Nigericin Calbiochem-Novabiochem, Bad Soden

Germany

Nitrocellulose membrane Schleicher & Schuell, Dassel, Germany

Paraformaldehyde Merck, Darmstadt, Germany

pEGFP(YFP-CFP)-N1 vector Clontech, Heidelberg, Germany

Penicillin Sigma-Aldrich, Steinheim, Germany

Phosphoric acid Merck, Darmstadt, Germany

Ponceau S Roche Diagnostics, Mannheim, Germany

Potassium cloride Sigma-Aldrich, Steinheim, Germany

Kaleidoscope prestained marker Bio Rad, Munich, Germany

PVDF-filters, Immobilon-P Millipore Corporation, Bedford, USA

Sodium chloride Sigma-Aldrich, Steinheim, Germany

Sodium citrate Merck, Darmstadt, Germany

Sodium hydroxide Carl Roth, Karlsruhe, Germany

Sodium phosphate Merck, Darmstadt, Germany

Sodium-dihydrogen phosphate Merck, Darmstadt, Germany

Sodiumdodecyl sulfate (SDS) Sigma-Aldrich, Steinheim, Germany

Sodim-hydrogen carbonate Sigma-Aldrich, Steinheim, Germany

Streptomycine Sigma-Aldrich, Steinheim, Germany

Tetramethylendiamine (TEMED) Sigma-Aldrich, Steinheim, Germany

Triton X-100 Sigma-Aldrich, Steinheim, Germany

Tween-20 Sigma-Aldrich, Steinheim, Germany

Zeocin Invitrogen, Leek, The Netherlands

### Antibodies:

monoclonal c-myc (9E10) antibody Roche Diagnostics, Mannheim, Germany

monoclonal FLAG (M2) antibody Sigma-Aldrich, Steinheim, Germany

monoclonal GFP (JL-8) antibody BD Biosciences

rabbit anti-NT-ET<sub>A</sub> antibody Dr.Jens Furkert, FMP-Berlin, Germany

rabbit anti-CT-ET<sub>A</sub> antibody Dr.Jens Furkert, FMP-Berlin, Germany

rabbit anti-NT-ET<sub>B</sub> antibody Dr.Jens Furkert, FMP-Berlin, Germany

rabbit anti-CT-ET<sub>B</sub> antibody Dr.Jens Furkert, FMP-Berlin, Germany

## 2. Materials and Experimental procedures

#### 2.1 Material

## 2.1.1 Chemicals, antibodies, cDNA's and receptor ligands

Acetic acid Merck, Darmstadt, Germany Acrylamide, Bisacrylamide Serva, Heidelberg, Germany

Ammoniumpersulfat (APS) Sigma-Aldrich, Steinheim, Germany

Aprotinin Merck, Darmstadt, Germany
Bacitracin Merck, Darmstadt, Germany

Bovine serum albumine (BSA) Sigma-Aldrich, Steinheim, Germany

Disodium hydrogenphosphate Merck, Darmstadt, Germany

Dithiothreitol (DTT) Sigma-Aldrich, Steinheim, Germany

Dried milk powder (low-fat)

Néstle AG, Frankfurt, Germany

Dulbecco's modified Eagle's medium Sigma-Aldrich, Steinheim, Germany

Ethanol J.T. Baker, Deventin, The Netherlands

Ethylendiamine-tetraacetate (EDTA) Sigma-Aldrich, Steinheim, Germany

Ethylene glycol bis-tetraacetate (EGTA) Sigma-Aldrich, Steinheim, Germany

Fetal calf serum (FCS)

Biochrom, Berlin, Germany

FuGENE 6 Transfection Reagent Roche Diagnostics, Mannheim, Germany

Glucose Sigma-Aldrich, Steinheim, Germany
Glutamine Sigma-Aldrich, Steinheim, Germany
Glycerin Sigma-Aldrich, Steinheim, Germany

Glycine Carl Roth, Karlsruhe, Germany Hydrochloric acid Merck, Darmstadt, Germany

IGEPAL CA-630 (NP-40) Sigma-Aldrich, Steinheim, Germany
Isopropanol J.T. Baker, Deventin, The Netherlands

LipofectAMINE Invitrogen

Lumi-Light solution Roche Diagnostics, Mannheim, Germany
Methanol J.T. Baker, Deventin, The Netherlands
Monensin Calbiochem-Novabiochem, Bad Soden,

Germany

Peroxidase-conjugated goat anti-rabbit

antibodies (F<sub>ab</sub> fragments) Dianova, Hamburg, Germany

Peroxidase-conjugated donkey anti-mouse

antibodies (F<sub>ab</sub> fragments) Dianova, Hamburg, Germany

cDNAs, constructs:

Epidermal growth factor receptor kindly provided by Dr. Alexander Sorkin,

cDNA (HER1) Health Science Center, University

of Colorado, Denver, USA

K44A.dynamin I kindly provided by Dr. S. L. Schmid, The

Scripps Research Institute, La Jolla, CA

**Endothelin receptor ligands:** 

BQ123 Alexis, Läufelfingen, Switzerland

BQ788 Calbiochem-Novabiochem, Bad Soden

Germany

BQ3020 was synthesized using the solid phase

method, Calbiochem-Novabiochem, Bad

Soden, Germany

ET-1 was synthesized using the solid phase

method, Calbiochem-Novabiochem, Bad

Soden, Germany

ET-3 Calbiochem-Novabiochem, Bad Soden

Germany

<sup>125</sup>I-ET-1 (2000 Ci/mmol) Amersham Pharmacia Biotech, Freiburg,

Germany

<sup>125</sup>I-ET-3 (2000 Ci/mmol) Amersham Pharmacia Biotech, Freiburg,

Germany

<sup>125</sup>I-PD151242 (2000 Ci/mmol) Amersham Pharmacia Biotech, Freiburg,

Germany

## 2.1.2 Apparatus and software

AA-gel cast-stand BIORAD Laboratories

AA-gel electrophoresis chamber BIORAD Laboratories

Centrifuges Beckmann TLK 100

Beckmann Optima L70

Sorvall RC 285

Haereus Biofuge pico

Lumi-Imager F1 Roche Diagnostics

Microscopes Zeiss 510 META inverted confocal laser

scanning microscope

Zeiss Axiovert 100

Photometer GeneQuantII, Pharmacia Biotech

Pipettes Eppendorf

Power supplies Bio-Rad, Amersham Pharmacia Biotech

Rotator Stuart Scientific, Blood tube rotator SB1

Scintillator Wallac 1409

Semi-dry Western Blot chamber Trans Blot SD, Bio-Rad

Sonicator Sonoplus UW 2040, Bandelin

Electronics

Thermomixer Eppendorf Thermomixer 5436

## Software:

Axio Vision Zeiss

Excel 2000, Excel X Microsoft

GraphPad Prism 3.02 GraphPad software

Photoshop 5.0, Photoshop 4.0 Adobe

Powerpoint 2000, Powerpoint X Microsoft

Word 2000, Word X Microsoft

#### **2.1.3** Cells

HEK293 cells were obtained from the Deutsche Sammlung von Mikroorganismen und Zellkulturen, Braunschweig, Germany.

1 x blot buffer:

## 2.2 Commonly used buffers

10 x blot buffer:

112 g glycine 100 ml 10x blot buffer

22 g tris-base 200 ml methanol

ad 1 liter de-ionized H<sub>2</sub>O ad 1 liter de-ionized H<sub>2</sub>O

10 x gel running buffer: 1 x gel running buffer:

60 g tris-base 100 ml 10x gel running buffer

280 g glycine ad 1 liter de-ionized H<sub>2</sub>O

10 g SDS

## 4 x Laemmli sample buffer (Laemmli buffer):

10% β-mercaptoethanol (w/v)

ad 1 liter de-ionized H<sub>2</sub>O

4% SDS (w/v)

2% bromphenol blue (w/v)

20% glycerol (w/v)

250 mM Tris, pH 6.8

## Native lysis buffer:

20 mM Tris-HCl, pH 7.5

1% (w/v) IGEPAL CA-630 (NP-40)

1 mM ethylene glycol tetraacetic acid

1 mM ethylenediamine tetraacetic acid

1 mM dithiothreitol

0.5% (w/v) sodium deoxycholate

0.1% (w/v) SDS

1.5 mM MgCl<sub>2</sub>

150 mM NaCl

0.5 mM phenyl methyl sulforyl fluoride

2 mg/ml soy bean trypsin inhibitor

1.43 mg/ml aprotinin

0.5 mM benzamidine

PBS: PBST:

137 mM NaCl PBS

2.7 mM KCl 0.1% (w/v) SDS

1.5 mM KH<sub>2</sub>PO<sub>4</sub> 1% (w/v) IGEPAL CA-630

 $8.0 \text{ mM Na}_2\text{HPO}_4$  (NP-40)

pH 7.4

## **Tris-BAME:**

50 mM Tris

2 mM EGTA

10 mM MgCl<sub>2</sub>

0.15 mM Bacitracin

0.0015 % (w/v) Aprotinin

# 2.1.3 Plasmids/constructs

Plasmid/Vector	Encoded protein	Source
ET <sub>A</sub> myc.CFP	ET <sub>A</sub> receptor with an N-	Plasmid kindly provided
(pEGFP-N1, BD	terminal c-myc epitope	by Dr. Alexander
Biosciences, Heidelberg,	and a C-terminal CFP	Oksche, FMP Berlin,
Germany)	moiety.	Germany.
ET <sub>B</sub> .CFP/ ET <sub>B</sub> .GFP/	ET <sub>B</sub> receptor with a C-	Plasmids kindly
ET <sub>B</sub> .YFP	terminal CFP, GFP or	provided by Dr.
(pEGFP-N1, BD	YFP moiety.	Alexander Oksche, FMP
Biosciences, Heidelberg,		Berlin, Germany.
Germany)		
ET <sub>B</sub> flag.YFP	ET <sub>B</sub> receptor with an N-	Plasmid kindly provided
(pEGFP-N1, BD	terminal FLAG epitope	by Dr. Alexander
Biosciences, Heidelberg,	and a C-terminal YFP	Oksche, FMP Berlin,
Germany)	moiety.	Germany.
ET <sub>A</sub> myc	ET <sub>A</sub> receptor with an N-	Plasmid kindly provided
(pcDNA3.1, Invitrogen,	terminal c-myc epitope.	by Dr. Alexander
Leek, The Netherlands)		Oksche, FMP Berlin,
		Germany.
ET <sub>B</sub> flag	ET <sub>B</sub> receptor with an N-	Plasmid kindly provided
(pcDNA3.1, Invitrogen,	terminal FLAG epitope.	by Dr. Alexander
Leek, The Netherlands)		Oksche, FMP Berlin,
		Germany.

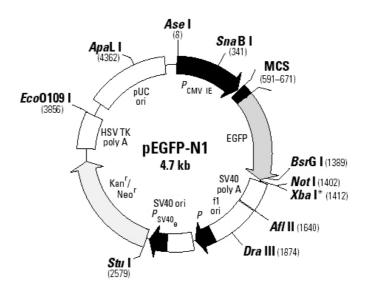


Fig. 4. Model of the pEGFP-N1 vector.

The figure above summarizes the features of the pEGFP-N1 vector.

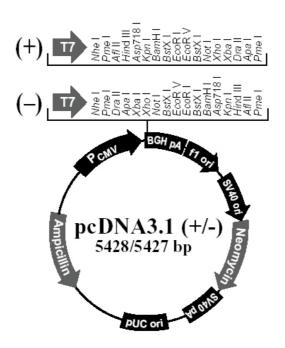


Fig. 5. Model of the pcDNA3.1 vector.

The figure above summarizes the features of the pcDNA3.1 (+) vector.

## 2.2 Experimental procedures

#### 2.2.1 Cell culture

HEK293 cells (obtained from Deutsche Sammlung von Mikroorganismen und Zellkulturen, Braunschweig, Germany) were maintained in DMEM medium supplemented with 10% fetal calf serum, 100 units/ml penicillin G, and 100 μg/ml streptomycin sulfate at 37°C in a humidified atmosphere of 95% air / 5% CO<sub>2</sub>. For fluorescence microscopy or laser scanning microscopy, cells were grown on glass cover slips for 48 h. For biochemical analyses, cells were grown for 48-72 h until 80% confluence was achieved.

## 2.2.2 Peptide synthesis and fluorescence labeling

ET-1 and BQ3020 (N-acetyl-[Ala<sup>11,15</sup>]-6-21-endothelin-1) were synthesized using the solid phase method (chlorotrityl-resin, 1.05 mmol/g; Calbiochem-Novabiochem) and standard 9-fluorenyl-methoxy-carbonyl chemistry (double couplings with 8 Eq of 9-fluorenyl-methoxy-carbonyl-amino acid derivates). After the final cleavage/deblocking, the crude peptide (50 mg) was dissolved in 500 ml of aqueous 4 mM NaHCO<sub>3</sub> solution and kept for two days at room temperature. The final purification was carried out by preparative HPLC (Polyencap A 300, 250 x 20 mm) applying a linear gradient 20 to 60% within 70 min [A, trifluoroacetic acid/water (0.1:99, v/v); B, trifluoroacetic acid/acetonitrile/water (0.1:80:19.9, v/v/v)]. The masses of purified ET-1 and BQ3020 were verified by electrospray mass spectrometry. Fluorescence labeling of ET-1 and BQ3020 was carried out by selective modification of the ε-amino group of Lys-9 of ET-1 and Lys-4 of BQ3020 using Cy3 monoreactive succinimidyl ester (Amersham Biosciences) in 0.1 M NaHCO<sub>3</sub> at pH 9.3 followed by preparative HPLC purification.

2.2.3 Transient and stable transfection of HEK293 cells

For transfertion, FuGENE6 was used according to the instructions of

the manufacturer (4 µl FuGENE 6/1 µg DNA). DNA if required was diluted in

sterile de-ionized water and pipetted into a reaction tube. FuGENE 6 was

diluted in serum- free media (in 100 µl per 25 mm or 700 µl per 92 mm culture

dish) and incubated for 5 min at room temperature before plasmid DNA was

added, followed by an incubation for 30 min at room temperature. Finally the

mix was added to cells at 50–80% confluency.

Stable transfection of HEK293 cells with LipoFECTAMINE:

1st. day: 200 000 cells were seeded in a 60 mm dish

2nd. day: 2 µg DNA was diluted in 100 µl serum-free medium without

antibiotics (SFM) and 7.5 µl LipoFECTAMINE was diluted in 100 µl SFM. Both

solutions were mixed and incubated for 30 min at room temperature. 1800 µl

SFM was added and the mixture was pipetted to cells (which were washed

before one time in SFM). Following incubation for 4 to 6 hours at 37°C the

medium was replaced by 5 ml of Dulbecco's modified Eagle's medium

supplemented with 10% fetal calf serum, 100 units/ml penicillin G, and 100

μg/ml streptomycin sulfate.

Cells were grown for 2-3 days before they were split in selection medium

containing G418 (400  $\mu$ g/ml) or Zeocin (17  $\mu$ g/ml).

The selection medium was changed every 3-4 days and after 10 days resistant

cell clones were picked with sterile Q-tips and transferred to 24 well plates

containing 1 ml selection medium per well. Cell clones were tested in binding

studies and/or fluorescence microscopy.

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2. Material and Experimental procedures

2.2.4 Generation and affinity-purification of polyclonal antibodies

A polyclonal NT-ET<sub>B</sub> serum was raised against a synthetic peptide

corresponding to the amino acids 19-37 in the N terminus of the ET<sub>B</sub> receptor

(P24530; CGLSRIWGEERGFPPDRATP) coupled to the carrier protein keyhole

limpet hemocyanin (KLH; Calbiochem-Novabiochem). NT-ET<sub>B</sub> antibody was

purified with the synthetic N-terminal peptide conjugated to protein Sepharose

6B (Pharmacia Biotech, Uppsala, Sweden) according to the manufacturers

protocol. The IgG fraction obtained was dialyzed against sodium phosphate

buffer (20 mM sodium phosphate buffer, 150 mM NaCl, pH 7.5) and stored in

aliquots at -20°C.

Buffers used for the affinity-purification of antibodies:

Binding buffer:

20 mM sodium phosphate buffer, pH 7-7.5

Elution buffer I:

0.2 M glycine

150 mM NaCl

pH 2.5

Elution buffer II:

0.2 M glycine

150 mM NaCl

pH 2.25

Strip buffer:

0.3 M NaHCO<sub>3</sub>

1 mM EDTA

pH 8.4

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Wash buffer: 0.1 M acetic acid

0.5 M NaCl

1 mM EDTA

50 mM HEPES, pH 8.0

PBS + 0.5% Tween 20

2M Tris-HCl, pH 9.0

Preparation of the sepharose for peptide binding:

250–500 mg thiopropyl sepharose 6B (Pharmacia Biotech AB, Uppsala, Sweden)

was mixed with 12 ml H<sub>2</sub>O in a 15 ml reaction tube and incubated on ice for 10

min. The sepharose was precipitated by centrifugation for 3 min at  $780 \times g$ .

Then the following washing steps were performed:

2 x 12 ml H<sub>2</sub>O

1 x 12 ml strip buffer.

After centrifugation the sepharose was incubated in 12 ml strip buffer for 40

min at room temperature on a rotator wheel, followed by two additional

washing steps with washing buffer. Prior to peptide binding the sepharose was

equilibrate with three washes of binding buffer (4°C).

Binding of the peptide:

5 mg of the peptide was dissolved in 400 μl sterile water and the pH was

titrated to pH 7-7.5 with 50 mM HEPES, pH 8.0. The sepharose was

resuspended in the peptide solution and incubated for 1 hour at 4°C in a rotator

wheel. The following wash steps were performed prior to the addition of the

antiserum:

1 x 12 ml binding buffer (4°C)

 $1 \times 12 \text{ ml}$  elution buffer, pH 2.5 (4°C)

3 x 12 ml binding buffer (4°C)

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## Binding of the antibodies:

5 to 10 ml of the antiserum were added to the sepharose and incubated in a rotator wheel at 4°C over night.

The following wash steps were performed:

5 x 10 ml PBS/0.5% Tween 20 (4°C)

Then the sepharose was dissolved in 10 ml PBS/0.5% Tween 20 and transfered to a column (BioRad) followed by an another wash with 10 ml PBS/0.5% Tween 20 prior to elution of the antibodies.

## Elution of the antibodies:

To elute the antibodies elution buffer I and elution buffer II were applied to the column as follows.

3x 1 ml elution buffer I (pH 2.5)

1x 4ml elution buffer I

3x 1 ml elution buffer II (pH 2.25)

The flow-through was collected in 1.5 ml reaction tubes containing 50  $\mu$ l 2 M Tris (pH 9) buffer to adjust the pH of the acidic eluate to pH 7–7.5.

## Concentration of the eluated antibodies:

To concentrate the antibodies the eluate (in 2 ml aliquots) was pipetted to Centricon-tubes (Centricon YM 30, Millipore, 30 kDa cut-off) and centrifuged at 4500 x g for 30 min at 4°C. Centricon concentrators can concentrate proteins on the basis of molecular weight. The buffer of the antibodies was changed by multiple centrifugation steps with binding buffer (approx. 10 ml). Finally the

concentrated antibodies were recovered by a reverse spin at 1000 x g for 15 min. at 4°C and 0.05% Na-acid was added.

## 2.2.5 Immunoblots for the detection of ET<sub>B</sub>.YFP and ET<sub>A</sub>myc.CFP

HEK293 cell clones stably expressing the fusion proteins were washed twice with PBS (137 mM NaCl, 2.7 mM KCl, 1.5 mM KH<sub>2</sub>PO<sub>4</sub>, 8.0 mM Na<sub>2</sub>HPO<sub>4</sub>, pH 7.4) and harvested with lysis buffer. To remove cell debris the lysates were centrifuged for 15 min at 14, 000 rpm. Lysates (30 µg protein per lane) were separated by SDS-PAGE (10% gels, 20 mA per gel) and transferred (100 mA per gel) to nitrocellulose filters (Schleicher und Schuell, Dassel, Germany). Filters were blocked for 1 h at room temperature in PBS/5% milk/5% FCS/0.05% Tween 20 and then probed for 1 h at room temperature or alternatively for 16 h at 4°C with polyclonal NT-ET<sub>B</sub> antibody (diluted 1 : 5000 in PBS/5% milk), monoclonal c-myc or monoclonal GFP antibodies (diluted 1 : 2000 in PBS/5% milk). Filters were then washed for 15 min with PBST (PBS + 0.1% SDS, 1% NP-40) followed by PBS wash and a final PBST wash. Primary antibodies were detected with horseradish peroxidase-conjugated goat anti-rabbit IgG or with horseradish peroxidase-conjugated donkey anti-mouse IgG (Jackson Immunoresearch Laboratories, USA). Secondary antibodies were diluted 1: 2000 in PBS + 5% milk, and filters were probed for 20 min at room temperature. Finally the filters were washed alternately for 15 min in PBST, PBS, PBS and antibodies were detected using Lumi-Light Western blotting substrate (Roche Diagnostics).

## 2.2.6 Immunoprecipitation experiments

HEK293 cell clones were grown in 75 cm $^2$  cell culture flasks for 48 h to near confluence. Cells were washed twice with PBS, and lysis buffer (1.2 ml) was added. The cells were then harvested with a rubber policeman, homogenized five times by passage through a 27G needle and centrifuged (800 x g) for 10 min

at 4°C. The supernatant was transferred to a new reaction tube and centrifuged (26,000 x g) for 30 min. The new supernatant was mixed with the NT-ET<sub>B</sub> antibody (diluted 1 : 2000 in PBS/5% milk) and protein A Sepharose (3.5 mg) in a final volume of 1.0 ml and incubated for 12 h in a shaker at 4°C. After 3 washes with lysis-buffer, the pellet was resuspended in 1 x Laemmli buffer and analyzed in immunoblot experiments.

## 2.2.7 Fluorescence resonance energy transfer (FRET)

FRET is a quantum mechanical process and is widely used as a biochemical tool to study protein-protein interactions. Excitation of a donor fluorophore can give rise to non-radiative transfer of the absorbed energy to an adjacent acceptor fluorophore provided that the emission spectrum of the donor overlaps sufficiently with the excitation spectrum of the acceptor. Direct energy transfer depends on the coupling of the respective dipole moments. Since the probability of energy transfer is proportional to orientation and inversely proportional to the sixth power of the distance  $(1/R^6)$  between the donor and the acceptor, FRET can be used to deduce the respective positional or rotational contexts of the two molecules. The positional constraints limit efficient FRET to distances of less than 10 nm and therefore sufficiently close to infer interaction. This makes FRET an ideal tool to study protein-protein interactions.

Cells were grown on glass coverslips for 48 h. Glass coverslips were mounted in a custom-made chamber and covered with incubation buffer (138 mM NaCl, 6 mM KCl, 1 mM MgCl<sub>2</sub>, 1 mM CaCl<sub>2</sub>, 5.5 mM glucose, 2 mg/ml BSA and 10 mM N-2-Hydroxyethylpiperazine-N'-2-ethanesulfonic acid (Hepes), pH 7.5). FRET analysis was performed as described (Brock et al. 2003), using an inverted microscope Axiovert 100 equipped with a Plan-Apochromat 63x/1.4 objective (both Carl Zeiss, Göttingen, Germany). In brief, CFP and YFP were alternately excited at 410 and 515 nm with a monochromator (Polychrome II; TILL Photonics, Gräfelfing, Germany) in combination with a dual reflectivity dichroic mirror (<460 nm and 500-520 nm; Chroma Technology, Rockingham,

VT, USA). Emitted light was filtered through 475-505 nm (CFP) or 535-565 nm (YFP) band pass filters changed by a motorized filter wheel (Lambda 10/2; Sutter Instruments, Novato, USA), and detected with a cooled CCD camera (Imago; TILL Photonics). For FRET analysis, CFP (40 ms) and YFP (8 ms) emission without acceptor bleach were recorded during 20 cycles with a 10-20 ms exposure per cycle. Then, 60 cycles were recorded with an additional 2 s illumination per cycle at 512 nm to bleach YFP. FRET was assessed as recovery of CFP (donor) fluorescence during YFP (acceptor) bleach. The FRET efficiency E can be expressed as  $E = 1 - (F_{DA} / F_{D})$ , where  $F_{DA}$  and  $F_{D}$  are intensities of the donor fluorescence in the presence and in the absence of the FRET acceptor, respectively. The increase in CFP fluorescence during acceptor photobleaching was plotted against the remaining YFP fluorescence, yielding a regression line, which served to extrapolate  $F_{DA}$  and  $F_{D}$ .

The influence of ligands on FRET efficiency was investigated by incubation of cells with 50 nM BQ3020, BQ788, BQ123 or ET-1 for 5 or 30 min at 37°C prior to FRET analysis. To clamp the pH at 7.5 or pH 5.0 in control experiments, monensin (10  $\mu$ M) and nigericin (10  $\mu$ M) were added. To block ligand-induced receptor sequestration, cells were co-transfected with ET<sub>A</sub>myc.CFP, ET<sub>B</sub>.YFP and K44A.dynamin or incubated with 450 mM sucrose.

## 2.2.8 Receptor sequestration assay

HEK293 cell clones (40,000 cells/well) were seeded into 96-well plates. After 24 h, cells were stimulated with ET-1 (50 nM) or BQ3020 (50 nM) for up to 2 h at 37°C. The cells were then washed twice with PBS and fixed for 15 min at room temperature (2.5% formaldehyde in 100 mM sodium cacodylate, 100 mM sucrose, pH 7.5). After two rinses with PBS, monoclonal peroxidase-conjugated FLAG antibody (diluted 1 : 2000, Sigma) was added for 1 h at 37°C. Cells were washed twice with PBS and stained with 95  $\mu$ l reaction buffer (200 mM citric acid, 200 mM K<sub>2</sub>HPO<sub>4</sub>, 0.01% H<sub>2</sub>O<sub>2</sub>) and 5  $\mu$ l/well substrate solution (20 mM tetramethylbenzidine in propanol) for 15 min at room temperature. The

reaction was terminated with 100  $\mu$ l/well H<sub>2</sub>SO<sub>4</sub> (1.5 M). The absorbance at 450 nm was determined in an ELISA bio-kinetiks reader (Bio-Tek instruments, Winooski, VT, USA).

## 2.2.9 Fluorescence microscopy and image analysis

Living HEK293 cells grown on glass coverslips were analyzed with a Zeiss 510 META inverted confocal laser scanning microscope equipped with a Plan-Apochromat 63x/1.4 objective. Excitation and emission wavelengths were:  $\lambda_{exc}$  = 458 nm and  $\lambda_{em}$  = 470-505 nm for CFP,  $\lambda_{exc}$  = 488 nm and  $\lambda_{em}$  = 515-550 nm for YFP, and  $\lambda_{exc}$  = 543 nm, and  $\lambda_{em}$  > 570 nm for Cy3.

## 2.2.10 <sup>125</sup>I-ET-1 displacement binding experiments

Radioligand experiments were performed as described (Oksche et al. 2000). In brief, membranes (0.1-0.5  $\mu$ g) were incubated in 200  $\mu$ l Tris/BAME buffer containing 50 pM of  $^{125}\text{I-ET-1}$  without or with increasing concentrations of unlabelled ligand (1 x 10-13 to 1 x 10-4 M) for 3 h at 25°C in a shaking water bath. The samples were then transferred to GF/C filters (Whatman, Maidstone, UK) pretreated with 0.1% (w/v) polyethylenimine and washed twice with PBS using a Brandel cell harvester. Filters were finally transferred into 5 ml vials and radioactivity was determined in a  $\gamma$ -counter. Data were analyzed with RadLig Software 4.0 (Cambridge, UK), and graphs were generated with Prism Software 2.01 (GraphPad, San Diego, USA).