Supplementary Information for:

Small Molecule Inhibition of Ligand-Stimulated

RAGE-DIAPH1 Signal Transduction

Michaele B. Manigrasso^{1@}, Jinhong Pan^{2@}, Vivek Rai^{1,@}, Jinghua Zhang¹, Sergey Reverdatto², Nosirudeen Quadri¹, Robert J. DeVita³, Ravichandran Ramasamy¹, Alexander Shekhtman^{2,^*} & Ann Marie Schmidt¹^{*}

(a) These authors contributed equally to this work.

^Co-Senior Authors

¹Diabetes Research Program, Department of Medicine, New York University Langone Medical Center, 550 First Avenue, New York, New York 10016

²Department of Chemistry, University at Albany, State University of New York,

1400 Washington Avenue, Albany, New York 12222

³RJD Medicinal Chemistry and Drug Discovery Consulting LLC, 332 W. Dudley Avenue, Westfield, New Jersey 07090





Supplementary Figure S1. Small molecule competitive inhibition of binding of ctRAGE to DIAPH1. (a-m). Testing of compounds 1-13 in the high throughput screening assays reveals a dose-dependent binding of fluorescent GFP-labeled ctRAGE to DIAPH1. Compounds were tested in duplicate at 10 μ M, 1 μ M, 0.1 μ M, and 0.01 μ M. Structures of compounds 1-13 are illustrated in the figure.



Supplementary Figure S2. Compounds 1, 2, 4, 5, 7, 8, 9, 10, 11, 12, and 13 bind to the structured part of ctRAGE. ${}^{1}H{}^{15}N{}$ HSQC NMR spectra of $[U{}^{-15}N{}]$ RAGE tail bound to compound 1(a), 2(b), 4(c), 5(d), 7(e), 8(f), 9(g), 10(h), 11(i), 12(j), and 13(k). The NMR peaks of ctRAGE residues that underwent either changes in the chemical shifts or extensive broadening due to interaction with the library compounds are labeled.



Supplementary Figure S3. Effects of compounds 1-13 on RAGE ligand- or PDGFstimulated migration of murine SMCs. Migration of SMCs after incubation with compounds

1-13 was assessed. After the addition of the compounds 1-13 (1 μ M) to the individual wells, a wound was immediately created down the center of the well using a p200 tip and a baseline photograph was taken. After 1.5 h incubation, the compounds were removed and murine aortic SMCs were stimulated with CML-AGE (10 μ g/ml) (a) or PDGF-BB (10 ng/ml) (b) for 7 h, and final images were taken. Each compound was tested in three independent wells and each well was photographed in 3 separate locations for a total of 9 images used. Representative images are shown.

Category	Parameter	Description
Assay	Type of assay	In vitro Binding Assay
	Target	GFP-ctRAGE – DIAPH1 binding
	Primary measurement	GFP Fluorescence
	Key reagents	GFP-ctRAGE and DIAPH1 (HeLa cell lysate)
	Assay protocol	High Throughput Binding Assay (See Methods)
	Additional comments	N/A
Library	Library size	58,000 compounds
	Library composition	Small molecule commercially-available library
	Source	DIVERSet library (CHEMBRIDGE), CT488A
	Additional comments	N/A
Screen	Format	384 well format
	Concentration(s) tested	10 μ M, final concentration dissolved in 0.1% DMSO
	Plate controls	Hela Lysate (DIAPH1) and GFP-ctRAGE
	Reagent/ compound dispensing system	Microplate Reader (PerkinElmer, Inc)
	software	
	Assay validation/QC	Z' score, 0.62; S/N, 8.3
	Correction factors	None
	Normalization	Plate background with no GFP
	Additional comments	N/A
Post-HTS analysis	Hit criteria	>50 % inhibition of binding of ctRAGE to DIAPH1
-	Hit rate	1.5 %
	Additional assay(s)	Secondary Assay – Positive hits above (N=777) subjected to 4 point dose response [10, 1, 0.1 and 0.01 μ M] \rightarrow (N=97) Compounds Subjected to Nuclear Magnetic Resonance (NMR) for Binding to ctRAGE and not DIAPH1 \rightarrow (N=13) Subjected to Fluorescence Titration for IC50
	Confirmation of hit purity and structure	Liquid chromatography-Mass Spectrometry and NMR
	Additional comments	N/A

Supplementary Table S1. Small molecule screening protocol and results

Supplementary Table S2. Compounds 1-13 Bind to the Cytoplasmic Domain of RAGE: Dissociation Constants

Compound	Dissociation Constant (nM)
1	9±4
2	3±1
3	18±4
4	2±1
5	1±0.5
6	0.3±0.05
7	3±1
8	1.5±0.5
9	3±1
10	2±0.5
11	2±0.6
12	32±10
13	6±3

	DMSO	C1	C2	C3	C4	C5	C6
n	6	6	6	6	6	5	6
11	0	0	0	0	0	5	0
Body	25.2±2.7	24.4±3.1	22.4±4.4	25.1±2.8	24.8±3.7	25.6±3.6	21.9±1.9
Glucose	333±76	367±83	357±58	362±65	379±74	285±40	347±63
(mg/dL) Days disbatio	40±10	44±5	44±5	39±7	42±7	41±8	38±11
ulabelic	C7	C8	CO	C10	C11	C12	C13
	C/	Co	69	C10	CII	C12	CIS
n	6	6					
	0	6	6	6	6	6	6
Body	0 21.6±1.4	6 23.67±1.9	6 25.1±5.0	6 24.2±2.9	6 25.1±2.4	6 24.8±2.3	6 24.0±3.0
Body Weight (g) Glucose	0 21.6±1.4 333±77	6 23.67±1.9 329±73	6 25.1±5.0 328±43	6 24.2±2.9 380±68	6 25.1±2.4 348±46	6 24.8±2.3 353±86	6 24.0±3.0 327±43
Body Weight (g) Glucose (mg/dL) Days	21.6±1.4 333±77 45+7	6 23.67±1.9 329±73 38+7	6 25.1±5.0 328±43 40+6	6 24.2±2.9 380±68 37+8	6 25.1±2.4 348±46 40+6	6 24.8±2.3 353±86 42+7	6 24.0±3.0 327±43 38+9

Supplementary Table S3. Characteristics of Mice Studied in the Isolated Perfused Heart

Gene Name (Taqman)	Gene Symbol	Assay ID/catalog number
TNF	Tnf	Mm00443258_m1
IL6	116	Mm00446190_m1
ACTB	Actb	4352341E
TNF	TNF	Hs01113624_g1
IL6	IL6	Hs00985639_m1
Eukaryotic 18S	18S	4310893E

Supplementary Table S4. Materials for performance of the real time PCR experiments