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1 **Neuregulin-1 Compensates for Endothelial Nitric Oxide**
2 **Synthase Deficiency**

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23 Running title: NRG1 Compensates for eNOS deficiency

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28 **Abbreviations**

29	Ang II	Angiotensin II
30	ECs	Endothelial cells
31	eNOS	Endothelial NO synthase
32	eNOS ^{-/-}	eNOS null, eNOS deficient, eNOS knock out
33	ERBB	Erythroblastic leukaemia viral oncogene homolog
34	ERK	Extracellular signal-regulated kinase (ERK)
35	NO	Nitric oxide
36	NRG1	Neuregulin-1
37	rhNRG1	Recombinant human neuregulin-1
38	WT	Wild type

39

40 **New & Noteworthy**

- 41 • ECs compensate for eNOS deficiency by increasing the secretion of NRG1.
- 42 • NRG1 administration prevents cardiac and renal hypertrophy and fibrosis caused by
- 43 Ang II infusion and eNOS deficiency.
- 44 • NRG1 expression is regulated by miR-134.

45

46

47 **Abstract**

48 Endothelial cells (ECs) secrete different paracrine signals that modulate the function of
49 adjacent cells; two examples of these paracrine signals are nitric oxide (NO) and neuregulin-1
50 (NRG1), a cardioprotective growth factor. Currently, it is undetermined whether one
51 paracrine factor can compensate for the loss of another. Herein, we hypothesized that NRG1
52 can compensate for endothelial NO synthase (eNOS) deficiency.

53 **Methods.** We characterized eNOS null and wild type (WT) mice by cardiac ultrasound and
54 histology and we determined circulating NRG1 levels. In a separate experiment, 8 groups of
55 mice were divided into 4 groups of eNOS null mice and wild type (WT) mice; half of the
56 mice received angiotensin II (Ang II) to induce a more severe phenotype. Mice were
57 randomized to daily injections with NRG1 or vehicle for 28 days.

58 **Results.** eNOS deficiency increased NRG1 plasma levels, indicating that ECs increase their
59 NRG1 expression when NO production is deleted. eNOS deficiency also increased blood
60 pressure, lowered heart rate, induced cardiac fibrosis, and affected diastolic function. In eNOS
61 null mice, Ang II administration increased cardiac fibrosis, but also induced cardiac
62 hypertrophy and renal fibrosis. NRG1 administration prevented the cardiac and renal
63 hypertrophy and fibrosis caused by Ang II infusion and eNOS deficiency. Moreover, *Nrg1*
64 expression in the myocardium is shown to be regulated by miR-134.

65 **Conclusion.** This study indicates that administration of endothelium-derived NRG1 can
66 compensate for eNOS deficiency in the heart and kidneys.

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68

69 **Key words:** Neuregulin-1, Fibrosis, Hypertrophy, Angiotensin II, NO signaling

70 **1 Introduction**

71 Morphologically, the endothelium is the largest continuous organ system and lines the vessels
72 in all tissues. (1, 2) Functionally, the endothelium regulates tissues by secreting multiple
73 paracrine factors, including nitric oxide (NO), which is produced by endothelial nitric oxide
74 synthase (eNOS). (3, 4) Decreased eNOS activity is the cornerstone of endothelial
75 dysfunction and is associated with different cardiovascular disorders, including high blood
76 pressure and cardiac hypertrophy. (4–10) Moreover, eNOS is crucial for normal kidney
77 function, as it has been shown that loss of NO causes glomerulosclerosis, tubular apoptosis,
78 and interstitial fibrosis. (11–14)

79 Besides NO, endothelial cells (ECs) produce many other paracrine factors—including small
80 molecules, peptides, and proteins—that modulate the function of adjacent cells in different
81 organs, including the heart and kidneys. (4) An important example is neuregulin-1 (NRG1),
82 an endothelium-derived growth factor, which activates ERBB3 (ERBB Receptor Tyrosine
83 Kinase 3) and ERBB4 receptors localized on adjacent tissue cells. (15–18) NRG1 plays
84 crucial roles in cardiovascular and renal development, and induces growth and differentiation
85 of different cell types. (5, 19, 28, 20–27) The cardioprotective effects of NRG1 are well
86 established, (9, 18, 37, 38, 29–36) and a phase 3 clinical trial is ongoing in patients with
87 systolic heart failure (NCT03388593). Recent studies in kidney physiology indicate that
88 disrupted ERBB signaling leads to fibrotic injury, polycystic kidney diseases, and kidney
89 cancer, (39–41) and that NRG1 has protective effects on the kidneys in a mouse model of
90 type-1 diabetes mellitus. (39)

91 Although the protective effects of many endothelium-derived factors have been studied in
92 detail separately, an open question remains whether ECs modulate secretion of one paracrine
93 factor in case another paracrine factor is abrogated; *i.e.*, if one paracrine factor is
94 downregulated, will ECs compensate by increased production of another paracrine factor? A
95 second question is whether exogenous administration of this other factor can compensate for
96 the initial defect.

97 To test the specific hypotheses that ECs respond to endothelial NO synthase (eNOS)
98 deficiency by increasing their production of NRG1, we first characterized cardiac function,
99 morphology, and NRG1 levels in eNOS null mice compared to wild type (WT) mice. To test
100 the hypothesis that exogenous administration NRG1 can compensate for eNOS deficiency, we

101 performed another experiment (in WT and eNOS null mice) in which we added Ang II
102 infusion to half of the groups to increase the severity of the model. Although the
103 cardioprotective effects of the endothelium-derived paracrine factors NO and NRG1 have
104 been well studied separately, the interaction between both has never been studied before.

105

106 **2 Materials & methods**

107 **2.1 Animal models.** All animal experiments were approved by the institutional ethics
108 committee of the University of Antwerp and conform to the Guide for the Care and Use of
109 Laboratory Animals published by the US National Institute of Health (Updated version 2011).
110 15–16-week-old eNOS knock-out mice (The Jackson Laboratory, male and female, eNOS null
111 mice; N=42) (42–44) and their wild type littermate control mice (WT, male and female;
112 N=42) were used in 2 separate experiments to determine cardiac function by ultrasound,
113 hypertrophy and fibrosis of the heart and kidneys, and NRG1 levels. In the second
114 experiment, osmotic minipumps (ALZET, model 2004) filled with Ang II (400 ng/kg.min,
115 Sigma Aldrich) (45) were implanted for 28 days—to induce a more severe phenotype—in
116 half of the eNOS null mice and WT mice under general anesthesia (3% isoflurane), while the
117 other half of the mice underwent sham surgery. Mice were randomized to receive either daily
118 recombinant human NRG1 (rhNRG1) injections (20 µg/kg.day, intraperitoneal, Peprotech) or
119 vehicle (PBS) for 4 weeks, starting on the day of surgery. (18) The dose of NRG1 was based
120 on previous studies. (18, 46, 47)

121 Cardiac ultrasound (Vevo 2100, VisualSonics) and blood pressure measurements (Coda, Kent
122 Scientific) were performed in week 4 of the experiment. Cardiac ultrasound was used to assess
123 systolic and diastolic function, as well as cardiac dimensions including LV mass, which was
124 calculated according to the formula: $LV\ mass = (0.8 * (1.04 * ((LVEDD + IVSd + LVPWd)^3 -$
125 $LVEDD^3))) + 0.6$. At the end of 4th week, mice were euthanized using pentobarbital (100
126 mg/kg, intraperitoneal, CEVA Santé Animale). Blood was collected from the periorbital
127 venous sinus. The apex of the heart was fixed in tissue embedding medium (NEG 50,
128 ThermoFischer Scientific), the mid part was fixed in 4% buffered formalin and embedded in
129 paraffin, and the basal segment of the heart was snap frozen for Western blot analyses. The
130 left kidney was cut longitudinally in 2 pieces and embedded in NEG 50 and paraffin. The
131 right kidney was snap frozen for Western blot analyses.

132 **2.2 Histology.** Paraffin embedded heart and kidney sections were stained with Masson's
133 trichrome to measure fibrosis. Fibrosis was measured on the total area of heart and kidney
134 sections, as well as in glomeruli and perivascular area; 10 glomeruli and 6 micro vessels were
135 randomly analyzed. Cardiomyocyte borders were visualized using both laminin staining
136 (Novus Biologicals) and fluorescent wheat germ agglutinin lectin (WGA, Invitrogen, Alexa
137 Fluor 594) and cardiomyocyte cross sectional area—at the level of the nucleus—was
138 calculated by averaging 25 random cardiomyocytes per image per animal (5 images per

139 animal). Isolectin staining (Isolectin IB, Alexa Fluor, Invitrogen) and DAPI staining was
140 performed on cardiac tissue for visualization of capillaries to determine capillary density,
141 which was calculated as the average number of capillaries/mm². A total of 3 images per
142 animal was used. Apoptosis was visualized by staining with antibodies specific for cleaved
143 caspase 3 (Cell Signaling) and with TUNEL assay kit (ApopTag Plus Peroxidase In situ
144 Apoptosis Detection kit, Merck). Images were acquired with an Olympus microscope and
145 Nikon eclipse Ti fluorescence microscope and analyzed with ImageJ software.

146 **2.3 Western blotting.** Heart and kidney sections were incubated in
147 Radioimmunoprecipitation assay lysis buffer (RIPA, Sigma) supplemented with protease and
148 phosphatase inhibitors (complete Mini and PhosSTOP, Roche) on ice for 5 min and mixed
149 using a Precellys 24 tissue homogenizer (Bertin Instruments). After centrifugation, to pellet
150 non-dissolved material, protein concentration of the supernatant of each sample was
151 determined using the bicinchoninic acid method (BCA, ThermoFischer). Equal amounts of
152 protein were separated on Bolt 4–12% bis-tris gels (Invitrogen) and blotted onto
153 polyvinylidene fluoride membranes. Membranes were blocked in Odyssey Blocking Buffer
154 (Li-Cor Biosciences) and probed with primary antibody (overnight, 4°C). Following primary
155 antibodies were used: rabbit anti p44/42 MAPK (Erk1/2, Cell Signaling 4695, 42,44 kDa,
156 1:1000) and rabbit anti phospho-p44/42 MAPK (Erk1/2, Thr202/Tyr204, Cell Signaling 4377,
157 42,44 kDa, 1:1000), and GAPDH (Cell Signaling 5174, 37 kDa, 1:2000). Subsequently,
158 infrared fluorescence (IR)-conjugated secondary antibodies (anti-rabbit: IgG926-32211 and
159 anti-mouse IgG926-68070; Li-COR Biosciences, 1:20000) were used for IR fluorescence
160 detection using an Odyssey SA IR imaging system (Li-COR Biosciences). Each Western blot
161 panel pictured in this article was derived from one single continuous blot.

162 **2.4 Urinary albumin analysis.** Urine was collected during a 24-hour period in metabolic
163 cages and stored at -80°C. Urinary albumin was measured using a mouse albumin ELISA kit
164 (Bethyl Laboratories) according to manufacturer's protocol and with a 10000 fold dilution.
165 Urinary creatinine concentration was measured using a clinical grade autoanalyzer (Siemens
166 Vista 1500, Laboratory of Clinical Biology at the Antwerp university hospital).

167 **2.5 ELISA of NRG1 in plasma.** Immediately after sacrifice, heparinized blood was
168 centrifuged at 14,000 rpm at 4°C for 15 min and plasma was separated from debris. Plasma
169 NRG1 was measured using a mouse NRG1 ELISA kit (Bethyl Laboratories) according to
170 manufacturer's protocol. Plasma samples were used without any dilution.

171 **2.6 miRNA analysis *in vivo*.** Two miRNA target gene prediction tools were used to find
172 miRNAs targeting *NRG1* in human and mouse: miRSystem and miRWalk. miRSystem is a
173 database which integrates seven miRNA target gene prediction programs: DIANA, miRanda,
174 miRBridge, PicTar, PITA, rna22, and TargetScan. (48–50) The miRWalk database contains
175 information produced with TarPmiR, a miRNA-mRNA binding site prediction tool that
176 applies a random-forest-based approach. Next, expression of the *NRG1*-targeting miRNAs
177 was checked in the TissueAtlas database, with focus on veins, arteries, hearts, and kidneys of
178 human biopsy samples. The 4 miRNAs—miR-16, miR-24, miR-134, and miR223—with the
179 highest mean expression in these 4 organs were evaluated in WT and eNOS-null mice with
180 qPCR. Total RNA was isolated (Mirvana kit, Invitrogen) from OCT embedded heart sections
181 of Ang II-injected animals (WT; N=8 and eNOS^{-/-}; N=7) using Precellys Homogenizer
182 device. Cardiac expression of the miRNA panel (miR-16-5p, miR-24-3p, miR-134-5p, miR-
183 223-3p) was quantified with qPCR, using TaqMan PreAmp Master Mix (Applied
184 Biosystems), TaqMan Universal Master Mix II no UNG (Applied Biosystems), and TaqMan
185 primers (Life Technologies). All samples were normalized to RNU6B and relative log fold
186 change expression was calculated using $\log(2^{-(\Delta\Delta Ct)})$ method in R.

187 **2.7 *In vitro* experiments with miRNA-134 inhibitor.** Primary mouse cardiac ECs were
188 obtained via magnetic bead isolation from C57BL/6 mice (Neonatal Cardiac Endothelial Cell
189 Isolation Kit, Miltenyi Biotec). Cells were cultured in DMEMF12 supplemented medium at
190 37°C, 5% CO₂. Cells were seeded at a density of 20,000 cells/well in a 96 well plate. Mmu-
191 mir-134-5p power inhibitor (700nM, Qiagen) was added to the cells according to the
192 manufacturer's protocol, followed by the addition of 1µM of Ang II (Sigma Aldrich, A9525).
193 Cells were treated for 48 hours, after which total RNA was isolated (MirVana Kit,
194 Invitrogen). mRNA expression levels of *Nrg1* were determined using RT-qPCR. All samples
195 were normalized to β-actin (Mm00607939-s1, Life Technologies), while relative fold change
196 expression was calculated using the $\log(2^{-(\Delta\Delta Ct)})$ method.

197 **2.8 Reverse transcription quantitative polymerase chain reaction of *Nppa* and *Nppb*.**
198 To determine expression of *Nppa* (coding for atrial natriuretic peptide, ANP) and *Nppb*
199 (coding for brain natriuretic peptide, BNP), cardiac tissue was homogenized using a Precellys
200 24 Homogenizer (Bertin Instruments). RNA was isolated with the Nucleospin RNA isolation
201 kit (Filter Service) according to the manufacturer's protocol. Reverse transcription was
202 performed using TaqMan™ Reverse Transcription Reagents (Invitrogen), followed by
203 quantification on PCR using TaqMan™ Universal PCR Master Mix (Applied Biosystems)

204 and TaqMan primers (*Nppa*: mm01255747_g1; *Nppb*:mm01255770_g1, Life Technologies)
205 on a Quant Studio 3 qPCR device. *Actb* (coding for beta-actin) was used as a house keeping
206 gene, while relative expression was calculated according to the $\log_2^{(-\Delta\Delta CT)}$ method.

207 **2.9 Statistical analysis.** All data were analyzed using Graph-pad Prism software version
208 8.0, and presented as Mean±SEM. Student's T test was used for comparison between two
209 genotypes (WT or eNOS^{-/-}) and Two-way ANOVA was used for each genotype separately
210 (WT or eNOS^{-/-}) and a Bonferroni correction for multiple comparison was used in post-hoc
211 tests. In case of non-normal distribution; logarithmic transformation of the data was
212 performed. Statistical significance was defined as P < 0.05.

213 **3 Results**

214 **3.1 eNOS deletion increases NRG1 plasma levels.** We first determined whether NRG1
215 increases upon eNOS deletion, by measuring NRG1 protein levels in the circulation. NRG1
216 protein levels were 5-fold higher in blood plasma of eNOS null mice (Fig. 1A). eNOS
217 deletion did not induce cardiomyocyte hypertrophy (Fig. 1B), but led to significant changes in
218 interstitial fibrosis in the heart (Fig. 1C), heart rate (Fig. 1D), and in a number of ultrasound
219 measurements of diastolic function: isovolumic relaxation time (IVRT), early (E), and late
220 (A) filling rate (Fig. 1I–J). eNOS deletion did not change ventricular ejection fraction or
221 cardiac mass (Fig. 1E–G).

222 **3.2 eNOS deletion, but not chronic infusion of a low dose of Ang II, increases blood**
223 **pressure.** Next, we aimed to study whether NRG1 could compensate for the loss of NO.
224 However, because the phenotype of eNOS null mice was mild and limited to myocardial
225 fibrosis without hypertrophy, we decided to include groups receiving a chronic infusion of
226 Ang II which induces a more severe phenotype with cardiac hypertrophy. We used a low dose
227 of Ang II—called a *subpressor* dose because of limited effects on blood pressure—to limit the
228 high mortality rate observed with higher doses of Ang II in eNOS null mice. (45) Although
229 eNOS null mice showed a higher blood pressure compared to their WT mice, *subpressor*
230 doses of Ang II or NRG1 treatment did not affect blood pressure (Fig. 2A–C); this indicates
231 that the effects observed in this study are largely blood pressure independent. The mortality
232 rate was higher in eNOS null animals compared to WT animals, when treated with Ang II
233 (Fig. 2D), but not statistically significant (33% in eNOS^{-/-} +Ang II vs. 11% in WT +Ang II
234 group). NRG1 did not significantly affect the mortality in this study. No mortality was
235 observed in animals without Ang II treatment, independent of their genotype.

236 **3.3 NRG1 administration compensates for eNOS deficiency in Ang II–induced cardiac**
237 **hypertrophy.** *Subpressor* doses of Ang II induced a more severe cardiac phenotype, and this
238 phenotype was still dependent on eNOS levels, *i.e.* it was milder in WT mice, allowing us to
239 study whether NRG1 could compensate for the loss of NO. Specifically, in WT mice, the
240 effects of *subpressor* doses of Ang II on cardiac hypertrophy were small and statistically non-
241 significant (Fig. 3B, C). However, when eNOS was deleted, *subpressor* doses of Ang II
242 induced a hypertrophic response, as shown by a significantly increased intraventricular
243 septum thickness (Fig. 3B) and increased post-mortem heart-to-body weight (Fig. 3C).
244 Importantly, daily administration of NRG1 completely compensated for the hypertrophic

245 responses induced by eNOS deficiency upon *subpressor* doses of Ang II (Fig. 3B, C). Left
246 ventricular end-diastolic volume was similar in all groups (Fig. 3D), and heart rate was slower
247 in eNOS null mice than in WT mice, but not affected by administration of either Ang II or
248 NRG1 (Fig. 3E). *Nppb*, coding for BNP, a marker of cardiomyocyte strain, was not
249 differentially expressed in cardiac tissues. In contrast, *Nppa*, coding for ANP, showed an
250 upregulation in eNOS null mice treated with NRG1 (Fig. 3F, G). Administration of NRG1
251 prevented the increase in cardiomyocyte cross sectional area (Fig. 3H, I), induced by the
252 combination of Ang II infusion and eNOS deficiency as defined by laminin staining (Fig.
253 3H). The same trend was observed with wheat germ agglutinin staining, with a significant
254 group effect of NRG1 in eNOS null mice ($P=0.02$; two-way ANOVA; Suppl. Fig. 1).

255 **3.4 NRG1 administration compensates for eNOS deficiency in Ang II–induced cardiac**
256 **fibrosis.** Deficiency of eNOS combined with Ang II administration also led to significantly
257 higher interstitial fibrosis (Fig. 4A, C)—and NRG1 treatment suppressed this fibrotic
258 response (+Ang II +NRG1, Fig. 4B). eNOS deficiency combined with Ang II administration
259 also led to significantly increased perivascular fibrosis, which was again attenuated by NRG1
260 treatment (Fig. 4C, D). No significant differences in myocardial capillary density were
261 observed between the different groups (Fig. 4E, F).

262 **3.5 NRG1 administration compensates for eNOS deficiency in Ang II–induced renal**
263 **albuminuria, hypertrophy, and fibrosis.** To determine whether the compensatory effects of
264 NRG1 to eNOS deficiency are cardiac-specific or system-wide, we also examined renal
265 function and morphology, because ECs have an important functional role in the kidney. First,
266 we measured albuminuria, because it often is the first sign of kidney dysfunction. eNOS
267 deficiency, even in the absence of Ang II, significantly increased albuminuria—quantified by
268 the relative urinary albumin concentration normalized to urinary creatinine concentration, and
269 this effect was accentuated when eNOS deficiency was combined with Ang II administration
270 (Fig. 5A). Here, NRG1 treatment only partly normalized albuminuria (Fig. 5A).

271 Second, NRG1 significantly reduced kidney weight in eNOS null mice, regardless of Ang II
272 administration (Fig. 5B). The glomerulus is a network of capillaries and histologically defined
273 as a tuft. An increase in glomerular tuft area corresponds to an increased glomerular volume
274 and is a key pathological finding in glomerulosclerosis. (51) Both eNOS deficiency and Ang
275 II administration increased glomerular tuft area significantly, with the highest areas measured
276 when combined (Fig. 5C, D). Administration of NRG1 compensated for eNOS deficiency,

277 both in the presence and absence of Ang II.

278 Third, eNOS deficiency and Ang II administration significantly increased fibrosis on total
279 sections of kidneys (Fig. 5E, F). Specifically, in glomeruli, the fibrotic response to eNOS
280 deficiency and Ang II was most pronounced (Fig. 5G), indicating most damage at this stage
281 primarily occurs in the glomeruli. Again, administration of NRG1 compensated for this
282 fibrosis induced by eNOS deficiency, Ang II administration, and the combination of both
283 (Fig. 5H). Together with the data in the heart, these data indicate that the compensatory
284 effects of NRG1 for eNOS deficiency on fibrosis are preserved in different organs.

285 **3.6 NRG1 administration compensates for eNOS deficiency–induced Erk**
286 **phosphorylation and vascular apoptosis.** The previous sections showed that administration
287 of NRG1 can compensate for the lack of NO in terms of tissue function, hypertrophy, and
288 fibrosis. Here, we questioned whether these compensatory effects could be traced to
289 subcellular signaling pathways and apoptotic pathways. First, the extracellular signal-
290 regulated kinase (Erk) pathway is a paramount pathway for tissue homeostasis in the heart.
291 (52) Western blot analysis of the myocardium showed that Ang II administration, eNOS
292 deficiency, or both together increased phosphorylation of Erk1/2 (Fig. 6A). Administration of
293 NRG1 compensated significantly for eNOS deficiency-induced phosphorylation of Erk1/2
294 (Fig. 6B). Second, eNOS deficiency induced a significantly higher level of apoptosis—
295 determined by cleaved caspase 3 staining (Fig. 6C)—in intramural cardiac arteries with or
296 without Ang II infusion. NRG1 treatment significantly reduced vascular apoptosis induced by
297 eNOS deficiency (Fig. 6D). No apoptosis was observed in cardiomyocytes. TUNEL staining
298 on cardiac tissue also showed no significant differences in cardiomyocyte apoptosis between
299 the different groups ($P=0.87$ in WT and $P=0.95$ in eNOS null mice). In summary, the
300 compensatory effects of NRG1 administration on eNOS deficiency can be traced to
301 subcellular signaling and apoptotic pathways.

302 **3.7 *Nrg1* expression is, in part, regulated by miR-134.** It is currently undetermined which
303 miRNAs regulate *Nrg1* expression in ECs. Therefore, two miRNA target gene prediction
304 tools were used to discover miRNAs targeting *NRG1* in human and mouse: miRSystem and
305 miRWalk. The miRSystem database resulted in 20 miRNAs predicted to bind both human and
306 mouse *NRG1* mRNA and the miRWalk database resulted in 53 miRNAs predicted to bind
307 both human and mouse *NRG1* mRNA; three of these miRNAs were identical. Expression of
308 these 70 miRNAs in veins, arteries, hearts, and kidneys—organs with a large fraction of

309 ECs—of human biopsy samples was checked in the TissueAtlas database. 27 miRNAs
310 showed significant expression in these organs and the four miRNAs—miR-16, miR-24, miR-
311 134, and miR223—with the highest mean expression in these organs were evaluated in
312 cardiac tissues of WT and eNOS null mice with qPCR. All four miRNAs were downregulated
313 in eNOS null hearts—which is consistent with the theory that these miRNAs inhibit *Nrg1*
314 expression—but only the downregulation of miR-134 was significant (Fig. 7A). Next, we
315 tested whether a miR-134 inhibitor increases *Nrg1* expression in ECs *in vitro*, which was not
316 the case in basal conditions (Fig. 7B). However, the miR-134 inhibitor did result in a mild
317 increase in *Nrg1* expression in ECs, when stimulated with Ang II (Fig. 7C).

318 Data supplements: <https://doi.org/10.5281/zenodo.4749643>

319 **4 Discussion**

320 The endothelial system occupies an unique position at the interface between the individual
321 organs and the circulating blood and regulates overall homeostasis by affecting all organs in
322 the body. (4, 53, 54) The most studied role of the endothelial system has been its control of
323 vasomotoricity and organ perfusion by interacting of ECs with subjacent smooth muscle cells.
324 More importantly, however, ECs are also present in capillaries, which constitute the largest
325 endothelial surface in every organ. In capillaries, ECs directly communicate with adjacent
326 tissue cells, such as neurons, alveolar cells, cardiomyocytes, tubular and glomerular cells in
327 the kidney, hepatocytes, and fat cells. (4, 53, 54) The endothelial system is an adaptive system
328 that can undergo dramatic changes in gene expression during development and disease. (55)

329 Although the protective effects of many endothelium-derived factors have been studied in
330 detail, an open question remains whether ECs modulate secretion of one paracrine factor in
331 case another paracrine factor is abrogated and whether supplementary administration of this
332 other factor can compensate for the initial defect. In this study, we tested the specific
333 hypotheses that ECs will respond to eNOS deficiency by increasing their production of NRG1
334 and that administration of NRG1 can compensate for eNOS deficiency. Summarized, this
335 study provides evidence that ECs increase the release of NRG1 upon eNOS deficiency, albeit
336 insufficient to prevent a pathophysiological phenotype, but that administration of *extra* NRG1
337 can compensate for the negative effects of diminished NO/eNOS signaling.

338 For this study, we selected two important endothelial-derived paracrine factors with proven
339 physiological relevance and with different underlying signaling pathways. First, we selected
340 NO, because it is well known that NO has crucial roles in the regulation of various
341 physiological processes. (56) NO is synthesized from L-arginine mediated by the NOS
342 family, which includes neuronal NOS (nNOS) in neural tissue, inducible NOS (iNOS) in
343 macrophages, and endothelial NOS (eNOS) in the endothelium. (57) Reduced NO
344 bioavailability is the hallmark of endothelial dysfunction, which contributes to different
345 cardiovascular diseases including high blood pressure, cardiac hypertrophy, cardiac
346 remodeling, and atherosclerosis. (4, 5, 58–62) A disrupted NO/eNOS signaling is also
347 associated with different kidney disorders, such as glomerulosclerosis, interstitial fibrosis,
348 diabetic nephropathy, and cardiorenal syndrome. (11) It has been shown previously that
349 eNOS null mice display normal cardiac function and ejection fraction when young, but
350 develop cardiac dysfunction at older age. (44) Our data are also consistent with earlier reports

351 showing decreased heart rate and increased blood pressure in eNOS null mice. (43) Second,
352 we selected NRG1, an epidermal growth factor that is produced by ECs and that exerts its
353 effect via ERBB tyrosine kinase receptors. (17) NRG1 plays a crucial role in cardiac
354 development and pathophysiology, and has anti-apoptotic and anti-fibrotic effects. (10, 16,
355 18) Moreover, NRG1 has been shown to attenuate heart failure in several animal models and
356 is currently under evaluation as a potential treatment for heart failure in a phase 3 clinical
357 trial. (36, 37)

358 We have previously reported that NRG1 attenuates Ang II-induced myocardial hypertrophy
359 and fibrosis in model of Ang II-induced hypertension in the mouse. (18) These observations
360 are partly recapitulated in this study. The difference of the present study is that we used a low
361 subpressor dose of Ang II, which did not induce the full picture of abnormalities in the heart
362 and kidneys observed with higher hypertension-inducing doses of Ang II. Only in mice where
363 eNOS was absent, subpressor Ang II doses induced myocardial fibrosis, myocardial
364 hypertrophy, kidney fibrosis, and albuminuria, which confirms the protective role of NO in
365 heart and kidney. An interesting observation in this study is that NRG1 stimulated *Nppa*
366 expression in cardiac tissue of eNOS null mice; because ANP has protective properties in the
367 heart and the kidneys, (63) one could speculate that part of the beneficial effects of NRG1
368 might be mediated by increased ANP levels. The main finding of the present study is that
369 NRG1 can compensate for NO when hypertrophic and fibrotic tissue responses are initiated
370 due to lack of NO.

371 Another conclusion of this study is that, although NRG1 levels were robustly induced in
372 eNOS null mice in the circulation, this upregulation is insufficient to prevent a
373 pathophysiological phenotype of hypertrophy and fibrosis. However, the fact that the
374 upregulation does not prevent a phenotype does not mean that the upregulation of NRG1 has
375 no physiological relevance. Compensatory pathways in biology almost never reverse
376 detrimental effects of deletion of an important signaling pathway. It is conceivable that,
377 without compensatory upregulation of NRG1, the phenotype of eNOS null mice would be
378 even more severe. An example supporting this claim is the finding that *Nrg1* deletion in
379 endothelial cells worsens the cardiac phenotype after ischemia/reperfusion injury. (64)

380 A next conclusion of this study is that the cardio- and nephro-protective effects of NRG1 are
381 independent of the eNOS pathway. To some extent, this finding is surprising because there
382 are indirect indications in the literature that some of the cardioprotective effects of NRG1 are

383 mediated by eNOS activation. (5, 35) Especially since NRG1 has been shown to induce
384 phosphorylation of endothelial NO synthase and nitrite production in cardiomyocytes. The
385 precise cell and tissue-protecting effects are still largely unknown, although anti-apoptotic and
386 anti-inflammatory effects have been postulated.

387 Interestingly, in this study we show that the compensatory effects of NRG1 administration on
388 eNOS deficiency can be traced to subcellular signaling and apoptotic pathways. We selected
389 the Erk pathway because it is involved in hypertrophy (65) and selected cleaved caspase-3
390 because it is involved in cell-death pathways. However, it is likely that the compensatory
391 effects of NRG1 go beyond these two pathways and affect many other subcellular signaling
392 pathways involved in hypertrophy and fibrosis. With regard to myocardial capillary density,
393 our results are in line with previous studies showing that eNOS null mice at adult age are
394 indistinguishable from their WT littermates. In contrast, neonatal eNOS null mice have a
395 lower capillary density. (42) It has been shown in a rat model of diabetic cardiomyopathy
396 that NRG1 can induce expression of vascular endothelial growth factor (VEGF), (66) which
397 can induce cardiac and renal protection (67) in a dose-dependent manner. (68) However, in
398 contrast to these earlier studies, we did not observe increased capillary density upon
399 exogenous NRG1 administration. The reason for this discrepancy is unclear, yet the lack of
400 angiogenic abnormalities in the animal model used in the present study might prevent the
401 detection of NRG1-associated proangiogenic activity. In relation to EC signaling, we also
402 show that miR-134 is a regulator of *Nrg1* expression in ECs, nevertheless it is likely that
403 many other miRNAs are involved as well. Despite the identification of the miR-134/*Nrg1*
404 regulatory axis at the RNA level, no results were obtained at the protein level, which is a
405 limitation of the present study.

406 Consistent with the data from the present paper, data from human studies have been published
407 showing that circulating NRG1 levels are upregulated in patients with coronary artery disease
408 and heart failure—two diseases characterized by endothelial dysfunction. (69–71) These data
409 indicate that our study could also be relevant in human disease. Wider ranging implications
410 could result from the fact that pharmacological stimulation of one endothelial pathway can
411 compensate for a defect in another endothelial pathway. The therapeutic implications could be
412 that—even when one pathway is the main driver of kidney failure or heart failure—it is still
413 worthwhile to try to compensate by targeting other pathways.

414 Finally, it has recently been claimed by Zurek et al. that NRG1 induces cardiac hypertrophy

415 in post-myocardial infarction rats. (72) This claim was based on a mere increase in
416 ventricular mass—calculated by MRI—at a single dose (2.5µg/kg/h for 7 days); both lower
417 and higher doses did not induce hypertrophy. The data of Zurek et al. contradict the anti-
418 hypertrophic response induced by NRG1 that was observed in the present study, as well as in
419 several previous studies. (18, 73) Importantly, in this study we demonstrated the anti-
420 hypertrophic effect of NRG1 using three well-established parameters of hypertrophy:
421 interventricular septum thickness using cardiac ultrasound, post-mortem heart weight, and
422 cardiomyocyte cross sectional area using histology.

423 **5 Conclusions**

424 This study indicates that endothelial-derived NRG1 can function as a compensatory pathway
425 for eNOS deficiency, but also that the natural upregulation of NRG1 is inadequate. The
426 compensatory effects of NRG1 for eNOS deficiency appear to be present in the endothelial
427 system throughout the organism. Translational implications are that administration of
428 endothelial-derived factors could be therapeutically effective, even if the underlying defect
429 involves a different paracrine factor.

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431

432 **6 Author Contributions**

433 VFM. Segers, G.W. De Keulenaer, P-J.D.F. Guns, and H. Shakeri designed research; H.
434 Shakeri, S. De Moudt, J.R.A. Boen, A.J. Leloup, J.O. Hendrickx, and G. Jacobs performed
435 research; H. Shakeri, J.R.A. Boen, S. De Moudt, G. Jacobs, and J. Hendrickx analyzed data;
436 V.F.M. Segers, P-J.D.F. Guns, G.W. De Keulenaer, G.R.Y. De Meyer, and H. Shakeri wrote
437 the paper; all authors read and approved the final version.

438

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446

447 **Figure legends**

448 **Figure 1. eNOS deletion increases NRG1 levels and myocardial fibrosis.** A) NRG1 levels
449 in plasma measured using ELISA for mouse NRG1 in eNOS deficient mice (eNOS^{-/-}) and
450 compared to the control group (N=5). B) Cardiomyocyte cross sectional area, C) cardiac
451 fibrotic area, and D) heart rate. Cardiac function was assessed by ultrasound: E) end-diastolic
452 interventricular septal thickness (IVS;d), F) left ventricular mass (LV mass), G) ejection
453 fraction (EF), H) isovolumic relaxation time (IVRT), I) left ventricular end-diastolic volume
454 (LV-EDV), J) early filling velocity (E), K) late filling velocity (A), and L) early diastolic
455 mitral annular tissue velocity (é). N=10 in B–L; Student's t-test; *P<0.05.

456 **Figure 2. Chronic infusion of a low dose of Ang II has no effect on blood pressure and**
457 **mortality.** A) Systolic blood pressure, B) diastolic blood pressure, and C) mean arterial
458 pressure (MAP) measured by tail-cuff method. D) The effect of Ang II treatment on mortality
459 in WT and eNOS^{-/-} animals. Graphs show both WT and eNOS^{-/-} animals with and without
460 Ang II infusion (AngII and Ctrl). Each group treated with and without NRG1 (NRG1 and
461 vehicle). N=8 per group; mean±SEM; two-way ANOVA.

462 **Figure 3. Administration of NRG1 compensates completely for Ang II- and eNOS**
463 **deficiency-induced cardiac hypertrophy.** A) Ejection fraction (EF) and B) diastolic
464 intraventricular septum thickness (IVS;d) measured by cardiac ultrasound. C) Post-mortem
465 heart weight normalized to body weight. D) Left ventricular end-diastolic volume measured
466 by ultrasound. E) Heart rate measured by CODA. mRNA expression levels of F) *Nppa* and G)
467 *Nppb*. H) Representative microscopic images of laminin staining of the heart used for
468 quantification of cross sectional area. I) Quantification of cardiomyocyte cross sectional area.
469 N=8 per group; mean±SEM; two-way ANOVA with Bonferroni correction for multiple
470 comparisons; *P<0.05; **P<0.01.

471 **Figure 4. Administration of NRG1 compensates for Ang II- and eNOS deficiency-**
472 **induced cardiac fibrosis.** A & C) Representative microscopic images of Masson's trichrome
473 staining (blue colour) of the heart to measure B) interstitial fibrosis in the total heart; and D)
474 perivascular fibrosis. E) Isolectin staining (green) and DAPI staining (blue) of the heart to
475 determine F) capillary density/mm². N=8 per group; Mean±SEM; two-way ANOVA with
476 Bonferroni correction for multiple comparisons; *P<0.05; ***P<0.001; #P<0.05 compared to
477 WT, ##P<0.01 compared to WT.

478 **Figure 5. Administration of NRG1 compensates for Ang II- and eNOS deficiency-**
479 **induced renal albuminuria, hypertrophy, and fibrosis.** A) Amount of urinary albumin
480 normalized by the urinary creatinine, measured from samples collected in 24 hours. B)
481 Kidney weight/body weigh at the end of the experiment. C) Representative images of
482 periodic-acid Schiff staining used to measure D) the size of the glomerular tuft area. E)
483 Representative images of Masson's trichrome staining on total kidney and G) glomeruli,
484 indicating F) renal interstitial fibrosis and H) glomerular fibrosis (blue color). N=8 per group;
485 Mean±SEM; two-way ANOVA with Bonferroni correction for multiple comparisons;
486 *P<0.05; **P<0.01; ***P<0.001.

487 **Figure 6. Administration of NRG1 compensates for eNOS deficiency-induced Erk**
488 **phosphorylation and vascular apoptosis.** A) Western blots of heart samples using
489 antibodies against Erk1/2 and phosphorylated Erk1/2 (p-Erk1/2); GAPDH is used as a protein
490 loading control. B) Analysis of relative band intensity of p-Erk1/2 normalized by the total
491 Erk1/2 protein. C) Representative microscopic images of cleaved-caspase 3 (Cl-casp3)
492 staining of the heart, to measure D) vascular apoptosis. N=8 per group; Mean±SEM; two-way
493 ANOVA with Bonferroni correction for multiple comparisons; *P<0.05.

494 **Figure 7. miR-134 regulates *Nrg1* expression.** A) Expression of four miRNAs in cardiac
495 tissue that were predicted to interact with NRG1 in human and mouse and with a high
496 expression in organs with a large fraction of ECs. N WT=7, N eNOS^{-/-}=8; Mean±SEM;
497 Unpaired T-test; *P<0.05. B) In primary cardiac ECs, the miR-134 inhibitor did not change
498 *Nrg1* expression in basal conditions. C) When Ang II was added to all groups (except to the
499 Untreated group), the miR-134 inhibitor induced a mild increase in *Nrg1* expression. N=9 per
500 group. Mean±SEM; one-way ANOVA with Tukey correction for multiple comparisons;
501 *P<0.05.

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