

Partial agonist behaviour depends upon the level of nociceptin/orphanin FQ receptor expression: studies using the ecdysone-inducible mammalian expression system

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1 Partial agonism is primarily dependent upon receptor density and coupling efficiency. As these parameters are tissue/model dependent, intrinsic activity in different tissues can vary. We have utilised the ecdysone-inducible expression system containing the human nociceptin/orphanin FQ (N/OFQ) peptide receptor (hNOP) expressed in Chinese hamster ovary cells (CHO_{INDhNOP}) to examine the activity of a range of partial agonists in receptor binding, GTP γ ³⁵S binding and inhibition of adenylyl cyclase studies.

2 Incubation of CHO_{INDhNOP} cells with ponasterone A (PON) induced hNOP expression ([leucyl-³H]N/OFQ binding) of 24, 68, 191 and 1101 fmol mg⁻¹ protein at 1, 2, 5 and 10 μ M PON, respectively. At 191 fmol mg⁻¹, protein hNOP pharmacology was identical to that reported for other traditional expression systems.

3 pEC₅₀ values for GTP γ ³⁵S binding ranged from 7.23 to 7.72 (2–10 μ M PON) for the partial agonist [Phe¹ ψ (CH₂-NH)Gly²]N/OFQ(1–13)-NH₂ ([F/G]N/OFQ(1–13)-NH₂) and 8.12–8.60 (1–10 μ M PON) for N/OFQ(1–13)-NH₂ and E_{max} values (stimulation factor relative to basal) ranged from 1.51 to 3.21 (2–10 μ M PON) for [F/G]N/OFQ(1–13)-NH₂ and 1.28–6.95 (1–10 μ M) for N/OFQ(1–13)-NH₂. Intrinsic activity of [F/G]N/OFQ(1–13)-NH₂ relative to N/OFQ(1–13)-NH₂ was 0.3–0.5. [F/G]N/OFQ(1–13)-NH₂ did not stimulate GTP γ ³⁵S binding at 1 μ M PON, but competitively antagonised the effects of N/OFQ(1–13)-NH₂ with a pK_B = 7.62.

4 pEC₅₀ values for cAMP inhibition ranged from 8.26 to 8.32 (2–10 μ M PON) for [F/G]N/OFQ(1–13)-NH₂ and 9.42–10.35 for N/OFQ(1–13)-NH₂ and E_{max} values (% inhibition) ranged from 19.6 to 83.2 for [F/G]N/OFQ(1–13)-NH₂ and 40.9–86.0 for N/OFQ(1–13)-NH₂. The intrinsic activity of [F/G]N/OFQ(1–13)-NH₂ relative to N/OFQ(1–13)-NH₂ was 0.48–0.97.

5 In the same cellular environment with receptor density as the only variable, we show that the profile of [F/G]N/OFQ(1–13)-NH₂ can be manipulated to encompass full and partial agonism along with antagonism.

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Abbreviations: CHO_{hNOP}, Chinese hamster ovary cells expressing human NOP; CHO_{INDhNOP}, Chinese hamster ovary cells expressing the ecdysone-inducible mammalian expression system containing the human NOP; [F/G]N/OFQ(1–13)-NH₂, [Phe¹ ψ (CH₂-NH)Gly²]N/OFQ(1–13)-NH₂; NalBzOH, naloxone benzoylhydrazone; N/OFQ, nociceptin/orphanin FQ; NOP, N/OFQ peptide receptor.

Introduction

Nociceptin/orphanin FQ (N/OFQ) is the endogenous peptide ligand for the G_i-coupled N/OFQ peptide receptor (NOP). The terminology used in this paper with respect to nomenclature is in line with recent IUPHAR guidelines (Cox *et al.*, 2000). At a cellular level, N/OFQ causes a reduction in cAMP formation, activation of potassium channels and inhibition of voltage-gated calcium channels, thereby reducing neuronal excitability and inhibiting transmitter release (Meunier *et al.*, 1995; Reinscheid *et al.*, 1995; Knoflach *et al.*, 1996; Vaughan &

Christie, 1996; Hawes *et al.*, 2000; Schlicker & Morari, 2000; Jennings, 2001; New & Wong, 2002). Central administration of N/OFQ has been shown to cause analgesia, hyperalgesia and allodynia, hypotension and bradycardia, diuresis and antinatriuresis and have anxiolytic properties (Calo *et al.*, 2000b; Meunier, 2000; Mogil & Pasternak, 2001).

As a result of structure–activity relationship studies and combinatorial library screens, several selective, potent agonists and antagonists have been described and these molecules have greatly enhanced our understanding of the physiological role(s) of the N/OFQ-NOP system. These molecules include the peptides [(pF)Phe⁴]N/OFQ(1–13)-NH₂, [Phe¹ ψ (CH₂-NH)Gly²]N/OFQ(1–13)-NH₂ ([F/G]N/OFQ(1–13)-NH₂), [Nphe¹]N/OFQ(1–13)-NH₂, UFP-101 and nonpeptides

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Ro65-6570/Ro64-6198 (agonists) and J-113397/JTC-801 (antagonists) (Guerrini *et al.*, 1998; 2000; Jenck *et al.*, 2000; Ozaki *et al.*, 2000; Bigoni *et al.*, 2002b; Calo *et al.*, 2002; McDonald *et al.*, 2002; Yamada *et al.*, 2002). From a combinatorial library of 52 million, Dooley *et al.* (1997) identified five hexapeptides with high affinity for NOP (Berger *et al.*, 2000a). Functionally, these peptides are all partial agonists with varying degrees of efficacy. The opioid antagonist naloxone benzoylhydrazone (NalBzOH) has been shown to possess low partial agonist activity at NOP, but activity at classical opioid receptors limits the use of this synthetic compound (Nicholson *et al.*, 1998; Bigoni *et al.*, 2002a).

Initial studies with [F/G]N/OFQ(1–13)–NH₂ in the mouse was deferens and guinea-pig ileum indicated that the peptide behaved as an NOP-selective antagonist (Guerrini *et al.*, 1998). However, subsequent studies reported variable intrinsic activity of this molecule from zero (pure antagonism) to one (full agonism) (Calo *et al.*, 1998; Grisel *et al.*, 1998; Meis & Pape, 1998; Menzies *et al.*, 1999; Okawa *et al.*, 1999; Mason *et al.*, 2001; Calo *et al.*, 2000a). Consensus is that [F/G]N/OFQ(1–13)–NH₂ is in fact a low-efficacy partial agonist (Calo *et al.*, 2000b). However, there are no detailed studies on NOP from one species at differing levels of expression using an identical cell background that specifically addresses this question.

To date, NOP has been expressed in a variety of mammalian cell lines including CHO and HEK-293 (Guerrini *et al.*, 2000; Dautzenberg *et al.*, 2001), which either utilise a transient expression strategy (hence relatively uncontrolled expression) or are used to generate stable clones (usually with high levels of expression). It would be desirable, with particular reference to the evaluation of agonist intrinsic activity, to have a range of lines available with differing levels of receptor expression. The ecdysone-inducible expression system represents a simple method allowing the production of cultures with differing expression levels of a receptor of interest. In this system, addition of ponasterone A (an ecdysone analogue) will produce a concentration-dependent increase in transcription and hence receptor expression. This has been demonstrated for sst2 (Cole *et al.*, 2001), 5-HT (h5-HT 1B, 1F, 4B) (Van Craenenbroeck *et al.*, 2001) and DOP receptors (Law *et al.*, 2000).

In this study, we have utilised the ecdysone-inducible expression system containing the hNOP receptor to examine the activity of a range of partial agonist molecules. Initially, we have characterised the system with reference to ponasterone A induction concentrations in (a) receptor binding, (b) GTP γ ³⁵S binding and (c) cAMP assays. A pharmacological characterisation of the receptor expressed at levels similar to those typically obtained in saturation studies of rat cerebrocortical membranes (~ 180 fmol mg⁻¹ protein (Okawa *et al.*, 1998)) is presented. Finally, we have examined the effects of a range of partial agonists including [F/G]N/OFQ(1–13)–NH₂, Ac-RYYRIK–NH₂ and Ac-RYYRWK–NH₂, and NalBzOH at differing levels of NOP receptor expression.

Methods

Sources of materials

N/OFQ, N/OFQ–NH₂, N/OFQ(1–13)–NH₂, [F/G]N/OFQ(1–13)–NH₂, J-113397, Ac-RYYRIK–NH₂ and

Ac-RYYRWK–NH₂ were synthesised at the Department of Pharmaceutical Sciences at the University of Ferrara as described previously (Guerrini *et al.*, 1997; De Risi *et al.*, 2001; Rizzi *et al.*, 2002). NalBzOH was purchased from Sigma (Poole, U.K.). Radioactivity, [*leucyl*-³H]N/OFQ (149–152 Ci mmol⁻¹) was obtained from Amersham Pharmacia Biotech (Buckinghamshire, U.K.), GTP γ ³⁵S (1250 Ci mmol⁻¹) and [2,8-³H]cAMP (28 Ci mmol⁻¹) were obtained from NEN DuPont (Boston, MA, U.S.A.). Pertussis toxin was obtained from Sigma. Ponasterone A, zeocin, all tissue culture media and supplements were obtained from Invitrogen (Paisley, U.K.). Chinese hamster ovary (CHO) cells containing the ecdysone-inducible expression system with the hNOP and CHO cells stably expressing the hNOP were kindly provided by Dr F. Marshall and Mrs N. Bevan of GSK (Stevenage, Herts, U.K.).

The ecdysone-inducible mammalian expression system

The system is based on the induction, molting and metamorphosis process utilised by *Drosophila*, which activates gene expression through the ecdysone receptor. This system has been modified for use in mammalian cell lines so that a chosen gene can be expressed by the application of a steroid promoter. The ecdysone system makes use of a heterodimeric nuclear receptor consisting of the ecdysone receptor (VgEcR) and the retinoid X receptor (RXR, modified from mammalian cells), which bind a hybrid response element (E/GRE) in the presence of the synthetic analogue of ecdysone, ponasterone A. The hybrid response element lies upstream of a minimal heat shock promoter, activation of which leads to transcription of the gene of interest (in this study, human NOP).

Cell culture and induction

CHO cells stably expressing the ecdysone-inducible mammalian expression system containing the hNOP (CHO_{INDhNOP}) were cultured in HAMS F12 supplemented with 10% foetal calf serum, penicillin (100 IU ml⁻¹), streptomycin (100 μ g ml⁻¹) and fungizone (2.5 μ g ml⁻¹). Stock cultures were further supplemented with geneticin (1 mg ml⁻¹) and zeocin (250 μ g ml⁻¹). CHO cells stably expressing the hNOP receptor (CHO_{hNOP}) were routinely cultured as described (Hashiba *et al.*, 2001). All cultures were maintained at 37°C with 5% carbon dioxide humidified air and subcultured as required using trypsin/EDTA. Cells were induced as they approached confluence for 20 h with the steroid ponasterone A, at concentrations of 1, 2, 5, 10 and 20 μ M. Non-induced CHO_{INDhNOP} cultures were used as negative controls in which the medium was replaced 20 h prior to use. In studies using pertussis toxin (PTx), 100 ng ml⁻¹ was added to media at the time of induction.

Membrane preparation

Membranes were prepared from freshly harvested cells, CHO_{hNOP} at confluence and CHO_{INDhNOP} 20 h postmedia change/induction. Cells were suspended in a homogenisation buffer of either Tris-HCl (50 mM), MgSO₄ (5 mM) pH 7.4 with KOH (saturation and displacement) or Tris-HCl (50 mM), EGTA (0.2 mM) pH 7.4 with NaOH (GTP γ ³⁵S). Suspensions were homogenised followed by centrifugation at 13,500 rpm,

for 10 min at 4°C. This was repeated three times in total. The membrane pellet was finally resuspended as appropriate for each experiment, the protein concentration was determined (Lowry *et al.*, 1951) and finally adjusted as required for the experimental procedure.

Saturation binding

The membrane protein (15–350 µg) (depending on induction level) was incubated in 0.5 ml of homogenisation buffer containing 0.5% BSA, 10 µM peptidase inhibitors (amastatin, bestatin, captopril, phosphoramidon) and various concentrations of [*leucyl*-³H]N/OFQ (~2 nM–0.002 µM) for 1 h at room temperature. Nonspecific binding (NSB) was defined in the presence of 1 µM unlabelled N/OFQ. Reactions were terminated by vacuum filtration through polyethylenimine (PEI) (0.5%)-soaked Whatman GF/B filters using a Brandel harvester.

Displacement binding

The membrane protein (35–70 µg) was incubated in the buffer used in saturation assays, but containing a fixed concentration of [*leucyl*-³H]N/OFQ (~200 pM) and varying concentrations of a range of displacers. NSB was determined in the presence of 1 µM unlabelled N/OFQ. Assays were incubated at room temperature for 1 h and reactions terminated *via* filtration through PEI (0.5%)-soaked Whatman GF/B filters using a Brandel harvester.

GTPγ³⁵S assays

CHO_{INDhNOP} (40 µg) or 20 µg CHO_{hNOP} membranes were incubated in 0.5 ml buffer containing Tris (50 mM), EGTA (0.2 mM), MgCl₂ (1 mM), NaCl (100 mM) BSA (1 mg ml⁻¹) pH 7.4 with NaOH to which bacitracin (0.15 mM), amastatin, bestatin, captopril and phosphoramidon (10 µM); GDP (5 µM/100 µM) and ~150 pM GTPγ³⁵S were added. NOP ligands were included in various combinations and at various concentrations. NSB was determined in the presence of 10 µM GTPγS. All receptor ligands were omitted when defining basal and NSB binding of GTPγ³⁵S. Reactions were incubated for 1 h at 30°C with gentle shaking and terminated by filtration through Whatman GF/B filters using a Brandel harvester.

In all cases, radioactivity was determined following filter extraction (8 h, Optiphase Safe, Wallac) using liquid scintillation spectroscopy.

Inhibition of forskolin-stimulated cAMP formation

Inhibition of forskolin-stimulated cAMP formation was measured using whole CHO cells induced at 1, 2, 5 and 10 µM ponasterone A. Confluent adherent cell cultures (grown in 24-well tissue culture trays) were incubated in the presence of 1 mM isobutylmethylxanthine (IBMX) and forskolin (1 µM) for 15 min. NOP ligands were included in various combinations and at different concentrations. Reactions were terminated using 10 M HCl and neutralised with 10 M NaOH/1 mM Tris, pH 7.4. The concentration of cAMP was measured using the protein-binding method set out by Brown *et al.* (1971).

Analysis of data

All data are expressed as mean ± s.e.m., from a minimum of three experiments performed as single points or in duplicate. Concentration–response curves and statistical analyses (paired/unpaired Students' *t*-test and ANOVA with Bonferroni correction for multiple comparison where appropriate) were performed using PRISM V3.0 (GraphPad, San Diego, U.S.A). p*K*_i values were calculated using the Cheng & Prusoff equation (log{IC₅₀/(1 + [Radiolabel]/K_D)} (Cheng & Prusoff, 1973). A K_D of 60.3 pM for [*leucyl*-³H]N/OFQ, measured from saturation binding using 5 µM-induced membranes was used. p*K*_B values were calculated using the formula p*K*_B = -log{(CR-1)/[antagonist]}, where CR is the ratio of the EC₅₀ of the agonist in the presence and absence of antagonist, assuming a slope value of unity. In GTPγ³⁵S binding studies, data are either presented as DPM ³⁵S bound (in studies where the GDP concentration is varied, as 'stimulation factor' is GDP dependent) or stimulation factor (i.e. the ratio between specific agonist-stimulated GTPγ³⁵S binding and basal specific binding). cAMP data are presented as percentage inhibition of the forskolin-stimulated response.

Results

Saturation binding assays

Incubation of CHO_{INDhNOP} cells with ponasterone A induced the expression of NOP, as measured by the binding of [*leucyl*-³H]N/OFQ. The total specific binding of [*leucyl*-³H]N/OFQ increased from 24 to 1101 fmol mg⁻¹ protein as the concentration of ponasterone A was increased (1–10 µM) (Table 1). In non-induced cultures, there was no significant specific binding despite the use of large quantities of membrane protein. Interestingly, the induction–expression relationship appeared to be bell-shaped, such that an apparent maximum was obtained at 10 µM ponasterone A, above which (i.e. 20 µM ponasterone A) binding decreased. In a simple series of trypan blue exclusion experiments (*n* = 6, data not shown), 20 µM ponasterone A did not cause any significant cytotoxicity.

The expression of NOP at 5 µM ponasterone A induction (~200 fmol mg⁻¹ protein) is similar to that measured in brain tissues, for example, in rat cerebral cortex membranes

Table 1 The binding of [*leucyl*-³H]N/OFQ to CHO_{INDhNOP} was ponasterone A dependent

Induction (ponasterone A) (µM)	p <i>K</i> _D	K _D (pM)	B _{max} (fmol mg protein ⁻¹)
0 (noninduced)	—	—	—
1	9.91 ± 0.04	123	23.5 ± 4.4
2	9.83 ± 0.09	148	68.3 ± 9.7
5	10.22 ± 0.15	60	190.6 ± 25.5
10	9.89 ± 0.14	129	1101.0 ± 145.3
20	9.89 ± 0.13	129	191.2 ± 33.9

Saturation analysis of log-transformed specific data was used to estimate B_{max} and p*K*_D. Data are mean ± s.e.m. for *n* ≥ 3 experiments.

(179.7 fmol [125 I]Tyr 14 -N/OFQ mg $^{-1}$ protein) (Okawa *et al.*, 1998; Hashiba *et al.*, 2001) and so this induction level has been used to perform a series of displacements and GTP γ^{35} S/cAMP studies in order to detail the pharmacology of the induced hNOP receptor.

Displacement binding assays

The binding of a fixed concentration of [*leucyl*- 3 H]N/OFQ was displaced in a concentration-dependent and saturable manner by a range of NOP peptide and non-peptide ligands in membranes prepared from CHO_{INDhNOP} cells induced with 5 μ M ponasterone A. pK_i values for these data are summarised in Table 2. The rank order pK_i is N/OFQ-NH $_2$ > N/OFQ(1-13)-NH $_2$ > N/OFQ = Ac-RYYRWK-NH $_2$ > [F/G]N/OFQ(1-13)-NH $_2$ > Ac-RYYRIK-NH $_2$ = J-113397 > NalBzOH.

GTP γ^{35} S and cAMP functional data

Initially, we made a comparison of GTP γ^{35} S binding stimulated by N/OFQ(1-13)-NH $_2$ and [F/G]N/OFQ(1-13)-NH $_2$ in CHO_{hNOP} and CHO_{INDhNOP} cells (5 μ M ponasterone A induction) at high (100 μ M) and low (5 μ M) concentrations of GDP. It has been hypothesised that low-efficacy (partial) agonists activate G proteins with vacant guanine nucleotide-binding sites more effectively. Hence, reducing the GDP concentration should lead to fewer occupied guanine nucleotide-binding sites and result in an increased intrinsic activity (Breivogel *et al.*, 1998; Berger *et al.*, 2000b; Bigoni *et al.*, 2002a).

The stable CHO_{hNOP} cell line used here expresses \sim 1.9 pmol mg $^{-1}$ protein NOP and has been used extensively by us in the past (Okawa *et al.*, 1998; Guerrini *et al.*, 2000; 2001; Hashiba *et al.*, 2001). At 100 μ M GDP, both N/OFQ(1-13)-NH $_2$ and [F/G]N/OFQ(1-13)-NH $_2$ stimulated GTP γ^{35} S binding to CHO_{hNOP} membranes in a concentration-dependent and saturable manner with pEC_{50} values of 9.11 and 8.28, respectively. N/OFQ(1-13)-NH $_2$ was a full agonist, E_{max} 10,117 net DPM, while [F/G]N/OFQ(1-13)-NH $_2$ displayed

partial agonist activity in this preparation with an E_{max} of 6221 net DPM (Figure 1, Table 3). At 5 μ M GDP, the potency of both [F/G]N/OFQ(1-13)-NH $_2$ (pEC_{50} 8.68) and N/OFQ(1-13)-NH $_2$ (pEC_{50} 9.57) increased as did the net stimulated binding of GTP γ^{35} S, 24,255 and 16,999 DPM, respectively. More importantly, the relative intrinsic activity of [F/G]N/OFQ(1-13)-NH $_2$ compared to N/OFQ(1-13)-NH $_2$ increased from 0.61 to 0.70. The same is true for N/OFQ(1-13)-NH $_2$ and [F/G]N/OFQ(1-13)-NH $_2$ using membranes from 5 μ M ponasterone A-induced cells (Table 3) with relative intrinsic activity increasing from 0.38 to 0.81. A similar increase in relative intrinsic activity can also be observed using Ac-RYYRWK-NH $_2$ and NalBzOH, the latter having been previously published (Bigoni *et al.*, 2002a).

In membranes prepared from CHO_{INDhNOP} cells incubated with 1, 2, 5 and 10 μ M ponasterone A, both N/OFQ(1-13)-NH $_2$ and [F/G]N/OFQ(1-13)-NH $_2$ stimulated the binding of GTP γ^{35} S in a concentration-dependent and saturable manner (Figure 2). As the induction concentration of ponasterone A was increased, the E_{max} (stimulation factor) of N/OFQ(1-13)-NH $_2$ increased from 1.28 (1 μ M) to 6.95 (10 μ M; see Table 4). The E_{max} of [F/G]N/OFQ(1-13)-NH $_2$ also increased as a function of the induction concentration, from 0.98, that is, basal (1 μ M) to 3.21 (10 μ M). However, the relative intrinsic activity of [F/G]N/OFQ(1-13)-NH $_2$ (relative to N/OFQ(1-13)-NH $_2$) remained similar at 0.37–0.55 for all induction levels.

In cAMP inhibition studies, the E_{max} of both N/OFQ(1-13)-NH $_2$ and [F/G]N/OFQ(1-13)-NH $_2$ also varied as a function of the induction concentration, from 41 to 86% and from 20 to 83% at low and high ponasterone A induction, respectively (Figure 3). The relative intrinsic activity of [F/G]N/OFQ(1-13)-NH $_2$ changed from 0.48 to 0.97 (Table 4), indicating that at 10 μ M ponasterone A this molecule, in this assay, behaved as a full agonist. At the lower 1 μ M ponasterone A induction, due to low expression of hNOP and sensitivity of this assay, data for cAMP studies could not be reliably analysed. These and GTP γ^{35} S binding data are shown as a function of receptor density in Figure 4.

Table 2 pK_i values for a range of NOP ligands measured in CHO_{INDhNOP} membranes induced with 5 μ M ponasterone A

Ligand Agonist	Class		pK_i
	Peptide (P) non-peptide (NP)		
N/OFQ	P		9.93 \pm 0.08
N/OFQ-NH $_2$	P		10.37 \pm 0.04
N/OFQ(1-13)-NH $_2$	P		10.35 \pm 0.04
Presumed partial agonists			
NalBzOH	NP		7.1 \pm 0.02
[F/G]N/OFQ(1-13)-NH $_2$	P		9.6 \pm 0.1
Ac-RYYRIK-NH $_2$	P		9.12 \pm 0.02
Ac-RYYRWK-NH $_2$	P		9.99 \pm 0.03
Antagonist			
J-113397	NP		9.09 \pm 0.11

pK_i values were calculated using the Cheng and Prusoff equation using a K_D of 60.3 pM for [*leucyl*- 3 H]N/OFQ, measured in saturation experiments for the same induction (Table 1). Data are mean \pm s.e.m. ($n = 4$).

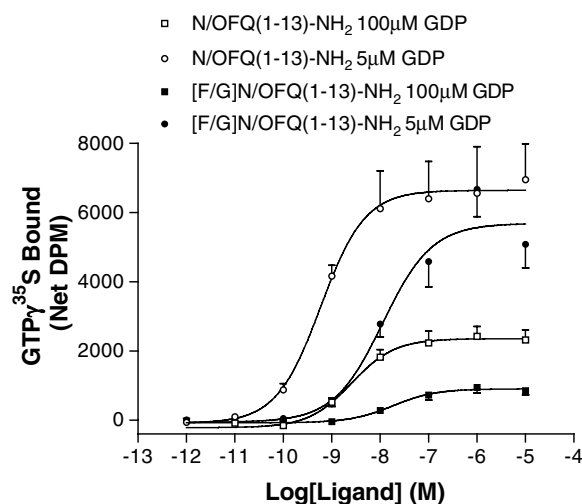


Figure 1 Net GTP γ^{35} S binding by N/OFQ(1-13)-NH $_2$ and [F/G]N/OFQ(1-13)-NH $_2$ in membranes from CHO_{INDhNOP} (5 μ M ponasterone A induction) with 100 and 5 μ M GDP. Data are mean \pm s.e.m. for $n = 4$.

Table 3 Effects of [F/G]N/OFQ(1–13)–NH₂ Ac-RYYRWK–NH₂ and NalBzOH on GTP γ ³⁵S binding in CHO_{hNOP} and CHO_{INDhNOP} (5 μ M ponasterone A) membranes in the presence of either 100 or 5 μ M GDP

	CHO _{hNOP}		CHO _{INDhNOP}		CHO _{hNOP}		CHO _{hNOP} (Bigoni <i>et al.</i> , 2002a)	
	N/OFQ(1–13)	F/G(1–13)	N/OFQ(1–13)	F/G(1–13)	N/OFQ	RYYRWK	N/OFQ	NalBzOH
<i>100 μM GDP</i>								
pEC ₅₀	9.11 ± 0.03	8.28 ± 0.01	8.65 ± 0.07	7.72 ± 0.05	8.30 ± 0.04	8.97 ± 0.03	8.53 ± 0.12	Inactive
E _{max} (DPM)	10117 ± 134	6221 ± 374	2368 ± 301	908 ± 104	9548 ± 86	5291 ± 78	4140 ± 40 ^a	Inactive
α	1.00	0.61	1.00	0.38	1.00	0.55	1.00	0
<i>5 μM GDP</i>								
pEC ₅₀	9.57 ± 0.02*	8.68 ± 0.12*	9.27 ± 0.12*	8.02 ± 0.03*	8.70 ± 0.01*	9.34 ± 0.09*	9.29 ± 0.02*	7.00 ± 0.10
E _{max} (DPM)	24255 ± 964*	16999 ± 242*	6635 ± 1080*	5388 ± 834*	14894 ± 317*	11775 ± 130*	18190 ± 65*	2278 ± 238
α	1.00	0.70	1.00	0.81	1.00	0.79	1.00	0.13

N/OFQ(1–13)–NH₂ and N/OFQ were used as reference full agonists. Data derived from PRISM-Fits as in Figure 1 and are mean \pm s.e.m. ($n \geq 3$). * $P < 0.05$ (paired t -test) increased pEC₅₀ or E_{max} compared with 100 μ M.

^aWhile it appears that there is some variation in the E_{max} (DPM) for N/OFQ and N/OFQ(1–13)NH₂ for CHO_{hNOP} cells reported in Bigoni *et al.* (2002a) relative to this study, it should be borne in mind that the study of Bigoni *et al.* (2002a) was performed with different batches of cells and GTP γ ³⁵S producing lower net DPM for basal, NSB and stimulated binding. However, the stimulation factors for N/OFQ(1–13)NH₂ in this study (100 μ M GDP; 10.26 \pm 0.72) are comparable to those for N/OFQ (100 μ M GDP; 8.12 \pm 0.29) in Bigoni *et al.* (2002a).

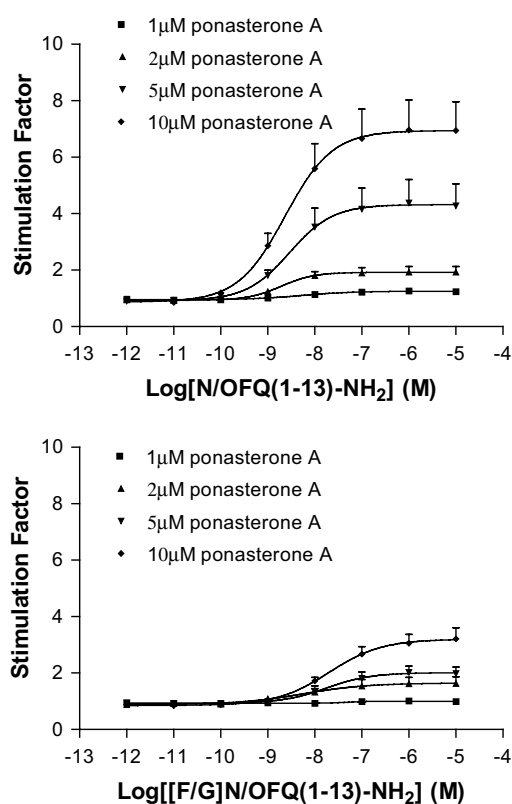


Figure 2 N/OFQ(1–13)–NH₂ (upper panel) and [F/G]N/OFQ(1–13)–NH₂ (lower panel)-stimulated GTP γ ³⁵S binding to membranes prepared from CHO_{INDhNOP} cells induced with 1, 2, 5 and 10 μ M ponasterone A. Data are mean \pm s.e.m. for $n \geq 4$.

Pertussis toxin (PTx) sensitivity

CHO_{INDhNOP} cells were induced for 20 h at 5 μ M ponasterone A in the absence and presence of PTx (100 ng ml⁻¹). Membrane fragments or whole cells were then tested for their ability to stimulate the binding of GTP γ ³⁵S or inhibit cAMP formation by a range of NOP-selective agonists (Table 5). While in this series of experiments the degree of GTP γ ³⁵S stimulation and inhibition of cAMP was slightly reduced, PTx treatment

clearly prevented agonist-stimulated GTP γ ³⁵S binding and CAMP inhibition by N/OFQ, N/OFQ(1–13)–NH₂, [F/G]N/OFQ(1–13)–NH₂ and confirms NOP action through either a G_i and/or G_o in CHO_{INDhNOP} cells.

In CHO_{INDhNOP} induced at 5 μ M ponasterone A, we examined the behaviour of a range of other NOP ligands including Ac-RYYRIK–NH₂, Ac-RYYRWK–NH₂ and NalBzOH. Both N/OFQ and N/OFQ–NH₂ produced concentration-dependent and saturable increases in the binding of GTP γ ³⁵S (Table 6). Both were full agonists since E_{max} values did not vary significantly from one another or N/OFQ(1–13)–NH₂. In the GTP γ ³⁵S assay at the same induction level, both Ac-RYYRIK–NH₂ and Ac-RYYRWK–NH₂ were clear partial agonists (E_{max} 1.66 \pm 0.02 and 2.16 \pm 0.08, respectively) with relative intrinsic activity values not significantly different from that of [F/G]N/OFQ(1–13)–NH₂ (Table 6). NalBzOH produced no measurable stimulation of GTP γ ³⁵S binding up to 100 μ M in membranes from cells induced at 1–10 μ M ponasterone A and is therefore classed as an antagonist in this assay system (Okawa *et al.*, 1999).

Antagonism studies

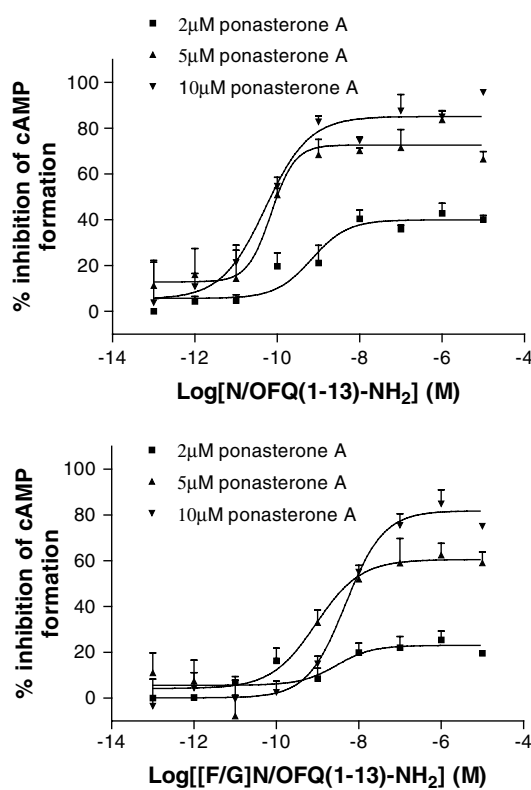
The NOP-selective, nonpeptide antagonist J-113397 was evaluated. J-113397 (100 nM) antagonised GTP γ ³⁵S binding stimulated by N/OFQ(1–13)–NH₂ with an apparent pK_B of 8.45 (Figure 5a). This value is essentially identical to the pA₂ for J-113397 of 8.53 measured previously in CHO_{hNOP} membranes (McDonald *et al.*, 2002).

At a 1 μ M ponasterone A induction, [F/G]N/OFQ(1–13)–NH₂ produced little or no measurable binding of GTP γ ³⁵S (Table 4), and hence this expression level was used to examine antagonist activity. At 1 μ M ponasterone A, the binding of GTP γ ³⁵S stimulated by N/OFQ(1–13)–NH₂ was competitively antagonised by 1 μ M [F/G]N/OFQ(1–13)–NH₂ with an apparent pK_B of 7.62 \pm 0.08 (Figure 5b). Using the same induction concentration in the GTP γ ³⁵S assay, the nonselective partial agonist NalBzOH (which was devoid of any agonist activity in this system, Table 6) was used to antagonise the actions of N/OFQ(1–13)–NH₂. NalBzOH (10 μ M) competitively antagonised GTP γ ³⁵S binding stimulated by N/OFQ(1–13)–NH₂ with an apparent pK_B of 7.02 \pm 0.13 (Figure 5c).

Table 4 N/OFQ(1–13)–NH₂ and [F/G]N/OFQ(1–13)–NH₂ stimulation of GTP γ ³⁵S binding and inhibition of cAMP formation in CHO_{INDhNOP} membranes and cells respectively, induced for 20 h with 1, 2, 5, 10 μ M ponasterone A

Induction (Ponasterone A)	GTP γ ³⁵ S			cAMP		
	<i>pEC</i> ₅₀ / <i>E</i> _{max} N/OFQ(1–13)NH ₂	<i>pEC</i> ₅₀ / <i>E</i> _{max} [F/G](1–13)NH ₂	N/OFQ(1–13)NH ₂ / [F/G](1–13)NH ₂	<i>pEC</i> ₅₀ / <i>E</i> _{max} N/OFQ(1–13)NH ₂	<i>pEC</i> ₅₀ / <i>E</i> _{max} [F/G](1–13)NH ₂	N/OFQ (1–13)NH ₂ / [F/G](1–13)NH ₂
1 μ M	8.12 \pm 0.32 1.28 \pm 0.03	Not analysable	—	Not analysable	Not analysable	—
2 μ M	8.68 \pm 0.11 1.93 \pm 0.20	7.23 \pm 0.38 1.51 \pm 0.15	0.55	9.42 \pm 0.49 40.9 \pm 2.2	8.26 \pm 0.87 19.6 \pm 4.8	0.48
5 μ M	8.52 \pm 0.06 4.33 \pm 0.80	7.68 \pm 0.10 2.01 \pm 0.23	0.30	9.72 \pm 0.40 79.5 \pm 4.1	8.99 \pm 0.18 59.37 \pm 5.8	0.75
10 μ M	8.60 \pm 0.07 6.95 \pm 1.05	7.72 \pm 0.06 3.21 \pm 0.38	0.37	10.35 \pm 0.22 86.0 \pm 3.7	8.32 \pm 0.13 83.23 \pm 4.0	0.97

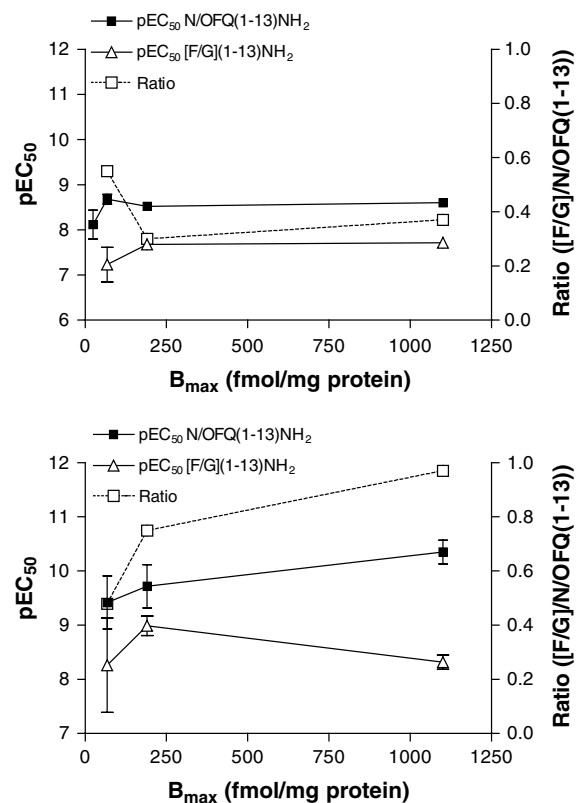
Data are mean \pm s.e.m. for $n \geq 3$ experiments. *pEC*₅₀ values for N/OFQ(1–13)–NH₂ and [F/G](1–13)–NH₂ did not differ ($P > 0.05$, ANOVA). There was a ponasterone concentration-dependent increase in *E*_{max} for N/OFQ(1–13)–NH₂ and [F/G](1–13)–NH₂ ($P < 0.05$, ANOVA).

**Figure 3** N/OFQ(1–13)–NH₂ (top panel) and [F/G]N/OFQ(1–13)–NH₂ (bottom panel) inhibition of forskolin-stimulated cAMP formation in whole CHO_{INDhNOP} cells induced with 2, 5 and 10 μ M ponasterone A. Data are mean \pm s.e.m. for $n \geq 3$.

Discussions

We show that the estimated intrinsic activity of a range of NOP partial agonists is dependent upon receptor density. In the ecdysone-inducible expression system, NOP not only displays the same pharmacology observed in different cell lines and tissues, but also allows reliable titration of receptor density.

To discriminate between antagonists and partial agonists, Berger *et al.* (2000b) described a method based upon decreasing the GDP concentration in GTP γ ³⁵S-binding studies. High GDP ($\geq 100 \mu$ M) concentration can mask the low

**Figure 4** Summary of *pEC*₅₀ values for N/OFQ(1–13)–NH₂ (top-GTP γ ³⁵S) and [F/G]N/OFQ(1–13)–NH₂ in CHO_{INDhNOP} (bottom-cAMP) as a function of receptor density (*B*_{max}). Also shown is the relative intrinsic activity (N/OFQ(1–13)–NH₂/[F/G]N/OFQ(1–13)–NH₂ ratio).

activity of partial agonists. Previously, we have shown that stimulation of GTP γ ³⁵S binding by the partial agonist NalBzOH depended on the GDP concentration (Bigoni *et al.*, 2002a). Here, we further describe this effect for [F/G]N/OFQ(1–13)–NH₂ and Ac-RYYWK–NH₂. Decreasing the GDP concentration to 5 μ M increased the net stimulated GTP γ ³⁵S binding. Moreover, the intrinsic activity of the partial agonists [F/G]N/OFQ(1–13)–NH₂ and Ac-RYYRWK–NH₂ relative to N/OFQ and N/OFQ(1–13)–NH₂ in both CHO_{hNOP} and CHO_{INDhNOP} (5 μ M ponasterone A) systems increased. This greater increase in intrinsic activity for

Table 5 PTx sensitivity of agonist-stimulated GTP γ^{35} S binding and cAMP inhibition for CHO_{INDhNOP} membranes and cells (5 μ M ponasterone A), respectively

Ligand	GTP γ^{35} S binding (stimulation factor)		cAMP inhibition (%)	
	Control	+ PTx	Control	+ PTx
N/OFQ	2.45 \pm 0.34	1.01 \pm 0.16*	43.1 \pm 8.9	0.0 \pm 9.6*
N/OFQ(1-13)-NH ₂	2.49 \pm 0.38	1.17 \pm 0.13*	50.3 \pm 8.8	1.7 \pm 7.0*
[F/G]N/OFQ(1-13)-NH ₂	1.48 \pm 0.13	0.74 \pm 0.20*	45.5 \pm 16.7	3.4 \pm 15.3*

Agonists were included at 10 μ M for GTP γ^{35} S experiments and 100 nM for cAMP measurements. Data are mean \pm s.e.m.; $n \geq 3$. *Values are significantly reduced compared with control, $P < 0.05$ (unpaired *t*-test).

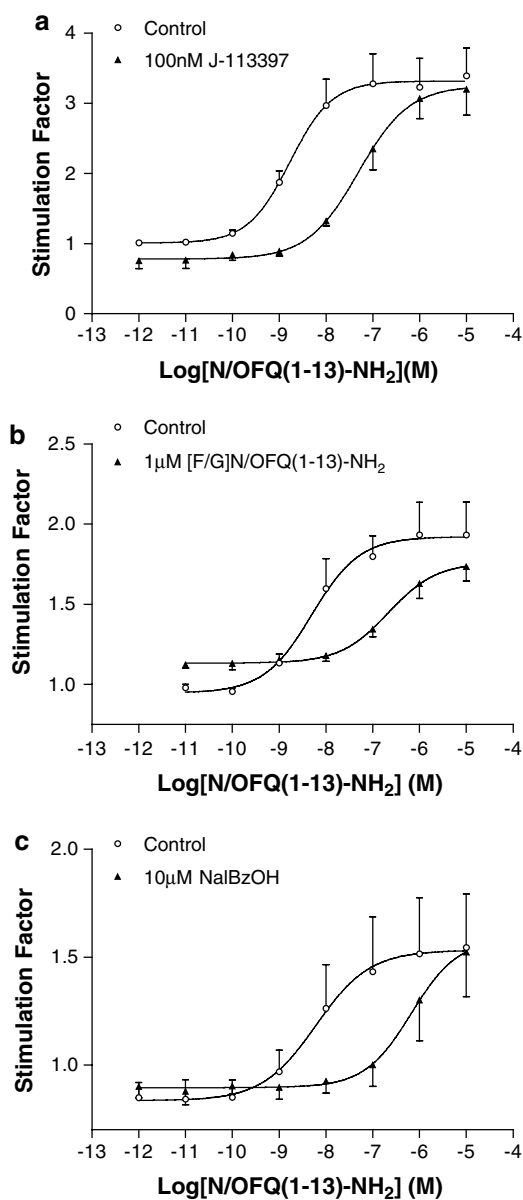
Table 6 Effects of NalBzOH, Ac-RYYRIK-NH₂ and Ac-RYYRWK-NH₂ at different induction concentrations on GTP γ^{35} S binding and comparison with N/OFQ and N/OFQ-NH₂

Ligand	[Induction] (ponasterone A) (μ M)	pEC_{50}	E_{max}
N/OFQ	5	8.26 \pm 0.01	3.72 \pm 1.01
N/OFQ-NH ₂	5	8.92 \pm 0.05	4.13 \pm 0.62
NalBzOH	1	No response at 100 μ M	
	5	No response at 100 μ M	
	10	No response at 100 μ M	
Ac-RYYRIK-NH ₂	1	7.76 \pm 0.26	1.10 \pm 0.04
	5	8.27 \pm 0.27	1.66 \pm 0.02*
Ac-RYYRWK-NH ₂	1	8.39 \pm 0.27	1.22 \pm 0.07
	5	8.69 \pm 0.11	2.16 \pm 0.08*

Data are mean \pm s.e.m. for $n \geq 3$ experiments. * $P < 0.05$ (unpaired *t*-test) significantly different compared with 1 μ M ponasterone A-induced cells.

[F/G]N/OFQ(1-13)-NH₂ and Ac-RYYRWK-NH₂ may suggest that partial agonists and full agonists differ in their dependency for GDP.

In order to carry out a more detailed study of the effects that differential expression of hNOP has on the intrinsic activity of different ligands in one system, the ecdysone expression system has been used (Van Craenenbroeck *et al.*, 2001). The higher concentration (10 μ M ponasterone A) produced receptor densities (~ 1 pmol mg⁻¹) similar to many commonly used transfected cell systems, for example, CHO_{hNOP} here used 1.9 pmol mg⁻¹ (Hashiba *et al.*, 2002), HEK 293 1.2 pmol mg⁻¹ (Dautzenberg *et al.*, 2001) and CHO_{hNOP} 0.9 pmol mg⁻¹ (Mason *et al.*, 2001). The 5 μ M induced receptor density (~ 200 fmol mg⁻¹) was similar to that reported in rat central tissue, for example, rat cortex 236 fmol mg⁻¹ (Berger *et al.*, 2000a), rat frontal cortex 246 fmol mg⁻¹ (Mason *et al.*, 2001) and rat cerebral cortex 180 fmol mg⁻¹ (Okawa *et al.*, 1998) and represents a pseudo-physiological level of receptor expression. Competition binding assays at this expression density indicated a pharmacology consistent with that reported in the literature. In GTP γ^{35} S and cAMP assays at 5 μ M induction, N/OFQ and N/OFQ-NH₂ were both full agonists with pEC_{50} values of 8.26 and 8.92 (GTP γ^{35} S), 9.38 and 9.66 (cAMP), respectively. Furthermore, in GTP γ^{35} S-binding assays at this induction concentration, Ac-RYYRIK-NH₂ and Ac-RYYRWK-NH₂ (Dooley *et al.*, 1997) were partial agonists and the effects of N/OFQ(1-13)-NH₂ were antagonised by J-113397 ($pK_B \sim 8.45$). In all assays, agonist effects were PTx sensitive, confirming the expected G_i/G_o coupling in this system.

**Figure 5** N/OFQ(1-13)-NH₂-stimulated GTP γ^{35} S binding is reversed by 100 nM J-113397 at 5 μ M ponasterone A induction (a), 1 μ M [F/G]N/OFQ(1-13)-NH₂ at 1 μ M ponasterone A induction (b) and 10 μ M NalBzOH at 1 μ M ponasterone A induction (c). Data are mean \pm s.e.m., $n = 3$.

In GTP γ ³⁵S binding, N/OFQ(1–13)–NH₂ was always a full agonist, while [F/G]N/OFQ(1–13)–NH₂ produced submaximal stimulation, possessing little or no response at 1 μ M induction. In cAMP measurements, N/OFQ(1–13)–NH₂ was again a full agonist in all preparations. However, [F/G]N/OFQ(1–13)–NH₂ displayed full agonism at the 10 μ M ponasterone A concentration, with the percentage inhibition being similar to that reported by us in our higher stable expressing CHO_{hNOP} transfects (see Okawa *et al.*, 1999), and partial agonism at all lower expressions of hNOP. This is typical of the amplification seen when measuring a downstream effector such as cAMP, that is, saturation of stimulus–response mechanisms becomes more evident the further down the stimulus–response chain the response is measured (Kenakin, 1997). In GTP γ ³⁵S assays at the lowest induction concentration (1 μ M ponasterone A), [F/G]N/OFQ(1–13)–NH₂ and NalBzOH acted as competitive antagonists with pK_B values of 7.62 and 7.02, similar to their pEC₅₀ values of 7.68 and 7.00 (Bigoni *et al.*, 2002a), respectively.

Conflicting data from different groups using similar and differing preparations reported agonism, partial agonism and antagonism for [F/G]N/OFQ(1–13)–NH₂ and also for Ac-RYYRIK–NH₂, Ac-RYYRWK–NH₂ and NalBzOH (Okawa *et al.*, 1999; Berger *et al.*, 2000a; Calo *et al.*, 2000a; Mason *et al.*, 2001). *In vitro* [F/G]N/OFQ(1–13)–NH₂ was a full agonist for inhibition of cAMP formation in CHO_{hNOP} cells and inhibition of glutamate release from synaptosomes (Okawa *et al.*, 1999). Following *i.c.v.* injection in rats, [F/G]N/OFQ(1–13)–NH₂ caused a decrease in heart rate, mean arterial pressure, urinary sodium excretion and a marked increase in urine flow, similar to N/OFQ but of longer duration (Kapusta *et al.*, 1999). Partial agonism was also reported for the stimulation of GTP γ ³⁵S binding in mouse N1E-115 cells (Olianas *et al.*, 1999). For a detailed review of the actions of [F/G]N/OFQ(1–13)–NH₂, see Calo *et al.* (2000a). This difference in signalling between central and peripheral NOP was explained by [F/G]N/OFQ(1–13)–NH₂ being a partial agonist with strong coupling in central tissue and high-expression transfected systems and weak coupling in peripheral tissue and low-expression systems (Okawa *et al.*, 1999). To date, the variable pharmacology of these partial agonists has not been carefully examined in the same expression system.

This problem has been addressed by only a few groups, using either cells transfected with different levels of NOP or using peripheral and central tissue (Mason *et al.*, 2001). A recent paper by Mason *et al.* (2001) showed differences in the relative intrinsic activities of [F/G]N/OFQ(1–13)–NH₂, Ac-RYYRIK–NH₂ and Ac-RYYRWK–NH₂ using transfected cells, central preparations and peripheral tissue. Differences in coupling efficiency are just one variable seen between different tissue preparations and native and recombinant NOP, which can affect values of relative intrinsic activity. Hence, differences in relative intrinsic activity may not be the result of changes in receptor number, but due to changes in coupling efficiency or other local cellular factors such as GDP concentration. Recent suggestions and data have shown that agonists differ in their efficacy for different cellular responses or subtypes of downstream effector (Berg *et al.*, 1998; Cordeaux *et al.*, 2000). Indeed, N/OFQ can stimulate PLC activity, with differential potency *via* a G_{z14}-mediated PTx-insensitive pathway (in G_{z14}-transfected cells, EC₅₀ 5 nM; Yung

et al., 1999) and *via* a G_{zi} PTx-sensitive pathway (EC₅₀ 0.4 nM; Reinscheid *et al.*, 1995). Therefore, different subtypes of effector or cellular pathways leading to a given response between different cell types or tissue preparations could give rise to differential efficacy/potency making conclusions about relative activities of ligands awkward, *i.e.*, differences in relative efficacy of a ligand between tissues may not be due to receptor density alone.

As noted, it has been suggested that the variable activity reported for the actions of [F/G]N/OFQ(1–13)–NH₂, that is, agonist, partial agonist and antagonist, was the result of different expressions of NOP at those sites assayed. Since we were able to control the expression of hNOP by changing the induction concentration of ponasterone A, we could measure the effect this had on the intrinsic activity of both N/OFQ(1–13)–NH₂ and [F/G]N/OFQ(1–13)–NH₂ in two functional assays. The efficacy of N/OFQ(1–13)–NH₂ with respect to its ability to stimulate GTP γ ³⁵S binding and inhibit adenylyl cyclase was full agonist in nature for all expression levels. However, the relative intrinsic activity of [F/G]N/OFQ(1–13)–NH₂ varied at different expression levels and between assays. In cAMP measurements, [F/G]N/OFQ(1–13)–NH₂ was a full agonist at 10 μ M ponasterone A and partial agonist at all other induction concentrations. An increased receptor reserve at 10 μ M ponasterone A induction and amplification steps in the pathway leading to the inhibition of adenylyl cyclase may explain this finding. However, it may be more fitting to suggest a coupling reserve (*i.e.* only a small proportion of the activated G protein is required to generate a full response), since no change in relative intrinsic activity is seen in GTP γ ³⁵S binding, suggesting that there is no receptor reserve. At expression levels greater than that induced by 1 μ M ponasterone A, that is, \sim 30 fmol mg^{–1}, [F/G]N/OFQ(1–13)–NH₂ was a partial agonist in GTP γ ³⁵S measurements. Below this expression, [F/G]N/OFQ(1–13)–NH₂ produced no response. This is due to the very low density of hNOP and [F/G]N/OFQ(1–13)–NH₂ reduced efficacy for the receptor. Indeed, it is tempting to suggest that in previous studies where [F/G]N/OFQ(1–13)–NH₂ behaved as an antagonist, this is due to similar low expression as that shown here, although as mentioned previously other factors can also play a role.

It can be seen from our data that intrinsic activity is a property of both the ligand and the tissue; hence relative values change with hNOP expression. Intrinsic activity is useful as a comparison of ligand efficacy in a sense of rank order (*i.e.* it can be said that N/OFQ(1–13)–NH₂ is more efficacious than [F/G]N/OFQ(1–13)–NH₂), but it is not possible to infer the molecular properties of agonism, such as intrinsic efficacy (*i.e.* the response per unit pharmacological receptor), from comparison of tissue maxima (Kenakin, 1997). However, in cases where fractional occupancy–response curves are of a more linear nature, in the absence of a receptor reserve or when the maximal tissue response has not been saturated, comparison of intrinsic activity (maximal tissue response) may represent a good measure of intrinsic efficacy, although this will need rigorous experimental validation (Kenakin, 2002). Since with GTP γ ³⁵S binding there appears to be no excess of receptor or saturation in response (suggested by the static nature of the relative intrinsic activity), for any receptor density, [F/G]N/OFQ(1–13)–NH₂ has between 0.37–0.55 relative intrinsic activity. Given that little or no change is seen in relative

intrinsic activity between [F/G]N/OFQ(1–13)–NH₂ and N/OFQ(1–13)–NH₂, it could be suggested that the latter ligand is not returning the system maximum response with regard to GTPγ³⁵S binding. Given a high density of available guanine nucleotide-binding sites (Albrecht *et al.*, 1998), it is not surprising that in this tissue and under these assay conditions (high GDP), GTPγ³⁵S binding is apparently not saturating; hence no clear receptor reserve and static relative intrinsic activity. However, in cAMP measurements at 2 μM ponasterone A induction where the maximal tissue response has not been reached even by the full agonist N/OFQ(1–13)–NH₂, the relative intrinsic activity is 0.48. Therefore, it can be suggested that the intrinsic efficacy of [F/G]N/OFQ(1–13)–NH₂ is ~0.4–0.5.

Finally, the issue that for a partial agonist the pEC₅₀ should predict its pA₂ or pK_B has been addressed. Using the lowest induction concentration of 1 μM ponasterone A, the ability of N/OFQ(1–13)–NH₂ to stimulate GTPγ³⁵S binding was competitively antagonised by [F/G]N/OFQ(1–13)–NH₂ with

a pK_B of 7.62, which is essentially identical to its pEC₅₀ of 7.68 (5 μM ponasterone A). This was also true for NalBzOH that antagonised N/OFQ(1–13)–NH₂-stimulated GTPγ³⁵S binding with a pK_B of 7.02 (pEC₅₀ 7.00; Bigoni *et al.*, 2002a). In this assay NalBzOH was devoid of agonist activity and is therefore a very-low-efficacy partial agonist. Ac-RYYRIK–NH₂ and Ac-RYYRWK–NH₂ displayed similar intrinsic activity to [F/G]N/OFQ(1–13)–NH₂ for GTPγ³⁵S binding. Overall, (at 5 μM ponasterone A) these partial agonists display a rank order intrinsic activity of Ac-RYYRWK–NH₂ > [F/G]N/OFQ(1–13)–NH₂ > Ac-RYYRIK–NH₂ > NalBzOH.

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