Gene	Primer	Sequence (5'to 3')
NDD 1	Forward	CTCCTGTTGTGTCTTCAGG
NKP-1	Reverse	CCCGATGAGGATCGGATTC
actro	Forward	CGTTACCTTCCTGTGCCATT
\$512	Reverse	CTCCATTTGCCAATCATGTG
STAI	Forward	AGTTGTGCATTTACGGGAGAG
512L	Reverse	GGATACTGCTTTCCACCACAG
н ээ	Forward	TGGCCTCACCATAAGAAAGG
11-33	Reverse	GCAAACGCTTGGATACTGC
UDDT	Forward	TTGCTTTCCTTGGTCAGGCA
пркі	Reverse	ATCCAACACTTCGTGGGGTC
R actin	Forward	GCCGCCAGCTCACCAT
p-actin	Reverse	TCGATGGGGTACTTCAGGGT
	Forward	ACCAGCCCCAGCAAGAGCACAAG
GADIN	Reverse	TTCAAGGGGTCTACATGGCAACTG

 Table S1: List of primers used for real-time PCR assays in the manuscript

Table S2. Proteome alterations induced by sST-2 in human cardiac fibroblasts

Protein name	Uniprot code	Gene name	Fold-change ST2 /Control	
Down-regulated proteins in ST-2-treated cells				
Histone H1.5	P16401	HIST1H1B	0.56029477	
Mitofusin-1	Q8IWA4	MFN1	0.6570627	
Actinfilament-associatedprotein 1	Q8N556	AFAP1	0.66072435	
Charged multivesicular body protein 4a	E9PSI1	CHMP4A	0.69763398	
SH3 domain-containing kinase-binding protein 1	Q5JPT2	SH3KBP1	0.70692753	
Small nuclear ribonucleoprotein E	P62304	SNRPE	0.71390947	
Na(+)/H(+) exchange regulatory cofactor NHE-RF2	Q15599-2	SLC9A3R2	0.73573919	
Tumor susceptibility gene 101 protein	F5H442	TSG101	0.75621962	
Prefoldinsubunit 2	Q9UHV9	PFDN2	0.75739442	
Synaptobrevinhomolog YKT6	015498	ҮКТ6	0.76696515	
Up-regulated proteins in ST-2-treated cells				
Nectin-2	Q92692-2	PVRL2	1.31688694	
Eukaryotic translation initiation factor 5B	A0A087WUT6	EIF5B	1.32187402	

Neuropilin-1	E9PEP6	NRP1	1.33730071	
E3 SUMO-protein ligase RanBP2	P49792	1.46901186		
ADP-ribosylation factor 5	P84085	ARF5	1.57810952	
Rab-likeprotein 6	Q3YEC7	RABL6	1.65312128	
Interleukin-1 receptor-like 1	E9PC41	IL1RL1	2.16884842	
Ig gamma-1 chain C region	A0A087X1C7	IGHG1	2.28690335	

FIGURE LEGENDS

Supplemental Figure 1. Representative western blot out of three. Stain free with 9 controls and 8 ST-2 conditions corresponding to 8 datapoints per condition graphed in the manuscript. Complete blots for α -SMA and Collagen 3 from main figure 1A, D.

Supplemental Figure 2. Representative western blot out of three. Stain free with 8 controls and 7 ST-2 conditions corresponding to 7 datapoints per condition graphed in the manuscript. Complete blots for Vimentin and CTGF from main figure 1A, F.

Supplemental Figure 3. Representative western blot out of two. Stain free with 5 controls and 5 ST-2 conditions corresponding to 5 datapoints per condition graphed in the manuscript. Complete blot for NRP-1 from main figure 2A.

Supplemental Figure 4. Representative western blot out of five. Stain free with 4 controls, 4 ST-2 and 4 ST-2+BAY conditions corresponding to 4 datapoints per condition graphed in the manuscript. Complete blots for α -SMA, Vimentin and CTGF from main figure 3B, D.

Supplemental Figure 5. Representative western blot out of three. Stain free with 5 controls, 4 ST-2 and 4 ST-2+KO NRP-1 conditions corresponding to 4 datapoints per condition graphed in the manuscript. Complete blots for α -SMA, Vimentin and CTGF from main figure 3H, K.

Supplemental Figure 6. Representative western blot out of five. Stain free with 6 controls, 2 NRP-1 10⁻¹⁰M, 2 NRP-1 10⁻⁹M and 2 NRP-1 10⁻⁸M conditions

corresponding to 2 datapoints per condition graphed in the manuscript. Complete blots for α -SMA and Vimentin from main figure 4A.

Supplemental Figure 7. Representative western blot out of two. Stain free with 3 controls, 4 NRP-1 10⁻¹⁰M, 4 NRP-1 10⁻⁹M and 4 NRP-1 10⁻⁸M conditions corresponding to 4 datapoints per condition graphed in the manuscript. Complete blot for CTGF from main figure 4E.

Supplemental Figure 8. Representative western blots out of five from each type. Stain free with 2 controls, 2 conditions of 10^{-8} M NRP-1 for 5 minutes, 2 conditions of 10^{-8} M NRP-1 for 10 minutes 2 conditions of 10^{-8} M NRP-1 for 15 minutes, 2 conditions of 10^{-8} M NRP-1 for 30 minutes and 2 conditions of 10^{-8} M NRP-1 for 60 minutes. These blots shown correspond to 2 datapoints per condition graphed in the manuscript. Complete blots for p-NF κ B, NF κ B, p-p42/44 and p42/44 from main figure 5A, B.

Supplemental Figure 9. Representative western blots out of five from each type. Stain free with 2 controls, 2 conditions of 10⁻⁸M NRP-1 for 5 minutes, 2 conditions of 10⁻⁸M NRP-1 for 10 minutes 2 conditions of 10⁻⁸M NRP-1 for 15 minutes, 2 conditions of 10⁻⁸M NRP-1 for 30 minutes and 2 conditions of 10⁻⁸M NRP-1 for 60 minutes. Complete blots for IRAK1/4, p-p38 and p38 from main figure 5C, D.

Supplemental Figure 10. Representative western blot out of two. Stain free with 5 controls, 5 NRP-1 and 5 NRP-1+BAY conditions, corresponding to 5 datapoints per condition graphed in the manuscript. Complete blot for α -SMA and Vimentin from main figure 5E.

Supplemental Figure 11. Representative western blot out of two. Stain free with 7 controls and 6 PO rats conditions corresponding to 6 datapoints per condition graphed in the manuscript. Complete blots for ST-2, IL-33, MyD88, TRAF6 and AP-1 from main figure 6A, F, G.

Supplemental Figure 12. Perivascular myocardial fibrosis in control and PO rats. Immunostaining for α -SMA, vimentin and fibronectin in control and PO rats





Supplemental Figure 2



Figure 2A - NRP-1 - 103 kDa



Supplemental Figure 4

	Control	Control	ST-2	ST-2	KD NRP-1	KD NRP-1	Control	ST-2	KD NRP-1	Control	Control	ST-2 H	D NRP-1
							1.	÷.		•			
				•									
										•			
	Control	Control S	\$1-2	ST-2	ST-2+	ST-2+ KD NRP-1	Control	ST-2	ST-2+ KD NRP-1	Control	Control	ST-2	ST-2+ KD NRP-1
Figure 3H - α-SMA – 42 kDa	-	-	-	-	ST-24	SL-2*	-	-	SL2+		-	-	SL2•
Figure 3H - Vimentin – 54 kDa	Control	Control S	81-2 • i	ST-2		KD NRP-1	Control	ST-2	KD NRP-1	Control	Control	ST-2	KD NRP-1
Figure 3K - CTGE- 38kDa	Control	Control S	\$T-2	ST-2	ST-2+ KD NRP-1	ST-2+ KD NRP-1	Control	ST-2	ST-2+ KD NRP-1	Control	Control	ST-2	ST-2+ KD NRP-1
				-		-	-	-					

Supplemental Figure 5



Figure 4A - α -SMA – 42 kDa



Figure 4E - CTGF- 38kDa



Figure 5A - p-NFκB – 65 kDa













Figure 5C - IRAK1/4 – 80 kDa

Control	5	10'	15'	30'	601	Control	5	10'	151	30"	601
			1	, Y					140.10		
1. 100 A	-	-	-	ener i	-	*****	1	-	-	-	





Figure 5D - p-p38 – 43 kDa



Supplemental Figure 10



