4), 5 mM Mg\textsubscript{2+}, 12 mM diethyl succinate, 0.2 mM 1-methoxy-5-methylphenazinium methyl sulfate and 1.2 mM nitro blue tetrazolium (NBT), in the absence and the presence of 12 mM malonate (a competitive inhibitor of succinate dehydrogenase). The reaction was started by adding 10 µl of cell-free homogenate, and the reduced NBT was evaluated spectrophotometrically at 600 nm. Succinate dehydrogenase activity was calculated by the slope of the curve.

Quantitative PCR (qPCR)

For RNA extraction, cells were seeded in six-well plates (10\textsuperscript{5} cells/well) and grown to 70% confluency. Then the media were removed, and cells were treated according to the experiments. After the treatments, the media were removed, and 500 µl of Trizol reagent (ThermoFisher, Carlsbad, CA, USA) was added to each well, followed by homogenisation by up and down pipetting. After
Table 1. Primers used for qPCR

| Gene          | Protein Description                  | Primer sequence | Amplicon length (bp) | Efficiency (%) |
|---------------|--------------------------------------|-----------------|----------------------|----------------|
| CCSER2        | Coiled-coil serine-rich protein 2    | TGAGCAGAGGCTCTCCCTCCA | 72                  | 99.5           |
| HK1           | Hexokinase 1                         | AGCAGCTCTCCAGGTCGGCCT | 108                 | 101.2          |
| HK2           | Hexokinase 2                         | GCTGCTCTCACTTCATGG | 361                 | 104.2          |
| HTR1A         | 5-hydroxytryptamine receptor 1A      | CGAATCTGCTGGCCTCCCT | 97                  | 97.3           |
| HTR1B         | 5-hydroxytryptamine receptor 1B      | CGTCGGATTTGATGATGTT | 91                  | 96.8           |
| HTR1D         | 5-hydroxytryptamine receptor 1D      | CTACACCAAGGCTCCACAC | 73                  | 98.3           |
| HTR1E         | 5-hydroxytryptamine receptor 1E      | GAACTGAGAGAGGCTGAGATG | 84                 | 95.4           |
| HTR1F         | 5-hydroxytryptamine receptor 1F      | GCTGAGAGGCTGAGATGAG | 100                | 97.8           |
| HTR2A         | 5-hydroxytryptamine receptor 2A      | GATTAGGAGGCTGAGATGAAG | 85                 | 101.1          |
| HTR2B         | 5-hydroxytryptamine receptor 2B      | TGGTTGGGTTTATGCATCAGC | 74                 | 99.3           |
| HTR2C         | 5-hydroxytryptamine receptor 2C      | CTACACCAAGGCTCCACAC | 92                  | 95.9           |
| HTR3A         | 5-hydroxytryptamine receptor 3A      | TACACCAAGGCTCCACACAC | 76                 | 96.5           |
| HTR3B         | 5-hydroxytryptamine receptor 3B      | GCTGAGAGGCTGAGATGAG | 83                  | 98.8           |
| HTR3C         | 5-hydroxytryptamine receptor 3C      | GCTGAGAGGCTGAGATGAG | 83                  | 100.8          |
| HTR3D         | 5-hydroxytryptamine receptor 3D      | TACACCAAGGCTCCACACAC | 70                 | 102.1          |
| HTR3E         | 5-hydroxytryptamine receptor 3E      | GCTGAGAGGCTGAGATGAG | 70                  | 99.2           |
| HTR4          | 5-hydroxytryptamine receptor 4       | GCTGAGAGGCTGAGATGAG | 100                 | 97.2           |
| HTR5A         | 5-hydroxytryptamine receptor 5A      | TACACCAAGGCTCCACACAC | 83                 | 95.9           |
| HTR5B         | 5-hydroxytryptamine receptor 5B      | GCTGAGAGGCTGAGATGAG | 70                  | 99.7           |
| HTR6          | 5-hydroxytryptamine receptor 6       | GCTGAGAGGCTGAGATGAG | 71                  | 96.9           |
| LDHA          | Lactate dehydrogenase A              | GCTGAGAGGCTGAGATGAG | 110                 | 97.8           |
| LDHB          | Lactate dehydrogenase B              | GCTGAGAGGCTGAGATGAG | 93                  | 96.4           |
| PKD1          | Pyruvate dehydrogenase 1             | GCTGAGAGGCTGAGATGAG | 103                 | 95.0           |
| PFKM          | PFK isoform M                        | GCTGAGAGGCTGAGATGAG | 171                 | 99.3           |
| PFKL          | PFK isoform L                        | GCTGAGAGGCTGAGATGAG | 118                 | 102.6          |
| PFKP          | PFK isoform P                        | GCTGAGAGGCTGAGATGAG | 187                 | 98.5           |
| PKCL          | PK L                                 | GCTGAGAGGCTGAGATGAG | 90                  | 97.7           |
| PKM1          | PK M                                 | GCTGAGAGGCTGAGATGAG | 110                 | 96.1           |
| PKM2          | PK M                                 | GCTGAGAGGCTGAGATGAG | 368                 | 97.8           |
| SLC16A1       | MCT1                                 | GCTGAGAGGCTGAGATGAG | 86                  | 98.8           |
| SDHA          | SDH A                                | GCTGAGAGGCTGAGATGAG | 105                 | 99.7           |
| SRT3          | SRT3                                 | GCTGAGAGGCTGAGATGAG | 78                  | 101.1          |

All the primers were optimised for 600 nM use in the experiments.
this, RNA extraction was performed by following the directions of the reagent. For cDNA synthesis, the high-capacity cDNA Reverse Transcription Kit (ThermoFisher, Carlsbad, CA, USA) was used. For qPCR, the GoTaq qPCR Master Mix (Promega, Fitchburg, WI, USA) was used and the reaction was performed in a QuantStudio 5 (ThermoFisher, Carlsbad, CA, USA). The programme for all amplifications was 2 min at 95 °C followed by 40 cycles of 15 s at 95 °C and 1 min at 60 °C. A dissociation curve was performed at the end of the experiment, and dissociation peak was analysed. The fold expression was calculated by the $2^{-\Delta\Delta Cq}$ method, as described previously. Expression levels are represented by the $2^{-\Delta Cq}$ method.

CCSER2 expression was used as reference gene (housekeeper), since its expression did not vary upon any of the used treatments (data not shown). Primers were designed by

![Graphs and images of cellular assays and gene expression](image)

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Cells were seeded in 24-well plates (4 × 10^4 cells/well) and expression, E
MCF-7 cells treated with different 5-HT concentrations in the absence or the presence of 0.1 µM ketanserin (5-HT2A/C receptor antagonist). These plotted values are the mean ± S.E.M. of six independent experiments (n = 6). For a, b, c and d, the data were analysed by two-way ANOVA, followed by Sidak’s post test and * means P < 0.05 compared with the black bar of the same group. For e, f and h, except for the results in the absence of 5-HT, the differences between each value in the absence and in the presence of ketanserin were significant (P < 0.05; two-way ANOVA, followed by Sidak’s post test). For PK activity, plotted values are the mean ± S.E.M. of six independent experiments (n = 6; * means P < 0.05, Student’s t test). k Representative western blot of evaluation of PKM2 expression. Quantification of the western blots of evaluation of PKM2 expression of three independent experiments (n = 3). Quantification (m) and representative western blots (o) of phosphorylation of ERK1/2 in MCF-7 cells treated with 5-HT and ketanserin (n = 3; * means P < 0.05 as compared with control in the absence of 5-HT, Student’s t test). q Pyruvate kinase activity in the absence or presence of 10 µM 5-HT and 50 nM LY3214996 (ERK1/2 inhibitor). The results for enzyme activity are from five independent experiments (n = 5, * means P < 0.05 as compared with control in the absence of LY3214996 and the bracket indicates P < 0.05 as compared with the absence of 5-HT; two-way ANOVA, followed by Sidak’s post test). Plotted results are expressed as mean ± S.E.M.

Confocal microscopy
For confocal microscopy experiments, MCF-7 cells were seeded on coverslips placed in 24-well plates and incubated to reach 70% confluency. The medium was removed, and cells were treated in the presence or absence of 10 µM 5-HT or 10 µM 5-HT plus 0.1 µM ketanserin for 24 h at 37°C. After treatment, 50 nM MitoTracker Green FM (Cat# M7514, Molecular Probes, Thermofisher, Carlsbad, CA, USA) and 100 nM MitoTracker Red CMXRos (Cat# M7512, Molecular Probes, Thermofisher, Carlsbad, CA, USA) were added to each well and incubated for 1 h at 37°C. After incubation, the coverslips were used to evaluate fluorescent staining inside the cells by using an A1 confocal microscope (Nikon, Tokyo, Japan) at ×63 magnification. Three independent fields from each coverslip were used for analyses of one experiment. Each image acquired was a z-stack of 12–16 (0.33-µm depth) sections. Maximum projections of three consecutive optical sections (corresponding to 1-µm total depth) were generated from the middle of the original z-stack. The complete analysis was taken by using three independent experiments (n = 3). Quantification of the images was performed by using the software Image J (http://imagej.nih.gov/ij, NIH, USA).

Mitochondrial evaluation by flow cytometry
Cells were seeded in 24-well plates (4 × 10^4 cells/well) and incubated for 3 h to adhere. After this short incubation, cells were treated accordingly, and then incubated for 24 h until reaching almost 70% confluency. Then, cells were treated with 50 nM MitoTracker Green FM (Cat# M7514, Molecular Probes, Thermofisher, Carlsbad, CA, USA) and 100 nM MitoTracker Red CMXRos (Cat® M7512, Molecular Probes, Thermofisher, Carlsbad, CA, USA) for 1 h at 37°C. Then, cells were detached and evaluated for the dye labelling by using a cell-sorting system (Muse® Cell Analyzer, Merck Millipore, Billerica, MA, USA).

Western blotting
For western blot, cells were seeded in six-well plates (10^5 cells/well) and grown to 70% confluency. Then the media were removed, and cells were treated according to the experiments. After the treatments, the media were removed and a mild-RIPA buffer supplemented with protease inhibitor cocktail (Sigma-Aldrich, St. Louis, MO, USA) was added for total protein extraction. Protein extracts were submitted to SDS-PAGE (8% gels), followed by overnight transfer to nitrocellulose membranes at 30 V. Membranes were stained with Ponceau S, processed and destained by washing with distilled water. Then, the membranes were incubated overnight with the following antibodies: anti-5-HT2C receptor (dilution 1:1000, Cat® ab197776, Abcam, Cambridge, UK), anti-5-HT7 receptor (dilution 1:100, Cat#ab61562, Abcam, Cambridge, UK), anti-ß-actin (dilution 1:1000, Cat# 4967, Cell Signaling Technology, Danvers, MA, USA), anti-Akt (dilution 1:1000, Cat# 9272, Cell Signaling Technology, Danvers, MA, USA), anti-phospho-Akt (T308) (dilution 1:1000, Cat# 9275, Cell Signaling Technology, Danvers, MA, USA), anti-CREB (dilution 1:1000, Cat# 9197, Cell Signaling Technology, Danvers, MA, USA), anti-phospho-CREB (S133) (dilution 1:1000, Cat# 9196, Cell Signaling Technology, Danvers, MA, USA), anti-Hif-1α (dilution 1:1000, Cat# NB 100–449, Novus Biologicals, Centennial, CO, USA), anti-Jak1 (dilution 1:1000, Cat# 3332, Cell Signaling Technology, Danvers, MA, USA), anti-phospho-Jak1 (Y1034/1035) (dilution 1:1000, Cat# 3331, Cell Signaling Technology, Danvers, MA, USA), anti-PGC1α (dilution 1:1000, Cat# 2178, Cell Signaling Technology, Danvers, MA, USA), anti-PKM2 (dilution 1:1000, Cat# D7844, Cell Signaling Technology, Danvers, MA, USA), anti-cPLA2 (dilution 1:1000, Cat# 2832, Cell Signaling Technology, Danvers, MA, USA), anti-phospho-cPLA2 (5505) (dilution 1:1000, Cat# 2831, Cell Signaling Technology, Danvers, MA, USA), anti-STAT3 (dilution 1:1000, Cat# 9139, Cell Signaling Technology, Danvers, MA, USA) and anti-phospho-STAT3 (Y705) (dilution 1:1000, Cat# 9131, Cell Signaling Technology, Danvers, MA, USA). After incubation with the primary antibodies, membranes were washed and treated for 1 h with the following secondary antibody according to the source of primary antibody: Peroxidase-AffiniPure Goat Anti-IgG (dilution 1:1000, Cat# 115-035-146, Jackson ImmunoResearch Labs, West Grove, PA, USA) and Peroxidase-AffiniPure Goat Anti-rabbit IgG (dilution 1:10000, Cat# 115-035-144, Jackson ImmunoResearch Labs, West Grove, PA, USA). After this incubation, membranes were washed and developed by using Amersham ECL Western Blotting Reagent (Cat# RPN2124, GE Healthcare Bio-Sciences, Pittsburgh, PA, USA). Staining was evaluated by using C-DiGit Blot Scanner (LI-COR, Lincoln, NE, USA) and quantifications of the blots were performed by using the software Image J (http://imagej.nih.gov/ij, NIH, USA).

Accession to public repository of mRNA library of human breast cancer
Data were collected by using the Gene Expression Browser (GXB) via Sidra Medical and Research Center Gateway, by assessing the Nag`s Reconstituted Public Data Set, which includes the data from 1839 breast cancer patients. mRNA array data were obtained by Affymetrix.

Data analyses and statistics
All graphics and statistical analyses were performed with software Prism 7 for Mac (GraphPad Software Inc., La Jolla, CA, USA). Student’s t test, one-way ANOVA followed by Tukey’s post test or two-way ANOVA followed by Sidak’s post test were used as appropriate.
RESULTS
Serotonin confers proliferative advantages to MCF-7 cells affecting their metabolic profile. When MCF-7 cells were grown in the presence of 10 µM 5-HT, we observed a 28% increase in the number of cells after 18 h of incubation as compared with control (Fig. 1a). This treatment decreased the permeability of cancer cells to 7-AAD (Fig. 1b) and reduced the staining for Annexin V (Fig. 1c). These effects are prevented by the presence of 0.1 µM ketanserin, an antagonist of the 5-HT2 receptor family. To identify 5-HT receptors expressed...
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Fig. 2 Serotonin modulates MCF-7 cells mitochondria biology. a Relative RNA expression of pyruvate dehydrogenase kinase 1 (PDK1), (b) enzymatic activity of succinate dehydrogenase (b), (c) relative RNA expression of succinate dehydrogenase A (SDHA) and (d) relative RNA expression of SIRT3. Plotted values are mean ± S.E.M. of seven independent experiments (n = 7; for a, c and d, * means P < 0.05 as compared with control in the absence of 5-HT, Student’s t test); for c * means P < 0.05 as compared with control in the absence of ketanserin and the bracket indicates P < 0.05 as compared with the absence of 5-HT; two-way ANOVA, followed by Sidak’s post test). e-m Representative confocal images of MCF-7 cells treated in the absence (e-g) or the presence of 5-HT (h-j) and 5-HT plus ketanserin (k-m), stained with MitoTracker green FM (e, h, k) and MitoTracker Red CMXRos staining. o Quantification of MitoTracker Red CMXRos staining. p Quantification of overlapping. Plotted values in panels N–P are mean ± S.E.M. of three independent experiments (n = 3; * means P < 0.05 compared with control; one-way ANOVA followed by Tukey’s post test).

by MCF-7 cells, we performed a qPCR for all the 5-HT receptor subtypes and found the expression of only 5-HT₁R₁, 5-HT₁R₅, 5-HT₂R₇ and 5-HT₇R (Fig. 1d). The 5-HT₆R₁D receptor presents a low affinity for ketanserin (Kᵢ = 15 µM), while 5-HT₂ is not antagonised by the drug. Therefore, these two receptors are not playing a role on the observed effects of 5-HT on MCF-7 proliferation. That being said, we performed the experiments that follow considering the effects of 5-HT observed as a consequence of activation of 5-HT₆R₁D receptors only.

In addition to the proliferative and anti-apoptotic effects reported above, 5-HT promoted a dose-dependent increase in glucose consumption by MCF-7 cells (Fig. 1e). This effect was accompanied by an increase in lactate production (Fig. 1f), suggesting that the hormone is activating glucose metabolism. However, the ratio of lactate production/glucose uptake decreased as a function of the hormone concentration (Fig. 1g), indicating that the fate of increased glucose metabolism is changing as a consequence of 5-HT action. In this context, our results revealed activation of MCF-7 mitochondrial metabolism by 5-HT (Fig. 1h), which suggests that mitochondrial oxidation might be the fate of increased glucose uptake by these cells. All these effects are prevented when the 5-HT₆R₁D receptor antagonist, ketanserin, is present in the assays, suggesting that these receptors are triggering the response of MCF-7 cells to 5-HT (Fig. 1a-h).

Serotonin modulates MCF-7 cells metabolic enzyme activity and expression

Next, we sought to identify the mechanism by which 5-HT is shifting MCF-7 metabolism towards oxidation. For this, we evaluated the activity of the major regulatory glycolytic enzymes, namely hexokinase (HK), phosphofructokinase (PFK) and pyruvate kinase (PK), within MCF-7 cells treated with 10 µM 5-HT. Although the three enzymes were stimulated by the hormone, the effects on HK and PFK were modest compared with the effects of 5-HT on PK activity that increased 100% upon 5-HT stimulation as shown in Supplementary Fig. S1A and S1B, the different isoforms of these two enzymes were downregulated upon 5-HT treatment. The expression of HK1 decreased approximately by 40% upon 5-HT treatment (Supplementary Fig. S1E), while the decrease in HK2 expression reached almost 70% (Supplementary Fig. S1F). This occurs in contrast to the 10% activation of this enzyme observed under the same treatment (Supplementary Fig. S1A). Similarly, the expression of the three PFK isoforms, PFKM (Supplementary Fig. S1G), PFKL (Supplementary Fig. S1H) and PFKP (Supplementary Fig. S1I) was reduced by around 10–20% upon treatment with 5-HT, despite the 10% activation of the enzyme (Supplementary Fig. S1B). The expression of the different PK isoforms upon 5-HT treatment followed a unique pattern. While there was no effect on PKL expression (Supplementary Fig. S1I), PKM1 expression decreased by 20% (Supplementary Fig. S1K) and PKM2 expression increased by 30% (Fig. 1j). This increase in PKM2 mRNA levels was also observed in the protein levels through western blot analysis (Fig. 1k, l). PKM2 represents 60% of the total PK expressed within these cells with no treatment (Supplementary Fig. S1L), which means that after 5-HT stimulation, this isoform represents nearly 80% of total PK activity within the cells (Supplementary Fig. S1M) and there is a net increase of ~15% in total PK expression. For PK activity, we observed a 2.1-fold increase upon 5-HT stimulation (Fig. 1i) and this cannot be explained solely by the increase in PK expression. In this context, it is well described that PKM2 isoform is the multiregulated isoform that can be activated by several signals, including ERK1/2-mediated phosphorylation of the enzyme. In addition, both 5-HT₂A and 5-HT₇C receptors activate ERK1/2 by MEK-mediated phosphorylation. Indeed, we showed that 5-HT promotes ERK1/2 phosphorylation (Fig. 1m, o), which is confirmed by evaluating the phosphorylation of ERK1/2 substrate, the cytosolic phospholipase A₂ (cPLA₂) on SS05 (Fig. 1n, p). Moreover, 5-HT-stimulated phosphorylation of both ERK1/2 and cPLA₂ is prevented by ketanserin (Fig. 1m–p), and the phosphorylation of cPLA₂ is prevented by LY3214995 (Fig. 1n, p).

In addition, the treatment of MCF-7 cells with LY3214996, an inhibitor of ERK1/2, partially prevented PK activation by 5-HT (Fig. 1q). To this end, we concluded that 5-HT-induced activation of PK might be due to an increased expression of the enzyme, especially PKM2 rather than the other isoforms, which makes the enzyme more responsive to ERK1/2 signalling. Yet, the only partial reversal of 5-HT-induced PK activation by ERK1/2 inhibition indicates that both increased expression of the enzyme and its activation by ERK1/2 are independently contributing to the increased PK activity induced by 5-HT on MCF-7 cells.

Serotonin triggers mitochondrial biogenesis in MCF-7 cells

The role of PKM2 in cancer has been scrutinised and reviewed by many authors, and it is clear that the expression of this PK isoform promotes cell growth and proliferation. Among the advantages of expression of PKM2 is the favouring of the Warburg effect over oxidation through promoting the expression of HIF-1α. Indeed, 5-HT promotes the expression of HIF-1α, as will be further shown and discussed. HIF-1α promotes the expression of pyruvate...
dehydrogenase kinase 1 (PDK1) that inhibits the pyruvate dehydrogenase complex (PDH). Therefore, we evaluated PDK1 expression in 5-HT-treated MCF-7 cells and found a 60% increase in the levels of PDK1 mRNA (Fig. 2a). Nevertheless, we observed here that 5-HT increases glucose consumption and mitochondria activity at the same time, suggesting that the fate of glucose is the oxidation. Indeed, 5-HT promotes a twofold increase in the levels of PDK1 mRNA (Fig. 2a). Nevertheless, we observed here that 5-HT increases glucose consumption and mitochondria activity at the same time, suggesting that the fate of glucose is the oxidation. Indeed, 5-HT promotes a twofold activation of succinate dehydrogenase (SDH) activity, which is also prevented by ketanserin (Fig. 2b). This explains our aforementioned observation of the increased mitochondrial metabolism upon 5-HT stimulation of MCF-7 cells (Fig. 1h).

However, even though 5-HT activates SDH, inhibition of PDH by ketanserin (0.1 μM) also prevented by ketanserin (Fig. 2b). This explains our aforementioned observation of the increased mitochondrial metabolism upon 5-HT stimulation of MCF-7 cells (Fig. 1h).
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PDK1 would avoid pyruvate oxidation in mitochondria. To investigate this, we first evaluated the SDHA expression and found that 5-HT treatment increased the mRNA levels of this enzyme by -1.5-fold (Fig. 2c), partially explaining the enzyme activation. This result also suggests that 5-HT may be promoting mitochondrial biogenesis, and hence the increased expression of PDK1 and SDHA may be due to this phenomenon. To test this possibility, we evaluated the levels of SirT3 (SIRT3), known to promote mitochondrial biogenesis. Indeed, SIRT3 mRNA levels significantly increased upon the treatment of MCF-7 cells with 5-HT (Fig. 2d). Moreover, upon staining of 5-HT-treated MCF-7 cells with MitoTracker Green (that stains mitochondria regardless of the oxidation state) and MitoTracker deep red (that stains metabolically active mitochondria), we observed an increase in total and active mitochondria in 5-HT-treated MCF-7 cells as compared with the control, and this effect was prevented by ketanserin (Fig. 2e–p). Therefore, our results strongly suggest that the increased oxidation of pyruvate by 5-HT-treated MCF-7 cells is due to the augmented number and activity of mitochondria in these cells.

Serotonin effects involve two distinct signalling pathways. So far, we observed that the stimulation of MCF-7 cells with serotonin activates glycolysis and shifts the fate of pyruvate from reduction to oxidation. Glycolysis activation is primarily due to the preferential expression of PKM2, which greatly activates this enzyme. Indeed, the Warburg effect has been always associated with reduced PKM2 activity, and thus it is reasonable to postulate that increased expression and activity of this enzyme promotes an augmented oxidation of pyruvate. Moreover, the shift of the fate of pyruvate might be also due to the increased mitochondrial biogenesis observed, which is related to the higher activity of the organelle within these cells. To test how 5-HT signalling is transduced into these effects, we tested a series of inhibitors of known 5-HT signalling pathways on the activities of PK and SDH, as indicators of increased glycolysis and mitochondrial biogenesis, respectively. Our results are shown in Fig. 3a, b for PK and SDH activities, respectively, and revealed different signalling pathways involved in each mechanism. Activation of PK by 5-HT was prevented by the presence of LY3009104 and WP1066, inhibitors of Jak1 and STAT3, respectively (Fig. 3a), whereas the increased SDH activity in 5-HT-treated cells was prevented by the presence of 2′,3′-dideoxyadenosine (DDA) and H89, inhibitors of adenyl cyclase and protein kinase A (PKA), respectively (Fig. 3b). Protein kinase C (PKC) and phospholipase C (PLC), two known transducers of 5-HT signal through 5-HT2A/2C receptors are not involved in these effects of the hormone, since their inhibition by using U73122 (PLC) and BIS I (PKC) has not affected activation of PK or SDH by 5-HT (Fig. 3a, b).

To further investigate the signalling pathways linking 5-HT stimulation and the activation of Jak1/STAT3 and cAMP/PKA pathways, we evaluated the effects of 5-HT treatment on the phosphorylation of Jak1 in MCF-7 cells. We found that 5-HT promoted the phosphorylation of Jak1 on Y1034/1035 (Fig. 3c, d), a known activation site of this kinase. Moreover, this phosphorylation was prevented by the presence of 5-HT2A/2C receptor antagonist, ketanserin (Fig. 3c, d). Indeed, Jak1 activation by 5-HT2A/2C receptor stimulation has been reported elsewhere, and this kinase has been shown previously to be associated with STAT3 phosphorylation in mammary cancer cells. We confirmed this association in the current work since we also observed an increased STAT3 phosphorylation on MCF-7 cells treated with 5-HT (Fig. 3e, f). In the context of glycolytic activation, it has been demonstrated that STAT3 regulates expression of Hif-1a in cancer cells and this impacts Akt phosphorylation. Therefore, we investigated whether these events are affected by 5-HT on MCF-7 cells. Our results revealed that upon 5-HT treatment, Hif-1a expression increased in MCF-7 cells, and this effect was prevented by the presence of STAT3 inhibitor, WP1066 (Supplementary Fig. S2A, S2B). Similar effects were observed for Akt phosphorylation on T308 (Supplementary Fig. S2C, S2D), revealing that STAT3 activation by Jak1 upon 5-HT treatment of MCF-7 cells increases both the expression of Hif-1a and phosphorylation of Akt. These two events are directly involved in activation of glycolysis and may partially explain the increased glucose consumption by 5-HT-treated MCF-7 cells. Hif-1a, for instance, is directly involved in the expression of PDK1, by explaining the results presented in Fig. 2a. Moreover, STAT3-activated Akt phosphorylation triggers the expression of PKM2 in cancer cells, which also explains our findings. Nonetheless, these results do not explain the decrease in lactate production/glucose consumption ratio or the increased mitochondrial biogenesis. However, it has been reported that STAT3-activated Akt triggers the expression of lactate dehydrogenase (LDH) B, an isoform of lactate dehydrogenase responsible for the mitochondrial oxidation of lactate into pyruvate. Therefore, we evaluated the expression of the two LDH isoforms, A and B, as well as the plasma membrane lactate transporter, MCT1. We found a slight reduction in LDHA expression (Supplementary Fig. S2E) and a substantial increase in LDHB expression (Fig. 3g). Besides, MCT1 expression was reduced upon 5-HT treatment (Fig. 3h), suggesting a difficult externalisation of lactate by the cells. These effects on LDHA, LDHB and MCT1 were prevented by ketanserin and Jak1 inhibitor LY3009104, supporting the involvement of 5-HT2A/2C and Jak1/STAT3 on this process (Fig. 3g, h; Supplementary Fig. S2E). Moreover, PKA inhibitor H89 did not alter the effects of 5-HT on LDHA, LDHB and MCT1 mRNA levels (Fig. 3g, h; Supplementary Fig. S2E). Cytosolic formed lactate might be taken up by mitochondria where it is oxidised by the increased LDHB within this organelle. This last effect is related to the increased mitochondria biogenesis observed in the current work. To assess this effect, we studied the effects of 5-HT on the phosphorylation of CREB on S133, a site for PKA action on this transcription factor. We observed an increased phosphorylation of CREB on S133 upon 5-HT treatment of MCF-7 cells, an effect fully prevented by ketanserin (Fig. 3i, j). The phosphorylation of CREB on S133 can be promoted by PKA and ERK1/2, which we have demonstrated to be activated by 5-HT (Fig. 1m–o). However, ERK1/2 inhibitor LY221976 also partially prevented the phosphorylation of CREB on S133 promoted by 5-HT (Fig. 3i, j). PKA inhibitor H89 also partially prevented 5-HT-induced CREB phosphorylation.
phosphorylation, confirming that 5-HT is promoting PKA activation in MCF-7 cells. Since PKA is reported to promote the expression of PGC1α, a nodal regulator of mitochondrial biogenesis, via CREB phosphorylation, we investigated the expression of this coregulator. Indeed, 5-HT treatment increased the expression of PGC1α in MCF-7 cells (Fig. 3k, l). This effect was not significantly affected by the ERK1/2 inhibitor, LY324996, but was prevented by both H89 and ketanserin, a PKA inhibitor and a 5-HTR2A/C antagonist, respectively (Fig. 3k, l). Altogether, these results strongly suggest that the effects of 5-HT promoting glucose consumption and mitochondrial biogenesis occur through two distinct signalling pathways, involving a Jak1/STAT3-induced glycolysis and cAMP/PKA-induced mitochondrial biogenesis. It is important to notice that U73122 and BIS I,
inhibitors of phospholipase C and protein kinase C, respectively, do not interfere with the effects of 5-HT on neither PK nor SDH activities, supporting the notion that these signal transducers are not involved in the effects reported in the current work.

JAK1/STAT3 activates glycolysis while cAMP/PKA triggers mitochondrial biogenesis

Aiming to prove that Jak1/STAT3 and cAMP/PKA pathways are distinctly involved in 5-HT-induced MCF-7 glucose metabolism, we treated the cells with the hormone in the absence and the presence of the Jak1 inhibitor, LY300914, and the adenylate cyclase inhibitor, DDA, and evaluated glucose consumption, lactate production and mitochondrial activity. The results showed that the effects of 5-HT on glucose consumption (Fig. 3m), lactate production (Fig. 3n) and lactate production/glucose consumption ratio (Fig. 3o) were completely hindered by LY300914, similar to ketanserin. DDA did not affect the stimulation of glucose consumption promoted by 5-HT on MCF-7 cells (Fig. 3m). However, the effects of 5-HT on lactate production by MCF-7 cells were enhanced in the presence of DDA (Fig. 3n). Moreover, both the Jak1 and the adenylate cyclase inhibitors reversed the effects of 5-HT on the ratio of lactate production and glucose consumption (Fig. 3o). In addition, by evaluating the effects of these inhibitors on 5-HT-stimulated mitochondrial activity of MCF-7 cells, we observed that LY300914 did not alter the effects of 5-HT, while DDA prevented these effects (Fig. 3p). These results corroborate that the activation of glycolysis by 5-HT on MCF-7 cells is mediated by Jak1/STAT3 signalling, while the augmented mitochondrial activity is mediated by cAMP/PKA. The fact that DDA enhances the effects of 5-HT on lactate production and reverses the effects of the hormone on the ratio between lactate production and glucose consumption confirms that the increased mitochondrial activity is responsible for the shift in the fate of pyruvate from fermentative to oxidative metabolism.

We have previously observed that STAT3 is involved in the 5-HT activation of glycolysis in skeletal muscle. Significance of the results for breast cancer

Significance of the results for breast cancer

Aimed to evaluate whether the results obtained are relevant to breast cancer biology, we tested the effects of 5-HT on the non-tumorigenic human breast cell line MCF10A and on the breast cancer metastatic cell line MDA-mb-231. It was previously reported that MCF-7, MCF10A and MDA-mb-231 have different patterns of 5-HT receptor expression. Here, we confirmed these results showing that MCF10A primarily expresses 5-HT<sub>2</sub> receptors, with very low levels of 5-HT<sub>2A</sub> and 5-HT<sub>2C</sub> (Fig. 4a), and MDA-mb-231 majorly expresses 5-HT<sub>2C</sub> with intermediate levels of 5-HT<sub>R</sub> (Fig. 4b). These data contrast with the pattern of 5-HT receptors expressed by MCF-7 cells, which primarily express 5-HT<sub>2C</sub> by presenting residual expression levels of the 5-HT<sub>R</sub> (Fig. 1d). The expression patterns of 5-HT<sub>2C</sub> and 5-HT<sub>R</sub> on the three cell lines used here were confirmed by measuring the proteins levels by western blot (Fig. 4c–e). This difference in 5-HT receptor expression might be responsible for the antagonistic effects of 5-HT on MCF10A, as compared with the two tumour cell lines. For instance, 5-HT decreased the viability of MCF10A cells (Fig. 4f) but increased the viability of MDA-mb-231 (Fig. 4g). Indeed, the effects of 5-HT on MCF10A were prevented by 50 nM spiperone, a 5-HT<sub>R</sub> antagonist, but not by ketanserin (Fig. 4f). On the other hand, spiperone was ineffective in preventing 5-HT effects on MDA-mb-231, while these effects were fully prevented by ketanserin (Fig. 4g). Similarly, 5-HT induced apoptosis on MCF10A cells (Fig. 4h), while it prevented this phenomenon on MDA-mb-231 (Fig. 4i), where the effects on the former and the latter cell lines were prevented by spiperone and ketanserin, respectively (Fig. 4h, i for MCF10A and MDA-mb-231, respectively). In addition, the effects of 5-HT on the expression of PKM2 of MCF-7 cells are similar in MDA-mb-231 cells (Fig. 4j). The treatment of these cells with 5-HT increased the expression of PKM2 (Fig. 4j), k, which we have correlated to an increased glycolytic flux within these cells. This effect was prevented by ketanserin, but not by spiperone (Fig. 4j, k). On the other hand, 5-HT did not affect the expression levels of PKM2 in MCF10A cells (Fig. 4j, k). Similarly, 5-HT did not affect neither the number nor the polarisation of mitochondria in MCF10A cells, as evaluated by MitoTracker Green (Fig. 4l) and MitoTracker Red (Fig. 4m), respectively. However, the effects of 5-HT on MDA-mb-231 were similar to those shown for MCF-7 (Fig. 2), increasing both the number (Fig. 4n) and the polarisation of mitochondria (Fig. 4o). These effects were prevented by ketanserin but not by spiperone, supporting the notion that they are mediated by 5-HT<sub>2A/C</sub> receptors.

Aimed to investigate the expression pattern of 5-HT<sub>2C</sub> and 5-HT<sub>R</sub> on different human breast cancers, we assessed the Gene Expression Browser (GXB), via Sidra Medical and Research Center Gateway, by assessing the Nagalla Reconstituted Public Data Set, which includes the data from 1839 breast cancer patients. Breast tumours can be classified according to the intrinsic molecular subtypes (IMS), which takes into account the expression patterns of important receptors related to breast cancer progression and physiology, i.e., the oestrogen receptor (ER), progesterone receptor (PR) and human epidermal growth factor receptor-type

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was not different between these two groups (Fig. 5c). The exact did not vary among breast cancers classifi

\[ \text{Expression level} = \text{LumA} < \text{LumB} < \text{HER2}^+ \] 

worsens, i.e., the expression levels are such as LumA < LumB < HER2 + and Basal (Fig. 5a). On the other hand, 5-HTR \(_2^C\) expression did not vary among breast cancers classified according to the IMS (Fig. 5a). Similarly, dividing the breast cancer between those whose patients showed cancer cells within their lymph nodes (LN +) or not (LN −), we observed a higher expression of 5-HTR \(_2^C\) in LN +, as compared with LN − (Fig. 5b), while 5-HTR \(_2^C\) expression was not different between these two groups (Fig. 5c).

The same result was obtained separating the breast cancers between those positive and negative for p53 (p53 + and p53 −, respectively), where 5-HTR \(_2^C\) was upregulated in p53 +, as compared with p53 − cancer (Fig. 5d), while 5-HTR \(_2^C\) remained unaltered between the two groups (Fig. 5e). Evaluation of lymph nodes and p53 expression is extremely important for breast cancer patients since these characteristics are directly associated with the cancer aggressiveness, the prognosis, the recurrence rate and survival.\(^{60,62}\) Our data here strongly indicate that the higher the 5-HTR \(_2^C\) expression, the worse is the prognosis for the breast cancer patients. Since we characterised that 5-HT signalling via this HTR receptor alters breast cancer cell metabolism towards highly glycolytic and highly oxidative, through upregulation of PKM2 and mitochondria biogenesis, we propose that these characteristics are directly associated with breast cancer aggressiveness and poor prognosis.

**DISCUSSION**

One of the hallmarks of cancer cells is a deregulated energy metabolism including the so-called aerobic glycolysis, or Warburg effect, where most of the consumed glucose is converted into lactate regardless of the oxygen supply to these cells.\(^{63}\) In fact, aerobic glycolysis is required for fast-growing cells due to the rapid supply of ATP and the diversion of glycolysis intermediates into various biosynthetic pathways.\(^{63}\) However, there are growing evidences exemplifying that aerobic glycolysis is not constitutively activated in cancer cells that constantly shift to oxidative metabolism, which contribute to cancer progression and metastasis.\(^{64}\) This shift involves the mitochondrial oxidation of pyruvate, decreasing the ratio of lactate formation.\(^{64}\) In the current paper, we show that serotonin confers proliferative advantages to breast cancer cells, which involves not only stimulation of glucose metabolism but also a shift from fermentative to oxidative metabolism. This effect is observed only in the cancer cells and not in the non-tumorigenic breast cell line, MCF10A.

Our data show that serotonin is signalling a proliferative advantage to breast cancer cells, by increasing the rate of cell proliferation and decreasing programmed cell death. Similar results have been published earlier by Sonier et al., where authors reported that 5-HT-treated MCF-7 cells proliferate faster than control, an effect that was attributed to the expression of 5-HT \(_{2A}\) by these cells.\(^{65}\) Here, through q-RT-PCR, we screened MCF-7 cells for the expression of several 5-HT receptors and found expression of only 5-HT \(_{1E}\), 5-HT \(_{1A}\), 5-HT \(_{2C}\) and 5-HT \(_{7}\) (Fig. 1d). Among these four 5-HT receptors, 5-HT \(_{2C}\) demonstrated the highest expression by MCF-7 cells (at least ten times more expressed than the other three receptors, Fig. 1d). Similar results were reported by Pai et al. who also showed that MCF-7 cells express 5-HT \(_{1E}\), 5-HT \(_{1A}\), and 5-HT \(_{7}\) but these authors used end-point PCR, which

\[ \text{Expression value} = \text{p53} - \text{p53} + \] 

Fig. 5 Correlation of the expression patterns of 5-HTR \(_{2C}\) and 5-HTR \(_{7}\) and prognosis markers in breast cancers. Data were collected by using the Gene Expression Browser (GXB) via Sidra Medical and Research Center Gateway, by assessing the Nagalla Reconstituted Public Data Set,\(^{27}\) which includes the data from 1839 breast cancer patients. mRNA array data were obtained by Affymetrix.

**a** Correlation of 5-HTR \(_{2C}\) and 5-HTR \(_{7}\) expression and the IMS classification of the breast cancer (\(n = 398\); different letters superscript means \(P < 0.05\) as compared with the other values; two-way ANOVA, followed by Sidak’s post test). **b**, **c** Comparison of 5-HTR \(_{2C}\) and 5-HTR \(_{7}\) expression in lymph-node-negative (LN −) or lymph-node-positive (LN +) primary breast cancer (\(n = 532\); * means \(P < 0.05\); Student’s t test). **d**, **e** Comparison of 5-HTR \(_{2C}\) and 5-HTR \(_{7}\) expression in p53-negative (p53 −) or p53-positive (p53 +) primary breast cancer (\(n = 275\); * means \(P < 0.05\); Student’s t test).
might have detected residual mRNA levels. Moreover, the proliferative effects of 5-HT on MCF-7 cells were hindered in the presence 0.1 µM ketanserin, which is at least ten times the Ki for 5-HTR2A and 5-HTR2C, by corroborating the results published by Sonier et al. In their paper, Sonier et al. have attributed the ketanserin effects to the antagonism of the drug on 5-HTR2A subtype. However, since ketanserin is also a 5-HTR2C antagonist and due to the predominance of 5-HTR2C over 5-HTR2A, we believed that the former is the major receptor responsible for 5-HT actions on MCF-7 cells. Nevertheless, we cannot discard a possible transduction by 5-HTR2A as proven by Sonier et al., since both are present in these cells (Fig. 1d) and represent similar affinities and responsiveness to 5-HT and ketanserin.

Altogether, our data indicate opposite effects of 5-HT on cancer and non-cancer cells. While conferring proliferative advantages to both tumoural cells (MCF-7 and MDA-mb-231), 5-HT decreased the non-cancer MCF10A proliferation and promoted apoptosis of this cell line. Indeed, these results are in accordance with previous work demonstrating the role of 5-HT on the evolution and regression of mammary cells, especially during lactation. The effects of 5-HT reported on the mammary gland also include modulations of glucose transport and metabolism, supporting our finding here. Accordingly, the differential effects of 5-HT on the mammary gland evolution might be due to the pattern of 5-HT receptor expression, whereas the 5-HT2A plays a role in the involution of the mammary gland possibly due to promoting cell apoptosis. Conversely, our data indicated that 5-HTR3A/C is related to promoting cells proliferation and inhibiting the apoptosis process, by conferring proliferative advantages to these cells. These data are strongly supported by another study where authors have proposed the proliferative action of 5-HT on breast cancer cells via 5-HTR2A. Pai et al. also reported that the pattern of 5-HT receptor expression varies with the type of breast tumours. Indeed, these authors also found expression of other 5-HT receptor subtypes among the same cell lines that we used here; however, their analyses were performed by end-point PCR, which detects also negligible levels of mRNA.

Conclusively, we propose that 5-HT, through 5-HT2A/2C receptors, phosphorylates Jak1, which in turn phosphorylates STAT3, promoting ERK1/2 activation, Akt phosphorylation and Hif-1α expression. Together, these effects promote glucose consumption by the preferential expression of PKM2. Simultaneously, 5-HT2A/2C receptor activation by 5-HT triggers adenylate cyclase, by activating PKA that promotes the expression of PGC1α. Finally, this coactivator induces mitochondrial biogenesis, thus shifting the fate of pyruvate formed due to increased glycolysis towards oxidation via mitochondrial metabolism (Fig. 6). Consequently, the breast cancer cells become more proliferative and less susceptible to apoptosis, being able to form more aggressive cancers.

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AUTHOR CONTRIBUTIONS
M.S.-P. and P.Z. designed research; M.S.-P., L.P.P., J.R.B., A.C.O., J.M.A., D.M.M., W.S.C., M.C.M. and P.Z. performed research; C.P.F. and D.B.-d.-S. contributed new analytical tools; M.S.-P. and P.Z. analysed the data and wrote the paper.

ADDITIONAL INFORMATION
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