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Sortilin binding compounds, formulations, and uses thereof

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(54) **SORTILIN BINDING COMPOUNDS, FORMULATIONS, AND USES THEREOF**

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A61K 31/40 (2006.01)
A61K 31/167 (2006.01)
A61K 31/381 (2006.01)
C07D 333/46 (2006.01)
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- (52) **U.S. Cl.**
CPC **A61K 31/40** (2013.01); **A61K 31/167** (2013.01); **A61K 31/381** (2013.01); **C07D 207/09** (2013.01); **C07D 333/46** (2013.01)
- (58) **Field of Classification Search**
CPC **A61K 31/40**; **A61K 31/167**; **A61K 31/381**
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See application file for complete search history.

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(57) **ABSTRACT**

Provided herein are compounds and formulations thereof that can be capable of binding to sortilin or an analog thereof. Also provided herein are methods of administering the compounds and formulation provided herein to a subject in need thereof.

17 Claims, 14 Drawing Sheets

Specification includes a Sequence Listing.

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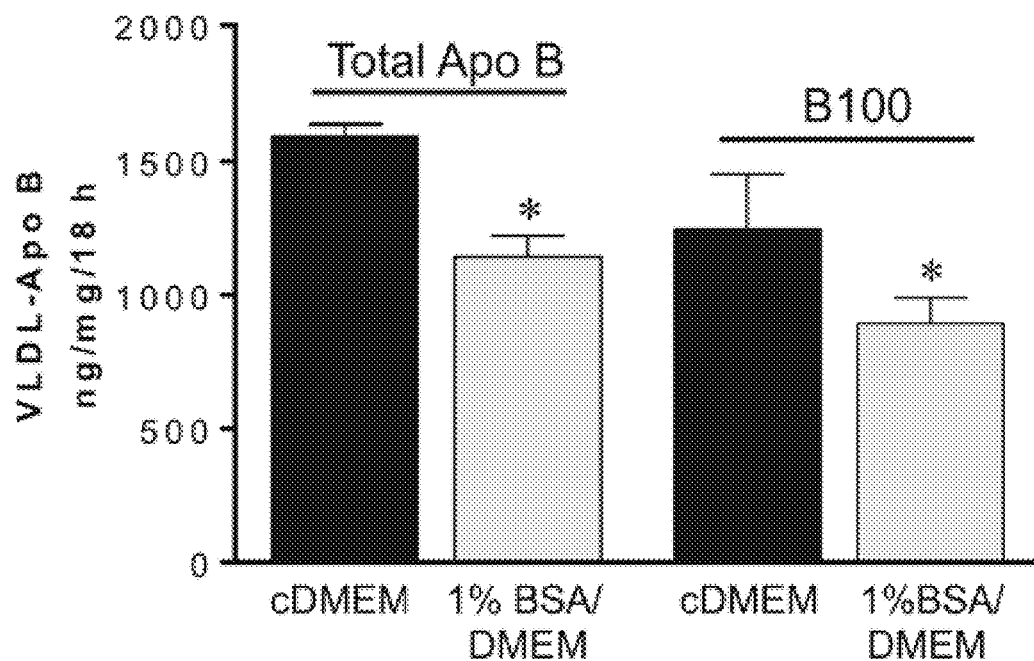


FIG. 1A

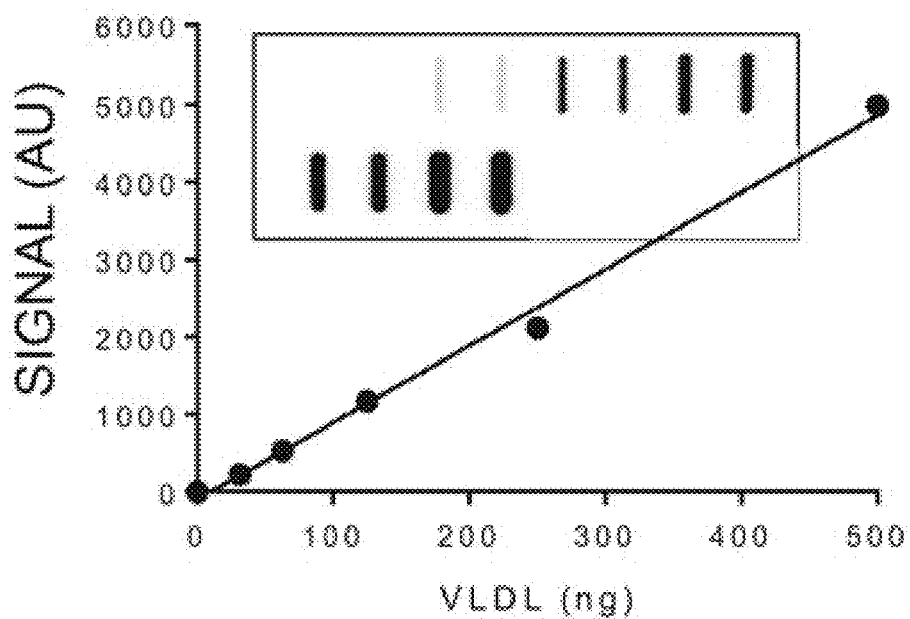


FIG. 1B

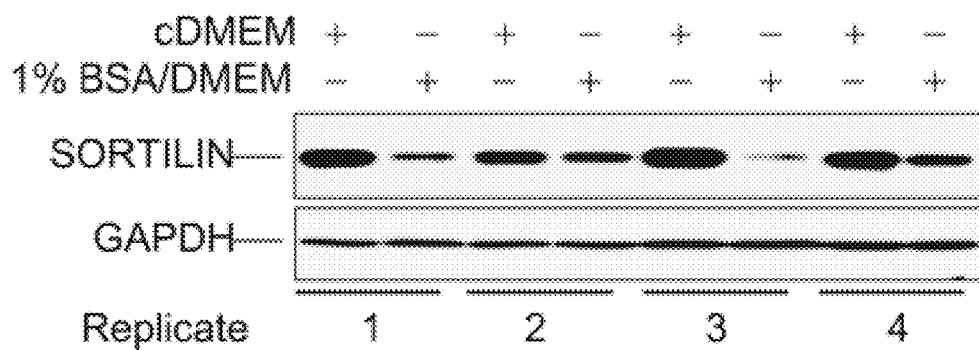


FIG. 1C

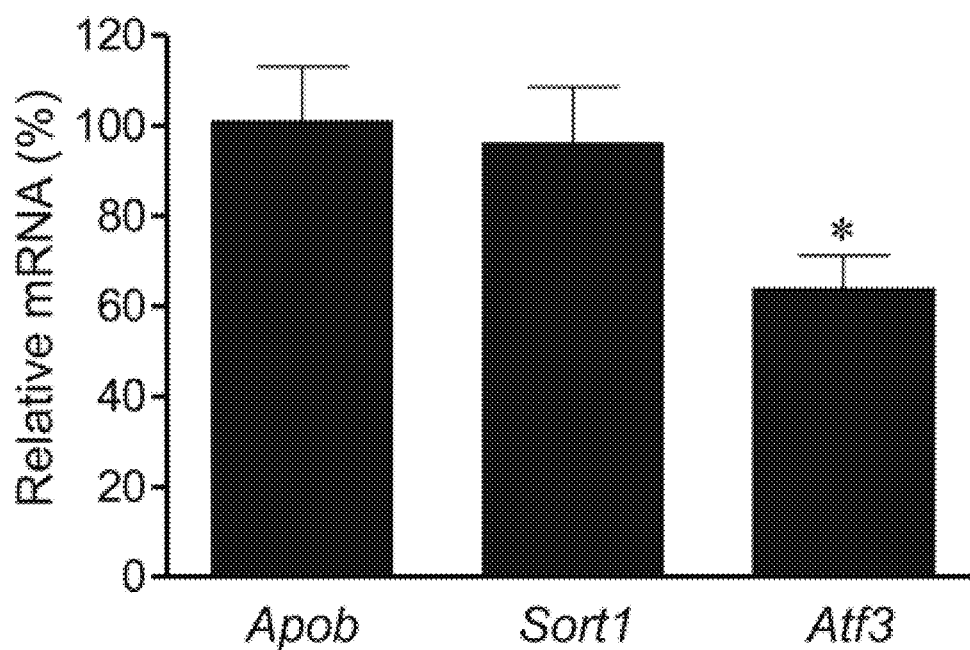


FIG. 1D

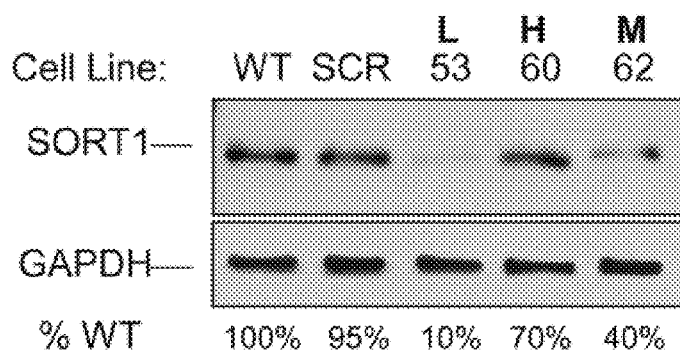


FIG. 2A

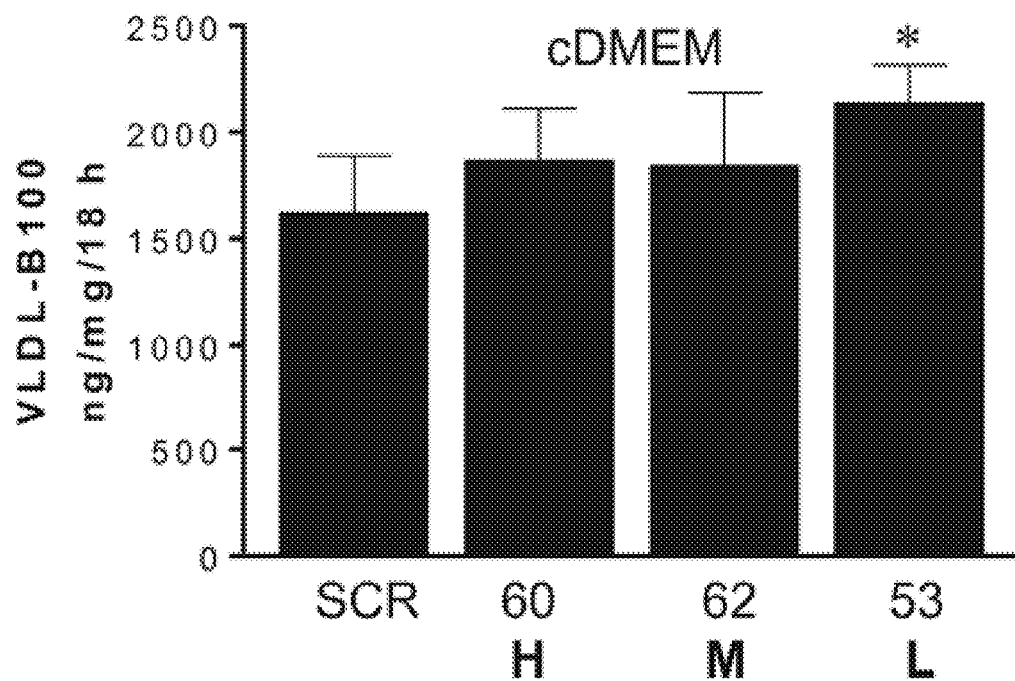


FIG. 2B

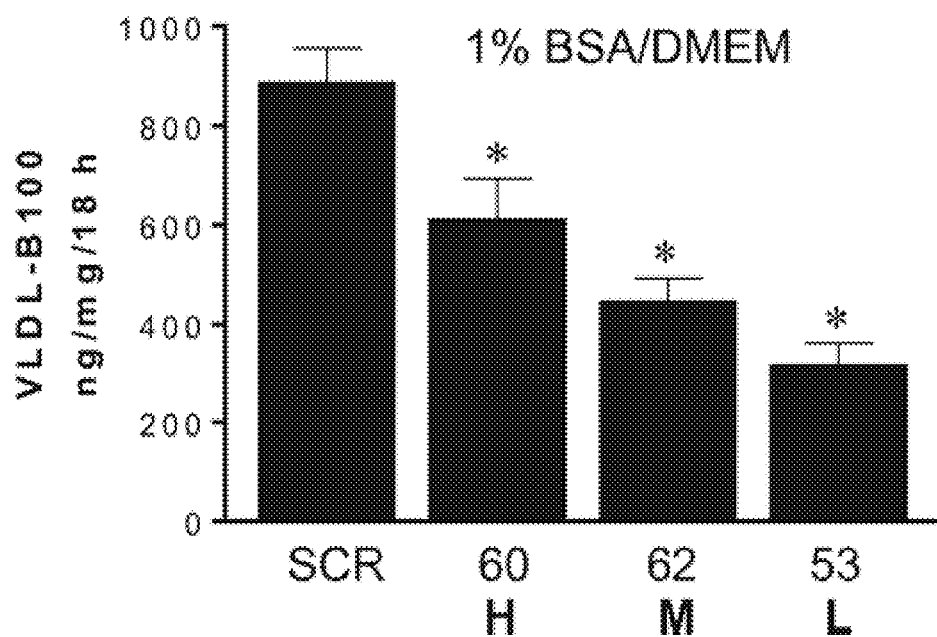


FIG. 2C

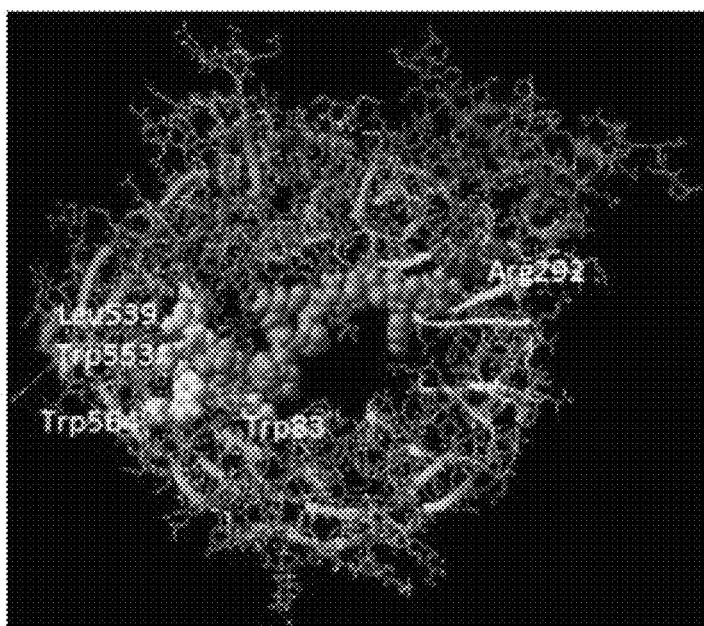


FIG. 3A

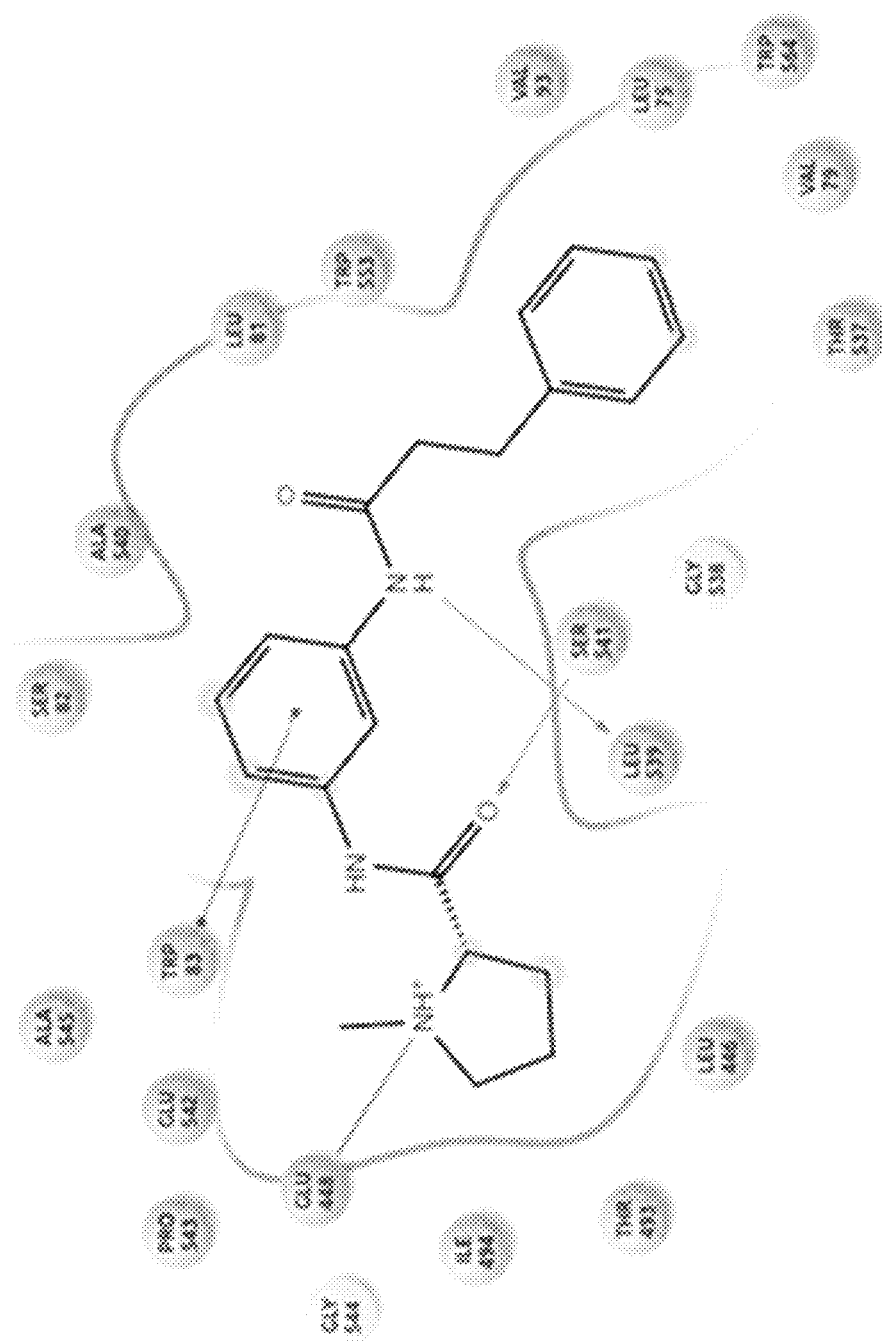


FIG. 3B

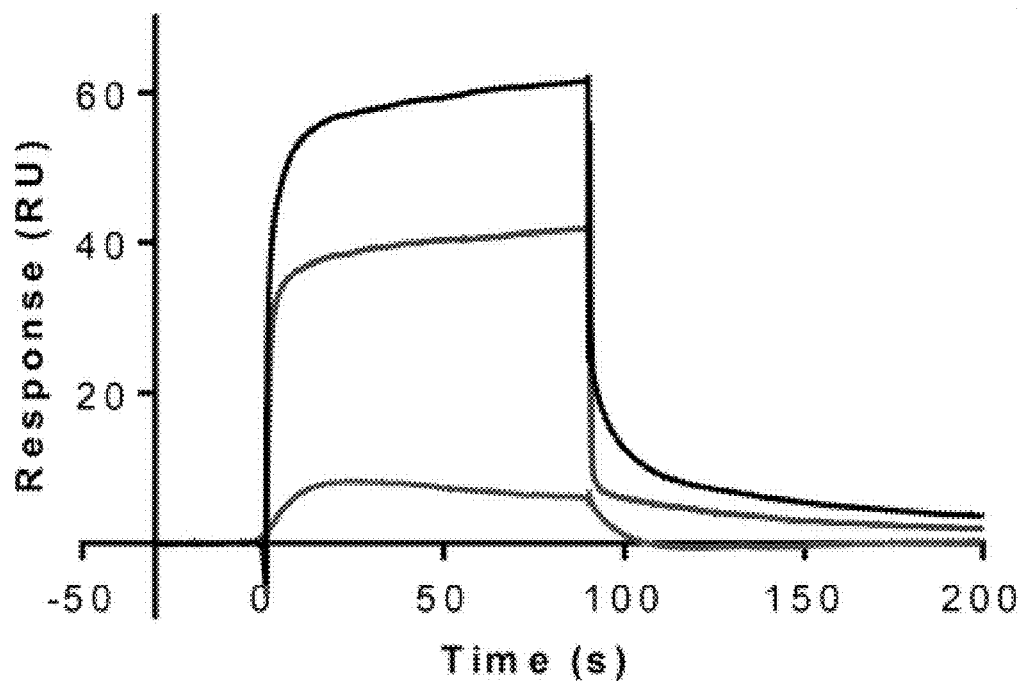


FIG. 3C

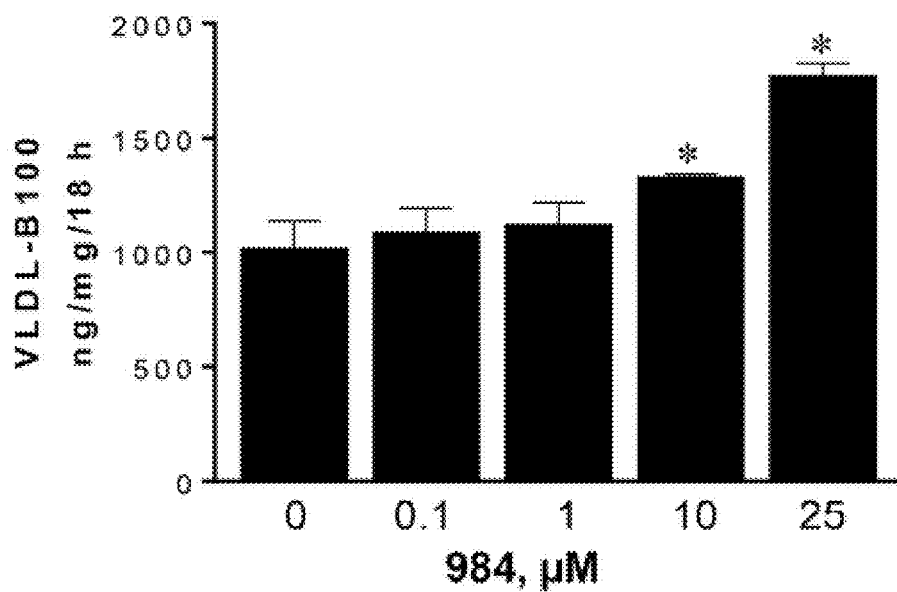


FIG. 3D

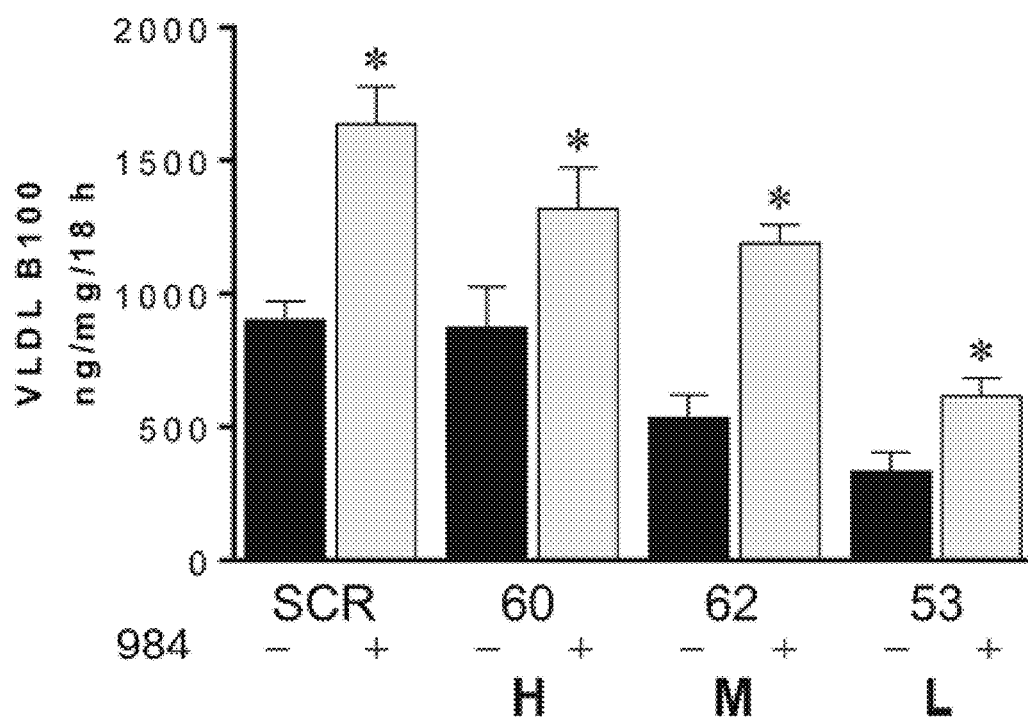


FIG. 4A

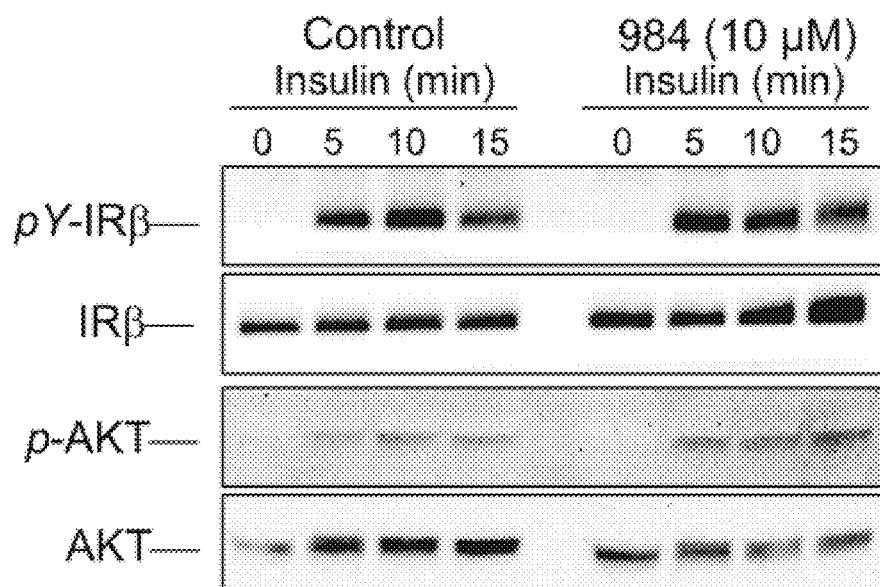


FIG. 4B

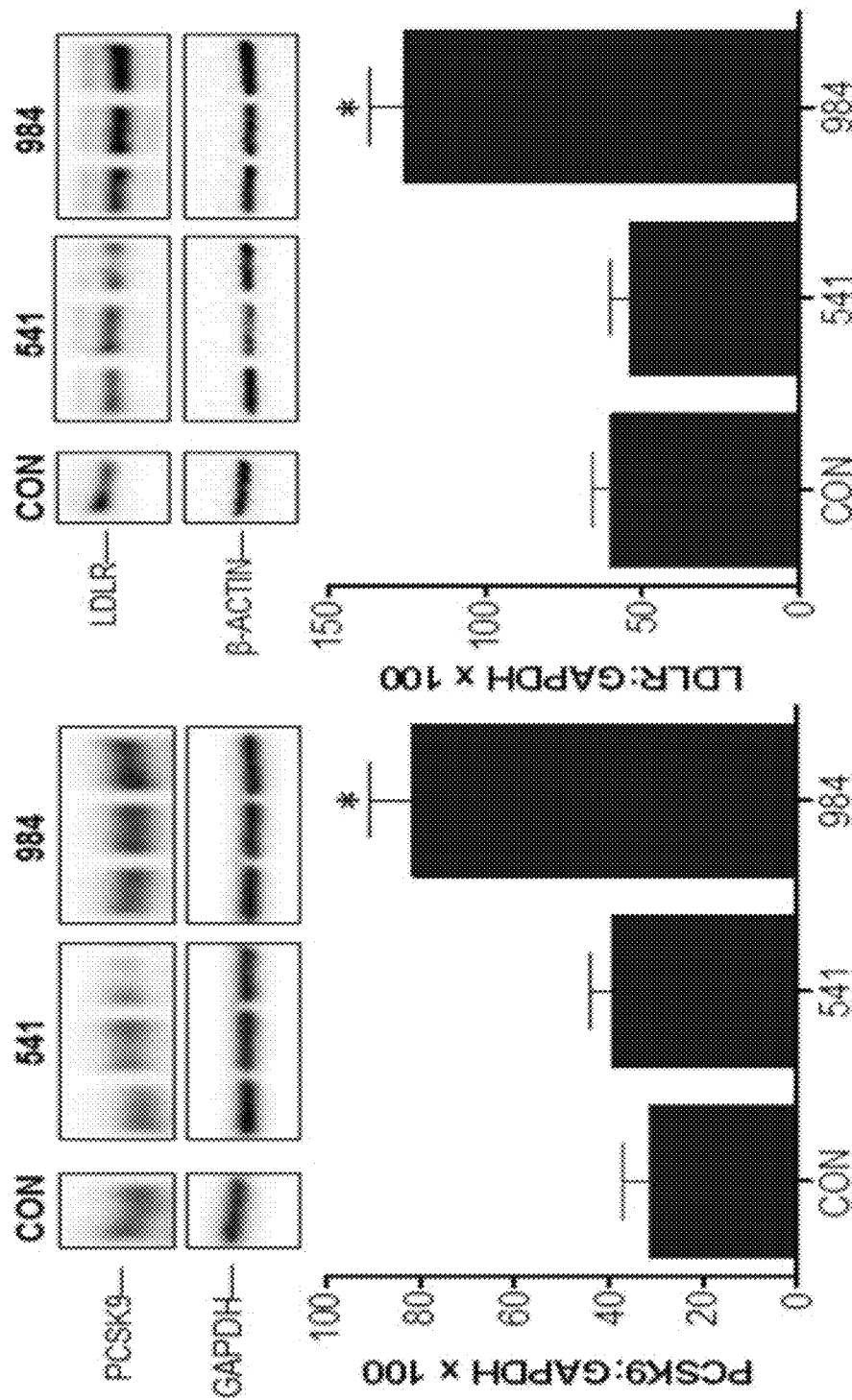


FIG. 5B

FIG. 5A

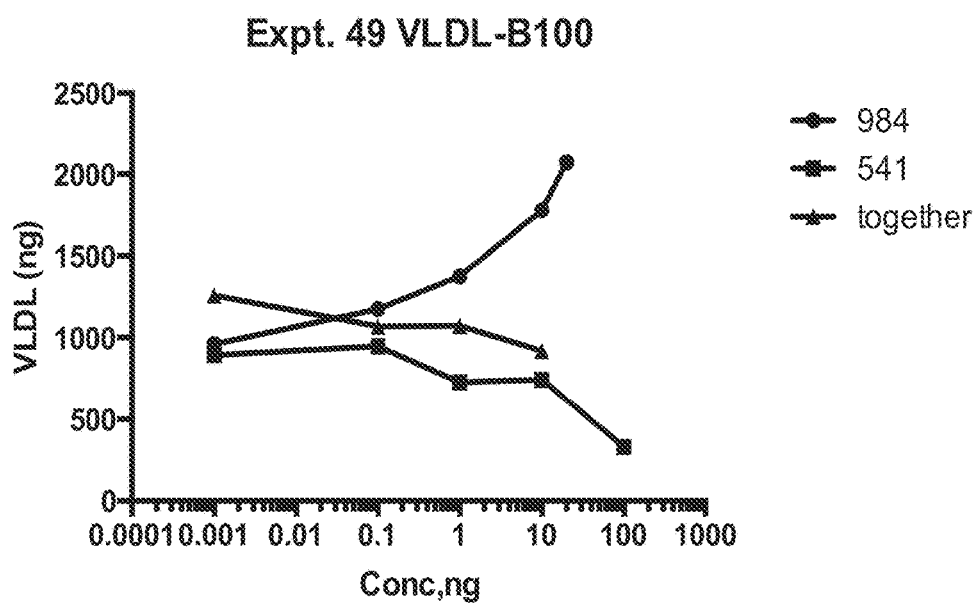


FIG. 6

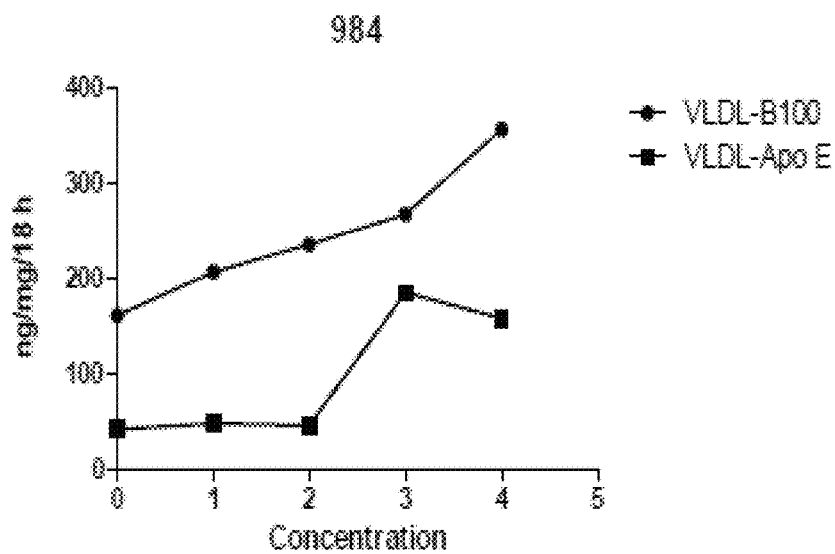


FIG. 7A

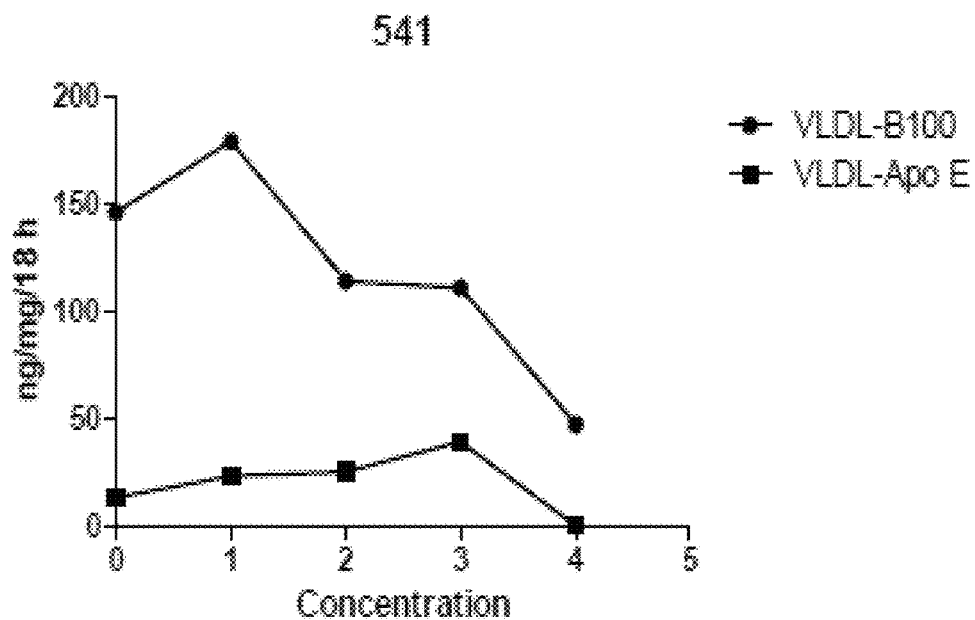


FIG. 7B

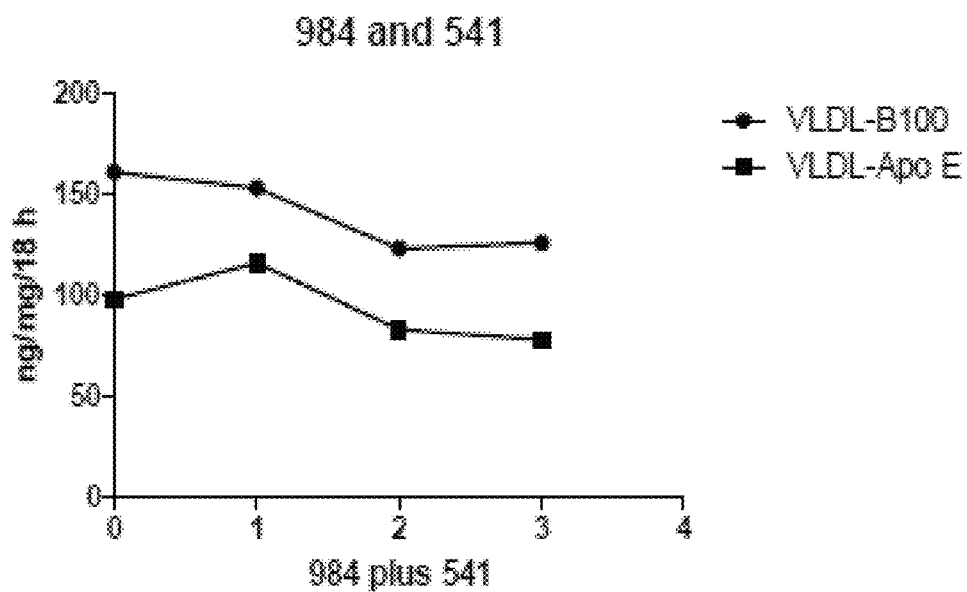


FIG. 7C

541 n=2 MST Buffer

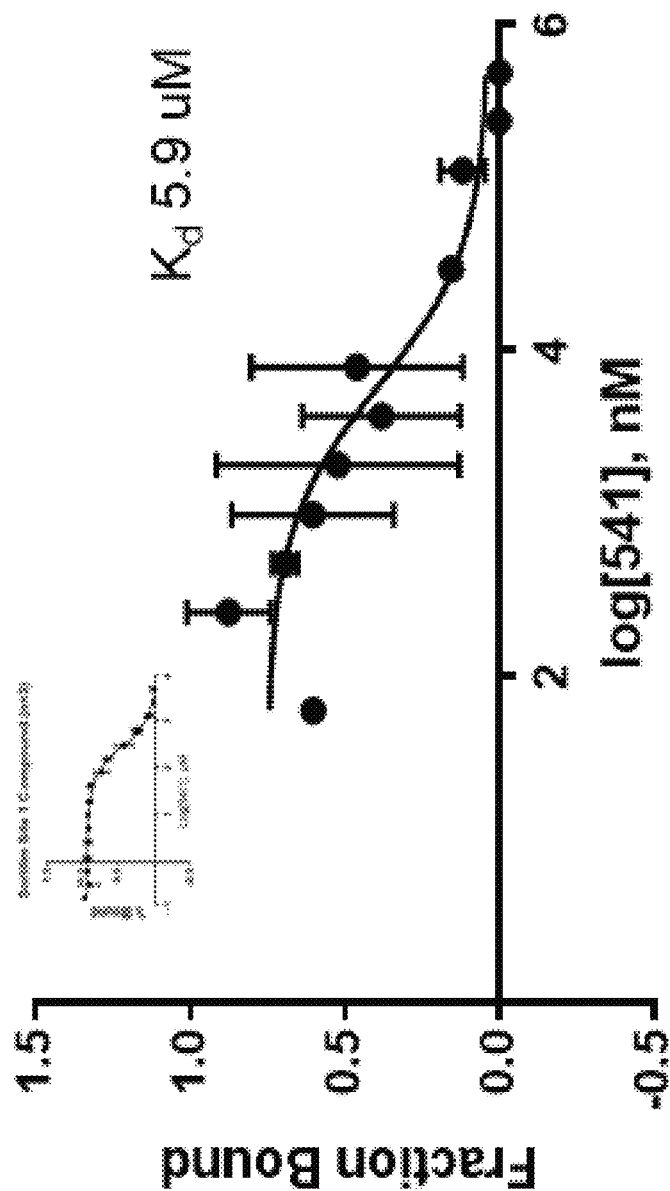


FIG. 8

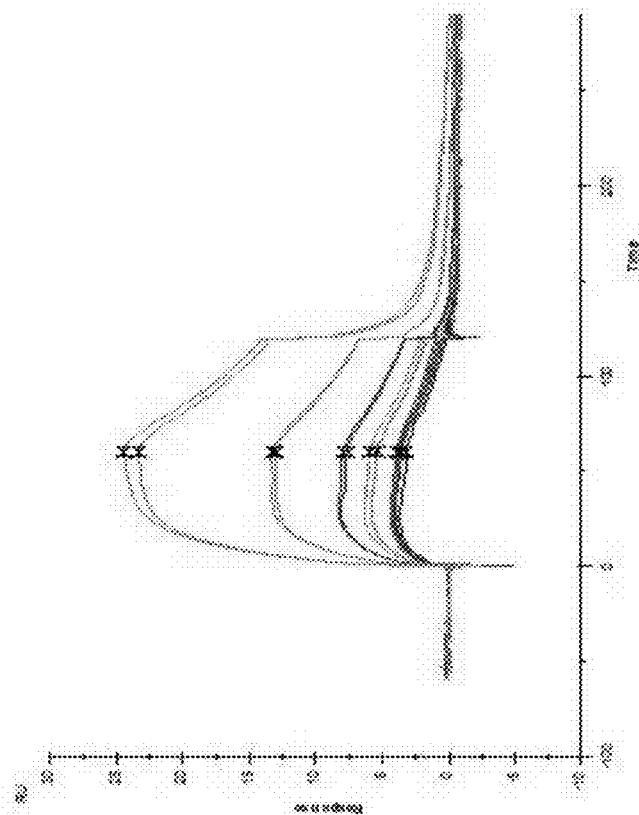
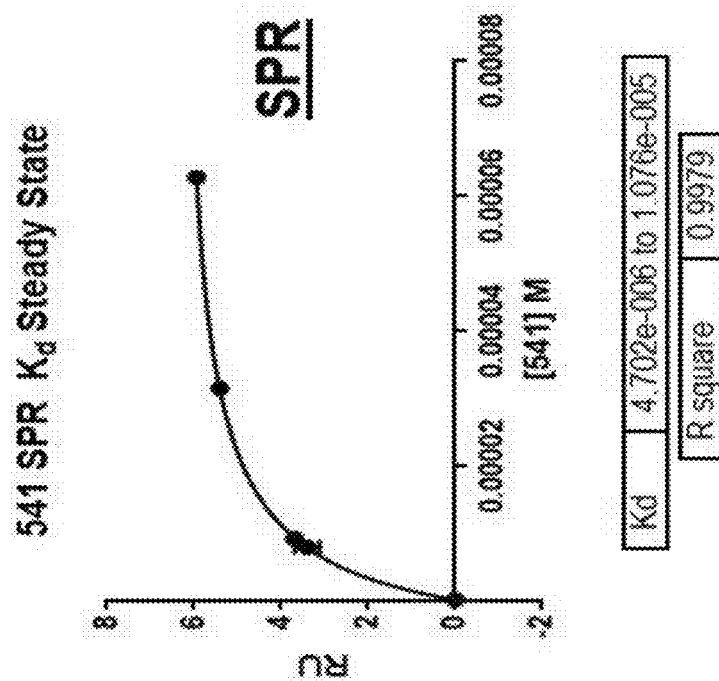


FIG. 9A

FIG. 9B

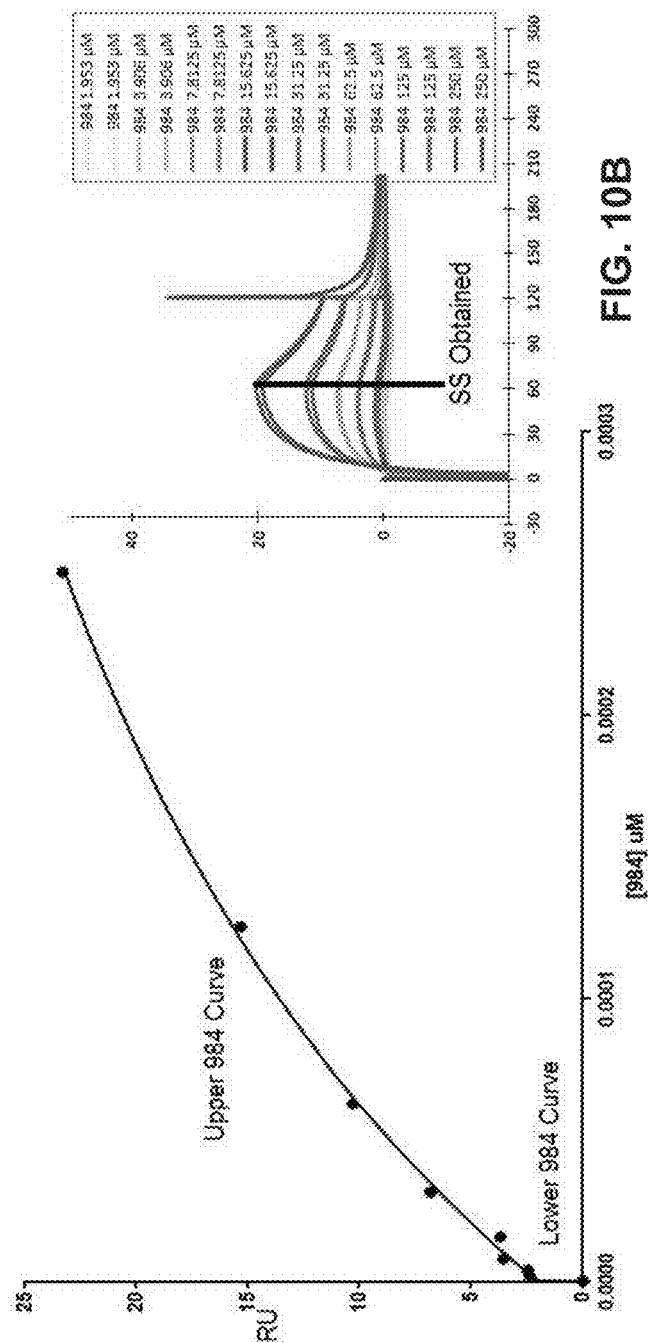


FIG. 10B

KdHi	2.602e-008	KdHi	0.0 to 1.813e-006
KdLo	0.0003068	KdLo	0.0001299 to 0.0004836

Percent Hi Affinity	0.7863
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R square	0.9980
Absolute Sum of Squares	0.9120
Sy x	0.4271

FIG. 10A

FIG. 11A

Cpd984 High Range

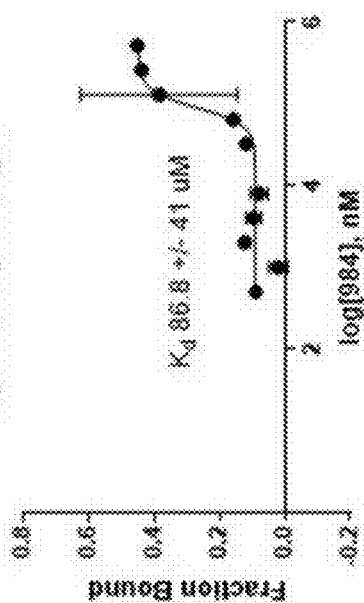
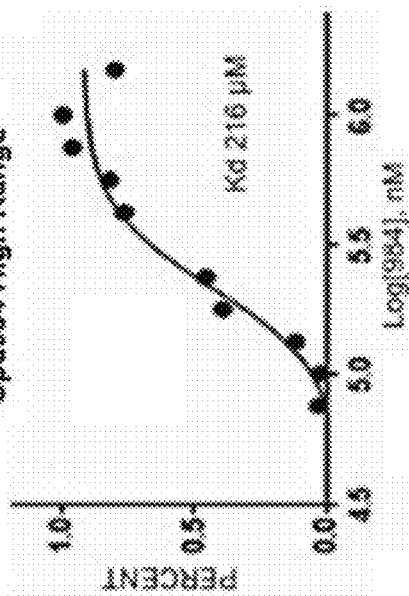


FIG. 11B

Cpd984 High Range



Cpd984 Low Range

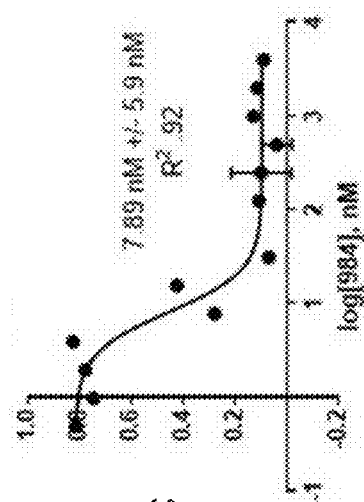


FIG. 11C

1

**SORTILIN BINDING COMPOUNDS,
FORMULATIONS, AND USES THEREOF****CROSS-REFERENCE TO RELATED
APPLICATIONS**

This application claims the benefit of and priority to U.S. Provisional Patent Application No. 62/294,412, filed on Feb. 12, 2016, entitled "LOWERING LDL CHOLESTEROL AND ASSOCIATED CARDIOVASCULAR DISEASE RISK BY INCREASING HEPATIC LDL RECEPTORS BY BLOCKING SECRETION OF PCSK9 THROUGH INTERFERENCE BY SMALL MOLECULES WITH PCSK9 INTERACTION WITH SORTILIN," the contents of which is incorporated by reference herein in its entirety.

**STATEMENT REGARDING FEDERALLY
SPONSORED RESEARCH OR DEVELOPMENT**

This invention was made with government support under grants number DK100163-01A1 and GM101132 awarded by the National Institutes of Health. The government has certain rights in the invention.

SEQUENCE LISTING

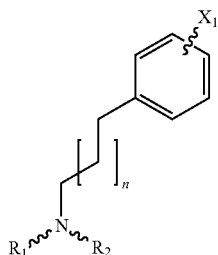
This application contains a sequence listing filed in electronic form as an ASCII.txt file entitled 292105-1110_ST25.txt, created on Feb. 13, 2017. The content of the sequence listing is incorporated herein in its entirety.

BACKGROUND

Proprotein convertase subtilisin/kexin type 9 (PCSK9) is an enzyme that is ubiquitously expressed in many tissues and binds to the receptor of low-density lipoprotein (LDL) particles. The LDL receptor (LDLR) binds and transports LDL-particles into cells and thus can reduce the concentration of LDL in circulation. While the LDLR has been the focus of much research and the target of many therapeutics aimed at treating high cholesterol, there still exists a need for improved and alternative treatments.

SUMMARY

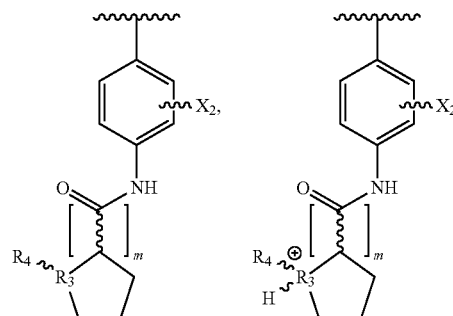
Provided herein are compounds that can have a structure according to Formula 2



where n can be 1-10, where X_1 can be selected from the group of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of

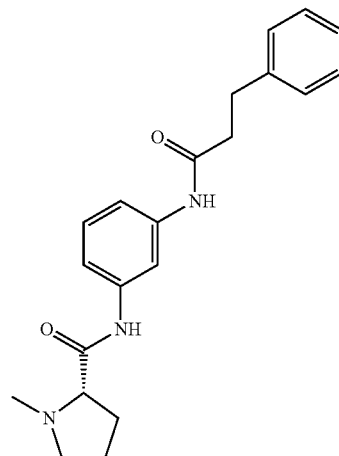
2

Formula 2, an acyl, an amino a sulfono, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof, where R_1 can be a carboxylic acid or an ester where R_2 can be a 5, 6, or 7 member heterocyclic ring that can be optionally further substituted with a suitable substituent, a sulfonium, a quaternary ammonium,



where X_2 can be selected from the group of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of Formula 2, an acyl, an amino a sulfono, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof, where R_3 can be N or S, where R_4 can be H, an C_1 - C_{10} alkyl, an amino, a carboxylic acid, an ester, an amino acid or positively charged derivative thereof, and where m represents the number of carbon atoms in an alkyl and can range from 0-10. The compound(s) can bind Site 2 on sortilin or analogue thereof. In some embodiments, the compound can bind to Site 2 on sortilin or analogue thereof and can increase binding of a compound to site 1 of sortilin or an analogue thereof. In some embodiments the compound can have a structure according to Formula 2A

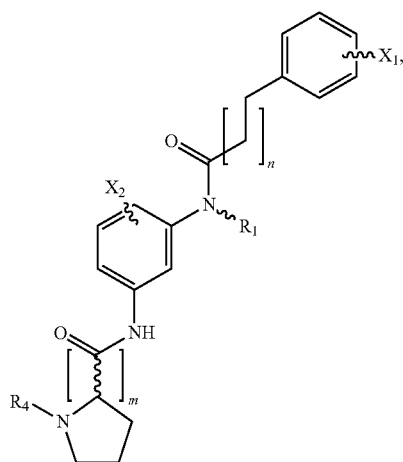
Formula 2A



Formula 2

In some embodiments the compound can have a structure according to Formula 2B

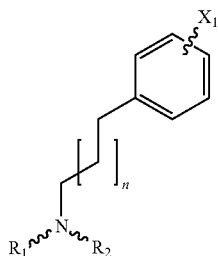
3



Formula 2B

where X_1 and X_2 can each be independently selected from the group of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of Formula 2B, an acyl, an amino a sulfono, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof, where R_1 can be a carboxylic acid or an ester, where R_4 can be H, an C_1 - C_{10} alkyl, an amino, a carboxylic acid, an ester, an amino acid or positively charged derivative thereof, where n can be 1-10, and where m represents the number of carbon atoms in an alkyl, where m can range from 0-10.

Also provided herein are pharmaceutical formulations that can include a compound having a structure according to Formula 2,

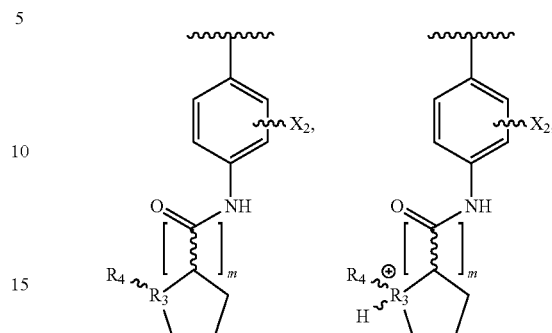


Formula 2

where n can be 1-10, where X_1 can be selected from the group of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of Formula 2, an acyl, an amino a sulfono, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof, where R_1 can be a carboxylic acid or an ester where R_2 can be a 5, 6, or 7 member heterocyclic ring that can be

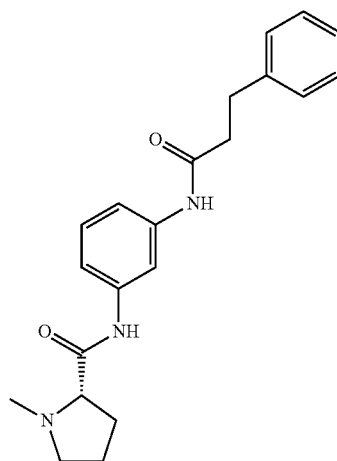
4

optionally further substituted with a suitable substituent, a sulfonium, a quaternary ammonium,



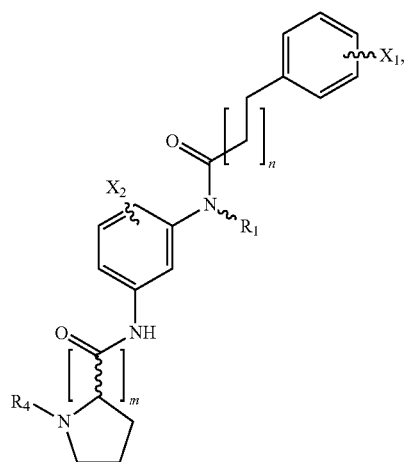
where X_2 can be selected from the group of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of Formula 2, an acyl, an amino a sulfono, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof, where R_3 can be N or S, where R_4 can be H, an C_1 - C_{10} alkyl, an amino, a carboxylic acid, an ester, an amino acid or positively charged derivative thereof, and where m represents the number of carbon atoms in an alkyl and can range from 0-10; and a pharmaceutically acceptable carrier. The compound(s) can bind Site 2 on sortilin or analogue thereof. In some embodiments, the compound can bind to Site 2 on sortilin or analogue thereof and can increase binding of a compound to site 1 of sortilin or an analogue thereof. In some embodiments the compound can have a structure according to Formula 2A

Formula 2A



In some embodiments the compound can have a structure according to Formula 2B

5



where X_1 and X_2 can each be independently selected from the group consisting of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of Formula 2B, an acyl, an amino, a sulfono, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof,

where R_1 can be a carboxylic acid or an ester,

where R_4 can be H, an C_1 - C_{10} alkyl, an amino, a carboxylic acid, an ester, an amino acid or positively charged derivative thereof,

where n can be 1-10, and

where m represents the number of carbon atoms in an alkyl, where m can range from 0-10.

Also provided herein are methods that can include the step of contacting sortilin or an analogue thereof with an amount of a compound having a structure according to Formula 2

Formula 2B

5

10

15

20

25

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35

40

45

50

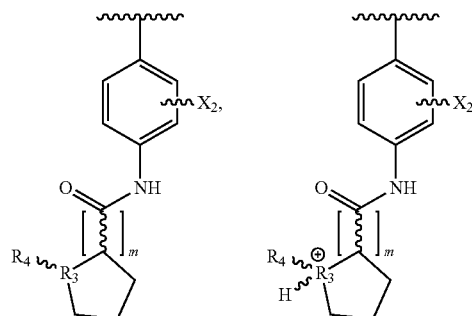
55

60

65

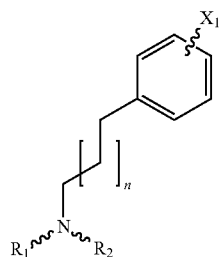
6

a iodo, a fluoro, and permissible combinations thereof, where R_1 can be a carboxylic acid or an ester where R_2 can be a 5, 6, or 7 member heterocyclic ring that can be optionally further substituted with a suitable substituent, a sulfonium, a quaternary ammonium,



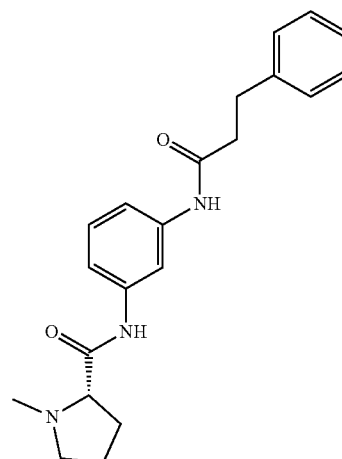
where X_2 can be selected from the group of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of Formula 2, an acyl, an amino, a sulfono, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof, where R_3 can be N or S, where R_4 can be H, an C_1 - C_{10} alkyl, an amino, a carboxylic acid, an ester, an amino acid or positively charged derivative thereof, and where m represents the number of carbon atoms in an alkyl and can range from 0-10. The compound(s) can bind Site 2 on sortilin or analogue thereof. In some embodiments, the compound can bind to Site 2 on sortilin or analogue thereof and can increase binding of a compound to site 1 of sortilin or an analogue thereof. In some embodiments the compound can have a structure according to Formula 2A

Formula 2A



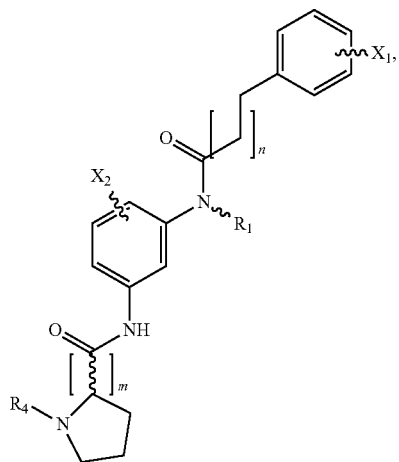
Formula 2

where n can be 1-10, where X_1 can be selected from the group of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of Formula 2, an acyl, an amino, a sulfono, a chloro, a bromo,



In some embodiments the compound can have a structure according to Formula 2B

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where X_1 and X_2 can each be independently selected from the group consisting of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of Formula 2B, an acyl, an amino, a sulfonyl, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof, where R_1 can be a carboxylic acid or an ester, where R_4 can be H, an C_1 - C_{10} alkyl, an amino, a carboxylic acid, an ester, an amino acid or positively charged derivative thereof, where n can be 1-10, and where m represents the number of carbon atoms in an alkyl, where m can range from 0-10. In some embodiments, the step of contacting can occur in vitro. In some embodiments, the step of contacting can occur in vivo. The method can further include the step of administering the compound having a structure according to Formula 2 to a subject, wherein the step of administering can occur prior to the step of contacting. In some embodiments, subject suffers from a disease whose pathology can involve a ligand of sortilin or analogue thereof. In some embodiments, the disease can be hypercholesterolemia or Alzheimer's disease. In some embodiments, the ligand of sortilin or analogue thereof is PCSK9, apolipoprotein B100, phosphatidylinositides, anionic phospholipids, anionic phospholipids contained on VLDL, amyloid beta, amyloid precursor protein, neurotensin or analogues of neurotensin, lipoprotein lipase, apolipoprotein AV or apolipoprotein E. In some embodiments, the ligand of sortilin or analogue thereof can be an apolipoprotein. In some embodiments the ligand of sortilin or analogue thereof can be PCSK9.

BRIEF DESCRIPTION OF THE DRAWINGS

Further aspects of the present disclosure will be readily appreciated upon review of the detailed description of its various embodiments, described below, when taken in conjunction with the accompanying drawings.

FIGS. 1A-1D show VLDL-apo B secretion and sortilin expression in McA cells. FIG. 1A shows a graph demonstrating VLDL-apo B and VLDL-B100 secretion were measured following 18 h incubation of McA cells in cDMEM (black bars) and in 1% BSA/DMEM (gray bars) by immuno slot blotting. FIG. 1B shows a representative standard curve generated from an immuno slot blot (duplicate standards) where VLDL protein is plotted against average chemiluminescence signal in arbitrary units (AU) generated using anti-rat B100 monoclonal antibody and HRP-anti-mouse

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IgG. FIG. 1C shows representative immunoblots of McA cell lysates from four independent experiments comparing McA cells incubated in cDMEM with 1% BSA/DMEM. Anti-GAPDH blots display equal protein loading. FIG. 1D shows a graph demonstrating relative expression of Apob, Sort1 and Atf3 mRNA in McA cells incubated in 1% BSA/DMEM as a percentage of mRNA present in McA cells incubated in cDMEM. Results are averages from 4 independent experiments.* Indicates means are significantly different.

FIGS. 2A-2C show the effect of variable sortilin expression on secretion of VLDL-B100 by McA cells. (FIG. 2A) Following siRNA-mediated KD of sortilin, three clones of McA cells were selected representing high (H, McA60), medium (M, McA62) and low (L, McA53) sortilin expression evaluated by immunoblotting compared with wild type (WT) McA cells. The effect of increasing sortilin KD on VLDL-B100 secretion by McA cells incubated in cDMEM (FIG. 2B) or incubated in 1% BSA/DMEM (FIG. 2C). Results in (FIG. 2B) and (FIG. 2C) are averages of triplicate plates for each condition ($n=2$ studies). * Indicates means are significantly different from the SCR McA cell line.

FIGS. 3A-3D show binding characteristics of NT and cpd984 to hsortilin. (FIG. 3A) Schrödinger's PRIME was run on PDB ID: 4PO7 predicting missing side chains of NT connecting two fragments of the peptide from the crystal structure across the length of the cavity of hsortilin. The predicted peptide XLYENKPRRPYL generated by connecting C-terminal fragment PYIL-OH and N-terminal fragment XLYEN-OH matches the amino acid sequence for NT and is the right length to connect exactly the N-terminus fitting into site 2 of hsortilin indicated by four of the residues present in the ligand binding diagram of cpd984 generated from the crystal structure. This is depicted in an atomic space filling model and colored yellow and the C-terminal end fitting into site 1 of hsortilin as indicated by the conserved residue R292 (R325 of hsortilin) of crystal structure 4PO7 is depicted in blue. (FIG. 3B) Ligand interaction diagram depicting binding of cpd984 binding to site 2 of hsortilin using Schrödinger's GLIDE indicating key amino acid residues likely to be involved in binding (blue, basic; red, acidic; green, hydrophobic amino acids) including glutamic acids 448 and 542 of pdb ID 4PO7 (Glu481 and Glu 575 of hsortilin). Hydrogen bonds are indicated with purple arrows and pi-stacking are indicated with green arrows. (FIG. 3C) Sensograms showing binding of NT to hsortilin (red, lower curve); cpd984 to hsortilin (green, middle curve) and NT binding in the presence of cpd984 (black, upper curve). The binding response in response units (RU) versus time (sec) was plotted of 100 nM NT injected alone less blank (red 4.5 and 5.5 RU) or with 500 μ M cpd984 (blue 64.1 and 64.7 RU) and 500 μ M cpd984 alone (green 45.1 and 44.8 RU). Subtracting the averaged RU of NT co-injected with 984 (64.4 RU) from 500 μ M cpd984 and 100 nM NT (50.0 RU) results in a 14.5 RU increase in binding to hsortilin. FIG. 3D) McA cells were incubated in 1% BSA/DMEM containing increasing concentrations of cpd984 for 18 h, and secreted VLDL-B100 was quantified by immuno slot blotting ($n=2$ studies). * Indicates means differ from the no cpd984 condition. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

FIGS. 4A-4B shows the effect of cpd984 on VLDL-B100 secretion and on insulin sensitivity in McA cells. (FIG. 4A) McA cell lines with variable sortilin KD were incubated with vehicle (black bars) or with 10 μ M cpd984 (gray bars) for 18 h (3-100 mm plates per condition). Viability of McA

cells was not compromised by incubations with cpd984 as there was no significant release of LDH into the medium compared with control incubations. VLDL was isolated from media of each plate and VLDL-B100 was quantified by immuno slot blotting. Results presented are averages of the 3 plates \pm S.D. (FIG. 4B) McA cells were incubated for 18 h in 1% BSA/DMEM \pm 10 μ M cpd984. McA cells were then stimulated with a time course of 250 nM insulin (0, 5, 10 and 15 min). IR β and AKT, pY-IR β , and p-AKT (S473) were evaluated by immunoblotting using protein and phospho-specific antibodies.

FIGS. 5A-5B demonstrate inhibition of the second site of sortilin by 984. McA cells were incubated in 1% BSA/DMEM with DMSO, 541 (10 μ M) or 984 (10 μ M) for 18 h. Cellular proteins were extracted and analyzed by IB for LDLR (LS-C146979, 1:2000) (FIG. 5A) or PCSK9 (Ab125251, 1:2000) (FIG. 5B) using HRP-linked secondary antibody and ECL detection. Loading controls included β -ACTIN (Rockland ph600-401-886) and GAPDH (cb1001mAb (6C5)). Band intensities were measured using ChemiDocXRS+ system and evaluated using ImageLab 5.1 software (BioRad). Results are average of triplicates \pm S.D.

FIG. 6 demonstrates an effect of cpd984 and cpd541 on secretion of VLDL-B100 by hepatocytes. McArdle rat hepatocytes were incubated with increasing concentrations of cpd984 or cpd541 for 14-18 h and media were collected. VLDL (d<0.19 g/ml) was isolated from media by ultracentrifugation and were slotted onto PVDF membranes. After blocking non-specific binding, blots were incubated with monoclonal anti-B100 antibody (24.05) followed by HRP-conjugated anti-mouse IgG. Binding was detected by chemiluminescence detection and quantified using Bio-Rad software. Results are the average of triplicate analysis in one study.

FIGS. 7A-7C show graphs demonstrating an effect of cpd984 and pd541 alone and in combination on the secretion of VLDL-B100 and VLDL-apo E by hepatocytes. McArdle rat hepatocytes were incubated with five media concentrations 0 (no cpd), 1 (0.1 μ M), 2 (1 μ M), 3 (10 μ M) and 4 (25 μ M) of cpd984 (FIG. 7A), cpd541 (FIG. 7B), or with a combination of both compounds (FIG. 7C) at indicated concentrations. Hepatocytes were incubated overnight (14-18 h) and media were collected and VLDL-B100 and VLDL-apo E were detected by slot blotting as described above. An anti-rat apoE antibody was used for measurement of VLDL-apo E secretion. Results are the average of triplicate analysis in one study.

FIG. 8 shows a graph demonstrating the results of two trials of Label Free Thermophoresis of 541 to human sortilin. Due to inherent fluorescence of 541 small molecule, thermophoresis was performed in two different buffer conditions (PBS n=3 and MST n=2) and at two sets of concentration ranges. Because thermophoresis performed in PBS at higher concentration 51 revealed substantial initial fluorescence, results in MST buffer are used for Kd estimation.

FIGS. 9A-9B show graphs representing the results of two Trials of SPR Indicating Binding of 541 to hsortilin at with Kd of 7.5 μ M.

FIGS. 10A-10B show graphs demonstrating the results of SPR of 984 which reveals two different binding ranges. SPR indicates a high and low affinity binding site of cpd984 to human sortilin of hsortilin with Kd of about 200 μ M and about 20 nM.

FIGS. 11A-11C show graphs demonstrating the results of thermophoresis of Cpd984, which revealed two different binding ranges with Kd of 7 nM and about 100 to 300 μ M.

DETAILED DESCRIPTION

Before the present disclosure is described in greater detail, it is to be understood that this disclosure is not limited to particular embodiments described, and as such may, of course, vary. It is also to be understood that the terminology used herein is for the purpose of describing particular embodiments only, and is not intended to be limiting.

Where a range of values is provided, it is understood that each intervening value, to the tenth of the unit of the lower limit unless the context clearly dictates otherwise, between the upper and lower limit of that range and any other stated or intervening value in that stated range, is encompassed within the disclosure. The upper and lower limits of these smaller ranges may independently be included in the smaller ranges and are also encompassed within the disclosure, subject to any specifically excluded limit in the stated range. Where the stated range includes one or both of the limits, ranges excluding either or both of those included limits are also included in the disclosure.

Unless defined otherwise, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which this disclosure belongs. Although any methods and materials similar or equivalent to those described herein can also be used in the practice or testing of the present disclosure, the preferred methods and materials are now described.

All publications and patents cited in this specification are herein incorporated by reference as if each individual publication or patent were specifically and individually indicated to be incorporated by reference and are incorporated herein by reference to disclose and describe the methods and/or materials in connection with which the publications are cited. The citation of any publication is for its disclosure prior to the filing date and should not be construed as an admission that the present disclosure is not entitled to antedate such publication by virtue of prior disclosure. Further, the dates of publication provided could be different from the actual publication dates that may need to be independently confirmed.

As will be apparent to those of skill in the art upon reading this disclosure, each of the individual embodiments described and illustrated herein has discrete components and features which may be readily separated from or combined with the features of any of the other several embodiments without departing from the scope or spirit of the present disclosure. Any recited method can be carried out in the order of events recited or in any other order that is logically possible.

Embodiments of the present disclosure will employ, unless otherwise indicated, techniques of biology, molecular biology, microbiology, physiology, organic chemistry, biochemistry, and the like, which are within the skill of the art. Such techniques are explained fully in the literature.

Definitions

As used herein, "about," "approximately," and the like, when used in connection with a numerical variable, can refer to the value of the variable and to all values of the variable that are within the experimental error (e.g., within the 95% confidence interval for the mean) or within \pm 10% of the indicated value, whichever is greater.

As used herein, "additive effect" can refer to an effect arising between two or more molecules, compounds, substances, factors, or compositions that is equal to or the same as the sum of their individual effects.

As used herein, "active derivative" and the like can refer to a compound that is capable of binding sortilin or an

analogue thereof as provided herein. The term “active derivative” and the like can also refer to a compound or analogue thereof provided herein that can be effective at altering the activity and/or abundance of a protein that is a ligand of sortilin or an analogue thereof, such as PCSK9. The term “active derivative” can also refer to a compound or analogue thereof that can be effective at treating a disease or symptom thereof whose pathology involves a ligand of sortilin or an analogue thereof. Assays for testing the ability of an active derivative to perform in this fashion are known to those of ordinary skill in the art and provided herein. The assays can include, but are not limited to, in vitro and in vivo assays.

As used herein, “administering” can refer to any administration route, including but not limited to, administration that is oral, topical, intravenous, subcutaneous, transcutaneous, transdermal, intramuscular, intra-articular, parenteral, intra-arterial, intradermal, intraventricular, intracranial, intraperitoneal, intralesional, intranasal, rectal, vaginal, by inhalation, or via an implanted reservoir. The term “parenteral” includes subcutaneous, intravenous, intramuscular, intra-articular, intra-synovial, intrasternal, intranasal, intrathecal, intrahepatic, intralesional, and intracranial injections or infusion techniques.

As used herein, “attached,” “attachment” and the like can refer to the formation of a covalent or non-covalent association (e.g. a bond) between two or more molecules or conjugation of two or more molecules. As used herein, “attached,” “attachment” and the like can refer to direct association of two or more molecules together with no intermediate molecules between those that are attached together or to the indirect attachment of two or more molecules together that is mediated via one or more linkers. Where the association is non-covalent, this can encompass charge interactions, affinity interactions, metal coordination, physical adsorption, host-guest interactions, hydrophobic interactions, π - π stacking interactions, hydrogen bonding interactions, van der Waals interactions, magnetic interactions, electrostatic interactions, dipole-dipole interactions, and/or combinations thereof. Where the association is covalent, this can encompass bonds where a pair of electrons is shared between one or more atoms in each molecule involved.

As used interchangeably herein, “biocompatible,” “biocompatibility,” and “biologically compatible” can refer to materials that are, with any metabolites or degradation products thereof, generally non-toxic to the recipient, and cause no significant adverse effects to the recipient. Generally speaking, biocompatible materials are materials which do not elicit a significant inflammatory or immune response when administered to a patient. In some embodiments, a biocompatible material elicits no detectable change in one or more biomarkers indicative of an immune response. In some embodiments, a biocompatible material elicits no greater than a 10% change, no greater than a 20% change, or no greater than a 40% change in one or more biomarkers indicative of an immune response.

As used herein, “a compound of formula (1), (2), (3), (4), (5), (6), (6a), (6b) (7), (8), (9), (10), (11), (12), (13), (14), (A), (B), (C), (D), and so forth and so on,” or “a compound having a structure according to formula (1), (2), (3), (4), (5), (6), (6a), (6b) (7), (8), (9), (10), (11), (12), (13), (14), (A), (B), (C), (D), etc.,” compound (1), (2), (3), (4), (5), (6), (6a), (6b) (7), (8), (9), (10), (11), (12), (13), (14), (A), (B), (C), (D), and so forth and so on,” or a “compound” can include all or any sub-group of solvates, complexes, polymorphs, derivatives thereof (including but not limited to, radiola-

beled derivatives (including deuterated derivatives where one or more H are replaced by D)), tautomers, stereoisomers, and optical isomers of the compound of the formulas listed above and salts thereof.

As used herein, “chemotherapeutic agent” or “chemotherapeutic” can refer to a therapeutic agent utilized to prevent or treat cancer.

As used herein, “control” can refer to an alternative subject or sample used in an experiment for comparison purposes and included to minimize or distinguish the effect of variables other than an independent variable. A control can be positive or negative. One of ordinary skill in the art will appreciate what are appropriate controls for a given context.

As used herein, “concentrated” can refer to an amount of a molecule, compound, or composition, including, but not limited to, a chemical compound, polynucleotide, peptide, polypeptide, protein, antibody, or fragments thereof, that indicates that the sample is distinguishable from its naturally occurring counterpart in that the concentration or number of molecules per volume is greater than that of its naturally occurring counterpart.

As used herein, “derivative” can refer to substituting, deleting, and/or adding one or more atoms or functional groups. The term “derivative” does not mean that the derivative is synthesized from the parent compound either as a starting material or intermediate, although this may be the case. The term “derivative” can include salts, prodrugs, or metabolites of the parent compound. Derivatives include compounds in which free amino groups in the parent compound have been derivatized to form amine hydrochlorides, p-toluene sulfonamides, benzoxycarbonylamides, t-butyloxycarbonylamides, thiourethane-type derivatives, trifluoroacetyl-amides, chloroacetyl-amides, or formamides. Derivatives include compounds in which carboxyl groups in the parent compound have been derivatized to form salts, methyl and ethyl esters, or other types of esters or hydrazides. Derivatives include compounds in which hydroxyl groups in the parent compound have been derivatized to form O-acyl or O-alkyl derivatives. Derivatives include compounds in which a hydrogen bond donating group in the parent compound is replaced with another hydrogen bond donating group such as OH, NH, or SH. Derivatives include replacing a hydrogen bond acceptor group in the parent compound with another hydrogen bond acceptor group such as esters, ethers, ketones, carbonates, tertiary amines, imines, thiones, sulfones, tertiary amides, and sulfides. “Derivatives” also includes extensions of the replacement of the cyclopentane ring with saturated or unsaturated cyclohexane or other more complex, e.g., nitrogen-containing rings, and extensions of these rings with various side groups.

As used herein, “diluted” can refer to an amount of a molecule, compound, or composition including but not limited to, a chemical compound, polynucleotide, peptide, polypeptide, protein, antibody, or fragments thereof, that indicates that the sample is distinguishable from its naturally occurring counterpart in that the concentration or number of molecules per volume is less than that of its naturally occurring counterpart.

As used herein, “deoxyribonucleic acid (DNA)” and “ribonucleic acid (RNA)” can generally refer to any polynucleotide or polydeoxyribonucleotide, which may be unmodified RNA or DNA or modified RNA or DNA. RNA may be in the form of a tRNA (transfer RNA), snRNA (small nuclear RNA), rRNA (ribosomal RNA), mRNA (messenger RNA), anti-sense RNA, RNAi (RNA interference construct), siRNA (short interfering RNA), or ribozymes.

As used herein, "DNA molecule" can include nucleic acids/polynucleotides that are made of DNA.

As used herein, "dose," "unit dose," or "dosage" can refer to physically discrete units suitable for use in a subject, each unit containing a predetermined quantity of a compound as provided herein and/or a pharmaceutical formulation thereof calculated to produce the desired response or responses in association with its administration.

As used herein, "effective amount" can refer to an amount sufficient to effect beneficial or desired results. An effective amount can be administered in one or more administrations, applications, or dosages. "Effective amount" can refer to an amount of a compound, derivative, and/or formulation thereof provided herein that can treat or prevent a disease or symptom thereof whose pathology involves a ligand of sortilin or an analogue thereof. Some diseases include, but are not limited to, whose pathology involves PCSK9, ApoB, binding of anionic phospholipids, binding of anionic phospholipids contained on VLDL, Amyloid Beta, Amyloid precursor protein (APP), neurotensin or analogues of neurotensin, receptor associated protein (RAP), Gal A, amyloid beta-protein, GLUT4 and/or ApoE. In some embodiments, the disease can be cholesterolemia, hypercholesterolemia, hepatic steatosis, diabetes, auto-immune neuroinflammation, Fabry disease, Alzheimer's disease, and combinations thereof. The term "effective amount" can refer to the amount of a compound provided herein to inhibit 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 97, 98, 99, and/or about 100% or more of the binding of a ligand to sortilin or analogue thereof. The term "effective amount" can refer to the amount of a compound provided herein to inhibit 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 97, 98, 99, and/or about 100% or more of the binding of a ligand to Site 1 and or Site 2 of sortilin or analogue thereof.

As used herein, "hydrate" can refer to a compound formed by the addition of water. Typically, but not always, this will be crystalline lattice structures that incorporate water molecules. Hydrates include stoichiometric hydrates, as well as compositions containing variable amounts of water.

As used herein, "identity," "identical to", can refer to the relationship between two or more nucleotide or polypeptide sequences, as determined by comparing the sequences. In the art, "identity" also refers to the degree of sequence relatedness between nucleotide or polypeptide as determined by the match between strings of such sequences. "Identity" can be readily calculated by known methods, including, but not limited to, those described in (Computational Molecular Biology, Lesk, A. M., Ed., Oxford University Press, New York, 1988; Biocomputing: Informatics and Genome Projects, Smith, D. W., Ed., Academic Press, New York, 1993; Computer Analysis of Sequence Data, Part I, Griffin, A. M., and Griffin, H. G., Eds., Humana Press, New Jersey, 1994; Sequence Analysis in Molecular Biology, von Heinje, G., Academic Press, 1987; and Sequence Analysis Primer, Gribskov, M. and Devereux, J., Eds., M Stockton Press, New York, 1991; and Carillo, H., and Lipman, D., SIAM J. Applied Math. 1988, 48: 1073. Preferred methods to determine identity are designed to give the largest match between the sequences tested. Methods to determine identity are codified in publicly available computer programs. The percent identity between two sequences can be determined by using analysis software (e.g., Sequence Analysis Software Package of the Genetics Computer Group, Madison Wis.) that incorporates the Needleman and Wunsch, (J. Mol. Biol., 1970, 48: 443-453,) algorithm (e.g., NBLAST, and

XBLAST). The default parameters are used to determine the identity for the polypeptides of the present disclosure, unless stated otherwise.

As used herein, "immunomodulator," can refer to an agent, such as a therapeutic agent, which is capable of modulating or regulating one or more immune function or response.

As used herein, "isolated" means separated from constituents, cellular and otherwise, in which the polynucleotide, peptide, polypeptide, protein, antibody, or fragments thereof, are normally associated with in nature. A non-naturally occurring polynucleotide, peptide, polypeptide, protein, antibody, or fragments thereof, do not require "isolation" to distinguish it from its naturally occurring counterpart.

As used herein, "mitigate" can refer to reducing a particular characteristic, symptom, or other biological or physiological parameter associated with a disease or disorder.

The term "molecular weight", as used herein, can generally refers to the mass or average mass of a material. If a polymer or oligomer, the molecular weight can refer to the relative average chain length or relative chain mass of the bulk polymer. In practice, the molecular weight of polymers and oligomers can be estimated or characterized in various ways including gel permeation chromatography (GPC) or capillary viscometry. GPC molecular weights are reported as the weight-average molecular weight (M_w) as opposed to the number-average molecular weight (M_n). Capillary viscometry provides estimates of molecular weight as the inherent viscosity determined from a dilute polymer solution using a particular set of concentration, temperature, and solvent conditions.

As used herein, "negative control" can refer to a "control" that is designed to produce no effect or result, provided that all reagents are functioning properly and that the experiment is properly conducted. Other terms that are interchangeable with "negative control" include "sham," "placebo," and "mock."

As used herein, "nucleic acid" and "polynucleotide" generally refer to a string of at least two base-sugar-phosphate combinations and refers to, among others, single- and double-stranded DNA, DNA that is a mixture of single- and double-stranded regions, single- and double-stranded RNA, and RNA that is mixture of single- and double-stranded regions, hybrid molecules comprising DNA and RNA that may be single-stranded or, more typically, double-stranded or a mixture of single- and double-stranded regions. In addition, polynucleotide as used herein refers to triple-stranded regions comprising RNA or DNA or both RNA and DNA. The strands in such regions may be from the same molecule or from different molecules. The regions may include all of one or more of the molecules, but more typically involve only a region of some of the molecules. One of the molecules of a triple-helical region often is an oligonucleotide. "Polynucleotide" and "nucleic acids" also encompasses such chemically, enzymatically or metabolically modified forms of polynucleotides, as well as the chemical forms of DNA and RNA characteristic of viruses and cells, including simple and complex cells, inter alia. For instance, the term polynucleotide includes DNAs or RNAs as described above that contain one or more modified bases. Thus, DNAs or RNAs comprising unusual bases, such as inosine, or modified bases, such as tritylated bases, to name just two examples, are polynucleotides as the term is used herein. "Polynucleotide" and "nucleic acids" also includes PNAs (peptide nucleic acids), phosphorothioates, and other variants of the phosphate backbone of native nucleic acids.

Natural nucleic acids have a phosphate backbone, artificial nucleic acids may contain other types of backbones, but contain the same bases. Thus, DNAs or RNAs with backbones modified for stability or for other reasons are “nucleic acids” or “polynucleotide” as that term is intended herein. As used herein, “nucleic acid sequence” and “oligonucleotide” also encompasses a nucleic acid and polynucleotide as defined above.

As used herein “peptide” refers to chains of at least 2 amino acids that are short, relative to a protein or polypeptide.

As used herein, “pharmaceutical formulation” can refer to the combination of an active agent, compound, or ingredient with a pharmaceutically acceptable carrier or excipient, making the composition suitable for diagnostic, therapeutic, or preventive use in vitro, in vivo, or ex vivo.

As used herein “pharmaceutically effective amount”, “effective amount” and the like can refer to an amount of a compound, derivative, or formulation thereof provided herein that can treat or prevent a disease or symptom thereof whose pathology involves a ligand of sortilin or an analogue thereof. Some diseases include, but are not limited to, whose pathology involves PCSK9, ApoB, binding of anionic phospholipids, binding of anionic phospholipids contained on VLDL, Amyloid Beta, Amyloid precursor protein (APP), neurotensin or analogues of neurotensin, receptor associated protein (RAP), Gal A, amyloid beta-protein, GLUT4 and/or ApoE. In some embodiments, the disease can be cholesterolemia, Alzheimer’s disease, hypercholesterolemia, hepatic steatosis, diabetes, auto-immune neuroinflammation, Fabry disease, and combinations thereof. The term “pharmaceutically effective amount” can refer to the amount of a compound provided herein to inhibit 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 97, 98, 99, and/or about 100% or more of the binding of a ligand to sortilin or analogue thereof. The term “pharmaceutically effective amount” can refer to the amount of a compound provided herein to inhibit 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 97, 98, 99, and/or about 100% or more of the binding of a ligand to Site 1 and or Site 2 of sortilin or analogue thereof. In embodiments, the “pharmaceutically effective amount” can be the least amount of a compound, derivative or formulation thereof provided herein needed to treat, prevent, or elicit the desired biological and/or medical effect in the response of a cell, tissue, organ, system, or subject that is being sought by the researcher, veterinarian, medical doctor or other clinician. In some embodiments, the “pharmaceutically effective amount” can be the least amount that can treat or prevent treat or prevent a disease or symptom thereof whose pathology involves a ligand of sortilin or an analogue thereof. Some diseases include, but are not limited to, whose pathology involves PCSK9, ApoB, binding of anionic phospholipids, binding of anionic phospholipids contained on VLDL, Amyloid Beta, Amyloid precursor protein (APP), neurotensin or analogues of neurotensin, receptor associated protein (RAP), Gal A, amyloid beta-protein, GLUT4 and/or ApoE. In some embodiments, the disease can be cholesterolemia, Alzheimer’s disease, hypercholesterolemia, hepatic steatosis, diabetes, auto-immune neuroinflammation, Fabry disease, and combinations thereof. The term “pharmaceutically effective amount” can refer to the least amount of a compound, derivative, or formulation thereof provided herein that can inhibit 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 97, 98, 99, and/or about 100% or more of the binding of a ligand to sortilin or analogue thereof. The term “pharmaceutically effective

amount” can refer to the least amount of a compound provided herein to inhibit 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 97, 98, 99, and/or about 100% or more of the binding of a ligand to Site 1 and or Site 2 of sortilin or analogue thereof. “Pharmaceutically effective amount” or “pharmaceutically effective dose,” can refer to the amount of a compound or formulation thereof provided herein that will elicit the biological and/or medical response of a cell, tissue, organ, system, or subject that is being sought by the researcher, veterinarian, medical doctor or other clinician. The pharmaceutically effective amount can vary depending on the compound, formulation the disorder or condition (normal or abnormal) and its severity, the route of administration, time of administration, rate of excretion, drug or compound, judgment of the researcher, veterinarian, medical doctor or other clinician, dosage form, and the age, weight, general health, sex and/or diet of the subject to be treated.

As used herein, “pharmaceutically acceptable” can refer to compounds, materials, compositions, and/or dosage forms which are, within the scope of sound medical judgment, suitable for use in contact with the tissues of human beings and animals without excessive toxicity, irritation, allergic response, or other problems or complications commensurate with a reasonable benefit/risk ratio, in accordance with the guidelines of agencies such as the Food and Drug Administration.

As used herein, “pharmaceutically acceptable carrier or excipient” can refer to a carrier or excipient that is useful in preparing a pharmaceutical composition that is generally safe, non-toxic and neither biologically nor otherwise undesirable, and includes a carrier or excipient that is acceptable for veterinary use as well as human pharmaceutical use. A “pharmaceutically acceptable carrier or excipient” as used herein also includes both one and more than one such carrier or excipient. Pharmaceutically acceptable carriers include, but are not limited to, diluents, preservatives, binders, lubricants, disintegrators, swelling agents, fillers, stabilizers, and combinations thereof.

As used herein, “pharmaceutically acceptable salt” can refer to any salt derived from organic and inorganic acids of a compound described herein. Pharmaceutically acceptable salt also refers to a salt of a compound described having an acidic functional group, such as a carboxylic acid functional group, and a base. Pharmaceutically acceptable salt also includes hydrates of a salt of a compound described herein.

As used herein, “positive control” can refer to a “control” that is designed to produce the desired result, provided that all reagents are functioning properly and that the experiment is properly conducted.

As used herein, “protein” as used herein can refer to a molecule composed of one or more chains of amino acids in a specific order. The term protein is used interchangeable with “polypeptide.” The order is determined by the base sequence of nucleotides in the gene coding for the protein. Proteins are required for the structure, function, and regulation of the body’s cells, tissues, and organs.

As used herein, “purified” or “purify” can be used in reference to a nucleic acid sequence, peptide, or polypeptide that has increased purity relative to the natural environment.

As used herein, “preventative,” “preventing,” “prevent” and the like can refer to partially or completely delaying and/or precluding the onset or recurrence of a disorder or conditions and/or one or more of its attendant symptoms or barring a subject from acquiring or reacquiring a disorder or condition or reducing a subject’s risk of acquiring or reacquiring a disorder or condition or one or more of its

attendant symptoms including, but not limited to, malaria, infection and/or transmission of a parasite of the genus *Plasmodium* or a symptom thereof.

As used herein, “purified” or “purify” can be used in reference to a nucleic acid sequence, peptide, or polypeptide that has increased purity relative to the natural environment.

As used herein, “separated” can refer to the state of being physically divided from the original source or population such that the separated compound, agent, particle, chemical compound, or molecule can no longer be considered part of the original source or population.

As used herein, “solvate” refers to a complex of variable stoichiometry formed by a solute (e.g. formulas (1)-(1) (A), (B), (C), (D), or any other compound herein or a salt thereof) and a solvent. Pharmaceutically acceptable solvates may be formed for crystalline compounds wherein solvent molecules are incorporated into the crystalline lattice during crystallization. The incorporated solvent molecules can be water molecules or non-aqueous molecules, such as but not limited to, ethanol, isopropanol, dimethyl sulfoxide, acetic acid, ethanolamine, and ethyl acetate molecules.

As used herein, the term “specific binding” can refer to non-covalent physical association of a first and a second moiety wherein the association between the first and second moieties is at least 2 times as strong, at least 5 times as strong as, at least 10 times as strong as, at least 50 times as strong as, at least 100 times as strong as, or stronger than the association of either moiety with most or all other moieties present in the environment in which binding occurs. Binding of two or more entities may be considered specific if the equilibrium dissociation constant, K_d , is 10^{-3} M or less, 10^{-4} M or less, 10^{-5} M or less, 10^{-6} M or less, 10^{-7} M or less, 10^{-8} M or less, 10^{-9} M or less, 10^{-10} M or less, 10^{-11} M or less, or 10^{-12} M or less under the conditions employed, e.g., under physiological conditions such as those inside a cell or consistent with cell survival. In some embodiments, specific binding can be accomplished by a plurality of weaker interactions (e.g., a plurality of individual interactions, wherein each individual interaction is characterized by a K_d of greater than 10^{-3} M). In some embodiments, specific binding, which can be referred to as “molecular recognition,” is a saturable binding interaction between two entities that is dependent on complementary orientation of functional groups on each entity. Examples of specific binding interactions include primer-polynucleotide interaction, aptamer-aptamer target interactions, antibody-antigen interactions, avidin-biotin interactions, ligand-receptor interactions, metal-chelate interactions, hybridization between complementary nucleic acids, etc.

As used interchangeably herein, “subject,” “individual,” or “patient,” can refer to a vertebrate, preferably a mammal, more preferably a human. Mammals include, but are not limited to, murines, simians, humans, farm animals, sport animals, and pets. The term “pet” includes a dog, cat, guinea pig, mouse, rat, rabbit, ferret, and the like. The term farm animal includes a horse, sheep, goat, chicken, pig, cow, donkey, llama, alpaca, turkey, and the like.

As used herein, “substantially pure” can mean an object species is the predominant species present (i.e., on a molar basis it is more abundant than any other individual species in the composition), and preferably a substantially purified fraction is a composition wherein the object species comprises about 50 percent of all species present. Generally, a substantially pure composition will comprise more than about 80 percent of all species present in the composition, more preferably more than about 85%, 90%, 95%, and 99%. Most preferably, the object species is purified to essential

homogeneity (contaminant species cannot be detected in the composition by conventional detection methods) wherein the composition consists essentially of a single species.

The terms “sufficient” and “effective,” as used interchangeably herein, can refer to an amount (e.g. mass, volume, dosage, concentration, and/or time period) needed to achieve one or more desired result(s). For example, a therapeutically effective amount refers to an amount needed to achieve one or more therapeutic effects.

As used herein, “synergistic effect,” “synergism,” or “synergy” can refer to an effect arising between two or more molecules, compounds, substances, factors, or compositions that that is greater than or different from the sum of their individual effects.

As used herein, “therapeutic,” “treating,” “treat,” and the like can refer to include partially or completely delaying, alleviating, mitigating or reducing the intensity of one or more attendant symptoms of a disease or condition including, but not limited to, those involving sortilin or analogue thereof, and/or a ligand of sortilin or analogue thereof.

As used herein, “variant” refers to a polypeptide that differs from a reference polypeptide, but retains essential properties. A typical variant of a polypeptide differs in amino acid sequence from another, reference polypeptide. Generally, differences are limited so that the sequences of the reference polypeptide and the variant are closely similar overall and, in many regions, identical. A variant and reference polypeptide may differ in amino acid sequence by one or more modifications (e.g., substitutions, additions, and/or deletions). A substituted or inserted amino acid residue may or may not be one encoded by the genetic code. A variant of a polypeptide may be naturally occurring such as an allelic variant, or it may be a variant that is not known to occur naturally. “Variant” includes functional and structural variants.

As used herein, “alkyl” and “alkylene” refer to a saturated hydrocarbon chain having the specified number of member atoms.

The term “alkyl” can also refer to the radical of saturated aliphatic groups (i.e., an alkane with one hydrogen atom removed), including straight-chain alkyl groups, branched-chain alkyl groups, cycloalkyl (alicyclic) groups, alkyl-substituted cycloalkyl groups, and cycloalkyl-substituted alkyl groups. “Alkyl” also refers to a saturated hydrocarbon chain having the specified number of atoms.

The term “alkyl” (or “lower alkyl”) as used herein can include both “unsubstituted alkyls” and “substituted alkyls,” the latter of which refers to alkyl moieties having one or more substituents replacing a hydrogen on one or more carbons of the hydrocarbon backbone. Such substituents include, but are not limited to, halogen, hydroxyl, carbonyl (such as a carboxyl, alkoxycarbonyl, formyl, or an acyl), thiocarbonyl (such as a thioester, a thioacetate, or a thioformate), alkoxyl, phosphoryl, phosphate, phosphonate, phosphinate, amino, amido, amidine, imine, cyano, nitro, azido, sulfhydryl, alkylthio, sulfate, sulfonate, sulfamoyl, sulfonamido, sulfonyl, heterocyclyl, aralkyl, or an aromatic or heteroaromatic moiety.

Unless the number of carbons is otherwise specified, “lower alkyl” as used herein can refer to an alkyl group, as defined above, but having from one to ten carbons in its backbone structure. Likewise, “lower alkenyl” and “lower alkynyl” have similar chain lengths.

It will be understood by those skilled in the art that the moieties substituted on the hydrocarbon chain can themselves be substituted, if appropriate. For instance, the substituents of a substituted alkyl may include halogen,

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hydroxy, nitro, thiols, amino, azido, imino, amido, phosphor-
 yl (including phosphonate and phosphinate), sulfonyl (in-
 cluding sulfate, sulfonamido, sulfamoyl and sulfonate), and
 silyl groups, as well as ethers, alkylthios, carbonyls (includ-
 ing ketones, aldehydes, carboxylates, and esters), $-\text{CF}_3$,
 $-\text{CN}$ and the like. Cycloalkyls can be substituted in the
 same manner.

As used herein, " C_{1-6} alkyl" can refer to an alkyl group
 having any number of member atoms from 1 to 6 member
 atoms, such as for example 1 to 4 atoms. Other alkyl groups
 may have any number of member atoms as indicated by the
 numbers given in the formula, which, like the previous
 example, can refer to an alkyl group having any number of
 member atoms within the specified range of member atoms.
 Alkyl groups may be straight or branched. Representative
 branched alkyl groups have one, two, or three branches.
 Alkyl includes methyl, ethyl, propyl (n-propyl and isopro-
 pyl), butyl (n-butyl, isobutyl, and t-butyl), pentyl (n-pentyl,
 isopentyl, and neopentyl), and hexyl.

As used herein, "heterocyclic group" can refer to a
 non-aromatic ring and having the specified number of mem-
 ber atoms being saturated or having one or more degrees of
 unsaturation and, unless otherwise specified, containing one
 or more heteroatoms.

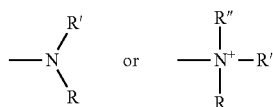
As used herein, "heteroaryl" can refer to an aromatic ring
 having the specified number of member atoms and, unless
 otherwise specified, containing one or more heteroatoms.
 Bicyclic and other polycyclic ring systems having a het-
 eroaryl ring are described as fused systems.

The term "heteroalkyl," as used herein, can refer to
 straight or branched chain, or cyclic carbon-containing rad-
 icals, or combinations thereof, containing at least one het-
 eroatom. Suitable heteroatoms include, but are not limited
 to, O, N, Si, P, Se, B, and S, wherein the phosphorous and
 sulfur atoms are optionally oxidized, and the nitrogen het-
 eroatom is optionally quaternized. Heteroalkyls can be sub-
 stituted as defined above for alkyl groups.

The term "heteroalkyl," as used herein, can refer to
 straight or branched chain, or cyclic carbon-containing rad-
 icals, or combinations thereof, containing at least one het-
 eroatom. Suitable heteroatoms include, but are not limited
 to, O, N, Si, P, Se, B, and S, wherein the phosphorous and
 sulfur atoms are optionally oxidized, and the nitrogen het-
 eroatom is optionally quaternized. Heteroalkyls can be sub-
 stituted as defined above for alkyl groups.

As used herein, "alkoxyl" or "alkoxy," as used herein, can
 refer to an alkyl group, as defined above, having an oxygen
 radical attached thereto. Representative alkoxy groups
 include methoxy, ethoxy, propyloxy, tert-butoxy and the
 like. An "ether" is two hydrocarbons covalently linked by an
 oxygen. Accordingly, the substituent of an alkyl that renders
 that alkyl is an ether or resembles an alkoxy, such as can be
 represented by one of $-\text{O-alkyl}$, $-\text{O-alkenyl}$, and $-\text{O-}$
 alkynyl . The terms "aroxy" and "aryloxy," as used inter-
 changeably herein, can be represented by $-\text{O-aryl}$ or O-het-
 eroaryl , wherein aryl and heteroaryl are as defined below.
 The alkoxy and aroxy groups can be substituted as described
 above for alkyl.

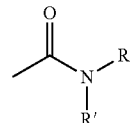
As used herein, "amine" and "amino" (and its protonated
 form) are art-recognized and refer to both unsubstituted and
 substituted amines, e.g., a moiety that can be represented by
 the general formula:



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wherein R, R', and R'' each independently represent a
 hydrogen, an alkyl, an alkenyl, $-(\text{CH}_2)_m-\text{R}_C$ or R and R'
 taken together with the N atom to which they are attached
 complete a heterocycle having from 4 to 8 atoms in the ring
 structure; R_C represents an aryl, a cycloalkyl, a cycloalkenyl,
 a heterocycle or a polycycle; and m is zero or an integer in
 the range of 1 to 8. In some embodiments, only one of R or
 R' can be a carbonyl, e.g., R, R' and the nitrogen together do
 not form an imide. In other embodiments, the term "amine"
 does not encompass amides, e.g., wherein one of R and R'
 represents a carbonyl. In further embodiments, R and R' (and
 optionally R'') each independently represent a hydrogen, an
 alkyl or cycloalkyl, an alkenyl or cycloalkenyl, or alkynyl.
 Thus, the term "alkylamine" as used herein means an amine
 group, as defined above, having a substituted (as described
 above for alkyl) or unsubstituted alkyl attached thereto, i.e.,
 at least one of R and R' is an alkyl group.

As used herein, "amido" is art-recognized as an amino-
 substituted carbonyl and includes a moiety that can be
 represented by the general formula:



wherein R and R' are as defined above.

As used herein, "Aryl" can refer to C_5 - C_{10} -membered
 aromatic, heterocyclic, fused aromatic, fused heterocyclic,
 biaromatic, or biheterocyclic ring systems. Broadly
 defined, "aryl", as used herein, includes 5-, 6-, 7-, 8-, 9-, and
 10-membered single-ring aromatic groups that may include
 from zero to four heteroatoms, for example, benzene, pyr-
 role, furan, thiophene, imidazole, oxazole, thiazole, triazole,
 pyrazole, pyridine, pyrazine, pyridazine, pyrimidine, and the
 like. Those aryl groups having heteroatoms in the ring
 structure may also be referred to as "aryl heterocycles" or
 "heteroaromatics." The aromatic ring can be substituted at
 one or more ring positions with one or more substituents
 including, but not limited to, halogen, azide, alkyl, aralkyl,
 alkenyl, alkynyl, cycloalkyl, hydroxyl, alkoxy, amino (or
 quaternized amino), nitro, sulphydryl, imino, amido, phos-
 phonate, phosphinate, carbonyl, carboxyl, silyl, ether, alkyl-
 thio, sulfonyl, sulfonamido, ketone, aldehyde, ester, hetero-
 cyclyl, aromatic or heteroaromatic moieties, $-\text{CF}_3$, $-\text{CN}$,
 and combinations thereof.

The term "aryl" can also include polycyclic ring systems
 having two or more cyclic rings in which two or more car-
 bons are common to two adjoining rings (i.e., "fused
 rings") wherein at least one of the rings is aromatic, e.g., the
 other cyclic ring or rings can be cycloalkyls, cycloalkenyls,
 cycloalkynyls, aryls and/or heterocycles. Examples of het-
 erocyclic rings include, but are not limited to, benzimida-
 zolyl, benzofuranyl, benzothiofuranyl, benzothiophenyl,
 benzoxazolyl, benzoxazolyl, benzthiazolyl, benztriazolyl,
 benztetrazolyl, benzisoxazolyl, benzisothiazolyl, benzimi-
 dazolyl, carbazolyl, 4aH carbazolyl, carbolinyl, chroma-
 nyl, chromenyl, cinnolinyl, decahydroquinolinyl, 2H,6H-1,
 5,2-dithiazinyl, dihydrofuro[2,3 b]tetrahydrofuran, furanyl,
 furazanyl, imidazolidinyl, imidazolyl, imidazolyl, 1H-in-
 dazolyl, indolenyl, indolyl, indolizyl, indolyl, 3H-indo-
 lyl, isatinoyl, isobenzofuranyl, isochromanyl, isoindazolyl,
 isoindolinyl, isoindolyl, isoquinolinyl, isothiazolyl, isox-
 azolyl, methylenedioxyphenyl, morpholinyl, naphthyridi-

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nyl, octahydroisoquinoliny, oxadiazolyl, 1,2,3-oxadiazolyl, 1,2,4-oxadiazolyl, 1,2,5-oxadiazolyl, 1,3,4-oxadiazolyl, oxazolidinyl, oxazolyl, oxindolyl, pyrimidinyl, phenanthridinyl, phenanthrolinyl, phenaziny, phenothiazinyl, phenoxathinyl, phenoxazinyl, phthalazinyl, piperazinyl, piperidinyl, piperidonyl, 4-piperidonyl, piperonyl, pteridinyl, purinyl, pyranyl, pyrazinyl, pyrazolidinyl, pyrazolinyl, pyrazolyl, pyridazinyl, pyridooxazole, pyridoimidazole, pyridothiazole, pyridinyl, pyridyl, pyrimidinyl, pyrrolidinyl, pyrrolinyl, 2H-pyrrolyl, pyrrolyl, quinazolinyl, quinolinyl, 4H-quinoliziny, quinoxaliny, quinuclidinyl, tetrahydrofuranyl, tetrahydroisoquinoliny, tetrahydroquinoliny, tetrazolyl, 6H-1,2,5-thiadiazinyl, 1,2,3-thiadiazolyl, 1,2,4-thiadiazolyl, 1,2,5-thiadiazolyl, 1,3,4-thiadiazolyl, thianthrenyl, thiazolyl, thienyl, thienothiazolyl, thienooxazolyl, thienoimidazolyl, thiophenyl, and xanthenyl. One or more of the rings can be substituted as defined above for "aryl."

As used herein, "aralkyl," can refer to an alkyl group substituted with an aryl group (e.g., an aromatic or heteroaromatic group).

As used herein, "aralkyloxy" can be represented by —O-aralkyl, wherein aralkyl is as defined above.

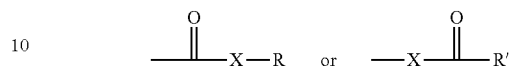
As used herein, "carbocycle," can refer to an aromatic or non-aromatic ring(s) in which each atom of the ring(s) is carbon.

As used herein, "heterocycle" or "heterocyclic" can refer to a monocyclic or bicyclic structure containing 3-10 ring atoms, and in some embodiments, containing from 5-6 ring atoms, wherein the ring atoms are carbon and one to four heteroatoms each selected from the following group of non-peroxide oxygen, sulfur, and N(Y) wherein Y is absent or is H, O, (C₁-C₁₀) alkyl, phenyl or benzyl, and optionally containing 1-3 double bonds and optionally substituted with one or more substituents. Examples of heterocyclic rings include, but are not limited to, benzimidazolyl, benzofuranyl, benzothiofuranyl, benzothiophenyl, benzoxazolyl, benzoxazolinyl, benzthiazolyl, benztriazolyl, benztetrazolyl, benzisoxazolyl, benzisothiazolyl, benzimidazoliny, carbazolyl, 4aH carbazolyl, carbolinyl, chromanyl, chromenyl, cinnoliny, decahydroquinoliny, 2H,6H-1,5,2-dithiazinyl, dihydrofuro[2,3 b]tetrahydrofuran, furanyl, furazanyl, imidazolidinyl, imidazoliny, imidazolyl, 1H-indazolyl, indoliny, indoliny, indoliziny, indolyl, 3H-indolyl, isatinoyl, isobenzofuranyl, isochromanyl, isoindazolyl, isoindoliny, isoindolyl, isoquinoliny, isothiazolyl, isoxazolyl, methyl-enedioxphenyl, morpholiny, naphthyridiny, octahydroisoquinoliny, oxadiazolyl, 1,2,3-oxadiazolyl, 1,2,4-oxadiazolyl, 1,2,5-oxadiazolyl, 1,3,4-oxadiazolyl, oxazolidinyl, oxazolyl, oxepanyl, oxetanyl, oxindolyl, pyrimidinyl, phenanthridinyl, phenanthroliny, phenaziny, phenothiazinyl, phenoxathinyl, phenoxazinyl, phthalazinyl, piperazinyl, piperidinyl, piperidonyl, 4-piperidonyl, piperonyl, pteridinyl, purinyl, pyranyl, pyrazinyl, pyrazolidinyl, pyrazolinyl, pyrazolyl, pyridazinyl, pyridooxazole, pyridoimidazole, pyridothiazole, pyridinyl, pyridyl, pyrimidinyl, pyrrolidinyl, pyrrolinyl, 2H-pyrrolyl, pyrrolyl, quinazolinyl, quinolinyl, 4H-quinoliziny, quinoxaliny, quinuclidinyl, tetrahydrofuranyl, tetrahydroisoquinoliny, tetrahydropyranyl, tetrahydroquinoliny, tetrazolyl, 6H-1,2,5-thiadiazinyl, 1,2,3-thiadiazolyl, 1,2,4-thiadiazolyl, 1,2,5-thiadiazolyl, 1,3,4-thiadiazolyl, thianthrenyl, thiazolyl, thienyl, thienothiazolyl, thienooxazolyl, thienoimidazolyl, thiophenyl, and xanthenyl. Heterocyclic groups can optionally be substituted with one or more substituents at one or more positions as defined above for alkyl and aryl, for example, halogen, alkyl, aralkyl, alkenyl, alkynyl, cycloalkyl, hydroxyl, amino, nitro, sulfhydryl, imino,

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amido, phosphate, phosphonate, phosphinate, carbonyl, carboxyl, silyl, ether, alkylthio, sulfonyl, ketone, aldehyde, ester, a heterocyclyl, an aromatic or heteroaromatic moiety, —CF₃, —CN, or the like.

The term "carbonyl" is art-recognized and includes such moieties as can be represented by the general formula:



wherein X is a bond or represents an oxygen or a sulfur, and R and R' are as defined above. Where X is an oxygen and R or R' is not hydrogen, the formula represents an "ester". Where X is an oxygen and R is as defined above, the moiety is referred to herein as a carboxyl group, and particularly when R is a hydrogen, the formula represents a "carboxylic acid." Where X is an oxygen and R' is hydrogen, the formula represents a "formate." In general, where the oxygen atom of the above formula is replaced by sulfur, the formula represents a "thiocarbonyl" group. Where X is a sulfur and R or R' is not hydrogen, the formula represents a "thioester." Where X is a sulfur and R is hydrogen, the formula represents a "thiocarboxylic acid." Where X is a sulfur and R' is hydrogen, the formula represents a "thioformate." On the other hand, where X is a bond, and R is not hydrogen, the above formula represents a "ketone" group. Where X is a bond, and R is hydrogen, the above formula represents an "aldehyde" group.

As used herein, "heteroatom" as used herein can refer to an atom of any element other than carbon or hydrogen. Exemplary heteroatoms include, but are not limited to, boron, nitrogen, oxygen, phosphorus, sulfur, silicon, arsenic, and selenium.

As used herein, "nitro" can refer to —NO₂; the term "halogen" designates —F, —Cl, —Br, or —I; the term "sulfhydryl" refers to —SH; the term "hydroxyl" refers to —OH; and the term "sulfonyl" refers to —SO₂—.

The term "substituted" as used herein, can refer to all permissible substituents of the compounds described herein. In the broadest sense, the permissible substituents include acyclic and cyclic, branched and unbranched, carbocyclic and heterocyclic, aromatic and nonaromatic substituents of organic compounds. Illustrative substituents include, but are not limited to, halogens, hydroxyl groups, or any other organic groupings containing any number of carbon atoms, e.g. 1-14 carbon atoms, and optionally include one or more heteroatoms such as oxygen, sulfur, or nitrogen grouping in linear, branched, or cyclic structural formats. Representative substituents include alkyl, substituted alkyl, alkenyl, substituted alkenyl, alkynyl, substituted alkynyl, phenyl, substituted phenyl, aryl, substituted aryl, heteroaryl, substituted heteroaryl, halo, hydroxyl, alkoxy, substituted alkoxy, phenoxy, substituted phenoxy, aroxy, substituted aroxy, alkylthio, substituted alkylthio, phenylthio, substituted phenylthio, arylthio, substituted arylthio, cyano, isocyano, substituted isocyano, carbonyl, substituted carbonyl, carboxyl, substituted carboxyl, amino, substituted amino, amido, substituted amido, sulfonyl, substituted sulfonyl, sulfonic acid, phosphoryl, substituted phosphoryl, phosphonyl, substituted phosphonyl, polyaryl, substituted polyaryl, C₃-C₂₀ cyclic, substituted C₃-C₂₀ cyclic, heterocyclic, substituted heterocyclic, amino acid, peptide, and polypeptide groups.

Heteroatoms, such as nitrogen, can have hydrogen substituents and/or any permissible substituents of organic

compounds described herein which satisfy the valences of the heteroatoms. It is understood that “substitution” or “substituted” includes the implicit proviso that such substitution is in accordance with permitted valence of the substituted atom and the substituent, and that the substitution results in a stable compound, i.e., a compound that does not spontaneously undergo transformation such as by rearrangement, cyclization, elimination, etc.

As used herein, “suitable substituent” can refer to a chemically and pharmaceutically acceptable group, i.e., a moiety that does not significantly interfere with the preparation of or negate the efficacy of the inventive compounds. Such suitable substituents may be routinely chosen by those skilled in the art. Suitable substituents include but are not limited to the following: a halo, C₁-C₆ alkyl, C₂-C₆ alkenyl, C₁-C₆ haloalkyl, C₁-C₆ alkoxy, C₁-C₆ haloalkoxy, C₂-C₆ alkynyl, C₃-C₈ cycloalkenyl, (C₃-C₈ cycloalkyl)C₁-C₆ alkyl, (C₃-C₈ cycloalkyl)C₂-C₆ alkenyl, (C₃-C₈ cycloalkyl)C₁-C₆ alkoxy, C₃-C₇ heterocycloalkyl, (C₃-C₇ heterocycloalkyl)C₁-C₆ alkyl, (C₃-C₇ heterocycloalkyl)C₂-C₆ alkenyl, (C₃-C₇ heterocycloalkyl)C₁-C₆ alkoxy, hydroxy, carboxy, oxo, sulfanyl, C₁-C₆ alkylsulfanyl, aryl, heteroaryl, aryloxy, heteroaryloxy, arylalkyl, heteroaralkyl, arylalkoxy, heteroaralkoxy, nitro, cyano, amino, C₁-C₆ alkylamino, di-(C₁-C₆ alkyl)amino, carbamoyl, (C₁-C₆ alkyl)carbonyl, (C₁-C₆ alkoxy)carbonyl, (C₁-C₆ alkyl)aminocarbonyl, di-(C₁-C₆ alkyl)aminocarbonyl, arylcarbonyl, aryloxycarbonyl, (C₁-C₆ alkyl)sulfonyl, and arylsulfonyl. The groups listed above as suitable substituents are as defined hereinafter except that a suitable substituent may not be further optionally substituted.

As used herein, “optionally substituted” can indicate that a group may be unsubstituted or substituted with one or more substituents as defined herein.

As used herein, the term “cardiovascular disease” can refer to diseases of large arteries (atherosclerosis). Unless otherwise defined herein, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art.

Discussion

Sortilin is a membrane glycoprotein that is ubiquitously expressed in many tissues and is most abundant in the central nervous system. Sortilin is involved with protein transport between various cell organelles and can be involved in many biological processes. Therefore, influencing the ability of sortilin to bind its ligands can influence the activity of those ligands. For example, PCSK9 is an enzyme that is ubiquitously expressed in many tissues and cell types and binds to the LDLR. PCSK9 is a ligand for sortilin, where sortilin acts as a secretory chaperone for PCSK9. Thus, sortilin can influence the activity of PCSK9, and amount of the LDLR. Currently, only one ligand binding site on sortilin is known. Sortilin therefore is currently an unexplored potential target for the development of compounds, e.g. small molecules, which are capable of binding sortilin for pharmaceutical development for a variety of diseases.

With that said, described herein are small molecule compounds and formulations thereof that can be capable of binding a first ligand binding site (Site 1) on sortilin or analogue thereof as defined by arginine 325 of SEQ ID NO: 1, or a newly observed second ligand binding site (Site 2) on sortilin or an analogue thereof. Sortilin Site 2 is defined by residues corresponding to SEQ ID NO: 1 present in FIG. 3B: where actual residues of sortilin are 33 residues in number greater than the number provided in the ligand interaction diagram. For example, glutamic acid 448 in the ligand interaction diagram corresponds to glutamic acid 481 of hsortilin and glutamic acid 542 in the ligand interaction diagram of FIG. 3B corresponds to glutamic acid 575 of hsortilin. Also provided herein are methods of altering ligand binding to sortilin by contacting sortilin or an analogue thereof, in vitro or in vivo, with a compound or formulation provided herein. Also provided herein are methods of altering ligand binding to sortilin or an analogue thereof in a subject in need thereof, by administering a compound or formulation thereof provided herein to the subject in need thereof. In some embodiments, the subject in need thereof can be suffering from a disease such as, but not limited to, high cholesterol or Alzheimer's. Other compositions, compounds, methods, features, and advantages of the present disclosure will be or become apparent to one having ordinary skill in the art upon examination of the following drawings, detailed description, and examples. It is intended that all such additional compositions, compounds, methods, features, and advantages be included within this description, and be within the scope of the present disclosure.

Sortilin Binding Compounds and Formulations Thereof

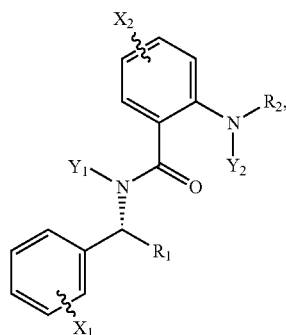
Sortilin Binding Compounds

Provided herein are small molecule compounds that can be capable of binding a ligand binding site on sortilin or an analogue thereof (collectively referred to as “sortilin binding compounds”). Sortilin is a member of the VPS10 family of proteins, which is a family of proteins that share the feature of a VPS10 domain and includes, but is not limited to, Sortilin, SorLA, SorCS1, SorCS2, and SorCS3 (Hampe et al. (2001) Hum. Genet. 108(6):529-536). As used herein, analogues of sortilin include proteins in the VPS10 family as well as structural analogues thereof that contain a binding site similar to the sortilin binding site 1 and/or 2 as described herein. In some embodiments, the sortilin analogue can have an amino acid sequence that can be 70, 75, 80, 85, 90, 95, 97, 98, 99, or about 100% identical to a human sortilin (SEQ ID NOs.: 1-3). In some embodiments, the sortilin analogue is SorLA or a protein having an amino acid sequence that can be 70, 75, 80, 85, 90, 95, 97, 98, 99, or about 100% identical to a human SorLA (SEQ ID NO.: 4).

Previously it was known that sortilin had a first ligand binding site (Site 1). As provided in the Examples below, a second site was characterized. Similar sites to sortilin Site 1 capable of binding at least some of the same compounds as sortilin Site 1 have been identified in analogues of sortilin (e.g. SorLA) (Kitago et al. (2015) Nat. Struct. Mol. Biol. 22:199-206). Provided herein are compounds that can bind to a binding site on sortilin or an analogue thereof. In some embodiments, the binding site can be Site 1 or a structural analogue thereof on sortilin or analogue thereof. In some embodiments, the binding site can be Site 2 or a structural analogue thereof on sortilin or an analogue thereof.

The compound that can bind sortilin or analogue thereof can have a structure according to Formula 1

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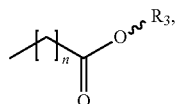
where X_1 and X_2 can each be independently selected from the group of: H, a C_1 to C_5 alkyl, a C_1 to C_5 cyclic alkyl that can be optionally further substituted with a C_1 to C_5 alkyl and where at least two carbons of the C_1 to C_5 cyclic alkyl are members of the benzene ring of Formula 1, an acyl, an amino, a sulfono, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof,

where Y_1 can be selected from the group of: H, a carboxylic acid or ester thereof, where the ester can be an alkyl ester,

where Y_2 can be selected from the group of: H, a carboxylic acid or ester thereof, where the ester can be an alkyl ester,

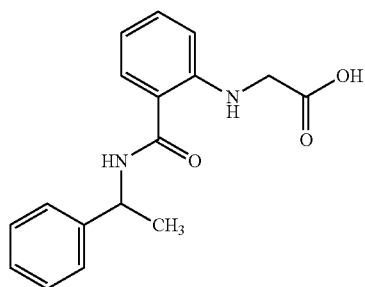
where R_1 can be a C_1 - C_5 alkyl or a C_1 - C_5 alkene,

where R_2 can be



where n can be 1-3 with each carbon being optionally substituted with a C_1 - C_5 alkyl and R_3 can be H or a C_1 to C_5 alkyl, or where R_2 , together with Y_2 , can form a tetrazole.

In some embodiments, the compound can have a structure according to Formula 1A



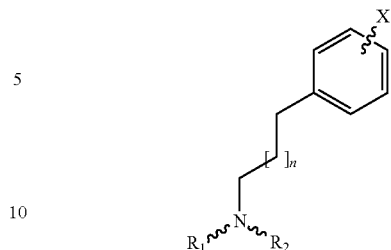
In some embodiments, the compounds according to Formula 1 and/or Formula 1A can bind Site 1 or a structural analogue thereof on sortilin or analogue thereof. The compound of Formula 1 or 1A can be further optionally substituted with a suitable substituent as desired and as is energetically permissible.

The compound that can bind sortilin or analogue thereof can have a structure according to Formula 2

Formula 1

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Formula 2

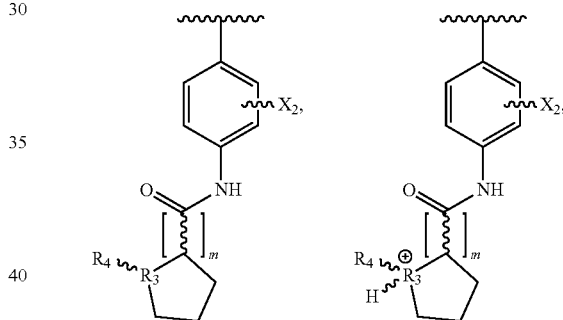


where n can be 1-10,

where X_1 can be selected from the group of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of Formula 2, an acyl, an amino, a sulfono, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof,

where R_1 can be a carboxylic acid or an ester

where R_2 can be a 5, 6, or 7 member heterocyclic ring that can be optionally further substituted with a suitable substituent, a sulfonium, a quaternary ammonium,



where X_2 can be selected from the group of: H, a C_1 - C_5 unsubstituted or substituted alkyl, a C_1 - C_5 cyclic alkyl where at least 2 carbon atoms of the C_1 - C_5 cyclic alkyl are members of the benzene ring of Formula 2, an acyl, an amino, a sulfono, a chloro, a bromo, a iodo, a fluoro, and permissible combinations thereof,

where R_3 can be N or S,

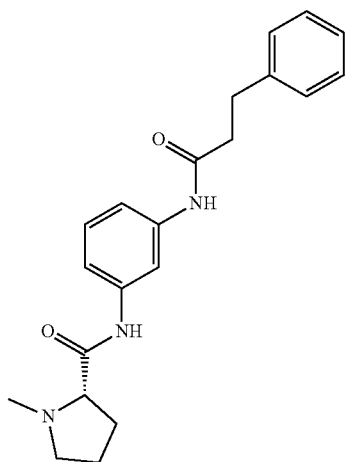
where R_4 can be H, a C_1 - C_{10} alkyl, an amino, a carboxylic acid, an ester, an amino acid or positively charged derivative thereof, and

where m represents the number of carbon atoms in an alkyl, where m can range from 0-10.

In some embodiments, the 5, 6, or 7 member heterocyclic ring that can be optionally substituted with a suitable substituent can be a pyridine, a pyrazole, an imidazole, a pyrrole, a pyrazine, a pyradiazine, a pyrimidine, or a thiazole. The 5, 6, or 7 member heterocyclic ring that can be optionally substituted with a suitable substituent can be positively charged. In some embodiments, the R_4 amino acid can be lysine, arginine, histidine, or a derivative thereof.

The compound that can bind sortilin or analogue thereof can have a structure according to Formula 2A

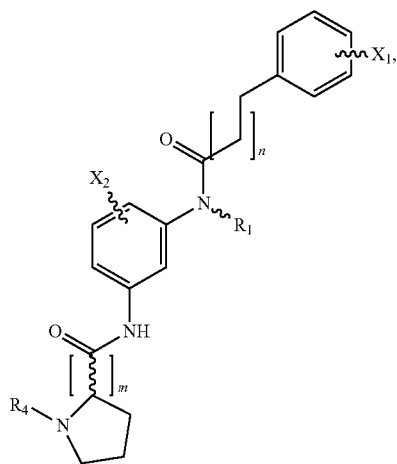
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Formula 2A

In some embodiments, the compounds according to Formula 2 and/or Formula 2A can bind Site 2 or a structural analogue thereof on sortilin or analogue thereof.

In some embodiments, the compound can have a structure according to Formula 2B,



Formula 2B

where X1 and X2 can each be independently selected from the group of: H, a C1-C5 unsubstituted or substituted alkyl, a C1-C5 cyclic alkyl where at least 2 carbon atoms of the C1-C5 cyclic alkyl are members of the benzene ring of Formula 2B, an acyl, an amino, a sulfono, a chloro, a bromo, an iodo, a fluoro, and permissible combinations thereof,

where R1 can be a carboxylic acid or an ester,

where R4 can be H, an C₁-C₁₀ alkyl, an amino, a carboxylic acid, an ester, an amino acid or positively charged derivative thereof,

where n can be 1-10, and where m represents the number of carbon atoms in an alkyl, where m can range from 0-10.

In some embodiments of a compound having a structure according to Formula 2B, the R₄ amino acid can be lysine, arginine, histidine, or a derivative thereof.

The compound of Formula 2, 2B, and/or 2B can be further optionally substituted with a suitable substituent as desired and as is energetically permissible.

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In some embodiments, the compound can be a derivative, including but not limited to, an active derivative of any one of the compounds having a structure according to Formulas 1, 1A, 2, 2A and/or 2B.

The compounds provided herein can compete with binding of a ligand (natural or synthetic) to sortilin or analogue thereof. In some embodiments, the ligand of sortilin or analogue thereof can be PCSK9, apolipoprotein B100, phosphatidylinositides, anionic phospholipids, anionic phospholipids contained on VLDL, amyloid beta, amyloid precursor protein, neurotensin or analogues of neurotensin, lipoprotein lipase, apolipoprotein AV and/or apolipoprotein E. The sortilin binding compounds provided herein can be capable of altering the amount of LDLR present. The sortilin binding compounds can be capable of reducing circulating levels of LDL and/or VLDL in the blood of a subject. The sortilin binding compounds provided herein can be capable of reducing or eliminating interactions of AB particles (e.g. AP-AB₁₋₃₂) in neuronal cells.

The sortilin binding compounds provided herein can be made using techniques and methods generally known to the skilled artisan in view of this disclosure. Such synthesis schemes are within the scope of this disclosure.

Pharmaceutical Formulations

Also provided herein are pharmaceutical formulations that can contain an amount of a compound capable of binding sortilin or an analogue thereof (collectively also referred to as "sortilin binding compounds") as provided elsewhere herein. The sortilin binding compounds described herein can be provided to a subject in need thereof alone or as such as an active ingredient, in a pharmaceutical formulation. In some embodiments, the pharmaceutical formulations contain an effective amount of a sortilin binding compounds. The pharmaceutical formulations described herein can be administered to a subject in need thereof. The subject in need thereof can have a disease or disorder whose pathology involves sortilin or an analogue thereof and/or a ligand of sortilin or an analogue thereof. In some embodiments, the subject can be a human. The term pharmaceutical formulation also encompasses pharmaceutically acceptable salts of the pharmaceutical formulations and/or active ingredients provided herein.

Pharmaceutically Acceptable Carriers and Auxiliary Ingredients and Agents

The pharmaceutical formulations containing an effective amount of a sortilin binding compound described herein can further include a pharmaceutically acceptable carrier. Suitable pharmaceutically acceptable carriers include, but are not limited to, water, salt solutions, alcohols, gum arabic, vegetable oils, benzyl alcohols, polyethylene glycols, gelatin, carbohydrates such as lactose, amylose or starch, magnesium stearate, talc, silicic acid, viscous paraffin, perfume oil, fatty acid esters, hydroxy methylcellulose, and polyvinyl pyrrolidone, which do not deleteriously react with the active composition.

The pharmaceutical formulations can be sterilized, and if desired, mixed with auxiliary agents, such as lubricants, preservatives, stabilizers, wetting agents, emulsifiers, salts for influencing osmotic pressure, buffers, coloring, flavoring and/or aromatic substances, and the like which do not deleteriously react with the active composition.

In addition to the effective amount of a sortilin binding compound described herein, the pharmaceutical formulation can also optionally include an effective amount of an auxiliary active agent, including but not limited to, DNA, RNA, amino acids, peptides, polypeptides, antibodies, aptamers, ribozymes, guide sequences for ribozymes that inhibit trans-

lation or transcription of essential tumor proteins and genes, hormones, immunomodulators, antipyretics, anxiolytics, antipsychotics, analgesics, antispasmodics, anti-inflammatories, anti-histamines, anti-infectives, chemotherapeutics, antihypertensives, anticoagulants, and antiarrhythmics.

Effective Amounts of the Sortilin Binding Compounds and Auxiliary Agents

The pharmaceutical formulations can contain an effective amount of a sortilin binding compound, and optionally, a therapeutically effective amount of an auxiliary agent. In some embodiments, the effective amount of the sortilin binding compound(s) can range from about 0.3 mg/kg body weight to about 30 mg/kg. The effective amount of the sortilin binding compound(s) can range from about 1 mg to about 10 g. For liquid formulations, some embodiments, the effective amount of the sortilin binding compound(s) or pharmaceutical formulation containing a sortilin binding compound(s) can range from about 10 μ L to about 10 mL. One of skill in the art will appreciate that the exact volume will depend on, inter alia, the age and size of the subject, as well as the location of administration. The effective concentration of the sortilin binding compound(s) can range from about 1 nM to 1M.

In embodiments where an optional auxiliary active agent is included in the pharmaceutical formulation, the therapeutically effective amount of the auxiliary active agent will vary depending on the auxiliary active agent. In some embodiments, the therapeutically effective amount of the optional auxiliary active agent can range from 0.001 micrograms to about 1000 milligram. In other embodiments, the therapeutically effective amount of the optional auxiliary active agent can range from about 0.01 IU to about 1000 IU. In further embodiments, the therapeutically effective amount of the auxiliary active agent can range from 0.001 mL to about 1 mL. In yet other embodiments, the therapeutically effective amount of the optional auxiliary active agent can range from about 1% w/w to about 50% w/w of the total pharmaceutical formulation. In additional embodiments, the therapeutically effective amount of the optional auxiliary active agent ranges from about 1% v/v to about 50% v/v of the total pharmaceutical formulation. In still other embodiments, the therapeutically effective amount of the optional auxiliary active agent ranges from about 1% w/v to about 50% w/v of the total pharmaceutical formulation.

Dosage Forms

In some embodiments, the pharmaceutical formulations described herein can be in a dosage form. The dosage forms can be adapted for administration by any appropriate route. Appropriate routes include, but are not limited to, oral (including buccal or sublingual), rectal, epidural, intracranial, intraocular, inhaled, intranasal, topical (including buccal, sublingual, or transdermal), vaginal, intraurethral, parenteral, intracranial, subcutaneous, intramuscular, intravenous, intraperitoneal, intradermal, intraosseous, intracardiac, intraarticular, intracavernous, intrathecal, intravireal, intracerebral, and intracerebroventricular and intradermal. Such formulations can be prepared by any method known in the art.

Dosage forms adapted for oral administration can be discrete dosage units such as capsules, pellets or tablets, powders or granules, solutions, or suspensions in aqueous or non-aqueous liquids; edible foams or whips, or in oil-in-water liquid emulsions or water-in-oil liquid emulsions. In some embodiments, the pharmaceutical formulations adapted for oral administration also include one or more agents which flavor, preserve, color, or help disperse the pharmaceutical formulation. Dosage forms prepared for oral

administration can also be in the form of a liquid solution that can be delivered as foam, spray, or liquid solution. In some embodiments, the oral dosage form can contain about 10 mg to 10 g of a pharmaceutical formulation containing an effective amount or an appropriate fraction thereof of the sortilin binding compound(s). The oral dosage form can be administered to a subject in need thereof by a suitable administration method.

Where appropriate, the dosage forms described herein can be microencapsulated. The dosage form can also be prepared to prolong or sustain the release of any ingredient. In some embodiments, the sortilin binding compound(s) can be the ingredient whose release is delayed. In other embodiments, the release of an optionally included auxiliary ingredient is delayed. Suitable methods for delaying the release of an ingredient include, but are not limited to, coating or embedding the ingredients in material in polymers, wax, gels, and the like. Delayed release dosage formulations can be prepared as described in standard references such as "Pharmaceutical dosage form tablets," eds. Liberman et. al. (New York, Marcel Dekker, Inc., 1989), "Remington—The science and practice of pharmacy", 20th ed., Lippincott Williams & Wilkins, Baltimore, Md., 2000, and "Pharmaceutical dosage forms and drug delivery systems", 6th Edition, Ansel et al., (Media, Pa.: Williams and Wilkins, 1995). These references provide information on excipients, materials, equipment, and processes for preparing tablets and capsules and delayed release dosage forms of tablets and pellets, capsules, and granules. The delayed release can be anywhere from about an hour to about 3 months or more.

Examples of suitable coating materials include, but are not limited to, cellulose polymers such as cellulose acetate phthalate, hydroxypropyl cellulose, hydroxypropyl methylcellulose, hydroxypropyl methylcellulose phthalate, and hydroxypropyl methylcellulose acetate succinate; polyvinyl acetate phthalate, acrylic acid polymers and copolymers, and methacrylic resins that are commercially available under the trade name EUDRAGIT® (Roth Pharma, Westerstadt, Germany), zein, shellac, and polysaccharides.

Coatings may be formed with a different ratio of water soluble polymer, water insoluble polymers, and/or pH dependent polymers, with or without water insoluble/water soluble non polymeric excipient, to produce the desired release profile. The coating is either performed on the dosage form (matrix or simple) which includes, but is not limited to, tablets (compressed with or without coated beads), capsules (with or without coated beads), beads, particle compositions, "ingredient as is" formulated as, but not limited to, suspension form or as a sprinkle dosage form.

Dosage forms adapted for topical administration can be formulated as ointments, creams, suspensions, lotions, powders, solutions, pastes, gels, sprays, aerosols, or oils. In some embodiments for treatments of the eye or other external tissues, for example the mouth or the skin, the pharmaceutical formulations are applied as a topical ointment or cream. When formulated in an ointment, the sortilin binding compound(s), optional auxiliary active ingredient, and/or pharmaceutically acceptable salt thereof can be formulated with a paraffinic or water-miscible ointment base. In other embodiments, the active ingredient can be formulated in a cream with an oil-in-water cream base or a water-in-oil base. Dosage forms adapted for topical administration in the mouth include lozenges, pastilles, and mouth washes.

Dosage forms adapted for nasal or inhalation administration include aerosols, solutions, suspension drops, gels, or dry powders. In some embodiments, the sortilin binding compound(s), the composition containing the sortilin bind-

ing compound(s), auxiliary active ingredient, and/or pharmaceutically acceptable salt thereof in a dosage form adapted for inhalation is in a particle-size-reduced form that is obtained or obtainable by micronization. In some embodiments, the particle size of the size reduced (e.g. micronized) compound or salt or solvate thereof, is defined by a D50 value of about 0.5 to about 10 microns as measured by an appropriate method known in the art. Dosage forms adapted for administration by inhalation also include particle dusts or mists. Suitable dosage forms wherein the carrier or excipient is a liquid for administration as a nasal spray or drops include aqueous or oil solutions/suspensions of an active ingredient, which may be generated by various types of metered dose pressurized aerosols, nebulizers, or insufflators.

In some embodiments, the dosage forms are aerosol formulations suitable for administration by inhalation. In some of these embodiments, the aerosol formulation contains a solution or fine suspension of the sortilin binding compound(s) and/or pharmaceutically acceptable salt thereof, and a pharmaceutically acceptable aqueous or non-aqueous solvent. Aerosol formulations can be presented in single or multi-dose quantities in sterile form in a sealed container. For some of these embodiments, the sealed container is a single dose or multi-dose nasal or an aerosol dispenser fitted with a metering valve (e.g. metered dose inhaler), which is intended for disposal once the contents of the container have been exhausted.

Where the aerosol dosage form is contained in an aerosol dispenser, the dispenser contains a suitable propellant under pressure, such as compressed air, carbon dioxide, or an organic propellant, including but not limited to a hydrofluorocarbon. The aerosol formulation dosage forms in other embodiments are contained in a pump-atomizer. The pressurized aerosol formulation can also contain a solution or a suspension of the sortilin binding compound(s) or a pharmaceutical formulation thereof. In further embodiments, the aerosol formulation also contains co-solvents and/or modifiers incorporated to improve, for example, the stability and/or taste and/or fine particle mass characteristics (amount and/or profile) of the formulation. Administration of the aerosol formulation can be once daily or several times daily, for example 2, 3, 4, or 8 times daily, in which 1, 2, or 3 doses or more are delivered each time.

For some dosage forms suitable and/or adapted for inhaled administration, the pharmaceutical formulation is a dry powder inhalable formulation. In addition to the sortilin binding compound(s), an optional auxiliary active ingredient, and/or pharmaceutically acceptable salt thereof, such a dosage form can contain a powder base such as lactose, glucose, trehalose, mannitol, and/or starch. In some of these embodiments, the sortilin binding compound(s), optional auxiliary active ingredient, and/or pharmaceutically acceptable salt thereof is in a particle-size reduced form. In further embodiments, a performance modifier, such as L-leucine or another amino acid, cellobiose octaacetate, and/or metals salts of stearic acid, such as magnesium or calcium stearate.

In some embodiments, the aerosol formulations are arranged so that each metered dose of aerosol contains a predetermined amount of an active ingredient, such as the one or more of the sortilin binding compound(s) described herein.

Dosage forms adapted for vaginal administration can be presented as pessaries, tampons, creams, gels, pastes, foams, or spray formulations. Dosage forms adapted for rectal administration include suppositories or enemas.

Dosage forms adapted for parenteral administration and/or adapted for any type of injection (e.g. intravenous, intraocular, intraperitoneal, subcutaneous, intramuscular, intradermal, intraosseous, epidural, intracardiac, intraarticular, intracavernous, intrathecal, intravitreal, intracerebral, and intracerebroventricular) can include aqueous and/or non-aqueous sterile injection solutions, which can contain anti-oxidants, buffers, bacteriostats, solutes that render the composition isotonic with the blood of the subject, and aqueous and non-aqueous sterile suspensions, which can include suspending agents and thickening agents. The dosage forms adapted for parenteral administration can be presented in a single-unit dose or multi-unit dose containers, including but not limited to sealed ampoules or vials. The doses can be lyophilized and resuspended in a sterile carrier to reconstitute the dose prior to administration. Extemporaneous injection solutions and suspensions can be prepared in some embodiments, from sterile powders, granules, and tablets.

Dosage forms adapted for ocular administration can include aqueous and/or non-aqueous sterile solutions that can optionally be adapted for injection, and which can optionally contain anti-oxidants, buffers, bacteriostats, solutes that render the composition isotonic with the eye or fluid contained therein or around the eye of the subject, and aqueous and non-aqueous sterile suspensions, which can include suspending agents and thickening agents.

For some embodiments, the dosage form contains a predetermined amount of the sortilin binding compound(s) per unit dose. In an embodiment, the predetermined amount of the sortilin binding compound(s) is an effective amount of the sortilin binding compound(s). In other embodiments, the predetermined amount of the sortilin binding compound(s) can be an appropriate fraction of the effective amount of the active ingredient. Such unit doses may therefore be administered once or more than once a day. Such pharmaceutical formulations may be prepared by any of the methods well known in the art.

Methods of Using the Sortilin Binding Compounds and Formulations Thereof

The sortilin binding compounds, derivatives, and formulations thereof provided herein can be used to bind sortilin or an analogue thereof, competitively inhibit a ligand (native or synthetic) of sortilin or an analogue thereof, and/or treat and/or prevent a disease whose pathology involves a ligand of sortilin or an analogue thereof. In some embodiments, the disease can be cholesteremia or Alzheimer's disease. In some embodiments, the ligand of sortilin or analogue thereof can be PCSK9, apolipoprotein B100, phosphatidylinositides, anionic phospholipids, anionic phospholipids contained on VLDL, amyloid beta, amyloid precursor protein, neurotensin or analogues of neurotensin, lipoprotein lipase, apolipoprotein AV and/or apolipoprotein E.

Provided herein are methods of inhibiting binding of a ligand to sortilin or analogue thereof that can include the step of contacting sortilin or an analogue thereof with a sortilin binding compound as provided herein. The step of contacting can occur in vitro or in vivo. The method can further include the step of determining binding of the sortilin binding compound to the sortilin or analogue thereof. One of ordinary skill in the art will appreciate ways that binding of sortilin or analogue thereof can be detected and measured.

Also provided herein are methods of inhibiting binding of a ligand to sortilin or analogue thereof and/or methods of treating and/or preventing a disease in a subject in need thereof that can include the step of administering an amount, such as an effective amount, of one or more sortilin binding

compounds to the subject. The subject can be suffering from a disease whose pathology involves a ligand of sortilin or an analogue thereof. In some embodiments, the subject in need thereof suffers from cholesterolemia and/or Alzheimer's disease. The method can include the step of administering an amount, such as an effective amount of one or more sortilin binding compounds to a subject such that the binding of PCSK9, apolipoprotein B100, phosphatidylinositides, anionic phospholipids, anionic phospholipids contained on VLDL, amyloid beta, amyloid precursor protein, neurotensin or analogues of neurotensin, lipoprotein lipase, apolipoprotein AV and/or apolipoprotein E to sortilin or analogue thereof is reduced. In some embodiments, the effective amount of the sortilin binding compound(s) administered to the subject in need thereof can inhibit binding of a ligand of sortilin or analogue thereof (including, but not limited to, PCSK9, apolipoprotein B100, phosphatidylinositides, anionic phospholipids, anionic phospholipids contained on VLDL, amyloid beta, amyloid precursor protein, neurotensin or analogues of neurotensin, lipoprotein lipase, apolipoprotein AV and apolipoprotein E.) by 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 97, 98, 99, and/or about 100%.

In some embodiments, the sortilin binding compounds can be used as a treatment for a cardiovascular disease. Cardiovascular diseases can occur in high numbers in patients with diabetes and lipoprotein disorders associated with diabetes. This diabetic dyslipidemia can be related to high levels of circulating triglyceride that are carried by very low density lipoproteins (VLDL-B100) which is a precursor of LDL in the circulating. Disorders of VLDL metabolism can also result in accumulation of triglyceride in the liver, a condition known as hepatic steatosis, which can lead to cirrhosis requiring liver transplantation. Hepatic steatosis can also occur in obesity including childhood obesity that is also associated with metabolic syndrome, a component of which, is high levels of blood triglyceride. Compounds having a structure according to formula 2, e.g. Formula 2A, can be a therapeutic agent for reducing hepatic steatosis and cardiovascular disease risk in obese insulin resistant and type 2 diabetic patients. Compounds having a structure according to formula 2, e.g. Formula 2A, can increase hepatic secretion of triglyceride contained in VLDL-B100 particles thereby reducing the liver content of triglyceride. The additional action of compounds having a structure according to formula 2, e.g. Formula 2A, of increasing LDL receptors through competition of compounds having a structure according to formula 2, e.g. Formula 2A, with PCSK9 at Site 2 of sortilin would ameliorate the high triglyceride levels by increasing the clearance of LDL cholesterol through LDL receptors. As compounds having a structure according to formula 2, e.g. Formula 2A, also increases PIP3 binding to site 1, compounds having a structure according to formula 2, e.g. Formula 2A, could also facilitate VLDL clearance by liver through interaction of sortilin at the cell surface.

In some embodiments, the sortilin binding compounds can be used to treat hypercholesterolemia in a subject. Patients with hypercholesterolemia, either acquired or genetic, are typically treated with "statins". Familial hypercholesterolemia (genetic), which is resistant to therapy, has been treated in clinical trials with a combination of "statins" and a monoclonal antibody based therapy to reduce circulating PCSK9 levels. The use of compounds having a structure according to Formula 2, e.g. Formula 2A, in all of these patients can therefore provide an alternative approach to lowering LDL cholesterol. The action of compounds having

a structure according to Formula 2, e.g. Formula 2A, can be to compete for binding to sortilin with PCSK9 blocking sortilin's chaperone function in secretion. Reduced secretion of PCSK9 can have a similar effect as the documented reduction of LDL cholesterol with the monoclonal antibody derivatives. In addition, compounds having a structure according to Formula 2, e.g. Formula 2A, can have a favorable effect on circulating VLDL-B100 by favoring cellular uptake of PIP3 containing VLDL by surface expressed sortilin receptors. The predominant effect of compounds having a structure according to Formula 2, e.g. Formula 2A, can be in liver where the majority of LDL receptors are expressed and mechanisms are available for disposing of the form of cholesterol taken up by LDL receptors and sortilin receptor.

In some embodiments, the sortilin binding compounds can be used to treat Alzheimer's disease in a subject. Alzheimer's disease is associated with accumulation of amyloid beta in neurons. A necessary pathway for removal of amyloid beta involves apolipoprotein E, a ligand that binds site 1 of sortilin. Increasing removal of amyloid beta and factors associated with amyloid beta including alpha beta dimer can interact with apo E in their removal. Compounds having a structure according to Formula 2, e.g. Formula 2A, can facilitate ligand interaction with sortilin at site 1 and should through interaction with apo E increase clearance of plaque components reducing symptoms. Sortilin is known to be associated with autophagic lysosomal degradation which would be a pathway for eliminating these toxic protein complexes.

In some embodiments, the sortilin binding compounds provided herein that can bind site 2 of sortilin or an analogue thereof can, via binding at site 2 on sortilin or analogue thereof, can alter binding of a compound to site 1 of sortilin. In some embodiments, a sortilin binding compound provided herein can, via binding at site 2 on sortilin or analogue thereof, enhance or increase binding of a compound to site 1 of sortilin or analogue thereof. In some embodiments, the sortilin binding compounds capable of binding site 2 of sortilin or analogue thereof can be used to alter (e.g. increase or enhance) the binding of a compound to site 1 of sortilin.

In some embodiments, the sortilin binding compounds provide herein can be used to alter the rate and/or amount degradation of one or more intracellular proteins. The interaction of sortilin with a ligand can lead to lysosomal degradation of intracellular proteins via autophagy. This can result in alterations of proteins secreted by the cell. In some embodiments, VLDL secretion can be altered in this manner.

Sortilin can be expressed on the plasma membrane of cells as well as on the membrane of organelles (e.g. the Golgi). By competing with endogenous ligands for sortilin expressed on the plasma membrane, the sortilin binding compounds provided herein can be capable of altering the interaction of plasma membrane expressed sortilin with circulating lipoproteins.

EXAMPLES

Now having described the embodiments of the present disclosure, in general, the following Examples describe some additional embodiments of the present disclosure. While embodiments of the present disclosure are described in connection with the following examples and the corresponding text and figures, there is no intent to limit embodiments of the present disclosure to this description. On the contrary, the intent is to cover all alternatives, modifications,

and equivalents included within the spirit and scope of embodiments of the present disclosure.

Example 1

Introduction

The role of sortilin in VLDL-B100 secretion is controversial [1], and inconsistencies have been summarized [2,3]. Some studies suggest sortilin serves as a chaperone where sortilin is hypothesized to facilitate secretion of VLDL-B100 [4]. Other studies demonstrate that sortilin inhibits VLDL-B100 secretion by trafficking towards degradation [5]. An obligatory inverse relationship is hypothesized to result in VLDL-B100 hypersecretion when sortilin is reduced under conditions of severe insulin resistance [6]. Understanding sortilin function is important considering that sortilin is involved in LDL metabolism [7], and is associated with cardiovascular disease [8]. Studies using surface plasmon resonance (SPR) indicate that human VLDL and LDL, both B100-containing lipoproteins, bind to sortilin [4,5]. We directly demonstrated enhanced binding of B100 to sortilin with insulin in insulin responsive McArdle RH7777 (McA) cells [9] suggesting the importance of insulin sensitivity in B100 sortilin interaction. Accordingly, the observed inconsistencies in the relationship between sortilin and VLDL-B100 secretion may relate to insulin sensitivity. Culture conditions of McA cells are known to affect insulin signaling and VLDL secretion, and under serum-enriched conditions McA cells are insulin resistant [10]. Using this model we have explored the relationship of cellular sortilin and VLDL-B100 secretion in insulin sensitive McA cells. Under these defined conditions, we show a role for sortilin in facilitating VLDL-B100 secretion. In contrast, in insulin resistant McA cells, VLDL-B100 secretion is relatively independent of sortilin. Results demonstrate the complexity of sortilin function and the importance of defining metabolic conditions when examining sortilin and VLDL-B100 relationships.

Materials and Methods

Cell Culture, Materials and Reagents.

McArdle RH-7777 cells (McA cells) were cultured as previously described in serum containing complete Dulbecco's Modified Eagle's Medium (cDMEM) [9, 10]. To induce insulin sensitivity, McA cells at 50-60% confluency were incubated for 18 h in serum-free media containing DMEM containing 1% (w/v) BSA (1% BSA/DMEM). Human sortilin (hsortilin) (Ser78-Asn755) with C-terminal 6-His tag was from R&D Systems, Inc., (Minneapolis, Minn.). Plasma from fasted Sprague Dawley rats (BioreclamationIVT, Westbury, N.Y.) was used to prepare VLDL apo B standards. Mouse monoclonal antibody to rat B100 was prepared in our laboratory and characterized previously [11]. Rabbit anti-sortilin antibody was from GeneTex (GTX54854, Irvine, Calif.) or from Abcam (ab16640, Cambridge, Mass.). Rabbit anti-p-AKT (9271), rabbit anti-AKT (9272), and rabbit anti-insulin receptor β -subunit (IR β , 3025) were from Cell Signaling Technology (Danvers, Mass.). Mouse anti-pY Platinum 4G10 was from EMD Millipore (Temecula, Calif.). (Horseradish peroxidase linked donkey anti-rabbit IgG (NA9340), sheep anti-mouse IgG (NXA931) and ECL Prime Western Blotting Detection Reagent (RPN2232) were from GE Healthcare (Buckinghamshire, UK). All other materials and reagents were essentially as described previously [9]. Compound 98,477,898 (2S)-1-methyl-N-{3-[(3-phenylpropanoyl)-amino]phenyl}pyrrolidine-2-carboxamide (cpd984) was obtained from ChemBridge Corp. (San

Diego, Calif.). A stock solution of cpd984 (10 mM) was prepared in DMSO, and stored in aliquots at -20° C. Knockdown of Sortilin in McArdle Cells Using siRNA.

McA cells were transfected using Fugene6 according to manufacturer's protocol (Promega Corp., Madison Wis.) using three different pGIPZ based vectors expressing shRNAi targeting rat Sort1 mRNA (V2LMM_58553, V3LMM_450660, V3LMM_450662), and one scrambled, non-silencing control (GE Healthcare Dharmacon, Lafayette, Colo.). McA cells with sortilin knockdown (KD) were selected using puromycin. Lysates from each cell line were examined by immunoblotting to assess sortilin expression relative to the non-silencing control cells (SCR).

Immunoblotting.

McA cell lysates were prepared and denatured proteins were separated by SDS-PAGE as described previously [10]. Following electrophoretic transfer to PVDF membranes and blocking non-specific binding, membranes were incubated with primary antibodies overnight at 4° C. in blocking buffer. Antibody binding was detected by incubation for 1-2 h at room temperature with species specific secondary HRP-linked antibodies and developed using Amersham™ Prime reagent (GE Healthcare). Insulin signaling to IR β and AKT were evaluated using nitrocellulose membranes (Bio-Rad) and phosphospecific (pY, p-AKT(Ser473) and mass specific antibodies [12]. Chemiluminescence was measured using the ChemiDocXRS+ system (Bio-Rad) and band intensities quantified using Image Lab 3.0.1 software (Bio-Rad).

RNA Isolation and mRNA Quantitation.

Total RNA was extracted using TriZol Reagent (Life Technologies, Grand Island, N.Y.) and mRNA was measured by quantitative polymerase chain reaction after reverse transcription as previously described [10] by the University of Rochester Genomic Research Center. TaqMan® gene expression primer probe sets used for fluorogenic quantification of rat mRNA transcripts were: apoB, Rn01499054; Sort1, Rn01521847, Atf3, Rn00563784 and Rplp0 (ARBP, 36B4), Rn00821065.

Immuno Slot Blotting.

Experimental media were adjusted to 1% (v/v) protease inhibitor cocktail I (EMD Millipore) and to a salt density of 1.019 g/ml by addition of a solution of NaBr (d=1.495 g/ml). VLDL was isolated by ultracentrifugation in a L-70 Ultracentrifuge (Beckman Coulter, Inc., Fullerton, Calif.) using a 50 Ti rotor (200,000 \times g, 18 h, 14° C.). Following centrifugation, the top 1.5-2.0 ml VLDL fraction was removed using a syringe and weighed to determine volume. VLDL samples were applied in triplicate wells (0.2-0.4 ml per well) in a Bio-Dot® SF apparatus (Bio-Rad). Two PVDF membranes were used together for blotting: the top was Immobilon-P (IPVH09120 SF) and the bottom was Immobilon-PSQ (ISEQ09120 SF); both were obtained from EMD Millipore. VLDL apo B standards were prepared from rat plasma VLDL and total apo B (B100 and B48) and B100 content were determined on stained gels following SDS-PAGE separation [13]. VLDL-apo B standards in TBS were slotted in duplicate alongside test samples. After filtration, 0.4 ml of TBS was added as a wash. After final filtration, membranes were air dried, rehydrated in methanol and incubated in TBS then in blocking buffer at 4° C. overnight. At this stage slot blots were evaluated similarly to immunoblotting. After chemiluminescence development and B100 quantitation, slot blots were stripped by incubation in Restore™ PLUS for 15 min at room temperature (Thermo Scientific, Rockford, Ill.) and were reblocked overnight at 4° C. Total apo B (B100 and B48) present in VLDL was then evaluated following incubation with rabbit polyclonal anti-rat apo B. The bottom

PVDF membrane was carried through the entire procedure to assure there was no “bleed through” of test samples. VLDL-apo B and VLDL-B100 content were calculated from standard curves generated by VLDL apo B standards. Recovery of rat VLDL added to unspent media averaged 94%±4.9% with a CV of 5.2% (n=6 replicates).

Surface Plasmon Resonance.

Surface plasmon resonance measurements were performed on a Biacore T200 instrument equipped with Ni-NTA sensor chips with ~6800 response units (RU) for hsortilin covalently immobilized to the surface. HBS-DMSO running buffer (10 mM Hepes pH 7.4, 150 mM NaCl, 1% DMSO) was used at a flow rate of 30 µl/min and injections performed with times for association of 90 s, and dissociation of 300 s, followed by injection of buffer to regenerate the hsortilin surface. Binding was expressed in relative RU; the difference in response between the immobilized protein flow cell and the corresponding control flow cell. NT was administered to the chip containing hsortilin at a concentration of 100 nM. NT at the same concentration of 100 nM was then administered in the presence of cpd984.

Computational Modeling and Compound Screening.

Schrödinger’s Maestro program (version 9.3.5) was used as the primary graphical user interface and Maestro version 10.2 (Schrödinger, LLC, New York, N.Y.) was used for ligand interaction diagramming. Virtual screening was performed on compounds contained in Chem-Bridge libraries (www.chembridge.com) that were prepared with Schrödinger’s LigPrep program [14]. The virtual screening method was performed using Schrödinger’s GLIDE software [15] on the hsortilin crystal structure PDB ID: 4PO7 [16]. Compounds were docked using GLIDE at the site where the N-terminal fragment of NT is found in the crystal structure and cpd984 was chosen for biological screening based on its docking score. Schrödinger’s PRIME software was used to generate missing side chains and loops of this crystal structure predicting the NT peptide spanning the cavity of hsortilin [17]. LigPrep was used on the N-terminal peptide XLYEN-OH from this crystal structure and it was then docked back into its respective site on the crystal structure. This self-docking task was able to reproduce the X-ray pose for this ligand.

Statistics.

Unless noted, results are expressed as the mean±S.E.M., where n equals the number of independent experiments in which replicate analyses were performed in each experiment. Significant differences were assessed using Student’s t-test with p-values≤0.05 being considered significant.

Results and Discussion

Reduced Sortilin Levels Associate with Lower VLDL Apo B Secretion.

Different levels of media accumulation of apo B by McA cells based on culture conditions have been reported [10]. Under serum enriched conditions (cDMEM), McA cells are insensitive to insulin activation of Class IA phosphatidylinositol 3-kinase (PI3K); insulin dependent translocation of activated PI3K to intracellular membranes, and fail to express insulin dependent apo B degradation (IDAD) [10]. However, following 18 h incubations in 1% BSA/DMEM, insulin sensitivity is restored while total medium apo B concentration is reduced [10]. We extended these results to measurement of VLDL apo B, and specifically to VLDL-B100, the major form of apo B secreted by McA cells [18], and the only form secreted by human liver. Using these conditions, we addressed the extent to which VLDL-apo B secretion was altered (FIG. 1A). Media VLDL-B100 was analyzed by immuno slot blotting following isolation by

ultracentrifugation at d<1.019 g/ml (FIG. 1B). Consistent with our previous report, both VLDL-apo B total and VLDL-B100 secretion were significantly reduced in insulin sensitive compared with insulin insensitive McA cells. Secreted VLDL-apo B total (ng/mg/18 h) averaged (mean±SD, n=5): 1595±40 in cDMEM vs 1142±79 in 1% BSA/DMEM. Secreted VLDL-B100 (ng/mg/18 h) averaged (mean±SD, n=5): 1245±208 in cDMEM vs 894±95 in 1% BSA/DMEM. Reduced VLDL secretion was not due to deficiency in fatty acid availability as lipid-enriched BSA was used in the 1% BSA/DMEM medium. Following 1% BSA/DMEM incubation, cellular sortilin levels were significantly depressed (FIG. 1C) averaging only 37%±7% (n=4, p<0.001) of that present in cells incubated in cDMEM. Reduced apo B secretion and cellular sortilin were not the consequence of lowered abundance of mRNA transcripts as Apob and Sort1 mRNA levels were unchanged by culture conditions (FIG. 1D). Considering the suggested role for transcriptional repression of sortilin by the endoplasmic reticulum stress target ATF3 [6], we also measured Atf3 mRNA levels. Atf3 mRNA was reduced on average by 30% in insulin sensitive compared with insensitive McA cells. Decreased ATF3 would be expected to correlate with increased cellular sortilin if there were transcriptional suppression. A corresponding reduction in VLDL-B100 secretion with reduced cellular sortilin suggests a direct relationship. Results are consistent with a model of genetic deletion of sortilin where VLDL-B100 secretion is also reduced [4].

Sortilin KD in Insulin Sensitive McA Cells Associates with Decreased VLDL-B100 Secretion.

If reduced VLDL-B100 secretion were related to lower cellular sortilin, it would be expected that additional reductions of sortilin would suppress VLDL-B100 even further. To address this premise, three stable cell lines of McA cells using siRNA were prepared with variable KD of sortilin compared with wildtype (WT) and McA cells transfected with scrambled shRNA (SCR) (FIG. 2A). The low (L), medium (M) and high (H) sortilin expressing McA cell lines were incubated in cDMEM, and VLDL-B100 secretion was measured (FIG. 2B). There was no reduction in VLDL-B100 secretion observed with progressively reduced sortilin. There was a small but significant increase in VLDL-B100 secretion in the lowest sortilin expressing McA53 cells (L) compared with SCR that averaged 32% (p<0.05). In contrast, in each McA cell line incubated in 1% BSA/DMEM, there was reduced VLDL-B100 secretion (FIG. 2C). Alterations in the relationship between VLDL-B100 secretion and sortilin under differing culture conditions suggest differences in sortilin function with insulin sensitivity.

Very low density lipoprotein (VLDL) whose integral protein is apolipoprotein B (B100) was studied in a mammalian hepatocyte cell line, McArdle RH7777 (McA) cells, and cellular sortilin expression was examined. Liver cells secrete VLDL-B100 into the blood and therefore the VLDL secreted by these cells represent a relevant human liver model to study the relationships of sortilin to VLDL secretion. It is well known that high levels of VLDL-B100 secretion occur in type 2 diabetes influencing levels of plasma triglyceride and cholesterol which may result in accelerated damage to arteries underlying diseases such as atherosclerosis resulting in heart attacks and stroke. In FIG. 1A, McA cells with enriched conditions (cDMEM) and insulin resistance, demonstrate increased VLDL-B100 and VLDL-total apo B secretion as occurs with humans with insulin resistance and type 2 diabetes. In FIG. 1C, under these same conditions, there are dramatic alterations of sortilin indicating sortilin as a potential target for therapeutic

intervention for small molecules as sortilin had been shown to be related to VLDL metabolism. [4] In liver cells sortilin levels can be reduced using molecular techniques involving siRNA, and high (H), middle (M) and low (L) levels of sortilin can be established. Under enriched conditions (cD-MEM) there is little relationship between cellular sortilin and VLDL-B100 secretion (FIG. 2B).

Identification of a Small Molecule that Binds the N-Terminal NT Binding Site on Hsortilin.

The complexity of sortilin function as a multi-ligand receptor may be related to different sites of ligand binding which direct different functions of sortilin. High resolution crystal structures of hsortilin have been reported, and a C-terminal NT binding site forming a salt bridge between the C-terminal carboxylate of NT and arginine 292 (R292) of the crystal structure has been identified (FIG. 3A) [16]. Small molecules that bind to this site have been characterized [19]. The N-terminus of NT (pyroGlu-Leu-Tyr-Glu-Asn) also interacts with hsortilin [16]. Using Schrödinger software, we explored the N-terminal NT binding site screening compounds from the ChemBridge library and further characterized NT fragments present in the hsortilin crystal structure PDB ID: 4PO7 (FIG. 3A) [16]. We identified cpd984 which had a high predicted affinity to the N-terminal NT binding site using GLIDE XP [15] with a gscore of -9.0 kcal/mol binding strength as compared with a gscore of -8.9 kcal/mol for the N-terminal NT sequence. The position of cpd984 relative to critical amino acid residues of this region is shown (FIG. 3B). Surface amino acids at the binding site for N-terminal NT correspond with amino acids in the interaction diagram presented for cpd984 indicating similar regions of binding. In contrast to the results demonstrated under enriched conditions shown in FIG. 2B, under basal conditions (1% BSA/DMEM), there is a direct relationship between cellular sortilin expression and VLDL-B100 secretion. In FIG. 3B, one such compound (984) is seen interacting with specific residues of sortilin. The primary ligand binding site on sortilin (SITE 1) has been well characterized by arginine 292 in sortilin binding the carboxy-terminal of neurotensin (NT).[16] SITE 2 is defined by interactions with sortilin of N-terminal NT depicted in FIG. 3A by residues 75, 82, 446, 448, 539, 553, and 564. These residues can be seen defining SITE 2 as shown in FIG. 3B and correspond to residues 108, 115, 479, 481, 572, 586, and 597 of hsortilin.

Cpd984 and NT Bind to Hsortilin.

To confirm that cpd984 was bound by hsortilin at a NT-related binding site, we performed SPR (FIG. 3C). We show that individually NT (red, lower curve) and cpd984 (green, middle curve) bind hsortilin. The presence of cpd984 resulted in an increase of 14.5 RU for co-injected NT (black, upper curve) over the RU of the combination of NT and cpd984 alone. Considering the larger size of NT, the increase in RU likely represents enhancement of NT binding to hsortilin by cpd984.

Sortilin is known to influence the fate of VLDL-B100.[5] The complexity of sortilin function has been summarized[1] and may be related to sites of ligand binding that direct different functions. The ectodomain of human sortilin (hsortilin) has been crystallized in the presence of NT, a natural ligand.[20] Using Schrödinger's PRIME, a binding structure for NT has been defined combining the C-terminal (C-term) and N-terminal (N-term) fragments to produce a predicted NT full length binding model. The C-term is the major NT binding site through a salt bridge with arginine 292 (R292) that is defined as SITE 1. The N-term interaction with sortilin is defined as SITE 2, which is a hydrophobic

binding site.[21] Schrödinger software was used to identify compounds in the ChemBridge library that bind hsortilin at SITE 1 (54122218 or 541) or SITE 2 (cpd98477898 or 984). VLDL interaction with sortilin at these two sites plays an important role in determining VLDL-B100 secretion or degradation. It is established that 984 binds sortilin by SPR through its ability to compete with NT binding as seen in FIG. 3C. Compounds 541 and 984 are suitable for in vitro cell culture studies and are non-toxic based on LDH release. Compound 984 increases VLDL-B100 secretion by McA cells (GIH. 3D) indicating the utility of this compound binding to SITE 2 and altering a known sortilin function.[21] In FIG. 4A, 984 is seen to enhance the secretion of VLDL-B100 at all concentrations of sortilin which were previously shown to directly relate to VLDL-B100 secretion in FIG. 2C. These results indicate the enhanced ability of 984 to increase VLDL-B100 secretion showing its potential utility as a chaperone to remove VLDL-B100 and its component triglyceride from liver cells as would be useful to ameliorate hepatic steatosis.

PIP3 is shown to bind SITE 1, and while PIP3 binding to SITE 1 increases the potential for VLDL degradation it does this only at higher PIP3 concentrations. Lower PIP3 levels may favor some binding with chaperone activity predominantly being mediated by SITE 2. Supporting evidence for this premise is presented in recent publications.[21],[22] First, it was shown that 984 increases the binding of NT to hsortilin (SITE 1). Second, using diC16-PIP3 liposomes it was demonstrated that 984 shows concentration dependent increases in binding to hsortilin. Third, the binding of the C-term NT (SITE 1) to hsortilin is increased 100-fold when incubated with 984. Fourth, using SPR, 541 (SITE 1) competes for NT binding and 984 enhances NT binding. At higher VLDL PIP3 content following insulin, there is initiation of "sortilin-VLDL-B100-PIP3" complex formation and entry into autophagic-lysosomal degradation. The closer contact of VLDL with sortilin created by SITE 2 favors interaction with PIP3 at SITE 1. Because of the complexity of VLDL structure, diC16-PIP3 nanodiscs⁷ were prepared showing competition with NT binding to sortilin which complements studies showing direct competition of diC4-PIP3 liposomes for NT binding.[22] Using diC16 liposomes to test binding to sortilin in "pull-down" assays, results indicate that B100 is not required for PIP3 binding. [22] Using PC/PE liposomes as a control, there is specificity for PIP3 for binding. In McA cells under basal conditions, 541 and 984 have opposing dose-dependent effects on the secretion of VLDL-B100. There is significant enhancement of secretion of VLDL-B100 in McA cells with reduced sortilin at 40% and 10% of control levels by 984. The hypothesis is that the binding of VLDL at SITE 2 enhances the binding to SITE 1 orienting the VLDL in position for further interactions mediated by PIP3. Multiple PIP3 interaction can result in recruitment of additional sortilins anchored in the Golgi membrane distorting the membrane closest to the VLDL particle and initiating autophagy.

Cpd984 Increases VLDL-B100 Secretion by Insulin Sensitive McA Cell.

We next tested whether cpd984 alters sortilin function in insulin sensitive McA cells (FIG. 3D). Cpd984 increased VLDL-B100 secretion with significant increases observed at 10 μ M and 25 μ M averaging 30% and 61%, respectively, over the no cpd984 control.

Cpd984 Increases VLDL-B100 Secretion by McA Cells Under-Expressing Sortilin.

To evaluate the effect of cpd984 on sortilin deficient McA cells, McA cell lines with sortilin KD were incubated with

10 μ M cpd984. Cpd984 significantly enhanced VLDL-B100 secretion in each cell line (FIG. 4B). Incubation of McA cells with cpd984 did not alter insulin sensitivity, as insulin-dependent pY of IR β 3 and stimulation of pAKT(S473) were not affected. Reduced VLDL-B100 secretion occurred in both M (McA62) and L (McA53) cell lines with and without cpd 984 added ($p < 0.05$). Although the mechanism of action of cpd984 is not fully established, its action is likely mediated through interaction with sortilin as although VLDL-B100 secretion is increased, it is still lower with reduced sortilin suggesting sortilin remains rate-limiting. The complex relationship between sortilin and VLDL-B100 secretion is demonstrated in the current study. We show a rate-limiting role for sortilin in VLDL-B100 secretion that is present only in insulin sensitive McA cells and correlates with sortilin expression. Previous studies have shown that binding of B100 to sortilin is enhanced by insulin [9], and precedes B100 degradation mediated by autophagy [12]. With insulin sensitivity, apo B mRNA is unchanged so the decrease in VLDL-B100 secretion observed in 1% BSA/DMEM is likely due to baseline degradation. Together, these data suggest that baseline B100 degradation is further enhanced by insulin which reverses the role of sortilin as a secretory chaperone and directs B100 to autophagy, a pathway seen only with insulin sensitivity. A reasonable explanation for divergent effects of sortilin on VLDL-B100 secretion is the presence of multiple binding sites on sortilin that with binding result in different functions. It is thought that cpd984 may occupy a site on sortilin that enhances VLDL-B100 secretion through the N terminal NT binding domain that favors the chaperone function of sortilin.

Inhibition of the Second Site of Sortilin by Cpd984.

FIGS. 5A-5B demonstrate inhibition of the second site of sortilin by 984. McA cells were incubated in 1% BSA/DMEM with DMSO, 541 (10 μ M) or 984 (10 μ M) for 18 h. Cellular proteins were extracted and analyzed by IB for LDLR (LS-C146979, 1:2000) (FIG. 5A) or PCSK9 (Ab125251, 1:2000) (FIG. 5B) using HRP-linked secondary antibody and ECL detection. Loading controls included β -ACTIN (Rockland ph600-401-886) and GAPDH (cb1001mAb (6C₅)). Band intensities were measured using ChemiDocXRS+ system and evaluated using ImageLab 5.1 software (BioRad). Results are average of triplicates \pm S.D. PCSK9 is a known ligand for sortilin where sortilin acts as a chaperone for PCSK9 secretion.

Evidence is presented in FIGS. 5A-5B that there is competition for binding by 984 leading to the accumulation of cellular PCSK9. It is likely that SITE 2 represents the binding site for PCSK9 to sortilin as there is no biological difference with administration of 541 that is a SITE 1 binding compound. PCSK9 is known to bind to LDL receptors after it is secreted. PCSK9 bound to LDL receptors results in degradation of LDL receptors that results in higher circulating LDL cholesterol, a risk factor for cardiovascular disease. The increase in LDL receptors observed in FIG. 5B is consistent with reduced PCSK9 secretion. In these experiments, 984 interaction with PCSK9 would be useful in reducing secretion of PCSK9, increasing LDL receptor expression and reducing cardiovascular risk associated with higher concentrations of LDL cholesterol.

Effect of Sortilin Binding Compounds on PIP3 and Treatment of Diseases

Binding of PIP3 to sortilin was documented. PIP3 is a well known downstream signaling molecule for insulin generated through activation of phosphatidylinositide 3-kinase and is a highly negatively charged phospholipid in the class of phosphatidylinositides. PIP3 generation following

insulin is reduced in diabetes and conditions of insulin resistance. Increased PIP3 binding induced by 984 can therefore be useful in treatment of diabetic disorders. This can be accomplished through the enhancement of PIP3 binding to sortilin as a result of 984 binding to SITE 2 and enhanced PIP3 binding to SITE 1. Since 984 increases NT binding to sortilin, it is likely that 984 binding to SITE 2 would enhance binding to SITE 1 for other ligands including apoAV, lipoprotein lipase and apo E. Documentation of PIP3 binding to SITE 1 is further secured by competition with NT by diC4-PIP3 as SITE 1 is the major binding site for NT. Relevance for PIP3 in VLDL-B100 binding was determined by showing the presence of PIP3 on circulating VLDL in the blood that is known to contain a monolayer phospholipids including phosphatidylinositides on its surface (data not shown). The presence of PIP3 on VLDL has the potential to alter intravascular metabolism as sortilin is present both intracellularly and also on the cell surface. Further evidence for the enhancement of SITE 2 on the binding of ligands at SITE 1 was evidenced by the observations that 984 increases sortilin binding to PIP3 liposomes (data not shown). 984 was observed to increase liposome binding of sortilin. 984 was observed to increase the binding affinity of the C-terminal NT fragment to immobilized sortilin. Since PIP3 is generated by insulin action, it would be expected that 984 would enhance the interaction of VLDL-B100-PIP3 with sortilin and this would be useful when there is resistance to insulin as occurs with type 2 diabetes.

Summary

Results presented in this Example demonstrate that there is not an obligatory reciprocal relationship between sortilin expression and VLDL B100 secretion, and that the function of sortilin in McA cells depends on insulin sensitivity. Resolution of the controversies related to sortilin and VLDL-B100 secretion may be facilitated by identification of small molecules that bind different sites on sortilin and promote different sortilin function.

Abbreviations Used in Example 1

Apo B, apolipoprotein B; B100, apo B derived from unedited Apob mRNA; IDAD, insulin dependent apolipoprotein B degradation; McA, McArdle RH7777 cells; PI3K, Class IA phosphatidylinositide 3-kinase; SPR, surface plasmon resonance; VLDL, very low density lipoprotein

References for Example 1

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Example 2

Introduction.

Sortilin is a multi-ligand binding protein with numerous binding partners [1]. Sortilin binds neurotensin (NT) at two sites: the carboxy terminus (C-term) site 1, and the amino terminus (N-term) site 2 [2]. Sortilin also binds B100-containing very low density lipoprotein (VLDL) and LDL [3,4]. It was previously demonstrated that sortilin facilitates the secretion of VLDL-B100 by insulin-sensitive McArdle RH7777 (McA) cells [5]. It has been further shown that insulin enhances B100 binding to sortilin in McA cells [6] preceding autophagic destruction [7]. Considering the association of sortilin with VLDL binding and with VLDL-B100 degradation in response to insulin, the possibility for insulin-generated phosphatidylinositol (3,4,5)-trisphosphate (PIP3) to interact directly with sortilin was explored. The potential of PIP3 binding to NT defined binding sites on sortilin was tested. The results in this Example demonstrate a role for PIP3 present on VLDL to directly modulate lipoprotein-sortilin interactions.

Materials and Methods.

Materials.

Plasma from fasted Sprague Dawley rats was obtained from Bioreclamation/VT, (Westbury, N.Y.) and used to prepare rat lipoproteins. Bovine serum albumin (BSA) (A7285) was from Sigma-Aldrich Corp. (St. Louis, Mo.). The following materials and reagents were obtained from Echelon Biosciences, Inc. (St. Lake City, Utah) including PIP3 PolyPIPosomes (Y-P039); dibutanoyl PIP3 (diC4-PIP3) (P-3904); dipalmitoyl PIP3 (diC16-PIP3) (P-3916); purified anti-PIP3 IgG (Z-P345b), and purified anti-PIP3 IgM (ZP345) monoclonal antibodies. Biotinylated goat anti-mouse IgM and goat serum (GS) were from Jackson ImmunoResearch Labs, Inc. (West Grove, Pa.). Streptavidin-horseradish peroxidase (HRP), HRP-conjugated sheep anti-mouse IgG (NXA931), HRP-conjugated donkey anti-rabbit IgG (NA9340V), ECLTM Prime western blotting detection reagent, CM7 (S series) and NiNTA (S series) sensor chips were from GE Healthcare (Buckinghamshire, UK). Human sortilin (hsortilin) recombinant protein (Ser78-Asp755) with C-terminal 6-His tag was from R&D Systems, Inc. (Minneapolis, Minn.). Neurotensin C-terminal fragment (P-Y-I-

L) (C-term NT) was synthesized by Biomartik Corp. (Wilmington, Del.). SuperBlock®, mouse anti-6×HIS monoclonal antibody and 1-Step™ Ultra TMB-ELISA (TMB) were from Thermo Scientific (Rockford, Ill.). Costar® Assay Plates were from Corning, Inc. (Kennebunk, Me.). Rabbit anti-hsotilin polyclonal antibody (GTx54854) was from Gene-Tex, Inc. (Irvine, Calif.). Compound 98477898 (2S)-1-methyl-N-{3-[(3-phenylpropanoyl)-amino]phenyl}pyrrolidine-2-carboxamide (cpd984) was obtained from ChemBridge Corp. (San Diego, Calif.). All other materials and reagents were essentially as described previously [5].

Computational Modeling.

Schrödinger's Maestro program (version 9.3.5) was used as the primary graphical user interface and Release 2015-2: Maestro, version 10.2 (Schrödinger, LLC, New York, N.Y.) was used for ligand interaction diagramming [8]. Docking of diC4-PIP3 on a grid centered at the C-term NT (site1) of hsotilin crystal structure at excess concentration of NT PDB ID: 4PO7 was performed using Schrödinger's Glide software [9] as described [5]. Docking additionally indicated a strong affinity of PIP3 for site 2 of hsotilin.

Surface Plasmon Resonance.

Surface plasmon resonance (SPR) measurements were performed on a Biacore T200 instrument equipped with two separate CM7 sensor chips, one with about 20,000 response units (RU) and another with about 27,000 RU of hsotilin covalently immobilized. HBS running buffer with and without 1% (v/v) DMSO (10 mM HEPES pH 7.4, 150 mM NaCl) was used at a flow rate of 30 µl/min and injections had an association time of 90 s, dissociation time of 300 s, and binding was measured in relative RU or difference in response between the immobilized protein flow cell and the corresponding control flow cell subtracting relevant blank injections of buffer alone [5]. Regeneration buffer of pH 3.0 glycine was injected after each injection for PIP3 nanodiscs and C-term NT flowed in the presence of cpd984. For all other SPR injections, regeneration consisted of injections of running buffer alone. GE BIAcore T200 evaluation software version 3.0 (BIAevaluate) was used to analyze and export results. Full length NT was injected using 2:1 NT dilutions on ~27,000 RU CM7 surface and kinetics fitted using BIAevaluate Langmuir 1:1 binding model using local Rmax exported from BIAevaluate to Microsoft Excel 2016 with fit indicated by the black line. Full length NT was flowed in the presence of diC4-PIP3 diluted directly in running buffer and RU (5 s before injection stop) plotted in GraphPad Prism version 6.05 according to log [inhibitor] versus response [three parameter] model to obtain IC50 of diC4-PIP3 with respect to full length NT. PIP3 containing nanodiscs were flowed at 2:1 dilutions to obtain kinetic data fitted using about 27,000 RU CM7 surface. Kinetic fit generated with BIAevaluate and local Rmax using Langmuir 1:1 binding model exported from BIAevaluate to Microsoft Excel 2016 with fit indicated by the black line. C-term NT was flowed using 2:1 dilutions alone and co-injected with 100 µM cpd984 and steady state Kd obtained for full length NT using RU (45 and 10 s after injection start, respectively) plotted in GraphPad according to One site-Total binding model.

Preparation of PIP3 Nanodiscs.

Lipid composition of nanodiscs consisted of 3.023 µmol dipalmitoyl phosphatidylcholine (PC), 0.098 µmol diC16-PIP3, and 0.78 µmol 1-palmitoyl, 2-oleoyl phosphatidylethanolamine (PE), which were combined, dried, and desiccated overnight. Lipids were then dissolved in 20 mM sodium deoxycholate in TBS (50 mM Tris-HCl, pH 7.4, 150 mM NaCl, 0.02% (w/v) sodium azide) and sonicated. Mem-

brane scaffold protein 1D1 (MSP1D1)) was then added in a ratio of 70:1 lipid to protein and detergent removed with Bio-Beads® SM-2 (Bio-Rad). Nanodiscs were isolated using size exclusion chromatography [10]. Nanodiscs were quantified using a NanoDrop and extinction coefficient of 21,000 l mol⁻¹ cm⁻¹ for MSP1D1, and the resultant mg/ml divided by two because there are two MSP proteins per nanodisc.

Liposome Floatation Assay.

Unilamellar liposomes containing specific lipid compositions were produced using lipids dissolved in chloroform that were mixed according to mole percentages of 2.6 µmoles total phospholipid [11, 12]. Lipids were dried and desiccated overnight followed by incubation with PBS (2.6 ml) for 1 h. Lipids were resuspended by vortexing and disrupted by sonication for 30 min. To measure binding, a liposome floatation assay was used by adding concentrations 50 nM or 100 nM hsotilin to liposomes (150 µl) containing 5% PIP3, 20% PE and 75% PC; 10% PIP3, 20% PE, 70% PC; or 0% PIP3, 80% PC, and 20% PE as described [12, 13]. Samples containing 50 nM or 100 nM protein and liposomes in the presence and absence of cpd984 were incubated for 10 min at 30° C. and 630 µl of 1.65 M sucrose/PBS was added. Samples were overlaid with 630 µl of 0.75 M sucrose/PBS, and PBS to the top of the tube and centrifuged (200,000×g, 90 min, 4° C.), and floated liposomes recovered from the 0.75 M sucrose layer. Liposomes at the interface of the top and second fraction were recovered and diluted in 2 ml of PBS and re-isolated by centrifugation (16,000×g, 20 min, 4° C.). 50 µl PBS and 20 µl 5×SDS sample buffer was added to the final liposome pellet. Bound proteins were resolved by SDS-PAGE and detected by western immunoblotting following transfer to PVDF membranes [5].

Assay of Immunoreactive PIP3 on Rat Plasma Lipoproteins.

Plasma VLDL, LDL and high density lipoprotein (HDL) from fasted rats were isolated by sequential density ultracentrifugation as previously described [14]. Immunoreactive PIP3 on plasma lipoproteins was evaluated by enzyme linked immunoassay and by immunoslot blotting. Enzyme linked immunoassay involved coating 96-well multi-well assay plates with 0.1 ml/well diluted lipoprotein or BSA in 0.05 M carbonate/bicarbonate buffer (25 µg protein/ml) overnight at 4° C. Wells were washed three times and post-coated with SuperBlock® for 1 h at room temperature to block non-specific binding. Anti-PIP3 (IgM) diluted in 1% (v/v) GS/TBS was added (1:1000 dilution) and incubated at 4° C. overnight. Wells were washed three times in 1% (v/v) GS/TBS and 0.1 ml biotinylated goat anti-mouse IgM antibody was then added and incubated for 30 min at 37° C. Wells were washed three times in 1% (v/v) GS/TBS and 0.1 ml of streptavidin-HRP in 1% (v/v) GS/TBS was added (1:500) and incubated for 30 min at 37° C. Wells were washed three times in 1% (v/v) GS/TBS and bound HRP was detected by addition of 0.1 ml TMB reagent with incubation at room temperature. Reaction was terminated by addition of 0.1 ml of 2 M sulfuric acid and absorbance was measured at 450 nm using the iMark™ Microplate Reader (Bio-Rad, Hercules, Calif.). The second method employed immunoslot blotting where rat plasma VLDL and LDL from three separate plasma pools were diluted in TBS (2.5-10 µg protein/ml) and slot blotted (200 µl/slot) onto PVDF membranes as previously described [5]. PolyPIP3osomes containing PIP3 diluted 1:2000 in PBS were used as positive control. Following filtration, membranes were air dried, quickly rehydrated in methanol, re-incubated in PBS and then incubated in SuperBlock® solution at 4° C. overnight.

Afterwards, membranes were incubated in 3% (w/v) BSA/PBS containing anti-PIP3 (IgG) (1:1000) overnight at 4° C. After washing three times in PBS/0.1% Tween 20 (PBST), membranes were incubated in HRP-conjugated sheep anti-mouse IgG in 3% (w/v) BSA/PBS for 90 min. Membranes were washed three times in PBST and then developed using ECL™ Prime western blotting detection reagent. Chemiluminescence was measured using the ChemoDocXRS+ system (Bio-Rad) and band intensities quantified using Image Lab 3.0.1 software (Bio-Rad).

Statistics.

Unless otherwise noted, results are expressed as mean±S.E.M. SPR experiments were analyzed using GE BIAevaluation Software, Microsoft Excel and GraphPad Prism.

Results and Discussion

Sortilin is a PIP3 Binding Protein.

Sortilin binding to NT has been characterized and involves N-term (site 2) and C-term (site 1) binding sites for binding to a single molecule of sortilin [2, 15]. A compound was identified with strong theoretical binding to site 2 (cpd984), and in biologic experiments cpd984 incubation increased VLDL-B100 secretion by McA cells [5]. N-term NT was depicted in site 2 located across the central space of sortilin from PIP3 computationally docked into site 1 of hsortilin (data not shown). The theoretical binding of diC4-PIP3 was modeled for binding to site 1 with a docking score (gscore) of -10.3 kcal/mol indicating a high predicted affinity for hsortilin [8]. DiC4-PIP3 is shown binding to site 1 in FIG. 1A with phosphate groups interacting with arginine 292 (R292). The interaction diagram provided in FIG. 5B indicates a number of additional interacting amino acids. To directly test PIP3 binding to sortilin, we prepared PIP3 containing nanodiscs and performed SPR using immobilized hsortilin (data not shown). Nanodiscs were used to simulate PIP3 in membranes or membrane-like surfaces as might be present on VLDL. The Kd for PIP3 nanodiscs was 17 nM. The theoretical prediction of sortilin as a PIP3 binding protein is supported by these results.

PIP3 Competes with NT for Binding to Sortilin.

To determine whether PIP3 binds to sortilin at similar sites as NT, diC4-PIP3 was flowed in the presence of full length NT on immobilized hsortilin and analyzed the response using SPR. In the absence of hsortilin was approximately 290 nM (data not shown). This value is higher than the nM value previously reported using isothermal titration calorimetry [2], potentially because of unproductive cross-linking of hsortilin to the CM7 chip. At 100 nM NT, there was progressive competition for NT-sortilin interaction by diC4-PIP3. An IC50 of approximately 750 nM was obtained for diC4-PIP3 inhibition of full length NT binding to sortilin (data not shown). Structural analysis of a sortilin-related receptor, sorLA, suggests that a site exists similar to site 2 on sortilin that binds mainly hydrophobic ligands [16]. While there was some predicted binding affinity of PIP3 to site 2 of sortilin, it is unlikely to be relevant to sortilin-lipoprotein interaction as hydrophobic components of PIP3 would be anticipated to be embedded within the lipoprotein. Based on the overall data, we hypothesize that sortilin binds PIP3 at a NT related site, with binding most likely occurring at the C-term site 1.

Immunoreactive PIP3 is Present on Circulating Lipoproteins.

Previous studies have demonstrated the presence of phosphatidylinositols on circulating lipoproteins [17]. PIP3 is present in serum bound to albumin [18]. It was tested whether PIP3 might also be present on plasma lipoproteins.

Plasma lipoproteins have a surface monolayer of phospholipids that would orient the phosphatidylinositol head group toward the aqueous environment. Qualitative studies were performed where freshly prepared rat plasma VLDL, LDL and HDL were adsorbed to solid phase plastic wells. After blocking non-specific binding, we detected PIP3 using an IgM monoclonal anti-PIP3 antibody. There was three times the immunoreactivity present in VLDL as compared with LDL, HDL or BSA. To confirm these findings, VLDL and LDL were slot blotted and PIP3 immunoreactivity determined using an IgG monoclonal anti-PIP3 antibody (data not shown). Results were compared with slotted PolyPIPisomes containing PIP3 as a positive control. Calculating the relative reactivity per microgram of three separate preparations of rat plasma VLDL and LDL, there was 4.3, 2.5 and 4.0 times more immunoreactivity present in VLDL compared with each matched LDL. These results indicate there is immunoreactive PIP3 on circulating lipoproteins. The presence of lipoprotein PIP3 could influence the interaction of lipoproteins with corresponding cellular receptors including sortilin.

Sortilin Binds PIP3 Liposomes and Binding is Enhanced by a Sortilin Binding Compound.

Using liposome floatation assay, sortilin was shown to bind PIP3 liposomes, but not PC/PE liposomes (data not shown). A sortilin binding compound (cpd984) was developed to bind site 2 [5], and was used to probe PIP3 and C-term NT binding to sortilin. In the presence of increasing concentrations of cpd984, there was increased sortilin binding to PIP3 liposomes without change in binding to PC/PE liposomes (data not shown). The enhanced binding of PIP3 liposomes to sortilin is confirmed using an anti-sortilin antibody (data not shown). A C-terminal NT tetrapeptide (P-Y-I-L) (C-term NT) was used in SPR experiments to examine the effect of cpd984 on site 1 binding to immobilized hsortilin. The C-term NT alone had a Kd of 138 nM. To test the effect of cpd984, the Kd for the C-term NT in the presence of 100 µM cpd984 was determined. The Kd was lowered from 138 nM to 662. These studies indicate cpd984 enhanced the affinity of binding of PIP3 liposomes and C-term NT to sortilin. The use of liposomes as lipoprotein surrogates containing PIP3 to show binding to sortilin is important, as liposomes do not contain other potential sortilin ligands that would be present on VLDL such as B100 and apoE. It is thought that VLDL containing multiple PIP3 molecules on its surface and additional ligands could bind both sites 1 and 2 on sortilin enhancing binding as the size of spherical VLDL is large enough to span the central region of sortilin. Considering the downstream role PIP3 plays in insulin signaling and the enigmatic role sortilin has in VLDL metabolism [19], the finding that sortilin is a PIP3 binding protein could have important implications for VLDL. The presence of PIP3 on VLDL could influence the relative binding of B100 and apoE to various receptors including sortilin. Furthermore, the presence of PIP3 within the secretory pathway on VLDL might affect cell transit if it were demonstrated to modulate sortilin-B100 binding. It appears from these results that there is an interaction between at least two sites on sortilin in response to binding [5]. Because there is such a strong increase with cpd984 in site 1 ligand binding, the data suggest that there could be a conformational change in sortilin similar to that reported for sorLA [16]. In summary, the data support that sortilin is a PIP3 binding protein that could influence VLDL interaction with sortilin based on PIP3 content. Considering PIP3 is a downstream insulin signaling molecule, the demonstration of

sortilin binding and presence of PIP3 on VLDL may represent a novel signaling pathway for insulin action.

Abbreviations for Example 2

PIP3, phosphatidylinositol (3,4,5)-trisphosphate; diC4-PIP3, dibutanoyl PIP3; hsrtilin, human recombinant sortilin; NT, neurotensin; C-term, carboxy terminus; N-term, amino terminus; SPR, surface plasmon resonance; RU, response units; VLDL, very low density lipoprotein; B100, apolipoprotein B derived from unedited APOB mRNA.

References for Example 2

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Example 3

This Example demonstrates the effect of cpd984 and cpd541 alone and in combination can have an effect on secretion of VLDL-B100 on hepatocytes and VLDL-apo E by hepatocytes. FIG. 6 demonstrates an effect of cpd984 and cpd541 on secretion of VLDL-B100 by hepatocytes. McArdle rat hepatocytes were incubated with increasing concentrations of cpd984 or cpd541 for 14-18 h and media were collected. VLDL (d<0.19 g/ml) was isolated from media by ultracentrifugation and were slotted onto PVDF membranes. After blocking non-specific binding, blots were incubated with monoclonal anti-B100 antibody (24.05) followed by HRP-conjugated anti-mouse IgG. Binding was detected by chemiluminescence detection and quantified using Bio-Rad software. Results are the average of triplicate analysis in one study.

FIGS. 7A-7C show graphs demonstrating an effect of cpd984 and pd541 alone and in combination on the secretion of VLDL-B100 and VLDL-apo E by hepatocytes. McArdle rat hepatocytes were incubated with five media concentrations 0 (no cpd), 1 (0.1 μ M), 2 (1 μ M), 3 (10 μ M) and 4 (25 μ M) of cpd984 (FIG. 7A), cpd541 (FIG. 7B), or with a combination of both compounds (FIG. 7C) at indicated concentrations. Hepatocytes were incubated overnight (14-18 h) and media were collected and VLDL-B100 and VLDL-apo E were detected by slot blotting as described above. An anti-rat apoE antibody was used for measurement of VLDL-apo E secretion. Results are the average of triplicate analysis in one study.

An important consideration for VLDL-B100 secretion and the effects of sortilin binding compounds is the possibility of combining compounds binding to SITE 1 and SITE 2. In FIG. 6 it is demonstrated that 984 increases VLDL-B100 secretion in contrast to 541 that decreases VLDL-B100 secretion. These results document different functions

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of SITE 1 and SITE 2 with influence on VLDL-B100. Apolipoprotein E (apoE) is present on VLDL-B100 particles as well as other apolipoproteins. In FIGS. 7A-7C, 984 and 541 alone and in combination demonstrate effects on secretion of VLDL-B100 and VLDL-apoE by liver cells. As can be seen in FIG. 7A, 984 increases VLDL-B100 and VLDL-apoE in contrast to 541 alone which decreases VLDL-B100 and VLDL-apoE. When given in combination, the effects of 984 are abrogated. The effect of 541 binding to SITE 1 therefore could be used in combination with 984 to alter ligand binding at SITE 2.

Example 4

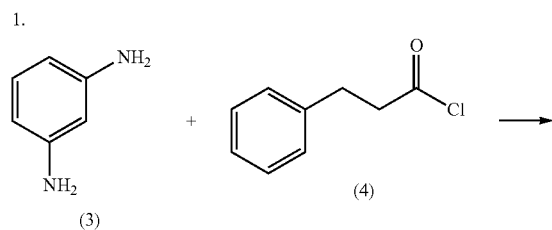
FIG. 8 shows a graph demonstrating the results of two trials of Label Free Thermophoresis of 541 to human sortilin. Due to inherent fluorescence of 541 small molecule, thermophoresis was performed in two different buffer conditions (PBS n=3 and MST n=2) and at two sets of concentration ranges. Because thermophoresis performed in PBS at higher concentration 51 revealed substantial initial fluorescence, results in MST buffer are used for K_d estimation. FIGS. 9A-9B show graphs representing the results of two trials of SPR Indicating Binding of 541 to hsortilin at with K_d of 7.5 μM. FIGS. 10A-10B show graphs demonstrating the results of SPR of 984 which reveals two different binding ranges. SPR indicates a high and low affinity binding site of cpd984 to human sortilin of hsortilin with K_d of about 200 μM and about 20 nM. FIGS. 11A-11C show graphs demonstrating the results of thermophoresis of Cpd984, which revealed two different binding ranges with K_d of 7 nM and about 100 to 300 μM.

FIGS. 8-9B confirm the binding of 541 to human sortilin using surface plasmon resonance (SPR) and thermophoresis. FIGS. 10A-10B confirms binding of 984 to human sortilin using SPR and demonstrates that this binding occurs through low and high affinity binding suggesting two binding sites. FIGS. 11A-11C demonstrate using thermophoresis the binding ranges of 984 for human sortilin including a very high affinity binding site with a K_d of 7 nM which should allow use of 984 at a very low concentration. All of the biologic results for 984 and 541 indicate that the compounds are not restricted with regards to their cellular uptake.

Example 5

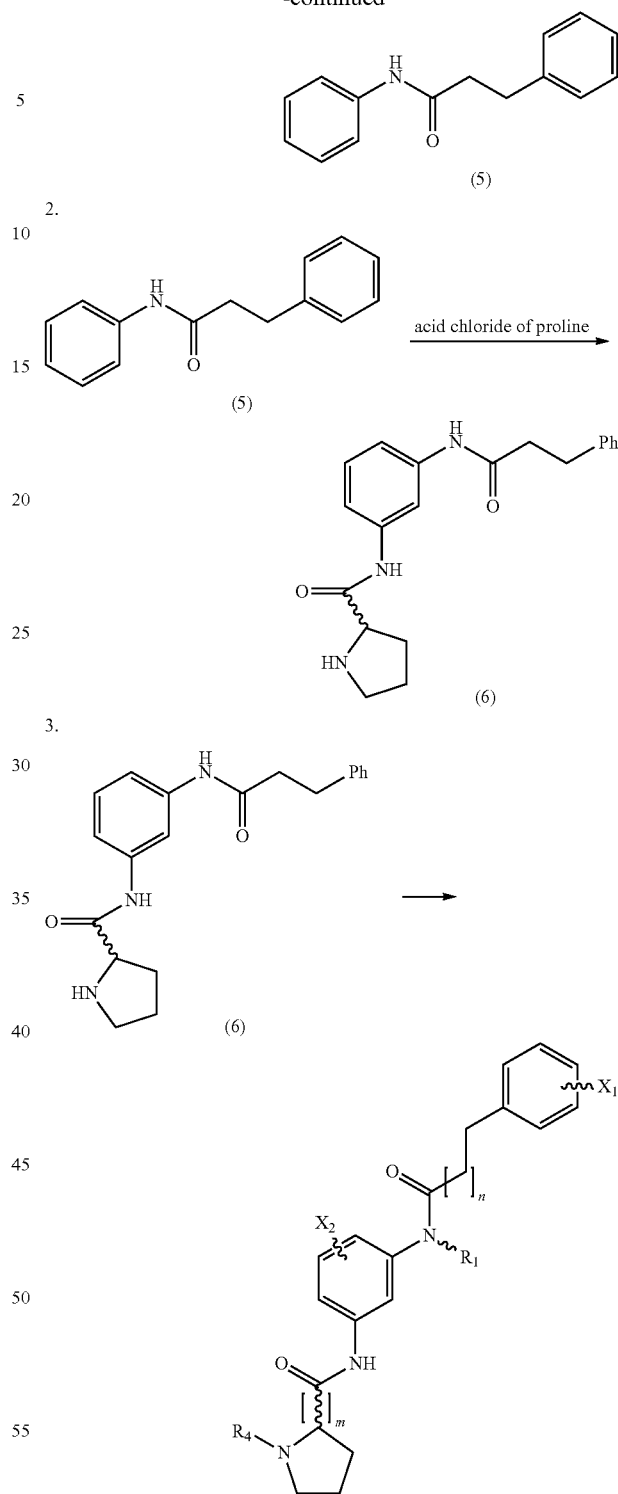
The sortilin binding compounds provided herein can be generated using techniques and methods generally known to one of ordinary skill in the art in view of this disclosure. An Exemplary synthesis Scheme is shown in Scheme 1.

Scheme 1



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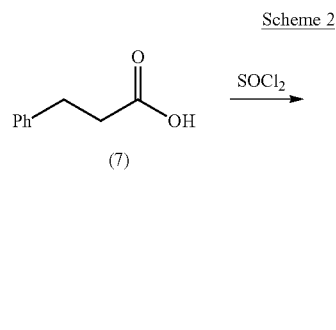
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In some embodiments, the synthesis of sortilin binding compounds can begin with a diamines, analogue, or derivative thereof combined with a phenyl alkanoyl containing a leaving group such as chloride (e.g. compound (4) in Scheme 1) to form an intermediate, such as compound (5). In instances where the diamine diacylates, one of ordinary skill in the art will appreciate ways to avoid, reduce and/or

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eliminate the dacylation at this step. The phenyl alkanoyl containing a leaving group can be produced via a synthesis scheme such as that shown in Scheme 2.



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A leaving group (e.g. Cl, Br, etc.) can be added to the phenyl alkanoyl, and the product can be used in Scheme 1. The resulting produce from step 1 in Scheme 1 (e.g. compound (5)) can be reacted with an acid chloride of proline, which can be made from reacting proline and SOCl₂ to form an intermediate compound (e.g. compound (6)).

Various substituents can be added to the any of the compounds or intermediates of Schemes 1 or 2 as need to generate Formulas 2, 2A, or 2B as needed. For example, substituents X₁ can be added using beta-phenyl propanoic acid analogs during Step 1. R₁ can be added by alkylating the product of Step 2. In embodiments, the alkylation reaction can be set up to favor alkylation at one position over another. X₂ can be added using m-phenylene diamine or analogue thereof in Step 1. R₄ can be added by using proline analogues that are alkylated at the proline nitrogen in Step 2. One of skill in the art will appreciate how to modify Synthesis Schemes 1 or 2 as needed depending on starting materials at any step and desired compounds. Such synthesis schemes are within the scope of this disclosure.

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Ala	Asn	Gly	Ser	Cys	Lys	Ala	Asp	Leu	Gly	Ala	Leu	Glu	Leu	Trp	Arg	275	280	285	
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 690 695 700
 Gly Arg Glu Glu His Leu Thr Thr Asn Gly Tyr Arg Lys Ile Pro Gly
 705 710 715 720
 Asp Lys Cys Gln Gly Gly Val Asn Pro Val Arg Glu Val Lys Asp Leu
 725 730 735
 Lys Lys Lys Cys Thr Ser Asn Phe Leu Ser Pro Glu Lys Gln Asn Ser
 740 745 750
 Lys Ser Asn Ser Val Pro Ile Ile Leu Ala Ile Val Gly Leu Met Leu
 755 760 765
 Val Thr Val Val Ala Gly Val Leu Ile Val Lys Lys Tyr Val Cys Gly
 770 775 780
 Gly Arg Phe Leu Val His Arg Tyr Ser Val Leu Gln Gln His Ala Glu
 785 790 795 800
 Ala Asn Gly Val Asp Gly Val Asp Ala Leu Asp Thr Ala Ser His Thr
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 Asn Lys Ser Gly Tyr His Asp Asp Ser Asp Glu Asp Leu Leu Glu
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 Glu Phe Gly Met Ala Ile Gly Pro Glu Asn Ser Gly Lys Val Val Leu
 35 40 45
 Thr Ala Glu Val Ser Gly Gly Ser Arg Gly Gly Arg Ile Phe Arg Ser
 50 55 60
 Ser Asp Phe Ala Lys Asn Phe Val Gln Thr Asp Leu Pro Phe His Pro
 65 70 75 80
 Leu Thr Gln Met Met Tyr Ser Pro Gln Asn Ser Asp Tyr Leu Leu Ala
 85 90 95
 Leu Ser Thr Glu Asn Gly Leu Trp Val Ser Lys Asn Phe Gly Gly Lys
 100 105 110
 Trp Glu Glu Ile His Lys Ala Val Cys Leu Ala Lys Trp Gly Ser Asp
 115 120 125

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Asn Thr Ile Phe Phe Thr Thr Tyr Ala Asn Gly Ser Cys Thr Asp Leu		
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Gly Ala Leu Glu Leu Trp Arg Thr Ser Asp Leu Gly Lys Ser Phe Lys		
145	150	155 160
Thr Ile Gly Val Lys Ile Tyr Ser Phe Gly Leu Gly Gly Arg Phe Leu		
	165 170	175
Phe Ala Ser Val Met Ala Asp Lys Asp Thr Thr Arg Arg Ile His Val		
	180 185	190
Ser Thr Asp Gln Gly Asp Thr Trp Ser Met Ala Gln Leu Pro Ser Val		
	195 200	205
Gly Gln Glu Gln Phe Tyr Ser Ile Leu Ala Ala Asn Asp Asp Met Val		
	210 215	220
Phe Met His Val Asp Glu Pro Gly Asp Thr Gly Phe Gly Thr Ile Phe		
225	230 235	240
Thr Ser Asp Asp Arg Gly Ile Val Tyr Ser Lys Ser Leu Asp Arg His		
	245 250	255
Leu Tyr Thr Thr Thr Gly Gly Glu Thr Asp Phe Thr Asn Val Thr Ser		
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Leu Arg Gly Val Tyr Ile Thr Ser Val Leu Ser Glu Asp Asn Ser Ile		
	275 280	285
Gln Thr Met Ile Thr Phe Asp Gln Gly Gly Arg Trp Thr His Leu Arg		
	290 295	300
Lys Pro Glu Asn Ser Glu Cys Asp Ala Thr Ala Lys Asn Lys Asn Glu		
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Cys Ser Leu His Ile His Ala Ser Tyr Ser Ile Ser Gln Lys Leu Asn		
	325 330	335
Val Pro Met Ala Pro Leu Ser Glu Pro Asn Ala Val Gly Ile Val Ile		
	340 345	350
Ala His Gly Ser Val Gly Asp Ala Ile Ser Val Met Val Pro Asp Val		
	355 360	365
Tyr Ile Ser Asp Asp Gly Gly Tyr Ser Trp Thr Lys Met Leu Glu Gly		
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Pro His Tyr Tyr Thr Ile Leu Asp Ser Gly Gly Ile Ile Val Ala Ile		
385	390 395	400
Glu His Ser Ser Arg Pro Ile Asn Val Ile Lys Phe Ser Thr Asp Glu		
	405 410	415
Gly Gln Cys Trp Gln Thr Tyr Thr Phe Thr Arg Asp Pro Ile Tyr Phe		
	420 425	430
Thr Gly Leu Ala Ser Glu Pro Gly Ala Arg Ser Met Asn Ile Ser Ile		
	435 440	445
Trp Gly Phe Thr Glu Ser Phe Leu Thr Ser Gln Trp Val Ser Tyr Thr		
	450 455	460
Ile Asp Phe Lys Asp Ile Leu Glu Arg Asn Cys Glu Glu Lys Asp Tyr		
465	470 475	480
Thr Ile Trp Leu Ala His Ser Thr Asp Pro Glu Asp Tyr Glu Asp Gly		
	485 490	495
Cys Ile Leu Gly Tyr Lys Glu Gln Phe Leu Arg Leu Arg Lys Ser Ser		
	500 505	510
Val Cys Gln Asn Gly Arg Asp Tyr Val Val Thr Lys Gln Pro Ser Ile		
	515 520	525
Cys Leu Cys Ser Leu Glu Asp Phe Leu Cys Asp Phe Gly Tyr Tyr Arg		
	530 535	540

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Pro Glu Asn Asp Ser Lys Cys Val Glu Gln Pro Glu Leu Lys Gly His
545                    550                    555                    560

Asp Leu Glu Phe Cys Leu Tyr Gly Arg Glu Glu His Leu Thr Thr Asn
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Gly Tyr Arg Lys Ile Pro Gly Asp Lys Cys Gln Gly Gly Val Asn Pro
                    580                    585                    590

Val Arg Glu Val Lys Asp Leu Lys Lys Lys Cys Thr Ser Asn Phe Leu
                    595                    600                    605

Ser Pro Glu Lys Gln Asn Ser Lys Ser Asn Ser Val Pro Ile Ile Leu
610                    615                    620

Ala Ile Val Gly Leu Met Leu Val Thr Val Val Ala Gly Val Leu Ile
625                    630                    635                    640

Val Lys Lys Tyr Val Cys Gly Gly Arg Phe Leu Val His Arg Tyr Ser
645                    650                    655

Val Leu Gln Gln His Ala Glu Ala Asn Gly Val Asp Gly Val Asp Ala
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Leu Asp Thr Ala Ser His Thr Asn Lys Ser Gly Tyr His Asp Asp Ser
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Asp Glu Asp Leu Leu Glu
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<210> SEQ ID NO 4
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<212> TYPE: PRT
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Gln Arg Leu His Gly Gly Ser Ala Pro Leu Pro Gln Asp Arg Gly Phe
35                    40                    45

Leu Val Val Gln Gly Asp Pro Arg Glu Leu Arg Leu Trp Ala Arg Gly
50                    55                    60

Asp Ala Arg Gly Ala Ser Arg Ala Asp Glu Lys Pro Leu Arg Arg Lys
65                    70                    75                    80

Arg Ser Ala Ala Leu Gln Pro Glu Pro Ile Lys Val Tyr Gly Gln Val
85                    90                    95

Ser Leu Asn Asp Ser His Asn Gln Met Val Val His Trp Ala Gly Glu
100                   105                   110

Lys Ser Asn Val Ile Val Ala Leu Ala Arg Asp Ser Leu Ala Leu Ala
115                   120                   125

Arg Pro Lys Ser Ser Asp Val Tyr Val Ser Tyr Asp Tyr Gly Lys Ser
130                   135                   140

Phe Lys Lys Ile Ser Asp Lys Leu Asn Phe Gly Leu Gly Asn Arg Ser
145                   150                   155                   160

Glu Ala Val Ile Ala Gln Phe Tyr His Ser Pro Ala Asp Asn Lys Arg
165                   170                   175

Tyr Ile Phe Ala Asp Ala Tyr Ala Gln Tyr Leu Trp Ile Thr Phe Asp
180                   185                   190

Phe Cys Asn Thr Leu Gln Gly Phe Ser Ile Pro Phe Arg Ala Ala Asp
195                   200                   205

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Leu	Leu	Leu	His	Ser	Lys	Ala	Ser	Asn	Leu	Leu	Leu	Gly	Phe	Asp	Arg
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Ser	His	Pro	Asn	Lys	Gln	Leu	Trp	Lys	Ser	Asp	Asp	Phe	Gly	Gln	Thr
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Trp	Ile	Met	Ile	Gln	Glu	His	Val	Lys	Ser	Phe	Ser	Trp	Gly	Ile	Asp
				245					250					255	
Pro	Tyr	Asp	Lys	Pro	Asn	Thr	Ile	Tyr	Ile	Glu	Arg	His	Glu	Pro	Ser
			260					265					270		
Gly	Tyr	Ser	Thr	Val	Phe	Arg	Ser	Thr	Asp	Phe	Phe	Gln	Ser	Arg	Glu
		275					280					285			
Asn	Gln	Glu	Val	Ile	Leu	Glu	Glu	Val	Arg	Asp	Phe	Gln	Leu	Arg	Asp
	290					295					300				
Lys	Tyr	Met	Phe	Ala	Thr	Lys	Val	Val	His	Leu	Leu	Gly	Ser	Glu	Gln
305					310					315					320
Gln	Ser	Ser	Val	Gln	Leu	Trp	Val	Ser	Phe	Gly	Arg	Lys	Pro	Met	Arg
			325						330					335	
Ala	Ala	Gln	Phe	Val	Thr	Arg	His	Pro	Ile	Asn	Glu	Tyr	Tyr	Ile	Ala
			340					345						350	
Asp	Ala	Ser	Glu	Asp	Gln	Val	Phe	Val	Cys	Val	Ser	His	Ser	Asn	Asn
		355					360					365			
Arg	Thr	Asn	Leu	Tyr	Ile	Ser	Glu	Ala	Glu	Gly	Leu	Lys	Phe	Ser	Leu
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Ser	Leu	Glu	Asn	Val	Leu	Tyr	Tyr	Ser	Pro	Gly	Gly	Ala	Gly	Ser	Asp
385					390					395					400
Thr	Leu	Val	Arg	Tyr	Phe	Ala	Asn	Glu	Pro	Phe	Ala	Asp	Phe	His	Arg
			405						410					415	
Val	Glu	Gly	Leu	Gln	Gly	Val	Tyr	Ile	Ala	Thr	Leu	Ile	Asn	Gly	Ser
			420					425					430		
Met	Asn	Glu	Glu	Asn	Met	Arg	Ser	Val	Ile	Thr	Phe	Asp	Lys	Gly	Gly
		435					440					445			
Thr	Trp	Glu	Phe	Leu	Gln	Ala	Pro	Ala	Phe	Thr	Gly	Tyr	Gly	Glu	Lys
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Ile	Asn	Cys	Glu	Leu	Ser	Gln	Gly	Cys	Ser	Leu	His	Leu	Ala	Gln	Arg
465					470					475					480
Leu	Ser	Gln	Leu	Leu	Asn	Leu	Gln	Leu	Arg	Arg	Met	Pro	Ile	Leu	Ser
			485						490					495	
Lys	Glu	Ser	Ala	Pro	Gly	Leu	Ile	Ile	Ala	Thr	Gly	Ser	Val	Gly	Lys
			500					505					510		
Asn	Leu	Ala	Ser	Lys	Thr	Asn	Val	Tyr	Ile	Ser	Ser	Ser	Ala	Gly	Ala
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Arg	Trp	Arg	Glu	Ala	Leu	Pro	Gly	Pro	His	Tyr	Tyr	Thr	Trp	Gly	Asp
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His	Gly	Gly	Ile	Ile	Thr	Ala	Ile	Ala	Gln	Gly	Met	Glu	Thr	Asn	Glu
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Leu	Lys	Tyr	Ser	Thr	Asn	Glu	Gly	Glu	Thr	Trp	Lys	Thr	Phe	Ile	Phe
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Ser	Glu	Lys	Pro	Val	Phe	Val	Tyr	Gly	Leu	Leu	Thr	Glu	Pro	Gly	Glu
			580					585					590		
Lys	Ser	Thr	Val	Phe	Thr	Ile	Phe	Gly	Ser	Asn	Lys	Glu	Asn	Val	His
			595				600					605			
Ser	Trp	Leu	Ile	Leu	Gln	Val	Asn	Ala	Thr	Asp	Ala	Leu	Gly	Val	Pro
	610					615					620				
Cys	Thr	Glu	Asn	Asp	Tyr	Lys	Leu	Trp	Ser	Pro	Ser	Asp	Glu	Arg	Gly

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625	630	635	640
Asn Glu Cys Leu Leu Gly His Lys Thr Val Phe Lys Arg Arg Thr Pro	645	650	655
His Ala Thr Cys Phe Asn Gly Glu Asp Phe Asp Arg Pro Val Val Val	660	665	670
Ser Asn Cys Ser Cys Thr Arg Glu Asp Tyr Glu Cys Asp Phe Gly Phe	675	680	685
Lys Met Ser Glu Asp Leu Ser Leu Glu Val Cys Val Pro Asp Pro Glu	690	695	700
Phe Ser Gly Lys Ser Tyr Ser Pro Pro Val Pro Cys Pro Val Gly Ser	705	710	715
Thr Tyr Arg Arg Thr Arg Gly Tyr Arg Lys Ile Ser Gly Asp Thr Cys	725	730	735
Ser Gly Gly Asp Val Glu Ala Arg Leu Glu Gly Glu Leu Val Pro Cys	740	745	750
Pro Leu Ala Glu Glu Asn Glu Phe Ile Leu Tyr Ala Val Arg Lys Ser	755	760	765
Ile Tyr Arg Tyr Asp Leu Ala Ser Gly Ala Thr Glu Gln Leu Pro Leu	770	775	780
Thr Gly Leu Arg Ala Ala Val Ala Leu Asp Phe Asp Tyr Glu His Asn	785	790	795
Cys Leu Tyr Trp Ser Asp Leu Ala Leu Asp Val Ile Gln Arg Leu Cys	805	810	815
Leu Asn Gly Ser Thr Gly Gln Glu Val Ile Ile Asn Ser Gly Leu Glu	820	825	830
Thr Val Glu Ala Leu Ala Phe Glu Pro Leu Ser Gln Leu Leu Tyr Trp	835	840	845
Val Asp Ala Gly Phe Lys Lys Ile Glu Val Ala Asn Pro Asp Gly Asp	850	855	860
Phe Arg Leu Thr Ile Val Asn Ser Ser Val Leu Asp Arg Pro Arg Ala	865	870	875
Leu Val Leu Val Pro Gln Glu Gly Val Met Phe Trp Thr Asp Trp Gly	885	890	895
Asp Leu Lys Pro Gly Ile Tyr Arg Ser Asn Met Asp Gly Ser Ala Ala	900	905	910
Tyr His Leu Val Ser Glu Asp Val Lys Trp Pro Asn Gly Ile Ser Val	915	920	925
Asp Asp Gln Trp Ile Tyr Trp Thr Asp Ala Tyr Leu Glu Cys Ile Glu	930	935	940
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Pro His Pro Tyr Ala Ile Ala Val Phe Lys Asn Glu Ile Tyr Trp Asp	965	970	975
Asp Trp Ser Gln Leu Ser Ile Phe Arg Ala Ser Lys Tyr Ser Gly Ser	980	985	990
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Arg Pro Cys Ser Leu Leu Cys Leu Pro Lys Ala Asn Asn Ser Arg	1025	1030	1035
Ser Cys Arg Cys Pro Glu Asp Val Ser Ser Ser Val Leu Pro Ser	1040	1045	1050

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Asn Thr 1070	Cys Val Lys Glu Glu 1075	Asn Thr Cys Leu Arg 1080	Asn Gln Tyr
Arg Cys 1085	Ser Asn Gly Asn Cys 1090	Ile Asn Ser Ile Trp 1095	Trp Cys Asp
Phe Asp 1100	Asn Asp Cys Gly Asp 1105	Met Ser Asp Glu Arg 1110	Asn Cys Pro
Thr Thr 1115	Ile Cys Asp Leu Asp 1120	Thr Gln Phe Arg Cys 1125	Gln Glu Ser
Gly Thr 1130	Cys Ile Pro Leu Ser 1135	Tyr Lys Cys Asp Leu 1140	Glu Asp Asp
Cys Gly 1145	Asp Asn Ser Asp Glu 1150	Ser His Cys Glu Met 1155	His Gln Cys
Arg Ser 1160	Asp Glu Tyr Asn Cys 1165	Ser Ser Gly Met Cys 1170	Ile Arg Ser
Ser Trp 1175	Val Cys Asp Gly Asp 1180	Asn Asp Cys Arg Asp 1185	Trp Ser Asp
Glu Ala 1190	Asn Cys Thr Ala Ile 1195	Tyr His Thr Cys Glu 1200	Ala Ser Asn
Phe Gln 1205	Cys Arg Asn Gly His 1210	Cys Ile Pro Gln Arg 1215	Trp Ala Cys
Asp Gly 1220	Asp Thr Asp Cys Gln 1225	Asp Gly Ser Asp Glu 1230	Asp Pro Val
Asn Cys 1235	Glu Lys Lys Cys Asn 1240	Gly Phe Arg Cys Pro 1245	Asn Gly Thr
Cys Ile 1250	Pro Ser Ser Lys His 1255	Cys Asp Gly Leu Arg 1260	Asp Cys Ser
Asp Gly 1265	Ser Asp Glu Gln His 1270	Cys Glu Pro Leu Cys 1275	Thr His Phe
Met Asp 1280	Phe Val Cys Lys Asn 1285	Arg Gln Gln Cys Leu 1290	Phe His Ser
Met Val 1295	Cys Asp Gly Ile Ile 1300	Gln Cys Arg Asp Gly 1305	Ser Asp Glu
Asp Ala 1310	Ala Phe Ala Gly Cys 1315	Ser Gln Asp Pro Glu 1320	Phe His Lys
Val Cys 1325	Asp Glu Phe Gly Phe 1330	Gln Cys Gln Asn Gly 1335	Val Cys Ile
Ser Leu 1340	Ile Trp Lys Cys Asp 1345	Gly Met Asp Asp Cys 1350	Gly Asp Tyr
Ser Asp 1355	Glu Ala Asn Cys Glu 1360	Asn Pro Thr Glu Ala 1365	Pro Asn Cys
Ser Arg 1370	Tyr Phe Gln Phe Arg 1375	Cys Glu Asn Gly His 1380	Cys Ile Pro
Asn Arg 1385	Trp Lys Cys Asp Arg 1390	Glu Asn Asp Cys Gly 1395	Asp Trp Ser
Asp Glu 1400	Lys Asp Cys Gly Asp 1405	Ser His Ile Leu Pro 1410	Phe Ser Thr
Pro Gly 1415	Pro Ser Thr Cys Leu 1420	Pro Asn Tyr Tyr Arg 1425	Cys Ser Ser
Gly Thr 1430	Cys Val Met Asp Thr 1435	Trp Val Cys Asp Gly 1440	Tyr Arg Asp

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Cys Ala	Asp Gly	Ser Asp	Glu	Glu Ala	Cys Pro	Leu	Leu Ala	Asn	
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1460			1465			1470			
Phe Glu	Phe Glu	Cys His	Gln	Pro Lys	Thr Cys	Ile	Pro Asn	Trp	
1475			1480			1485			
Lys Arg	Cys Asp	Gly His	Gln	Asp Cys	Gln Asp	Gly	Arg Asp	Glu	
1490			1495			1500			
Ala Asn	Cys Pro	Thr His	Ser	Thr Leu	Thr Cys	Met	Ser Arg	Glu	
1505			1510			1515			
Phe Gln	Cys Glu	Asp Gly	Glu	Ala Cys	Ile Val	Leu	Ser Glu	Arg	
1520			1525			1530			
Cys Asp	Gly Phe	Leu Asp	Cys	Ser Asp	Glu Ser	Asp	Glu Lys	Ala	
1535			1540			1545			
Cys Ser	Asp Glu	Leu Thr	Val	Tyr Lys	Val Gln	Asn	Leu Gln	Trp	
1550			1555			1560			
Thr Ala	Asp Phe	Ser Gly	Asp	Val Thr	Leu Thr	Trp	Met Arg	Pro	
1565			1570			1575			
Lys Lys	Met Pro	Ser Ala	Ser	Cys Val	Tyr Asn	Val	Tyr Tyr	Arg	
1580			1585			1590			
Val Val	Gly Glu	Ser Ile	Trp	Lys Thr	Leu Glu	Thr	His Ser	Asn	
1595			1600			1605			
Lys Thr	Asn Thr	Val Leu	Lys	Val Leu	Lys Pro	Asp	Thr Thr	Tyr	
1610			1615			1620			
Gln Val	Lys Val	Gln Val	Gln	Cys Leu	Ser Lys	Ala	His Asn	Thr	
1625			1630			1635			
Asn Asp	Phe Val	Thr Leu	Arg	Thr Pro	Glu Gly	Leu	Pro Asp	Ala	
1640			1645			1650			
Pro Arg	Asn Leu	Gln Leu	Ser	Leu Pro	Arg Glu	Ala	Glu Gly	Val	
1655			1660			1665			
Ile Val	Gly His	Trp Ala	Pro	Pro Ile	His Thr	His	Gly Leu	Ile	
1670			1675			1680			
Arg Glu	Tyr Ile	Val Glu	Tyr	Ser Arg	Ser Gly	Ser	Lys Met	Trp	
1685			1690			1695			
Ala Ser	Gln Arg	Ala Ala	Ser	Asn Phe	Thr Glu	Ile	Lys Asn	Leu	
1700			1705			1710			
Leu Val	Asn Thr	Leu Tyr	Thr	Val Arg	Val Ala	Ala	Val Thr	Ser	
1715			1720			1725			
Arg Gly	Ile Gly	Asn Trp	Ser	Asp Ser	Lys Ser	Ile	Thr Thr	Ile	
1730			1735			1740			
Lys Gly	Lys Val	Ile Pro	Pro	Pro Asp	Ile His	Ile	Asp Ser	Tyr	
1745			1750			1755			
Gly Glu	Asn Tyr	Leu Ser	Phe	Thr Leu	Thr Met	Glu	Ser Asp	Ile	
1760			1765			1770			
Lys Val	Asn Gly	Tyr Val	Val	Asn Leu	Phe Trp	Ala	Phe Asp	Thr	
1775			1780			1785			
His Lys	Gln Glu	Arg Arg	Thr	Leu Asn	Phe Arg	Gly	Ser Ile	Leu	
1790			1795			1800			
Ser His	Lys Val	Gly Asn	Leu	Thr Ala	His Thr	Ser	Tyr Glu	Ile	
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Ser Ala	Trp Ala	Lys Thr	Asp	Leu Gly	Asp Ser	Pro	Leu Ala	Phe	
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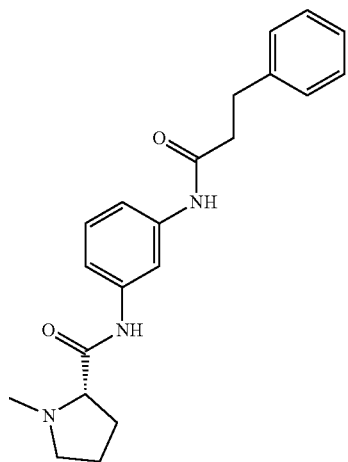
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Asn Lys Thr Val Ile Val Ser Lys Asp Glu Gln Tyr Leu Phe Leu 1895 1900 1905		
Val Arg Val Val Val Pro Tyr Gln Gly Pro Ser Ser Asp Tyr Val 1910 1915 1920		
Val Val Lys Met Ile Pro Asp Ser Arg Leu Pro Pro Arg His Leu 1925 1930 1935		
His Val Val His Thr Gly Lys Thr Ser Val Val Ile Lys Trp Glu 1940 1945 1950		
Ser Pro Tyr Asp Ser Pro Asp Gln Asp Leu Leu Tyr Ala Ile Ala 1955 1960 1965		
Val Lys Asp Leu Ile Arg Lys Thr Asp Arg Ser Tyr Lys Val Lys 1970 1975 1980		
Ser Arg Asn Ser Thr Val Glu Tyr Thr Leu Asn Lys Leu Glu Pro 1985 1990 1995		
Gly Gly Lys Tyr His Ile Ile Val Gln Leu Gly Asn Met Ser Lys 2000 2005 2010		
Asp Ser Ser Ile Lys Ile Thr Thr Val Ser Leu Ser Ala Pro Asp 2015 2020 2025		
Ala Leu Lys Ile Ile Thr Glu Asn Asp His Val Leu Leu Phe Trp 2030 2035 2040		
Lys Ser Leu Ala Leu Lys Glu Lys His Phe Asn Glu Ser Arg Gly 2045 2050 2055		
Tyr Glu Ile His Met Phe Asp Ser Ala Met Asn Ile Thr Ala Tyr 2060 2065 2070		
Leu Gly Asn Thr Thr Asp Asn Phe Phe Lys Ile Ser Asn Leu Lys 2075 2080 2085		
Met Gly His Asn Tyr Thr Phe Thr Val Gln Ala Arg Cys Leu Phe 2090 2095 2100		
Gly Asn Gln Ile Cys Gly Glu Pro Ala Ile Leu Leu Tyr Asp Glu 2105 2110 2115		
Leu Gly Ser Gly Ala Asp Ala Ser Ala Thr Gln Ala Ala Arg Ser 2120 2125 2130		
Thr Asp Val Ala Ala Val Val Val Pro Ile Leu Phe Leu Ile Leu 2135 2140 2145		
Leu Ser Leu Gly Val Gly Phe Ala Ile Leu Tyr Thr Lys His Arg 2150 2155 2160		
Arg Leu Gln Ser Ser Phe Thr Ala Phe Ala Asn Ser His Tyr Ser 2165 2170 2175		
Ser Arg Leu Gly Ser Ala Ile Phe Ser Ser Gly Asp Asp Leu Gly 2180 2185 2190		
Glu Asp Asp Glu Asp Ala Pro Met Ile Thr Gly Phe Ser Asp Asp 2195 2200 2205		
Val Pro Met Val Ile Ala 2210		

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We claim:

1. A method comprising:
contacting sortilin or an analogue thereof with an amount
of a compound having a structure according to Formula
2A

Formula 2A



2. The method of claim 1, wherein the step of contacting occurs in vitro.

3. The method of claim 1, further comprising the step of administering the compound having a structure according to Formula 2A to a subject, wherein the step of administering occurs prior to the step of contacting.

4. The method of claim 1, wherein the subject suffers from a disease whose pathology involves a ligand of sortilin or analogue thereof.

5. The method of claim 4, wherein the disease is hypercholesteremia or Alzheimer's disease.

6. The method of claim 4, wherein the ligand of sortilin or analogue thereof is PCSK9, apolipoprotein B100, phosphatidylinositides, anionic phospholipids, anionic phospholipids contained on VLDL, amyloid beta, amyloid precursor protein, neurotensin or analogues of neurotensin, lipoprotein lipase, apolipoprotein AV or apolipoprotein E.

7. The method of claim 4, wherein the ligand of sortilin or analogue thereof is an apolipoprotein.

8. The method of claim 4, wherein the ligand of sortilin or analogue thereof is PCSK9.

9. The method of claim 8, wherein the compound is a competitive inhibitor of PCSK9 for sortilin.

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10. The method of claim 6, wherein the compound is an allosteric enhancer of the ligand of sortilin or analogue thereof.

11. The method of claim 1, wherein the subject suffers a disease that involves sortilin in its pathology.

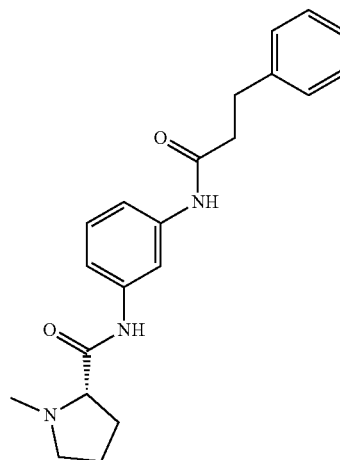
12. The method of claim 1, wherein the compound is capable of specifically binding sortilin.

13. The method of claim 1, wherein the compound is a competitive inhibitor of a sortilin ligand or an analogue thereof.

14. The method of claim 1, wherein the compound is an allosteric enhancer of a sortilin ligand or an analogue thereof.

15. A method of inhibiting binding of a sortilin ligand to sortilin in a subject in need thereof, the method comprising:
administering to the subject an effective amount of a compound according to Formula 2A

Formula 2A



wherein the sortilin ligand is selected from the group consisting of: PCSK9, apolipoprotein B100, phosphatidylinositides, anionic phospholipids, anionic phospholipids contained on VLDL, amyloid beta, amyloid precursor protein, neurotensin or analogues of neurotensin, lipoprotein lipase, apolipoprotein AV and apolipoprotein E.

16. The method of claim 13, wherein the subject suffers from hypercholesteremia or Alzheimer's disease.

17. The method of claim 13, wherein the subject suffers from hepatic steatosis.

* * * * *