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(54) **METHODS FOR REDUCING RISKS
ASSOCIATED WITH HEART FAILURE AND
FACTORS ASSOCIATED THEREWITH**

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- (63) Continuation of application No. 17/838,766, filed on Jun. 13, 2022, now abandoned, which is a continuation of application No. 16/724,076, filed on Dec. 20, 2019, now abandoned, which is a continuation of application No. 14/437,325, filed on Apr. 21, 2015, filed as application No. PCT/US2013/066228 on Oct. 22, 2013, now abandoned.
- (60) Provisional application No. 61/716,867, filed on Oct. 22, 2012, provisional application No. 61/822,752, filed on May 13, 2013, provisional application No.

61/839,743, filed on Jun. 26, 2013, provisional application No. 61/839,750, filed on Jun. 26, 2013.

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(52) **U.S. Cl.**

CPC *A61K 38/07* (2013.01); *A61K 38/06* (2013.01); *A61K 45/06* (2013.01); *C07K 5/1019* (2013.01)

(57) **ABSTRACT**

The disclosure provides methods of preventing or treating heart failure in a mammalian subject, reducing risk factors associated with heart failure, and/or reducing the likelihood or severity of heart failure. The disclosure also provides methods of preventing, or treating LV remodeling in a mammalian subject, and/or reducing the likelihood or severity of LV remodeling. The methods comprise administering to the subject an effective amount of an aromatic-cationic peptide. In some embodiments, the methods comprise administering to the subject an effective amount of an aromatic cationic peptide to reduce levels of C-reactive protein, tumor necrosis factor alpha, interleukin 6, reactive oxygen species, Nt-pro BNP, and/or cardiac troponin I, and/or reduce expression levels of MLCL AT1 and/or ALCAT 1 in subjects in need thereof.

Specification includes a Sequence Listing.

FIG. 1

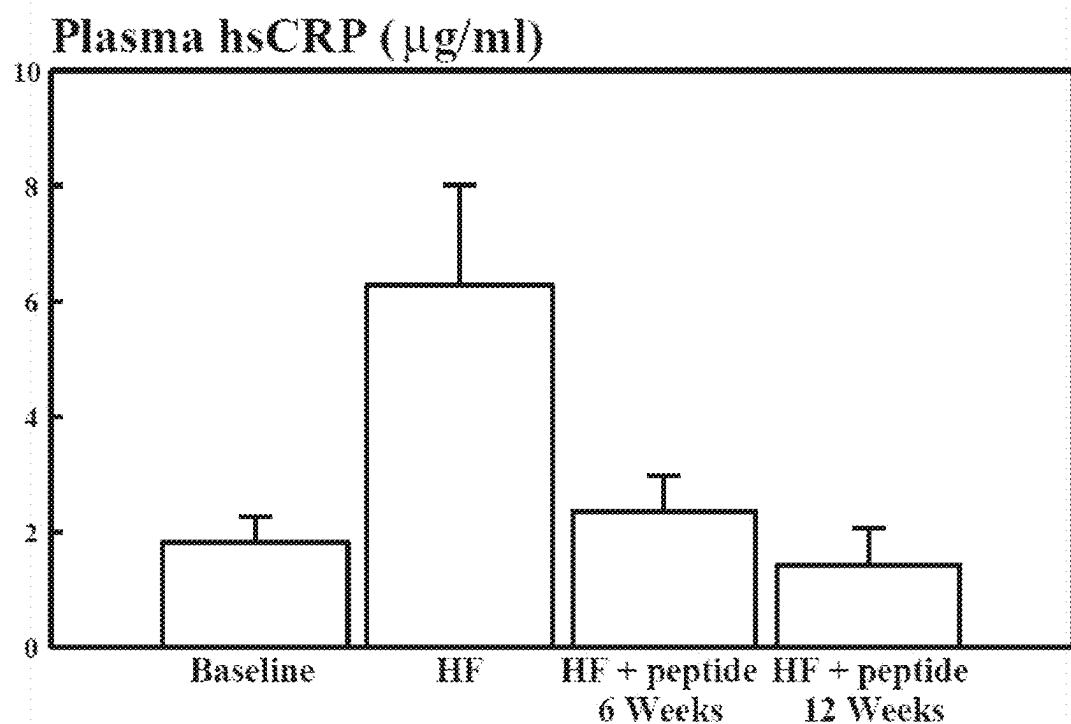


FIG. 2

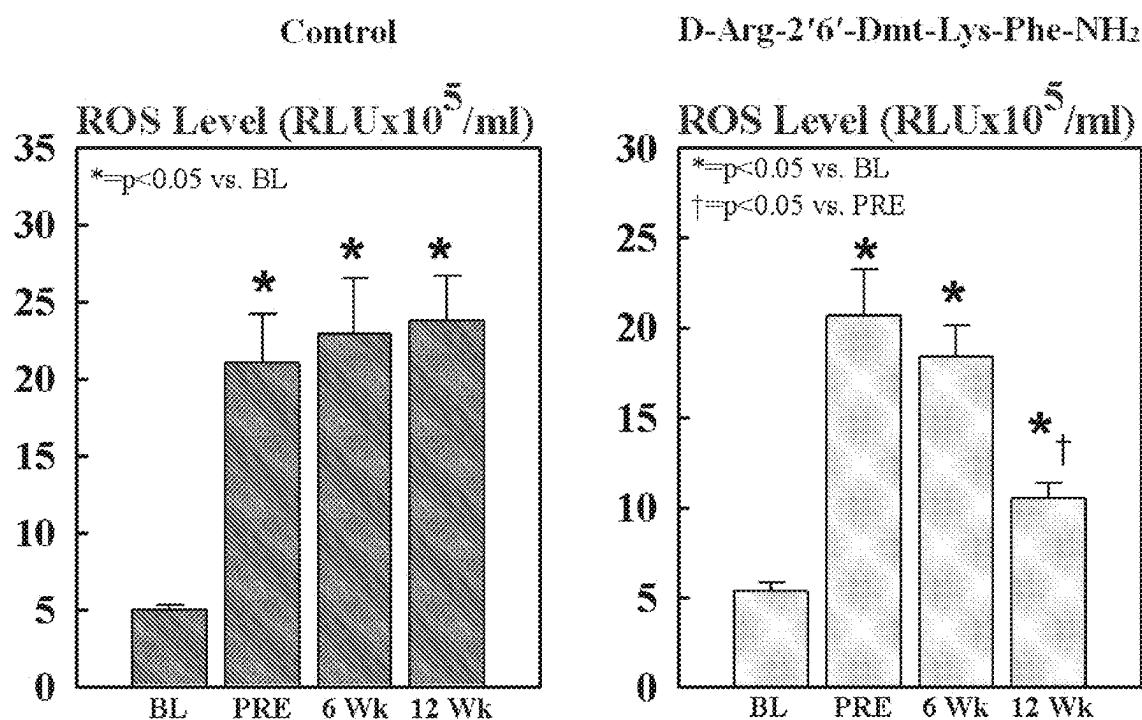


FIG. 3

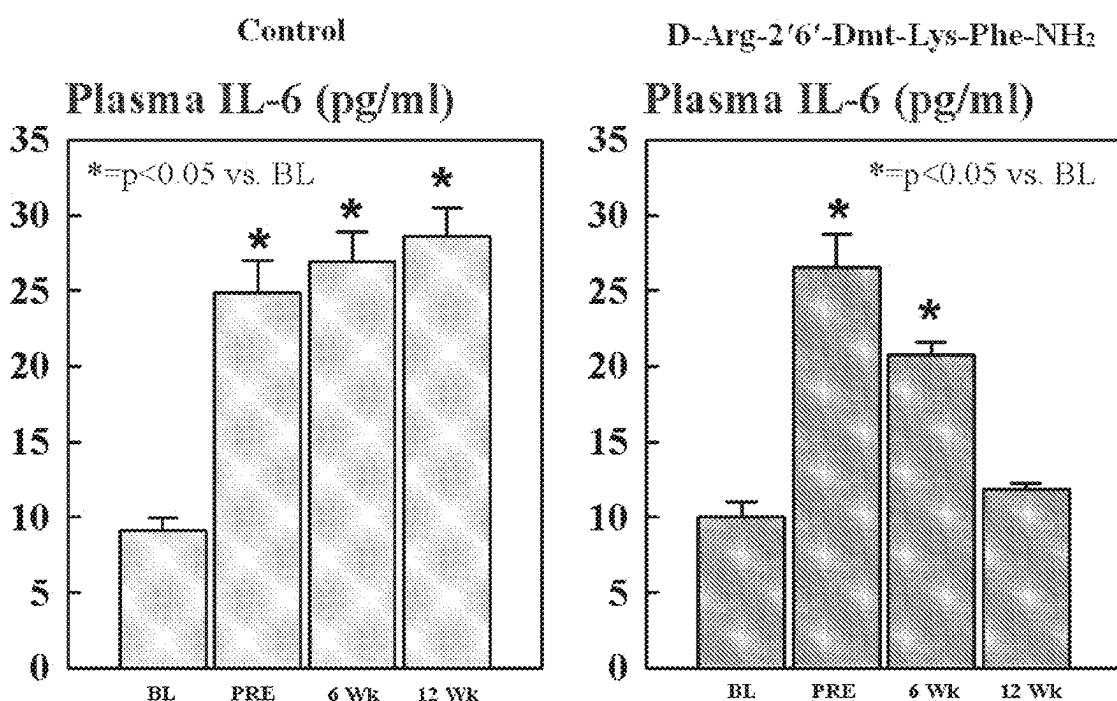


FIG. 4

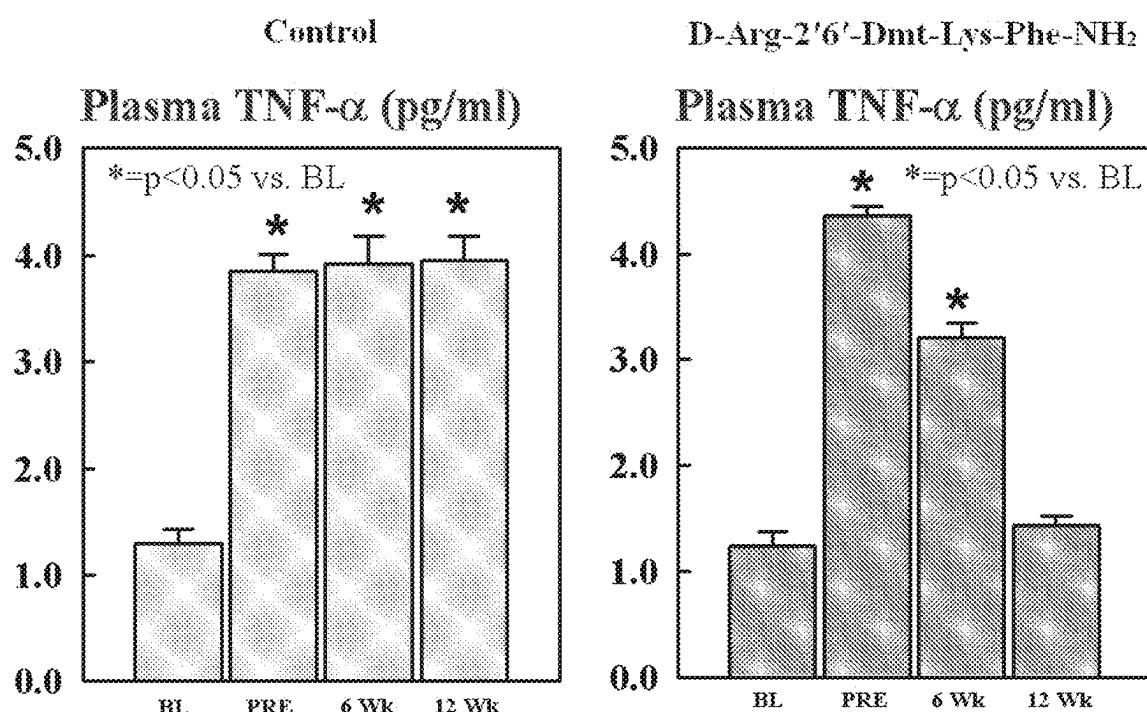


FIG. 5

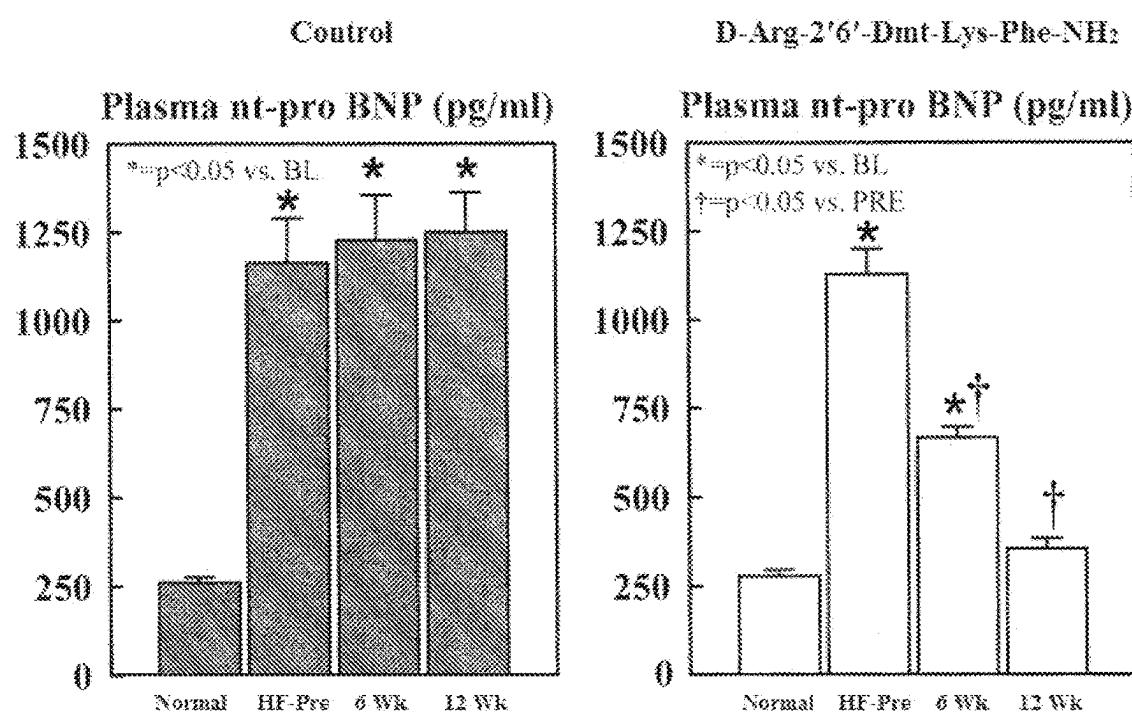


FIG. 6

Cardiolipin - 18:2-18:2-18:2-18:2

(nmol/mg non-collagen protein)

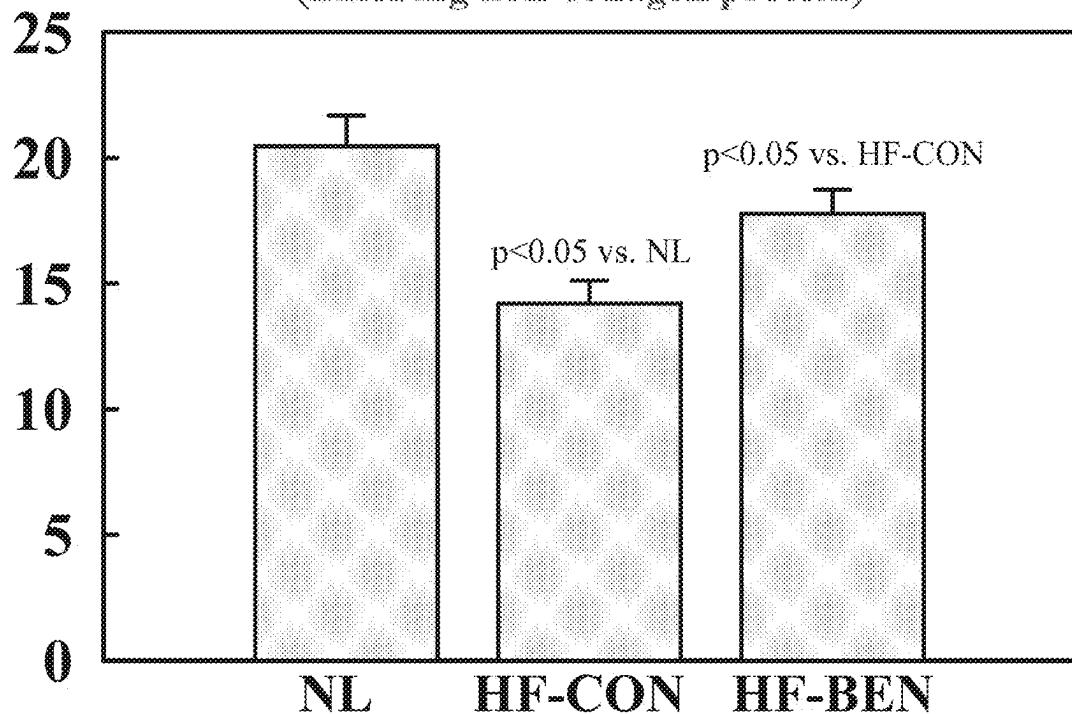


FIG. 7A

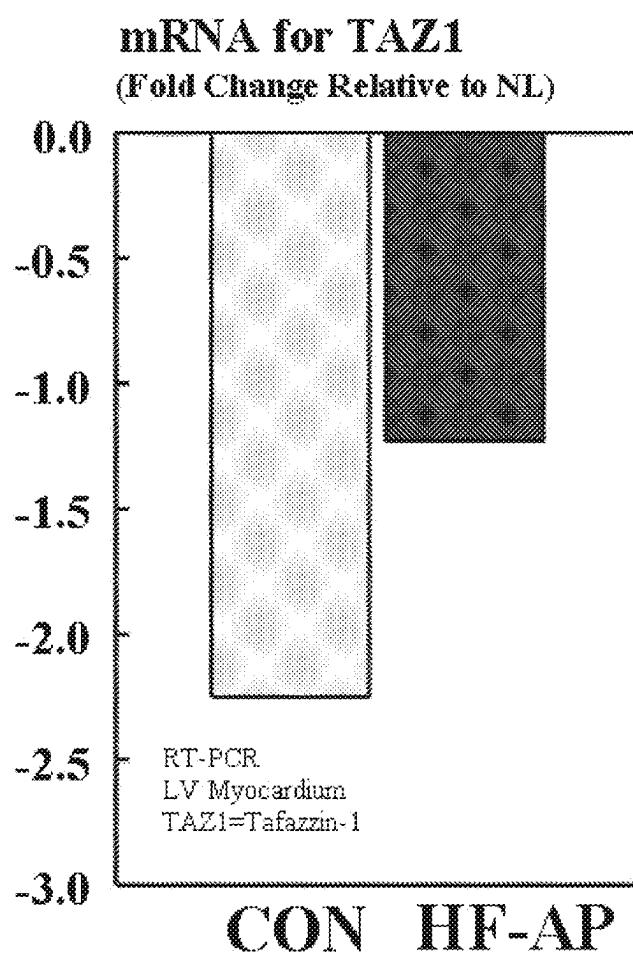


FIG. 7B

**mRNA for MLCLAT1
(Fold Change Relative to NL)**

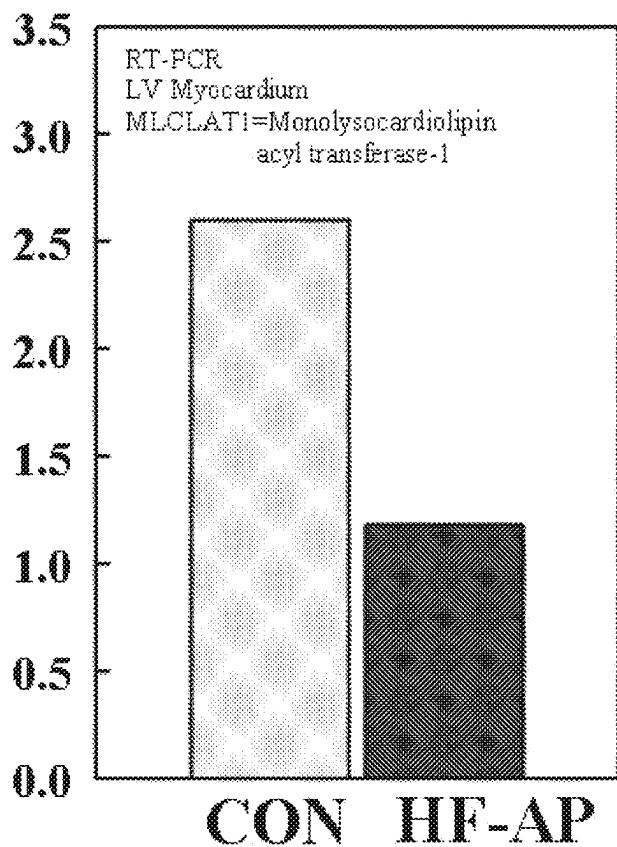
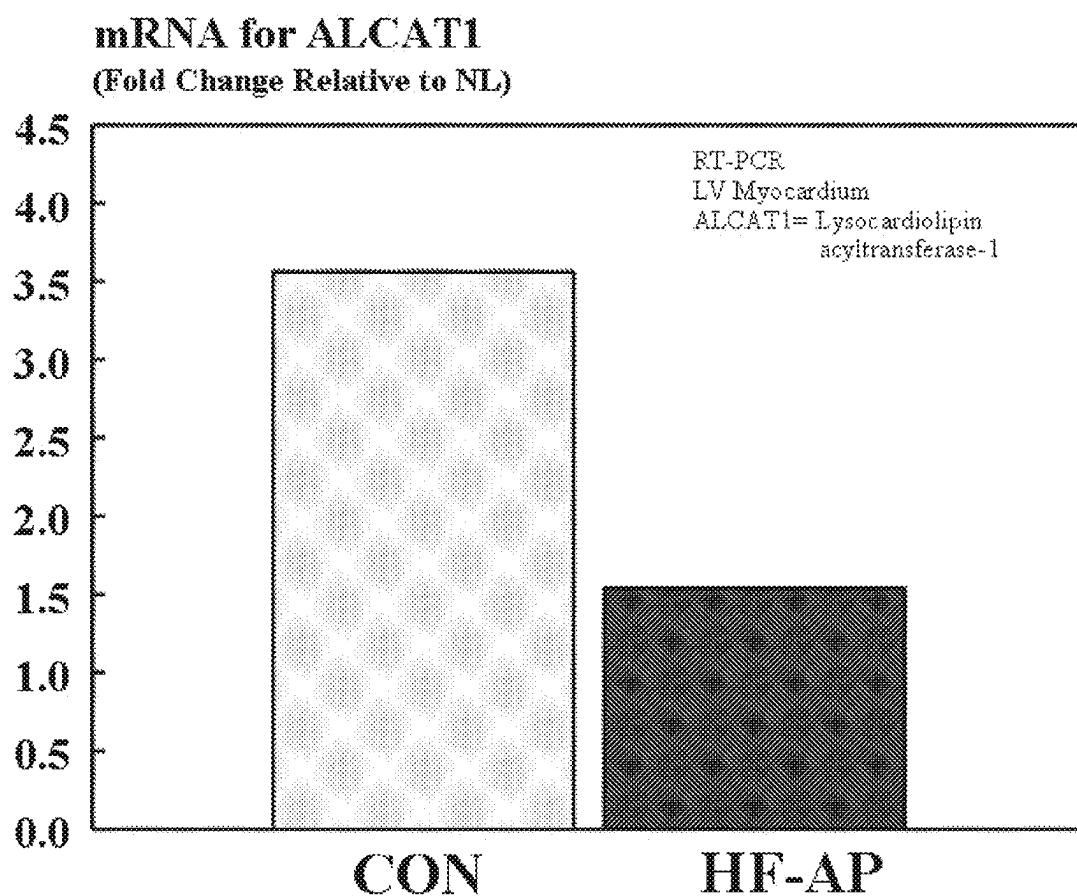


FIG. 7C



METHODS FOR REDUCING RISKS ASSOCIATED WITH HEART FAILURE AND FACTORS ASSOCIATED THEREWITH

CROSS-REFERENCE TO RELATED APPLICATIONS

[0001] This application is a continuation of U.S. patent application Ser. No. 17/838,766, filed Jun. 13, 2022, which is a continuation of U.S. patent application Ser. No. 16/724,076, filed Dec. 20, 2019, which is a continuation of U.S. patent application Ser. No. 14/437,325, filed Apr. 21, 2015, which is the U.S. National Stage of International Patent Application No. PCT/US2013/066228, filed Oct. 22, 2013, which claims priority to U.S. Application No. 61/716,867, filed Oct. 22, 2012, U.S. Application No. 61/822,752, filed May 13, 2013, U.S. Application No. 61/839,743, filed Jun. 26, 2013, and U.S. Application No. 61/839,750, filed Jun. 26, 2013. The entire contents of these applications are incorporated herein by reference in their entireties.

SEQUENCE LISTING

[0002] The instant application contains a Sequence Listing which has been submitted electronically in XML format and is hereby incorporated by reference in its entirety. Said XML copy, created on May 4, 2025, is named 091151-1613_SL.xml and is 213,175 bytes in size.

TECHNICAL FIELD

[0003] The present technology relates generally to compositions and methods for preventing or treating heart failure, reducing risk factors associated with heart failure, and/or reducing the likelihood (risk) or severity of heart failure and/or preventing or treating left ventricular remodeling. In particular, the present technology relates to administering aromatic-cationic peptides in effective amounts to reduce or normalize levels of C-reactive protein, TNF-alpha, IL-6, or reactive oxygen species ("ROS"), brain natriuretic peptide, cardiac troponin I, and/or reduce expression of ALCAT 1, MLCL AT1 in mammalian subjects. The present technology also relates to administering aromatic-cationic peptides in effective amounts to increase and/or normalize expression of Taz1 and/or mitochondrial ATP-sensitive potassium channel activity.

BACKGROUND

[0004] The following description is provided to assist the understanding of the reader. None of the information provided or references cited is admitted to be prior art to the present invention.

[0005] Heart failure is a leading cause of mortality and morbidity worldwide. In the United States, it affects nearly 5 million people and is the only major cardiovascular disorder on the rise. It is estimated that 400,000 to 700,000 new cases of heart failure are diagnosed each year in the U.S. and the number of deaths in the U.S. attributable to this condition has more than doubled since 1979, currently averaging 250,000 annually. Although heart failure affects people of all ages, the risk of heart failure increases with age and is most common among older people. Accordingly, the number of people living with heart failure is expected to increase significantly as the elderly population grows over the next few decades. The causes of heart failure have been linked to various disorders including coronary artery dis-

ease, atherosclerosis, past myocardial infarction, hypertension, abnormal heart valves, cardiomyopathy or myocarditis, congenital heart disease, severe lung disease, diabetes, severe anemia, hyperthyroidism, arrhythmia or dysrhythmia.

[0006] Heart failure (HF), also called congestive heart failure, is commonly characterized by decreased cardiac output, decreased cardiac contractility, abnormal diastolic compliance, reduced stroke volume, and pulmonary congestion. The clinical manifestations of heart failure reflect a decrease in the myocardial contractile state and a reduction in cardiac output. Apart from deficiencies in cardiac contractility, the HF disease state may arise from left ventricular failure, right ventricular failure, biventricular failure, systolic dysfunction, diastolic dysfunction, and pulmonary effects. A progressive decrease in the contractile function of cardiac muscle, associated with heart disease, often leads to hypoperfusion of critical organs.

[0007] Following myocardial infarction there is a dynamic and progressive left ventricle (LV) remodeling that contributes to LV dilation, heart failure, and death. LV remodeling increases LV wall stress, which leads to an increase in oxygen demand. To help compensate for the loss of myocardium and reduced stroke volume, the LV develops global dilation and the non-infarcted wall of the LV develops eccentric hypertrophy. As the ventricle dilates, the dilation process initially helps to compensate for reduced stroke volume. However, eventually progressive dilatation and hypertrophy lead to congestive heart failure. One of the strongest predictors of death one year post myocardial infarction is the volume of the left ventricle.

SUMMARY

[0008] The present technology relates generally to the use of aromatic-cationic peptides to treat or prevent heart failure. The present technology also relates to reducing the level of C-reactive protein, TNF-alpha, IL-6, or reactive oxygen species ("ROS"), brain natriuretic peptide, and cardiac troponin I in a subject in need thereof by administering a therapeutically effective amount of an aromatic-cationic peptide, or a pharmaceutically acceptable salt thereof, e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3). In some embodiments, reducing the level of C-reactive protein, TNF-alpha, IL-6, ROS, or cardiac troponin I is useful for the treatment or prevention of heart failure, reducing risk factors associated with heart failure, and/or reducing the likelihood (risk) or severity of heart failure in the subject.

[0009] The present technology also relates to the treatment or prevention of left ventricular remodeling in mammals through administration of therapeutically effective amounts of aromatic-cationic peptides to subjects in need thereof. In some embodiments, the aromatic-cationic peptides stabilize mitochondrial biogenesis in cardiac tissues. In some embodiments, administration of aromatic-cationic peptides to a subject in need thereof leads to a decrease in brain natriuretic peptide (as measure by a decrease in NT-pro BNP), which correlates to a reduction in LV remodeling. In some embodiments, administration of aromatic-cationic peptides to a subject in need thereof leads to a decrease in cardiac troponin I, which correlates to a reduction in LV remodeling.

[0010] In one aspect, the disclosure provides a treating or preventing heart failure comprising administering to the mammalian subject a therapeutically effective amount of an

aromatic-cationic peptide or a pharmaceutically acceptable salt thereof, e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3). In some embodiments, the aromatic-cationic peptide is a peptide having:

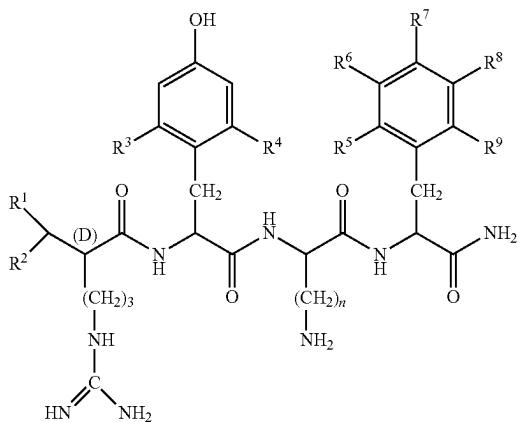
- [0011] at least one net positive charge;
- [0012] a minimum of four amino acids;
- [0013] a maximum of about twenty amino acids;
- [0014] a relationship between the minimum number of net positive charges (p_m) and the total number of amino acid residues (r) wherein $3p_m$ is the largest number that is less than or equal to $r+1$; and a relationship between the minimum number of aromatic groups (a) and the total number of net positive charges (p_r) wherein $2a$ is the largest number that is less than or equal to p_r+1 , except that when a is 1, p_r may also be 1. In particular embodiments, the mammalian subject is a human.

[0015] In some embodiments, $2p_m$ is the largest number that is less than or equal to $r+1$, and a may be equal to p_r . The aromatic-cationic peptide may be a water-soluble peptide having a minimum of two or a minimum of three positive charges.

[0016] In some embodiments, the peptide comprises one or more non-naturally occurring amino acids, for example, one or more D-amino acids. In some embodiments, the C-terminal carboxyl group of the amino acid at the C-terminus is amidated. In certain embodiments, the peptide has a minimum of four amino acids. The peptide may have a maximum of about 6, a maximum of about 9, or a maximum of about 12 amino acids.

[0017] In some embodiments, the peptide comprises a tyrosine or a 2',6'-dimethyltyrosine (dimethyltyrosine is represented by Dmt) residue at the N-terminus. For example, the peptide may have the formula Tyr-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 4) or 2',6'-Dmt-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 5). In another embodiment, the peptide comprises a phenylalanine or a 2',6'-dimethylphenylalanine residue at the N-terminus. For example, the peptide may have the formula Phe-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 6) or 2',6'-Dmp-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 7). In a particular embodiment, the aromatic-cationic peptide has the formula D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3).

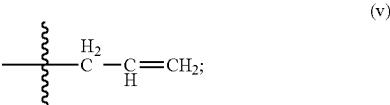
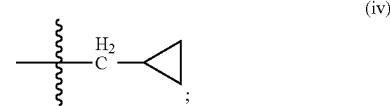
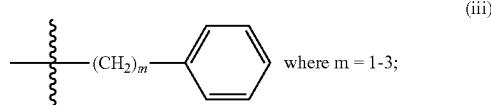
[0018] In some embodiments, the peptide is defined by formula I:



[0019] wherein R¹ and R² are each independently selected from

[0020] (i) hydrogen;

[0021] (ii) linear or branched C₁-C₆ alkyl;



[0022] R³ and R⁴ are each independently selected from

[0023] (i) hydrogen;

[0024] (ii) linear or branched C₁-C₆ alkyl;

[0025] (iii) C₁-C₆ alkoxy;

[0026] (iv) amino;

[0027] (v) C₁-C₄ alkylamino;

[0028] (vi) C₁-C₄ dialkylamino;

[0029] (vii) nitro;

[0030] (viii) hydroxyl;

[0031] (ix) halogen, where "halogen" encompasses chloro, fluoro, bromo, and iodo;

[0032] R⁵, R⁶, R⁷, R⁸, and R⁹ are each independently selected from

[0033] (i) hydrogen;

[0034] (ii) linear or branched C₁-C₆ alkyl;

[0035] (iii) C₁-C₆ alkoxy;

[0036] (iv) amino;

[0037] (v) C₁-C₄ alkylamino;

[0038] (vi) C₁-C₄ dialkylamino;

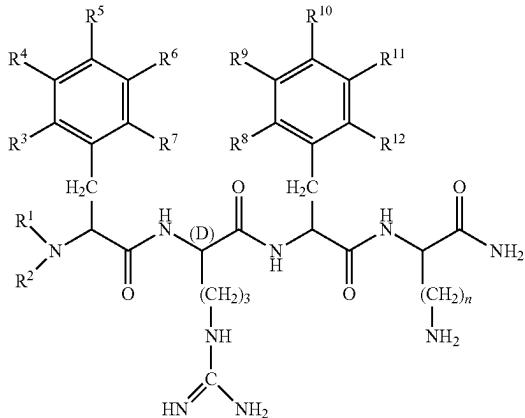
[0039] (vii) nitro;

[0040] (viii) hydroxyl;

[0041] (ix) halogen, where "halogen" encompasses chloro, fluoro, bromo, and iodo; and n is an integer from 1 to 5.

[0042] In some embodiments, R¹ and R² are hydrogen; R³ and R⁴ are methyl; R⁵, R⁶, R⁷, R⁸, and R⁹ are all hydrogen; and n is 4.

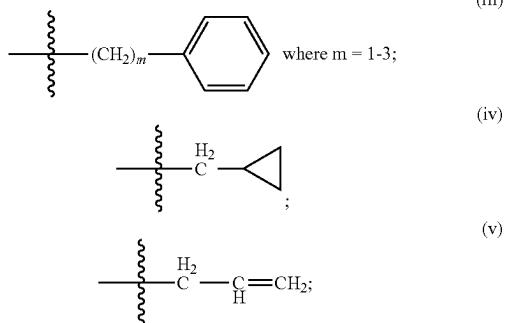
[0043] In some embodiments, the peptide is defined by formula II:



[0044] wherein R¹ and R² are each independently selected from

[0045] (i) hydrogen;

[0046] (ii) linear or branched C₁-C₆ alkyl;



R³, R⁴, R⁵, R⁶, R⁷, R⁸, R⁹, R¹⁰, R¹¹ and R¹² are each independently selected from

[0047] (i) hydrogen;

[0048] (ii) linear or branched C₁-C₆ alkyl;

[0049] (iii) C₁-C₆ alkoxy;

[0050] (iv) amino;

[0051] (v) C₁-C₄ alkylamino;

[0052] (vi) C₁-C₄ dialkylamino;

[0053] (vii) nitro;

[0054] (viii) hydroxyl;

[0055] (ix) halogen, where "halogen" encompasses chloro, fluoro, bromo, and iodo; and n is an integer from 1 to 5.

[0056] In some embodiments, R¹, R², R³, R⁴, R⁵, R⁶, R⁷, R⁸, R⁹, R¹⁰, R¹¹, and R¹² are all hydrogen; and n is 4. In another embodiment, R¹, R², R³, R⁴, R⁵, R⁶, R⁷, R⁸, R⁹, and R¹¹ are all hydrogen; R⁸ and R¹² are methyl; R¹⁰ is hydroxyl; and n is 4.

[0057] In one aspect, the present technology provides methods for reducing the level of one or more of C-reactive protein, TNF-alpha, interleukin-6, reactive oxygen species, and cardiac troponin I in a mammalian subject in need thereof. In some embodiments, the method includes admin-

istering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof.

[0058] Additionally or alternatively, in another aspect the present technology provides methods for preventing, treating or ameliorating heart failure in a mammalian subject having an increased level of one or more of C-reactive protein, TNF-alpha, interleukin-6, reactive oxygen species, and cardiac troponin I. In some embodiments, the method includes administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof.

[0059] In some embodiments, the subject has been diagnosed with heart failure. In some embodiments, the heart failure results from hypertension; ischemic heart disease; exposure to a cardiotoxic compound; myocarditis; thyroid disease; viral infection; gingivitis; drug abuse; alcohol abuse; pericarditis; atherosclerosis; vascular disease; hypertrophic cardiomyopathy; acute myocardial infarction; left ventricular systolic dysfunction; coronary bypass surgery; starvation; an eating disorder; or a genetic defect.

[0060] In some embodiments, the subject has at least one risk factor associated with heart failure selected from the group consisting of high blood pressure; coronary artery disease; heart attack; irregular heartbeats; diabetes; taking diabetes medications rosiglitazone or pioglitazone; sleep apnea; congenital heart defects; viral infection; alcohol use; obesity; smoking; sedentary lifestyle; high cholesterol; family history of heart failure; stress; and kidney conditions.

[0061] In some embodiments, the peptide is administered orally, topically, systemically, intravenously, subcutaneously, intraperitoneally, or intramuscularly.

[0062] Additionally or alternatively, in some embodiments, the peptide is administered separately, sequentially or simultaneously administering a cardiovascular agent to the subject. In some embodiments, the cardiovascular agent is one or more of the following agents: an anti-arrhythmia agent, a vasodilator, an anti-anginal agent, a corticosteroid, a cardiotropine, a diuretic, a sedative, an angiotensin converting enzyme (ACE) inhibitor, an angiotensin II antagonist, a thrombolytic agent, a calcium channel blocker, a thromboxane receptor antagonist, a radical scavenger, an anti-platelet drug, a β-adrenalin receptor blocking drug, α-receptor blocking drug, a sympathetic nerve inhibitor, a digitalis formulation, an inotrope, and an antihyperlipidemic drug.

[0063] In some embodiments, the peptide is D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) in the form of a pharmaceutically acceptable salt. In some embodiments, the salt comprises acetate or trifluoroacetate salt.

[0064] Additionally or alternatively, in some aspects, a method is provided for improving LV function in a mammalian subject. In some embodiments, the subject has an increased level of one or more of C-reactive protein, reactive oxygen species, interleukin-6, TNF-alpha, and cardiac troponin I. In some embodiments, the method comprises: administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof.

[0065] In some embodiments, the mammalian subject has suffered or is likely to suffer heart failure, myocardial infarction, or other stenotic or vascular event.

[0066] In some embodiments, the peptide is D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) in the form of a pharmaceutically acceptable salt. In some embodiments, the salt comprises acetate or trifluoroacetate salt.

[0067] In one aspect, the present technology provides methods for reducing the level of Nt-pro BNP and/or cardiac troponin I in a mammalian subject in need thereof, the method comprising: administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof. In some embodiments, the subject has suffered acute myocardial infarction. In some embodiments, a reduction of Nt-pro BNP and/or cardiac troponin I is an indicator of an effective prevention, treatment, or amelioration of LV remodeling.

[0068] In one aspect, the present technology provides methods for reducing the expression of MLCL AT1 or ALCAT1 in a mammalian subject in need thereof. In some embodiments, the method includes administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof. In some embodiments, the subject has been diagnosed with heart failure.

[0069] In another aspect, the present technology provides methods increasing the expression of Taz1 in a mammalian subject in need thereof. In some embodiments, the method includes administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof. In some embodiments, the subject has been diagnosed with heart failure.

[0070] In another aspect, the present technology provides methods for reducing the risk of heart failure in a mammalian subject having an increased expression of MLCL AT1 or ALCAT1 and/or decreased expression of Taz1. In some embodiments, the method includes administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof.

[0071] In another aspect, the present technology provides methods for stabilizing cardiolipin remodeling in a mammalian subject having or suspected of having heart failure. In some embodiments, the mammalian subject has an increased expression of MLCL AT1 or ALCAT1 and/or decreased expression of Taz1. In some embodiments, the cardiolipin is 18:2 species of cardiolipin.

[0072] In one aspect, the present technology provides methods for reducing the level of cardiac troponin I in a mammalian subject in need thereof. In some embodiments, the method includes administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof.

[0073] Additionally or alternatively, in another aspect the present technology provides methods for preventing, treating or ameliorating heart failure in a mammalian subject having an increased level of cardiac troponin I. In some embodiments, the method includes administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof.

[0074] In some embodiments, the subject has been diagnosed with heart failure. In some embodiments, the heart failure results from hypertension; ischemic heart disease;

exposure to a cardiotoxic compound; myocarditis; thyroid disease; viral infection; gingivitis; drug abuse; alcohol abuse; pericarditis; atherosclerosis; vascular disease; hypertrophic cardiomyopathy; acute myocardial infarction; left ventricular systolic dysfunction; coronary bypass surgery; starvation; an eating disorder; or a genetic defect.

[0075] In some embodiments, the subject has at least one risk factor associated with heart failure selected from the group consisting of high blood pressure; coronary artery disease; heart attack; irregular heartbeats; diabetes; taking diabetes medications rosiglitazone or pioglitazone; sleep apnea; congenital heart defects; viral infection; alcohol use; obesity; smoking; sedentary lifestyle; high cholesterol; family history of heart failure; stress; and kidney conditions.

[0076] In some embodiments, the peptide is administered orally, topically, systemically, intravenously, subcutaneously, intraperitoneally, or intramuscularly.

[0077] Additionally or alternatively, in some embodiments, the peptide is administered separately, sequentially or simultaneously administering a cardiovascular agent to the subject. In some embodiments, the cardiovascular agent is one or more of the following agents: an anti-arrhythmia agent, a vasodilator, an anti-anginal agent, a corticosteroid, a cardiotropine, a diuretic, a sedative, an angiotensin converting enzyme (ACE) inhibitor, an angiotensin II antagonist, a thrombolytic agent, a calcium channel blocker, a thromboxane receptor antagonist, a radical scavenger, an anti-platelet drug, a β-adrenaline receptor blocking drug, α-receptor blocking drug, a sympathetic nerve inhibitor, a digitalis formulation, an inotrope, and an antihyperlipidemic drug.

[0078] In some embodiments, the peptide is D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) in the form of a pharmaceutically acceptable salt. In some embodiments, the salt comprises acetate or trifluoroacetate salt.

[0079] In another aspect, the present technology provides methods for increasing mitochondrial ATP-sensitive potassium channel (mK ATP) activity in a subject in need thereof, the method including administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof.

[0080] Additionally or alternatively, in some aspects, the present technology provides methods for reducing the risk of heart failure in a mammalian subject having a decreased mK ATP activity, the method including administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof.

[0081] In some embodiments, the subject has been diagnosed with heart failure. In some embodiments, the heart failure results from hypertension; ischemic heart disease; exposure to a cardiotoxic compound; myocarditis; thyroid disease; viral infection; gingivitis; drug abuse; alcohol abuse; pericarditis; atherosclerosis; vascular disease; hypertrophic cardiomyopathy; acute myocardial infarction; left ventricular systolic dysfunction; coronary bypass surgery; starvation; an eating disorder; or a genetic defect. In some embodiments, the peptide is administered orally, topically, systemically, intravenously, subcutaneously, intraperitoneally, or intramuscularly. In some embodiments, the pharmaceutically acceptable salt comprises acetate or trifluoroacetate salt.

[0082] In another aspect, the present technology provides methods for stabilizing mitochondria in a mammalian subject having or suspected of having heart failure. In some embodiments, the mammalian subject has a decreased activity of mK ATP.

BRIEF DESCRIPTION OF THE FIGURES

[0083] FIG. 1 is a graph showing levels of C-reactive protein as determined by high-sensitivity assay after 6 weeks or 12 weeks of treatment with the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) compared to baseline levels and untreated controls.

[0084] FIG. 2 is a graph showing levels of ROS after 6 weeks or 12 weeks of treatment with the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) compared to baseline levels and untreated controls.

[0085] FIG. 3 is a graph showing levels of interleukin-6 (IL-6) after 6 weeks or 12 weeks of treatment with the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) compared to baseline levels and untreated controls.

[0086] FIG. 4 is a graph showing levels of tumor necrosis factor alpha (TNF- α) after 6 weeks or 12 weeks of treatment with the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) compared to baseline levels and untreated controls.

[0087] FIG. 5 is a graph showing levels of NT-pro BNP (N-terminal pro-brain natriuretic peptide) after 6 weeks or 12 weeks of treatment with the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) compared to baseline levels and untreated controls.

[0088] FIG. 6 is a chart showing the effects of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on cardiolipin content, cardiolipin species 18:2-18:2-18:2-18:2, in a heart failure model.

[0089] FIG. 7A is a chart showing the effect of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on Taz1 expression in a heart failure model.

[0090] FIG. 7B is a chart showing the effect of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on MLCL AT1 expression in a heart failure model.

[0091] FIG. 7C is a chart showing the effect of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on ALCAT1 expression in a heart failure model.

DETAILED DESCRIPTION

[0092] It is to be appreciated that certain aspects, modes, embodiments, variations and features of the invention are described below in various levels of detail in order to provide a substantial understanding of the present invention. The definitions of certain terms as used in this specification are provided below. Unless defined otherwise, all technical and scientific terms used herein generally have the same meaning as commonly understood by one of ordinary skill in the art to which this invention belongs.

[0093] As used in this specification and the appended claims, the singular forms “a”, “an” and “the” include plural referents unless the content clearly dictates otherwise. For example, reference to “a cell” includes a combination of two or more cells, and the like.

[0094] As used herein, the “administration” of an agent, drug, or peptide to a subject includes any route of introducing or delivering to a subject a compound to perform its

intended function. Administration can be carried out by any suitable route, including orally, intranasally, parenterally (intravenously, intramuscularly, intraperitoneally, or subcutaneously), or topically. Administration includes self-administration and the administration by another.

[0095] As used herein, the term “amino acid” includes naturally-occurring amino acids and synthetic amino acids, as well as amino acid analogs and amino acid mimetics that function in a manner similar to the naturally-occurring amino acids. Naturally-occurring amino acids are those encoded by the genetic code, as well as those amino acids that are later modified, e.g., hydroxyproline, γ -carboxyglutamate, and O-phosphoserine. Amino acid analogs refers to compounds that have the same basic chemical structure as a naturally-occurring amino acid, i.e., an α -carbon that is bound to a hydrogen, a carboxyl group, an amino group, and an R group, e.g., homoserine, norleucine, methionine sulfoxide, methionine methyl sulfonium. Such analogs have modified R groups (e.g., norleucine) or modified peptide backbones, but retain the same basic chemical structure as a naturally-occurring amino acid. Amino acid mimetics refers to chemical compounds that have a structure that is different from the general chemical structure of an amino acid, but that functions in a manner similar to a naturally-occurring amino acid. Amino acids can be referred to herein by either their commonly known three letter symbols or by the one-letter symbols recommended by the IUPAC-IUB Biochemical Nomenclature Commission.

[0096] As used herein, the term “chronic,” with reference to administration, refers to administration of a therapeutic agent, such as an aromatic-cationic peptide, for about 3 days, about 4 days, about 5 days, about 6 days, about 1 week, about 2 weeks, about 3 weeks, 4 weeks, 5 weeks 6 weeks, about 2 months, about 3 months, about 6 months, about 9 months, about 1 year or longer. In some embodiments, chronic administration includes administration once per day, twice per day, 3-5 times per day, every other day, every third day, once per week or once per month.

[0097] As used herein, the term “effective amount” or “therapeutically effective amount” refers to a quantity sufficient to achieve a desired therapeutic and/or prophylactic effect, e.g., an amount which results in the decrease of (e.g., normalization of) levels of one or more of, e.g., C-reactive protein, interleukin 6, ROS, TNF-alpha, cardiac troponin I, NT-pro BNP, MLCL AT1, or ALCAT1 in a subject, and/or an amount which is sufficient to prevent, ameliorate, or treat left ventricle (LV) remodeling and/or improvement of LV function and/or an amount which results in the increase of (e.g., normalization of) expression levels of, e.g., Taz1 and/or increase of mK ATP in a subject in need thereof. In the context of therapeutic or prophylactic applications, in some embodiments, the amount of a composition administered to the subject will depend on the levels of C-reactive protein, interleukin 6, ROS, TNF-alpha, NT-pro BNP, or cardiac troponin I, the expression of MLCL AT1, ALCAT1, or Taz1, and/or the activity of mK ATP in the subject, the type and severity of the disease and on the characteristics of the individual, such as general health, age, sex, body weight and tolerance to drugs. In some embodiments, it will also depend on the degree, severity and type of disease. The skilled artisan will be able to determine appropriate dosages depending on these and other factors. The compositions can also be administered in combination with one or more additional therapeutic compounds. In the methods described

herein, the aromatic-cationic peptides may be administered to a subject having one or more signs or symptoms or risk factors of heart failure, such as cardiomegaly, tachypnea, hepatomegaly, and myocardial infarction. For example, in some embodiments, a therapeutically effective amount of the aromatic-cationic peptides includes levels in which the level of C-reactive protein, interleukin 6, ROS, TNF-alpha, and/or cardiac troponin I is reduced in a subject after administration. Additionally or alternatively, in some embodiments, a therapeutically effective amount prevents, ameliorates, or treats LV remodeling and/or improves LV function. Additionally or alternatively, in some embodiments, a therapeutically effective amount of the aromatic-cationic peptides includes levels in which the expression of MLCL AT1 or ALCAT1 is reduced in a subject in need thereof after administration. Additionally or alternatively, in some embodiments, a therapeutically effective amount of an aromatic-cationic peptide includes levels in which the expression of Taz1 is increased in a subject in need thereof after administration. Additionally or alternatively, in some embodiments, a therapeutically effective amount of the aromatic-cationic peptides includes levels in which the activity of mK ATP is increased. In some embodiments, a therapeutically effective amount also reduces or ameliorates the physiological effects of a heart failure and/or the risk factors of heart failure, and/or the likelihood of heart failure. In some embodiments, an effective amount of an aromatic-cationic peptide is an amount sufficient to decrease levels of brain natriuretic peptide in a subject, e.g., to a normal or control level, for that subject.

[0098] As used herein, the terms “congestive heart failure” (CHF), “chronic heart failure,” “acute heart failure,” and “heart failure” are used interchangeably, and refer to any condition characterized by abnormally low cardiac output in which the heart is unable to pump blood at an adequate rate or in adequate volume. When the heart is unable to adequately pump blood to the rest of the body, or when one or more of the heart valves becomes stenotic or otherwise incompetent, blood can back up into the lungs, causing the lungs to become congested with fluid. If this backward flow occurs over an extended period of time, heart failure can result. Typical symptoms of heart failure include shortness of breath (dyspnea), fatigue, weakness, difficulty breathing when lying flat, and swelling of the legs, ankles or abdomen (edema). Causes of heart failure are related to various disorders including coronary artery disease, systemic hypertension, cardiomyopathy or myocarditis, congenital heart disease, abnormal heart valves or valvular heart disease, severe lung disease, diabetes, severe anemia hyperthyroidism, arrhythmia or dysrhythmia and myocardial infarction. The primary signs of congestive heart failure are cardiomegaly (enlarged heart), tachypnea (rapid breathing; occurs in the case of left side failure) and hepatomegaly (enlarged liver; occurs in the case of right side failure).

[0099] As used herein, the term “hypertensive cardiomyopathy” refers to a weakened heart caused by the effects of hypertension (high blood pressure). Over time, uncontrolled hypertension causes weakness of the heart muscle. As hypertensive cardiomyopathy worsens, it can lead to congestive heart failure. Early symptoms of hypertensive cardiomyopathy include cough, weakness, and fatigue. Additional symptoms of hypertensive cardiomyopathy include leg swelling, weight gain, difficulty breathing when

lying flat, increasing shortness of breath with activity, and waking in the middle of the night short of breath.

[0100] As used herein, the term “left ventricle (LV) remodeling” has the meaning known to those of skill in the art, and refers to a condition, typically following myocardial infarction. Following myocardial infarction there is a dynamic and progressive LV remodeling that contributes to LV dilation, heart failure, and death. Within the first week of a myocardial infarction (MI) the necrotic zone thins and stretches (infarct expansion) contributing to regional dilation of the infarct zone. This phenomenon increases LV wall stress, thus, increasing oxygen demand. To help compensate for the loss of myocardium and reduced stroke volume, the LV develops global dilation and the non-infarcted wall of the LV develops eccentric hypertrophy whereby sarcomeres are added on in a circumferential or lengthwise fashion. As the ventricle dilates this process initially helps to compensate for reduced stroke volume, but eventually progressive dilation and hypertrophy lead to congestive heart failure. One of the strongest predictors of death one year post MI is the volume of the left ventricle; the more dilated, the greater the chance of death. Metabolic and functional abnormalities of the non-infarcted myocardium and myocardium at the infarct border zone may contribute to the LV remodeling phenomenon. Abnormalities in mitochondrial structure and function can lead to reduced production of ATP in the very muscle needed to support the weakened heart. Therefore, in some embodiments, aromatic-cationic peptides, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), are useful to treat, ameliorate and/or prevent LV remodeling, and/or stabilize and/or enhance the function of remaining viable myocardium in a heart failure subject. By way of example, but not by way of limitation, in some embodiments, the aromatic-cationic peptide reduces Nt-pro BNP and/or cardiac troponin I, wherein the reduction of Nt-pro BNP and/or cardiac troponin I correlates to the decrease or reversal of LV remodeling.

[0101] As used herein, a “normalized” level of CRP, or TNF-alpha, or IL-6, or ROS, or Nt-pro BNP, or cardiac troponin I levels, and/or MLCL AT1, ALCAT 1, or Taz1 expression, and/or mK ATP activity refers to reducing a subject's CRP level, or TNF-alpha level, or IL-6 level, or ROS level, or Nt-pro BNP levels, or cardiac troponin I, or MLCL AT1, or ALCAT 1 levels and/or increasing Taz1 expression or mK ATP activity to the subject's baseline level or baseline range, or reducing the subject's level to a level or range determined as “normal” or “control” level, e.g., via control studies and/or control sampling of the subject over time, or of an appropriate population (e.g., matched by age, ethnicity, disease state, drug treatment regime, weight, sex, etc.). As used herein “control level” refers to a level considered average or normal for the subject, or for an appropriate population of subjects.

[0102] As used herein “reducing” a subject's CRP level, or TNF-alpha level, or IL-6 level, or ROS level, or Nt-pro BNP level, or cardiac troponin I level, or MLCL AT1 expression level, or ALCAT1 expression level means lowering the level of CRP, or TNF-alpha, or IL-6, or ROS, or Nt-pro BNP, or cardiac troponin I, or MLCL AT1, or ALCAT1 expression level in the subject (e.g., a subject's blood CRP level). In some embodiments, reducing CRP level or TNF-alpha level or IL-6 level or ROS level, or Nt-pro BNP level, or cardiac troponin I level, or MLCL AT1, or ALCAT1 level is a reduction by about 1%, about 5%, about 10%, about 15%,

about 20%, about 25%, about 30%, about 35%, about 40%, about 45%, about 50%, about 55%, about 60%, about 65%, about 70%, about 75%, about 80%, about 85%, about 90%, about 95%, or more.

[0103] As used herein “increasing” a subject’s Taz1 expression level or mK ATP activity means increasing the level of Taz1 (e.g., a subject’s Taz1 expression level in left ventricular myocardium) or increasing the activity of mK ATP in the subject. In some embodiments, increasing Taz1 expression level and/or mK ATP activity is an increase by about 1%, about 5%, about 10%, about 15%, about 20%, about 25%, about 30%, about 35%, about 40%, about 45%, about 50%, about 55%, about 60%, about 65%, about 70%, about 75%, about 80%, about 85%, about 90%, about 95%, about 100%, or about 110% or more, e.g., from a baseline or control level. Alternatively, or additionally, in some embodiments, increasing Taz1 expression level is measured as attenuating the reduction of Taz1 by about 0.25 fold to about 0.5 fold, or about 0.5 fold to about 0.75 fold, or about 0.75 fold to about 1.0 fold, or about 1.0 fold to about 1.5 fold, e.g., as compared to a baseline or control level.

[0104] An “isolated” or “purified” polypeptide or peptide is substantially free of cellular material or other contaminating polypeptides from the cell or tissue source from which the agent is derived, or substantially free from chemical precursors or other chemicals when chemically synthesized. For example, an isolated aromatic-cationic peptide would be free of materials that would interfere with diagnostic or therapeutic uses of the agent. Such interfering materials may include enzymes, hormones and other proteinaceous and nonproteinaceous solutes.

[0105] As used herein, the terms “polypeptide,” “peptide,” and “protein” are used interchangeably herein to mean a polymer comprising two or more amino acids joined to each other by peptide bonds or modified peptide bonds, i.e., peptide isosteres. Polypeptide refers to both short chains, commonly referred to as peptides, glycopeptides or oligomers, and to longer chains, generally referred to as proteins. Polypeptides may contain amino acids other than the 20 gene-encoded amino acids. Polypeptides include amino acid sequences modified either by natural processes, such as post-translational processing, or by chemical modification techniques that are well known in the art.

[0106] As used herein, the term “simultaneous” therapeutic use refers to the administration of at least two active ingredients by the same route and at the same time or at substantially the same time.

[0107] As used herein, the term “separate” therapeutic use refers to an administration of at least two active ingredients at the same time or at substantially the same time by different routes.

[0108] As used herein, the term “sequential” therapeutic use refers to administration of at least two active ingredients at different times, the administration route being identical or different. More particularly, sequential use refers to the whole administration of one of the active ingredients before administration of the other or others commences. It is thus possible to administer one of the active ingredients over several minutes, hours, or days before administering the other active ingredient or ingredients. There is no simultaneous treatment in this case.

[0109] As used herein, the terms “treating” or “treatment” or “alleviation” refers to therapeutic treatment, wherein the object is to prevent or slow down (lessen) the targeted

pathologic condition or disorder. A subject is successfully “treated” for heart failure if, after receiving a therapeutic amount of the aromatic-cationic peptides according to the methods described herein, the subject shows observable and/or measurable reduction in or absence of one or more signs and symptoms of heart failure, such as, e.g., cardiac output, myocardial contractile force, cardiomegaly, tachycardia, and/or hepatomegaly. Treating heart failure, as used herein, also refers to treating any one or more of the conditions underlying heart failure, including, without limitation, decreased cardiac contractility, abnormal diastolic compliance, reduced stroke volume, pulmonary congestion, and decreased cardiac output. The terms also apply to a reduction in C-reactive protein, interleukin 6, ROS, TNF-alpha, cardiac troponin I levels, Nt-pro BNP, MLCL AT1 and/or ALCAT 1 in those subjects having higher than a control or “normal” level of C-reactive protein, interleukin 6, ROS, TNF-alpha, cardiac troponin I levels, Nt-pro BNP, MLCL AT1 and/or ALCAT 1. The terms also apply to an increase in Taz1 expression and/or increased mK ATP activity in those subjects having lower than a control or “normal” level of Taz1 or lower activity of mK ATP. Additionally, or alternatively, the terms apply to an observable and/or measurable reduction in or absence of one or more signs and symptoms associated with LV remodeling, such as, e.g., LV stroke volume, improved LV ejection fraction, improved fractional shortening, reduced infarct expansion, improved hemodynamics, reduced scar formation in LV myocardium, and reduced lung volumes. It is also to be appreciated that the various modes of treatment or prevention of medical conditions as described are intended to mean “substantial,” which includes total but also less than total treatment or prevention, and wherein some biologically or medically relevant result is achieved.

[0110] As used herein, “prevention” or “preventing” of a disorder or condition refers to a compound that, in a statistical sample, reduces the occurrence of the disorder or condition in the treated sample relative to an untreated control sample, or delays the onset or reduces the severity of one or more symptoms of the disorder or condition relative to the untreated control sample. As used herein, preventing heart failure includes preventing the initiation of heart failure, delaying the initiation of heart failure, preventing the progression or advancement of heart failure, slowing the progression or advancement of heart failure, delaying the progression or advancement of heart failure, and reversing the progression of heart failure from an advanced to a less advanced stage. As used herein, prevention of heart failure also includes preventing a recurrence of heart failure. As used herein, preventing LV remodeling includes preventing the initiation of LV remodeling, delaying the initiation of LV remodeling, preventing the progression or advancement of LV remodeling, slowing the progression or advancement of LV remodeling, delaying the progression or advancement of LV remodeling, and reversing the progression of LV remodeling from an advanced to a less advanced stage.

[0111] As used herein, “net charge” refers to the balance of the number of positive charges and the number of negative charges carried by the amino acids present in the peptide. In this specification, it is understood that net charges are measured at physiological pH. The naturally occurring amino acids that are positively charged at physiological pH include L-lysine, L-arginine, and L-histidine. The naturally

occurring amino acids that are negatively charged at physiological pH include L-aspartic acid and L-glutamic acid.

[0112] As used herein, the term “pharmaceutically acceptable salt” refers to a salt prepared from a base or an acid which is acceptable for administration to a patient, such as a mammal (e.g., salts having acceptable mammalian safety for a given dosage regime). However, it is understood that the salts are not required to be pharmaceutically acceptable salts, such as salts of intermediate compounds that are not intended for administration to a patient. Pharmaceutically acceptable salts can be derived from pharmaceutically acceptable inorganic or organic bases and from pharmaceutically acceptable inorganic or organic acids. In addition, when a peptide contains both a basic moiety, such as an amine, pyridine or imidazole, and an acidic moiety such as a carboxylic acid or tetrazole, zwitterions may be formed and are included within the term “salt” as used herein.

C-Reactive Protein

[0113] As used herein “C-reactive protein” (CRP) refers to a pentameric polypeptide composed of five identical subunits, which is a member of the pentraxin family of proteins. The CRP subunit is expressed as a 224-amino acid pro-polypeptide; an 18 amino-acid leader sequence is removed to form a mature 206-amino acid CRP unit. Exemplary, non-limiting sequences of the 224-amino acid CRP precursor are provided below in Table 1 (Accession Numbers NP_000558 and CAA39671, respectively).

TABLE 1

Exemplary CRP precursor amino acid sequences
SEQ ID NO: 1 mekllcflvl ts1shafgqt dmsrkafvfp kesdtsyvsl kapl1kplka ftvc1hfyt1 lsstrgysif syatkrqdne ilifwskd1g ysftvggsei lf1evpevtva pvhictswes as1giefwvd gkpvrks1k kgytvgaeas i1lgqeqd1f ggnfegsqsl vgdignvnwmw df1vlspe1n tiylggpfsp nn1nwralky evqgevft1k1 q1wp
SEQ ID NO: 2 mekllcflvl ts1shafgqt dmsrkafvfp kesdtsyvsl kapl1kplka ftvc1hfyt1 lsstrg1v1s rmppr1ktmr ffifwskd1g ysftvggsei lf1evpevtva pvhictswes as1giefwvd gkpvrks1k kgytvgaeas i1lgqeqd1f ggnfegsqsl vgdignvnwmw df1vlspe1n tiylggpfsp nn1nwralky evqgevft1k1 q1wp

[0114] CRP is an acute phase reactant, which is produced by the liver in response to inflammatory stimuli and which circulates in the blood. CRP levels rise in response to acute or chronic inflammation, such as but not limited to inflammation due to infection (e.g., bacterial, viral or fungal), rheumatic and other inflammatory diseases, malignancy, tissue injury or necrosis. Plasma CRP levels of can increase (e.g., 100-fold or more) after severe trauma, bacterial infection, inflammation, surgery or neoplastic proliferation. CRP levels rapidly increase within hours after tissue injury, and it is suggested that CRP is part of the innate immune system and contributes to host defense. Regarding CRP function and the immune system, CRP has been shown to increase LDL uptake into macrophages and enhance the ability of macrophages to form foam cells; inhibit endothelial nitric oxide synthase expression in endothelial cells; increase plasminogen activator inhibitor-1 expression and activity; activate macrophages to secrete tissue factor; up regulate the

expression of adhesion molecules in endothelial cells to attract monocytes to the site of injury. Thus, CRP levels have been utilized as a marker for inflammation and immune response.

[0115] In addition, CRP levels have been correlated with vascular sclerosis and cardiovascular disease or cardiovascular events, such as heart failure and myocardial infarction. Since cardiovascular disease is at least in part an inflammatory process, CRP has been investigated, for example, in the context of arteriosclerosis and subsequent vascular disorders. For example, a chronic, low-level increase of CRP was found to be predictive of the risk of future cardiovascular events, including myocardial infarction, ischemic stroke, peripheral vascular disease and sudden cardiac death. Based on multiple epidemiological and intervention studies, minor CRP elevation (as determined by high-sensitivity CRP assays, “hsCRP”) has been shown to be associated with future cardiovascular risk. HsCRP is discussed in more detail below.

[0116] In addition to its role as a cardiovascular risk marker, CRP has also been shown to participate directly in atherosogenesis, and high levels of CRP mRNA have been found in atherosclerotic plaques. It has been shown that CRP is produced by human artery smooth muscle cells of atherosclerotic lesions in response to inflammatory cytokines; thus, locally produced CRP may participate directly in aspects of atherosogenesis, promoting the development of cardiovascular complications.

[0117] In some embodiments, treatment with an aromatic-cationic peptide, such as, e.g., D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), decreases C-reactive protein levels in the myocardium in mammalian subjects that have suffered or are at risk of suffering heart failure.

[0118] In some embodiments, C-reactive protein levels are decreased by about 0.25 fold to about 0.5 fold, or about 0.5 fold to about 0.75 fold, or about 0.75 fold to about 1.0 fold, or about 1.0 fold to about 1.5 fold, or about 1.5 fold to about 2.0 fold, or about 2.0 fold to about 3.0 fold, or about 3.0 fold to about 5.0 fold, or about 5.0 fold to about 6.0 fold.

Determination of CRP Levels

[0119] There are three broad categories of CRP assays as recognized by the U.S. Food and Drug Administration (FDA): (1) Conventional C-Reactive Protein (CRP) assays; (2) High sensitivity C-Reactive Protein (hsCRP) assays; and (3) Cardiac C-Reactive Protein (cCRP) assays.

Conventional CRP Assays

[0120] Conventional CRP assays typically include qualitative, semi-quantitative and quantitative assays, with indications for use for evaluation of infection, tissue injury, and inflammatory disorders. These assays provide information for the diagnosis, therapy, and monitoring of inflammatory diseases. As discussed previously, CRP is one of the cytokine-induced “acute-phase” proteins whose blood levels rise during a general, unspecific response to infections and non-infectious inflammatory processes. For conventional CRP assays, test values are typically considered clinically significant at levels above 10 mg/L. In apparently healthy person's blood CRP levels are below 5 mg/L, while in various conditions this threshold is often exceeded within four to eight hours after an acute inflammatory event, with CRP values reaching approximately 20 to 500 mg/L. CRP is

a more sensitive and more reliable indicator of acute inflammatory processes than the erythrocyte sedimentation rate (ESR) and leukocyte count. Blood CRP levels rise more rapidly than ESR, and after the disease has subsided CRP values rapidly fall and reach the reference interval often days before ESR has returned to normal.

High Sensitivity CRP Assays

[0121] High sensitivity CRP assays have a range of measurement that extends below the measurement range typical of most conventional CRP assays. This lower range of measurement expands the indications for use to include, by way of example but not by way of limitation, the evaluation of conditions thought to be associated with inflammation in otherwise seemingly healthy individuals, and to evaluate cardiac risk in subjects suffering from or at risk of heart failure. Typically, hsCRP assays measure CRP levels from less than 1 mg/L (e.g., as low as 0.04 mg/ml) to greater than or equal to 10 mg/L. As used herein and as is common in the art "high sensitivity CRP" or "hsCRP" refers to the assay, and also refers to the CRP levels as determined by hsCRP assay. Thus, the statement "a subject's hsCRP level is less than 1 mg/L" means that the subject's CRP level, as determined by a high sensitivity CRP assay, is less than 1 mg/L. Thus, the subject's CRP level is less than 1 mg/L.

Cardiac C-Reactive Protein (cCRP) Assays

[0122] Pursuant to FDA guidelines, cardiac CRP assays are indicated for use as an aid in the identification and stratification of individuals at risk for future cardiovascular disease. When used in conjunction with traditional clinical laboratory evaluation of acute coronary syndromes, cCRP may be useful as an independent marker of prognosis for recurrent events in patients with stable coronary disease or acute coronary syndrome. Cardiac CRP assays, like hsCRP assays, have measurement ranges that extend below the measurement range typical of most conventional CRP assays. The difference between hsCRP and cCRP is not the analyte itself or even the method of the assay, but the additional performance validation required by the FDA to support the expanded intended use in the evaluation of coronary disease. Accordingly, cCRP assays are a species of hsCRP assay. While hsCRP assays are useful to correlate cardiac risk and are used in the art to do so, they are simply not recognized by the FDA for such uses.

Sample Source C-Reactive Protein

[0123] While C-reactive protein levels are typically tested via a subject's blood sample (e.g., whole blood, plasma, serum), the present disclosure is not intended to be limited by sample type. For example, in some embodiments, C-reactive protein levels are determined by evaluating any fluid or tissue sample of a subject, known to or suspected of containing C-reactive protein. Non-limiting examples include plasma, serum, whole blood, urine, sputum, semen, cerebrospinal fluid, pericardial fluid, peritoneal fluid, pleural fluid, synovial fluid, stool samples and nasal aspirates.

Interleukin-6

[0124] Interleukin-6 (IL-6) belongs to a family of pleiotropic and evolutionary conserved cytokines involved in the regulation of stem cells, hematopoiesis, thrombopoiesis, macrophage function, neuron function, acute phase response, bone metabolism, and cardiac hypertrophy. The

family includes, besides IL-6, the cytokines IL-11, cardiotrophin-1, oncostatin-M, leukemia-inhibitory factor, and ciliary neurotrophic factor, which all utilize a common signal-transducing component named gp130 besides their cytokine specific receptors. Recently, elevated soluble IL-6 receptor (IL-6R) levels in heart failure have been reported. Circulating IL-6 exerts a negative inotropic influence on isolated papillary muscle preparations in the hamster model and in atrial strips in humans, potentially modulated by the nitric oxide, β -adrenoceptor pathway, and the ceramide-sphingomyelin pathway. Elevated serum IL-6 levels correlate with reduced contractility, elevated preload, elevated heart rate, and reduced afterload in patients with impaired left ventricular function.

[0125] Regarding the source of IL-6 production, recent murine transgene and knockout data suggest that an intracardiac IL-6/gp130 system exists and is an essential component in the compensatory response to hemodynamic overload. Human myocardium has been suggested to be a source of IL-6 during myocardial infarction, ischemia, reperfusion, rejection, and heart failure. See Wollert, K. and Drexler, H., "The role of interleukin-6 in the failing heart", *Heart Fail Rev.*, 6(2): 95-103 (2001).

[0126] In some embodiments, treatment with an aromatic-cationic peptide, such as, e.g., D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), decreases IL-6 levels in the myocardium in mammalian subjects that have suffered or are at risk of suffering heart failure.

[0127] In some embodiments, IL-6 levels are decreased by about 0.25 fold to about 0.5 fold, or about 0.5 fold to about 0.75 fold, or about 0.75 fold to about 1.0 fold, or about 1.0 fold to about 1.5 fold, or about 1.5 fold to about 2.0 fold, or about 2.0 fold to about 3.0 fold, or about 3.0 fold to about 5.0 fold, or about 5.0 fold to about 8.0 fold.

Tumor Necrosis Factor Alpha

[0128] Tumor necrosis factor alpha (TNF-alpha or TNF- α) is a cytokine involved in systemic inflammation and is a member of a group of cytokines that stimulate the acute phase reaction. It is produced chiefly by activated macrophages, although it can be produced by many other cell types as CD4+ lymphocytes, NK cells and neurons.

[0129] In patients with advanced congestive heart failure (CHF), elevated levels of circulating TNF-alpha and soluble TNF receptors have been found. The pathophysiological implications of activation of the TNF system in CHF seem to rely mainly on its effects on the heart and the endothelium. TNF-alpha exerts a negative inotropic effect both directly and indirectly, this latter being mediated by enhancement of nitric oxide production. Moreover, TNF-alpha has been suggested to trigger the apoptotic process in cardiac myocytes. There is consensus on the detrimental role played by TNF-alpha in CHF further supported by the evidence of a temporal association between TNF activation and transition from asymptomatic to symptomatic CHF. See Ceconi et al., "Tumor necrosis factor in congestive heart failure: a mechanism of disease for the new millennium?" *Prog. Cardiovasc. Dis.*, 41(Suppl 1): 25-30 (1998).

[0130] In some embodiments, treatment with an aromatic-cationic peptide, such as, e.g., D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), decreases TNF-alpha levels in the myocardium in mammalian subjects that have suffered or are at risk of suffering heart failure.

[0131] In some embodiments, TNF-alpha levels are decreased by about 0.25 fold to about 0.5 fold, or about 0.5 fold to about 0.75 fold, or about 0.75 fold to about 1.0 fold, or about 1.0 fold to about 1.5 fold, or about 1.5 fold to about 2.0 fold, or about 2.0 fold to about 3.0 fold, or about 3.0 fold to about 5.0 fold, or about 5.0 fold to about 8.0 fold.

Reactive Oxygen Species

[0132] Reactive oxygen species (ROS) are chemically reactive molecules containing oxygen. Examples include oxygen ions and peroxides. ROS form as a natural byproduct of the normal metabolism of oxygen and have important roles in cell signaling and homeostasis. However, during times of environmental stress ROS levels can increase dramatically. The increased of ROS may result in significant damage to cell structures, cumulatively known as oxidative stress.

[0133] There is evidence suggesting that oxygen-derived free radicals are involved in the pathogenesis of CHF. Studies suggest that the highly toxic radical species damage sub-cellular membranes leading to the disruption in excitation-contractile coupling and eventually the dysfunction of the myocardium. In addition, these radicals destroy nitric oxide, a potent signaling molecule responsible for maintaining cardiovascular tone. Antioxidants hold great promise in minimizing the damage occurring as a result of the excessive generation of the free radicals during ischemia/reperfusion injury and CHF.

[0134] In some embodiments, treatment with an aromatic-cationic peptide, such as, e.g., D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), decreases ROS levels in the myocardium in mammalian subjects that have suffered or are at risk of suffering heart failure.

[0135] In some embodiments, ROS is decreased by about 0.25 fold to about 0.5 fold, or about 0.5 fold to about 0.75 fold, or about 0.75 fold to about 1.0 fold, or about 1.0 fold to about 1.5 fold, or about 1.5 fold to about 2.0 fold, or about 2.0 fold to about 3.0 fold, or about 3.0 fold to about 5.0 fold, or about 5.0 fold to about 8.0 fold.

Brain Natriuretic Peptide

[0136] Brain natriuretic peptide (BNP) is a 32-amino acid polypeptide secreted by the ventricles of the heart in response to excessive stretching of cardiomyocytes. The release of BNP is modulated by calcium ions. BNP in humans is produced mainly in the cardiac ventricles.

[0137] BNP is synthesized as a 134-amino acid prepro-hormone (preproBNP), encoded by the human gene NPPB. The removal of the 25-residue N-terminal signal peptide generates the prohormone, proBNP, which is stored intracellularly as an O-linked glycoprotein. ProBNP is subsequently cleaved by a specific convertase, i.e., furin or corin, into NT-pro BNP, which is a 76-amino acid biologically inactive polypeptide, and BNP, the biologically active 32-amino acid polypeptide. ProBNP and NT-pro BNP are secreted into the blood in equimolar amounts.

[0138] BNP binds to and activates the atrial natriuretic factor in a fashion similar to atrial natriuretic peptide (ANP) but with 10-fold lower affinity. The biological half-life of BNP, however, is twice as long as that of ANP, and that of NT-proBNP is even longer, making these peptides better targets than ANP for diagnostic blood testing.

[0139] The physiologic actions of BNP are similar to those of ANP and include decrease in systemic vascular resistance and central venous pressure as well as an increase in natriuresis. The net effect of BNP and ANP is a decrease in blood volume, which lowers systemic blood pressure and afterload, yielding an increase in cardiac output, partly due to a higher ejection fraction.

[0140] BNP or NT-pro BNP can be used for screening and prognosis of heart failure. Elevated BNP or NT-pro BNP levels can indicate risk for heart failure or indicate occurrence of acute heart failure. Additionally, both are typically increased in patients with left ventricular dysfunction reduction of BNP concentration after treatments, e.g., with ACE inhibitors or R blockers, may reflect the reversal or prevention of the LV remodeling process.

[0141] In some embodiments, treatment with an aromatic-cationic peptide, such as, e.g., D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), decreases NT-pro BNP in the myocardium in mammalian subjects that have suffered or are at risk of suffering heart failure.

[0142] In some embodiments, NT-pro BNP is decreased by about 0.25 fold to about 0.5 fold, or about 0.5 fold to about 0.75 fold, or about 0.75 fold to about 1.0 fold, or about 1.0 fold to about 1.5 fold, or about 1.5 fold to about 2.0 fold.

Cardiac Troponin I

[0143] Troponin is a complex of three regulatory proteins (troponin C, troponin I, and troponin T) that is integral to muscle contraction in skeletal and cardiac muscle, but not smooth muscle. Troponin T and troponin I isoforms from cardiac muscle are structurally different from the corresponding isoforms found in skeletal muscle.

[0144] Troponin T binds to tropomyosin, interlocking them to form a troponin-tropomyosin complex. Troponin I binds to actin in thin myofilaments to hold the troponin-tropomyosin complex in place and decreases troponin C affinity for calcium, thus inhibiting troponin-tropomyosin interactions. Troponin C binds to calcium ions and plays the main role in Ca²⁺ dependent regulation of muscle contraction.

[0145] Since cardiac troponin (cTn) is structurally different from skeletal troponin, the cTn subunits can be measured as a specific biomarker for myocardial injury. cTn is released as a result of myocyte necrosis, myocyte apoptosis, or cardiac troponin degradation, all of which would indicate the worsening of cardiac dysfunction and/or progression of heart failure. Multiple studies have evaluated the association between elevated circulating cTn and adverse clinical outcomes in various HF populations. Despite variations in study design, patient populations, and assay characteristics, there has been a consistent association between cTn elevation and worsened outcomes. Kociol et al., *Journal of the American College of Cardiology*, 56(14): 1071-78 (September 2010).

[0146] Additionally, studies showed that elevated levels of cardiac troponin I may contribute to the progression of heart failure. See, e.g., Göser et al., *Circulation*, 114: 1693-1702 (2006). Elevated levels of cardiac troponin I lead to the formation of cardiac troponin autoantibodies. Increased cardiac troponin autoantibodies increased cardiac inflammation and increased expression of inflammatory cytokines. Increased cardiac troponin autoantibodies produced cardiomyopathic phenotype characterized by myocardial fibrosis, left ventricular dilation, and impaired cardiac function.

[0147] In some embodiments, treatment with an aromatic cationic peptide, such as, e.g., D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), decreases cardiac troponin I in the myocardium in mammalian subjects that have suffered or are at risk of suffering heart failure.

[0148] In some embodiments, cardiac troponin I is decreased by about 0.25 fold to about 0.5 fold, or about 0.5 fold to about 0.75 fold, or about 0.75 fold to about 1.0 fold, or about 1.0 fold to about 1.5 fold, or about 1.5 fold to about 2.0 fold.

Mitochondrial ATP-Sensitive Potassium Channel (mK ATP)

[0149] The reduced form of nicotinamide adenine dinucleotide phosphate (NADPH) is increased in the failing heart and leads to reduced activation of the mitochondrial ATP-sensitive potassium channels (mK ATP). The reduced mK ATP activity leads to ionic deregulation in the mitochondrial environment with subsequent matrix contraction and reduced ATP production. Increased activity or “opening” of mK ATP improves oxidative phosphorylation by promoting matrix swelling, maintains the structure of the inner mitochondrial membrane, preserves the low permeability of the outer membrane to ADP, and permits “efficient” energy transfers between mitochondrial and myofibrillar ATPase.

[0150] In some embodiments, treatment with an aromatic cationic peptide, such as, e.g., D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), increases the mK ATP activity in the myocardium in mammalian subjects that have suffered or are at risk of suffering heart failure.

[0151] In some embodiments, mK ATP activity is increased by about 0.25 fold to about 0.5 fold, or about 0.5 fold to about 0.75 fold, or about 0.75 fold to about 1.0 fold, or about 1.0 fold to about 1.5 fold, or about 1.5 fold to about 2.0 fold.

Cardiolipin

[0152] Cardiolipin (cardiolipin) is an important component of the inner mitochondrial membrane, where it constitutes about 20% of the total lipid composition. In mammalian cells, cardiolipin is found almost exclusively in the inner mitochondrial membrane where it is essential for the optimal function of enzymes involved in mitochondrial metabolism.

[0153] Cardiolipin is a species of diphosphatidylglycerol lipid comprising two phosphatidylglycerols connected with a glycerol backbone to form a dimeric structure. It has four alkyl groups and potentially carries two negative charges. As there are four distinct alkyl chains in cardiolipin, the molecule has the potential for great complexity. However, in most animal tissues, cardiolipin contains 18-carbon fatty alkyl chains with 2 unsaturated bonds on each of them. It has been proposed that the (18:2) in the four acyl chain configuration is an important structural requirement for the high affinity of cardiolipin to inner membrane proteins in mammalian mitochondria. However, studies with isolated enzyme preparations indicate that its importance may vary depending on the protein examined.

[0154] Each of the two phosphates in the molecule can capture one proton. Although it has a symmetric structure, ionization of one phosphate happens at different levels of acidity than ionizing both, with pK1=3 and pK2>7.5. Hence, under normal physiological conditions (a pH of approximately 7.0), the molecule may carry only one negative charge. Hydroxyl groups (—OH and —O—) on the phos-

phate form stable intramolecular hydrogen bonds, forming a bicyclic resonance structure. This structure traps one proton, which is conducive to oxidative phosphorylation.

[0155] During the oxidative phosphorylation process catalyzed by Complex IV, large quantities of protons are transferred from one side of the membrane to another side causing a large pH change. Without wishing to be bound by theory, it has been suggested that cardiolipin functions as a proton trap within the mitochondrial membranes, strictly localizing the proton pool and minimizing pH in the mitochondrial intermembrane space. This function is thought to be due to the unique structure of cardiolipin, which, as described above, can trap a proton within the bicyclic structure while carrying a negative charge. Thus, cardiolipin can serve as an electron buffer pool to release or absorb protons to maintain the pH near the mitochondrial membranes.

[0156] In addition, cardiolipin has been shown to play a role in apoptosis. An early event in the apoptosis cascade involves cardiolipin. As discussed in more detail below, a cardiolipin-specific oxygenase produces cardiolipin-hydroperoxides which causes the lipid to undergo a conformational change. The oxidized cardiolipin then translocates from the inner mitochondrial membrane to the outer mitochondrial membrane where it is thought to form a pore through which cytochrome c is released into the cytosol. Cytochrome c can bind to the IP3 receptor stimulating calcium release, which further promotes the release of cytochrome c. When the cytoplasmic calcium concentration reaches a toxic level, the cell dies. In addition, extra-mitochondrial cytochrome c interacts with apoptotic activating factors, causing the formation of apoptosomal complexes and activation of the proteolytic caspase cascade.

[0157] Other roles proposed for cardiolipin are: 1) participation in stabilization of the physical properties of the membrane (Schlame et al., 2000; Koshkin and Greenberg, 2002; Ma et al., 2004), for example, membrane fluidity and osmotic stability and 2) participation in protein function via direct interaction with membrane proteins (Schlame et al., 2000; Palsdottir and Hunte, 2004). Cardiolipin has been found in tight association with inner membrane protein complexes such as the cytochrome bc1 complex (complex III). As well, it has been localized to the contact sites of dimeric cytochrome c oxidase, and cardiolipin binding sites have also been found in the ADP/ATP carrier (AAC; for review see Palsdottir and Hunte, 2004). Recent work also suggests a role of cardiolipin in formation of respiratory chain super complexes (respirasomes).

[0158] The major tetra-acyl molecular species are 18:2 in each of the four fatty acyl positions of the cardiolipin molecule (referred to as the 18:2-18:2-18:2-18:2 cardiolipin species). Remodeling of cardiolipin is essential to obtain this enrichment of cardiolipin with linoleate because cardiolipin synthase has no molecular species substrate specificity for cytidine-5'-diphosphate-1,2-diacyl-sn-glycerol. In addition, the species pattern of cardiolipin precursors is similar enough to imply that the enzymes of the cardiolipin synthetic pathway are not molecular species-selective. Alterations in the molecular composition of cardiolipin are associated with various disease states.

[0159] Remodeling of cardiolipin occurs via at least three enzymes. Mitochondrial cardiolipin is remodeled by a deacylation-reacylation cycle in which newly synthesized cardiolipin is rapidly deacylated to monolysocardiolipin

(MLCL) and then reacylated back to cardiolipin. MLCL AT1 is responsible for the deacylation and ALCAT1 is responsible for the reacylation. In addition to these mitochondrial and microsomal acyltransferase activities, mitochondrial cardiolipin may be remodeled by a mitochondrial cardiolipin transacylase. Tafazzin (TAZ1) is a cardiolipin transacylase that specifically remodels mitochondrial cardiolipin with linoleic acid.

Taz1

[0160] Tafazzin (Taz1) is a protein that in humans is encoded by the TAZ gene. Taz1 functions as a phospholipid-lysophospholipid transacylase. Taz1 is highly expressed in cardiac and skeletal muscle and is involved in the metabolism of cardiolipin.

[0161] Taz1 is involved in the maintenance of the inner membrane of mitochondria. These proteins are involved in maintaining levels of cardiolipin, which is essential for energy production in the mitochondria.

[0162] Some mutations in the TAZ gene cause a condition called X-linked dilated cardiomyopathy. This is a condition in which the heart becomes so weakened and enlarged that it cannot pump blood efficiently, leading to heart failure. The decreased heart function can negatively affect many body systems and lead to swelling in the legs and abdomen, fluid in the lungs, and an increased risk of blood clots.

[0163] Another mutation in the TAZ gene causes a condition called isolated non-compaction of left ventricular myocardium (INVM). This condition occurs when the lower left chamber of the heart (left ventricle) does not develop correctly. The heart muscle is weakened and cannot pump blood efficiently, often leading to heart failure. Sometimes abnormal heart rhythms (arrhythmias) can also occur.

[0164] In some embodiments, treatment with an aromatic-cationic peptide, such as, e.g., D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), increases the expression of Taz1 in the myocardium in mammalian subjects that have suffered or are at risk of suffering heart failure.

[0165] In some embodiments, Taz1 expression level is increased by about 0.25 fold to about 0.5 fold, or about 0.5 fold to about 0.75 fold, or about 0.75 fold to about 1.0 fold, or about 1.0 fold to about 1.5 fold.

MLCL AT1

[0166] Monolysocardiolipin acyltransferase (MLCL AT1) catalyzes the acylation of MLCL to cardiolipin in mammalian tissues.

[0167] In some embodiments, treatment with an aromatic-cationic peptide, such as, e.g., D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), decreases the expression of MLCL AT1 in the myocardium in mammalian subjects that have suffered or are at risk of suffering heart failure.

[0168] In some embodiments, reducing MLCL AT1 expression level is a reduction measured by about 1 fold to about 1.5 fold reduction, or about 1.5 fold to about 2.0 fold reduction, or about 2.0 fold to about 2.5 fold reduction, or about 2.5 fold to about 3.0 fold reduction.

ALCAT1

[0169] Acyl-CoA lysocardiolipin acyltransferase 1 (ALCAT1) was initially identified as a microsomal lysocardio-

lipin acyltransferase. ALCAT1 possesses acyltransferase activities toward lysophosphatidylinositol (LPI) and lysophosphatidylglycerol (LPG).

[0170] ALCAT1 recognizes both monolysocardiolipin and dilyso-cardiolipin as substrates with a preference for linoleoyl-CoA and oleoyl-CoA as acyl donors. ALCAT1 acts as a remodeling enzyme for cardiolipin.

[0171] In some embodiments, treatment with an aromatic-cationic peptide, such as, e.g., D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), decreases the expression of ALCAT1 in the myocardium in mammalian subjects that have suffered or are at risk of suffering heart failure.

[0172] In some embodiments, reducing ALCAT1 expression level is a reduction measured by about 1 fold to about 1.5 fold reduction, or about 1.5 fold to about 2.0 fold reduction, or about 2.0 fold to about 2.5 fold reduction, or about 2.5 fold to about 3.0 fold reduction.

Aromatic-Cationic Peptides

[0173] The present technology relates to methods for decreasing the level of one or more of CRP, TNF-alpha, IL-6, ROS, cardiac troponin I, Nt-pro BNP, MLCL AT1, and ALCAT1 and/or increasing Taz1 expression and/or increasing mK ATP activity in a subject in need thereof, by administering aromatic-cationic peptides as disclosed herein. In some embodiments, reducing one or more of the level of CRP, TNF-alpha, IL-6, ROS, cardiac troponin I, Nt-pro BNP, MLCL AT1, and ALCAT1 and/or increasing Taz1 expression and/or increasing mK ATP activity is useful for the treatment or prevention of heart failure and related conditions, reducing risk factors associated with heart failure, and/or reducing the likelihood (risk) or severity of heart failure in the subject.

[0174] The present technology also relates to methods for preventing, ameliorating, or treating LV remodeling in a subject in need thereof, by administering aromatic-cationic peptides as disclosed herein. In some embodiments, the subject has an increased level of one or more of C-reactive protein, reactive oxygen species, interleukin-6, TNF-alpha, cardiac troponin I, Nt-pro BNP, MLCL AT1, and ALCAT1. In some embodiments, a decrease in Nt-pro BNP and/or cardiac troponin I is used as a biomarker to indicate the reduction or prevention of LV remodeling. In some embodiments, the subject has a decreased level of mK ATP activity or decreased expression of Taz1.

[0175] The present technology also relates to methods for increasing the activity of mitochondrial ATP sensitive potassium channels (mK ATP) or expression of Taz1 in a subject in need thereof, by administering aromatic-cationic peptides as disclosed herein. In some embodiments, increasing the activity of mK ATP or expression of Taz1 is useful for the treatment or prevention of heart failure and related conditions, reducing risk factors associated with heart failure, and/or reducing the likelihood (risk) or severity of heart failure in the subject.

[0176] The aromatic-cationic peptides are water-soluble and highly polar. Despite these properties, the peptides can readily penetrate cell membranes. The aromatic-cationic peptides typically include a minimum of three amino acids or a minimum of four amino acids, covalently joined by peptide bonds. The maximum number of amino acids present in the aromatic-cationic peptides is about twenty amino acids covalently joined by peptide bonds. Suitably, the

maximum number of amino acids is about twelve, more preferably about nine, and most preferably about six.

[0177] The amino acids of the aromatic-cationic peptides can be any amino acid. As used herein, the term "amino acid" is used to refer to any organic molecule that contains at least one amino group and at least one carboxyl group. Typically, at least one amino group is at the α position relative to a carboxyl group. The amino acids may be naturally occurring. Naturally occurring amino acids include, for example, the twenty most common levorotatory (L) amino acids normally found in mammalian proteins, i.e., alanine (Ala), arginine (Arg), asparagine (Asn), aspartic acid (Asp), cysteine (Cys), glutamine (Gln), glutamic acid (Glu), glycine (Gly), histidine (His), isoleucine (Ile), leucine (Leu), lysine (Lys), methionine (Met), phenylalanine (Phe), proline (Pro), serine (Ser), threonine (Thr), tryptophan, (Trp), tyrosine (Tyr), and valine (Val). Other naturally occurring amino acids include, for example, amino acids that are synthesized in metabolic processes not associated with protein synthesis. For example, the amino acids ornithine and citrulline are synthesized in mammalian metabolism during the production of urea. Another example of a naturally occurring amino acid includes hydroxyproline (Hyp).

[0178] The peptides optionally contain one or more non-naturally occurring amino acids. Optimally, the peptide has no amino acids that are naturally occurring. The non-naturally occurring amino acids may be levorotary (L-), dextrorotatory (D-), or mixtures thereof. Non-naturally occurring amino acids are those amino acids that typically are not synthesized in normal metabolic processes in living organisms, and do not naturally occur in proteins. In addition, the non-naturally occurring amino acids suitably are also not recognized by common proteases. The non-naturally occurring amino acid can be present at any position in the peptide. For example, the non-naturally occurring amino acid can be at the N-terminus, the C-terminus, or at any position between the N-terminus and the C-terminus.

[0179] The non-natural amino acids may, for example, comprise alkyl, aryl, or alkylaryl groups not found in natural amino acids. Some examples of non-natural alkyl amino acids include α -aminobutyric acid, β -aminobutyric acid, γ -aminobutyric acid, δ -aminovaleric acid, and ϵ -aminocaproic acid. Some examples of non-natural aryl amino acids include ortho-, meta-, and para-aminobenzoic acid. Some examples of non-natural alkylaryl amino acids include ortho-, meta-, and para-aminophenylacetic acid, and γ -phenyl- β -aminobutyric acid. Non-naturally occurring amino acids include derivatives of naturally occurring amino acids.

The derivatives of naturally occurring amino acids may, for example, include the addition of one or more chemical groups to the naturally occurring amino acid.

[0180] For example, one or more chemical groups can be added to one or more of the 2', 3', 4', 5', or 6' position of the aromatic ring of a phenylalanine or tyrosine residue, or the 4', 5', 6', or 7' position of the benzo ring of a tryptophan residue. The group can be any chemical group that can be added to an aromatic ring. Some examples of such groups include branched or unbranched C₁-C₄ alkyl, such as methyl, ethyl, n-propyl, isopropyl, butyl, isobutyl, or t-butyl,

C₁-C₄ alkyloxy (i.e., alkoxy), amino, C₁-C₄ alkylamino and C₁-C₄ dialkylamino (e.g., methylamino, dimethylamino), nitro, hydroxyl, halo (i.e., fluoro, chloro, bromo, or iodo). Some specific examples of non-naturally occurring derivatives of naturally occurring amino acids include norvaline (Nva) and norleucine (Nle).

[0181] Another example of a modification of an amino acid in a peptide is the derivatization of a carboxyl group of an aspartic acid or a glutamic acid residue of the peptide. One example of derivatization is amidation with ammonia or with a primary or secondary amine, e.g. methylamine, ethylamine, dimethylamine or diethylamine. Another example of derivatization includes esterification with, for example, methyl or ethyl alcohol. Another such modification includes derivatization of an amino group of a lysine, arginine, or histidine residue. For example, such amino groups can be acylated. Some suitable acyl groups include, for example, a benzoyl group or an alkanoyl group comprising any of the C₁-C₄ alkyl groups mentioned above, such as an acetyl or propionyl group.

[0182] The non-naturally occurring amino acids are suitably resistant or insensitive to common proteases. Examples of non-naturally occurring amino acids that are resistant or insensitive to proteases include the dextrorotatory (D-) form of any of the above-mentioned naturally occurring L-amino acids, as well as L- and/or D-non-naturally occurring amino acids. The D-amino acids do not normally occur in proteins, although they are found in certain peptide antibiotics that are synthesized by means other than the normal ribosomal protein synthetic machinery of the cell. As used herein, the D-amino acids are considered to be non-naturally occurring amino acids.

[0183] In order to minimize protease sensitivity, the peptides should have less than five, preferably less than four, more preferably less than three, and most preferably, less than two contiguous L-amino acids recognized by common proteases, irrespective of whether the amino acids are naturally or non-naturally occurring. Optimally, the peptide has only D-amino acids, and no L-amino acids. If the peptide contains protease sensitive sequences of amino acids, at least one of the amino acids is preferably a non-naturally occurring D-amino acid, thereby conferring protease resistance. An example of a protease sensitive sequence includes two or more contiguous basic amino acids that are readily cleaved by common proteases, such as endopeptidases and trypsin. Examples of basic amino acids include arginine, lysine and histidine.

[0184] The aromatic-cationic peptides should have a minimum number of net positive charges at physiological pH in comparison to the total number of amino acid residues in the peptide. The minimum number of net positive charges at physiological pH will be referred to below as (p_m). The total number of amino acid residues in the peptide will be referred to below as (r). The minimum numbers of net positive charges discussed below are all at physiological pH. The term "physiological pH" as used herein refers to the normal pH in the cells of the tissues and organs of the mammalian body. For instance, the physiological pH of a human is

normally approximately 7.4, but normal physiological pH in mammals may be any pH from about 7.0 to about 7.8.

[0185] Typically, a peptide has a positively charged N-terminal amino group and a negatively charged C-terminal carboxyl group. The charges cancel each other out at physiological pH. As an example of calculating net charge, the peptide Tyr-Arg-Phe-Lys-Glu-His-Trp-D-Arg (SEQ ID NO: 8) has one negatively charged amino acid (i.e., Glu) and four positively charged amino acids (i.e., two Arg residues, one Lys, and one His). Therefore, the above peptide has a net positive charge of three.

[0186] In one embodiment, the aromatic-cationic peptides have a relationship between the minimum number of net positive charges at physiological pH (p_m) and the total number of amino acid residues (r) wherein $3p_m$ is the largest number that is less than or equal to $r+1$. In this embodiment, the relationship between the minimum number of net positive charges (p_m) and the total number of amino acid residues (r) is as follows:

TABLE 2

		Amino acid number and net positive charges ($3p_m \leq p + 1$)																	
(r)		3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
(p_m)		1	1	2	2	2	3	3	3	4	4	4	5	5	5	6	6	6	7

[0187] In another embodiment, the aromatic-cationic peptides have a relationship between the minimum number of net positive charges (p_m) and the total number of amino acid residues (r) wherein $2p_m$ is the largest number that is less than or equal to $r+1$. In this embodiment, the relationship between the minimum number of net positive charges (p_m) and the total number of amino acid residues (r) is as follows:

TABLE 3

		Amino acid number and net positive charges ($2p_m \leq p + 1$)																	
(r)		3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
(p_m)		2	2	3	3	4	4	5	5	6	6	7	7	8	8	9	9	10	10

[0188] In one embodiment, the minimum number of net positive charges (p_m) and the total number of amino acid residues (r) are equal. In another embodiment, the peptides have three or four amino acid residues and a minimum of one net positive charge, suitably, a minimum of two net positive charges and more preferably a minimum of three net positive charges.

[0189] It is also important that the aromatic-cationic peptides have a minimum number of aromatic groups in comparison to the total number of net positive charges (p_t). The minimum number of aromatic groups will be referred to below as (a). Naturally occurring amino acids that have an aromatic group include the amino acids histidine, tryptophan, tyrosine, and phenylalanine. For example, the hexapeptide Lys-Gln-Tyr-D-Arg-Phe-Trp (SEQ ID NO: 9) has a net positive charge of two (contributed by the lysine and arginine residues) and three aromatic groups (contributed by tyrosine, phenylalanine and tryptophan residues).

[0190] The aromatic-cationic peptides should also have a relationship between the minimum number of aromatic groups (a) and the total number of net positive charges at physiological pH (p_t) wherein 3a is the largest number that is less than or equal to p_t+1 , except that when p_t is 1, a may also be 1. In this embodiment, the relationship between the minimum number of aromatic groups (a) and the total number of net positive charges (p_t) is as follows:

TABLE 4

		Aromatic groups and net positive charges ($3a \leq p_t + 1$ or $a = p_t = 1$)																			
(p_t)		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
(a)		1	1	1	1	2	2	2	3	3	3	4	4	4	5	5	5	6	6	6	7

[0191] In another embodiment, the aromatic-cationic peptides have a relationship between the minimum number of aromatic groups (a) and the total number of net positive charges (p_t) wherein 2a is the largest number that is less than or equal to p_t+1 . In this embodiment, the relationship between the minimum number of aromatic amino acid residues (a) and the total number of net positive charges (p_t) is as follows:

TABLE 5

		Aromatic groups and net positive charges ($2a \leq p_t + 1$ or $a = p_t = 1$)																			
(p_t)		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
(a)		1	1	2	2	3	3	4	4	5	5	6	6	7	7	8	8	9	9	10	10

[0192] In another embodiment, the number of aromatic groups (a) and the total number of net positive charges (p_+) are equal.

[0193] Carboxyl groups, especially the terminal carboxyl group of a C-terminal amino acid, are suitably amidated with, for example, ammonia to form the C-terminal amide. Alternatively, the terminal carboxyl group of the C-terminal amino acid may be amidated with any primary or secondary amine. The primary or secondary amine may, for example, be an alkyl, especially a branched or unbranched C₁-C₄ alkyl, or an aryl amine. Accordingly, the amino acid at the C-terminus of the peptide may be converted to an amido, N-methylamido, N-ethylamido, N,N-dimethylamido, N,N-diethylamido, N-methyl-N-ethylamido, N-phenylamido or

N-phenyl-N-ethylamido group. The free carboxylate groups of the asparagine, glutamine, aspartic acid, and glutamic acid residues not occurring at the C-terminus of the aromatic-cationic peptides may also be amidated wherever they occur within the peptide. The amidation at these internal positions may be with ammonia or any of the primary or secondary amines described above.

[0194] In one embodiment, the aromatic-cationic peptide is a tripeptide having two net positive charges and at least one aromatic amino acid. In a particular embodiment, the aromatic-cationic peptide is a tripeptide having two net positive charges and two aromatic amino acids.

[0195] Aromatic-cationic peptides include, but are not limited to, the following peptide examples:

TABLE 6

EXEMPLARY PEPTIDES
2', 6'-Dmp-D-Arg-2', 6'-Dmt-Lys-NH ₂ (SEQ ID NO: 10)
2', 6'-Dmp-D-Arg-Phe-Lys-NH ₂ (SEQ ID NO: 7)
2', 6'-Dmt-D-Arg-PheOrn-NH ₂
2', 6'-Dmt-D-Arg-Phe-Ahp (2-aminoheptanoicacid)-NH ₂
2', 6'-Dmt-D-Arg-Phe-Lys-NH ₂ (SEQ ID NO: 5)
2', 6'-Dmt-D-Cit-PheLys-NH ₂
Ala-D-Phe-D-Arg-Tyr-Lys-D-Trp-His-D-Tyr-Gly-Phe (SEQ ID NO: 11)
Arg-D-Leu-D-Tyr-Phe-Lys-Glu-D-Lys-Arg-D-Trp-Lys-D-Phe-Tyr-D-Arg-Gly (SEQ ID NO: 12)
Asp-Arg-D-Phe-Cys-Phe-D-Arg-D-Lys-Tyr-Arg-D-Tyr-Trp-D-His-Tyr-D-Phe-Lys-Phe (SEQ ID NO: 13)
Asp-D-Trp-Lys-Tyr-D-His-Phe-Arg-D-Gly-Lys-NH ₂ (SEQ ID NO: 14)
D-Arg-2', 6'-Dmt-Lys-Phe-NH ₂ (SEQ ID NO: 3)
D-Glu-Asp-Lys-D-Arg-D-His-Phe-Phe-D-Val-Tyr-Arg-Tyr-D-Tyr-Arg-His-Phe-NH ₂ (SEQ ID NO: 15)
D-His-Glu-Lys-Tyr-D-Phe-Arg (SEQ ID NO: 16)
D-His-Lys-Tyr-D-Phe-Glu-D-Asp-D-Asp-D-His-D-Lys-Arg-Trp-NH ₂ (SEQ ID NO: 17)
D-Tyr-Trp-Lys-NH ₂
Glu-Arg-D-Lys-Tyr-D-Val-Phe-D-His-Trp-Arg-D-Gly-Tyr-Arg-D-Met-NH ₂ (SEQ ID NO: 18)
Gly-Ala-Lys-Phe-D-Lys-Glu-Arg-Tyr-His-D-Arg-D-Arg-Asp-Tyr-Trp-D-His-Trp-His-D-Lys-Asp. (SEQ ID NO: 19)
Gly-D-Phe-Lys-His-D-Arg-Tyr-NH ₂ (SEQ ID NO: 20)
His-Tyr-D-Arg-Trp-Lys-Phe-D-Asp-Ala-Arg-Cys-D-Tyr-His-Phe-D-Lys-Tyr-His-Ser-NH ₂ (SEQ ID NO: 21)
Lys-D-Arg-Tyr-NH ₂
Lys-D-Gln-Tyr-Arg-D-Phe-Trp-NH ₂ (SEQ ID NO: 22)
Lys-Trp-D-Tyr-Arg-Asn-Phe-Tyr-D-His-NH ₂ (SEQ ID NO: 23)
Met-Tyr-D-Arg-Phe-Arg-NH ₂ (SEQ ID NO: 24)
Met-Tyr-D-Lys-Phe-Arg (SEQ ID NO: 25)
Phe-Arg-D-His-Asp (SEQ ID NO: 26)
Phe-D-Arg-2', 6'-Dmt-Lys-NH ₂ (SEQ ID NO: 27)

TABLE 6-continued

EXEMPLARY PEPTIDES

Phe-D-Arg-His

Phe-D-Arg-Lys-Trp-Tyr-D-Arg-His (SEQ ID NO: 28)

Phe-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 6)

Phe-Phe-D-Tyr-Arg-Glu-Asp-D-Lys-Arg-D-Arg-His-Phe-NH₂ (SEQ ID NO: 29)

Phe-Tyr-Lys-D-Arg-Trp-His-D-Lys-Glu-Arg-D-Tyr-Thr (SEQ ID NO: 30)

Thr-Gly-Tyr-Arg-D-His-Phe-Trp-D-His-Lys (SEQ ID NO: 31)

Thr-Tyr-Arg-D-Lys-Trp-Tyr-Glu-Asp-D-Lys-D-Arg-His-Phe-D-Tyr-Gly-Val-Ile-D-His-Arg-Tyr-Lys-NH₂ (SEQ ID NO: 32)

Trp-D-Lys-Tyr-Arg-NH₂ (SEQ ID NO: 33)

Trp-Lys-Phe-D-Asp-Arg-Tyr-D-His-Lys (SEQ ID NO: 34)

Tyr-Asp-D-Lys-Tyr-Phe-D-Lys-D-Arg-Phe-Pro-D-Tyr-His-Lys (SEQ ID NO: 35)

Tyr-D-Arg-Phe-Lys-Glu-NH₂ (SEQ ID NO: 36)

Tyr-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 4)

Tyr-D-His-Phe-D-Arg-Asp-Lys-D-Arg-His-Trp-D-His-Phe (SEQ ID NO: 37)

Tyr-His-D-Gly-Met (SEQ ID NO: 38)

Val-D-Lys-His-Tyr-D-Phe-Ser-Tyr-Arg-NH₂ (SEQ ID NO: 39)

[0196] In one embodiment, the peptides have mu-opioid receptor agonist activity (i.e., they activate the mu-opioid receptor). Peptides, which have mu-opioid receptor agonist activity, are typically those peptides that have a tyrosine residue or a tyrosine derivative at the N-terminus (i.e., the first amino acid position). Suitable derivatives of tyrosine include 2'-methyltyrosine (Mmt); 2',6'-dimethyltyrosine (2'6'-Dmt); 3',5'-dimethyltyrosine (3'5'Dmt); N,2',6'-trimethyltyrosine (Tmt); and 2'-hydroxy-6'-methyltyrosine (Hmt). **[0197]** In one embodiment, a peptide that has mu-opioid receptor agonist activity has the formula Tyr-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 4). Tyr-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 4) has a net positive charge of three, contributed by the amino acids tyrosine, arginine, and lysine and has two aromatic groups contributed by the amino acids phenylalanine and tyrosine. The tyrosine of Tyr-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 4) can be a modified derivative of tyrosine such as in 2',6'-dimethyltyrosine to produce the compound having the formula 2',6'-Dmt-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 5). 2',6'-Dmt-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 5) has a molecular weight of 640 and carries a net three positive charge at physiological pH. 2',6'-Dmt-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 5) readily penetrates the plasma membrane of several mammalian cell types in an energy-independent manner (Zhao et al., *J. Pharmacol Exp Ther.*, 304:425-432, 2003).

[0198] Alternatively, in other instances, the aromatic-cationic peptide does not have mu-opioid receptor agonist activity. For example, during long-term treatment, such as in a chronic disease state or condition, the use of an aromatic-cationic peptide that activates the mu-opioid receptor may be contraindicated. In these instances, the potentially adverse or addictive effects of the aromatic-cationic peptide may preclude the use of an aromatic-cationic peptide that

activates the mu-opioid receptor in the treatment regimen of a human patient or other mammal. Potential adverse effects may include sedation, constipation and respiratory depression. In such instances an aromatic-cationic peptide that does not activate the mu-opioid receptor may be an appropriate treatment. Peptides that do not have mu-opioid receptor agonist activity generally do not have a tyrosine residue or a derivative of tyrosine at the N-terminus (i.e., amino acid position 1). The amino acid at the N-terminus can be any naturally occurring or non-naturally occurring amino acid other than tyrosine. In one embodiment, the amino acid at the N-terminus is phenylalanine or its derivative. Exemplary derivatives of phenylalanine include 2'-methylphenylalanine (Mmp), 2',6'-dimethylphenylalanine (2'6'-Dmp), N,2',6'-trimethylphenylalanine (Tmp), and 2'-hydroxy-6'-methylphenylalanine (Hmp).

[0199] An example of an aromatic-cationic peptide that does not have mu-opioid receptor agonist activity has the formula Phe-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 6). Alternatively, the N-terminal phenylalanine can be a derivative of phenylalanine such as 2',6'-dimethylphenylalanine (2'6'-Dmp). Tyr-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 4) containing 2',6'-dimethylphenylalanine at amino acid position 1 has the formula 2',6'-Dmp-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 7). In one embodiment, the amino acid sequence of 2',6'-Dmt-D-Arg-Phe-Lys-NH₂ (SEQ ID NO: 5) is rearranged such that Dmt is not at the N-terminus. An example of such an aromatic-cationic peptide that does not have mu-opioid receptor agonist activity has the formula D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3).

[0200] Suitable substitution variants of the peptides listed herein include conservative amino acid substitutions. Amino acids may be grouped according to their physicochemical characteristics as follows:

- [0201] (a) Non-polar amino acids: Ala(A) Ser(S) Thr(T) Pro(P) Gly(G) Cys (C);
- [0202] (b) Acidic amino acids: Asn(N) Asp(D) Glu(E) Gln(Q);
- [0203] (c) Basic amino acids: His(H) Arg(R) Lys(K);
- [0204] (d) Hydrophobic amino acids: Met(M) Leu(L) Ile(I) Val(V); and
- [0205] (e) Aromatic amino acids: Phe(F) Tyr(Y) Trp(W) His (H).

[0206] Substitutions of an amino acid in a peptide by another amino acid in the same group is referred to as a conservative substitution and may preserve the physico-chemical characteristics of the original peptide. In contrast, substitutions of an amino acid in a peptide by another amino acid in a different group are generally more likely to alter the characteristics of the original peptide.

[0207] Examples of peptides that activate mu-opioid receptors include, but are not limited to, the aromatic-cationic peptides shown in Table 7.

TABLE 7

Peptide Analogs with Mu-Opioid Activity						
Amino Acid Position 1	Amino Acid Position 2	Amino Acid Position 3	Amino Acid Position 4	C-Terminal Modification	SEQ ID NO	
Tyr	D-Arg	Phe	Lys	NH ₂	4	
Tyr	D-Arg	Phe	Orn	NH ₂		
Tyr	D-Arg	Phe	Dab	NH ₂		
Tyr	D-Arg	Phe	Dap	NH ₂		
2',6'Dmt	D-Arg	Phe	Lys	NH ₂	5	
2',6'Dmt	D-Arg	Phe	Lys-NH(CH ₂) ₂ -NH ₂	NH-dns	40	
2',6'Dmt	D-Arg	Phe	Lys-NH(CH ₂) ₂ -NH ₂	NH-attn	41	
2',6'Dmt	D-Arg	Phe	dnsLys	NH ₂	42	
2',6'Dmt	D-Cit	Phe	Lys	NH ₂		
2',6'Dmt	D-Cit	Phe	Ahp	NH ₂		
2',6'Dmt	D-Arg	Phe	Orn	NH ₂		
2',6'Dmt	D-Arg	Phe	Dab	NH ₂		
2',6'Dmt	D-Arg	Phe	Dap	NH ₂		
2',6'Dmt	D-Arg	Phe	Ahp(2-aminoheptanoic acid)	NH ₂		
Bio-	D-Arg	Phe	Lys	NH ₂	43	
2',6'Dmt	D-Arg	Phe	Lys	NH ₂	44	
3',5'Dmt	D-Arg	Phe	Lys	NH ₂	45	
3',5'Dmt	D-Arg	Phe	Orn	NH ₂		
3',5'Dmt	D-Arg	Phe	Dab	NH ₂		
3',5'Dmt	D-Arg	Phe	Dap	NH ₂		
Tyr	D-Arg	Tyr	Lys	NH ₂		
Tyr	D-Arg	Tyr	Orn	NH ₂		
Tyr	D-Arg	Tyr	Dab	NH ₂		
Tyr	D-Arg	Tyr	Dap	NH ₂		
2',6'Dmt	D-Arg	Tyr	Dap	NH ₂		
2',6'Dmt	D-Arg	2',6'Dmt	Lys	NH ₂	46	
2',6'Dmt	D-Arg	2',6'Dmt	Orn	NH ₂		
2',6'Dmt	D-Arg	2',6'Dmt	Dab	NH ₂		
2',6'Dmt	D-Arg	2',6'Dmt	Dap	NH ₂		
3',5'Dmt	D-Arg	3',5'Dmt	Arg	NH ₂	48	
3',5'Dmt	D-Arg	3',5'Dmt	Lys	NH ₂	49	
3',5'Dmt	D-Arg	3',5'Dmt	Orn	NH ₂		
3',5'Dmt	D-Arg	3',5'Dmt	Dab	NH ₂		
Tyr	D-Lys	Phe	Dap	NH ₂	50	
Ty	D-Lys	Phe	Arg	NH ₂	51	
Tyr	D-Lys	Phe	Lys	NH ₂	52	
Tyr	D-Lys	Phe	Orn	NH ₂		
2',6'Dmt	D-Lys	Phe	Dab	NH ₂		
2',6'Dmt	D-Lys	Phe	Dap	NH ₂		

TABLE 7-continued

Peptide Analogs with Mu-Opioid Activity						
Amino Acid Position 1	Amino Acid Position 2	Amino Acid Position 3	Amino Acid Position 4	C-Terminal Modification	SEQ ID NO	
2',6'Dmt	D-Lys	Phe	Arg	NH ₂	52	
2',6'Dmt	D-Lys	Phe	Lys	NH ₂	53	
3',5'Dmt	D-Lys	Phe	Orn	NH ₂		
3',5'Dmt	D-Lys	Phe	Dab	NH ₂		
3',5'Dmt	D-Lys	Phe	Dap	NH ₂		
3',5'Dmt	D-Lys	Phe	Arg	NH ₂	54	
Tyr	D-Lys	Tyr	Lys	NH ₂	55	
Tyr	D-Lys	Tyr	Orn	NH ₂		
Tyr	D-Lys	Tyr	Dab	NH ₂		
Tyr	D-Lys	Tyr	Dap	NH ₂		
2',6'Dmt	D-Lys	Tyr	Lys	NH ₂	56	
2',6'Dmt	D-Lys	Tyr	Orn	NH ₂		
2',6'Dmt	D-Lys	Tyr	Dab	NH ₂		
2',6'Dmt	D-Lys	Tyr	Dap	NH ₂		
2',6'Dmt	D-Lys	2',6'Dmt	Lys	NH ₂	57	
2',6'Dmt	D-Lys	2',6'Dmt	Orn	NH ₂		
2',6'Dmt	D-Lys	2',6'Dmt	Dab	NH ₂		
2',6'Dmt	D-Lys	2',6'Dmt	Dap	NH ₂		
2',6'Dmt	D-Arg	Phe	dnsDap	NH ₂		
2',6'Dmt	D-Arg	Phe	atnDap	NH ₂		
3',5'Dmt	D-Lys	3',5'Dmt	Lys	NH ₂	58	
3',5'Dmt	D-Lys	3',5'Dmt	Orn	NH ₂		
3',5'Dmt	D-Lys	3',5'Dmt	Dab	NH ₂		
3',5'Dmt	D-Lys	3',5'Dmt	Dap	NH ₂		
Tyr	D-Lys	Phe	Arg	NH ₂	59	
Tyr	D-Orn	Phe	Arg	NH ₂	52	
Tyr	D-Dab	Phe	Arg	NH ₂		
Tyr	D-Dap	Phe	Arg	NH ₂		
2',6'Dmt	D-Arg	Phe	Arg	NH ₂	60	
3',5'Dmt	D-Lys	Phe	Arg	NH ₂	54	
3',5'Dmt	D-Orn	Phe	Arg	NH ₂		
Tyr	D-Lys	Tyr	Arg	NH ₂	61	
Tyr	D-Orn	Tyr	Arg	NH ₂		
Tyr	D-Dab	Tyr	Arg	NH ₂		
Tyr	D-Dap	Tyr	Arg	NH ₂		
2',6'Dmt	D-Arg	2',6'Dmt	Arg	NH ₂	62	
2',6'Dmt	D-Orn	2',6'Dmt	Arg	NH ₂	63	
2',6'Dmt	D-Dab	2',6'Dmt	Arg	NH ₂		
2',6'Dmt	D-Dap	2',6'Dmt	Arg	NH ₂		
2',6'Dmt	D-Arg	2',6'Dmt	Arg	NH ₂		
3',5'Dmt	D-Arg	3',5'Dmt	Arg	NH ₂	64	
3',5'Dmt	D-Lys	3',5'Dmt	Arg	NH ₂		
3',5'Dmt	D-Orn	3',5'Dmt	Arg	NH ₂	65	
Mmt	D-Arg	Phe	Lys	NH ₂		
Mmt	D-Arg	Phe	Orn	NH ₂		
Mmt	D-Arg	Phe	Dab	NH ₂		
Mmt	D-Arg	Phe	Dap	NH ₂		
Tmt	D-Arg	Phe	Lys	NH ₂	66	
Tmt	D-Arg	Phe	Orn	NH ₂		
Tmt	D-Arg	Phe	Dab	NH ₂		
Tmt	D-Arg	Phe	Dap	NH ₂		
Hmt	D-Arg	Phe	Lys	NH ₂	67	
Hmt	D-Arg	Phe	Orn	NH ₂		
Hmt	D-Arg	Phe	Dab	NH ₂		
Hmt	D-Arg	Phe	Dap	NH ₂		
Mmt	D-Lys	Phe	Lys	NH ₂	68	
Mmt	D-Lys	Phe	Orn	NH ₂		
Mmt	D-Lys	Phe	Dab	NH ₂		
Mmt	D-Lys	Phe	Dap	NH ₂		
Tmt	D-Lys	Phe	Lys	NH ₂	69	
Tmt	D-Lys	Phe	Orn	NH ₂		
Tmt	D-Lys	Phe	Dab	NH ₂		
Tmt	D-Lys	Phe	Dap	NH ₂		
Tyr	D-Lys	Phe	Dap	NH ₂	70	
Ty	D-Lys	Phe	Arg	NH ₂		
Tyr	D-Lys	Phe	Lys	NH ₂	51	
Tyr	D-Lys	Phe	Orn	NH ₂		
Tyr	D-Lys	Phe	Dab	NH ₂		
Tyr	D-Lys	Phe	Dap	NH ₂		
2',6'Dmt	D-Lys	Phe	Dap	NH ₂	71	
2',6'Dmt	D-Lys	Phe	Dap	NH ₂	72	

TABLE 7-continued

Peptide Analogs with Mu-Opioid Activity					
Amino Acid Position 1	Amino Acid Position 2	Amino Acid Position 3	Amino Acid Position 4	C-Terminal Modification	SEQ ID NO
Hmt	D-Lys	Phe	Orn	NH ₂	
Hmt	D-Lys	Phe	Dab	NH ₂	
Hmt	D-Lys	Phe	Dap	NH ₂	
Hmt	D-Lys	Phe	Arg	NH ₂	73
Mmt	D-Lys	Phe	Arg	NH ₂	69
Mmt	D-Orn	Phe	Arg	NH ₂	
Mmt	D-Dab	Phe	Arg	NH ₂	
Mmt	D-Dap	Phe	Arg	NH ₂	
Mmt	D-Arg	Phe	Arg	NH ₂	74
Tmt	D-Lys	Phe	Arg	NH ₂	71
Tmt	D-Orn	Phe	Arg	NH ₂	
Tmt	D-Dab	Phe	Arg	NH ₂	
Tmt	D-Dap	Phe	Arg	NH ₂	
Tmt	D-Arg	Phe	Arg	NH ₂	75
Hmt	D-Lys	Phe	Arg	NH ₂	73
Hmt	D-Orn	Phe	Arg	NH ₂	
Hmt	D-Dab	Phe	Arg	NH ₂	
Hmt	D-Dap	Phe	Arg	NH ₂	
Hmt	D-Arg	Phe	Arg	NH ₂	76

Dab = diaminobutyric acid

Dap = diaminopropionic acid

Dmt = dimethyltyrosine

Mmt = 2'-methyltyrosine

Tmt = N,2',6'-trimethyltyrosine

Hmt = 2'-hydroxy,6'-methyltyrosine

dnsDap = β-dansyl-L-α,β-diaminopropionic acid

atnDap = β-anthraniloyl-L-α,β-diaminopropionic acid

Bio = biotin

[0208] Examples of peptides that do not activate mu-opioid receptors include, but are not limited to, the aromatic-cationic peptides shown in Table 8.

TABLE 8

Peptide Analogs Lacking Mu-Opioid Activity					
Amino Acid Position 1	Amino Acid Position 2	Amino Acid Position 3	Amino Acid Position 4	C-Terminal Modification	SEQ ID NO
D-Arg	Dmt	Lys	Phe	NH ₂	77
D-Arg	Dmt	Phe	Lys	NH ₂	78
D-Arg	Phe	Lys	Dmt	NH ₂	79
D-Arg	Phe	Dmt	Lys	NH ₂	80
D-Arg	Lys	Dmt	Phe	NH ₂	81
D-Arg	Lys	Phe	Dmt	NH ₂	82
Phe	Lys	Dmt	D-Arg	NH ₂	83
Phe	Lys	D-Arg	Dmt	NH ₂	84
Phe	D-Arg	Phe	Lys	NH ₂	6
Phe	D-Arg	Dmt	Lys	NH ₂	85
Phe	D-Arg	Lys	Dmt	NH ₂	86
Phe	Dmt	D-Arg	Lys	NH ₂	87
Phe	Dmt	Lys	D-Arg	NH ₂	88
Lys	Phe	D-Arg	Dmt	NH ₂	89
Lys	Phe	Dmt	D-Arg	NH ₂	90
Lys	Dmt	D-Arg	Phe	NH ₂	91
Lys	Dmt	Phe	D-Arg	NH ₂	92
Lys	D-Arg	Phe	Dmt	NH ₂	93
Lys	D-Arg	Dmt	Phe	NH ₂	94
D-Arg	Dmt	D-Arg	Phe	NH ₂	95
D-Arg	Dmt	D-Arg	Dmt	NH ₂	96
D-Arg	Dmt	D-Arg	Tyr	NH ₂	97
D-Arg	Dmt	D-Arg	Trp	NH ₂	98
Trp	D-Arg	Phe	Lys	NH ₂	99
Trp	D-Arg	Tyr	Lys	NH ₂	100
Trp	D-Arg	Trp	Lys	NH ₂	101
Trp	D-Arg	Dmt	Lys	NH ₂	102
D-Arg	Trp	Lys	Phe	NH ₂	103

TABLE 8-continued

Peptide Analogs Lacking Mu-Opioid Activity					
Amino Acid Position 1	Amino Acid Position 2	Amino Acid Position 3	Amino Acid Position 4	C-Terminal Modification	SEQ ID NO
D-Arg	Trp	Phe	Lys	NH ₂	104
D-Arg	Trp	Lys	Dmt	NH ₂	105
D-Arg	Trp	Dmt	Lys	NH ₂	106
D-Arg	Lys	Trp	Phe	NH ₂	107
D-Arg	Lys	Trp	Dmt	NH ₂	108
Cha	D-Arg	Phe	Lys	NH ₂	
Ala	D-Arg	Phe	Lys	NH ₂	109

Cha = cyclohexyl alanine

[0209] The amino acids of the peptides shown in Table 7 and 8 may be in either the L- or the D-configuration.

[0210] The peptides may be synthesized by any of the methods well known in the art. Suitable methods for chemically synthesizing the protein include, for example, those described by Stuart and Young in *Solid Phase Peptide Synthesis*, Second Edition, Pierce Chemical Company (1984), and in *Methods Enzymol.*, 289, Academic Press, Inc., New York (1997).

Prophylactic and Therapeutic Uses of Aromatic-Cationic Peptides

[0211] General. The aromatic-cationic peptides described herein are useful to prevent or treat disease. Specifically, the disclosure provides for both prophylactic and therapeutic methods of treating a subject having or suspected of having one or more of an elevated CRP level, e.g., as determined by hsCRP assay, an elevated TNF-alpha level, an elevated IL-6 level, or an elevated ROS level, an elevated cardiac troponin I level, an elevated Nt-pro BNP level, an elevated MLCL AT1 expression level, an elevated ALCAT 1 expression level, and/or a decrease in mK ATP activity and/or decrease in Taz1 expression. Accordingly, in some embodiments, the disclosure provides for both prophylactic and therapeutic methods of treating a subject having or at risk of (susceptible to) heart failure, and having or suspected of having an elevated CRP level, e.g., as determined by hsCRP, an elevated TNF-alpha level, an elevated IL-6 level, an elevated ROS level, an elevated cardiac troponin I level, an elevated Nt-pro BNP level, an elevated MLCL AT1 expression level, an elevated ALCAT 1 expression level, and/or a decrease in mK ATP activity and/or decrease in Taz1 expression. Accordingly, the present methods provide for the prevention and/or treatment of heart failure in a subject by administering an effective amount of an aromatic-cationic peptide to a subject in need thereof to reduce or normalize one or more of the CRP level, the TNF-alpha level, the IL-6 level, the ROS level, the cardiac troponin I level, MLCL AT1 expression level, the ALCAT 1 expression level, and/or Nt-pro BNP, and/or to increase or normalize the Taz1 expression level and/or the mK ATP activity of the subject.

[0212] Additionally, the disclosure provides for both prophylactic and therapeutic methods of treating a subject at risk of LV remodeling, or suffering from LV remodeling, e.g., due to elevated CRP, TNF-alpha, IL-6, ROS, cardiac troponin I, MLCL AT1, or ALCAT 1 and/or a decrease in mK ATP activity and/or decrease in Taz1 expression. Accordingly, the present methods provide for the prevention and/or

treatment of LV remodeling in a subject by administering an effective amount of an aromatic-cationic peptide to a subject in need thereof.

[0213] In some embodiments, elevation of one or more of CRP level, TNF-alpha level, IL-6 level, or ROS level, or Nt-pro BNP level, or cardiac troponin I level, or MLCL AT1 expression, or ALCAT 1 expression serve to indicate future vascular events in subjects without any previous history of cardiovascular disease. Additionally, or alternatively, in some embodiments, the decreased activity of mK ATP or decreased Taz1 expression serves to indicate future vascular events in subjects without any previous history of cardiovascular disease. In some embodiments, the determination of one or more of CRP level, TNF-alpha level, IL-6 level, or ROS, or Nt-pro BNP level, or cardiac troponin I level, or MLCL AT1 expression, or ALCAT 1 expression, mK ATP activity, or Taz1 expression enhances risk assessment and therapeutic outcomes in primary cardiovascular disease prevention. In some embodiments, one or more of CRP level, TNF-alpha level, IL-6 level, or ROS level, or Nt-pro BNP level, or cardiac troponin I level, or MLCL AT1 expression, or ALCAT 1 expression, or Taz1 expression, or mK ATP activity serve as an independent marker of for evaluating the possibility of recurrent cardiac events, such as myocardial infarction or restenosis, e.g., after percutaneous coronary intervention. In some embodiments, one or more of CRP level, TNF-alpha level, IL-6 level, or ROS level, or Nt-pro BNP level, or cardiac troponin I level, or mK ATP activity, or MLCL AT1 expression, or ALCAT 1 expression, or Taz1 expression are used in cardiac risk stratification and assessment, and are prognostic factors in conditions such as acute coronary syndrome, stroke peripheral artery disease, and post myocardial infarction complication such as cardiac failure. In some embodiments, one or more of CRP level, TNF-alpha level, IL-6 level, and ROS level, or Nt-pro BNP level, or cardiac troponin I level or mK ATP activity, or MLCL AT1 expression, or ALCAT 1 expression, or Taz1 expression is used to determine the probability of recurrence of cardiac events in patients with stable coronary heart disease and/or acute coronary syndrome. In some embodiments, one or more of CRP level, TNF-alpha level, IL-6 level, or ROS level, or Nt-pro BNP level, or cardiac troponin I level or mK ATP activity, or MLCL AT1 expression, or ALCAT 1 expression, or Taz1 expression is a predictor of early complications and late clinical stenosis and mortality in subjects undergoing percutaneous coronary interventions. In some embodiments, one or more of CRP level, TNF-alpha level, IL-6 level, or ROS level, or Nt-pro BNP level, or cardiac troponin I level, or mK ATP activity, or MLCL AT1 expression, or ALCAT 1 expression, or Taz1 expression add prognostic value in acute coronary syndrome, stable angina, unstable angina, and acute myocardial infarction. In some embodiments, measuring one or more of CRP level, TNF-alpha level, IL-6 level, or ROS level, or Nt-pro BNP level, or cardiac troponin I level, or mK ATP activity, or MLCL AT1 expression, or ALCAT 1 expression, or Taz1 expression add prognostic value in LV remodeling.

[0214] Accordingly, in some embodiments, therapeutic and/or prophylactic treatment of subjects having one or more of elevated CRP level, TNF-alpha level, IL-6 level, ROS level, cardiac troponin I level, or MLCL AT1 expression, and/or ALCAT 1 expression with an aromatic cationic peptide as disclosed herein, such as, e.g., D-Arg-2',6'Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically accept-

able salt thereof, such as acetate or trifluoroacetate salt, reduces one or more of the elevated CRP level, the TNF-alpha level, the IL-6 level, ROS level, the cardiac troponin I level, the MLCL AT1 expression, and/or the ALCAT 1 expression thereby reducing the risk of any of the cardiac, stenotic/vascular events, e.g., LV remodeling, described above. Additionally, or alternatively, in some embodiments, therapeutic and/or prophylactic treatment of subjects having decreased mK ATP activity and/or decreased Taz1 expression with an aromatic cationic peptide as disclosed herein, such as, e.g., D-Arg-2',6'Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof, such as acetate or trifluoroacetate salt, increases mK ATP activity and/or Taz1 expression, thereby reducing the risk of any of the cardiac, stenotic/vascular events, e.g., LV remodeling, described above. In some embodiments, one or more of the CRP level, the TNF-alpha level, the IL-6 level, the ROS level, Nt-pro BNP level, or cardiac troponin I level, or Nt-pro BNP level, or mK ATP activity, or MLCL AT1 expression, or ALCAT 1 expression, or Taz1 expression is normalized after treatment with the aromatic-cationic peptide.

[0215] Therapeutic Methods. The following discussion is presented by way of example only, and is not intended to limit the disclosed methods and compositions to a specific disease or disease state. It is understood that lowering one or more of a subject's CRP level, or TNF-alpha level, or IL-6 level, or ROS level, cardiac troponin I level, MLCL AT1 expression, or ALCAT 1 expression and/or increasing mK ATP activity and/or increasing Taz1 expression in the subject will reduce the risk of any number of negative cardiac, stenotic or vascular events, e.g., LV remodeling. In some embodiments, the present technology includes methods for treating LV remodeling in a subject having or suspected of having an elevated CRP level, or TNF-alpha level, or IL-6 level, or ROS level, or cardiac troponin I level, or MLCL AT1 expression, or ALCAT 1 expression and/or decreased mK ATP activity and/or decreased Taz1 expression for therapeutic purposes. In therapeutic applications, compositions or medicaments are administered to a subject suspected of, or already suffering from such a disease in an amount sufficient to cure, or at least partially arrest, the symptoms of the disease, including its complications and intermediate pathological phenotypes in development of the disease. As such, in some embodiments, the present technology provides methods of treating an individual having or suspected of having an elevated CRP level, TNF-alpha level, IL-6 level, ROS level, cardiac troponin I level, MLCL AT1 expression, and/or ALCAT 1 expression and/or decreased mK ATP and/or decreased Taz1 expression afflicted with heart failure or LV remodeling.

[0216] Subjects suffering from heart failure can be identified by any or a combination of diagnostic or prognostic assays known in the art. For example, typical symptoms of heart failure include shortness of breath (dyspnea), fatigue, weakness, difficulty breathing when lying flat, and swelling of the legs, ankles or abdomen (edema). In some embodiments, the subject may also exhibit elevated CRP level, TNF-alpha level, IL-6 level, ROS level, NT-pro BNP level, cardiac troponin I level, MLCL AT1 expression, and/or ALCAT 1 expression and/or decreased mK ATP and/or decreased Taz1 expression. The subject may also be suffering from other disorders including coronary artery disease, systemic hypertension, cardiomyopathy or myocarditis, con-

genital heart disease, abnormal heart valves or valvular heart disease, severe lung disease, diabetes, severe anemia hyperthyroidism, arrhythmia or dysrhythmia and myocardial infarction. The primary signs of congestive heart failure are cardiomegaly (enlarged heart), tachypnea (rapid breathing; occurs in the case of left side failure) and hepatomegaly (enlarged liver; occurs in the case of right side failure). Acute myocardial infarction ("AMI") due to obstruction of a coronary artery is a common initiating event that can lead ultimately to heart failure. However, a subject that has AMI does not necessarily develop heart failure. Likewise, subjects that suffer from heart failure did not necessarily suffer from an AMI.

[0217] Subjects suffering from LV remodeling can be identified by any or a combination of diagnostic or prognostic assays known in the art. For example, typical symptoms of LV remodeling include decreased LV stroke volume, reduced LV ejection fraction, poor fractional shortening, increased infarct expansion, poor hemodynamics, increased scar formation in LV myocardium, and increased lung volumes. Symptoms of LV remodeling also include symptoms associated with heart failure such as, e.g., shortness of breath, fatigue, and swelling of the extremities. In some embodiments, a "therapeutically effective amount" of aromatic-cationic peptides include levels in which the physiological effects of elevated CRP level, TNF-alpha level, IL-6 level, ROS level, cardiac troponin I level, MLCL AT1 expression, and/or ALCAT 1 expression are, at a minimum, ameliorated. Additionally, or alternatively, in some embodiments, a therapeutically effective amount of the aromatic-cationic peptides include levels in which the physiological effects of decreased mK ATP levels and/or decreased Taz1 expression are, at a minimum, ameliorated. Additionally, or alternatively, in some embodiments, a therapeutically effective amount of the aromatic-cationic peptides include levels in which the physiological effects of LV remodeling are, at a minimum, ameliorated.

[0218] Prophylactic Methods. In some aspects, the invention provides a method for preventing or reducing the likelihood or severity of heart failure in a subject having or suspected of having one or more of an elevated CRP level, TNF-alpha level, or IL-6 level, or ROS level, cardiac troponin I level, MLCL AT1 expression, and/or ALCAT 1 expression and/or decreased mK ATP and/or decreased Taz1 expression by administering to the subject an aromatic-cationic peptide that reduces (e.g., normalizes) one or more of the CRP level, TNF-alpha level, IL-6 level, ROS level, cardiac troponin I level, MLCL AT1 expression, ALCAT 1 expression and/or increases (e.g., normalizes) mK ATP and/or increases Taz1 expression. Subjects at risk for heart failure can be identified by, e.g., any or a combination of diagnostic or prognostic assays as described herein. In prophylactic applications, pharmaceutical compositions or medicaments of aromatic-cationic peptides are administered to a subject susceptible to, or otherwise at risk of a disease or condition in an amount sufficient to eliminate or reduce the risk, lessen the severity, or delay the onset of the disease, including biochemical, histologic and/or behavioral symptoms of the disease, its complications and intermediate pathological phenotypes presenting during development of the disease. Administration of a prophylactic aromatic-cationic can occur prior to the manifestation of symptoms characteristic of the disease or disorder, such that the disease or disorder is prevented or, alternatively, delayed in its

progression. The appropriate compound can be determined based on screening assays described above.

[0219] Subjects diagnosed with or at risk for heart failure may exhibit one or more of the following non-limiting risk factors: high blood pressure; coronary artery disease; heart attack; irregular heartbeats; diabetes; some diabetes medications (e.g., rosiglitazone and pioglitazone have been found to increase the risk of heart failure); sleep apnea; congenital heart defects; viral infection; alcohol use; obesity, lifestyle (e.g., smoking, sedentary lifestyle), high cholesterol, family history, stress, and kidney conditions.

Improvement of Left Ventricular Function

[0220] Patients with HF often have elevated levels of pro-inflammatory cytokines and chemokines, compounds that are involved in adverse left ventricular (LV) remodeling, neurohormonal activation, impaired autonomic and vascular tone, and progression of coronary atherosclerosis (Sola et al. "Atorvastatin Improves Left Ventricular Systolic Function and Serum Markers of Inflammation in Nonischemic Heart Failure." *J. Am. Coll. Cardiol.* 47(2): 332-337 (2006)). Higher levels of these inflammatory markers, including TNF- α , IL-6, and CRP, are also associated with adverse cardiovascular morbidity and mortality. In some embodiments, the reduction of pro-inflammatory cytokines, e.g., TNF-alpha, IL-6, and CRP, by aromatic-cationic peptides, e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), decreases LV remodeling.

[0221] In some embodiments, the reduction of pro-inflammatory cytokines, e.g., TNF-alpha, IL-6, and CRP, by aromatic-cationic peptides, e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), improves LV function in subjects at risk for HF, or suffering from HF, or other stenotic or vascular events. In some embodiments, improvement in LV function (as compared to untreated subjects) is measured by decreased LV stroke volume, increased LV ejection fraction, improved fractional shortening, decreased infarct expansion, good hemodynamics, decreased scar formation in LV myocardium, decreased lung volumes, or a combination thereof.

[0222] In acute myocardial infarction and HF, ROS is generated in the ischemic myocardium, especially after reperfusion. ROS directly injure the cell membrane and cause cell death. ROS also stimulates signal transduction to elaborate inflammatory cytokines, e.g., TNF-alpha, IL-1 beta and -6, in the ischemic region and surrounding myocardium as a host reaction. Inflammatory cytokines also regulate cell survival and cell death in the chain reaction with ROS.

[0223] In some embodiments, the reduction of ROS, by aromatic-cationic peptides, e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), improves LV function after myocardial infarction or in subjects at risk for HF. In some embodiments, improvement in LV function (as compared to untreated subjects) is measured by decreased LV stroke volume, increased LV ejection fraction, improved fractional shortening, decreased infarct expansion, good hemodynamics, decreased scar formation in LV myocardium, decreased lung volumes, or a combination thereof.

[0224] In acute myocardial infarction and HF, cardiac troponin I is generated in the ischemic myocardium. Cardiac troponin I can lead to the increase in cardiac troponin autoantibodies. Increased cardiac troponin autoantibodies can increase cardiac inflammation and increase expression of inflammatory cytokines.

[0225] In some embodiments, the reduction of cardiac troponin I by aromatic-cationic peptides, e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), improves LV function after myocardial infarction or in subjects at risk for HF. In some embodiments, improvement in LV function (as compared to untreated subjects) is measured by decreased LV stroke volume, increased LV ejection fraction, improved fractional shortening, decreased infarct expansion, good hemodynamics, decreased scar formation in LV myocardium, decreased lung volumes, or a combination thereof.

[0226] In acute myocardial infarction and HF, a decrease in the reduced form of NADPH leads to decreased mK ATP activity. The decrease in mK ATP activity leads to ionic deregulation in mitochondrial environment with subsequent matrix contraction and reduced ATP production.

[0227] In some embodiments, the increase in mK ATP activity by aromatic-cationic peptides, e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), improves LV function after myocardial infarction or in subjects at risk for HF. In some embodiments, improvement in LV function (as compared to untreated subjects) is measured by decreased LV stroke volume, increased LV ejection fraction, improved fractional shortening, decreased infarct expansion, good hemodynamics, decreased scar formation in LV myocardium, decreased lung volumes, or a combination thereof.

Determination of the Biological Effect of the Aromatic-Cationic Peptide-Based Therapeutic

[0228] In various embodiments, suitable in vitro or in vivo assays are performed to determine the effect of a specific aromatic-cationic peptide-based therapeutic and whether its administration is indicated for treatment. In various embodiments, in vitro assays can be performed with representative animal models, to determine if a given aromatic-cationic peptide-based therapeutic exerts the desired effect in reducing one or more of CRP levels, TNF-alpha levels, IL-6 levels, ROS levels, cardiac troponin I level, MLCL AT1 expression, and/or ALCAT 1 expression and/or increasing mK ATP activity and/or increasing Taz1 expression, thereby preventing or treating heart failure or LV remodeling. Compounds for use in therapy can be tested in suitable animal model systems including, but not limited to rats, mice, chicken, cows, monkeys, rabbits, and the like, prior to testing in human subjects. Similarly, for in vivo testing, any of the animal model system known in the art can be used prior to administration to human subjects.

[0229] HF has been induced in different species with volume overload, pressure overload, fast pacing, myocardial ischemia, cardiotoxic drugs, or genetically modified models. Models using pressure overload have been most commonly used. Hypertension is associated with an increased risk for the development of HF. In one mouse model, angiotensin II (Ang II) increases blood pressure and induces cardiomyocyte hypertrophy, increased cardiac fibrosis, and impaired cardiomyocyte relaxation. Infusion of angiotensin to mice by mini osmotic pump increases systolic and diastolic blood pressure, increases heart weight and left ventricular thickness (LVMI), and impaired myocardial performance index (MPI). CRP levels, TNF-alpha levels, IL-6 levels, ROS levels, and/or cardiac troponin I levels, MLCL AT1, ALCAT1, and/or Taz1 expression levels, and/or mK ATP activity are monitored at various time points before, during and after HF induction.

[0230] In a second illustrative mouse model, sustained high level expression of Gαq can lead to marked myocyte apoptosis, resulting in cardiac hypertrophy and heart failure by 16 weeks of age (D'Angelo et al., 1998). The β-adrenergic receptors (βARs) are primarily coupled to the heterotrimeric G protein, Gs, to stimulate adenylyl cyclase activity. This association generates intracellular cAMP and protein kinase A activation, which regulate cardiac contractility and heart rate. Overexpression of Gαq leads to decreased responsiveness to β-adrenergic agonists and results in HF. CRP levels, TNF-alpha levels, IL-6 levels, ROS levels, and/or cardiac troponin I levels, MLCL AT1, ALCAT1, and/or Taz1 expression levels, and/or mK ATP activity are monitored at various time points before, during and after HF induction.

[0231] Experimental constriction of the aorta by surgical ligation is also widely used as a model of HF. Transaortic constriction (TAC) results in pressure overload induced HF, with increase in left ventricular (LV) mass. TAC is performed as described by Tamavski O et al. (2004) using a 7-0 silk double-knot suture to constrict the ascending aorta. After TAC, mice develop HF within a period of 4 weeks. CRP levels, TNF-alpha levels, IL-6 levels, ROS levels, and/or cardiac troponin I levels, MLCL AT1, ALCAT1, and/or Taz1 expression levels, and/or mK ATP activity are monitored at various time points before, during and after HF induction.

Prophylactic and Therapeutic Uses for CRP

[0232] As noted above, numerous studies have linked CRP levels, as determined in high sensitivity assays, to cardiac risk. One exemplary risk assessment scale is shown in Table 9:

TABLE 9

hsCRP levels correlated with cardiovascular risk	
hsCRP measurement	Cardiovascular risk
<1 mg/L	low risk
1-3 mg/L	intermediate risk
3-10 mg/L	high risk
>10 mg/L	unspecific elevation

[0233] In some embodiments, a normal, normalized or control CRP level is less than about 1 mg/L. In some embodiment a control, normal or normalized CRP level is between about 1 and 3 mg/L.

[0234] In some embodiments, a lower CRP level is determined by hsCRP assay. In some embodiments, a reduced or lowered CRP level is from a high cardiac risk level (e.g., 3 mg/L or greater) to a lower risk level (e.g., 1-3 mg/L or less than 1 mg/L). In some embodiments, reducing CRP levels is a reduction by about 1 mg/L, about 2 mg/L, about 3 mg/L, about 4 mg/L, about 5 mg/L, about 6 mg/L, about 7 mg/L, about 8 mg/L or about 9 mg/L.

Prophylactic and Therapeutic Uses for ROS

[0235] In some embodiments, a normal, normalized, or control ROS level is less than about 3 RLU×10⁵/ml. In some embodiments, a normal, normalized, or control ROS level is between about 3 RLU×10⁵/ml to about 13 RLU×10⁵/ml.

[0236] In some embodiments, reducing ROS levels is a reduction between about 1 RLU×10⁵/ml to about 5 RLU×

10^5 /ml, between about 5 RLU $\times 10^5$ /ml to about 10 RLU $\times 10^5$ /ml, between about 10 RLU $\times 10^5$ /ml to about 15 RLU $\times 10^5$ /ml, between about 15 RLU $\times 10^5$ /ml to about 20 RLU $\times 10^5$ /ml, or between about 20 RLU $\times 10^5$ /ml to about 25 RLU $\times 10^5$ /ml.

Prophylactic and Therapeutic Uses for IL-6

[0237] In some embodiments, a normal, normalized, or control IL-6 level is less than about 10 pg/ml. In some embodiments, a normal, normalized, or control IL-6 level is between about 5 pg/ml to about 13 pg/ml.

[0238] In some embodiments, reducing IL-6 levels is a reduction between about 1 pg/ml to about 5 pg/ml, between about 5 pg/ml to about 10 pg/ml, between about 10 pg/ml to about 15 pg/ml, or between about 15 pg/ml to about 20 pg/ml.

Prophylactic and Therapeutic Uses for TNF-Alpha

[0239] In some embodiments, a normal, normalized, or control TNF-alpha level is less than about 1.5 pg/ml. In some embodiments, a normal, normalized, or control TNF-alpha level is between about 1 pg/ml to about 2 pg/ml.

[0240] In some embodiments, reducing TNF-alpha levels is a reduction about 1 pg/ml, or about 2 pg/ml, or about 3 pg/ml, or about 4 pg/ml, or 5 pg/ml.

Prophylactic and Therapeutic Uses for MLCL AT1, ALCAT1, and Taz1

[0241] Therapeutic Methods: The following discussion is presented by way of example only, and is not intended to limit the disclosed methods and compositions to a specific disease or disease state. It is understood that lowering the expression of MLCL AT1 or ALCAT1 and/or raising the expression Taz1 in a subject in need thereof will reduce the risk of any number of negative cardiac, stenotic or vascular events. One aspect of the present technology includes methods of treating heart failure in a subject having or suspected of having an elevated MLCL AT1 or ALCAT1 expression and/or lowered Taz1 expression for therapeutic purposes. In therapeutic applications, compositions or medicaments are administered to a subject suspected of, or already suffering from such a disease in an amount sufficient to cure, or at least partially arrest, the symptoms of the disease, including its complications and intermediate pathological phenotypes in development of the disease. As such, in some embodiments, the present technology provides methods of treating an individual having or suspected of having an elevated MLCL AT1 or ALCAT1 expression level afflicted with heart failure. Alternatively, or additionally, in some embodiments, the present technology provides methods of treating an individual having or suspected of having an decreased Taz1 expression afflicted with heart failure.

[0242] Subjects suffering from heart failure can be identified by any or a combination of diagnostic or prognostic assays known in the art. For example, typical symptoms of heart failure include shortness of breath (dyspnea), fatigue, weakness, difficulty breathing when lying flat, and swelling of the legs, ankles or abdomen (edema). In some embodiments, the subject may also exhibit elevated MLCL AT1 or ALCAT1 expression and/or lowered Taz1 expression. The subject may also be suffering from other disorders including coronary artery disease, systemic hypertension, cardiomyopathy or myocarditis, congenital heart disease, abnor-

mal heart valves or valvular heart disease, severe lung disease, diabetes, severe anemia hyperthyroidism, arrhythmia or dysrhythmia and myocardial infarction. The primary signs of congestive heart failure are cardiomegaly (enlarged heart), tachypnea (rapid breathing; occurs in the case of left side failure) and hepatomegaly (enlarged liver; occurs in the case of right side failure). Acute myocardial infarction ("AMI") due to obstruction of a coronary artery is a common initiating event that can lead ultimately to heart failure. However, a subject that has AMI does not necessarily develop heart failure. Likewise, subjects that suffer from heart failure did not necessarily suffer from an AMI.

[0243] Prophylactic Methods: In one aspect, the present technology provides a method for preventing heart failure in a subject having or suspected of having one or more of an elevated MLCL AT1 or ALCAT1 expression and/or decreased Taz1 expression, by administering to the subject an aromatic-cationic peptide that normalizes one or more of the MLCL AT1, ALCAT1, or Taz1 expression levels. Subjects at risk for heart failure can be identified by, e.g., any or a combination of diagnostic or prognostic assays as described herein. In prophylactic applications, pharmaceutical compositions or medicaments of aromatic-cationic peptides are administered to a subject susceptible to, or otherwise at risk of a disease or condition in an amount sufficient to eliminate or reduce the risk, lessen the severity, or delay the onset of the disease, including biochemical, histologic and/or behavioral symptoms of the disease, its complications and intermediate pathological phenotypes presenting during development of the disease. Administration of a prophylactic aromatic-cationic can occur prior to the manifestation of symptoms characteristic of the disease or disorder, such that the disease or disorder is prevented or, alternatively, delayed in its progression. The appropriate compound can be determined based on screening assays described above.

[0244] Subjects diagnosed with or at risk for heart failure may exhibit one or more of the following non-limiting risk factors: high blood pressure; coronary artery disease; heart attack; irregular heartbeats; diabetes; some diabetes medications (e.g., rosiglitazone and pioglitazone have been found to increase the risk of heart failure); sleep apnea; congenital heart defects; viral infection; alcohol use; obesity, lifestyle (e.g., smoking, sedentary lifestyle), high cholesterol, family history, stress, and kidney conditions.

C-Reactive Protein as a Biomarker for Peptide Dosage.

[0245] In some embodiments, C-reactive protein levels are used to determine the dosage of aromatic-cationic peptide (e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3)) to administer to a subject. For example, in some embodiments, if the level of C-reactive protein in a subject is less than about 1 mg/L, no or low levels of an aromatic cationic peptide of the present disclosure (e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3)) is administered to the subject. In some embodiments, if a subject's C-reactive protein level is between about 1-3 mg/L, a medium level of an aromatic-cationic peptide of the present disclosure is administered, and if a subject's level of C-reactive protein is between about 3-10 mg/L, a high level of an aromatic-cationic peptide of the present disclosure is administered to the subject. In some embodiments, a low dose of an aromatic-cationic peptide, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), results in a plasma concentration of less

than about 50 ng/ml. In some embodiments, a low dose of aromatic-cationic peptide results in a plasma concentration of about 1, about 5, about 10, about 20, about 30, or about 40 ng/ml. In some embodiments, a low dose or aromatic-cationic peptide results in a plasma concentration of between about 10-40 ng/ml, between about 10-30 ng/ml, between about 10-20 ng/ml, between about 20-40 ng/ml, between about 20-30 ng/ml, between about 30-40 ng/ml, between about 30-50 ng/ml, or between about 40-50 ng/ml. In some embodiments, a medium dose or an aromatic-cationic peptide, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), results in a plasma concentration of between about 50-500 ng/ml. In some embodiments, a medium dose of aromatic-cationic peptide results in a plasma concentration of about 50 ng/ml, about 100 ng/ml, about 150 ng/ml, about 200 ng/ml, about 250 ng/ml, about 300 ng/ml, about 350 ng/ml, about 400 ng/ml or about 450 ng/ml. In some embodiments, a medium dose of aromatic-cationic peptide results in a plasma concentration of between about 50-100 ng/ml, between about 50-200 ng/ml, between about 50-300 ng/ml, between about 50-400 ng/ml, between about 100-200 ng/ml, between about 100-300 ng/ml, between about 100-400 ng/ml, between about 100-500 ng/ml, between about 200-300 ng/ml, between about 200-400 ng/ml, between about 200-500 ng/ml, between about 300-400 ng/ml, between about 300-500 ng/ml or between about 400-500 ng/ml. In some embodiments, a high dose of an aromatic-cationic peptide results in a plasma concentration of between about 500-5000 ng/ml. In some embodiments, a high dose of aromatic-cationic peptide results in a plasma concentration of about 1000 ng/ml, about 1500 ng/ml, about 2000 ng/ml, about 2500 ng/ml, about 3000 ng/ml, about 3500 ng/ml, about 4000 ng/ml or about 4500 ng/ml. In some embodiments, a high dose of aromatic-cationic peptide results in a plasma concentration of between about 500-1000 ng/ml, between about 500-2000 ng/ml, between about 500-3000 ng/ml, between about 500-4000 ng/ml, between about 1000-2000 ng/ml, between about 1000-3000 ng/ml, between about 1000-4000 ng/ml, between about 1000-5000 ng/ml, between about 2000-3000 ng/ml, between about 2000-4000 ng/ml, between about 2000-5000 ng/ml, between about 3000-4000 ng/ml, between about 3000-5000 ng/ml or between about 4000-5000 ng/ml. In some embodiments, a high dose of an aromatic-cationic peptide, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), results in a plasma concentration of greater than 5000 ng/ml. For example, in some embodiments, a high dose of an aromatic-cationic peptide results in a plasma concentration of about 6000 ng/ml, about 7000 ng/ml, about 8000 ng/ml, about 9000 ng/ml, or about 10,000 ng/ml. In some embodiments, a high dose of an aromatic-cationic peptide results in a plasma concentration of between about 5000-10,000 ng/ml, between about 5000-9000 ng/ml, between about 5000-8000 ng/ml, between about 5000-7000 ng/ml, between about 5000-6000 ng/ml, between about 6000-7000 ng/ml, between about 6000-8000 ng/ml, between about 6000-9000 ng/ml, between about 6000-10,000 ng/ml, between about 7000-8000 ng/ml, between about 7000-9000 ng/ml, between about 7000-10,000 ng/ml, between about 8000-9000 ng/ml, or between about 9000-10000 ng/ml. As described below, those skilled in the art will understand that dosage, toxicity and therapeutic efficacy of the aromatic-cationic peptides disclosed herein can be determined by standard pharmaceutical procedures in cell cultures or experimental animals.

[0246] For peptide dosage information, prophylactic or therapeutic treatment, CRP levels can be determined by any or a combination of methods known in the art. For example, in some embodiments, a hsCRP or cCRP assay is performed. In some embodiments, a subject's CRP level is between about 1-3 mg/L prior to peptide administration. In some embodiments, the subject's CRP level is between about 3-10 mg/L before peptide administration. In some embodiments, the subject's CRP level is greater than 10 mg/L before peptide administration. In some embodiments, the subject's CRP level is about 2 fold greater than a control level before peptide administration. In some embodiments, the subject's CRP level is about 3 fold greater than a control level before peptide administration. In some embodiments, the subject's CRP level is about 4, 5 or 6 fold greater than a control before peptide administration.

Ros, TNF-Alpha, or IL-6 as Biomarkers for Peptide Dosage.

[0247] TNF- α and IL-6 levels in plasma are typically reported in pg/ml and can range between about 0.1 pg/ml to about 2000 pg/ml depending on a pathophysiological condition one is working with. The total ROS amount in plasma is typically reported in RLU (relative light units/ml). The range of ROS level can be as low as about 1.0 RLU $\times 10^5$ /ml and as high as about 200 RLU $\times 10^5$ /ml depending on the physiological condition one is working with. Thus, in some embodiments, ROS levels, TNF-alpha and/or IL-6 levels are used to determine the dosage of aromatic-cationic peptide (e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3)) to administer to a subject. For example, in some embodiments, if the level of ROS, TNF-alpha, or IL-6 protein in a subject is low (e.g., about 0.1-500 pg/ml for TNF-alpha, or IL-6 or about 1-70 RLU $\times 10^5$ /ml for ROS), then a lower dose of aromatic-cationic peptide would be administered. Additionally or alternatively, in some embodiments, if the level of ROS, TNF-alpha, or IL-6 protein in a subject is in a medium range (e.g., about 500-1000 pg/ml for TNF-alpha, or IL-6 or about 70-130 RLU $\times 10^5$ /ml for ROS), then a medium dose of aromatic-cationic peptide would be administered. Additionally or alternatively, in some embodiments, if the level of ROS, TNF-alpha, or IL-6 protein in a subject is high (e.g., about 1000-2000 pg/ml for TNF- α or IL-6 or about 130-200 RLU $\times 10^5$ /ml for ROS), then a high dose of aromatic-cationic peptide would be administered.

[0248] For peptide dosage information, prophylactic or therapeutic treatment, ROS levels can be determined by any or a combination of methods known in the art. For example, in some embodiments, an absorbance, a fluorescence, or a luminescence assay is performed. In some embodiments, a subject's ROS level is between about 1 RLU $\times 10^5$ /ml to about 5 RLU $\times 10^5$ /ml prior to peptide administration. In some embodiments, the subject's ROS level is between about 5 RLU $\times 10^5$ /ml to about 10 RLU $\times 10^5$ /ml, or between about 10 RLU $\times 10^5$ /ml to about 20 RLU $\times 10^5$ /ml, or between about 20 RLU $\times 10^5$ /ml to about 30 RLU $\times 10^5$ /ml before peptide administration. In some embodiments, the subject's ROS level is greater than 30 RLU $\times 10^5$ /ml before peptide administration. In some embodiments, the subject's ROS level is about 1.5 fold greater than a control level before peptide administration. In some embodiments, the subject's ROS level is about 2 fold greater than a control level before peptide administration. In some embodiments, the subject's ROS level is about 3, 4, 5, 6, 7, 8 fold or more greater than a control before peptide administration.

[0249] For peptide dosage information, prophylactic or therapeutic treatment, IL-6 levels can be determined by any or a combination of methods known in the art. For example, in some embodiments, an ELISA or HPLC assays is performed. In some embodiments, a subject's IL-6 level is about 8 pg/ml prior to peptide administration. In some embodiments, the subject's IL-6 level is between about 5-13 pg/ml before peptide administration. In some embodiments, the subject's IL-6 level is greater than 13 pg/ml before peptide administration. In some embodiments, the subject's IL-6 level is about 1.5 fold greater than a control level before peptide administration. In some embodiments, the subject's IL-6 level is about 2 fold greater than a control level before peptide administration. In some embodiments, the subject's IL-6 level is about 3, 4, 5, 6, 7, 8 fold or more greater than a control before peptide administration.

[0250] For peptide dosage information, prophylactic or therapeutic treatment, TNF-alpha levels can be determined by any or a combination of methods known in the art. For example, in some embodiments, an ELISA or HPLC assays is performed. In some embodiments, a subject's TNF-alpha level is about 1 pg/ml prior to peptide administration. In some embodiments, the subject's TNF-alpha level is between about 1.0-1.5 pg/ml before peptide administration. In some embodiments, the subject's TNF-alpha level is greater than 1.5 pg/ml before peptide administration. In some embodiments, the subject's TNF-alpha level is about 2 fold greater than a control level before peptide administration. In some embodiments, the subject's TNF-alpha level is about 3 fold greater than a control level before peptide administration. In some embodiments, the subject's TNF-alpha level is about 4, 5, 6, 7 or 8 fold or more greater than a control before peptide administration.

Nt-Pro BNP as a Biomarker for Peptide Dosage

[0251] Nt-pro BNP in plasma can be reported in pg/ml and can range between about 1 pg/ml to about 3000 pg/ml depending on a pathophysiological condition one is working with. Thus, in some embodiments, Nt-pro BNP is used to determine the dosage of aromatic-cationic peptide (e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3)) to administer to a subject. For example, in some embodiments, if the level of Nt-pro BNP in a subject is low (e.g., about 1-1000 pg/ml), then a lower dose of aromatic-cationic peptide would be administered. Additionally or alternatively, in some embodiments, if the level of Nt-pro BNP in a subject is in a medium range (e.g., about 1000-2000 pg/ml), then a medium dose of aromatic-cationic peptide would be administered. Additionally or alternatively, in some embodiments, if the level of Nt-pro BNP in a subject is high (e.g., about 2000-3000 pg/ml), then a high dose of aromatic-cationic peptide would be administered.

[0252] For peptide dosage information, Nt-pro BNP levels can be determined by any or a combination of methods known in the art. For example, in some embodiments, an ELISA or HPLC assays is performed. In some embodiments, a subject's Nt-pro BNP level is about 250 pg/ml prior to peptide administration. In some embodiments, the subject's Nt-pro BNP level is between about 150-400 pg/ml before peptide administration. In some embodiments, the subject's Nt-pro BNP level is greater than 400 pg/ml before peptide administration. In some embodiments, the subject's Nt-pro BNP level is about 5 fold greater than a control level before peptide administration. In some embodiments, the subject's

Nt-pro BNP level is about 6 fold greater than a control level before peptide administration. In some embodiments, the subject's Nt-pro BNP level is about 3, 4, 5, 6, 7, 8 fold or more greater than a control before peptide administration.

[0253] In some embodiments, Nt-pro BNP levels are measured as a biomarker for the decrease or prevention of LV remodeling after administration of aromatic-cationic peptide to a subject in need thereof. For example, in some embodiments, administration with aromatic-cationic peptides disclosed herein, e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), reduces Nt-pro BNP by about 30%, about 40%, about 50% about 60%, about 70%, or about 80%. In some embodiments, administration with aromatic-cationic peptides returns Nt-pro BNP to or near baseline levels.

Cardiac Troponin I as a Biomarker for Peptide Dosage

[0254] In some embodiments, the subject's Nt-pro BNP level is about 3, 4, 5, 6, 7, 8 fold or more greater than a control before peptide administration.

[0255] In some embodiments, cardiac troponin I levels are measured as a biomarker for the decrease or prevention of LV remodeling after administration of aromatic-cationic peptide to a subject in need thereof. For example, in some embodiments, administration with aromatic-cationic peptides disclosed herein, e.g., D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), reduces cardiac troponin I by about 10%, about 20%, about 30%, about 40%, about 50% about 60%, about 70%, about 80%, or about 90%. In some embodiments, administration with aromatic-cationic peptides returns cardiac troponin I to or near baseline levels.

Modes of Administration and Effective Dosages

[0256] Any method known to those in the art for contacting a cell, organ or tissue with a peptide may be employed. Suitable methods include in vitro, ex vivo, or in vivo methods. In vivo methods typically include the administration of an aromatic-cationic peptide, such as those described above, to a mammal, suitably a human. When used in vivo for therapy, the aromatic-cationic peptides are administered to the subject in effective amounts (i.e., amounts that have desired therapeutic effect). The dose and dosage regimen will depend upon the degree of the infection in the subject, the characteristics of the particular aromatic-cationic peptide used, e.g., its therapeutic index, the subject, and the subject's history.

[0257] The effective amount may be determined during pre-clinical trials and clinical trials by methods familiar to physicians and clinicians. An effective amount of a peptide useful in the methods may be administered to a mammal in need thereof by any of a number of well-known methods for administering pharmaceutical compounds. The peptide may be administered systemically or locally.

[0258] The peptide may be formulated as a pharmaceutically acceptable salt. Salts derived from pharmaceutically acceptable inorganic bases include ammonium, calcium, copper, ferric, ferrous, lithium, magnesium, manganese, potassium, sodium, and zinc salts, and the like. Salts derived from pharmaceutically acceptable organic bases include salts of primary, secondary and tertiary amines, including substituted amines, cyclic amines, naturally-occurring amines and the like, such as arginine, betaine, caffeine, choline, N,N'-dibenzylethylenediamine, diethylamine, 2-diethylaminoethanol, 2-dimethylaminoethanol,

ethanolamine, ethylenediamine, N-ethylmorpholine, N-ethylpiperidine, glucamine, glucosamine, histidine, hydramidine, isopropylamine, lysine, methylglucamine, morpholine, piperazine, piperidine, polyamine resins, procaine, purines, theobromine, triethylamine, trimethylamine, tripropylamine, tromethamine and the like. Salts derived from pharmaceutically acceptable inorganic acids include salts of boric, carbonic, hydrohalic (hydrobromic, hydrochloric, hydrofluoric or hydroiodic), nitric, phosphoric, sulfamic and sulfuric acids. Salts derived from pharmaceutically acceptable organic acids include salts of aliphatic hydroxyl acids (e.g., citric, gluconic, glycolic, lactic, lactobionic, malic, and tartaric acids), aliphatic monocarboxylic acids (e.g., acetic, butyric, formic, propionic and trifluoroacetic acids), amino acids (e.g., aspartic and glutamic acids), aromatic carboxylic acids (e.g., benzoic, p-chlorobenzoic, diphenylacetic, gentisic, hippuric, and triphenylacetic acids), aromatic hydroxyl acids (e.g., o-hydroxybenzoic, p-hydroxybenzoic, 1-hydroxynaphthalene-2-carboxylic and 3-hydroxynaphthalene-2-carboxylic acids), ascorbic, dicarboxylic acids (e.g., fumaric, maleic, oxalic and succinic acids), glucuronic, mandelic, mucic, nicotinic, orotic, pamoic, pantothenic, sulfonic acids (e.g., benzenesulfonic, camphorsulfonic, edisyllic, ethanesulfonic, isethionic, methanesulfonic, naphthalenesulfonic, naphthalene-1,5-disulfonic, naphthalene-2,6-disulfonic and p-toluenesulfonic acids), xinafoic acid, and the like. In some embodiments, the salt is an acetate or trifluoroacetate salt.

[0259] The aromatic-cationic peptides described herein can be incorporated into pharmaceutical compositions for administration, singly or in combination, to a subject for the treatment or prevention of a disorder described herein. Such compositions typically include the active agent and a pharmaceutically acceptable carrier. As used herein the term "pharmaceutically acceptable carrier" includes saline, solvents, dispersion media, coatings, antibacterial and antifungal agents, isotonic and absorption delaying agents, and the like, compatible with pharmaceutical administration. Supplementary active compounds can also be incorporated into the compositions.

[0260] Pharmaceutical compositions are typically formulated to be compatible with its intended route of administration. Examples of routes of administration include parenteral (e.g., intravenous, intradermal, intraperitoneal or subcutaneous), oral, inhalation, transdermal (topical), intraocular, iontophoretic, and transmucosal administration. Solutions or suspensions used for parenteral, intradermal, or subcutaneous application can include the following components: a sterile diluent such as water for injection, saline solution, fixed oils, polyethylene glycols, glycerine, propylene glycol or other synthetic solvents; antibacterial agents such as benzyl alcohol or methyl parabens; antioxidants such as ascorbic acid or sodium bisulfite; chelating agents such as ethylenediaminetetraacetic acid; buffers such as acetates, citrates or phosphates and agents for the adjustment of tonicity such as sodium chloride or dextrose. pH can be adjusted with acids or bases, such as hydrochloric acid or sodium hydroxide. The parenteral preparation can be enclosed in ampoules, disposable syringes or multiple dose vials made of glass or plastic. For convenience of the patient or treating physician, the dosing formulation can be provided in a kit containing all necessary equipment (e.g., vials of drug, vials of diluent, syringes and needles) for a treatment course (e.g., 7 days of treatment).

[0261] Pharmaceutical compositions suitable for injectable use can include sterile aqueous solutions (where water soluble) or dispersions and sterile powders for the extemporaneous preparation of sterile injectable solutions or dispersion. For intravenous administration, suitable carriers include physiological saline, bacteriostatic water, Cremophor EL™ (BASF, Parsippany, N.J.) or phosphate buffered saline (PBS). In all cases, a composition for parenteral administration must be sterile and should be fluid to the extent that easy syringability exists. It should be stable under the conditions of manufacture and storage and must be preserved against the contaminating action of microorganisms such as bacteria and fungi.

[0262] The aromatic-cationic peptide compositions can include a carrier, which can be a solvent or dispersion medium containing, for example, water, ethanol, polyol (for example, glycerol, propylene glycol, and liquid polyethylene glycol, and the like), and suitable mixtures thereof. The proper fluidity can be maintained, for example, by the use of a coating such as lecithin, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants. Prevention of the action of microorganisms can be achieved by various antibacterial and antifungal agents, for example, parabens, chlorobutanol, phenol, ascorbic acid, thiomersol, and the like. Glutathione and other antioxidants can be included to prevent oxidation. In many cases, it will be preferable to include isotonic agents, for example, sugars, polyalcohols such as mannitol, sorbitol, or sodium chloride in the composition. Prolonged absorption of the injectable compositions can be brought about by including in the composition an agent that delays absorption, for example, aluminum monostearate or gelatin.

[0263] Sterile injectable solutions can be prepared by incorporating the active compound in the required amount in an appropriate solvent with one or a combination of ingredients enumerated above, as required, followed by filtered sterilization. Generally, dispersions are prepared by incorporating the active compound into a sterile vehicle, which contains a basic dispersion medium and the required other ingredients from those enumerated above. In the case of sterile powders for the preparation of sterile injectable solutions, typical methods of preparation include vacuum drying and freeze drying, which can yield a powder of the active ingredient plus any additional desired ingredient from a previously sterile-filtered solution thereof.

[0264] Oral compositions generally include an inert diluent or an edible carrier. For the purpose of oral therapeutic administration, the active compound can be incorporated with excipients and used in the form of tablets, troches, or capsules, e.g., gelatin capsules. Oral compositions can also be prepared using a fluid carrier for use as a mouthwash. Pharmaceutically compatible binding agents, and/or adjuvant materials can be included as part of the composition. The tablets, pills, capsules, troches and the like can contain any of the following ingredients, or compounds of a similar nature: a binder such as microcrystalline cellulose, gum tragacanth or gelatin; an excipient such as starch or lactose, a disintegrating agent such as alginic acid, Primogel, or corn starch; a lubricant such as magnesium stearate or Sterotes; a glidant such as colloidal silicon dioxide; a sweetening agent such as sucrose or saccharin; or a flavoring agent such as peppermint, methyl salicylate, or orange flavoring.

[0265] For administration by inhalation, the compounds can be delivered in the form of an aerosol spray from a

pressurized container or dispenser, which contains a suitable propellant, e.g., a gas such as carbon dioxide, or a nebulizer. Such methods include those described in U.S. Pat. No. 6,468,798.

[0266] Systemic administration of a therapeutic compound as described herein can also be by transmucosal or transdermal means. For transmucosal or transdermal administration, penetrants appropriate to the barrier to be permeated are used in the formulation. Such penetrants are generally known in the art, and include, for example, for transmucosal administration, detergents, bile salts, and fusidic acid derivatives. Transmucosal administration can be accomplished through the use of nasal sprays. For transdermal administration, the active compounds are formulated into ointments, salves, gels, or creams as generally known in the art. In one embodiment, transdermal administration may be performed by iontophoresis.

[0267] A therapeutic protein or peptide can be formulated in a carrier system. The carrier can be a colloidal system. The colloidal system can be a liposome, a phospholipid bilayer vehicle. In one embodiment, the therapeutic peptide is encapsulated in a liposome while maintaining peptide integrity. One skilled in the art would appreciate, there are a variety of methods to prepare liposomes. (See Lichtenberg et al., *Methods Biochem. Anal.*, 33:337-462 (1988); Anselem et al., *Liposome Technology*, CRC Press (1993)). Liposomal formulations can delay clearance and increase cellular uptake (See Reddy, *Ann. Pharmacother.*, 34(7-8):915-923 (2000)). An active agent can also be loaded into a particle prepared from pharmaceutically acceptable ingredients including, but not limited to, soluble, insoluble, permeable, impermeable, biodegradable or gastroretentive polymers or liposomes. Such particles include, but are not limited to, nanoparticles, biodegradable nanoparticles, microparticles, biodegradable microparticles, nanospheres, biodegradable nanospheres, microspheres, biodegradable microspheres, capsules, emulsions, liposomes, micelles and viral vector systems.

[0268] The carrier can also be a polymer, e.g., a biodegradable, biocompatible polymer matrix. In one embodiment, the therapeutic peptide can be embedded in the polymer matrix, while maintaining protein integrity. The polymer may be natural, such as polypeptides, proteins or polysaccharides, or synthetic, such as poly α -hydroxy acids. Examples include carriers made of, e.g., collagen, fibronectin, elastin, cellulose acetate, cellulose nitrate, polysaccharide, fibrin, gelatin, and combinations thereof. In one embodiment, the polymer is poly-lactic acid (PLA) or copoly lactic/glycolic acid (PGLA). The polymeric matrices can be prepared and isolated in a variety of forms and sizes, including microspheres and nanospheres. Polymer formulations can lead to prolonged duration of therapeutic effect. (See Reddy, *Ann. Pharmacother.*, 34(7-8):915-923 (2000)). A polymer formulation for human growth hormone (hGH) has been used in clinical trials. (See Kozarich and Rich, *Chemical Biology*, 2:548-552 (1998)).

[0269] Examples of polymer microsphere sustained release formulations are described in PCT publication WO 99/15154 (Tracy et al.), U.S. Pat. Nos. 5,674,534 and 5,716,644 (both to Zale et al.), PCT publication WO 96/40073 (Zale et al.), and PCT publication WO 00/38651 (Shah et al.). U.S. Pat. Nos. 5,674,534 and 5,716,644 and PCT publication WO 96/40073 describe a polymeric matrix

containing particles of erythropoietin that are stabilized against aggregation with a salt.

[0270] In some embodiments, the therapeutic compounds are prepared with carriers that will protect the therapeutic compounds against rapid elimination from the body, such as a controlled release formulation, including implants and microencapsulated delivery systems. Biodegradable, biocompatible polymers can be used, such as ethylene vinyl acetate, polyanhydrides, polyglycolic acid, collagen, polyorthoesters, and polylactic acid. Such formulations can be prepared using known techniques. The materials can also be obtained commercially, e.g., from Alza Corporation and Nova Pharmaceuticals, Inc. Liposomal suspensions (including liposomes targeted to specific cells with monoclonal antibodies to cell-specific antigens) can also be used as pharmaceutically acceptable carriers. These can be prepared according to methods known to those skilled in the art, for example, as described in U.S. Pat. No. 4,522,811.

[0271] The therapeutic compounds can also be formulated to enhance intracellular delivery. For example, liposomal delivery systems are known in the art, see, e.g., Chonn and Cullis, "Recent Advances in Liposome Drug Delivery Systems," *Current Opinion in Biotechnology* 6:698-708 (1995); Weiner, "Liposomes for Protein Delivery: Selecting Manufacture and Development Processes," *Immunomethods*, 4(3): 201-9 (1994); and Gregoriadis, "Engineering Liposomes for Drug Delivery: Progress and Problems," *Trends Biotechnol.*, 13(12):527-37 (1995). Mizguchi et al., *Cancer Lett.*, 100: 63-69 (1996), describes the use of fusogenic liposomes to deliver a protein to cells both in vivo and in vitro.

[0272] Dosage, toxicity and therapeutic efficacy of the therapeutic agents can be determined by standard pharmaceutical procedures in cell cultures or experimental animals, e.g., for determining the LD₅₀ (the dose lethal to 50% of the population) and the ED₅₀ (the dose therapeutically effective in 50% of the population). The dose ratio between toxic and therapeutic effects is the therapeutic index and it can be expressed as the ratio LD₅₀/ED₅₀. Compounds that exhibit high therapeutic indices are preferred. While compounds that exhibit toxic side effects may be used, care should be taken to design a delivery system that targets such compounds to the site of affected tissue in order to minimize potential damage to uninjected cells and, thereby, reduce side effects.

[0273] The data obtained from the cell culture assays and animal studies can be used in formulating a range of dosage for use in humans. The dosage of such compounds lies preferably within a range of circulating concentrations that include the ED₅₀ with little or no toxicity. The dosage may vary within this range depending upon the dosage form employed and the route of administration utilized. For any compound used in the methods, the therapeutically effective dose can be estimated initially from cell culture assays. A dose can be formulated in animal models to achieve a circulating plasma concentration range that includes the IC₅₀ (i.e., the concentration of the test compound which achieves a half-maximal inhibition of symptoms) as determined in cell culture. Such information can be used to determine useful doses in humans accurately. Levels in plasma may be measured, for example, by high performance liquid chromatography.

[0274] Typically, an effective amount of the aromatic-cationic peptides, sufficient for achieving a therapeutic or prophylactic effect, range from about 0.000001 mg per

kilogram body weight per day to about 10,000 mg per kilogram body weight per day. Suitably, the dosage ranges are from about 0.0001 mg per kilogram body weight per day to about 100 mg per kilogram body weight per day. For example dosages can be 1 mg/kg body weight or 10 mg/kg body weight every day, every two days or every three days or within the range of 1-10 mg/kg every week, every two weeks or every three weeks. In one embodiment, a single dosage of peptide ranges from 0.001-10,000 micrograms per kg body weight. In one embodiment, aromatic-cationic peptide concentrations in a carrier range from 0.2 to 2000 micrograms per delivered milliliter. An exemplary treatment regime entails administration once per day or once a week. In therapeutic applications, a relatively high dosage at relatively short intervals is sometimes required until progression of the disease is reduced or terminated, and preferably until the subject shows partial or complete amelioration of symptoms of disease. Thereafter, the patient can be administered a prophylactic regime.

[0275] In some embodiments, a therapeutically effective amount of an aromatic-cationic peptide may be defined as a concentration of peptide at the target tissue of 10^{-12} to 10^{-6} molar, e.g., approximately 10^{-7} molar. This concentration may be delivered by systemic doses of 0.001 to 100 mg/kg or equivalent dose by body surface area. The schedule of doses would be optimized to maintain the therapeutic concentration at the target tissue, most preferably by single daily or weekly administration, but also including continuous administration (e.g., parenteral infusion or transdermal application).

[0276] The skilled artisan will appreciate that certain factors may influence the dosage and timing required to effectively treat a subject, including but not limited to, the severity of the disease or disorder, previous treatments, the general health and/or age of the subject, and other diseases present. Moreover, treatment of a subject with a therapeutically effective amount of the therapeutic compositions described herein can include a single treatment or a series of treatments.

[0277] The mammal treated in accordance present methods can be any mammal, including, for example, farm animals, such as sheep, pigs, cows, and horses; pet animals, such as dogs and cats; laboratory animals, such as rats, mice and rabbits. In a preferred embodiment, the mammal is a human.

Combination Therapy with an Aromatic-Cationic Peptide and Other Therapeutic Agents

[0278] In some embodiments, the aromatic-cationic peptides may be combined with one or more additional agents for the prevention or treatment of heart failure. Drug treatment for heart failure typically involves diuretics, ACE inhibitors, digoxin (also called *digitalis*), calcium channel blockers, and beta-blockers. In mild cases, thiazide diuretics, such as hydrochlorothiazide at 25-50 mg/day or chlorothiazide at 250-500 mg/day, are useful. However, supplemental potassium chloride may be needed, since chronic diuresis causes hypokalemia alkalosis. Moreover, thiazide diuretics usually are not effective in patients with advanced symptoms of heart failure. Typical doses of ACE inhibitors include captopril at 25-50 mg/day and quinapril at 10 mg/day.

[0279] In one embodiment, the aromatic-cationic peptide is combined with an adrenergic beta-2 agonist. An "adrenergic beta-2 agonist" refers to adrenergic beta-2 agonists and analogues and derivatives thereof, including, for example,

natural or synthetic functional variants, which have adrenergic beta-2 agonist biological activity, as well as fragments of an adrenergic beta-2 agonist having adrenergic beta-2 agonist biological activity. The term "adrenergic beta-2 agonist biological activity" refers to activity that mimics the effects of adrenaline and noradrenaline in a subject and which improves myocardial contractility in a patient having heart failure. Commonly known adrenergic beta-2 agonists include, but are not limited to, clenbuterol, albuterol, formoterol, levalbuterol, metaproterenol, pирbutерол, salmeterol, and terbutaline.

[0280] In one embodiment, the aromatic-cationic peptide is combined with an adrenergic beta-1 antagonist. Adrenergic beta-1 antagonists and adrenergic beta-1 blockers refer to adrenergic beta-1 antagonists and analogues and derivatives thereof, including, for example, natural or synthetic functional variants which have adrenergic beta-1 antagonist biological activity, as well as fragments of an adrenergic beta-1 antagonist having adrenergic beta-1 antagonist biological activity. Adrenergic beta-1 antagonist biological activity refers to activity that blocks the effects of adrenaline on beta receptors. Commonly known adrenergic beta-1 antagonists include, but are not limited to, acebutolol, atenolol, betaxolol, bisoprolol, esmolol, and metoprolol.

[0281] Clenbuterol, for example, is available under numerous brand names including Spiropent® (Boehringer Ingelheim), Broncodil® (Von Boch I), Broncotrol® (Quimedical PT), Cesbron® (Fidelis PT), and Clenbuter® (Biomedica Foscama). Similarly, methods of preparing adrenergic beta-1 antagonists such as metoprolol and their analogues and derivatives are well-known in the art. Metoprolol, in particular, is commercially available under the brand names Lopressor® (metoprolol tartate) manufactured by Novartis Pharmaceuticals Corporation, One Health Plaza, East Hanover, N.J. 07936-1080. Generic versions of Lopressor® are also available from Mylan Laboratories Inc., 1500 Corporate Drive, Suite 400, Canonsburg, Pa. 15317; and Watson Pharmaceuticals, Inc., 360 Mt. Kemble Ave. Morristown, N.J. 07962. Metoprolol is also commercially available under the brand name Toprol XL®, manufactured by Astra Zeneca, LP.

[0282] In one embodiment, an additional therapeutic agent is administered to a subject in combination with an aromatic cationic peptide, such that a synergistic therapeutic effect is produced. Therefore, lower doses of one or both of the therapeutic agents may be used in treating heart failure, resulting in increased therapeutic efficacy and decreased side-effects.

[0283] In any case, the multiple therapeutic agents may be administered in any order or even simultaneously. If simultaneously, the multiple therapeutic agents may be provided in a single, unified form, or in multiple forms (by way of example only, either as a single pill or as two separate pills). One of the therapeutic agents may be given in multiple doses, or both may be given as multiple doses. If not simultaneous, the timing between the multiple doses may vary from more than zero weeks to less than four weeks. In addition, the combination methods, compositions and formulations are not to be limited to the use of only two agents.

EXAMPLES

[0284] The present invention is further illustrated by the following examples, which should not be construed as limiting in any way.

Example 1—Effects of Aromatic-Cationic Peptides on C-Reactive Protein in a Dog Model of Heart Failure

[0285] In this Example, the effect of the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on reducing C-reactive protein levels in dogs with coronary microembolization-induced heart failure was investigated.

Methods

[0286] Heart failure was induced in dogs via multiple sequential intracoronary microembolizations as described in Sabbah, et al., *Am J Physiol.* (1991) 260:H1379-84, herein incorporated by reference in its entirety.

[0287] Half the dogs were subsequently treated with the mitochondrial peptide; the other half were treated with drug vehicle and served as controls. Peptide treatment was started upon induction of heart failure (HF), defined as left ventricular ejection fraction of approximately 30%. Peripheral venous blood samples were obtained from all dogs at the following time points: 1) when the dogs were normal (baseline, prior to the induction of heart failure); 2) when the dogs were induced into heart failure; 3) at 6 weeks after initiating therapy with the mitochondrial peptide or vehicle (HF+peptide, or HF+vehicle) both administered once daily as subcutaneous injections; and 4) at 12 weeks after initiating therapy with the mitochondrial peptide or vehicle. The daily dose of the peptide was 0.5 mg/kg/day administered intravenously. Blood samples were drawn on EDTA anticoagulant and were centrifuged at 2,500 RPM and the plasma extracted, aliquoted in 1 ml volumes into cryostat tubes and stored at -70° C. until assayed. Once the follow-up was completed in all dogs, the plasma samples were thawed to room temperature and CRP was assayed using a canine specific ELISA kit (ALPCO: Cat #41-CRPCA-E01). The calorimetric method was used to assess the amount of CRP in plasma. CRP concentration was expressed as g/ml.

Results

[0288] The results are shown in the Tables 10 and 11 below and in FIG. 1. In the Tables, “BL” is baseline; “HF” is heart failure prior to initiating therapy; “6 Wk” is 6 weeks after initiating therapy; “12 Wk” is 12 weeks after initiating therapy and marked the end of the study; and “SEM” is standard error of the mean. Values in the table are g/ml CRP. As shown in the tables, in dogs treated with vehicle, plasma CRP increased in HF compared to baseline and tended to increase further at 6 week and 12 weeks after initiating subcutaneous injection with vehicle. In dogs treated with the mitochondrial peptide, CRP also increased when the dogs were induced into HF but treatment with the mitochondrial peptide reduced CRP at 6 weeks and reduced or normalized its concentration in plasma at 12 week.

TABLE 10

C-reactive protein concentration in control animals				
Dog #	BL	Vehicle (Control)		
		Pre	6 Wk	12 Wk
1	1.04	6.27	5.27	5.6
2	1.11	5.47	7.13	10.76
3	0.63	10.94	11.72	16.41

TABLE 10-continued

C-reactive protein concentration in control animals				
Dog #	BL	Vehicle (Control)		
		Pre	6 Wk	12 Wk
Mean	0.93	7.56	8.04	10.92
SEM	0.15	1.71	1.92	3.13

TABLE 11

C-reactive protein concentration in animals treated with peptide				
Dog #	BL	Mitochondrial peptide D-Arg-2',6'-Dmt-Lys-Phe-NH ₂ (SEQ ID NO: 3)		
		Pre	6 Wk	12 Wk
1	1.02	5.07	1.12	0.51
2	2.54	9.71	3.23	2.69
3	1.89	4.06	2.69	1.07
Mean	1.82	6.28	2.35	1.42
SEM	0.44	1.74	0.63	0.65

[0289] As shown in FIG. 1, plasma CRP levels as determined using a high-sensitivity assay increased about 3 fold in the heart failure subjects. Treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) for six weeks reduced CRP levels, while treatment for 12 weeks reduced CRP levels in the treated heart failure subjects to normal or near normal levels.

[0290] The results show that D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful for lowering CRP levels in heart failure subjects, thereby reducing the risk of a future heart failure event or recurrence, reducing the severity of future heart failure, and/or preventing heart failure in an undiagnosed subjects. As such, the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful for reducing CRP levels, thereby preventing or treating heart failure in mammalian subjects.

Example 2—Treatment or Prevention of Heart Failure in Human

[0291] Forty human subjects diagnosed with heart failure and having elevated CRP levels in the range of 3-10 mg/L, as determined by either a hsCRP or cCRP assay, will be randomly divided into four groups and will be treated with vehicle (control subject group), or aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) at a dosage of 1 mg/kg/d, 3 mg/kg/d or 5 mg/kg/d. Peptide or vehicle will be administered subcutaneously in three separate doses per day, every 8 hours. CRP levels will be evaluated during the course of 12-week treatment.

Results

[0292] Treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is predicted to reduce CRP levels in treated subjects. It is anticipated that treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) will reduce CRP levels to normal range (e.g., less than 1 mg/L), and/or will reduce the level of CRP from a high risk level (e.g., 3-10 mg/L) to an intermediate risk level (e.g., 1-3 mg/L), thereby reducing the risk of a future heart failure event or recurrence, reducing the severity of heart failure, and/or preventing heart failure

in an undiagnosed subjects or subjects at risk of heart failure. As such, results are anticipated to show that aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful in methods for lowering CRP levels in humans, and for preventing or treating heart failure in human subjects.

Example 3—Effects of Aromatic-Cationic Peptides on ROS in a Dog Model of Heart Failure

[0293] In this example, the effect of the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on ROS levels in dogs with coronary microembolization-induced heart failure was investigated.

Methods

[0294] Heart failure was induced as described in Example 1. Ten dogs were used in the experiment to determine the effect of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on ROS levels. Half the dogs were treated with the peptide; the other half were treated with drug vehicle and served as controls. Peptide treatment was started upon induction of heart failure (HF), defined as left ventricular ejection fraction of approximately 30%. The daily dose of the peptide was 0.5 mg/kg/day administered intravenously. Blood samples were drawn on EDTA anticoagulant and were centrifuged at 2,500 RPM and the plasma extracted, aliquoted in 1 ml volumes into crystat tubes and stored at -70° C. until assayed. Once the follow-up was completed in all dogs, the plasma samples were thawed to room temperature and ROS was assayed. To assess ROS levels in the plasma, luminol (5-amino-2,3 dihydro-1,4 phtalazindione) was added to the samples. ROS levels were then assessed by chemiluminescence activity.

Results

[0295] The results are shown in the Tables 12 and 13 below and in FIG. 2. In the tables, "BL" is baseline; "PRE" is heart failure prior to initiating therapy; "6 Wk" is 6 weeks after initiating therapy; "12 Wk" is 12 weeks after initiating therapy and marked the end of the study; and "SEM" is standard error of the mean. "ND" indicates that no plasma was assayed. Values in the table are RLU×10⁵/ml ROS, wherein RLU stands for relative light units. As shown in the tables, in dogs treated with vehicle, plasma ROS increased in HF compared to baseline and tended to increase further at 6 weeks and 12 weeks after initiating subcutaneous injection with vehicle. In dogs treated with the mitochondrial peptide, ROS also increased when the dogs were induced into HF but treatment with the mitochondrial peptide reduced ROS at 6 weeks and reduced or normalized its concentration in plasma at 12 week.

TABLE 12

Reactive oxygen species concentration in control animals				
Dog #	Vehicle (Control)			
	BL	PRE	6 Wk	12 Wk
1	4.6	ND	ND	20.1
2	5.5	14.4	14.7	16.8
3	3.8	22.5	26.0	27.3
4	5.4	18.4	19.5	21.5

TABLE 12-continued

Reactive oxygen species concentration in control animals				
Dog #	Vehicle (Control)			
	BL	PRE	6 Wk	12 Wk
5	5.6	29.0	31.3	33.2
Mean	5.0	21.1	22.9	23.8
SEM	0.3	3.1	3.6	2.9

TABLE 13

Reactive oxygen species concentration in animals treated with peptide				
Dog #	Mitochondrial peptide D-Arg-2',6'-Dmt-Lys-Phe-NH ₂			
	BL	PRE	6 Wk	12 Wk
1	6.8	ND	ND	9.0
2	5.7	14.6	15.1	12.4
3	4.7	26.0	23.3	10.5
4	5.5	23.6	18.4	12.4
5	3.9	18.5	16.6	8.3
Mean	5.3	20.7	18.4	10.5
SEM	0.5	2.6	1.8	0.8

[0296] As shown in FIG. 2, plasma ROS levels increased over 4 fold in the heart failure subjects. Treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) for six weeks reduced ROS levels, while treatment for 12 weeks reduced ROS levels in the treated heart failure subjects to near normal levels.

[0297] The results show that D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful for lowering ROS levels in heart failure subjects, thereby reducing the risk of a future heart failure event or recurrence, reducing the severity of future heart failure, and/or preventing heart failure in an undiagnosed subjects. As such, the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ TD NO: 3) is useful for reducing ROS levels, and preventing or treating heart failure in mammalian subjects.

Example 4—Effects of Aromatic-Cationic Peptides on IL-6 in a Dog Model of Heart Failure

[0298] In this example, the effect the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on interleukin-6 (IL-6) levels in dogs with coronary microembolization-induced heart failure was investigated.

Methods

[0299] The methods of Example 1 were used to induce heart failure, administer peptide or vehicle and collect plasma. Fourteen dogs were used in this example. Half the dogs were treated with the mitochondrial peptide; the other half were treated with drug vehicle and served as controls. Peptide treatment was started in both groups upon induction of heart failure (HF), defined as left ventricular ejection fraction of approximately 30%.

[0300] IL-6 was quantified in EDTA-plasma using the double antibody sandwich ELISA technique. Briefly, IL-6 (Canine IL-6 Duo Set, Cat #DY1609) was purchased from R&D Systems. The assay was performed based on instructions provided by the supplier with minor modifications. The

amount of EDTA-plasma incubated with the antibody plate was 150 µl and incubation time was 18 hours at 4° C. Using standard curves and software, the concentration of IL-6 was expressed as pg/ml.

Results

[0301] The results are shown in the Tables 14 and 15 below and in FIG. 3. In the tables, “BL” is baseline; “PRE” is heart failure prior to initiating therapy; “6 Wk” is 6 weeks after initiating therapy; “12 Wk” is 12 weeks after initiating therapy and marked the end of the study; and “SEM” is standard error of the mean. Values in the table are pg/ml IL-6. As shown in the tables, in dogs treated with vehicle, plasma IL-6 increased in HF compared to baseline and tended to increase further at 6 week and 12 weeks after initiating subcutaneous injection with vehicle. In dogs treated with the mitochondrial peptide, IL-6 also increased when the dogs were induced into HF but treatment with the mitochondrial peptide reduced IL-6 at 6 weeks and reduced or normalized its concentration in plasma at 12 week.

TABLE 14

IL-6 concentration in control animals				
Dog #	Vehicle (Control)			
	BL	PRE	6 Wk	12 Wk
1	10.3	28.4	30.1	31.0
2	9.0	17.6	22.1	24.7
3	11.6	30.6	34.3	34.3
4	7.0	18.5	20.3	21.4
5	8.2	21.3	22.6	24.7
6	11.8	28.5	29.2	30.2
7	5.8	29.2	29.8	33.8
Mean	9.1	24.9	26.9	28.6
SEM	0.9	2.1	2.0	1.9

TABLE 15

IL-6 concentration in animals treated with peptide				
Mitochondrial peptide D-Arg-2',6'-Dmt-Lys-Phe-NH ₂ (SEQ ID NO: 3)				
Dog #	BL	PRE	6 Wk	12 Wk
1	10.3	31.2	19.6	12.5
2	12.5	29.4	20.5	11.9
3	13.0	30.4	19.6	11.4
4	9.1	29.3	18.8	11.7
5	5.8	18.5	23.4	11.3
6	7.0	29.2	24.3	13.6
7	12.6	18.0	19.5	11.5
Mean	10.1	26.6	20.8	12.0
SEM	1.1	2.2	0.8	0.3

[0302] As shown in FIG. 3, plasma IL-6 levels, as determined using a high-sensitivity assay, increased about 3 fold in the heart failure subjects. Treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) for six weeks reduced IL-6 levels, while treatment for 12 weeks reduced IL-6 levels in the treated heart failure subjects to near normal levels.

[0303] The results show that D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful for lowering IL-6 levels in heart failure subjects, thereby reducing the risk of a future heart failure event or recurrence, reducing the severity of

future heart failure, and/or preventing heart failure in an undiagnosed subjects. As such, the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ TD NO: 3) is useful for reducing TL-6 levels, and preventing or treating heart failure in mammalian subjects.

Example 5—Effects of Aromatic-Cationic Peptides on TNF-Alpha in a Dog Model of Heart Failure

[0304] In this example, the effect of the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on tumor necrosis factor alpha levels in dogs with coronary microembolization-induced heart failure was investigated.

Methods

[0305] The methods of Example 1 was used induce heart failure, administer peptide or vehicle, and to collect plasma samples for analysis. Fourteen dogs were used in this example. Half the dogs were treated with the peptide; the other half were treated with drug vehicle and served as controls. Peptide treatment was started in both groups upon induction of heart failure (HF), defined as left ventricular ejection fraction of approximately 30%.

[0306] TNF-alpha was quantified in EDTA-plasma using a double antibody sandwich ELISA. Briefly, TNF-alpha (Canine TNF α Duo Sett, Cat #DY1507) was purchased from R&D Systems. The assay was performed based on instructions provided by the supplier with minor modifications. The amount of EDTA-plasma incubated with antibody plate was 150 µl and incubation time was 18 hours at 4° C. Using standard curves and software, the concentration of TNF α was expressed as pg/ml.

Results

[0307] The results are shown in the Tables 16 and 17 below and in FIG. 4. In the tables, “BL” is baseline; “PRE” is heart failure prior to initiating therapy; “6 Wk” is 6 weeks after initiating therapy; “12 Wk” is 12 weeks after initiating therapy and marked the end of the study; and “SEM” is standard error of the mean. Values in the table are pg/ml TNF-alpha. As shown in the tables, in dogs treated with vehicle, plasma TNF-alpha increased in HF compared to baseline and tended to increase further at 6 week and 12 weeks after initiating subcutaneous injection with vehicle. In dogs treated with the peptide, TNF-alpha also increased when the dogs were induced into HF but treatment with the mitochondrial peptide reduced TNF-alpha at 6 weeks and reduced or normalized its concentration in plasma at 12 week.

TABLE 16

TNF- α concentration in control animals				
Dog #	Vehicle (Control)			
	BL	PRE	6 Wk	12 Wk
1	0.99	3.78	4.37	4.80
2	0.84	3.86	4.55	3.93
3	1.72	3.68	3.95	3.68
4	1.15	3.47	3.86	2.94
5	1.85	3.54	2.53	3.76
6	1.30	4.74	3.73	4.00

TABLE 16-continued

TNF- α concentration in control animals				
Dog #	Vehicle (Control)			
	BL	PRE	6 Wk	12 Wk
7	1.17	3.91	4.48	4.57
Mean	1.29	3.85	3.92	3.95
SEM	0.14	0.16	0.26	0.23

TABLE 17

TNF- α concentration in animals treated with peptide				
Dog #	Mitochondrial peptide D-Arg-2',6'-Dmt-Lys-Phe-NH ₂ (SEQ ID NO: 3)			
	BL	PRE	6 Wk	12 Wk
1	1.30	4.59	3.72	1.69
2	1.11	4.31	3.12	1.53
3	1.54	4.42	2.60	1.35
4	0.90	4.26	3.27	1.14
5	0.97	3.96	3.08	1.42
6	1.02	4.26	3.50	1.18
7	1.85	4.69	3.16	1.75
Mean	1.24	4.36	3.21	1.44
SEM	0.13	0.09	0.13	0.09

[0308] As shown in FIG. 4, plasma TNF-alpha levels increased about 4 fold in the heart failure subjects. Treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) for six weeks reduced TNF-alpha levels, while treatment for 12 weeks reduced TNF-alpha levels in the treated heart failure subjects to near normal levels.

[0309] The results show that D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful for lowering TNF-alpha levels in heart failure subjects, thereby reducing the risk of a future heart failure event or recurrence, reducing the severity of future heart failure, and/or preventing heart failure in an undiagnosed subjects. As such, the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful for reducing TNF-alpha levels, and preventing or treating heart failure in mammalian subjects.

Example 6—D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) Administered Post-Myocardial Infarction Improves LV Function

[0310] This study will demonstrate that chronic therapy with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), begun at 2 hours post induction of heart failure by a transmural, non reperfused infarct in the rat, can improve outcome. Since D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) treatment will start at two hours after permanent coronary occlusion, any benefit will be independent of phenomena such as no-reflow reduction. Two hours after coronary occlusion, all or nearly all cells destined to die due to ischemic necrosis have died in the rat model. This study measures the ability of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) to reduce LV volumes, improve fractional shortening and ejection fraction, reduce infarct expansion, improve survival, improve hemodynamics, and reduce lung volumes.

Methods

[0311] Rats are anesthetized, ventilated, and a thoracotomy performed in the left 4th intercostal space. Tempera-

ture is maintained at 36° C. by placing the rats on a heating pad during the procedure. The pericardium is excised and the proximal left coronary artery is isolated and permanently occluded with a suture. Coronary artery occlusion is confirmed by cyanosis and akinesis of the anterior wall of the ventricle. The chest is closed, air evacuated, and the rats are allowed to recover. Analgesia is administered per the veterinarian. An echocardiogram is obtained at approximately 15 minutes post coronary artery occlusion. At 2 hours rats are randomized to receive chronic daily D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) (delivered subcutaneously by an Alzet Osmotic Pump—3 mg/kg/day) or water. The Osmotic Pump delivers approximately 0.15 μ l/hr for 6 weeks (model 2006; 200 μ l). The Alzet pump is implanted subcutaneously between the shoulder blades while the rat is still anesthetized. After 6 weeks the rats are re-anesthetized, weighed, and a second echocardiogram is obtained under anesthesia. Cut downs are performed to isolate the carotid artery and jugular vein. Heart rate and blood pressure are measured. A Millar catheter is inserted into the left ventricle and LV systolic pressure, LV end diastolic pressure, +dP/dt, and -dP/dt are measured. A left ventriculogram is performed using IV fluoroscopic contrast in order to determine LV stroke volume and ejection fraction. Under deep anesthesia, the heart is excised, weighed, and pressure fixed at 11 mmHg with formalin. The lungs are also excised and weighed. Postmortem LV volume is measured by filling the LV cavity with fluid and measuring the total fluid. The hearts are sliced into four transverse sections and histologic slides are prepared and stained with hematoxylin and eosin and with picrosirius red, which stains collagen. Quantitative histologic analysis includes: total circumference, scar circumference, non-infarcted wall circumference, total LV area, total LV cavity area, LV wall thickness (at several points), non-infarcted wall thickness; myocardial infarct expansion index.

Results

LV Fractional Shortening by Echocardiography

[0312] It is anticipated that the left ventricular fractional shortening (LVFS) will improve in the D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) treated group, as compared to the untreated group.

LV Stroke Volume and Ejection Fraction by LV Ventriculography

[0313] It is anticipated that there will be a higher LV stroke volume and LV ejection fraction in the D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) treated group, as compared to the untreated group.

Hemodynamics

[0314] No significant differences are anticipated in heart rate, systolic and diastolic blood pressure between the two groups at 6 weeks after treatment.

Post-Mortem LV Volumes

[0315] It is anticipated that there will be a lower post-mortem LV volume in the D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3)-treated group as compared to the control group.

Scar Circumference, Scar Thickness, and Expansion Index

[0316] It is anticipated that the LV non-scar circumference will be longer in the D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) treated group as compared to the water group. Additionally, the scar circumference is anticipated to be smaller in the D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) treated group as compared to the water group. It is anticipated that the scar circumference, expressed as percentage of total LV circumference, will be smaller in the D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) group as compared to the water. The scar thickness, septum thickness and expansion index expressed as: [LV cavity area/Total LV area×Septum thickness/Scar thickness], is anticipated to be comparable between the two groups.

Lung Weights (a Measure of Fluid Overload)

[0317] The lung dry and wet weight is measured, and the ratio of dry/wet is anticipated to be similar in between the two groups.

[0318] These results will show that aromatic-cationic peptides of the present technology, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), or a pharmaceutically acceptable salt thereof, such as acetate or trifluoroacetate salt, are useful in the prevention of LV remodeling and improvement of LV function. In particular, these results will show that aromatic-cationic peptides of the present invention, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), or a pharmaceutically acceptable salt thereof, such as acetate or trifluoroacetate salt, are useful in methods comprising administration of the peptide to subjects in need of decreased left ventricular remodeling and improved LV function.

Example 7—Effects of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on Post-Infarction Remodeling and Cardiac Function in a Rodent Model of Heart Failure

[0319] In this study, D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is tested to see if it will improve cardiac function and result in beneficial mitochondrial gene expression in a post-infarct model of heart failure.

Methods

[0320] Rats will undergo the permanent coronary artery ligation, as described in Example 6. The rats will be split into two groups and treated for six weeks with either 200-300 ng/ml of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or 0.9% NaCl (saline) continuously through mini-osmotic pumps, which are implanted into each animal.

[0321] After the six week period, LV function is assessed with echocardiography. Additionally, the hearts are excised and the heart tissue is analyzed for LV chamber volume using tetrazolium salt staining. Heart tissue in the border zone and remote areas around the infarct are also harvested and undergo gene array analysis to determine the expression levels of genes involved in mitochondrial metabolism.

Results

[0322] It is anticipated that chronic treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) will reduce LV dilation in a post-infarction model of heart failure.

[0323] These results will show that aromatic-cationic peptides, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), or a pharmaceutically acceptable salt thereof, such as acetate or trifluoroacetate salt are useful in the treatment of LV remodeling and heart failure. Accordingly, the peptides are useful in methods comprising administering aromatic-cationic peptides to a subject suffering from heart failure or post myocardial infarct.

Example 8—Effects of Aromatic-Cationic Peptides on NT-Pro BNP Levels in a Dog Model of Heart Failure

[0324] In this Example, the effect of the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on reducing NT-pro BNP levels in dogs with coronary microembolization-induced heart failure was investigated.

Methods

[0325] Heart failure was induced in dogs via multiple sequential intracoronary microembolizations as described in Sabbah, et al., *Am J Physiol.* (1991) 260:H1379-84, herein incorporated by reference in its entirety and summarized in Example 1.

[0326] Heart failure was induced as described in Example 1. Ten dogs were used in the experiment to determine the effect of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on NT-pro BNP levels. Half the dogs were treated with the peptide; the other half were treated with drug vehicle and served as controls. Peptide treatment was started upon induction of heart failure (HF), defined as left ventricular ejection fraction of approximately 30%. The daily dose of the peptide was 0.5 mg/kg/day administered intravenously. Blood samples were drawn on EDTA anticoagulant and were centrifuged at 2,500 RPM and the plasma extracted, aliquoted in 1 ml volumes into crystat tubes and stored at -70° C. until assayed. Once the follow-up was completed in all dogs, the plasma samples were thawed to room temperature and NT-pro BNP was assayed.

[0327] Nt-pro BNP (pg/ml) was determined in EDTA-plasma on the principle of the double antibody sandwich Enzyme-linked immunosorbent assay (ELISA). The assay was performed based upon the instructions came along with the assay kit (NT-pro BNP, Kamiya Biomedical Company, Cat #KT-23770). Using standard curves with the help of Software (MasterPlex-2010), the concentration of each biomarker was determined.

Results

[0328] The results are shown in the Tables 18 and 19 below and in FIG. 5. In the tables, normal is baseline before heart failure; "HF-Pre" is heart failure prior to initiating therapy; "6 Wk" is 6 weeks after initiating therapy; "12 Wk" is 12 weeks after initiating therapy and marked the end of the study; "AVG" is the average; "SD" is standard deviation; "SEM" is standard error of the mean; and "NS" is not significant. Values in the table are pg/ml NT-pro BNP. As shown in the tables, in dogs treated with vehicle, plasma NT-pro BNP, on average, increased in HF compared to baseline and tended to increase further at 6 week and 12 weeks after initiating subcutaneous injection with vehicle. In dogs treated with the mitochondrial peptide, NT-pro BNP also increased when the dogs were induced into HF but

treatment with the mitochondrial peptide reduced NT-pro BNP at 6 weeks and reduced or normalized its concentration in plasma at 12 week.

TABLE 18

Nt-pro BNP levels (pg/ml) in Plasma of CHF Dogs in control animals				
Dog #	Vehicle (Control)			
	Normal	HF-Pre	6 Wk	12 Wk
1	324	1814	1492	1363
2	224	1418	1588	1568
3	211	997	1363	1372
4	267	1060	1464	1588
5	244	950	871	930
6	294	991	1104	1082
7	271	911	712	850
AVG	262	1163	1228	1250
SD	39	333	337	298
SEM	15	126	127	113
P-value vs. Normal		<.05	<.05	<.05
P-value vs. HF-Pre			NS	NS

TABLE 19

C-reactive protein concentration in animals treated with peptide				
Mitochondrial peptide D-Arg-2',6'-Dmt-Lys-Phe-NH ₂ (SEQ ID NO: 3)				
Dog #	BL	Pre	6Wk	12 Wk
1	289	1018	695	417
2	265	1090	593	393
3	346	941	654	423
4	257	1512	695	299
5	203	1223	559	422
6	271	1078	796	234
7	320	1034	680	294
AVG	278	1128	667	354
SD	46	190	77	77
SEM	17	72	29	29
P-value vs. Normal		<.05	<.05	NS
P-value vs. HF-Pre			<.05	<.05

[0329] As shown in FIG. 5, plasma NT-pro BNP levels as determined using double antibody ELISA increased about 4 fold in the heart failure subjects. Treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) for six weeks reduced NT-pro BNP levels, while treatment for 12 weeks reduced NT-pro BNP levels in the treated heart failure subjects to normal or near normal levels.

[0330] The results show that D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) lowers NT-pro BNP levels in heart failure subjects. The decrease in NT-pro BNP correlates to a decrease in BNP, since both are release in equimolar concentration after the cleavage of proBNP, thereby indicating a decrease in the stretching of cardiomyocytes. As such, the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful for preventing or treating LV remodeling in mammalian subjects.

Example 9—Effects of Aromatic-Cationic Peptides on Mitochondria ATP-Sensitive Potassium Channel (mK ATP) in a Dog Model of Heart Failure

[0331] In this Example, the effect of the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on

increasing mK ATP activity in dogs with coronary microembolization-induced heart failure was investigated.

Methods

[0332] Heart failure was induced in dogs via multiple sequential intracoronary microembolizations as described in Sabbah, et al., *Am J Physiol.* (1991) 260:H1379-84, herein incorporated by reference in its entirety.

[0333] Heart failure was induced as described in Example 1. Ten dogs were used in the experiment to determine the effect of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on mK ATP activity. Half the dogs were treated with the peptide (n=5); the other half were treated with drug vehicle and served as controls (n=5). Peptide treatment was started upon induction of heart failure (HF), defined as left ventricular ejection fraction of approximately 30%. The daily dose of the peptide was 0.5 mg/kg/day administered intravenously for three months. Subcutaneous daily injections of saline were administered to the controls. Left ventricular tissue was harvested at the end of the three months of treatment. Mitochondria were isolated from the tissue. MK ATP activation was measured using the thallium-sensitive fluorophore assay kit and quantified in relative fluorescence units (RFU) per mg protein. Mitochondrial ATP to ADP ratio was measured using the bioluminescent ApoSENSOR™ assay kit (Enzo Life Sciences, Farmingdale, NY).

Results

[0334] Treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) resulted in a significant increase in the ATP/ADP (0.38±0.04 vs. 1.16±0.15, p<0.05). Treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) also resulted in a significant increase in activation of mK ATP (1372±112 vs. 2775±254, p<0.05).

[0335] The results show that D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) increases mK ATP activity in heart failure subjects, thereby reducing the risk of a future heart failure event or recurrence, reducing the severity of future heart failure, and/or preventing heart failure in an undiagnosed subjects. As such, the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful for increasing mK ATP activity, and preventing or treating heart failure in mammalian subjects.

Example 10—Effects of Aromatic-Cationic Peptides on Cardiac Troponin I Levels in a Dog Model of Heart Failure

[0336] In this example, the effect of the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on cardiac troponin I levels in dogs with coronary microembolization-induced heart failure is investigated.

Methods

[0337] Heart failure is induced as described in Example 1. Ten dogs are used in the experiment to determine the effect of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on cardiac troponin I levels. Half the dogs are treated with the peptide; the other half are treated with drug vehicle and served as controls. Peptide treatment is started upon induction of heart failure (HF), defined as left ventricular ejection fraction of approximately 30%. The daily dose of the peptide was 0.5 mg/kg/day is administered intravenously. Blood samples are drawn on EDTA anticoagulant and are centri-

fuged at 2,500 RPM and the plasma extracted, aliquoted in 1 ml volumes into crystat tubes and stored at -70° C. until assayed. Cardiac troponin I levels are measured for a baseline, at six weeks after treatment, and at twelve weeks after treatment. Once the assay is completed in all dogs, the plasma samples are thawed to room temperature and cardiac troponin I is assayed. Baseline, six weeks treatment, and twelve weeks treatment are compared to normal controls.

Results

[0338] It is anticipated that treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) will reduce cardiac troponin I levels after six weeks of treatment as compared to untreated and that after 12 weeks of treatment it is anticipated that cardiac troponin I levels will be near normal levels.

[0339] The results will show that D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful for lowering cardiac troponin I levels in heart failure subjects, thereby reducing the risk of a future heart failure event or recurrence, reducing the severity of future heart failure, and/or preventing heart failure in an undiagnosed subjects. As such, the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) is useful for reducing cardiac troponin I levels, and preventing or treating heart failure in mammalian subjects.

Example 11—Effects of Aromatic-Cationic Peptides on Cardiolipin in a Dog Model of Heart Failure

[0340] In this Example, the effect of aromatic-cationic peptide such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on cardiolipin levels in dogs with coronary microembolization-induced heart failure was investigated.

Methods

[0341] Heart failure was induced in dogs via multiple sequential intracoronary microembolizations as described in Sabbah, et al., *Am J Physiol.* (1991) 260: H1379-84, herein incorporated by reference in its entirety. Half the dogs were subsequently treated with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3); the other half were treated with drug vehicle and served as controls. Peptide treatment was started upon induction of heart failure (HF), defined as left ventricular ejection fraction of approximately 30%. A daily dose of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) (0.5 mg/kg/day) was administered intravenously.

[0342] At the end of the treatment phase (12 weeks), dogs in both the vehicle and treatment groups were sacrificed and a sample of heart muscle from the left ventricle was removed, washed with saline, and immediately frozen and stored at -80° C.

[0343] For cardiolipin analysis, lipids were extracted from the heart tissue sample with a chloroform/methanol solution (Bligh Dyer extraction). Individual lipid extracts were reconstituted with chloroform: methanol (1:1), flushed with N₂, and then stored at -20° C. before analysis via electrospray ionization mass spectroscopy using a triple-quadruple mass spectrometer equipped with an automated nanospray apparatus. Enhanced multidimensional mass spectrometry-based shotgun lipidomics for cardiolipin was performed as described by Han, et al. (2006) *J Lipid Res* 47(4): 864-879.

Results

[0344] The 18:2 cardiolipin species was reduced in untreated HF dogs (HF-CON) ($p<0.05$) as compared to normal cardiac tissue from normal dogs (NL). FIG. 1. However, HF dogs treated with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) (HF-AP) had levels of 18:2 cardiolipin that were similar to NL dogs and greater than HF-CON ($p<0.05$). FIG. 6.

[0345] The 18:2 cardiolipin species is reduced in HF. The reduction of 18:2 cardiolipin leads to poor oxidative phosphorylation and subsequent LV dysfunction. Chronic treatment with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) normalized 18:2 cardiolipin, which leads to improved LV function and rate of mitochondrial ATP synthesis.

[0346] These results show that aromatic-cationic peptides of the present invention, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), or a pharmaceutically acceptable salt thereof, such as acetate or trifluoroacetate salt, are useful in the prevention and treatment of diseases and conditions associated with aberrant cardiolipin levels. In particular, these results show that aromatic-cationic peptides of the present invention, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), or a pharmaceutically acceptable salt thereof, such as acetate or trifluoroacetate salt, are useful in increasing cardiolipin levels and for treating heart failure and related conditions.

Example 12—Effects of Aromatic-Cationic Peptides on MLCL AT1, ALCAT1, and Taz1 Expression in a Dog Model of Heart Failure

[0347] In this Example, the effect of the aromatic-cationic peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) on cardiolipin remodeling enzymes, MLCL AT1, ALCAT1, and Taz1 in dogs with coronary microembolization-induced heart failure was investigated.

Methods

[0348] Heart failure was induced in dogs via multiple sequential intracoronary microembolizations as described in Sabbah, et al., *Am J Physiol.* (1991) 260: H1379-84, herein incorporated by reference in its entirety.

[0349] Twelve dogs were subject to coronary microembolization-induced heart failure (LV ejection fraction ~30%) as described in Example 1. Subjects were randomized into D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3)-treated and control groups for a three-month trial. Subjects received 3 months of therapy with subcutaneous injections of D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) (0.5 mg/kg once daily, n=6) or saline (Untreated-HF Control, n=6). RNA was prepared from LV tissue of all 12 dogs at the end of the treatment phase and from the LV of 6 normal (NL) dogs for comparison. mRNA levels of Taz1, MLCL AT1 and ALCAT1 were determined by real-time PCR. Changes in mRNA expression were expressed as fold changes using the CT Method with normalization to glyceraldehyde 1,3 diphosphate dehydrogenase (GAPDH) as internal control.

Results

[0350] Compared to normal level (NL), mRNA levels of Taz1 in untreated HF dogs decreased 2.25-fold (FIG. 7A) while mRNA of MLCL AT1 and ALCAT1 increased 2.60-fold and 3.56-fold, respectively. FIGS. 7B-C. Treatment

with D-Arg-2'6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) attenuated the decrease of Taz1 1.23-fold and reduced the increase in MLCL AT1 and ALCAT1 1.18-fold and 1.54-fold, respectively. FIGS. 7A-C.

[0351] HF is associated with deregulation of cardiolipin remodeling enzymes that can lead to pathologic remodeling of cardiolipin and to structural and functional mitochondrial abnormalities. Chronic therapy with D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) partially reverses these maladaptations thus allowing for resumption of physiologic post-biosynthesis remodeling of cardiolipin.

[0352] These results show that aromatic-cationic peptides of the present invention, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), or a pharmaceutically acceptable salt thereof, such as acetate or trifluoroacetate salt, are useful in the prevention and treatment of diseases and conditions associated with aberrant cardiolipin remodeling enzyme levels, e.g., MLCL AT1, ALCAT 1, and Taz1. In particular, these results show that aromatic-cationic peptides of the present invention, such as D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3), or a pharmaceutically acceptable salt thereof, such as acetate or trifluoroacetate salt, are useful in methods for normalization of cardiolipin remodeling enzyme levels, e.g., decreasing MLCL AT 1, ALCAT 1, and increasing Taz1 expression levels, and for treating heart failure and related conditions.

EQUIVALENTS

[0353] The present invention is not to be limited in terms of the particular embodiments described in this application, which are intended as single illustrations of individual aspects of the invention. Many modifications and variations of this invention can be made without departing from its spirit and scope, as will be apparent to those skilled in the art. Functionally equivalent methods and apparatuses within the scope of the invention, in addition to those enumerated herein, will be apparent to those skilled in the art from the foregoing descriptions. Such modifications and variations are intended to fall within the scope of the appended claims.

The present invention is to be limited only by the terms of the appended claims, along with the full scope of equivalents to which such claims are entitled. It is to be understood that this invention is not limited to particular methods, reagents, compounds compositions or biological systems, which can, 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.

[0354] In addition, where features or aspects of the disclosure are described in terms of Markush groups, those skilled in the art will recognize that the disclosure is also thereby described in terms of any individual member or subgroup of members of the Markush group.

[0355] As will be understood by one skilled in the art, for any and all purposes, particularly in terms of providing a written description, all ranges disclosed herein also encompass any and all possible sub-ranges and combinations of sub-ranges thereof. Any listed range can be easily recognized as sufficiently describing and enabling the same range being broken down into at least equal halves, thirds, quarters, fifths, tenths, etc. As a non-limiting example, each range discussed herein can be readily broken down into a lower third, middle third and upper third, etc. As will also be understood by one skilled in the art all language such as "up to," "at least," "greater than," "less than," and the like, include the number recited and refer to ranges which can be subsequently broken down into sub-ranges as discussed above. Finally, as will be understood by one skilled in the art, a range includes each individual member. Thus, for example, a group having 1-3 cells refers to groups having 1, 2, or 3 cells. Similarly, a group having 1-5 cells refers to groups having 1, 2, 3, 4, or 5 cells, and so forth.

[0356] All patents, patent applications, provisional applications, and publications referred to or cited herein are incorporated by reference in their entirety, including all figures and tables, to the extent they are not inconsistent with the explicit teachings of this specification.

[0357] Other embodiments are set forth within the following claims.

SEQUENCE LISTING

```

Sequence total quantity: 109
SEQ ID NO: 1      moltype = AA  length = 224
FEATURE          Location/Qualifiers
source           1..224
                  mol_type = protein
                  organism = Homo sapiens
SEQUENCE: 1
MEKLLCFLVL TSLSHAFGQT DMSRKAFVFP KESDTSYVSL KAPLTKPLKA FTVCLHFYTE 60
LSSTRGYSIF SYATKRQDNE IILIFWSKDIG YSFTVGGSEI LFEVPEVTVA PVHICTSWES 120
ASGIVEFWVD GKPRVRKSLK KGTVGAEAS IILGQEQQDSF GGNFEGGSQSL VGDIGNVNMMW 180
DFVLSPDEIN TIYLGGPFSP NVLNWRALKY EVQGEVFTKP QLWP                224

SEQ ID NO: 2      moltype = AA  length = 224
FEATURE          Location/Qualifiers
source           1..224
                  mol_type = protein
                  organism = Homo sapiens
SEQUENCE: 2
MEKLLCFLVL TSLSHAFGQT DMSRKAFVFP KESDTSYVSL KAPLTKPLKA FTVCLHFYTE 60
LSSTRGTVFS RMPPRDKTMR FFFIFWSKDIG YSFTVGGSEI LFEVPEVTVA PVHICTSWES 120
ASGIVEFWVD GKPRVRKSLK KGTVGAEAS IILGQEQQDSF GGNFEGGSQSL VGDIGNVNMMW 180
DFVLSPDEIN TIYLGGPFSP NVLNWRALKY EVQGEVFTKP QLWP                224

SEQ ID NO: 3      moltype = AA  length = 4
FEATURE          Location/Qualifiers

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-continued

```

source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = D-amino acid
SITE           2
               note = 2',6'-dimethyltyrosine
SITE           4
               note = Amidated residue
SEQUENCE: 3
RYFK                                     4

SEQ ID NO: 4      moltype = AA  length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 4
YRFK                                     4

SEQ ID NO: 5      moltype = AA  length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = 2',6'-dimethyltyrosine
SITE           2
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 5
YRFK                                     4

SEQ ID NO: 6      moltype = AA  length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 6
FRFK                                     4

SEQ ID NO: 7      moltype = AA  length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = 2',6'-dimethylphenylalanine
SITE           2
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 7
FRFK                                     4

SEQ ID NO: 8      moltype = AA  length = 8
FEATURE
source          1..8
               mol_type = protein
               organism = synthetic construct
SITE           8
               note = D-amino acid
SEQUENCE: 8
YRFKEHWR                                         8

SEQ ID NO: 9      moltype = AA  length = 6
FEATURE

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-continued

```

source          1..6
               mol_type = protein
               organism = synthetic construct
SITE           4
               note = D-amino acid
SEQUENCE: 9
KQYRFW                                                 6

SEQ ID NO: 10      moltype = AA  length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = 2',6'-dimethylphenylalanine
SITE           2
               note = D-amino acid
SITE           3
               note = 2',6'-dimethyltyrosine
SITE           4
               note = Amidated residue
SEQUENCE: 10
FRYK                                                 4

SEQ ID NO: 11      moltype = AA  length = 10
FEATURE
source          1..10
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           3
               note = D-amino acid
SITE           6
               note = D-amino acid
SITE           8
               note = D-amino acid
SEQUENCE: 11
AFRYKWHYGF                                              10

SEQ ID NO: 12      moltype = AA  length = 14
FEATURE
source          1..14
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           3
               note = D-amino acid
SITE           7
               note = D-amino acid
SITE           9
               note = D-amino acid
SITE           11
               note = D-amino acid
SITE           13
               note = D-amino acid
SEQUENCE: 12
RLYFKEKRWK FYRG                                         14

SEQ ID NO: 13      moltype = AA  length = 16
FEATURE
source          1..16
               mol_type = protein
               organism = synthetic construct
SITE           3
               note = D-amino acid
SITE           6
               note = D-amino acid
SITE           7
               note = D-amino acid
SITE           10
               note = D-amino acid
SITE           12
               note = D-amino acid
SITE           14

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-continued

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SEQUENCE: 13          note = D-amino acid
DRFCFRKYRY WHYFKF                           16

SEQ ID NO: 14          moltype = AA  length = 9
FEATURE
source
1..9
mol_type = protein
organism = synthetic construct
SITE
2
note = D-amino acid
SITE
5
note = D-amino acid
SITE
8
note = D-amino acid
SITE
9
note = D-amino acid
note = Amidated residue
SEQUENCE: 14
DWKYHFRGK                           9

SEQ ID NO: 15          moltype = AA  length = 15
FEATURE
source
1..15
mol_type = protein
organism = synthetic construct
SITE
1
note = D-amino acid
SITE
4
note = D-amino acid
SITE
5
note = D-amino acid
SITE
8
note = D-amino acid
SITE
12
note = D-amino acid
SITE
15
note = D-amino acid
note = Amidated residue
SEQUENCE: 15
EDKRHFFVYR YYRHF                           15

SEQ ID NO: 16          moltype = AA  length = 6
FEATURE
source
1..6
mol_type = protein
organism = synthetic construct
SITE
1
note = D-amino acid
SITE
5
note = D-amino acid
SEQUENCE: 16
HEKYFR                           6

SEQ ID NO: 17          moltype = AA  length = 11
FEATURE
source
1..11
mol_type = protein
organism = synthetic construct
SITE
1
note = D-amino acid
SITE
4
note = D-amino acid
SITE
6
note = D-amino acid
SITE
7
note = D-amino acid
SITE
8
note = D-amino acid
SITE
9
note = D-amino acid
SITE
11
note = D-amino acid
note = Amidated residue
SEQUENCE: 17
HKYFEDDHKR W                           11

SEQ ID NO: 18          moltype = AA  length = 13
FEATURE
Location/Qualifiers

```

-continued

```

source          1..13
               mol_type = protein
               organism = synthetic construct
SITE           3
               note = D-amino acid
SITE           5
               note = D-amino acid
SITE           7
               note = D-amino acid
SITE           10
              note = D-amino acid
SITE           13
               note = Amidated D-amino acid
SEQUENCE: 18
ERKYVFWHRG YRM                                13

SEQ ID NO: 19          moltype = AA length = 19
FEATURE
source          Location/Qualifiers
               1..19
               mol_type = protein
               organism = synthetic construct
SITE           5
               note = D-amino acid
SITE           10
               note = D-amino acid
SITE           11
               note = D-amino acid
SITE           15
               note = D-amino acid
SITE           18
               note = D-amino acid
SEQUENCE: 19
GAKFKERYHR RDYWHWHKD                                19

SEQ ID NO: 20          moltype = AA length = 6
FEATURE
source          Location/Qualifiers
               1..6
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           5
               note = D-amino acid
SITE           6
               note = Amidated residue
SEQUENCE: 20
GFKHRY                                6

SEQ ID NO: 21          moltype = AA length = 17
FEATURE
source          Location/Qualifiers
               1..17
               mol_type = protein
               organism = synthetic construct
SITE           3
               note = D-amino acid
SITE           7
               note = D-amino acid
SITE           11
               note = D-amino acid
SITE           14
               note = D-amino acid
SITE           17
               note = Amidated residue
SEQUENCE: 21
HYRWKFDARC YHFKYHS                                17

SEQ ID NO: 22          moltype = AA length = 6
FEATURE
source          Location/Qualifiers
               1..6
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           5
               note = D-amino acid
SITE           6

```

-continued

SEQUENCE: 22 KQYRFW	note = Amidated residue	
		6
SEQ ID NO: 23 FEATURE source	moltype = AA length = 8 Location/Qualifiers 1..8	
SITE	mol_type = protein organism = synthetic construct	
SITE	3	
SITE	note = D-amino acid	
SITE	8	
SEQUENCE: 23 KWYRNFYH	note = Amidated D-amino acid	
		8
SEQ ID NO: 24 FEATURE source	moltype = AA length = 5 Location/Qualifiers 1..5	
SITE	mol_type = protein organism = synthetic construct	
SITE	3	
SITE	note = D-amino acid	
SITE	5	
SEQUENCE: 24 MYRFR	note = Amidated residue	
		5
SEQ ID NO: 25 FEATURE source	moltype = AA length = 5 Location/Qualifiers 1..5	
SITE	mol_type = protein organism = synthetic construct	
SITE	3	
SEQUENCE: 25 MYKFR	note = D-amino acid	
		5
SEQ ID NO: 26 FEATURE source	moltype = AA length = 4 Location/Qualifiers 1..4	
SITE	mol_type = protein organism = synthetic construct	
SITE	3	
SEQUENCE: 26 FRHD	note = D-amino acid	
		4
SEQ ID NO: 27 FEATURE source	moltype = AA length = 4 Location/Qualifiers 1..4	
SITE	mol_type = protein organism = synthetic construct	
SITE	2	
SITE	note = D-amino acid	
SITE	3	
SITE	note = 2',6'-dimethyltyrosine	
SITE	4	
SEQUENCE: 27 FRYK	note = Amidated residue	
		4
SEQ ID NO: 28 FEATURE source	moltype = AA length = 7 Location/Qualifiers 1..7	
SITE	mol_type = protein organism = synthetic construct	
SITE	2	
SITE	note = D-amino acid	
SITE	6	
SEQUENCE: 28 FRKWyRH	note = D-amino acid	
		7
SEQ ID NO: 29 FEATURE	moltype = AA length = 11 Location/Qualifiers	

-continued

```

source          1..11
               mol_type = protein
               organism = synthetic construct
SITE           3
               note = D-amino acid
SITE           7
               note = D-amino acid
SITE           9
               note = D-amino acid
SITE           11
               note = Amidated residue
SEQUENCE: 29
FFYREDKRRH F                         11

SEQ ID NO: 30      moltype = AA length = 12
FEATURE
source          1..12
               mol_type = protein
               organism = synthetic construct
SITE           4
               note = D-amino acid
SITE           7
               note = D-amino acid
SITE           8
               note = D-amino acid
SITE           11
               note = D-amino acid
SEQUENCE: 30
FYKRWHKKER YT                         12

SEQ ID NO: 31      moltype = AA length = 9
FEATURE
source          1..9
               mol_type = protein
               organism = synthetic construct
SITE           5
               note = D-amino acid
SITE           8
               note = D-amino acid
SEQUENCE: 31
TGYRHFWHK                         9

SEQ ID NO: 32      moltype = AA length = 20
FEATURE
source          1..20
               mol_type = protein
               organism = synthetic construct
SITE           4
               note = D-amino acid
SITE           9
               note = D-amino acid
SITE           10
               note = D-amino acid
SITE           13
               note = D-amino acid
SITE           17
               note = D-amino acid
SITE           20
               note = Amidated residue
SEQUENCE: 32
TYRKWYEDKR HFYGVIHRYK                         20

SEQ ID NO: 33      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 33
WKYR                                         4

SEQ ID NO: 34      moltype = AA length = 8
FEATURE

```

-continued

```

source          1..8
               mol_type = protein
               organism = synthetic construct
SITE           4
               note = D-amino acid
SITE           7
               note = D-amino acid
SEQUENCE: 34
WKFDRYHK                                              8

SEQ ID NO: 35          moltype = AA  length = 12
FEATURE
source          1..12
               mol_type = protein
               organism = synthetic construct
SITE           3
               note = D-amino acid
SITE           6
               note = D-amino acid
SITE           7
               note = D-amino acid
SITE           10
               note = D-amino acid
SEQUENCE: 35
YDKYFKRFPY HK                                              12

SEQ ID NO: 36          moltype = AA  length = 5
FEATURE
source          1..5
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           5
               note = Amidated residue
SEQUENCE: 36
YRFKE                                              5

SEQ ID NO: 37          moltype = AA  length = 11
FEATURE
source          1..11
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           4
               note = D-amino acid
SITE           7
               note = D-amino acid
SITE           10
               note = D-amino acid
SEQUENCE: 37
YHPRDKRHW F                                              11

SEQ ID NO: 38          moltype = AA  length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           3
               note = D-amino acid
SEQUENCE: 38
YHGM                                              4

SEQ ID NO: 39          moltype = AA  length = 8
FEATURE
source          1..8
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           5
               note = D-amino acid
SITE           8
               note = Amidated residue
SEQUENCE: 39

```

-continued

VKHYFSYR

8

```

SEQ ID NO: 40      moltype = AA  length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1
note = 2',6'-dimethyltyrosine
SITE
2
note = D-amino acid
SITE
4
note = N6-(2-(dansyl)ethyl)lysine amide
SEQUENCE: 40
YRFK

```

4

```

SEQ ID NO: 41      moltype = AA  length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1
note = 2',6'-dimethyltyrosine
SITE
2
note = D-amino acid
SITE
4
note = N6-(2-(anthraniloyl)ethyl)lysine amide
SEQUENCE: 41
YRFK

```

4

```

SEQ ID NO: 42      moltype = AA  length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1
note = 2',6'-dimethyltyrosine
SITE
2
note = D-amino acid
SITE
4
note = N6-(dansyl)lysine amide
SEQUENCE: 42
YRFK

```

4

```

SEQ ID NO: 43      moltype = AA  length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1
note = 2',6'-dimethyltyrosine
SITE
2
note = D-amino acid
SITE
4
note = Amidated residue
SEQUENCE: 43
YRFK

```

4

```

SEQ ID NO: 44      moltype = AA  length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1
note = 3',5'-dimethyltyrosine
SITE
2
note = D-amino acid
SITE
4
note = Amidated residue
SEQUENCE: 44
YRFK

```

4

```

SEQ ID NO: 45      moltype = AA  length = 4
FEATURE
source
1..4
mol_type = protein

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-continued

```

SITE organism = synthetic construct
      2
      note = D-amino acid
SITE      4
      note = Amidated residue
SEQUENCE: 45
YRYK                                         4

SEQ ID NO: 46      moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
      mol_type = protein
      organism = synthetic construct
SITE      1
      note = 2',6'-dimethyltyrosine
SITE      2
      note = D-amino acid
SITE      4
      note = Amidated residue
SEQUENCE: 46
YRYK                                         4

SEQ ID NO: 47      moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
      mol_type = protein
      organism = synthetic construct
SITE      1
      note = 2',6'-dimethyltyrosine
SITE      2
      note = D-amino acid
SITE      3
      note = 2',6'-dimethyltyrosine
SITE      4
      note = Amidated residue
SEQUENCE: 47
YRYK                                         4

SEQ ID NO: 48      moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
      mol_type = protein
      organism = synthetic construct
SITE      1
      note = 3',5'-dimethyltyrosine
SITE      2
      note = D-amino acid
SITE      3
      note = 3',5'-dimethyltyrosine
SITE      4
      note = Amidated residue
SEQUENCE: 48
YRYR                                         4

SEQ ID NO: 49      moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
      mol_type = protein
      organism = synthetic construct
SITE      1
      note = 3',5'-dimethyltyrosine
SITE      2
      note = D-amino acid
SITE      3
      note = 3',5'-dimethyltyrosine
SITE      4
      note = Amidated residue
SEQUENCE: 49
YRYK                                         4

SEQ ID NO: 50      moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
      mol_type = protein
      organism = synthetic construct
SITE      2

```

-continued

```

SITE          note = D-amino acid
        4
        note = Amidated residue

SEQUENCE: 50
YKFR

SEQ ID NO: 51      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            2
note = D-amino acid
SITE            4
note = Amidated residue

SEQUENCE: 51
YKFK

SEQ ID NO: 52      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = 2',6'-dimethyltyrosine
SITE            2
note = D-amino acid
SITE            4
note = Amidated residue

SEQUENCE: 52
YKFR

SEQ ID NO: 53      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = 2',6'-dimethyltyrosine
SITE            2
note = D-amino acid
SITE            4
note = Amidated residue

SEQUENCE: 53
YKFK

SEQ ID NO: 54      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = 3',5'-dimethyltyrosine
SITE            2
note = D-amino acid
SITE            4
note = Amidated residue

SEQUENCE: 54
YKFR

SEQ ID NO: 55      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            2
note = D-amino acid
SITE            4
note = Amidated residue

SEQUENCE: 55
YKYK

SEQ ID NO: 56      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein

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SITE organism = synthetic construct
1
SITE note = 2',6'-dimethyltyrosine
2
SITE note = D-amino acid
3
SITE note = Amidated residue
SEQUENCE: 56
YKYK 4

SEQ ID NO: 57 moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
mol_type = protein
organism = synthetic construct
SITE 1
note = 2',6'-dimethyltyrosine
2
SITE note = D-amino acid
3
SITE note = 2',6'-dimethyltyrosine
4
SITE note = Amidated residue
SEQUENCE: 57
YKYK 4

SEQ ID NO: 58 moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
mol_type = protein
organism = synthetic construct
SITE 1
note = 3',5'-dimethyltyrosine
2
SITE note = D-amino acid
3
SITE note = 3',5'-dimethyltyrosine
4
SITE note = Amidated residue
SEQUENCE: 58
YKYK 4

SEQ ID NO: 59 moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
mol_type = protein
organism = synthetic construct
SITE 1
note = 2',6'-dimethyltyrosine
2
SITE note = D-amino acid
4
SITE note = Amidated residue
SEQUENCE: 59
YRFR 4

SEQ ID NO: 60 moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
mol_type = protein
organism = synthetic construct
SITE 1
note = 3',5'-dimethyltyrosine
2
SITE note = D-amino acid
4
SITE note = Amidated residue
SEQUENCE: 60
YRFR 4

SEQ ID NO: 61 moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
mol_type = protein
organism = synthetic construct
2

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-continued

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SITE          note = D-amino acid
        4
SEQUENCE: 61 note = Amidated residue
KYKR

SEQ ID NO: 62      moltype = AA length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = 2',6'-dimethyltyrosine
SITE            2
note = D-amino acid
SITE            3
note = 2',6'-dimethyltyrosine
SITE            4
note = Amidated residue
SEQUENCE: 62
YRYR

SEQ ID NO: 63      moltype = AA length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = 2',6'-dimethyltyrosine
SITE            2
note = D-amino acid
SITE            3
note = 2',6'-dimethyltyrosine
SITE            4
note = Amidated residue
SEQUENCE: 63
YKYR

SEQ ID NO: 64      moltype = AA length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = 3',5'-dimethyltyrosine
SITE            2
note = D-amino acid
SITE            3
note = 3',5'-dimethyltyrosine
SITE            4
note = Amidated residue
SEQUENCE: 64
YKVR

SEQ ID NO: 65      moltype = AA length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = 2'-methyltyrosine
SITE            2
note = D-amino acid
SITE            4
note = Amidated residue
SEQUENCE: 65
YRFK

SEQ ID NO: 66      moltype = AA length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = N,2',6'-trimethyltyrosine
SITE            2

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-continued

```
SITE          note = D-amino acid
        4
SEQUENCE: 66          note = Amidated residue
YRFK                                     4

SEQ ID NO: 67      moltype = AA  length = 4
FEATURE          Location/Qualifiers
source           1..4
                  mol_type = protein
                  organism = synthetic construct
SITE            1
                  note = 2'-hydroxy-6'-methyltyrosine
SITE            2
                  note = D-amino acid
SITE            4
                  note = Amidated residue
SEQUENCE: 67
YRFK                                     4

SEQ ID NO: 68      moltype = AA  length = 4
FEATURE          Location/Qualifiers
source           1..4
                  mol_type = protein
                  organism = synthetic construct
SITE            1
                  note = 2'-methyltyrosine
SITE            2
                  note = D-amino acid
SITE            4
                  note = Amidated residue
SEQUENCE: 68
YKFK                                     4

SEQ ID NO: 69      moltype = AA  length = 4
FEATURE          Location/Qualifiers
source           1..4
                  mol_type = protein
                  organism = synthetic construct
SITE            1
                  note = 2'-methyltyrosine
SITE            2
                  note = D-amino acid
SITE            4
                  note = Amidated residue
SEQUENCE: 69
YKFR                                     4

SEQ ID NO: 70      moltype = AA  length = 4
FEATURE          Location/Qualifiers
source           1..4
                  mol_type = protein
                  organism = synthetic construct
SITE            1
                  note = N,2',6'-trimethyltyrosine
SITE            2
                  note = D-amino acid
SITE            4
                  note = Amidated residue
SEQUENCE: 70
YKFK                                     4

SEQ ID NO: 71      moltype = AA  length = 4
FEATURE          Location/Qualifiers
source           1..4
                  mol_type = protein
                  organism = synthetic construct
SITE            1
                  note = N,2',6'-trimethyltyrosine
SITE            2
                  note = D-amino acid
SITE            4
                  note = Amidated residue
SEQUENCE: 71
YKFR                                     4
```

-continued

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SEQ ID NO: 72      moltype = AA length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1
note = 2'-hydroxy-6'-methyltyrosine
SITE
2
note = D-amino acid
SITE
4
note = Amidated residue
SEQUENCE: 72
YKFK
4

SEQ ID NO: 73      moltype = AA length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1
note = 2'-hydroxy-6'-methyltyrosine
SITE
2
note = D-amino acid
SITE
4
note = Amidated residue
SEQUENCE: 73
YKFR
4

SEQ ID NO: 74      moltype = AA length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1
note = 2'-methyltyrosine
SITE
2
note = D-amino acid
SITE
4
note = Amidated residue
SEQUENCE: 74
YRFR
4

SEQ ID NO: 75      moltype = AA length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1
note = N,2',6'-trimethyltyrosine
SITE
2
note = D-amino acid
SITE
4
note = Amidated residue
SEQUENCE: 75
YRFR
4

SEQ ID NO: 76      moltype = AA length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1
note = 2'-hydroxy-6'-methyltyrosine
SITE
2
note = D-amino acid
SITE
4
note = Amidated residue
SEQUENCE: 76
YRFR
4

SEQ ID NO: 77      moltype = AA length = 4
FEATURE
source
1..4
mol_type = protein
organism = synthetic construct
SITE
1

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-continued

```

SITE          note = D-amino acid
              2
SITE          note = Dimethyltyrosine
              4
note = Amidated residue
SEQUENCE: 77
RYKF                                     4

SEQ ID NO: 78      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = D-amino acid
2
note = Dimethyltyrosine
SITE            4
note = Amidated residue
SEQUENCE: 78
RYFK                                     4

SEQ ID NO: 79      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = D-amino acid
4
note = Amidated Dimethyltyrosine
SEQUENCE: 79
RFKY                                     4

SEQ ID NO: 80      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = D-amino acid
3
note = Dimethyltyrosine
SITE            4
note = Amidated residue
SEQUENCE: 80
RFKY                                     4

SEQ ID NO: 81      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = D-amino acid
3
note = Dimethyltyrosine
SITE            4
note = Amidated residue
SEQUENCE: 81
RKYF                                     4

SEQ ID NO: 82      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers
1..4
mol_type = protein
organism = synthetic construct
SITE            1
note = D-amino acid
4
note = Amidated Dimethyltyrosine
SEQUENCE: 82
RKFY                                     4

SEQ ID NO: 83      moltype = AA  length = 4
FEATURE
source           Location/Qualifiers

```

-continued

```

source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           3
               note = Dimethyltyrosine
SITE           4
               note = Amidated D-amino acid
SEQUENCE: 83
FKYR                                     4

SEQ ID NO: 84      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           3
               note = D-amino acid
SITE           4
               note = Amidated Dimethyltyrosine
SEQUENCE: 84
FKRY                                     4

SEQ ID NO: 85      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           3
               note = Dimethyltyrosine
SITE           4
               note = Amidated residue
SEQUENCE: 85
FRYK                                     4

SEQ ID NO: 86      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           4
               note = Amidated Dimethyltyrosine
SEQUENCE: 86
FRKY                                     4

SEQ ID NO: 87      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = Dimethyltyrosine
SITE           3
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 87
FYRK                                     4

SEQ ID NO: 88      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = Dimethyltyrosine
SITE           4
               note = Amidated D-amino acid
SEQUENCE: 88
FYKR                                     4

SEQ ID NO: 89      moltype = AA length = 4
FEATURE

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-continued

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source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           3
               note = D-amino acid
SITE           4
               note = Amidated Dimethyltyrosine
SEQUENCE: 89
KFRY                                     4

SEQ ID NO: 90      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           3
               note = Dimethyltyrosine
SITE           4
               note = Amidated D-amino acid
SEQUENCE: 90
KFYR                                     4

SEQ ID NO: 91      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = Dimethyltyrosine
SITE           3
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 91
KYRF                                     4

SEQ ID NO: 92      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = Dimethyltyrosine
SITE           4
               note = Amidated D-amino acid
SEQUENCE: 92
KYFR                                     4

SEQ ID NO: 93      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           4
               note = Amidated Dimethyltyrosine
SEQUENCE: 93
KRYF                                     4

SEQ ID NO: 94      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           3
               note = Dimethyltyrosine
SITE           4
               note = Amidated residue
SEQUENCE: 94
KRYF                                     4

SEQ ID NO: 95      moltype = AA length = 4
FEATURE

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-continued

```

source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = D-amino acid
SITE           2
               note = Dimethyltyrosine
SITE           3
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 95
RYRF                                     4

SEQ ID NO: 96      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = D-amino acid
SITE           2
               note = Dimethyltyrosine
SITE           3
               note = D-amino acid
SITE           4
               note = Amidated Dimethyltyrosine
SEQUENCE: 96
RYRY                                     4

SEQ ID NO: 97      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = D-amino acid
SITE           2
               note = Dimethyltyrosine
SITE           3
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 97
RRYR                                     4

SEQ ID NO: 98      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = D-amino acid
SITE           2
               note = Dimethyltyrosine
SITE           3
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 98
RYRW                                     4

SEQ ID NO: 99      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 99
WRFK                                     4

SEQ ID NO: 100     moltype = AA length = 4
FEATURE

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-continued

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source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 100
WRYK                                         4

SEQ ID NO: 101      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 101
WRWK                                         4

SEQ ID NO: 102      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           2
               note = D-amino acid
SITE           3
               note = Dimethyltyrosine
SITE           4
               note = Amidated residue
SEQUENCE: 102
WRYK                                         4

SEQ ID NO: 103      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 103
RWFK                                         4

SEQ ID NO: 104      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = D-amino acid
SITE           4
               note = Amidated residue
SEQUENCE: 104
RWFK                                         4

SEQ ID NO: 105      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein
               organism = synthetic construct
SITE           1
               note = D-amino acid
SITE           4
               note = Amidated Dimethyltyrosine
SEQUENCE: 105
RWKY                                         4

SEQ ID NO: 106      moltype = AA length = 4
FEATURE
source          1..4
               mol_type = protein

```

-continued

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SITE organism = synthetic construct
      1
SITE note = D-amino acid
      3
SITE note = Dimethyltyrosine
      4
SITE note = Amidated residue

SEQUENCE: 106
RWYK

SEQ ID NO: 107 moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
mol_type = protein
organism = synthetic construct
      1
SITE note = D-amino acid
      4
SITE note = Amidated residue

SEQUENCE: 107
RKWF

SEQ ID NO: 108 moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
mol_type = protein
organism = synthetic construct
      1
SITE note = D-amino acid
      4
SITE note = Amidated Dimethyltyrosine

SEQUENCE: 108
RKWY

SEQ ID NO: 109 moltype = AA length = 4
FEATURE Location/Qualifiers
source 1..4
mol_type = protein
organism = synthetic construct
      2
SITE note = D-amino acid
      4
SITE note = Amidated residue

SEQUENCE: 109
ARFK

```

4

4

4

4

1.-67. (canceled)

68. A method for reducing the level of cardiac troponin I in a mammalian subject in need thereof, the method comprising: administering to the subject a therapeutically effective amount of the peptide D-Arg-2',6'-Dmt-Lys-Phe-NH₂ (SEQ ID NO: 3) or a pharmaceutically acceptable salt thereof.

69. The method of claim **68**, wherein the subject has been diagnosed with heart failure.

70. The method of claim **69**, wherein the heart failure results from hypertension; ischemic heart disease; exposure to a cardiotoxic compound; myocarditis; thyroid disease; viral infection; gingivitis; drug abuse; alcohol abuse; pericarditis; atherosclerosis; vascular disease; hypertrophic cardiomyopathy; acute myocardial infarction; left ventricular systolic dysfunction; coronary bypass surgery; starvation; an eating disorder; or a genetic defect.

71. The method of claim **68**, wherein the peptide is administered orally, topically, systemically, intravenously, subcutaneously, intraperitoneally, or intramuscularly

72. The method of claim **68**, further comprising separately, sequentially or simultaneously administering a cardiovascular agent to the subject.

73. The method of claim **72**, wherein the cardiovascular agent is selected from the group consisting of: an anti-arrhythmia agent, a vasodilator, an anti-anginal agent, a corticosteroid, a cardiotropine, a diuretic, a sedative, an angiotensin converting enzyme (ACE) inhibitor, an angiotensin II antagonist, a thrombolytic agent, a calcium channel blocker, a thromboxane receptor antagonist, a radical scavenger, an anti-platelet drug, a β-adrenaline receptor blocking drug, α-receptor blocking drug, a sympathetic nerve inhibitor, a *digitalis* formulation, an inotrope, and an antihyperlipidemic drug.

74. The method of claim **68**, wherein the pharmaceutically acceptable salt comprises acetate or trifluoroacetate salt.

75.-117. (canceled)

* * * * *