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Melittin stimulates fatty acid release through non-phospholipase-mediated mechanisms and interacts with the dopamine transporter and other membrane spanning proteins

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Abstract

Phospholipase A₂ releases the fatty acid arachidonic acid from membrane phospholipids. We used the purported phospholipase A₂ stimulator, melittin, to examine the effects of endogenous arachidonic acid signaling on dopamine transporter function and trafficking. In HEK-293 cells stably transfected with the dopamine transporter, melittin reduced uptake of [³H]dopamine. Additionally, measurements of fatty acid content demonstrated a melittin-induced release of membrane-incorporated arachidonic acid, but inhibitors of phospholipase C, phospholipase D, and phospholipase A₂ did not prevent the release. Subsequent experiments measuring [¹²⁵I]RTI-55 binding to the dopamine transporter demonstrated a direct interaction of melittin, or a melittin-activated endogenous compound, with the transporter to inhibit antagonist binding. This effect was not specific to the dopamine transporter, as [³H]spiperone binding to the recombinant dopamine D₂ receptor was also inhibited by melittin treatment. Finally, melittin stimulated an increase in internalization of the dopamine transporter, and this effect was blocked by pretreatment with cocaine. Thus, melittin acts through multiple mechanisms to regulate cellular activity, including release of membrane-incorporated fatty acids and interaction with the dopamine transporter.

Keywords

endocytosis; arachidonic acid; phospholipase; dopamine transporter; dopamine receptor

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1. Introduction

The arachidonic acid cascade is a complex biological signaling pathway that is found in most cell types. Free arachidonic acid is generated by phospholipase A₂-mediated hydrolysis of membrane phospholipids. Altered phospholipase A₂ activity is implicated in many neurological disorders such as dyslexia (MacDonell et al., 2000), Parkinson's disease (Ross et al., 2001) and schizophrenia (Horrobin et al., 1994). Additionally phospholipase A₂ activators can induce (Reid et al., 1996) while inhibitors of phospholipase A₂ (Reid et al., 1996) and of arachidonic acid metabolism (Reid et al., 2002) can inhibit, psychostimulant sensitization. Considering that hyperactivity of the dopamine system may be a component of the etiology of schizophrenia, and the dopamine transporter is the primary mechanism of clearing released dopamine from the extra-synaptic space (Parsons and Justice, Jr., 1994; Wightman and Zimmerman, 1990) and is the site of action of psychostimulants such as cocaine and amphetamines (Ritz and Kuhar, 1989), understanding the effects of the arachidonic acid signaling cascade on the dopamine transporter could be crucial to understanding symptoms of addiction and other neuropsychiatric disorders.

In purified synaptosomes from rat striatum, arachidonic acid dose-dependently increases spontaneous release of dopamine and decreases dopamine synthesis and uptake. This effect is mediated by arachidonic acid itself rather than by a metabolite (L'hirondel et al., 1995). Arachidonic acid also inhibits uptake of glutamate (Volterra et al., 1992, 1994) and alters proton conductance by the rat glutamate transporter, EAAT4 (Fairman et al., 1998; Tzingounis et al., 1998). Arachidonic acid also directly stimulates a cocaine-sensitive cation conductance in the dopamine transporter that is distinct from substrate associated current (Ingram and Amara, 2000).

Phospholipase A₂, which can be stimulated by G-protein coupled dopamine D₂ receptors, hydrolyzes membrane phospholipids to cause a release of platelet-activating factor and arachidonic acid (Piomelli et al., 1991; Vial and Piomelli, 1995). In bovine cerebellar cells arachidonic acid can activate protein kinase C_γ (Shearman et al., 1989), yet Zhang and Reith (1996) have shown that arachidonic acid effects on dopamine transporter function are not mediated by protein kinase C. Thus, the mechanisms mediating arachidonic acid's effects on the dopamine transporter are not clear.

Melittin is a 26 amino acid amphiphilic peptide isolated from the venom of the honeybee, *Apis mellifera*. Melittin activates arachidonic acid signaling pathways in neurons (Geddis et al., 2004; Muzzio et al., 2001) and in multiple cell lines including rat PC12 and mouse L1210 cells (Lee et al., 2001; Palomba et al., 2004), possibly through phospholipase A₂ stimulation. Additionally melittin acts through non-phospholipase A₂-mediated mechanisms to release fatty acids and alter cell function (Lee et al., 2001). While melittin has been used to examine the effects of phospholipase A₂ signaling on dopamine cell function (Reid et al., 1996; Zhang and Reith, 1996), the mechanism of action has been presumed to be release of arachidonic acid. The potential contributions of extra- arachidonic acid effects of melittin to dopamine cell function have not been explored. In this study we used HEK cells stably transfected with the dopamine transporter to test the hypothesis that melittin stimulates phospholipases and affects dopamine transporter function and trafficking.

2. Material and Methods

2.1 Materials

Melittin was purchased from Sigma-Aldrich (St. Louis, MO USA). The radioligands [³H] dopamine (59 Ci/mmol), [³H] arachidonic acid (60 Ci/mmol), and [³H]spiperone (73 Ci/mmol) were purchased from Amersham Biosciences (Piscataway, NJ USA), and

[¹²⁵I]RTI-55 (2200 Ci/mmol) was purchased from Perkin Elmer (Wellesley, MA USA). Rat anti-human dopamine transporter antibody was purchased from Chemicon (Temecula, CA USA) and goat anti-rat secondary conjugated to alkaline phosphatase was purchased from Santa Cruz (Santa Cruz, CA USA). Other reagents were purchased from commercial sources (Sigma Aldrich, St Louis, MO USA)

2.2 Stable Cell Line Transfection

The cDNAs were cloned and human embryonic kidney-293 (HEK) cells were cotransfected with the cDNAs for the human dopamine transporter (hDAT) (in pcDNA1), the long form of the dopamine D₂ receptor (in pcDNA1) and the antibiotic resistance vector pBabepuro at stoichiometries of 7:7:1 using the calcium phosphate precipitation method (Chen and Okayama, 1988). The cloning and characterization of the human dopamine transporter and D₂ receptor cDNAs were described previously (Eshleman et al., 1995; Neve et al., 1989). Cells were selected for resistance to puromycin (2 µg/ml), and grown in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum, 0.05 U penicillin/streptomycin, 2 µg/ml puromycin, in a humidified 10% CO₂ incubator at 37°C

2.3 [¹²⁵I]RTI-55 binding

Experiments were performed as previously described (Eshleman et al., 1999). Briefly, cells were grown to confluence on 15 cm tissue culture plates. The cells were rinsed with Ca²⁺, Mg²⁺-free phosphate buffered saline (PBS; 0.1 M H₂PO₄, 150 mM NaCl), and lysis buffer (2 mM HEPES, 1 mM EDTA) was added. Cells were then removed from the plate, centrifuged for 20 min at 30,000g, and the pellet was homogenized in 5 ml of 0.32 M sucrose with a Polytron homogenizer for 5 s. Membrane aliquots (10–30 µg of protein) were added to each assay, which resulted in binding of <10% of the total radioactivity added. Assays were performed in duplicate in Krebs-Hepes buffer (25 mM HEPES, 122 mM NaCl, 5 mM KCl, 1.2 mM MgSO₄, 2.5 mM CaCl₂, 1 µM pargyline, 100 µM tropolone, 2 mg glucose/ml, 1 mM ascorbic acid, pH 7.4), in a final volume of 250 µl. [¹²⁵I]RTI-55 (40 pM) was added and membranes were incubated for 90 min at room temperature in the dark. Nonspecific binding was determined in the presence of 10 µM mazindol. The assay was terminated by filtration through Perkin Elmer filtermat A filters (Boston, MA USA) using a 96-well Tomtec cell harvester (Hamden, CT USA). Scintillation fluid was added to the filters and remaining radioactivity was determined using a Perkin Elmer 1205 betaplate scintillation counter. For the saturation binding experiments [¹²⁵I]RTI-55 was diluted with unlabeled ligand at concentrations ranging from 0.04–10 nM.

2.4 [³H]spiperone binding

Cells were grown to confluence on 10 cm tissue culture plates then lysed with lysis buffer, centrifuged at 31,000g for 20 min, re-suspended and homogenized with a polytron (setting 6 for 10 seconds) in Tris-buffered saline (TBS; 20 mM Tris, 137 mM NaCl, pH 7.6). [³H]Spiperone binding assays were performed as previously described (Neve et al., 1989). Briefly, cell homogenates were incubated with the radioligand (0.0125 nM to 0.4 nM) for 60 min at 37°C. The assay was performed in TBS plus bovine serum albumin (BSA; 1 mg/100 mL) in a total volume of 1mL. Nonspecific binding was determined in the presence of 2 µM butaclamol. The assay was terminated and radiation quantified as described above. Experiments were conducted with duplicate determinations at each radioligand concentration.

2.5 [³H]Dopamine uptake

For attached cell assays, cells were plated in 24-well tissue culture plates coated with poly-D-lysine (1 mg/ml in H₂O). Cells were incubated at 37°C with drugs in Krebs-Hepes buffer

in a total volume of 500 μL . The uptake assay was initiated by addition of 50 nM [^3H]dopamine at 37°C. Nonspecific uptake was determined in the presence of 10 μM mazindol, added 5 min before the radioligand. Uptake continued for 10 min and was terminated by washing the cells twice with ice-cold buffer. Radioactivity was extracted by rupturing the cellular membranes with 3% trichloroacetic acid for 30 min. Extracted radioactivity was measured using a Beckman LS 3801 scintillation counter (Fullerton, CA USA). The average values of all experiments for the maximal amount of radioactivity transported into the cells was $14,700 \pm 1700$ cpms and for the nonspecific cpms was 890 ± 120 . Cells from one well of each assay plate were used to determine protein concentrations using the BCA protein assay kit (Pierce, Rockford, IL USA).

2.6 Fatty acid analysis by gas-liquid chromatography (GLC)

Cells grown on tissue culture plates were treated with drug in PBS at 37°C and the reaction was stopped by rapidly cooling the cells on ice. Cells were then rinsed with fatty-acid-free bovine serum albumin (0.1mg/ml in PBS) followed by a rinse in PBS. The cells were scraped from the plates and centrifuged at 200 \times g for 10 min. The pellets were retained for analysis. Fatty acid extraction and quantification was performed by the Lipid Atherosclerosis Laboratory at Oregon Health & Science University using the chloroform-methanol procedure described previously (Rapp et al., 1983) with some modifications. The cell extracts were mixed with chloroform:methanol (2:1), centrifuged, and the top layer removed. Free cholesteryl ester was saponified with ethanol and KOH at 37°C for 60 min. The cholesterol was extracted with hexane. Acidifying the aqueous phase and re-extracting with hexane recovered the cholesteryl ester fatty acids. The fatty acids were methylated with boron trifluoride-methanol and percents of total fatty acid were determined by gas-liquid chromatography using a 30 meter Supelco SP 2330 fused silica capillary column (run at 190°) attached to a Perkin-Elmer AutoSystem chromatograph and a Perkin-Elmer Turbochrome 41 integrator. Fatty acid standards (Supelco, Inc., Bellefonte, PA USA) were analyzed in each experiment. The 17-carbon fatty acid (C17:0) was added to each sample to determine μg amounts of each fatty acid analyzed based on relative abundance to this standard. Levels of C17:0 are extremely low in naïve cells.

2.7 [^3H]Arachidonic acid release

Cells were grown on poly-D-lysine coated, 24-well tissue culture plates and incubated in DMEM with 1nM [^3H]arachidonic acid (106,000 dpm / well) for 18–24 h. DMEM supplemented with 0.2% fatty acid free BSA was used for all rinses and assay conditions. Following [^3H]arachidonic acid incubation the cells were rinsed twice to remove unincorporated [^3H]arachidonic acid. The treatments were performed in a total volume of 500 μL . Following treatments the medium was removed and centrifuged at 12,000 \times g for 1 min. Released [^3H]arachidonic acid in 400 μl of supernatant was measured using a Beckman LS 3801 scintillation counter. Trypsin-EDTA was added to each well to remove cells remaining on the plates. This sample was also counted to measure incorporated (non-released) [^3H]arachidonic acid. The value for non-released [^3H]arachidonic acid was combined with the value for released [^3H]arachidonic acid to determine the total amount of [^3H]arachidonic acid incorporated into cellular membranes in each well. Cells from one well of each assay plate were used to determine protein concentrations using the BCA protein assay kit (Pierce). Experiments were conducted in triplicate for each time point or drug concentration.

To address the contribution of the dopamine D₂ receptor to [^3H]arachidonic acid release, additional assays were conducted using a modification of a method optimized to detect [^3H]arachidonic acid release following activation of the G-protein coupled 5-hydroxytryptamine 2A receptor (Kurrasch-Orbaugh et al., 2003). Briefly, cells were plated

as above and incubated with [³H]arachidonic acid (0.5 μCi) for 4 h in DMEM. Medium was removed, the wells were washed 3 times with supplemented DMEM, and cells were treated with vehicle or the dopamine D₂ agonist quinpirole for 1 h. The reaction was terminated and radioactivity measured as described above.

2.8 Phospholipase activity measurements

To determine the ability of melittin to activate phospholipase A₂ activity in HEK- D₂ -hDAT cells, phospholipase A₂ activity was determined using the cytoplasmic-phospholipase A₂ Assay Kit (Cayman Chemical, Ann Arbor, MI). In cell preparations that contain more than one type of phospholipase A₂, the kit measures total phospholipase A₂ activity. Cells were plated on 10 cm cell culture plates and tested 4 days later. Cells were treated with vehicle or melittin (10 μM, for 30 min) with or without quinacrine (10 μM). Cells were then rinsed with phosphate-buffered saline, removed from plate and centrifuged for 10 min at 4°C. The pellet was homogenized in 0.5 ml cold buffer (50 mM Hepes pH 7.4, 1 mM EDTA) and centrifuged at 14,300×g for 15 min at 4°C. The resulting supernatant was stored at -80°C until assayed. On the day of assay, the supernatant was thawed, and centrifuged as above. The cytoplasmic-phospholipase A₂ kit was used as directed with 30–60 μg protein, results were expressed as μmol product/min/mg protein using bee venom cytoplasmic-phospholipase A₂ (2–16 ng) as the internal standard.

Activity of phospholipase D was measured using the Amplex Red Phospholipase D assay kit (Molecular Probes). Plated cells were treated with melittin (1 μM) for 30 min either alone, or following a 30 min pretreatment with propanolol (1–100 μM). Cells were scraped from the plates and suspended in kit assay buffer supplemented with 1% triton-X-100, protease inhibitors, and PMSF. Cells were homogenized by rotating for 30 min at 4°C and lysates were collected following centrifugation (15,000 rcf, 10 min). Lysates were added directly to the assay according to company directions. The assay product was detected using a fluorescence plate reader (Molecular Devices).

To determine the ability of melittin to activate phospholipase C activity in HEK- D₂ -hDAT cells, inositol phosphate formation was measured using the inositol phosphate-One Elisa Kit (CisBio, Bedford, MA). Cells were plated in 24 well plates at a density of 400,000 cells per well in DMEM supplemented with charcoal-stripped FetalClone. The next day, cells were rinsed and incubated for 1 h with unsupplemented DMEM without or with 30 μM neomycin. After aspiration of medium, cells were treated with stimulation buffer from the kit with or without 30 μM neomycin, and either buffer, 1.3 μM melittin, 1 μM quinpirole, or 1 μM dopamine. Following 1 hour incubation, the cells were lysed and processed according to kit protocols.

2.9 Biotinylation of membrane proteins

To measure dopamine transporter that was internalized following drug treatment, cells were incubated with 0.3 mg/ml NHS-SS-biotin (Pierce) in PBS at 4°C with gentle agitation for 40 min. Unbound biotin was removed by rinsing with 0.1 M glycine in PBS. Cells were then incubated with drug or vehicle in DMEM at 37°C and the reaction was stopped by removal of the media and rapid cooling of the plates on ice. Remaining cell surface biotin was stripped with glutathione (150 mM glutathione, 150 mM NaCl, pH 8.75). Cells were then rinsed with 50 mM iodoacetamide in PBS to neutralize the glutathione, and solubilized in CHAPS buffer (1% CHAPS, 25mM Tris, pH 7.4, 150 mM NaCl), and lysed using a glass-glass homogenizer. The samples were centrifuged at 14,000×g, and the supernatants were reserved. The protein concentration of each sample was determined using the BCA Assay kit (Pierce). Biotinylated proteins were isolated from nonbiotinylated proteins by incubation with ImmunoPure™ Immobilized streptavidin (Pierce) for 40 min at 4°C with gentle

agitation. Proteins were eluted from the streptavidin beads with 30 μ l of Laemmli sample buffer (Sigma) and constant mixing for 20 min. Samples were separated by SDS-PAGE and transferred to PVDF membranes for western blotting and dopamine transporter detection with anti-dopamine transporter antibody (Chemicon). Dopamine transporter immunoreactivity was quantified by densitometry on a Typhoon phosphorimager using ImageQuant software (Molecular Dynamics; Sunnyvale CA USA).

2.10 Data Analysis

Dose-response curves were analyzed by nonlinear regression using Prism 3.0 (GraphPad Software, San Diego, CA USA). Results were analyzed using Student's t-test or one-way ANOVA followed by Tukey's multiple comparison test, as appropriate.

3. RESULTS

3.1 Characterization of transfected HEK cells

To produce a cell model for examining the effects of melittin on dopamine transporter trafficking and regulation, HEK cells were stably transfected with the human dopamine transporter and the dopamine D₂ receptor. Saturation binding of the recombinant dopamine transporter was determined in the presence of 40 pM [¹²⁵I]RTI-55 diluted with unlabeled ligand. The B_{max} of the binding site was 2.42 ± 0.12 pmol/mg and the K_D was 1.71 ± 0.23 nM (Fig 1A). Saturation binding experiments to characterize recombinant D₂ receptor expression were conducted using [³H]spiperone concentrations of 0.0125nM to 0.4nM. The B_{max} for radioligand binding was 261.1 ± 33.5 fmol/mg with a K_D of 0.10 ± 0.03 nM (Fig 1B).

3.2 Melittin treatment decreases dopamine transporter function

To determine the effects of phospholipase A₂ activation on dopamine transporter function, HEK-D₂-hDAT cells were incubated with melittin for 10–120 min (Table 1) followed by incubation with 50 nM [³H]dopamine for 10 min. Functional activity of the dopamine transporter was quantified by measuring the [³H]dopamine taken up by the cells. Melittin treatment times longer than 5 min reduced uptake almost completely. The IC₅₀ concentration for melittin's effects was 1.3 μ M at the 30 min time point. This concentration and treatment time were used in subsequent assays. Melittin was significantly less potent following a 5 min treatment, as compared to any other time point (one-way ANOVA, P<0.001). To eliminate possible effects of feedback by the D₂ receptor on dopamine uptake, butaclamol (100 nM) was added to the cells 10 min before uptake was initiated. There was no difference in [³H]dopamine uptake between butaclamol-treated and untreated cells during this time period (Fig 2).

These findings agree with previous reports demonstrating a melittin-induced decrease in dopamine transporter-mediated uptake of dopamine following 15 min of treatment of rat C6 glioma cells transfected with the dopamine transporter. The reported decrease in uptake coincided with a decrease in binding of the cocaine analogue WIN 35,428 to the dopamine transporter in whole, attached cells (Zhang and Reith, 1996). The goal of the present study was to further characterize this effect and determine the precise mechanism of melittin's action.

3.3 Melittin releases arachidonic acid from cellular membranes

To examine the ability of melittin to activate phospholipase A₂, the levels of fatty acids remaining in cellular membranes following treatments were quantified by GLC. Cells were treated with melittin (1.3 μ M) for 30 min then rinsed with fatty acid-free BSA in PBS to bind and remove any free, released arachidonic acid. Following treatment, the membrane

bound arachidonic acid was $75.33 \pm 0.06\%$ of vehicle-treated levels (Fig 3). There was also a 20% reduction in the arachidonic acid precursor linoleic acid. Both values are significant ($P < 0.001$) reductions. In addition to these expected reductions, there were also reductions in the saturated fatty acids myristic, palmitic, and stearic acid of 24%, 20%, and 18% respectively, and a 21% reduction in the unsaturated fatty acid oleic acid. These results demonstrate a melittin-induced release of arachidonic acid. Release of the other fatty acids suggests additional actions of melittin on membrane lipids, not specific to phospholipase A₂ stimulation.

While melittin is typically utilized to stimulate phospholipase A₂, recent studies have revealed that melittin simultaneously stimulates multiple phospholipases in mouse L1210 cells (Lee et al., 2001). To determine the specific phospholipase stimulated by melittin in transfected HEK cells, [³H]arachidonic acid was incorporated into lipid membranes, the cells were treated with melittin and inhibitors of phospholipase C, phospholipase D, or phospholipase A₂, and the amount of [³H]arachidonic acid released following treatment was quantified. Initially, cells were treated with melittin for 30 min, demonstrating a dose-dependent release of [³H]arachidonic acid. Following treatment with 1.3 μM melittin the amount of released [³H]arachidonic acid was $324.1 \pm 34.2\%$ of baseline release (Fig 4A), $28.4 \pm 2.5\%$ of total incorporated [³H]arachidonic acid (Fig 4B), and 77.4 ± 8.0 fmol [³H]arachidonic acid/mg of protein (data not shown). These data using preloaded arachidonic acid correlate with GLC quantification of endogenous arachidonic acid released following melittin treatment.

Arachidonic acid can be released directly in a single reaction by phospholipase A₂. While not releasing arachidonic acid directly, phospholipase C and phospholipase D generate free fatty acids that contain arachidonate and action by subsequent enzymes can produce free arachidonic acid (for review see Piomelli, 1995). To determine if these phospholipases were involved in [³H]arachidonic acid release, cells were preloaded with [³H]arachidonic acid and treated for 60 min with phospholipase inhibitors alone or in combination with 1.3 μM melittin during the final 30 min of treatment. Treatment with neomycin, a phospholipase C inhibitor (Carney et al., 1985), did not prevent melittin-induced [³H]arachidonic acid release (Fig 4C). While propranolol is an antagonist of β-adrenergic receptors (Mehvar and Brocks, 2001), it has also been used to inhibit phospholipase D pathway-mediated release of arachidonic acid in a variety of cells, including stimulated aortic cells (Shinoda et al., 1997), epithelial keratinocytes (Lefkowitz and Smith, 2002), and myocytes (Albert et al., 2005). Treatment with propranolol did not inhibit melittin-induced [³H]arachidonic acid release (Fig 4D). Surprisingly the general phospholipase A₂ inhibitor quinacrine (Demuth et al., 2005) was also without effect on melittin-induced [³H]arachidonic acid release (Fig 4E). Phospholipase A₂ inhibitors successfully block melittin-induced release of arachidonic acid in rabbit proximal tubule cells (Han et al., 2002), but quinacrine is unable to inhibit melittin-induced release of arachidonic acid in rat Leydig cells (Ronco et al., 2002).

Since the effect of melittin may be cell type specific we additionally chose inhibitors of specific subtypes of phospholipase A₂ to determine if these enzymes were stimulated by melittin to cause release of arachidonic acid. Phospholipase A₂ can be divided into extracellular and intracellular types. The extracellular type is a secreted enzyme (secreted phospholipase A₂) that does not exhibit preference for particular fatty acids, contains disulfide bonds, and is sensitive to reducing agents, such as dithiothreitol. The intracellular types can be further divided into Ca²⁺-dependent phospholipase A₂, requiring 0.1–1 μM of free Ca²⁺ for activation, and Ca²⁺-independent phospholipase A₂ (for review see Dennis, 1994; Glaser et al., 1993). To inhibit secreted phospholipase A₂, cells were treated with the reducing agent dithiothreitol prior to the application of melittin. This agent was unable to inhibit the melittin-induced release of [³H]arachidonic acid (Fig 4F). The Ca²⁺-independent

phospholipase A₂ subtype is selectively inhibited by 4-bromo-enol lactone (Yang et al., 1999), and application of this inhibitor did not inhibit the effect of melittin (Fig 4G). Arachidonyl trifluoromethyl ketone is a potent and selective inhibitor of Ca²⁺-dependent phospholipase A₂ (Street et al., 1993), yet this compound was also unable to inhibit melittin-induced release of [³H]arachidonic acid in these cells (Fig 4H).

To determine if the dopamine D₂ receptors co-expressed in these cells contributed to [³H]arachidonic acid release, cells were loaded with [³H]arachidonic acid for either 4 or 24 hours. Treatment with the dopamine D₂ agonist quinpirole at concentrations ranging from 1 nM to 10 μM released less than 0.4% of loaded [³H]arachidonic acid under both conditions (data not shown), indicating minimal contribution of D₂ receptor activity to [³H]arachidonic acid release.

To determine if we could directly measure melittin-induced phospholipase A₂ activity, we treated cells with vehicle or melittin (10 μM) with or without the general phospholipase A₂ inhibitor quinacrine (10 μM) for 30 min and then measured cellular phospholipase A₂ activity. Cellular control phospholipase A₂ activity ranged from 0.114–0.123 μmol product/min/mg protein, and there was no significant effect of melittin (90 ± 16%), quinacrine (134 ± 19%) or melittin + quinacrine (97 ± 8%) compared to control (n=3, one way ANOVA, P>0.05).

To further clarify the effect melittin has on phospholipase D activity we treated cells with vehicle or melittin (1.3 μM) for 30 min with or without a pretreatment of propranolol (10 μM, 30 min). Following melittin treatment phospholipase D activity was 79.8 ± 10% of the activity detected in vehicle treated cells. Phospholipase D activity following propranolol alone was 62.8±14.7% of vehicle treatment and following propranolol + melittin was 73.1 ± 22% of vehicle treatment. None of these differences were significant (n=5, ANOVA, P>0.05) indicating that melittin does not activate phospholipase D in these cells.

To determine the ability of melittin to activate phospholipase C in HEK-D₂-hDAT cells, we pretreated cells with vehicle or neomycin (30 μM) and then incubated cells with melittin (1.3 μM), quinpirole (1 μM) or dopamine (1 μM) for one hour. The basal inositol phosphate-1 concentration was 8.3 ± 2.5 nM, and there was no significant effect of melittin (9.5 ± 3.7 nM), quinpirole (11.1 ± 2.0 nM) or dopamine (8.8 ± 3.0 nM, n=3, one way ANOVA, P>0.05). There was also no significant difference between phospholipase C activity in the presence of melittin without or with neomycin. These data indicate that neither melittin nor the D₂ receptor activates phospholipase C in these cells.

3.4 Melittin decreases antagonist binding to the transporter

To examine possible mechanisms for melittin inhibition of dopamine transporter-mediated uptake, binding of the cocaine analogue [¹²⁵I]RTI-55 to the transporter was examined. Membranes from homogenized cells were treated with varying concentrations of melittin (Fig 5A) in the presence of 40 pM [¹²⁵I]RTI-55. The IC₅₀ value for melittin inhibition of [¹²⁵I]RTI-55 binding was 0.88 μM, and the Hill slope was -1.75 ± 0.21. In saturation binding assays, concurrent addition of melittin (1.3 μM; Fig 5B) to the [¹²⁵I]RTI-55 binding assay significantly reduced the B_{max} value (Table 2; P<0.05, one-way ANOVA, followed by Dunnett's multiple comparison test) while there was no significant effect on the K_D value (P>0.05). The decrease in B_{max} suggests that melittin acts like an apparent noncompetitive inhibitor of RTI-55 binding to the dopamine transporter (Limbird, 2005; Tomlinson, 1988). A higher concentration of melittin (10 μM) further reduced the B_{max} value while the K_D value was not significantly altered, as compared to untreated cells (Table 2).

To determine if decreased membrane fluidity played a role in melittin's inhibition of antagonist binding to the dopamine transporter, RTI-55 binding experiments were conducted at 4°C. Parallel experiments were conducted at room temperature and at 4°C in the presence of melittin (1.3 μM). The experiments continued for 16 h to allow both temperature conditions to come to equilibrium. All samples contained protease inhibitors to prevent protein degradation at room temperature. For experiments conducted at 4°C, all reagents, drugs, and membranes were chilled on ice before the assays were initiated. Melittin treatment significantly reduced the B_{\max} from 6.7 ± 0.8 to 2.9 ± 0.3 pmol/mg at 4°C and from 7.2 ± 1.3 to 2.4 ± 1.0 pmol/mg at room temperature ($P < 0.05$, Fig 5C). Melittin did not significantly shift the binding affinity (K_D 8.3 ± 1.8 to 4.5 ± 0.9 nM at 4°C and 9.4 ± 2.8 to 4.0 ± 1.0 at room temperature, $P > 0.05$). Additionally, RTI-55 binding in the absence of melittin at 4°C was not significantly different from binding at room temperature, and binding in the presence of melittin at 4°C was not significantly different from RTI-55 binding in the presence of melittin at room temperature ($P > 0.05$). Thus, membrane fluidity does not appear to play a role in melittin's effect on the dopamine transporter.

Additional studies were conducted in these cells to determine whether melittin interacts with the dopamine D_2 receptor (Fig 5D). Saturation binding experiments were conducted using [3 H]spiperone concentrations of 0.0125 nM to 0.4 nM alone, or in the presence of 1.3 μM melittin. Melittin caused a significant reduction in the B_{\max} from 261.1 ± 33.5 fmol/mg to 158.5 ± 26.0 fmol/mg ($P < 0.05$). Melittin did not significantly shift the binding affinity (K_D 0.10 ± 0.03 to 0.11 ± 0.05 nM, $P > 0.05$). Thus melittin also acted like an apparent noncompetitive inhibitor of spiperone binding to the dopamine D_2 receptor, and so the effect of melittin on radioligand binding is not specific to the dopamine transporter.

3.5 Melittin increases dopamine transporter internalization

To determine whether the decrease in dopamine transporter function following melittin treatment was due to an increase in transporter internalization, biotinylation of cell surface proteins was performed. Extracellular proteins were labeled with a reducible form of biotin prior to treatments. Cell surface biotin remaining after treatment was cleaved. Thus biotin-labeled proteins internalized by treatment were protected from the reducing agent. Following treatment with melittin (1.3 μM) or vehicle for 30 min at 37°C melittin-induced internalization was significantly ($P < 0.01$) increased to $137.9 \pm 7.6\%$ of baseline levels (Fig 6). Interestingly, this effect was blocked by 15 min pretreatment with 1 μM cocaine ($97.8 \pm 6.2\%$ of control, $P < 0.05$ compared to melittin alone), while cocaine treatment alone did not alter dopamine transporter trafficking ($93.9 \pm 8.4\%$ of control). As a control, cells were biotinylated at 4°C, rinsed with cold glycine, and placed on ice while in DMEM. These cells demonstrate a non-permissive trafficking condition. Additionally, these cells demonstrate the removal of the cell surface biotin with the glutathione buffer. This condition reduced dopamine transporter internalization to $30.4 \pm 7.2\%$ of control and $34.3 \pm 16.5\%$ of control in the presence of melittin.

4. DISCUSSION

We generated a stable HEK cell line that expresses the recombinant human dopamine transporter and dopamine D_2 receptor. While dopamine transporter function and trafficking have been examined following phorbol ester stimulation of protein kinase C, few studies have explored the role of phospholipase A_2 pathways in dopamine transporter regulation. We used the phospholipase A_2 stimulator melittin throughout these experiments, and found that it not only stimulated the release of membrane-incorporated arachidonic acid, but also acted directly on the dopamine transporter. While these experiments are instrumental in the initial investigation of melittin's effects, further examination of the actions of melittin in an

appropriate physiological context, such as cultured dopamine neurons, will be beneficial to understanding melittin-induced phenomena.

Melittin incorporates into outer cell membranes and alters the characteristics of the membranes and of the proteins embedded within (for review see Bernheimer and Rudy, 1986). Based on structural information and the melittin-induced shift in the B_{\max} and Hill slope for [125 I]RTI-55 binding without a change in K_D (Fig 5B & C), it is likely that melittin did not bind directly to the [125 I]RTI-55 recognition site. Instead, it acted as an apparent non-competitive inhibitor, probably binding to a region of the transporter that alters the conformation of the dopamine transporter, making it unable to bind RTI-55. This designation of melittin as a noncompetitive inhibitor is in a phenomenological sense analogous to enzyme kinetics (Limbird, 2005; Tomlinson, 1988). We cannot exclude the possibility that an endogenous compound activated or released by melittin could be interacting with the dopamine transporter to decrease binding.

Alternatively, melittin's inhibition of [125 I]RTI-55 binding to the dopamine transporter could be due to nonspecific effects such as altered membrane structure. The effects of melittin on sarcoplasmic reticulum membrane fluidity are comparable to decreasing the temperature by 5°C, thus decreasing the rotational mobility of fluid lipids and reducing the rotational ability of integral Ca^{2+} ATPase (Mahaney et al., 1992). We observed that maximal binding of RTI-55 to the dopamine transporter, and the effects of melittin on RTI-55 binding, were not significantly different at 4°C as compared to room temperature (Fig 5C). Thus fluidity of the membrane does not alter the binding of the direct blocker RTI-55 to the dopamine transporter, and melittin's effect on RTI-55 binding is probably not due to disruption of membrane structure.

In addition to decreasing ligand binding to the dopamine transporter, melittin was an inhibitor of [^3H]spiperone binding to the dopamine D_2 receptor (Fig 5D), suggesting that melittin may interact more generally with membrane-incorporated proteins. Thus melittin appears to act in a previously unreported pharmacological manner, which may serve as a confounding variable for experiments that use melittin to stimulate phospholipase A_2 . Furthermore, melittin binds to a variety of cellular proteins including calmodulin (Maulet and Cox, 1983), bovine mitochondrial F1-ATPase (Gledhill and Walker, 2005) and to the ubiquitous transcription factor NF- κ B (Park et al., 2004). While melittin has been used as a tool to examine the peptide binding properties of the previously mentioned proteins, this is the first examination of melittin's effects on trafficking and antagonist binding to cell surface membrane-incorporated proteins.

Melittin treatment caused a strong reduction in [^3H]dopamine uptake that occurred rapidly, within 5–10 min of treatment (Table 1) and biotinylation experiments revealed an increased internalization that contributes to this effect. Melittin induces pore formation in erythrocytes (reviewed in Raghuraman and Chattopadhyay, 2007) and inhibits NaKATPase activity (Cuppoletti and Abbott, 1990), activities that could be factors in decreased transport. Additional experiments, such as electrophysiological recordings, could be done to determine whether melittin affects Na^+ or Cl^- gradients across cellular membranes.

As discussed above, the observed decrease in cellular [^3H]dopamine uptake *via* the dopamine transporter following melittin treatment may be due to the multiple effects of melittin on the cells, including transporter internalization, inhibition of the binding of dopamine, and possible pore formation. These mechanisms may explain the disparities between the effects of direct application of arachidonic acid and of melittin (Zhang and Reith, 1996). Studies using rotating disk electrode voltammetry demonstrate an arachidonic acid-induced decrease in dopamine uptake. Additionally the *cis*-unsaturated fatty acids oleic

and linoleic acid inhibit dopamine transporter-mediated uptake, although to a lesser degree (Chen et al., 2003), which suggests that there may be a property common to several fatty acids that alters dopamine transporter function. Consistent with these results, the other fatty acids released by melittin (Fig 3) could have contributed to the decrease in dopamine transporter-mediated uptake.

While no studies have specifically explored the effects of arachidonic acid on dopamine transporter trafficking, C6 cells expressing the dopamine transporter demonstrate a decrease in B_{\max} with no change in K_D following exposure to arachidonic acid in attached, whole cell preparations (Zhang and Reith, 1996), suggesting transporter internalization. It is possible that direct actions of arachidonic acid on the dopamine transporter initiate trafficking pathways. Arachidonic acid ethyl ester, a non-metabolizable arachidonic acid analogue, incorporates into the cell membrane, but does not mimic the effects of arachidonic acid on dopamine transporter function (Chen et al., 2003; Ingram and Amara, 2000). This result suggests that arachidonic acid acts from within the lipid phase of the membrane, not by simply altering the lipid microenvironment surrounding membrane-spanning proteins and may involve a specific fatty acid binding domain on the dopamine transporter. Further studies will need to be performed to distinguish between effects mediated by direct actions of melittin and indirect effects of melittin-induced fatty acid release.

Melittin stimulates phospholipase A_2 , phospholipase C and phospholipase D activity in L1210 cells. The effects of melittin on fatty acid release are completely blocked only with simultaneous pretreatment with inhibitors of all three phospholipases, with phospholipase A_2 activity accounting for about 10% (Lee et al., 2001). We found that melittin stimulated the release of arachidonic acid from cellular membranes (Fig 3 & 4), but this effect was not blocked by treatment with inhibitors of phospholipase C, phospholipase D, or phospholipase A_2 , although we did not test the effect of simultaneous treatment with the three inhibitors. Thus, melittin did not act as a specific pharmacological stimulator of phospholipases. This effect may be particular to the chosen cell type, as phospholipase A_2 inhibitors successfully block melittin-induced release of arachidonic acid in rabbit proximal tubule cells (Han et al., 2002), but quinacrine is unable to inhibit melittin-induced release of arachidonic acid in rat Leydig cells (Ronco et al., 2002). Further, it is suggested that melittin has no inherent phospholipase activity, and that fatty acids are released subsequent to the membrane perturbations following melittin incorporation and pore formation (Raghuraman and Chattopadhyay, 2007). At the very least, while some effects of melittin on cellular activity may be mediated through the arachidonic acid signaling pathway, they may not be directly or exclusively produced by phospholipase A_2 . If melittin is found to directly interact with the transporter, then any effects of melittin-liberated fatty acids on the transporter may be masked by this interaction, and thus prove difficult to distinguish.

Since phospholipase inhibitors did not block melittin-stimulated arachidonic acid release, there may be some aspect of the peptide that interacts with the membrane to cause release of fatty acids. The 14 amino acid peptide mastoparan, found in wasp venom, is used to stimulate GTP binding regulatory proteins indirectly (Higashijima et al., 1988). Mastoparan disrupts the membrane lipid rafts where many receptors and second messenger proteins are localized, thus releasing these proteins, lipids, and arachidonic acid to the cytoplasm (Sugama et al., 2005; Nakamura et al., 2004). Experiments using nuclear magnetic resonance measurements demonstrate an interaction between melittin and ganglioside components of lipid rafts (Chatterjee and Mukhopadhyay, 2002). Thus, it is possible that melittin stimulates fatty acid release and activates signaling pathways by direct interactions with the membrane.

Cocaine (1 μM , 45 min) pretreatment had no effect by itself but prevented melittin-induced internalization (Fig 6), which may be due to a steric hindrance of the dopamine transporter by cocaine that prevented melittin's interaction with the transporter. In contrast, cocaine (10 μM for 10 min and 1 μM for 24 h) causes an increase in dopamine transporter surface expression in human dopamine transporter-expressing HEK and N2A cells, respectively (Daws et al., 2002; Little et al., 2002), differences that may be due to drug concentration or cells used. Regardless, melittin does not affect dopamine transporter surface expression the way that cocaine does. It is not known whether cocaine also blocks the effects of arachidonic acid applied to the dopamine transporter. In addition, melittin is likely operating through a different mechanism than phorbol esters, since cocaine treatment does not alter phorbol ester-induced dopamine transporter internalization (Daniels and Amara, 1999).

Interestingly, phospholipase A₂ activity may play a role in psychostimulant sensitization. Direct injections of melittin into rat ventral tegmental area cause sensitization, seen as increased locomotor activity, stereotypy, and dopamine release in the nucleus accumbens, to subsequent cocaine administration. This sensitization is blocked by pretreatment with the phospholipase A₂ inhibitor quinacrine (Reid et al., 1996). Thus, the influence of phospholipase activity and arachidonic acid signaling on dopamine system functioning may be important for understanding stimulant addiction.

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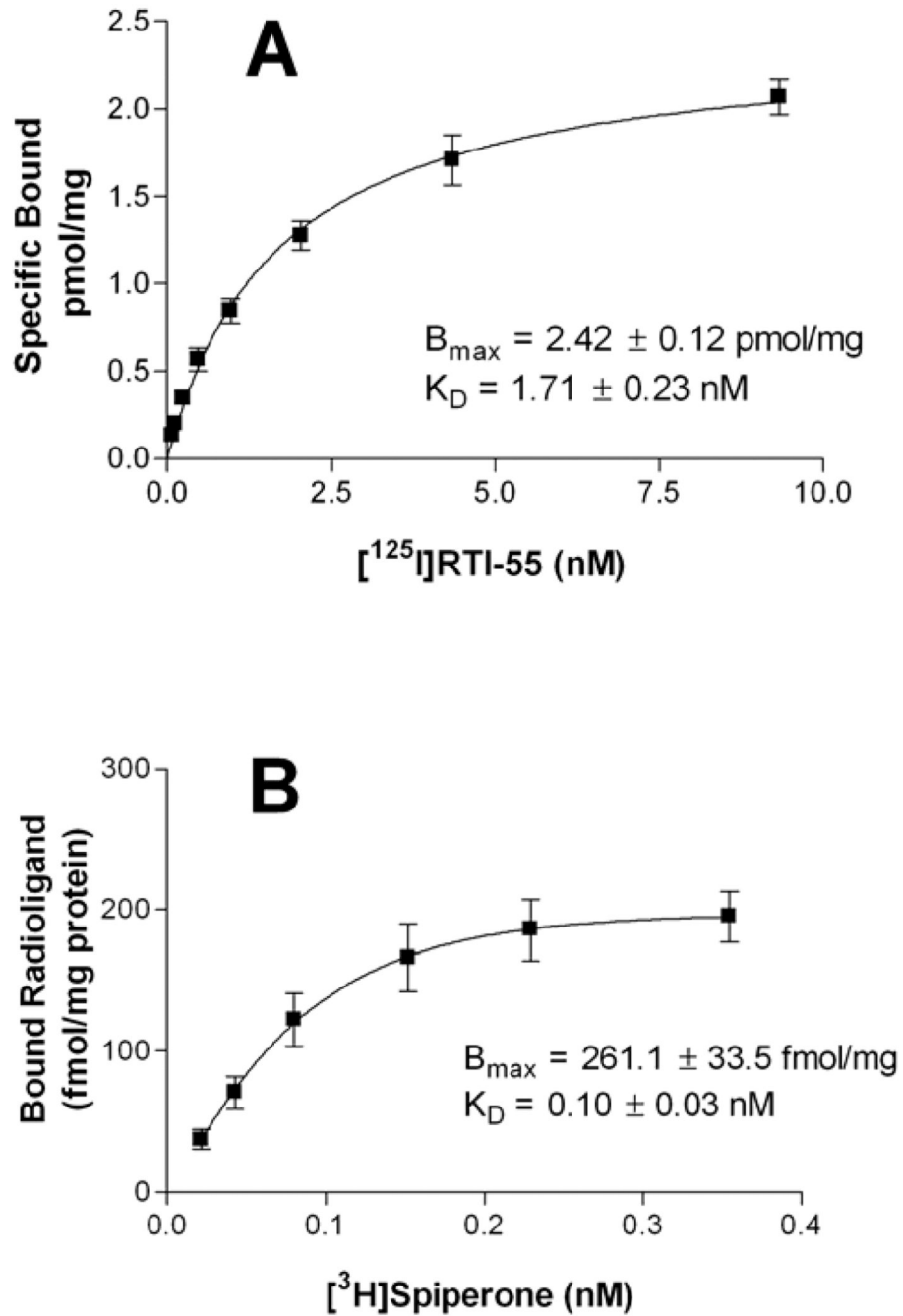


Fig 1. Characterization of stable transfections in HEK cells

Saturation binding experiments were performed as described in the text. $[^{125}\text{I}]\text{RTI-55}$ binding to the dopamine transporter, 1A, is a composite of 6 independent experiments and, $[^3\text{H}]\text{spiperone}$ binding to the D_2 receptor, 1B, is a composite of 5 independent experiments each conducted with duplicate determinations.

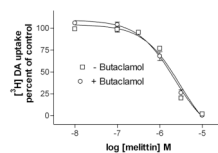


Fig 2. Melittin causes a dose dependent decrease in dopamine transporter-mediated uptake
HEK-D₂-hDAT cells were treated with varying doses of melittin for 30 min, rinsed, and the uptake assay was initiated by the addition of 50 nM [³H]dopamine at 37°C. Experiments to block the D₂ receptor were conducted in parallel using identical melittin treatments followed by the inclusion of butaclamol (100 nM) 10 min before the addition of [³H]dopamine. Results are presented as a percent of vehicle treated control cells. The curves are mean ± S.E.M. of three independent experiments, each conducted with triplicate determinations.

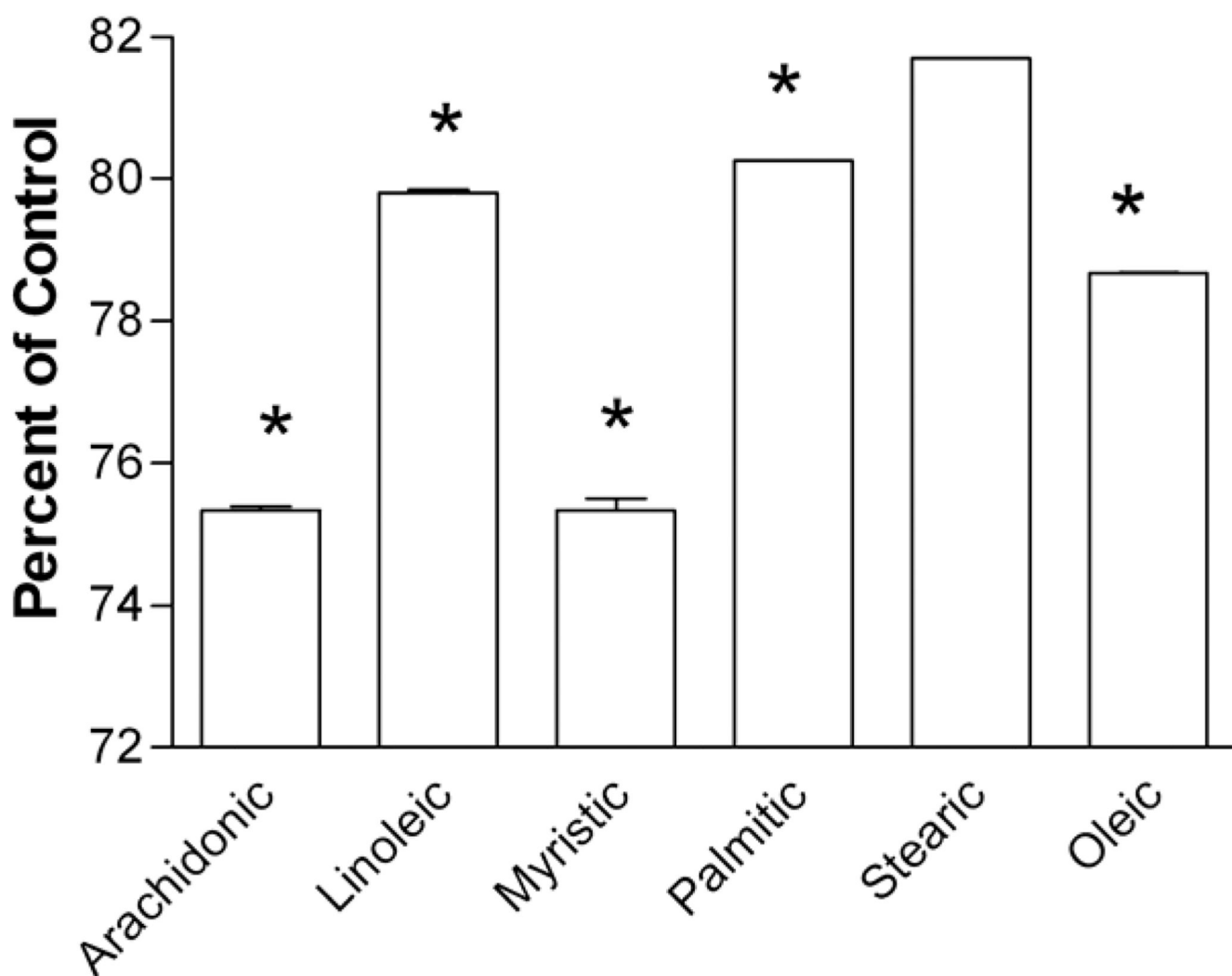


Fig 3. Melittin releases arachidonic acid from cell membranes

Cells were treated with melittin (1.3 μM) for 30 min and fatty acids remaining in the membrane after treatment were isolated for GLC analysis as described in the text. Levels of fatty acids remaining after treatment are presented as a percent of vehicle treated controls. Data are mean ± S.E.M. of three independent experiments (***, $P < 0.001$ by Student's *t* test).

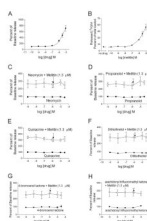


Fig 4. Melittin-stimulated release of arachidonic acid is through non-phospholipase-mediated mechanisms

HEK-D₂-hDAT cells were incubated with [³H]arachidonic acid for 18–20 h. Stimulated [³H]arachidonic acid release was analyzed as described in the text. Cells were treated with melittin for 30 min and the percent of baseline release (A) and the percent of total cell incorporated [³H]arachidonic acid released (B) were quantified. To determine the ability of melittin to stimulate phospholipases, cells were treated with neomycin (C), propranolol (D), quinacrine (E), dithiothreitol (F), 4-bromo-enol lactone (G), or arachidonyl trifluoromethyl ketone (H) for 60 min either individually (closed squares) or with the addition of a single dose of melittin (1.3 μM, open squares) during the final 30 min of the treatment. The single closed circle in graphs C–H represents the effect of melittin alone (1.3 μM). Released [³H]arachidonic acid is presented as a percent of untreated baseline release levels. Data are the mean ± S.E.M. of three experiments, each conducted with triplicate determinations.

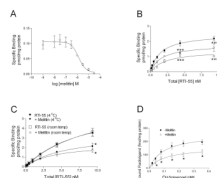


Fig 5. Melittin inhibits binding of RTI-55 to the dopamine transporter

To generate dose-response binding curves for melittin (A) HEK- D₂-hDAT cells were treated with varying concentrations of melittin in the presence of 40 pM [¹²⁵I] RTI-55 as described in the text. Equilibrium saturation binding experiments (B) were also conducted. The graph depicts [¹²⁵I]RTI-55 binding in the absence of melittin (closed squares), and [¹²⁵I] RTI-55 binding in the presence of 1.3µM (open circles) or 10µM (open squares) melittin. Data are the mean ± S.E.M. of 4–6 independent experiments (***,P<0.001 by two-way ANOVA followed by Bonferroni post test). The role of temperature in melittin's effects on RTI-55 equilibrium saturation binding is shown in panel (C). The curves are averages of 4 independent experiments. [¹²⁵I] RTI-55 binding at 4°C (closed square) and at room temperature (open square), and [¹²⁵I] RTI-55 binding in the presence of melittin (1.3µM) at 4°C (closed circle) and room temperature (open circle) are depicted (*,P<0.05, by two-way ANOVA followed by Bonferroni post-test comparing treated to vehicle treated samples). The effect of melittin on [³H]spiperone binding to the D₂ receptor was performed as described in the text (D) (*, P<0.05, ANOVA followed by Student's t test)

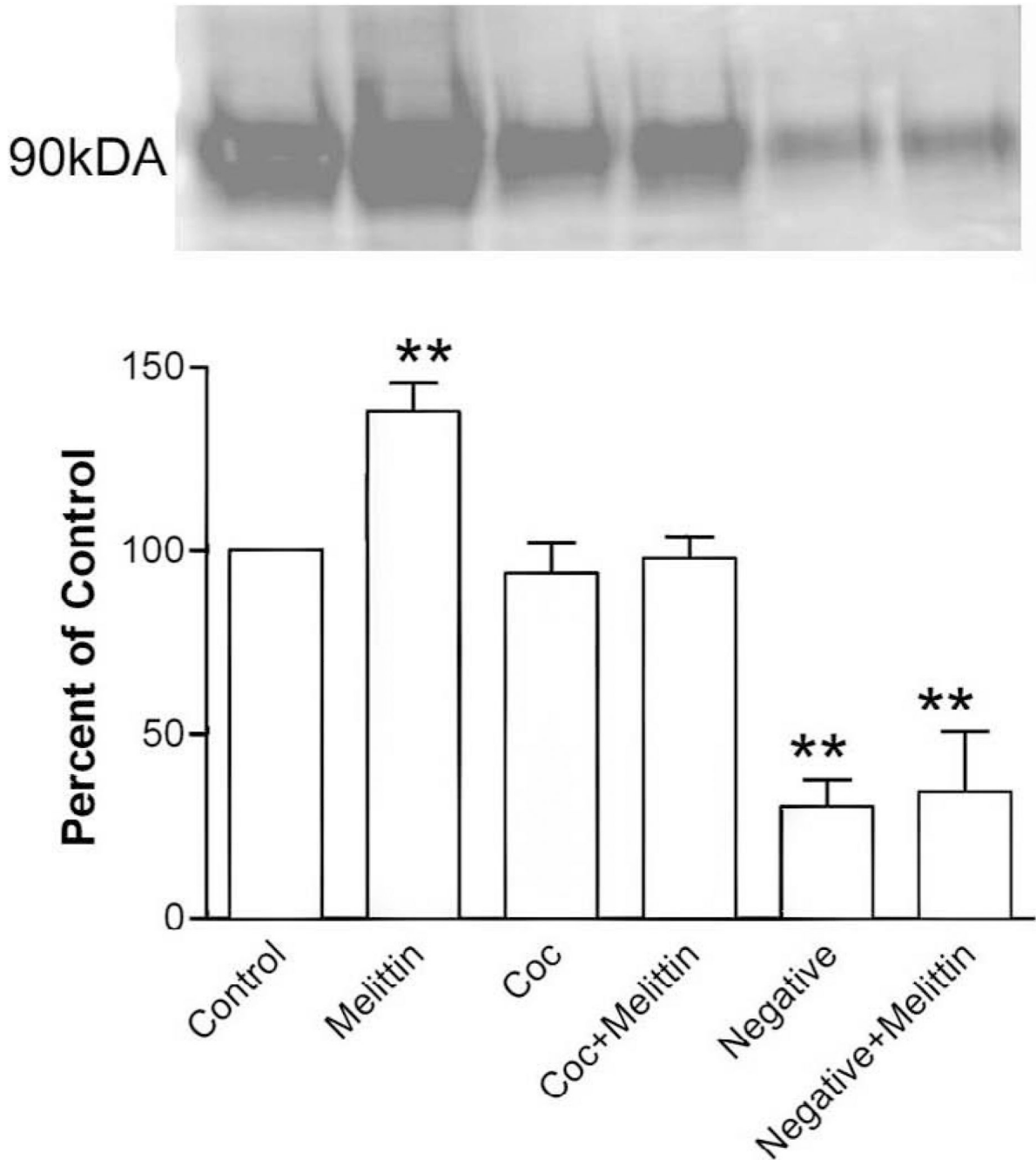


Fig 6. Melittin increases dopamine transporter internalization

HEK- D₂-hDAT cells were biotinylated and then treated with vehicle or melittin (1.3 μ M) for 30 min, cocaine (1 μ M) for 45 min, or cocaine (1 μ M) 15 min prior and during 30 min melittin (1.3 μ M) treatment. Untreated, biotinylated negative control cells remained at 4°C for 30 min (cold). Following treatment the cells were rapidly chilled and the media removed. The remaining cell surface biotin was stripped and the internalized dopamine transporter was isolated by avidin purification and separated by SDS-PAGE. The immunoblot was detected with anti-dopamine transporter antibody. The representative blot shows biotinylated, internalized product for control and treated cells. Total biotinylated sample is also shown. Band optical density was quantified and expressed as a percent of vehicle

treated control. Graph: dopamine transporter internalization expressed as means \pm S.E.M. from 3–6 experiments (*, $P < 0.05$; **, $P < 0.01$, one way ANOVA followed by Tukey's posthoc test).

Table 1

Potency of melittin on inhibition of dopamine transporter function

Treatment Time	[³ H]Dopamine uptake Melittin IC ₅₀ (μM)
5 min	4.2 ± 0.1
10 min	0.82 ± 0.01
15 min	1.76 ± 0.1
30 min	1.3 ± 0.14
60 min	0.8 ± 0.1
120 min	1.79 ± 0.12

Cells were treated with melittin for the times indicated, followed by the addition of 50 nM [³H]dopamine. Uptake continued for 10 min. Data represent mean ± S.E.M. of three independent experiments, each conducted with triplicate determinations. The IC₅₀ value for melittin at 5 min is significantly different than all other times (P<0.001, one-way ANOVA and Tukey's Multiple Comparison Test), but IC₅₀ values for melittin treatment times from 10–120 min do not differ significantly (P>0.05, one-way ANOVA).

Table 2

Effects of melittin on RTI-55 saturation binding.

Treatment	[¹²⁵ I]RTI-55 B _{max} (pmol/mg)	[¹²⁵ I]RTI-55 K _D (nM)
None	2.49 ± 0.14	1.48 ± 0.23
Melittin 1.3 μM	1.78 ± 0.13 ^a	1.62 ± 0.33
Melittin 10 μM	1.42 ± 0.16 ^a	2.38 ± 0.7

Homogenized cell membranes were incubated for 90 min with 40 pM [¹²⁵I]RTI-55, varying concentrations of non-radiolabeled ligand, and melittin as indicated. Nonspecific binding was determined in the presence of 10 μM mazindol. The assay was terminated by filtration. Data are mean ± S.E.M. from at least 3 experiments.

^aP<0.05, one-way ANOVA followed by Dunnett's Multiple Comparison Test.