Structure-Activity Relationship Study Identifies A Novel Lipophilic Amiloride Derivative That Induces Lysosome-Dependent Cell Death in Therapy-Resistant Breast Cancer Cells

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Abstract

Amiloride and its derivatives have long attracted attention as potential anticancer therapeutic agents. Several early studies characterized amilorides as inhibitors of sodium-proton antiporter-dependent tumor growth and urokinase plasminogen activator-mediated metastasis. However, more recent observations indicate that amiloride derivatives are specifically cytotoxic toward tumor cells relative to normal cells and have the capacity to target tumor cell populations resistant to currently-employed therapies. A major barrier to clinical translation of the amilorides is their modest cytotoxic potency, with EC_{50} values in the high micromolar to low millimolar range. Here we report structure-activity relationship observations that underscore the importance of the guanidinium group and the presence of lipophilic substituents at the C(5) position of the amiloride pharmacophore in promoting cytotoxicity. Moreover, we demonstrate that our most potent derivative called LLC1 is specifically cytotoxic toward mouse mammary tumor organoids and drug-resistant populations of various breast cancer cell lines, and induces lysosomal membrane permeabilization as a prelude to lysosome-dependent cell death. Our observations offer a roadmap for the future development of amiloride-based cationic amphiphilic drugs that engage the lysosome to specifically kill breast tumor cells.

1   Introduction

Amiloride is a commonly employed oral antikaliuretic and diuretic used in the treatment of hypertension and congestive heart failure [1]. Structurally, amiloride is a substituted (pyrazinecarbonyl) guanidine; its relatively high pKa of 8.7 ensures its protonation and function at physiological pH [2-4]. Its renal mechanism of action involves inhibition of both the Na^+/H^+ exchange

system of the proximal tubule and the epithelial Na^+ channel of the distal tubule to increase sodium

excretion without affecting potassium excretion [3, 4]. In addition to its uses in hypertension and

congestive heart failure, amiloride and its numerous derivatives have been studied for a variety of

indications, from multiple sclerosis and cystic fibrosis to cancer and chronic pain [5-9].

While an array of studies since the 1980s have highlighted Na^+/H^+ exchange and urokinase

plasminogen activator (uPA) inhibition mechanisms in amiloride suppression of tumor aggressiveness

and metastasis [10], more recent observations indicate that amiloride and its derivatives also exhibit

tumor cell-selective cytotoxicity via a non-apoptotic mechanism. For example, our studies with the

derivative 5-(N,N-hexamethylene)amiloride (HMA) underscore the notion that the properties of

amiloride derivatives may be ideally suited for targeting particularly aggressive or therapy-refractory

tumors [11, 12]. HMA efficiently kills bulk breast tumor cells independent of subtype, species of

origin, or proliferation state, but does not efficiently kill non-transformed cells derived from a variety

of tissues at the same concentrations. Indeed, cell lines derived from a variety of tumor types are

similarly susceptible to HMA, suggesting that its mechanism of cytotoxic action may be dependent on

cellular transformation rather than patient-specific genetic alterations. Moreover, HMA is cytotoxic

toward breast cancer cell populations that are resistant to currently-employed therapies, including the

highly refractory cancer stem cell population. Finally, HMA induces morphological changes within

the lysosomes of tumor cells, and cytotoxicity is rescued by an inhibitor of the lysosomal protease

cathepsin [11, 12], pointing a central role for the lysosome in its mechanism of action.

While previous *in vitro* studies have revealed many attractive properties of amiloride derivatives, and

suggest that derivatization of the C(5) position can markedly enhance the tumor-specific cytotoxic

properties of the drug, further translation of HMA is hampered by its relatively modest potency and

poor pharmacokinetic properties in mice [13]. In this study we aim to glean further insight into

structure-activity relationships within the amiloride pharmacophore by comparing the cytotoxic

potential of derivatives modified at the C(2) and C(5) positions. Moreover, we aim to identify more

potent derivatives that optimize tumor cell-specific cytotoxicity and preserve the ability to target

therapy-resistant populations. Finally, we aim to confirm the role of the lysosome in amiloride

derivative mechanism of action by characterizing drug impact on lysosomal membrane

permeabilization.

2   Materials and methods

2.1   Cell Culture and Drug Treatments

MDA-MB-231, MCF7, SKBR3, T47D, and 4T1 cells were purchased from American Type Culture

Collection (ATCC, Manassas, VA, USA) and maintained at 37°C and 10% CO_2 in media supplemented

with 10% fetal bovine serum (FBS, Genesee Scientific) and antibiotics (penicillin/streptomycin; Gibco

- Thermo Fisher), as recommended by ATCC. MCF7-B7-TS (MCF7 TS), MCF7-G11-TR-1 (MCF7

TR-1), MCF7-G11-TR-5 (MCF7 TR-5), and MCF7 MX-100 cells were obtained from the Physical

Sciences-Oncology Network (PS-ON) Bioresource Core Facility (PBCF) at ATCC or gifted from Dr.

A.M. Yu, UC Davis. The tamoxifen (TAM) resistant lines were maintained in MCF7 base media

supplemented with 1 μM or 5 μM TAM [14]. The mitoxantrone resistant line (MX-100) was

maintained as described [15]. Cell line attributes are summarized in Supplementary Table S1.

Amiloride, HMA, benzamil, and 10357 were sourced as described in Supplementary Table S2, and

applied to cells to cells at 50-70% confluency. Amiloride derivatives were added 24hr before analysis

of cell viability, while chemotherapeutic agents (docetaxel, doxorubicin, and mitoxantrone) were

added at specified timepoints to develop short term resistance in parental cell lines. Vehicle was added

as a negative control for each experiment.

2.2 Design and Synthesis of Novel Amiloride Derivatives

The new amiloride derivatives were designed to introduce novel hydrophobic groups at the C(5) position to improve cell permeability and potential *in vivo* availability. Synthesis of LLC1 was carried out as shown in Supplementary Fig. S1. To a suspension of methyl 3-amino-5,6-dichloro-2-pyrazinecarboxylate (compound 1; 444 mg, 2.0 mmol) in anhydrous DMF (3 mL) was added 4-(4-fluorophenyl)piperidine (compound 2; 358.5 mg, 2.0 mmol) and N,N-*Diisopropylethylamine (*DIPEA, 697 µL, 4.0 mmol), the reaction mixture was stirred at room temperature overnight and then added to cold water. The precipitate was collected by centrifuge and washed with water. The crude product (compound 3) was used for the next step without further purification. 0.5 M NaOCH$_3$ solution in CH$_3$OH (20 mL) was added to guanidine hydrochloride (955.3 mg, 10 mmol) and the resulting mixture was stirred at room temperature for 0.5 hr. The white precipitate was removed by filtration, and the free guanidine solution in methanol was added to a suspension of methyl ester (compound 3; 364.8mg, 1.0 mmol) in DMF (5 mL) and stirred at room temperature overnight. Brine (20 mL) was added, and the mixture extracted with ethyl acetate (3 × 30 mL). The organic layer was washed with 10% NaCl (2 × 30 mL), dried over anhydrous Na$_2$SO$_4$, and concentrated under reduced pressure. The crude mixture was dissolved in 5 mL of 30% CH$_3$CN/70% H$_2$O and 0.1% trifluoroacetic acid and purified by a reverse phase HPLC. The collected eluent was lyophilized to yield the designed product LLC1. The structure was verified by Orbitrap HR-ESI MS: 392.1391 [M+1] (Supplementary Fig. S2A). The purity was >98% as shown in HPLC (Supplementary Fig. S2B).

2.3 MTT Assay

Cells were seeded on 24-well plates and cultured to 50-70% confluency prior to treatment with various compounds for 24-72hr depending on the assay context. Following treatments, media was aspirated from cells and replaced with 5 mg/mL 3-(4,5-Dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide (MTT, #M5655, Sigma Aldrich) solution in base media. Cells were incubated at 37°C for 1-

2hr to induce formazan crystal formation, then washed with PBS to remove impurities. Crystals were

dissolved using isopropyl alcohol (IPA, 0.5% 1N HCl in isopropanol), and sample absorbance (λ_{ex} 570

nm, λ_{em} 655 nm) was measured with a FilterMax F5 microplate reader (Molecular Devices) and Multi-

Mode Analysis software (Version 3.4.0.27, Beckman Coulter).

2.4   Galectin Translocation Assay

MCF7 cells were plated on glass coverslips in 12-well plates and cultured to 70-80% confluency. Cells

were treated with vehicle, 2 mM LLOMe, 40 μM HMA, or 10 μM LLC1 for 12hr and then fixed with

4% paraformaldehyde for 10min. After fixation, detection of ruptured lysosomes by galectin-3

translocation was evaluated by immunocytochemistry as previously described [16]. Samples were

incubated with 50 mM ammonium chloride in PBS for 10min, washed three times with PBS, and then

permeabilized and blocked in 1 mL blocking buffer (1% BSA, 0.02% sodium azide, 0.2% NP40, 5%

goat serum) at room temperature for 1hr. Cells were incubated with purified mouse anti-human

Galectin-3 (1:100 dilution, #556904, BD Biosciences) diluted in blocking buffer for 1hr at room

temperature. Cells were washed three times with PBS and then incubated with goat anti-mouse IgG1

cross-absorbed secondary antibody Alexa Fluor 488 (1:1,000 dilution, #A-21121, Invitrogen) at room

temperature for 1hr. Cells were again washed three times with PBS before incubation with DAPI

(1:10,000) in PBS for 10min in the dark and then rinsed twice with PBS twice before mounting with

Fluoromount G. Slides were imaged by confocal microscopy using a ZEISS LSM 800 with Airyscan,

and galectin-3 puncta formation was analyzed from ten replicate fields with FIJI

(https://imagej.net/Fiji) software.

2.5   Ex Vivo Drug Cytotoxicity Organoid Assay

Tumor samples were generated from 1 x 10^6 4T1 cells engrafted into Balb/cJ mice (#000651, The

Jackson Laboratory) at the 4th mammary fat pad. Tumors (1.0-1.5 cm in diameter) or pooled mammary

glands from Balb/cJ mice were harvested and chopped into fine pieces using a razor blade before addition to digestion buffer (DMEM/F12 #SH30023, HyClone;1% PenStrep, Gibco - Thermo Fisher; 2% FBS, Genesee Scientific; collagenase/hyaluronidase (5%: tumors & 10%: mammary glands), #07912, StemCell Technologies; 5ug/mL insulin, Gibco - Thermo Fisher) and vortexed. Samples were digested for 2hr at 37°C under gentle shaking, and samples vortexed every 30min, before they were processed. After neutralization of the digestion buffer with DMEM (Gibco - Thermo Fisher) containing 10% FBS, organoids were centrifuged at 700rpm to remove cellular debris. ACK lysis buffer (Gibco - Thermo Fisher) was used to remove red blood cells from the organoids. A quick spin was performed to remove additional debris and to pellet organoids. Once resuspended in PBS, organoids were embedded into MatriGel (#354230, Corning) and cultured in organoid growth media (OGM) on 24-well plates for 24hr as described previously [17]. After overnight incubation in OGM, organoids were treated with vehicle or 10 µM LLC1 for up to 72hr, and viability was measured using RealTime Glo (#G9711, Promega) according to manufacturer's instructions. Images were taken before drug treatment (0hr) and every 24hr over the course of drug treatment. Representative brightfield images were taken with an Olympus IX81 microscope with CellSens Entry software. Chemiluminescent images were taken with a ChemiDoc Touch Imaging System (BioRad) and analyzed with FIJI (https://imagej.net/Fiji) software to quantify RealTime Glo signal.

2.6 Statistical, Data, and Image Analysis

A minimum of three replicates per experiment was performed with values calculated and expressed as averages, unless otherwise stated. P-values of less than 0.05 were considered statistically significant using a student's t-test. Data analysis was performed using Microsoft Excel or GraphPad Prism version 9.3.1 for Windows (GraphPad Software; San Diego, CA, USA; www.graphpad.com). Images were compiled in Microsoft PowerPoint and only brightness and contrast were altered for clarity.

3 Results

3.1 Addition of lipophilic substituents to amiloride C(2) and C(5) positions increases cytotoxic potency

Amiloride is often modified at the C(2), C(5), and C(6) positions of the pyrazine ring (see Fig. 1A) to improve its transporter inhibition activity and selectivity [7, 18], and such derivatives have been employed to assess the contribution of ion channel function to tumor growth and progression [8, 10, 11, 13, 19-28]. However, our recent studies indicating that the amiloride derivative HMA can provoke the non-apoptotic death of tumor cells relative to non-cancer cells by acting via the lysosome [11, 12] prompt the question of whether these cytotoxic properties might be further optimized.

As HMA adds a lipophilic hexamethylene group to the core pharmacophore at the C(5) position, we first asked whether lipophilic modification increases the cytotoxic potency of amiloride using immediately available amiloride derivatives. Fig. 1 depicts the structures of parent drug amiloride (Fig. 1A), C(2) derivative benzamil (Fig. 1B), C(5) derivative HMA (Fig. 1C), and C(5) derivative 10357 (Fig. 1D) [26], and summarizes the cytotoxic potency of these derivatives toward cell lines representing ER/PR-positive (MCF7), HER2-positive (SKBR3) and triple-negative (MDA-MB-231) breast tumor subtypes in an MTT cell viability assay (Fig. 1E). Consistent with previous observations [11, 22, 23, 27, 29, 30], we observed that amiloride modification with various lipophilic substituents at the C(2) or C(5) position markedly augments cytotoxicity, strongly suggesting that access to an intracellular target(s) is key to drug mechanism of action.

3.2 Novel amiloride derivatives underscore the importance of the guanidinium group

To more rigorously test the link between amiloride derivative lipophilicity and cytotoxic potential, we synthesized nine amiloride derivatives, all modified with various substituents at the C(5) position. The structures and EC_{50} values for cytotoxicity toward the three breast cancer cell line models are

summarized in Table 1, and the chemical characteristics of all derivatives ordered by EC_{50} are

summarized in Table 2. Three key points emerge from comparisons of these derivatives. First, Table 2

suggests a strong correlation between cytotoxic efficacy and lipophilicity, reflected in partition

coefficient (logP) values of the derivatives. For example, LLC1 and 10357, which exhibit some of the

highest logP values (2.46 and 2.88, respectively) among the group, also exhibit the most potent

cytotoxicity toward the breast cancer cell lines (5-13 µM and 9-17 µM, respectively). On the other

hand, the low logP values for amiloride and LLC10 (-0.89 and -0.28, respectively) correlate with

particularly poor cytotoxicity (148-995 µM and 883-2732 µM, respectively). Second, a direct

comparison of derivatives LLC1 and LLC3 (Table 1) illustrates the importance of maintaining the

integrity of the C(2) guanidinium structure. Both derivatives contain an identical lipophilic C(5)

substituent, but the removal of one of the guanidinium amines correlates with a 6-16 fold decrease in

cytotoxic potency. Finally, to a first approximation derivatives tend to act with similar potency toward

all three cell lines (Table 1). This suggests that, consistent with previous observations [11, 12],

amiloride derivatives act with similar efficacy toward the different breast cancer subtypes. These

observations reveal new parameters for the optimization of amiloride derivatives, and highlight LLC1

as a novel lead.

3.3   LLC1 reduces the viability of murine mammary tumor organoids ex vivo

We have previously demonstrated that HMA reduces the viability of organoids derived from the PymT

and NDL genetically-engineered mouse models of breast cancer [12]. To determine whether LLC1 is

selectively cytotoxic toward mammary tumors *ex vivo*, we compared its impact on tumor and normal

mammary gland organoids generated from Balb/cJ mice orthotopically engrafted with 4T1 (ER⁻/PR⁻

/HER2⁻), a murine mammary cancer cell line. The approximate EC_{50} value of LLC1 toward human

tumor cell lines (10 µM, Table 1) was used to compare the efficiency of cell death as a function of

time. We observed that while tumor organoids markedly shrink over 72 hours of treatment (Figs. 2A

and 2B), normal mammary gland organoids are resistant to LLC1 treatment (Figs. 2C and 2D). These observations suggest that, similar to its predecessor HMA [11, 12], LLC1 exhibits significant selectivity toward cancer cells and leaves untransformed cells largely unharmed.

3.4 HMA and LLC1 act on therapy-resistant breast cancer cell populations

Our previous observations indicate that HMA is cytotoxic toward 'persister' populations of breast cancer cells that survive 3-9 day exposure to conventional chemotherapeutic agents cisplatin, docetaxel and doxorubicin [12]. To confirm HMA's capabilities and test LLC1's properties, we examined the impact of both drugs on the viability of various breast cancer cell line models of short-term and established therapeutic resistance. We first examined the response of therapy-resistant populations of MDA-MB-231 cells toward the derivatives. In the experiment shown in Fig. 3, cells were first pretreated for 48 hours with DMSO vehicle, and then further treated for 24 hours with DMSO, 250 nM DTX, 5 μM DOXO, and either 40 μM HMA (Fig. 3A) or 10 μM LLC1 (Fig. 3D). In each case, drug treatment tended to lower cell viability, confirming that the drugs behave as expected. However, in cells pretreated for 48 hours with either DTX (Figs. 3B and 3E) or DOXO (Figs. 3C and 3F), a second challenge with the pretreating drug did not reduce viability while HMA or LLC1 treatment was cytotoxic.

We further observed that roughly half of T47D cells, an ER/PR$^+$ breast cancer cell line, exhibit resistance to cell death induced by 72 hour treatment with doxorubicin (DOXO, Fig. 4A) and docetaxel (DTX, Fig. 4D), reflected in the inability of high doses of either drug to further impact viability in a titration. However, we found that both HMA and LLC1 are cytotoxic toward the residual this population that remains after chemotherapeutic treatment, and exhibit similar EC$_{50}$ values as cell lines representing a variety of solid tumor types (Figs. 3B, 3C and 3E, 3F; refs [11, 12]).

Finally, to assess amiloride derivative effect on a model of sustained drug resistance, we examined the

impact of HMA and LLC1 on MCF7 cells selected for long-term resistance to tamoxifen (TAM), an

estrogen modulator, and mitoxantrone (MX), a DNA damaging agent [31-34]. Parental MCF7 and

MCF7 MX-100, a derivative selected for stable MX resistance that overexpresses ABCG2 (aka breast

cancer resistance pump, BCRP) by 30-fold, and MCF TS (tamoxifen sensitive derivative) and MCF7

TR-1 and MCF7 TR-5 (tamoxifen resistant derivatives), were treated with varying concentrations of

TAM, DTX, DOXO, HMA, and LLC1 and cytotoxicity EC_{50} values were determined (Table 3). We

observed that parental MCF7 cells are 3-13 fold more sensitive to chemotherapeutic agents DTX and

DOXO than MX-100, but both cell lines are similarly sensitive to HMA and LLC1. Likewise, MCF7

TS cells are far more sensitive to tamoxifen than MCF7 TR-1 and MCF7 TR-5 cells, but all three lines

ae similarly sensitive to HMA and LLC1. Collectively, these observations provide strong evidence that

amiloride derivatives act on breast tumor cell populations that are resistant to clinically pertinent anti-

cancer agents.

3.5   HMA and LLC1 cause lysosomal membrane permeabilization in resistant breast cancer

lines

We have previously demonstrated that HMA induces disrupts lysosomal morphology and that drug-

induced cell death is rescued by inhibition of the lysosomal protease cathepsin [11]. These observations

raise the possibility that amiloride derivatives act by inducing lysosomal membrane permeabilization

(LMP), and that the resulting release of cathepsin into the cytosol triggers the non-apoptotic lysosomal

cell death pathway. To determine whether the amilorides might act via this mechanism we employed

a sensitive galectin translocation assay [16], which measures the accumulation of diffusely distributed

cytosolic and nuclear galectin protein to punctate lysosomes upon LMP [35], to assess drug-induced

lysosomal disruption. In the experiments depicted in Fig. 5, MCF7 TR-5 cells were treated with for 12

hours with 2mM LLOMe (a potent LMP inducer used here as a positive control, refs [[35, 36]), 40 μM

HMA, or 10 μM LLC1, and galectin localization was imaged by confocal fluorescence microscopy.

We observed that both HMA and LLC1 induce the relocalization of galectin to lysosomes in MCF7

TR-5 cells (Fig. 5A), quantitatively reflected in an increase in the total number of puncta per cell (Fig.

5B), a loss in the percentage of cells exhibiting 0 puncta, and gains in the percentages of cells exhibiting

either 1-5 puncta or 6 or more puncta (Fig. 5C). A quantitative comparison of galectin relocalization

in MCF7 TS and MCF7 TR-1 (see Supplementary Figs. S3 & S4) to MCF7 TR-5 cells revealed that

both HMA and LLC1 similarly induce LMP in all three lines (Figs. 5D-5G). These observations

indicate that amiloride derivatives compromise the integrity of lysosomes, which in turn could

significantly contribute to their cytotoxic mechanism. Moreover, this process appears to be similarly

engaged in sensitive and drug-resistant cells, possibly accounting for the activity of these agents toward

resistant populations.

4   Discussion

Amiloride derivatives exhibit significant promise as cancer therapeutics because they have the potential

to eradicate therapy-resistant populations of breast cancer through permeabilization of the lysosome

and engagement of lysosome-dependent cell death (LDCD). The optimization of amiloride structure is

an enticing area for medicinal chemists and biologists alike because the relative ease of synthesis and

chemical modification allows the straightforward development of new derivatives, while optimizing

its unique anticancer properties. This study reveals that LLC1, a novel lipophilic C(5) amiloride

derivative, induces cell death in the single digit micromolar range, selectively kills tumor organoids

relative to normal mammary gland organoids, and acts to induce LMP. These observations suggest that

LLC1 has potential clinical applicability and warrants further testing and optimization as a lead

compound.

Our structure-activity relationship analysis provides key insight into building more effective cancer-

selective cytotoxic amiloride derivatives. First, we observe a strong relationship between amiloride

lipophilicity, reflected in more positive logP values, and drug cytotoxicity. Indeed, our most potent

derivative LLC1 exhibits one of the highest logP values of the compounds we have examined,

suggesting that improving lipophilicity further could result in even more effective drug leads. These

conclusions dovetail with current trends in drug development that prioritize more lipophilic drugs

because of their higher likelihood of passing safety and ADME concerns during Phase 1 clinical studies

[37]. At the same time, disruption of the guanidinium moiety at the C(2) position appears deleterious

to cytotoxic function. LLC1 and LLC3 are identical except for the deletion of one of the guanidinium

amino groups in LLC3, and this change leads to an order of magnitude loss of potency. In this regard

it might be interesting to determine whether lipophilic derivatives of other existing guanidinium-

containing drugs such as metformin might also exhibit tumor cell-selective cytotoxicity.

More broadly, this concept can be further applied to cationic amphiphilic drugs (CADs) which are

defined by their ability to accumulate in lysosomes because of their lipophilic nature and amine-

containing structures [38, 39]. As amiloride derivatives fall under the CAD family, it would be of

interest to perform screens such as that described by Pagliero et al. to assess the efficiency of LMP

initiation by newly synthesized C(5) amiloride derivatives relative to cytotoxicity EC_{50} values [40].

The use of omics and biochemical approaches could also prove fruitful in finding putative direct targets

of amiloride derivative action.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material,

further inquiries may be directed to the corresponding author.

Author contributions

Conceptualization: KLC; methodology: MH, RL; validation: MH; formal analysis: MH;

investigation: MH; resources: RL, KSL; writing-original draft preparation: MH; writing-review and

editing: MH, RL, KLC; visualization: MH; supervision: KSL, KLC; funding acquisition: KLC. All

authors have read and agreed to the published version of the manuscript.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial

relationships that could be construed as a potential conflict of interest.

Supplementary material

The Supplementary Material for this article may be found online at:

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TABLES

Name	MDA-MD-231 EC$_{50}$	MCF7 EC$_{50}$	SKBR3 EC$_{50}$	Structure		
LLC1	7 µM ± 0.4	13 µM ± 2	5 µM ± 0.6	R$_1$=	R$_2$=	
LLC2	43 µM ± 5	54 µM ± 5	97 µM ± 10	R$_1$=	R$_2$=	
LLC3	114 µM ± 25	75 µM ± 28	46 µM ± 6	R$_1$=	R$_2$=	
LLC4	40 µM ± 5	56 µM ± 17	34 µM ± 4	R$_1$=	R$_2$=	
LLC5	18 µM ± 1	29 µM ± 4	10 µM ± 0.6	R$_1$=	R$_2$=	
LLC7	111 µM ± 41	138 µM ± 58	129 µM ± 23	R$_1$=	R$_2$=	
LLC8	90 µM ± 11	75 µM ± 9	74 µM ± 7	R$_1$=	R$_2$=	
LLC9	52 µM ± 7	55 µM ± 11	53 µM ± 7	R$_1$=	R$_2$=	
LLC10	3 mM ± 1	1 mM ± 3	0.9 mM ± 0.4	R$_1$=	R$_2$=	

436 **Table 1. Potencies of novel 2- & 5-substituted amiloride derivatives.** The core pharmacophore with
437 positions of the two R groups is depicted to the right, while EC$_{50}$ data (from MDA-MB-231, MCF7,
438 and SKBR3 cells) and specific R groups for each compound are represented in the table. Data are
439 compiled from three biological replicates of each cell line.
440

Name	Rule of 5	PSA (Å²)	Pol (Å³)	MR (cm³/mol)	Acidic pKa	Basic pKa	PI	LogP	HLB	Sol (mg/ml)	Sol Level	IC₅₀ (µM)
LLC1	Y	136.51	37.29	103.42	14.85	7.12	10.98	2.46	11.477	0.015	Moderate	5-13
10357	Y	142.5	39.64	102.68	14.85	5.46	10.15	2.877	11.712	0.001	Low	9-17
LLC5	Y	136.51	33.88	94.34	14.85	7.1	10.97	1.811	12.4	0.037	Moderate	10-29
HMA	Y	136.51	29.94	83.16	14.85	7.1	10.97	1.342	13.109	0.188	High	18-36
LLC4	N	158.21	41.7	112.85	14.85	7.15	10.99	1.241	20.051	0.06	Moderate	34-56
LLC9	Y	174.84	41.1	111.49	14.5	7.12	10.76	1.21	17.68	0.121	High	52-55
LLC2	Y	136.51	31.01	86.18	14.85	7.12	10.98	0.905	12.697	0.152	High	43-97
Benzamil	Y	145.3	31.01	86.08	14.68	7.42	11.04	1.112	13.059	0.073	High	50-114
LLC3	Y	110.16	34.8	97.56	12.47	2.68	7.57	2.834	16.875	0.018	Moderate	46-114
LLC8	Y	136.51	27.86	79.52	14.85	7.13	10.99	0.691	13.7	0.263	High	74-90
LLC7	Y	171.31	31.46	86.21	14.47	9.89	12.13	0.029	18.857	0.215	High	111-138
Amiloride	Y	159.29	19.74	56.69	14.85	7.29	11.06	-0.89	15.933	2.768	High	148-995
LLC10	Y	162.53	31.46	86.62	14.85	10.22	12.53	-0.282	19.225	1.1	High	883-2732

441
442
Table 2. Chemical and cytotoxicity characteristics of amiloride derivatives. The chemical
properties of the novel amiloride derivatives are ordered according to increasing collective cytotoxicity
EC₅₀ toward MCF7, SKBR3 and MDA-MB-231 breast cancer cell lines. All of the chemical data [Rule
of 5, topological polar surface area (PSA), polarizability (Pol), molar refractivity (MR), acidic pKa,
basic pKa, isoelectric point (PI), logP, hydrophilic-lipophilic balance (HLB), intrinsic solubility (Sol),
and solubility level (Sol level)], were obtained with Chemicalize [July 2022]; https://chemicalize.com/,
developed by ChemAxon.
450

	DTX (µM)	DOXO (µM)	HMA (µM)	LLC1 (µM)
MCF7	1 ± 0.6	2 ± 0.3	27 ± 2	13 ± 2
MCF7 MX-100	13 ± 3	7 ± 0.5	39 ± 3	12 ± 1

	TAM (µM)	HMA (µM)	LLC1 (µM)
MCF7 TS	13 ± 1	68 ± 3	25 ± 0.9
MCF7 TR-1	>10*	67 ± 4	26 ± 2
MCF7 TR-5	>10*	59 ± 5	19 ± 2

Table 3. MCF7 chemoresistant cells are susceptible to HMA and LLC1 but not conventional
chemotherapeutics. A table illustrating the variety of MCF7-chemoresistant cells treated with their
respective chemotherapeutic (tamoxifen (TAM) or mitoxantrone (MX)), HMA, or LLC1 for 24hr to
determine their EC50 values. An asterisk denoted by a value indicated the EC_{50} value could not be
calculated and is the high dose used to treat the cells. Data is represented from a minimum of three
independent studies.

FIGURES

Figure 1. Modification of amiloride with lipophilic substituents enhances its cytotoxic potency.

EC_{50}	Amiloride (µM)	Benzamil (µM)	HMA (µM)	10357 (µM)
MDA-MB-231	398 ± 44	114 ± 32	36 ± 2	17 ± 2
MCF7	995 ± 341	56 ± 12	27 ± 2	15 ± 2
SKBR3	148 ± 15	50 ± 6	18 ± 1	9 ± 0.7

463 **Figure 1. Modification of amiloride with lipophilic substituents enhances its cytotoxic potency.**

464 **(A)** Structure of amiloride with its key carbons numbered. **(B-D)** 2- and 5-substituted amiloride

465 derivatives. **(E)** Cytotoxic potency of amilorides toward cultured breast cancer cell lines.

Figure 2. LLC1 selectively reduces viability of mouse mammary tumors *ex vivo*. Representative

images of organoids generated from mammary tumors **(A)** and normal mammary glands **(C)** of 4T1-

engrafted Balb/cJ mice are shown. Scale bar = 100 μm. **(B, D)** Organoids were treated with 10 μM

LLC1, and viability was assessed using RealTime Glo over 72hr and normalized to vehicle (DMSO).

Tumor **(B)** or mammary fat pad (MFP) **(D)** organoid viability was monitored every 24hr. Data was

compiled from three biological replicates of organoids derived from tumors or mammary glands of

independent mice.

Figure 3. LLC1 is cytotoxic toward chemoresistant triple-negative breast cancer cell populations.

(A,D) MDA-MD-231 cells were initially treated with DMSO for 48hr, followed by a second challenge

with DMSO, 250 nM docetaxel (DTX), 5 µM doxorubicin (DOXO), and either 40 µM HMA **(A)** or

10 µM LLC1 **(D)** for 24hr, and then cell viability was determined. **(B,E)** Following 48hr 250 nM DTX

(B, E) or 5 µM DOXO **(C, F)** pretreatment, the residual population was treated with vehicle, initial

chemotherapy treatment (DOXO or DTX), HMA **(B, C)**, or LLC1 **(E, F)** for an additional 24hr. Data

is represented from a minimum of four independent studies.

Figure 4. LLC1 is cytotoxic toward chemoresistant ER/PR-positive breast cancer cell

populations. Dose response curves of T47D cells treated with doxorubicin (DOXO) **(A)** or docetaxel

(DTX) **(D)** for 72h are depicted. Populations of cells that persist at drug concentrations higher than

650 nM DOXO and 800 nM DTX (arrows) are considered drug resistant. Residual T47D cells that

survive treatment with these levels of drugs for 72h were then titrated with HMA **(B, E)** or LLC1 **(C,**

F) for an additional 24h and viability assessed. Data represent at least three biological replicates.

Figure 5. LLC1 induces lysosomal membrane permeabilization in drug-resistant cells. (A-C)
MCF7 cells resistant to 5 μM tamoxifen (MCF7 TR-5) cells were treated with vehicle (DMSO), 2
mM LLOMe positive control, 40 μM HMA, or 10 μM LLC1 for 12hr, and galectin translocation into
puncta was assessed by confocal fluorescence microscopy. **(A)** Representative images of treated
MCF7 TS cells are depicted. Scale bar = 100μm. **(B)** Average puncta per cell was calculated and
plotted. **(C)** Percentages of cells exhibiting 0, 1-5 and 6 or more puncta for each drug condition was
plotted. **(D-G)** A similar analysis of MCF7 TR-1 and MCF7 TR-5 cells allowed comparisons of
average puncta/cell **(D)**, and percentages exhibiting 0 **(E)**, 1-5 **(F)**, or 6 or more **(G)** puncta. Data is

502   compiled from a minimum of four biological replicates from each cell line. *, $P < 0.05$; **, $P < 0.01$;

503   ***, $P < 0.001$; ****, $P < 0.0001$.