

## Supplementary Information for

## Phosphodiesterase 2 inhibition preferentially promotes NO-guanylyl cyclase-cGMP signaling to reverse the development of heart failure

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Figure S1. Design and verification of GC-1a and GC-A gene deletion. (A) The targeting vector comprised a LacZ-Neo<sup>r</sup> cassette flanked by sequences on either side of exons 6 and 7 resulting in deletion of the cyclase domain. (B) Immunoblot of GC-1a and GC-1\beta subunits in soluble lung extracts showing complete deletion of the GC-1a protein (and significant reduction in GC-1\beta). (C) Guanylyl cyclase activity is markedly reduced in soluble lung extracts from sGC1a<sup>-/-</sup> and GC-A<sup>-/-</sup> mice stimulated with DETA-NONOate (100  $\mu$ M) and ANP (1  $\mu$ M), respectively. (D) The vasorelaxant potency of the NO-donor Spermine-NONOate is significantly reduced in aortic rings from male and female GC-1a<sup>-/-</sup> mice. (E) Mean arterial blood pressure (MABP) is increased in male, but not female, GC-1a<sup>-/-</sup> animals. (F) Heart rate (HR) is similar in WT and GC-1a<sup>-/-</sup> mice. Data are expressed as mean±sem with analysis by one-way ANOVA with Bonferroni post hoc test or two-way ANOVA. \*\*\*p<0.001 v basal, ###p<0.001 v WT + DETA-NONOate or ANP, <sup>\$\$\$\$\$</sup>p<0.001 v corresponding WT. n=6.



[A]



Figure S2. Effects of PDE2 inhibition on cardiac fibrosis, GC-1a and PDE2A expression. (A) Representative images of whole heart fibrotic burden and (B) GC-1a expression in sham mice and animals undergoing abdominal aortic constriction (AAC) for 6 weeks (6w) in the absence and presence of BAY 60-7550 (10 mg/kg/day; p.o.; initiated at 3w). (C) PDE2A expression in sham mice and animals administered isoproterenol (20mg/kg/day; s.c.) for 2 weeks in the absence and presence of BAY 60-7550 (10 mg/kg/day; p.o.; initiated at 1w). Data are expressed as mean $\pm$ sem. n=3.

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Figure S3. Characteristic cardiac dysfunction in Langendorff hearts from mice subjected to AAC. (A) Coronary perfusion pressure (CPP), left ventricular developed pressure (LVDP) and heart rate (HR) in WT, GC-1 $\alpha^{-/-}$  or GC-A<sup>-/-</sup> sham animals or mice subjected to 6 weeks of abdominal aortic constriction (AAC) in the absence and presence of BAY 60-7550 (10 mg/kg/day; p.o.; initiated at 3w). (B) Changes ( $\Delta$ ) in CPP in response to BAY 60-7550 (100 nM) in WT, GC-1 $\alpha^{-/-}$  or GC-A<sup>-/-</sup> sham animals or mice subjected to 6 weeks of abdominal aortic constriction (AAC). Data are expressed as mean±sem. n=6-12.



Figure S4. The increased cardiomyocyte size in GC-1 $\alpha^{-/-}$  mice is not due to apoptosis. Representative images depicting terminal deoxynucleotidyl transferase (TdT) dUTP Nick-End Labeling (TUNEL)-staining of heart sections from wild type (WT) and GC-1 $\alpha^{-/-}$  sham mice and animals undergoing abdominal aortic constriction (AAC) for 6 weeks. Positive apoptotic control shown in the upper panel.



Figure S5. Treatment with a NOS inhibitor ameliorates the beneficial effects of PDE2 inhibition in HF. (A) Echocardiographic indices of heart structure & function, (B) cardiac fibrosis, and (C) cardiomyocyte size in sham mice and animals undergoing abdominal aortic constriction (AAC) for 6 weeks (6w) with administration of the NOS inhibitor L-N<sup>G</sup>-nitroarginine methylester (L-NAME; 100 mg/kg/day; p.o.) in the absence and presence of BAY 60-7550 (10 mg/kg/day; p.o.; initiated at 3w). Data are expressed as mean±sem and analyzed by one-way ANOVA with Bonferroni post hoc test. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001 v sham. n=6-12.



**Figure S6. Comparison of the effects of BAY 60-7550 across genotypes following pressure overload.** Echocardiographic indices of heart structure & function, cardiac fibrosis, and cardiomyocyte size in sham mice and animals undergoing abdominal aortic constriction (AAC) for 3 weeks (3w) or 6 weeks (6w) in the absence and presence of BAY 60-7550 (10mg/kg/day; p.o.; initiated at 3w). Data are expressed as mean±sem. n=6-12.



Figure S7. The cellular localization of PDE2 does not change overly in response to pressure overload. Representative images of wheatgerm agglutinin (cell membrane, red) and PDE2 (green) stained heart sections from wild type (WT), GC-A<sup>-/-</sup> and GC-1 $\alpha^{-/-}$  sham mice and animals undergoing abdominal aortic constriction (AAC) for 6 weeks.



Figure S8. Expression of pro-hypertrophic and pro-fibrotic markers is diminished by PDE2i following AAC. Quantitate PCR analysis of an array of established pro-hypertrophic and pro-fibrotic markers in whole heart homogenates from sham mice or animals subjected to abdominal aortic constriction (AAC) for 6 weeks in the absence and presence of BAY 60-7550 (10 mg/kg/day; p.o.; initiated at 3w). Data are expressed as mean±sem with analysis by one-way ANOVA with Bonferroni post-hoc test. \*p<0.05 v sham, #p<0.05 v AAC (6w). n=6.

Gene target	Primer sequence (5'-3')
$WT(CC_{1}c^{+/+})$	Forward: TTACCCCAGTAGCCATTTCC
WI (GC-1a)	Reverse: CTTCAGCTTCCCCAACCTCT
CC 1=-/-	Forward: AGGTAGCCGGATCAAGCGTAT
GC-1a	Reverse: GATGGATTGCACGCAGGTTCT
$WT (CC A^{+/+})$	Forward: GCATGGTTCAGCTCTAAGAC
$WI(GC-A^{+})$	Reverse: CCTTCAGTTATCTACATCTGC
CC 4-/-	Forward: CTAACCCTGTGAACTGTAAGC
GC-A	Reverse: CCTTCAGTTATCTACATCTGC

Table S1. Primer sequences for genotyping

Gene target	Primer sequence (5'-3')			
PDF?	Forward: ATCTTTGCCTTGTTTATTTCCTG			
FDE2	Reverse: CAGCCAGCACAGATTTCG			
AND	Forward: GGATTTCAAGAACCTGCTAGACC			
Alvr	Reverse: GCAGAGCCCTCAGTTTGCT			
σινο	Forward: TGGGCTGTAACGCACTGAA			
DIVE	Reverse: TGTTGTGGCAAGTTTGTGCTT			
aMHC	Forward: GCCCAGTACCTCCGAAAGTC			
amine	Reverse: GCCTTAACATACTCCTCCTTGTC			
<i>PMHC</i>	Forward: ACTGTCAACACTAAGAGGGTCA			
рмнс	Reverse: TTGGATGATTTGATCTTCCAGGG			
Collagon Tune 1 g1	Forward: TCTGACTGGAAGAGCGGAGAG			
Coulagen Type I al	Reverse: AGACGGCTGAGTAGGGAACA			
Collagon Tune 1 a?	Forward: TGGATACGCGGACTCTGTTG			
Collagen Type 1 a.2	Reverse: CCCTTTCGTACTGATCCCGATT			
Collagon Tune 4 gl	Forward: CTGGAGAAAAGGGCCAGAT			
Conagen Type 4 a.1	Reverse: TCCTTAACTTGTGCCTGTCC			
CTCE	Forward: GGGCCTCTTCTGCGATTC			
CIGF	Reverse: ATCCAGGCAAGTGCATTGGTA			
Fibronactin	Forward: CCGGTGGCTGTCAGTCAGA			
Fibronecun	Reverse: CCGTTCCCACTGCTGATTTATC			
CSK2a	Forward: GGAGAAGAAGGACGAGCTGTA			
USN30	Reverse: TGTCAGGGTCCACAAGCAAA			
CSK3R	Forward: CAAAAGGAGTGAAAAGCCAAGAG			
USKSp	Reverse: TTGCTGCCATCTTTATCTCTGCT			
MMD 2	Forward: GACAAGTTCTGGAGATACAATGAAGTG			
1 <b>VIIVIF - 2</b>	Reverse: CAGGTTATCAGGGATGGCATTC			
NEATe3	Forward: AACCCTATCGAGTGTTCCCA			
MPAICS	Reverse: TCCCGGTCAGTCTTTGCTTC			
NER	Forward: CCTACGGAACTGGGCAAATGT			
	Reverse: TCCCCTCTGTTTTGGTTGCT			
<b>PDI</b> _10	Forward: GCTTGCCTCTAGTGTCCTCC			
KI L-17	Reverse: TTGGCGATTTCATTGGTCTCA			
SERCA_2	Forward: TGGAACCTTTGCCGCTCATTT			
SERCA-2	Reverse: CAGAGGCTGGTAGATGTGTT			
TGE-R	Forward: TCAGACATTCGGGAAGCAGT			
101-p	Reverse: GCCCTGTATTCCGTCTCCTTG			
TIMP_7	Forward: GATTCAGTATGAGATCAAGCAGATAAAGA			
111/11 -2	Reverse: GCGAGACCCCGCACACT			
aSM4	Forward: ACTACTGCCGAGCGTGA			
	Reverse: ATAGGTGGTTTCGTGGATGC			
R_actin	Forward: CACAGCTTCTTTGCAGCTCCTT			
p-ucun	Reverse: TCAGGATACCTCTCTTGCTCT			

 Table S2. Primer sequences for qPCR analysis

	WT			GC-A <sup>-/-</sup>			GC-1α <sup>-/-</sup>		
	Sham	AAC (6w)	AAC (6w) + PDE2i	Sham	AAC (6w)	AAC (6w) + PDE2i	Sham	AAC (6w)	AAC (6w) + PDE2i
MABP	97.7	*114.1	*113.1	101.5	*118.7	*119.3	107.4	*117.1	*115.5
(mmHg)	±1.9	±3.4	$\pm 1.3$	±2.9	±3.9	±1.3	±3./	$\pm 2.0$	±2.8

**Table S3.** Effect of BAY 60-7550 (10 mg/kg/day) on mean arterial blood pressure (MABP) in pressure overload (AAC; 6w)-induced HF.

\*p<0.05 v sham, n=6-12

**Table S4.** Effect of BAY 60-7550 (10 mg/kg/day) on heart rate (HR) in pressure overload (AAC; 6w)- and isoproterenol (ISO; 2w)- induced HF.

	Sham	AAC (6w)	AAC (6w) + PDE2i	Sham	ISO (2w)	ISO (2w) + PDE2i		
HR (bpm)	529±48	524±11	514±28	515±32	631±25*	663±17*		
$*\pi < 0.05$ w show $\pi = 6.12$								

<sup>\*</sup>p<0.05 v sham, n=6-12