Development and Characterization of a Novel, High-Affinity, Specific, Radiolabeled Ligand for BRS-3 Receptors

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ABSTRACT

Bombesin (Bn) receptor subtype 3(BRS-3) is an orphan G-protein-coupled receptor of the Bn family, which does not bind any natural Bn peptide with high affinity. Receptor knockout studies show that the animals develop diabetes, obesity, altered temperature control, and other central nervous system (CNS)/endocrine/gastrointestinal changes. It is present in CNS, peripheral tissues, and tumors; however, its role in normal physiology/pathophysiology, as well as its receptor localization/pharmacology is largely unknown, in part due to the lack of a convenient, specific, direct radiolabeled ligand. This study was designed to address this problem and to develop and characterize a specific radiolabeled ligand for BRS-3. The peptide antagonist Bantag-1 had >10,000-fold selectivity for human BRS-3 (hBRS-3) over other mammalian Bn receptors (BnRs) [i.e., gastrin-releasing peptide receptor (GRPR) and neuromedin

Introduction

Bombesin (Bn) receptor subtype 3 (BRS-3) is an orphan G-protein-coupled receptor (GPCR) included in the Bn family of receptors (BnRs) because of its high homology (47%-51%) to the mammalian members of this receptor family [i.e., gastrin-releasing peptide (GRP) receptor (GRPR) and neuromedin B (NMB) receptor (NMBR)] (Gorbulev et al., 1992; Fathi et al., 1993; Jensen et al., 2008). BRS-3 is receiving increased attention (Majumdar and Weber, 2012b; Gonzalez et al., 2015b; Xiao and Reitman, 2016) because receptor knockout studies in mice show numerous alterations in both energy metabolism/satiety (Ohki-Hamazaki et al., 1997; Feng et al., 2011; Majumdar and Weber, 2012a; Gonzalez et al., 2015b; Xiao and Reitman, 2016) and glucose/insulin control (Ohki-Hamazaki et al., 1997; Gonzalez et al., 2015b; Xiao and

B receptor (NMBR)]. Using iodogen and basic conditions, it was radiolabeled to high specific activity (2200 Ci/mmol) and found to bind with high affinity/specificity to hBRS-3. Binding was saturable, rapid, and reversible. The ligand only interacted with known BRS-3 ligands, and not with other specific GRPR/NMBR ligands or ligands for unrelated receptors. The magnitude of ¹²⁵I-Bantag-1</sup> binding correlated with BRS-3 mRNA expression and the magnitude of activation of phospholipase C in lung cancer cells, as well as readily identifying BRS-3 in lung cancer cells and normal tissues, allowing the direct assessment of BRS-3 receptor pharmacology/numbers on cells containing BRS-3 with other BnRs, which is usually the case. This circumvents the need for subtraction assays, which are now frequently used to assess BRS-3 indirectly using radiolabeled panligands, which interact with all BnRs.

Reitman, 2016). These alterations include the development of obesity, diabetes, hypertension, impaired glucose metabolism, reduced metabolic rates, and altered behavior, including changes in feeding with hyperphagia and decreased movement; and wide expression of hyperleptinemia and altered growth hormone secretion in both the central nervous system (CNS), especially in hypothalamic nuclei, amygdala, and the caudate nucleus, and peripheral tissues, such as the gastrointestinal (GI) tract; is frequently overexpressed in human tumors; and its activation has a growth-promoting effect on tumors, similar to GRPR and NMBR (Fathi et al., 1993; Moody et al., 2003; Sancho et al., 2011; Gonzalez et al., 2015b; Moreno et al., 2016; Yamada et al., 2000a,b).

The exploration of the role of BRS-3 in normal physiology as well as in pathologic processes has been hindered by the fact it remains an orphan receptor, by the lack of availability of selective agonists/antagonists until recently, and by confusion in its pharmacology, including the localization of the receptors in both the CNS and peripheral tissue (Jensen et al., 2008; Gonzalez et al., 2015b; Xiao and Reitman, 2016). The latter problem is due to the lack of a convenient specific BRS-3 ligand

ABBREVIATIONS: BALB 3T3 or BALB, mouse MIC embryonic fibroblast cells; Bantag-1, Boc-Phe-His-4-amino-5-cyclohexyl-2,4,5-trideoxypentonyl-Leu-(3-dimethylamino) benzylamide *N*-methylammonium trifluoroacetate; Bn, bombesin; BnR, bombesin receptors; BRS-3, bombesin receptor subtype 3; BSA, bovine serum albumin; CNS, central nervous system; DMEM, Dulbecco's minimum essential medium; GI, gastrointestinal; GPCR, G-protein-coupled receptor; GRP, gastrin-releasing peptide; GRPR, gastrin-releasing peptide receptor; hBRS-3, human bombesin receptor subtype 3; hGRPR, human gastrin-releasing peptide receptor; hNMBR, neuromedin B receptor; IP, inositol phosphate; ¹²⁵I-Peptide-1, ¹²⁵I-[D-Tyr⁶, β-Ala¹¹, Phe¹³, NIe¹⁴]Bn-(6-14); ME, (Bn-(6-13) methyl ester; MK-5046 [(2S)-1, 1,1-trifluoro-2-[4-(1H-pyrazoI-1-yl)phenyl]-3-(4-[[1-(trifluoromethyl)cyclopropyl]methyl]-1H-imidazoI-2-yl)propan-2-o]; NMB, neuromedin B; NMBR, neuromedin B receptor; PD, PD168368.

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that can be used for receptor localization/characterization in different tissues as well as the lack of agreement on the reliability of BRS-3 antibodies, with one recent review (Xiao and Reitman, 2016) concluding that existing BRS-3 receptor antibodies are of uncertain specificity. This is a particular issue with the CNS, allowing the correlation of BRS-3 location with the results of recent studies attempting to define the exact CNS areas responsible for many of the changes in energy homeostasis, food intake, metabolic rate, body temperature control, and body weight (Gonzalez et al., 2015b; Xiao and Reitman, 2016; Maruyama et al., 2017; Xiao et al., 2017; Pinol et al., 2018).

The current study was an attempt to address this latter problem by developing and characterizing a radiolabel ligand that could be readily available; had high selectivity for the BRS-3 receptor over the closely related GRPR/NMBR, which are frequently coexpressed on the same cells (Fathi et al., 1993; Jensen et al., 2008; Moreno et al., 2016); and could be used for BRS-3 pharmacological characterization and localization, both in the CNS and peripheral tissues. In this study, evidence is presented that this was accomplished by preparing iodinated Bantag-1 [Boc-Phe-His-4-amino-5-cyclohexyl-2,4,5trideoxypentonyl-Leu-(3-dimethylamino)benzyl amide-N-methyl ammonium trifluoroacetate)], a BRS-3 antagonist that had a >10,000 selectivity over GRPR/NMBR for the BRS-3, demonstrating that it had a high degree of saturable binding to BRS-3 with rapid kinetics; interacted only with BRS-3 reported ligands, thus demonstrating high specificity for BRS-3; and could be used to directly assess the presence or number of BRS-3 receptors in normal or tumoral tissues. The availability of this ligand should markedly facilitate the ability to study the presence and pharmacology of this increasing important receptor in both normal physiologic and pathophysiological conditions.

Materials and Methods

Materials

BALB 3T3 (mouse fibroblast) cells; NCI-N417 and NCI-H69 (human small cell lung cancer cells); and NCI-H520, NCI-H460, NCI-H720, and NCI-H727 (human non-small cell lung cancer cells) were purchased from the American Type Culture Collection (Gaithersburg, MD); Dulbecco's minimum essential medium (DMEM), RPMI 1640, PBS, FBS, Dulbecco's phosphate-buffered saline, trypsin-EDTA $1\times$, penicillin streptomycin, and GENETICIN-selective antibiotic (G418 sulfate) were from Invitrogen (Carlsbad, CA); GRP and NMB were from Bachem (Torrance, CA); the Bn analogs $[D-Tyr^6, \beta-Ala^{11}, Phe^{13}, Nle^{14}]Bn-(6-14)$ (Peptide-1) (Mantey et al., 1997; Pradhan et al., 1998; Moreno et al., 2013) and other BnR ligands were obtained from Dr. David H. Coy, Tulane University (New Orleans, LA); Bantag-1, a BRS-3 antagonist (Guan et al., 2010; Feng et al., 2011; Nakamura et al., 2016) was from Sigma-Aldrich (Allentown, PA); MK-5046 ([(2S)-1,1,1-trifluoro-2-[4-(1H-pyrazol-1-yl)phenyl]-3-(4-[[1-(trifluoromethyl)cyclopropyl]methyl]-1H-imidazol-2-yl)propan-2-ol]), a BRS-3 agonist (Sebhat et al., 2010; Moreno et al., 2013), was from MedChi Express (Monmouth Junction, NJ); and iodine-125 radionuclide [125I] (10 mCi) (378 MEq) was from Perkin Elmer Life Science (Boston, MA).

Methods

Stable Transfection. Human BRS-3 (hBRS-3), human GRPR (hGRPR), and human NMBR (hNMBR) stably transfected into BALB 3T3 cells were grown as described previously (Benya et al., 1992, 1993, 1994, 1995; Ryan et al., 1998a). BALB 3T3 cells were grown overnight

at 37°C in 10-cm tissue culture plates at a density of 2×10^6 cells per plate. Twenty-four micrograms of hBRS-3, hGRPR, or hNMBR plasmid DNA was transfected into the BALB 3T3 cells by the cationic lipid-mediated method using 60 μ l of LipofectAMINE 2000 in OPTI-MEM 1–reduced serum medium for 3 hours at 37°C. At the end of the incubation period, the medium was replaced with regular growth medium. Three days after transfection, cells were split into very dilute concentrations, and 800 mg/l selection antibiotic G418 (Geneticin, Thermo Fisher Scientific) was added to the regular growth medium. In about 2 weeks, single colonies were selected and expanded. The best clones, as determined by binding, were further expanded, frozen, and used for our experiments.

Cell Culture. BALB 3T3 cells stably expressing hGRPR, hNMBR, and hBRS-3 were grown in DMEM supplemented with supplemented 10% FBS and 300 mg/l G418 sulfate. Lung cancer cells NCI-N417, H69, H520, H460, and H727, which natively express BnRs including BRS-3 (Moreno et al., 2018), were grown in RPMI 1640 supplemented with 10% FBS, and NCI-H720 was grown in DMEM/F12 Medium supplemented with 5% FBS. All the cells were grown in their respective propagation medium supplemented with supplemented 10% FBS and incubated at 37°C in a 5% CO₂ atmosphere.

Preparation of ¹²⁵I-Labeled Peptides. ¹²⁵I-Bantag-1, with specific activity of 2200 Ci/mmol, was prepared by a modification of methods described previously (Jensen et al., 1983; von Schrenck et al., 1990; Mantey et al., 1993, 1997). Briefly, 0.8 µg of IODO-GEN (in 0.02 µg/ml chloroform) was transferred to a 5-ml glass test tube, dried under a stream of nitrogen, and washed with 100 μ l of 0.5 M KH₂PO₄ (pH 7.4). To the reaction tube, $20 \,\mu l$ of $0.5 \,M \,KH_2 PO_4$ (pH 8.0), $10 \,\mu g$ of peptide in 5 μ l of water, and 2 mCi (20 μ l) Na ¹²⁵I were added, mixed gently, and incubated at room temperature for 6 minutes. The incubation was stopped by the addition of 300 μ l of distilled water. The free 125 I was separated by applying the reaction mixture onto a Sep-Pak and washing with 5 ml of distilled water and 5 ml of 0.1% trifluoroacetic acid. The radiolabeled peptide was eluted with 60% acetonitrile in 0.1% trifluoroacetic acid (10 \times 200 μ l sequential elutions). The radioligand was purified by taking the two peaks with the highest radioactivity and applying them to a reverse-phase highperformance liquid chromatography using a µBondapak column $(0.46 \times 25 \text{ cm})$ from Waters. The column was eluted with a linear gradient of acetonitrile in 0.1% trifluoroacetic acid from 16% to 64% acetonitrile in 60 minutes, with a flow of 1.0 ml/min, and collecting 1-ml fractions. Samples of the fractions were counted, and the highest peaks were tested for radioligand binding to hBRS-3-transfected cells (Ryan et al., 1998a; Moreno et al., 2013). The best binding fractions were stored with 0.5% BSA at -20° C. ¹²⁵I-Peptide-1 (¹²⁵I-[D-Tyr⁶, β -Ala¹¹, Phe¹³, Nle¹⁴] Bn-(6-14)), with a specific activity of 2200 Ci/mmol, was prepared as described previously (Mantey et al., 1997; Moreno et al., 2013; Nakamura et al., 2016).

Binding Studies. hBRS-3/BALB $(1 \times 10^6 \text{ cells/ml})$, hGRPR/BALB $(0.5 \times 10^6$ cells/ml), hNMBR/BALB (0.05 \times 10^6 cells/ml), NCI-N417 $(3 \times 10^7 \text{ cells/ml})$, NCI-H69 $(6 \times 10^6 \text{ cells/ml})$, NCI-H460 $(6 \times 10^6 \text{ cells/ml})$ cells/ml), NCI-H520 (6 \times 10⁶ cells/ml), NCI-H720 (3 \times 10⁶ cells/ml), NCI-H727 (4 \times 10⁶ cells/ml) cells were incubated for 40–60 minutes at 22°C with ¹²⁵I-labeled ligand and with or without ligand in 300 μ l of binding buffer, as described previously (Pradhan et al., 1998; Ryan et al., 1998a,b; Moreno et al., 2013). The standard binding buffer contained 24.5 mM HEPES (pH 7.4), 98 mM NaCl, 6 mM KCl, 2.5 mM KH₂PO₄, 5 mM sodium pyruvate, 5 mM sodium fumarate, 5 mM sodium glutamate, 2 mM glutamine, 11.5 mM glucose, 0.5 mM CaCl₂, 1.0 mM MgCl₂, 0.01% (w/v) soybean trypsin inhibitor, 0.2% (v/v) amino acid mixture, 1% (w/v) BSA, and 0.05% (w/v) bacitracin. After the incubation, 100 μ l of each sample was removed and added to 300 μ l of washing buffer 2% (w/v) BSA in PBS in 400-µl microfuge tubes and centrifuged for 1 minute at 10,000g (Microfuge E; Beckman, Fullerton, CA) to separate the bound radioligand from the unbound radioligand. The supernatant was aspirated, and the pelleted cells were rinsed twice with a washing buffer. The amount of radioactivity bound to the

cells was measured in a Wizard 1470 Automatic Gamma Counter (Packard Instruments, Meriden, CT). Total binding was expressed as the percentage of total radioactivity that was associated with the cell pellet. Nonsaturable binding was <15% of the total binding in all experiments. Each point was measured in duplicate, and each experiment was replicated at least four times. The calculation of affinity was performed by determining the IC₅₀ using the curve-fitting program GraphPad Prism 4.0 (GraphPad Software, La Jolla, CA).

Preparation of Membranes from Various Tissues. Cell membranes were prepared as described previously (Mantey et al., 1993; Tsuda et al., 1997), using a homogenizing buffer containing 50 nM Tris (pH 7.4), 0.2 mg/ml soybean inhibitor, 0.1% bacitracin, and 0.2 mg/ml benzamidine. A concentration of 1×10^7 cell/ml lung cancer cells or rat tissues (cerebellum, cortex, mid-brain, testis, duodenum) were homogenized for 30 seconds with a polytron (Brinkmann Instruments, Inc.) at speed 6 and were centrifuged at 1500 rpm for 10 minutes. The supernatant was removed and recentrifuged at 15,000 rpm to pellet the membranes. The membranes were suspended in binding buffer and stored at -70° C until used.

Binding of ¹²⁵I-Peptide-1 and ¹²⁵I-Bantag-1 to Membranes. The standard membrane-binding buffer contained 10 mM HEPES (pH 7.4, 118 mM NaCl, 4.7 mM KCl, 5 mM MgCl₂, 1 mM EGTA, 0.2 mg/ml benzamidine, 0.2 mg/ml soybean trypsin inhibitor, 0.1% bacitracin, and 0.2% (w/v) BSA. Binding was as stated above with cells, except that membranes were diluted with binding buffer to a concentration equivalent to 6×10^6 cells/ml and that after incubation membranes were spun through 4% BSA for 2 minutes to pellet and wash.

Dissociation of Bound Ligand from hBRS-3/BALB Membranes and Cells. Dissociation studies were performed in both hBRS-3 BALB cells and membranes. Membranes were incubated with 50 pM radioligand (i.e., ¹²⁵I-Bantag-1 or ¹²⁵I-Peptide-1) at room temperature for 60 minutes with or without 1 μ M peptide-1 in binding buffer as described above. hBRS-3/BALB (1 × 10⁶ cells/ml) cells were incubated for 60 minutes at 37°C with 50 pM ¹²⁵I-labeled ligand (¹²⁵I-Bantag-1 or ¹²⁵I-Peptide-1) and with or without 1 μ M peptide-1 in binding buffer, as described above. Then a final concentration of 1 μ M peptide-1 was added, and the membranes or cells were incubated at room temperature or 37°C, respectively. At various times, 100- μ l aliquots were taken and spun through 4% BSA for 2 minutes to pellet. The pellet was washed, and the amount of radioactivity saturably bound was determined with a gamma counter.

Internalization of ¹²⁵I-Peptide-1 and ¹²⁵I-Bantag-1. Internalization of ligands was determined as described previously (Benya et al., 1993; Mantey et al., 1993). Briefly, after incubation of hBRS-3/BALB cells with 50 pM ¹²⁵I-Peptide-1 or ¹²⁵I-Bantag-1 for the indicated times and temperatures, samples (100 μ l) were added to acid-stripping buffer [0.2 M acetic acid with 0.5 M NaCl (pH 2.5)] to remove the surface-bound radioligand or were added to standard washing buffer to assess total saturable bounding. After 5 minutes, at 4°C, the cells were centrifuged and washed two times with washing buffer, and the amount of saturable radioactivity left was determined on the gamma counter.

Measurement of [³**H]-Inositol Phosphates.** For lung cancer cells not previously reported (Moreno et al., 2018) (H838,A549,H1299), the ability of stimulation of BRS-3 to activate phospholipase C resulting in [3H]-inositol phosphate (IP) production was determined as described previously (Rowley et al., 1990; Moreno et al., 2018). Briefly, all adherent cells were subcultured into 24-well plates (5 × 10⁵ cells/ml) in regular propagation media and then incubated for 24 hours at 37°C in a 5% CO₂ atmosphere. Then, the cells were incubated with 3 μ Ci/ml myo-[2-³H] inositol in growth media supplemented with 2% FBS for an additional 24 hours. After the incubation, the 24-well plates were washed by incubating for 30 minutes at 37°C with 1 ml/well PBS (pH 7.0) containing 20 mM lithium chloride. Cells that grew in suspension were centrifuged to removed RPMI 1640 medium and

incubated $(2 \times 10^6$ cell/ml) directly with myo-[2-³H] inositol in RPMI 1640 with 2% FBS for 24 hours (Moreno et al., 2018). The cells were then centrifuged to remove the myo-[2-³H]-inositol, washed with wash buffer containing 20 mM LiCl for 10 minutes at 4°C, recentrifuged to removed wash buffer, and distributed to 5-ml tubes, where they were incubated without (control) or with peptides in 500 μ l of IP assay buffer for 60 minutes at 37°C. In both types of cells, the incubation was terminated by the addition of 1 ml of ice-cold 1% (v/v) hydrochloric acid in methanol. Total [³H]IP was isolated by anion exchange chromatography, as described previously (Rowley et al., 1990; Ryan et al., 1998a; Uehara et al., 2011; Moreno et al., 2018).

Cytosolic Ca²⁺

For lung cancer cells not previously reported (Moreno et al., 2018) (H838, A549, H1299), the ability of stimulation of BRS-3 to stimulate cytosolic Ca²⁺ was investigated with cells with adherent growth treated with trypsin-EDTA and harvested (2.5×10^{6} cells/ml). After centrifugation, the cells were resuspended in SIT-medium (RPMI-1640 containing 3×10^{-8} M sodium selenite, 5 µg/ml insulin, and 10 µg/ml transferrin) (Ryan et al., 1998a,b; Moreno et al., 2018) containing 1 µM Fura-2/acetoxymethyl ester (Calbiochem, La Jolla, CA) at 37°C for 30 minutes. Subsequently, the cells were centrifuged at 1000g for 5 minutes, resuspended, and changed in cytosolic Ca²⁺, assessed as described previously (Moreno et al., 2018).

Statistical Analysis. The results are the mean and S.E.M. from at least three separate experiments. The K_i (kinetic constant) values were calculated from the IC_{50} values obtained from competitive inhibition binding curves using different Bn analogs and antagonists (50 pM ¹²⁵I-Peptide-1 or 50 pM ¹²⁵I-Bantag-1) using the equation of Cheng and Prusoff (1973). IC₅₀ values from the binding data were obtained by curve fitting using GraphPad Prism 7.0 (nonlinear curve fitting). An analysis of variance was used to determine the statistical significance of differences in the affinity of each Bn agonist/antagonist with Bantag-1 with changes showing $a \ge 2$ -fold difference. In all experiments, cell concentrations were set such that <15% of total radioactivity was bound and the amount of saturable ligand bound was similar for all compounds, and, therefore, in the statistical analysis only two variables (i.e., each Bn agonist/antagonist and its own control, binding a similar amount of ligand) were analyzed. Correlation coefficients were calculated using GraphPad Prism 7.0.



Fig. 1. Time course of binding of the radiolabeled nonselective Bn analog agonist (Peptide-1) and the BRS-3 peptide antagonist Bantag-1. BALB 3T3 cells transfected with hBRS-3 were incubated with 50 pM ¹²⁵I-Bantag-1 or ¹²⁵I-Peptide-1 alone or with 1 μ M unlabeled ligand at 22°C for the indicated times. Results are expressed as the percentage of the added counts bound. In each experiment, each value was determined in duplicate, and the results given are from at least four separate experiments.

Time Course of Association and Dissociation of Binding of the Radiolabeled Nonselective Bn Analog Agonist (¹²⁵I-Peptide-1) and the Radiolabeled BRS-3–Selective Peptide Antagonist ¹²⁵I-Bantag-1

The binding association of both radioligands ¹²⁵I-Peptide-1 and ¹²⁵I-Bantag-1 to hBRS-3 BALB 3T3 cells was time dependent, with a rapid increase, reaching a maximal increase after 30 minutes (¹²⁵I-Peptide-1: 9.24% ± 0.27% of control; ¹²⁵I-Bantag-1: 8.10% ± 0.23% of control) (Fig. 1) and was maintained for up to 90 minutes (¹²⁵I-Peptide-1: 10.30% ± 0.05% of control; ¹²⁵I-Bantag-1: 8.10% ± 0.10% of control) (Fig. 1). Although both radioligands bound well to hBRS-3/BALB cells, the universal ligand ¹²⁵I-Peptide-1 demonstrated 12% greater steady-state binding than was seen with the peptide antagonist ¹²⁵I-Bantag-1 (Fig. 1). When each radioligand was incubated with a saturating concentration of Peptide-1 or Bantag-1, the total amount of binding was reduced by 91% and 85%, respectively (At 30 minutes: ¹²⁵I-Peptide-1 + 1 μ M Peptide-1:



0.83% \pm 0.05% of control; 125 I-Bantag-1 + 1 μM Bantag-1: 1.20% \pm 0.11% of control (Fig. 1).

The time course of dissociation for each ligand was similar for membranes and cells. With both membranes and cells, 125 I-Bantag-1 binding dissociated more rapidly than 125 I-Peptide-1 binding with 70%–80% of 125 I-Bantag-1 dissociated by 10 minutes, whereas only 40%–50% of 125 I-Peptide-1 was dissociated by this time (Fig. 2).

Ability of Peptide-1 or Bantag-1 to Interact with BRS-3s, GRPRs, or NMBRs

In separate BALB 3T3 cells containing hBRS-3, hGRPR, or hNMBR, ¹²⁵I-Peptide-1 bound to all three receptors with high affinity and the universal ligand Peptide-1 had a high affinity for the three BnRs (K_i : 1.88, 0.67, and 3.66 nM, respectively) (Fig. 3A; Table 1), similar to reported previously (Mantey et al., 1997; Pradhan et al., 1998; Sancho et al., 2010). However, the peptide antagonist Bantag-1 only had high affinity for hBRS-3/BALB (K_i , 0.72 nM) (Fig. 3B; Table 1) and did not interact with hGRPR/BALB or hNMBR/BALB, even at concentrations of 3000 nM (K_i , >10,000 nM) (Fig. 3B; Table 1). These results demonstrate that Peptide-1 binds with



Fig. 2. Time course of dissociation of the radiolabeled nonselective Bn analog agonist (Peptide-1) and the BRS-3 peptide antagonist Bantag-1 from hBRS-3/BALB 3T3 cell membranes (top) or intact cells (bottom). hBRS-3/BALB (1 × 10⁶ cells/ml) were incubated for 60 minutes at 37°C with 50 pM ¹²⁵I-labeled ligand (¹²⁵I-Bantag-1 or ¹²⁵I-Peptide-1) and with or without unlabeled 1 μ M peptide-1 in binding buffer. Membranes were incubated with 50 pM radio-ligand (i.e., ¹²⁵I-Bantag-1 or ¹²⁵I-Peptide-1) at room temperature for 60 minutes with or without unlabeled 1 μ M peptide-1 ligand in binding buffer. Then a final concentration of 1 μ M peptide-1 was added, and the membranes or cells were incubated at room temperature or 37°C, respectively. At various times, 100- μ I aliquots were taken and saturable binding determined as described in *Methods*. In each experiment, each value was determined in duplicate, and results are from at least four separate experiments.

Fig. 3. The ability of the nonselective Bn analog agonist (Peptide-1) (A) and the BRS-3-selective peptide antagonist Bantag-1 (B) to inhibit the binding of either ligand to cells containing BRS-3, GRPR, or NMBR. BALB 3T3 cells stably transfected with hBRS-3 (1×10^6 cells/ml), hGRPR (0.5×10^6 cells/ml), or hNMBR (0.05×10^6 cells/ml) were incubated with 50 pM ¹²⁵I-Peptide-1 for 60 minutes at 22°C, and the saturable binding was determined as described in *Materials and Methods*. The results are expressed as the percentage of saturable binding without unlabeled peptide added (percentage control). The results are the mean and S.E.M. from at least three separate experiments, and in each experiment the data points were determined in duplicate. BALB, BALB 3T3 cells. For other abbreviations, see the Fig. 1 legend.

TABLE 1

Comparison of the affinities of the peptide agonist Peptide-1 and the BRS-3 peptide antagonist Bantag-1 for members of the Bn receptor family (hBRS-3, hGRPR, and hNMBR)

Transfected cells hBRS-3 (1 × 10⁶ cells/ml), hNMBR (0.05 × 10⁶ cells/ml), or hGRPR (0.5 × 10⁶ cells/ml) were incubated with 50 pM ¹²⁶I-Peptide-1 for 60 minutes at 22°C. In each experiment, each value was determined in duplicate, and values given are the mean and S.E.M. from at least three separate experiments. Data are calculated from dose-inhibition curves shown in Fig. 1 for each peptide using a nonlinear regression curve-fitting program (Prism). Each value is the mean ± S.E.M. from at least three experiments.

Ligand	Ki			
	hBRS-3	hGRPR	hNMBR	
		nM		
Peptide-1 Bantag-1	$\begin{array}{c} 1.58 \pm 0.09 \\ 0.72 \pm 0.01 \end{array}$	$\begin{array}{c} 0.67 \pm 0.04 \\ {>}10{,}000 \end{array}$	$3.66 \pm 0.02 \ > 10,000$	

high affinity with all three human BnRs and is therefore nonselective, whereas Bantag-1 has >10,000-fold higher affinity for BRS-3 and, under these experimental conditions, only interacts with hBRS-3.

Comparison of the Ability of the Radiolabeled BRS-3 Antagonist, ¹²⁵I-Bantag-1, and the Radiolabeled Nonselective Agonist ¹²⁵I-Peptide-1 to Bind with Native BALB 3T3 Cells or BALB 3T3 Cells Stably Transfected with hBRS-3, hGRPR, or hNMBR

To demonstrate directly by binding studies that, similar to unlabeled Peptide-1 and Bantag-1, the radiolabel agonist ¹²⁵I-Peptide-1 and the radiolabel antagonist ¹²⁵I-Bantag-1, also differed in their abilities to interact with cells containing the different BnRs [hBRS-3/BALB, hGRPR/BALB and hNMBR/BALB, and BALB 3T3 cells], we incubated under similar conditions both radiolabels with the different cells. The agonist ¹²⁵I-Peptide-1 and the antagonist ¹²⁵I-Bantag-1 bound to similar extent to hBRS-3/BALB cells (3.9% and 4.3%



Fig. 4. Comparison of the ability of the radiolabeled BRS-3 antagonist ¹²⁵I-Bantag and the radiolabeled agonist ¹²⁵I-Peptide-1 to bind to native BALB 3T3 cells or BALB 3T3 cells stably transfected with hBRS-3, hGRPR, or hNMBR. Cells were incubated with 50 pM ¹²⁵I-Peptide-1 or 50 pM ¹²⁵I-Bantag-1 for 60 minutes at 22°C, and the saturable binding was determined as described in *Materials and Methods*. The results are expressed as the percentage of added saturable counts bound. Cell concentrations were the same as listed in Fig. 3, with 1 × 10⁶ cells/ml BALB cells. The results are the mean and S.E.M. from at least three separate experiments and in each experiment the data points were determined in duplicate. For abbreviations, see the legends of Figs. 1 and 3.

added counts, respectively) (Fig. 4). However, only ¹²⁵I-Peptide-1 bound to GRPR or NMBR (Fig. 4). In contrast neither ligand bound to native BALB 3T3 cells, which do not contain BnRs (Fig. 4).

Time Course and Temperature Dependence of Internalization and the Radiolabeled Antagonist ¹²⁵I-Bantag-1 or Agonist ¹²⁵I-Peptide 1 by hBRS-3/BALB 3T3 Cells

Previous studies demonstrate that BRS-3, as well as other BnRs, can internalized ligands (Benya et al., 1992, 1993, 1994, 1995; Kumagai et al., 2015). To assess the internalization of each ligand, we performed acid-stripping studies, which have been shown to remove surface-bound ligand (Benya et al., 1992, 1993, 1994, 1995; Kumagai et al., 2015). We found that both ¹²⁵I-Bantag-1 and ¹²⁵I-Peptide-1 were internalized in a time-dependent manner, with a significant increase after 10 minutes (40% and 33%, respectively) (Fig. 5A), and maximum internalization occurred after 30 minutes (59% and 43%, respectively) (Fig. 5A). The increase in internalization of



Fig. 5. Comparison of the time course and extent of internalization of the radiolabeled antagonist 125 I-Bantag-1 or radiolabeled agonist, 125 I-Peptide-1 with hBRS-3/BALB 3T3 cells. hBRS-3 stably transfected (top/bottom panels) into BALB 3T3 cells (1 \times 10⁶ cells/ml) were incubated with 50 pM 125 I-Bantag-1 or 50 pM 125 I-Peptide-1 at 37°C or 4°C (bottom panel) for the indicated times. The internalization of bound ligand was determined by assessing the effect of acid-stripping medium (0.2 M acetic acid and 0.5 M NaCl, pH 2.5), which removes surface-bound ligand. "Binding times" refers to the time the cells were incubated with the radioligand before acid stripping. Results are expressed as the percentage of total bound ligand remaining after acid stripping (percentage internalized) and are the mean and S.E.M. from at least three separate experiments, and in each experiment the data points were determined in duplicate. (Top panel) The percentage of total ligand bound that is internalized at the different incubation times is shown. (Bottom panel) Both the total saturable binding and the percentage remaining after acid wash (percentage internalized) are shown after incubation at 4°C or 37°C for the indicated times. For abbreviations, see the Fig. 1 legend.

¹²⁵I-Bantag-1 was maintained in time after 60 minutes (64%) (Fig. 5A). However, the amount of ¹²⁵I-Peptide-1 internalization decreased slightly over time but was still present after 60 minutes (33%) (Fig. 5A). The internalization of ¹²⁵I-Bantag-1 and ¹²⁵I-Peptide-1 was also temperature dependent. At 4°C, no internalization was seen with either of the two radioligands (Fig. 5B). However, both were internalized at 60 minutes at 37°C (Fig. 5B).

Specificity of Binding of ¹²⁵I-Bantag-1 to hBRS-3 Receptor

To assess the specificity of the binding of ¹²⁵I-Bantag-1 to hBRS-3 receptor, the ability of various ligands that interact preferentially with BRS-3 (Bantag-1, MK-5046) (Sebhat et al., 2010; Moreno et al., 2013), only the other Bn receptors [selective GRPR antagonists [Tyr⁴,D-Phe¹²]Bn; [[Leu¹³. $\psi(CH_2NH)$, Leu¹⁴]Bn; [D-Phe⁶, Leu¹³, $\Psi(CH_2NH)$, Cpa¹⁴]Bn-(6-14) D-Phe⁶]Bn-(6-13)methyl ester; D-Phe⁶]Bn-(6-13)ethyl ester); selective NMBR antagonist PD168368 (PD), selective GRPR (GRP), or NMBR (NMB) agonists] (Heinz-Erian et al., 1987; Coy et al., 1988, 1989; Wang et al., 1990a,b; Ryan et al., 1999; Gonzalez et al., 2009; Uehara et al., 2011; Ramos-Álvarez et al., 2015), ligands of a Drosophila receptor related to mammalian BRS-3 (i.e., CCHamides) (Matsufuji et al., 2014, 2015) or ligands interacting with receptors different from the Bn receptor family, were tested (Tables 2 and 3). At concentrations that cause a maximal effect at their receptors, none of these agents inhibited the binding of ¹²⁵I-Bantag-1 to hBRS-3-transfected BALB 3T3 cells except for MK-5046, which is a known BRS-3 agonist (Sebhat et al., 2010; Moreno et al., 2013). These results demonstrate the specificity of ¹²⁵I-Bantag-1 binding.

Ability of Various BRS-3 Ligands to Inhibit Binding of ¹²⁵I-Peptide-1 or ¹²⁵I-Bantag-1 to hBRS-3/BALB Cells

A number of various selective BRS-3 ligands have been reported recently in various studies using both functional

TABLE 2

hBRS-3 stably transfected Balb 3T3 cells (1 \times 10⁶ cells/ml) were incubated with 50 pM ¹²⁵T-Bantag-1 either alone or with the indicated concentration of the unlabeled peptide listed. Results are expressed as the percentage of the saturable binding with no unlabeled peptide. Results are the mean \pm S.E.M. from at least three experiments, and in each experiment each point was determined in duplicate.

Agent Added	¹²⁵ I-Bantag-1 Bound
	% Control
Peptide-1 (10 μ M)	8.50 ± 1.20
Bantag-1 (0.01 nM)	80.0 ± 4.1
Bantag-1 (10 μ M)	0
MK-5046 (10 μ M)	62.5 ± 3.4
Bn (10 μ M)	96.0 ± 9.2
CCH Amide I (10 μ M)	87.0 ± 4.3
CCH Amide II (10 μ M)	99.0 ± 5.2
Substance P (10 μ M)	93.0 ± 4.0
CCK-8 (10 μ M)	100.0 ± 3.2
Gastrin-1 (10 μ M)	94.0 ± 2.6
Somatostatin-14 (10 μ M)	100.0 ± 3.3
$[Tyr^4, D-Phe^{12}]Bn (1 \mu M)$	98 ± 3.8
[Leu ¹³ , Ψ (CH ₂ NH), Leu ¹⁴]Bn-(6-14) (1 μ M)	100.0 ± 4.1
$[D-Phe^{6}, Leu^{13}, \Psi(CH_{2}NH),$	100.0 ± 2.9
Cpa ¹⁴]Bn-(6-14) (10 µM)	
[D-Phe ⁶]Bn-(6-13)methyl ester (1 µM)	96.3 ± 1.2
$[D-Phe^{6}]Bn-(6-13)ethyl ester (1 \mu M)$	100.0 ± 4.2
PD-1676252 (1 µM)	99.1 ± 3.0

CCH, carbamylcholine; CCK, cholecystokin.

assays and/or various binding studies with ¹²⁵I-Peptide-1 (Mantey et al., 1997, 2001, 2004; Weber et al., 2002, 2003). To compare the results of the ability of these ligands to interact with BRS-3 receptors directly assessed by binding studies using ¹²⁵I-Peptide-1 or ¹²⁵I-Bantag-1 as the radioligand, a number were compared under identical binding conditions (Fig. 6; Table 3). In general, there was a very close comparison between the affinity of each ligand for the hBRS-3 receptor assessed by the two different radioligands (Fig. 5; Table 3). Specifically, Bantag-1 had the highest affinity (0.27-0.40 nM) (Fig. 6: Table 3) for hBRS-3 expressed in BALB 3T3 cells with both radioligands. Peptide-1 had a 2-fold to 4-fold lower affinity than Bantag-1 (Fig. 6A; Table 3), and Peptide-7 had a 7-fold lower affinity than Bantag-1 (K_i, 2.0–2.5 nM) (Fig. 6; Table 3). Peptide-2, which differed from peptide 1 in having a D-Phe⁶ substitution for D-Tvr⁶, had a 13-fold to 29-fold decrease in affinity compared with Bantag-1 and Peptide-6, with a similar structure to Peptide-7 but with an (S) configuration, was 28-fold to 117-fold lower than Bantag-1 for both radioligands (Fig. 6A; Table 3). Peptide-15, which differed from Peptide-6 by the removal of a 4-Cl group on Apa¹¹, had a 50-fold to 118-fold lower affinity than Bantag-1 with both radioligands (K_i, 20–31.5 nM) (Fig. 6A; Table 3). The remaining peptides (peptides 34, 54, and 68) had little to no affinity for BRS-3 (>400 nM) assessed by both radiolabeled ligands, as did the selective GRPR agonist GRP or the selective NMBR agonist NMB (Fig. 6; Table 3).

Determination of Relative Receptor Numbers for Various BRS-3 Receptors on Various Cancer Cells Using ¹²⁵I-Peptide-1 and ¹²⁵I-Bantag-1

Assessing the presence of BRS-3s is complicating by the occurrence of multiple BnRs on most tissues and the nonselectivity of ¹²⁵I-Peptide 1 for hBRS-3 receptors, which has been used in most studies (Pradhan et al., 1998; Reubi et al., 2002; Jensen et al., 2008). Until this present study, the only readily available method to directly determine hBRS-3 presence/number on tissues was to use ¹²⁵I-Peptide-1 with a GRPR- and NMBR-selective ligand to selectively block binding to the NMBR/GRPR receptor, which might also be present in significant numbers to BRS-3-3 (DeMichele et al., 1994; Reubi et al., 2002; Weber, 2009; Moreno et al., 2018). To assess the possible increased usefulness of ¹²⁵I-Bantag-1, compared with the presently used method, we compared the results of ¹²⁵I-Bantag-1 binding with the previously generally used subtraction method with ¹²⁵I-Peptide-1 to identify and quantitate BRS-3 on a number of lung cancer cells, which can contain multiple BnRs (Moreno et al., 2018). We studied six different lung cancer cell lines (N417, H69 are small cell lung cancer cells; and H520, H720, H727, and H460, which are non-small cell lung cancer) because it has been previously reported they contained different amounts of BnRs including hBRS-3 (Fathi et al., 1993; DeMichele et al., 1994; Siegfried et al., 1999; Moreno et al., 2018). To address this study, we compared the binding of saturating concentrations of radioligand ¹²⁵I-Peptide-1 (2.5 nM) and ¹²⁵I-Bantag-1 (2.5 nM) (Fig. 7) alone or in combination with a GRPR-selective ligand, methyl ester Bn-(6-13) (ME) (Wang et al., 1990b), and a NMBR-selective ligand, PD (Ryan et al., 1999; Tokita et al., 2001; Gonzalez et al., 2009), to inhibit binding to hGRPR and hNMBR (Fig. 7B).

Ability of various agents to alter binding of $^{125}\mbox{I-Bantag-1}$ to hBRS-3 receptor

TABLE 3

Comparison of affinities of various BnR agonists and antagonists for hBRS-3 containing cells determined from ¹²⁵I-Peptide-1 or ¹²⁵I-Bantag-1 binding

hBRS-3 stably transfected into BALB 3T3 cells (1×10^6 cells/ml) (Ryan et al., 1998b; Sancho et al., 2010) were incubated with 50 pM ¹²⁵I-Peptide-1 or ¹²⁵I-Bantag-1 for 60 minutes at 22°C. In each experiment, each value was determined in duplicate, and values given are the mean and S.E.M. from at least three separate experiments. Data are calculated from dose-inhibition curves shown in Fig. 5 for each peptide using a nonlinear regression curve-fitting program (Prism). Each value is the mean \pm S.E.M. from at least three experiments. Bantag-1 was described in Moreno et al. (2013), Guan et al. (2010), Feng et al. (2011); #1 in Mantey et al. (1997); #2 in Mantey et al. (1997); #6 in Mantey et al. (2004); #77 in Mantey et al. (2004); #34 in Boyle et al. (2005); #54 in Weber et al. (2005); and #68 in Weber et al. (2002, 2003).

Dentide No	Peptide Structure	<i>K</i>	Ki	
Pepude No.		¹²⁵ I-Peptide-1	¹²⁵ I-Bantag-1	
		nl	A.	
Bantag-1	Boc-Phe-His-4-amino-5-cyclohexyl-2,4,5-trideoxypentonyl- Leu-(3-dimethylamino) benzylamide N-methylammonium trifluoroacetate	0.40 ± 0.25	0.27 ± 0.01	
1	$[D-Tyr^{6}, \beta-Ala^{11}, Phe^{13}, Nle^{14}]Bn-(6-14)$	1.58 ± 0.09	0.68 ± 0.02	
2	$[D-Phe^{6}, \beta-Ala^{11}, Phe^{13}, Nle^{14}]Bn-(6-14)$	5.50 ± 0.29	7.94 ± 0.38	
6	[D-Tyr ⁶ (S), Apa ¹¹ -4Cl, Phe ¹³ , Nle ¹⁴]Bn-(6-14)	110 ± 6	31.6 ± 0.6	
7	[D-Tyr ⁶ (R), Apa ¹¹ -4Cl, Phe ¹³ , Nle ¹⁴]Bn-(6-14)	2.52 ± 0.13	2.04 ± 0.06	
15	[D-Tyr ⁶ (S), Apa ¹¹ , Phe ¹³ , Nle ¹⁴]Bn-(6-14)	20.0 ± 1.1	31.5 ± 1.1	
34	Ac-Phe, Trp, Ala, His (t-Bzl), Nip, Gly, Arg-NH ₂	162 ± 8	74 ± 3	
54	H-D-Phe, Gln, D-Trp, Phe-NH ₂	>1000	> 1000	
68	3-phenyl-propionyl-Ala, D-Trp, NH (CH ₂) C ₆ H ₅	>1000	467 ± 11	
NMB	Neuromedin B	>1000	> 1000	
GRP	Gastrin-releasing peptide	>1000	>1000	

In binding studies with ¹²⁵I-Peptide-1 and the peptide antagonist Bantag-1 to block the hBRS-3 binding site, the receptor number of hGRPR/hNMBR in H720 and H727 was 8-fold higher than in N417, H69, H520, or H460 (hGRPR/hNMBR receptor numbers: H720, 1.34; H727, 1.45; N417, 0.06; H69, 0.02; H520, 0.07; and H460, 0.14 fmol/10⁶ cells) (Fig. 7A). When these lung cancer cells were incubated with ¹²⁵I-Peptide-1, and the hGRPR/hNMBR inhibitors ME and PD, coinhibitors that block binding selective to GRPR/NMBR, allowing assessment only of BRS-3-receptor interaction, the receptor number of BRS-3 was 16-fold and 6-fold higher in N417 and H69 than in H520, H720, H727, or H460 (hBRS-3 receptor numbers: N417, 0.720; H69, 0.15; H520, 0.04; H720, 0.01; H727, 0.04; and H460, 0.03 fmol/10⁶ cells) (Fig. 7B). In contrast, when the lung cancer cells were incubated only with the ¹²⁵I-Bantag-1, N417, H69, and H520 had higher amounts of BRS-3 receptor number than H720. H727, and H460 (hBRS-3 receptor numbers: N417, 1.13; H69, 0.63; H520, 0.40; H720, 0.32; H727, 0.38; and H460, 0.43 fmol/10⁶ cells) (Fig. 7C). The results demonstrate the advantage of the ease of use of the selective BRS-3 ligand ¹²⁵I-Bantag-1, which does not require the indirect determination by the subtraction of GRPR/NMBR.

Assess Whether BRS-3 Receptor Number Determined Using ¹²⁵I-Bantag-1 Correlates with Magnitude of Expression of BRS-3 mRNA, or Maximal Increase in Phospholipase C Activation (in Cytosolic Ca²⁺ or [3H]IP) in Lung Cancer Cell Lines

In a recent study (Moreno et al., 2018), the BRS-3 agonist MK-5046 was reported to activate phospholipase C, causing increases in cytosolic Ca^{2+} and IP generation in lung cancer cells possessing BRS-3. To determine whether the BRS-3 receptor number correlated with the magnitude of phospholipase C activation or the mBRS-3 mRNA expression, we correlated BRS-3 receptor number assessed using ¹²⁵I-Bantag-1 with each of these variables in a number of lung cancer cell lines. As seen in Fig. 8, the BRS-3 receptor number determined in eight lung

cancer cell lines correlated closely (r = 93-99, P < 0.001) with the expression level of BRS-3 mRNA determined by quantitative PCR (Moreno et al., 2018) (Fig. 8A), the maximal increase in stimulated [³H]IP (Fig. 8B) and the maximal stimulated increase in cytosolic Ca²⁺ (Fig. 8C).

Assess Whether Using ¹²⁵I-Bantag-1, BRS-3 Receptor Number Can Be Determined in Lung Cancer Cell Line Membranes and How These Values Compare to Results in Various Normal Rat Tissues Reported to Have High BRS-3 Expression of BRS-3 Either in Rats or Other Species

Four different human lung cancer cell lines having different levels of expression of BRS-3 were studied (Table 4). In each case using ¹²⁵I-Bantag-1, BRS-3 receptors could be detected on the cell membranes (Table 4). However, the BRS-3 receptor densities varied markedly from 1.34 ± 0.81 fmol/mg protein for H838 cells to 64.4 ± 14.2 fmol/mg protein in N417 membranes, which is similar to densities seen in the whole-cell studies shown in Fig. 7.

In rats and other species, high levels of BRS-3 expression are reported by various methods (immunohistochemistry, mRNA studies, and one binding study in the CNS using [³H]-Bag-2) (Fathi et al., 1993; Jennings et al., 2003; Sano et al., 2004; Porcher et al., 2005; Guan et al., 2010; Zhang et al., 2013) in various brain nuclei, testis, and small intestine (especially duodenum). We examined five of these (brain cortex, mid-brain, cerebellum, testis, and duodenum) using 125I-Bantag-1 binding and found that they all possessed BRS-3 receptors (Table 4). In these tissues, the BRS-3 densities overlapped with the levels seen in many of the lung cancer cell lines.

Discussion

The purpose of this study was to attempt to develop and characterize a specific, high-affinity radioligand that could be used for BRS-3 localization and pharmacologic studies in both normal and pathophysiological conditions. This is important



Fig. 6. The ability of various Bn receptor ligands to inhibit binding of the radiolabeled agonist 125I-Peptide-1 (A) or the radiolabeled antagonist ¹²⁵I-Bantag-1 (B) to hBRS-3/BALB (1×10^6 cells/ml). Cells were incubated with 50 pM ¹²⁵I-Peptide-1 or ¹²⁵I-Bantag-1 with or without increasing the concentrations of various unlabeled Bn receptor ligands for 60 minutes at 22°C. Results are expressed as the percentage of saturable binding with no addition. The results are the mean and S.E.M. from at least three separate experiments, and in each experiment the data points were determined in duplicate. For abbreviations, see the legends of Figs. 1–3 and Table 3.

for a number of reasons. First, the BRS-3 receptor is receiving increased attention (Gonzalez et al., 2015b; Maruyama et al., 2017; Nio et al., 2017; Xiao et al., 2017; Pinol et al., 2018) because of its prominent role in causing diabetes, hypertension, and obesity by regulating body weight and energy maintenance, body temperature control, and control of metabolic homeostasis, particularly insulin/glucose control, as evidenced from studies of BRS-3 knockout mice (Ohki-Hamazaki et al., 1997; Ladenheim et al., 2008; Majumdar and Weber, 2012a; Lateef et al., 2014; Gonzalez et al., 2015b). Studies using primarily non-ligand receptor localization methods (immunohistochemistry, BRS-3 mRNA methods) (Fathi et al., 1993; Jennings et al., 2003; Sano et al., 2004; Porcher et al., 2005; Guan et al., 2010; Zhang et al., 2013; Maruyama et al., 2017; Xiao et al., 2017) provide evidence that BRS-3 is highly expressed in hypothalamic nuclei associated with metabolic maintenance, and that glutamatergic neurons in this area (Xiao et al., 2017) are particularly important. Furthermore, recent studies (Das et al., 2009; Moreno et al., 2013, 2018) demonstrate that BRS-3 is overexpressed in



Fig. 7. The determination of relative receptor numbers for various Bn receptors on lung cancer cells using the radiolabel agonist ¹²⁵I-Peptide-1 or the radiolabeled antagonist ¹²⁵I-Bantag-1. The lung cancer cells (5×10^6 cells/ml) N417, H69, H520, H720, H727, and H460 were incubated with 2.5 nM radioligand with or without 1 μ M Peptide-1 or Bantag-1. In (A), the cells were incubated with ¹²⁵I-Peptide-1, which binds to all three BnRs, in the presence of 1 μ M Bantag-1 to block bind to BRS-3, thus showing only the hNMBR/hGRPR cell receptor number. In (B), ¹²⁵I-Peptide-1 was incubated with 1 μ M methyl ester Bn-(6-13) (a selective GRPR ligand) and 1 μ M PD (a selective NMBR ligand) to inhibit binding to hGRPR and hNMBR, and this reflects only binding to the BRS-3 receptor. In (C), only 2.5 nM ¹²⁵I-Bantag-1 was added, which only binds to hBRS-3 and thus directly assesses hBRS-3 receptor number. The results are the mean and S.E.M. from at least three separate experiments, and in each experiment the data points were determined in duplicate. For abbreviations, see the legends of Figs. 1, 3, and 6.



Fig. 8. The correlation of BRS3 receptor number with the ability to activate phospholipase C (Ca²⁺, [³H]IP) or the magnitude of BRS3 mRNA in various lung cancer cell lines. The magnitude of BRS3 mRNA (A), MK-5046-stimulated increases in [³H]IP (B), or increases in MK-5046-stimulated Ca²⁺ were correlated with BRS3 receptor number, as determined by saturation binding. The lung cancer cells (5 imes 10⁶ cells/ml) N417, H69, H520, H720, H727, A549, H838, and H1299 were incubated with 2.5 nM 125 I-Bantag-1 with or without 1 μ M Bantag-1, and the total BRS-3 receptor number was calculated using the specific activity. For lung cancer cells not previously reported (Moreno et al., 2018) (H838, A549, H1299), the maximal stimulation of BRS-3 by the BRS-3-specific agonist MK-5046 (10 nM) was determined (Moreno et al., 2018) and correlated with the BRS-3 receptor number, as described in Methods. For the remaining lung cancer cells, the data for change in Ca²⁺ [³H]IP as well as the BRS-3 mRNA expression determined by quantitative PCR data in all of the cancer cell lines was from Moreno et al. (2018). The results are the mean and S.E.M. from at least three separate experiments, and in each experiment the data points were determined in duplicate.

TABLE 4

BRS-3 receptor density in selected human lung cancer cell lines and normal rat tissues

Membranes were made from the indicated tissues as described in *Methods*. The density of BRS-3 binding sites was determined as described in the Fig. 7 legend using 2.5 nM 125 I-Bantag-1 binding. Each value is the mean \pm S.E.M. from at least three experiments.

Membrane tissue	BRS-3
	fmol/mg protein
Lung cancer cell lines	
N417	64.4 ± 14.2
H69	45.0 ± 10.1
H520	$7.4~\pm~0.6$
H838	1.34 ± 0.81
Normal rat tissues	
Cortex (brain)	12.0 ± 2.9
Mid-brain	21.0 ± 2.5
Cerebellum	15.0 ± 1.5
Testis	18.1 ± 0.6
Duodenum	39.1 ± 0.7

numerous tumors and has prominent tumoral growth effects, similar to those seen with the other mammalian BnRs (i.e., GRPR, NMBR) (Fathi et al., 1993; Reubi et al., 2002; Ramos-Álvarez et al., 2015; Moreno et al., 2016, 2018). In addition, recent studies demonstrate that the BRS-3 has important roles in a number of different tissues, including regulation of behavioral changes in the CNS (Yamada et al., 2000a,b), the regulation of metabolic changes in skeletal muscle (Ramos-Álvarez et al., 2014; Gonzalez et al., 2015a), important metabolic effects in adipocytes (Nakamichi et al., 2004), the maintenance of bronchial epithelial homeostasis (Liu et al., 2018), and motility effects in the GI tract (Porcher et al., 2005). Second, at present no generally available specific, highaffinity ligand is available to study the expression, location, or pharmacology of the BRS-3 receptor. A ligand, ¹²⁵I-Peptide-1 (in this study) [(¹²⁵I-D-Tyr⁶, β -Ala¹¹, Phe¹³, Nle¹⁴]Bn-(6–14)], which interacts with high affinity with hBRS-3 has been described and used in some receptor localization studies (Fleischmann et al., 2000; Reubi et al., 2002; Waser et al., 2007), but it has a number of disadvantages that limit its usefulness. This ligand also interacts with high affinity with GRPR and NMBR (Mantey et al., 1997; Pradhan et al., 1998), which frequently coexist with BRS-3 on the same cells (Fathi et al., 1993; Ryan et al., 1998b; Moreno et al., 2018). In addition, it has low affinity for mouse and rat BRS-3 and thus is not useful in these commonly used experimental animals (Liu et al., 2002; Uehara et al., 2011; Ramos-Álvarez et al., 2016). The high affinity of this ligand for all three BnRs requires that in receptor studies for BRS that a subtraction approach be used by adding specific ligands for the GRPR and NMBR to block the binding to these receptors (Fleischmann et al., 2000; Reubi et al., 2002; Waser et al., 2007), leaving the BRS-3 binding only; however, this is cumbersome and increases the inaccuracy. In one study (Guan et al., 2010), a tritiated BRS-3 agonist, [³H]-Bag-2, was described, but it is not well characterized and is not generally available. Third, studies, especially in the CNS in nonhypothalamic areas, have shown important differences between the results of possible BRS-3 locations assessed by BRS-3 mRNA studies and BRS-3 antibodies, leading authorities in recent reviews to question the specificity of available BRS-3 antibodies (Gonzalez et al., 2015b; Xiao and Reitman, 2016). Last, even though the endogenous ligand remains unknown at

present, nevertheless, because of the important roles of BRS-3 outlined above in energy, glucose/insulin homeostasis, and obesity, a number of pharmaceutical companies/investigators have recently developed selective agonists/antagonists (e.g., MK-5046, chiral diazepines, Bag-1/2, Bantag-1) (Guan et al., 2010; Feng et al., 2011; Matsufuji et al., 2014, 2015; Gonzalez et al., 2015b; Moody et al., 2015; Kiyotsuka et al., 2016; Ramos-Álvarez et al., 2016). These new compounds increasingly are being used to investigate the role of BRS-3 in normal physiologic function as well in pathophysiological conditions. This is already starting to increase the need for specific radioligands to study the location/pharmacology of the BRS-3 in these processes.

A number of our results support the conclusion that in this study we have developed a highly selective, specific ligand, 125-Bantag-1, that should be generally useful for the characterization and study of the pharmacology of the BRS-3 receptor. First, we find that Bantag-1 is highly selective (>10,000-fold) for hBRS-3 over the other closely related mammalian BnRs (i.e., GRPR, NMBR), which is consistent with the results of previous studies (Guan et al., 2010; Moreno et al., 2018). Furthermore, this selectivity is preserved for the radiolabeled Bantag-1, because we find that 125 I-Bantag-1 does not bind to either GRPR or NMBR. Second, we find high specificity of ¹²⁵I-Bantag-1 for BRS-3. This result is supported by findings that ligands of other unrelated GI hormones/neurotransmitters, ligands specific for GRPR/NMBR or ligands of a Drosophila receptor related to mammalian BRS-3 (i.e., CCHamides) (Matsufujiet al., 2014, 2015) all do not alter the interaction of ¹²⁵I-Bantag-1 with the BRS-3 receptor. Third, dose-inhibition curves and BRS-3 receptor affinities for a number of different ligands reported to interact with the BRS-3, computed from ¹²⁵I-Bantag-1 binding, show very close agreement with those calculated from the well-characterized panBnR peptide agonist ligand ¹²⁵I-Peptide-1 (Mantey et al., 1997; Ryan et al., 1998b; Gonzalez et al., 2015b). Fourth, previous studies have demonstrated that the peptide antagonist Bantag-1, in contrast to the peptide agonist Peptide-1, has high affinity for mouse and rat BRS-3 receptors (Guan et al., 2010; Feng et al., 2011). These results demonstrate that ¹²⁵I-Bantag-1 will also be a valuable tool for assessing BRS-3 in the most commonly used laboratory animals, rats/mice, whereas¹²⁵I-peptide-1 is not. Fifth, that ¹²⁵I-Bantag-1 binding studies could be pharmacologically useful for characterizing BRS-3 receptors was shown by the ability to allow clear quantification of BRS-3 receptor expression. This allowed us to establish a close correlation of the BRS-3 receptor number with the amount of BRS-3 mRNA, as well as the maximal magnitude of BRS-3 activation of phospholipase C in lung cancer cells. In addition, it allowed BRS-3 receptor expression to be studied in both normal tissue membranes as well as in those from cancer cells.

In general, in this study the interaction of the radiolabeled antagonist ¹²⁵I-Bantag-1 with BRS-3 receptors was similar to that seen with the radiolabeled agonist ¹²⁵I-Peptide 1, which has similarities and differences from other studies comparing the binding of these two classes of ligands on other BnRs, as well as to other GPCRs. This conclusion is supported by the similar kinetics of association of both BRS-3 ligands with the BRS-3; their similar maximal binding under identical conditions to these cells; their similar dissociation kinetics; as well as the similar dose-inhibition curves of various putative BRS-3 ligands with both BRS-3 ligands. This result differs from a previous study (Mantey et al., 1997) comparing a radiolabeled GRPR-radiolabeled peptide antagonist and agonist binding to GRPR receptors, where the kinetics differed between the two, with the antagonist showing more rapid association kinetics, and more rapid and greater time-dependent dissociation than the agonist. However, these GRPR study results were similar to those of the present study in showing that both radiolabeled ligands gave similar affinities for the GRPR (Mantey et al., 1997). The results that similar affinities for BRS-3 for each of the eight putative selective BRS-3 agonists examined were found with both ligands in the present study differs from findings of radiolabeled agonist/antagonist binding with cholecystokinin, gastrin, adrenergic, and luteinizing hormone-releasing hormone receptors (Mantey et al., 1997).

A somewhat surprising finding in the present study is that the radiolabeled antagonist is internalized to the same degree as the radiolabeled agonist. Generally, GPCR peptide antagonists are poorly and slowly internalized, if at all, whereas agonists are extensively and rapidly internalized (Mantey et al., 1993; Ginj et al., 2006; Moreno et al., 2016; Dalm et al., 2017). The above generalization is also true for BnRs, where numerous studies have reported that most BnR peptide antagonists are poorly internalized, if at all (Mantey et al., 1993; Cescato et al., 2008; Varasteh et al., 2013; Nock et al., 2017; Popp et al., 2017; Cheng et al., 2018). The internalization of radiolabeled Bantag-1 in our studies was not due to the possibility that Bantag-1 was functioning as a partial agonist, because in previous studies it has been reported to have no agonist activity at the BRS-3 receptor for the activation of phospholipase C, as assessed by either stimulation of the generation of phosphoinositides (Moreno et al., 2013) or mobilization of cellular calcium (Guan et al., 2010). Furthermore, Bantag-1 functioned as fully competitive antagonist of the BRS-3 receptor activation for stimulating IP generation (Moreno et al., 2013).

In addition to its use for pharmacological studies of the BRS-3 in normal physiologic and pathologic conditions, radiolabeled Bantag-1 may prove of value in tumor localization and for receptor-directed cytotoxicity for tumors. Recently, numerous studies have demonstrated that a number of common neoplasms overexpress BnRs (i.e., prostate, breast, lung, pancreas, and CNS tumors) (Reubi et al., 2002; Reubi, 2013; Moreno et al., 2016), and in some tumors such as lung cancers, BRS-3 is the most frequently overexpressed and is expressed in the highest numbers (Moreno et al., 2018). A number of recent studies report the ability of BnR radioligands to image various human tumors in humans, particularly breast and prostate, using high-affinity GRPR ligands (Sancho et al., 2011; Moreno et al., 2016; Ferreira et al., 2017; Cheng et al., 2018; Zang et al., 2018), and with the high expression of BRS-3 in lung tumors (Moreno et al., 2018) a similar approach might be used with radiolabeled Bantag-1. Furthermore, using ¹⁷⁷Lu- or ⁹⁰Y-labeled peptides, peptide-radioreceptor therapy has been shown to be effective and has been recently approved for the treatment of neuroendocrine tumors with radiolabeled somatostatin receptor ligands (Strosberg et al., 2016; Ito and Jensen, 2017). Because of this, there is now considerable interest in using a similar approach with radiolabeled GRP agonists/antagonists for a number of cancers overexpressing these receptors, particularly, prostate and breast cancer (Moreno et al., 2016; Reynolds et al., 2016; Maina et al., 2017). This raises the possibility that a similar approach be

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used for radiolabeled Bantag-1 for lung cancer, because of its frequent high expression in these tumors (Moreno et al., 2018).

In conclusion, we describe and characterize a novel, highaffinity, selective, and specific radiolabeled ligand, ¹²⁵I-Bantag-1, that can be easily prepared and thus should be generally useful for pharmacological characterization of BRS-3 receptors, both in peripheral tissues and the CNS. The availability of this ligand should be useful in exploring the role of this orphan receptor in energy/insulin-glucose homeostasis, body temperature control, GI motility, and allergic lung diseases, and their possible theranostic use to selectively image and deliver cytotoxic agents to neoplasms overexpressing this receptor.

Authorship Contributions

Participated in research design: Ramos-Alvarez, Lee, Mantey, and Jensen

Conducted experiments: Ramos-Alvarez, Lee, Mantey, and Jensen. Contributed new reagents or analytic tools: Ramos-Alvarez, Lee, Mantey, and Jensen.

Performed data analysis: Ramos-Alvarez, Lee, Mantey, and Jensen.

Wrote or contributed to the writing of the manuscript: Ramos-Alvarez, Lee, Mantey, and Jensen.

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