

Original Article

A high-throughput screening system for G-protein-coupled receptors using β -lactamase enzyme complementation technology

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Aim: To establish a system for monitoring the activation of G-protein-coupled receptors (GPCRs) using β -lactamase enzyme fragment complementation (EFC) technology.

Methods: Two inactive β -lactamase deletion fragments, bla(a) and bla(b), were fused to β -arrestin and GPCR, respectively. A stable cell line named HEK/293- β 2a2, which expressed two fusion proteins, GPCR/bla(b) and β -arrestin2/bla(a), was generated under antibiotic selection. A natural compound library of high performance liquid chromatography (HPLC)-fractionated samples from the ethanol extracts of Chinese medicinal herbs was used for high-throughput screening (HTS) of β 2-adrenoceptor (β 2AR) agonists against the cell line HEK/293- β 2a2. The interested hits were validated by the measurement of second-messenger cyclic adenosine monophosphate (cAMP) production.

Results: The stable cell line HEK/293- β 2a2 responded to β 2AR agonist/antagonist in a dose-dependent manner. The EC₅₀ value obtained for isoproterenol was 15.5 nmol/L, and the IC₅₀ value obtained for propranolol was 51.9 nmol/L. Furthermore, HTS was performed to identify β 2AR agonists from the natural compound library we established. The Z' factor value was determined to be 0.68. Three hits were identified from primary screening and found to be as potent as isoproterenol in a cAMP assay.

Conclusion: A cell-based high-throughput functional assay was established to directly monitor the activation of GPCRs based on the interaction between agonist-activated GPCR and β -arrestin using β -lactamase EFC technology, which can be used to search for leads in the natural compound library.

Keywords: G-protein-coupled receptor; β -lactamase; enzyme fragment complementation; high-throughput screening

Acta Pharmacologica Sinica (2010) 31: 1618–1624; doi: 10.1038/aps.2010.154; published online 22 Nov 2010

Introduction

G-protein-coupled receptors (GPCRs) are the largest known family of transmembrane receptors. They bind to and transduce signals for a huge variety of endogenous ligands, and they are involved in most physiological functions, such as metabolism, neuronal transmission, embryonic development, hormonal homeostasis and cancer invasion^[1,2]. Of all currently marketed drugs, over 30% target GPCRs^[3], and the development of new technologies for GPCR-based drug discovery promises to keep this field active for years to come.

Recently, several cell-based drug discovery strategies were developed using β -arrestin as a surrogate marker for GPCR

function. β -arrestins are ubiquitously expressed in all cells and tissues. They bind with high affinity to the agonist-activated GPCR carboxyl-terminal tail. Assays developed based on GPCR- β -arrestin interaction, such as transfluor assay^[4,5], bioluminescence resonance energy transfer assay^[6], PathHunter™ β -arrestin Assays^[7] and Tango assay^[8] were widely used in the investigation of many GPCRs^[9–15]. These achievements have demonstrated that monitoring GPCR- β -arrestin interaction can offer a direct indication of GPCR activation.

Here we describe a cell-based protein-protein interaction assay based on β -lactamase complementation technology that uses GPCR and β -arrestin as protein partners to monitor GPCR activation. The β -lactamase-based enzyme fragment complementation (EFC) assay was pioneered in 2002^[16,17]. The properties of β -lactamase provide significant advantages in mammalian cells. β -lactamase is monomeric and relatively

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Received 2010-05-16 Accepted 2010-07-29

cells per well in a 96-well black matrigel-coated microplate with a clear-bottom. The cells were cultured for 24 h before treatment. Culture media were replaced with serum-free DMEM containing 0.01 g/L diluted testing compounds. Positive and negative controls were designed in each plate. After incubation at 37 °C for 5 min, β -lactamase activity was quantitated with CCF4/AM.

cAMP assay

HEK/293- β 2a2 cells were plated (4×10^4 cells/well) in a 96-well plate for 24 h. Compounds were diluted with phosphate-buffered saline and incubated with the cells for 15 min prior to measuring the cAMP level with cAMP-Glo™ Assay (Promega, Madison, WI, USA), according to product instructions. Luminescence was determined using an Envision 2100 (PerkinElmer, Waltham, MA, USA).

Data analysis and calculations

Data were presented as mean \pm SD. Statistical significance was evaluated with Student's *t*-test, and the difference was considered statistically significant at $P < 0.05$. The Z' factor was calculated as described previously by Zhang *et al*^[24]. For concentration-response determinations, a curve was fitted to calculate EC_{50} and IC_{50} using GraphPad Prism 5 software.

Results

Recombinant cell line generation

To study GPCR activation with a β -lactamase EFC assay, stable cell line HEK/293- β 2a2 was established by the co-transfection of plasmids encoding human β -arrestin2/bla(a) and β 2AR/bla(b), two fusion constructs (Figure 1A), into HEK-293.

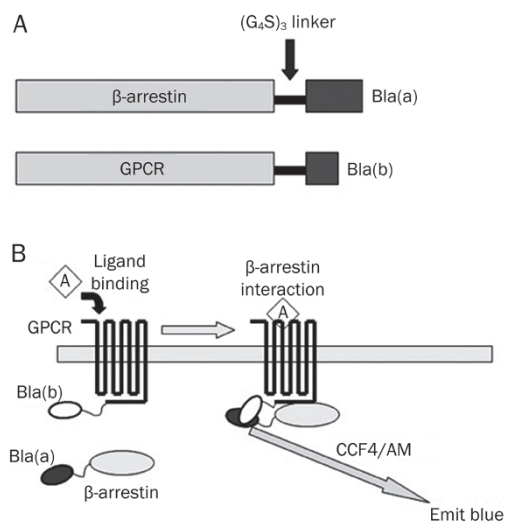


Figure 1. β -lactamase EFC based receptor/ β -arrestin interaction assay. (A) Vector construction. β -lactamase fragment was fused to target protein with a flexible $(G_4S)_3$ linker. (B) Illustration of β -lactamase EFC based receptor/ β -arrestin interaction assay. Agonist (A) activated GPCR recruits β -arrestin to the cell membrane, which brings two β -lactamase fragments proximity to reconstitute and recover its activity. The recovery of β -lactamase activity was then determined with CCF4/AM assay.

They were selected as described in materials and methods. The cell clone with the best signal to background ratio was chosen and named HEK/293- β 2a2. In this clone, β 2AR/bla(b) and β -arrestin2/bla(a), two fusion constructs, were highly expressed. The two β -lactamase deletion mutants, bla(a) and bla(b), are individually inactive and have a low affinity for each other. Ligand stimulation caused the conformation change of β 2AR and recruited β -arrestins to the cell membrane. The interaction between β 2AR and β -arrestin brought the two inactive β -lactamase fragments together to reconstitute a functional enzyme. The recovery of β -lactamase activity, as determined with the CCF4/AM assay, was a direct indicator of agonist-induced GPCR activation via the β -arrestin recruitment (Figure 1B).

Characteristics of the stable cell line HEK/293- β 2a2

Agonist stimulation in cells co-expressing β 2AR/bla(b) and β -arrestin2/bla(a) caused the complementation of two β -lactamase fragments. The reconstituted β -lactamase hydrolyzed the substrate CCF4/AM and emitted blue fluorescence when excited at 409 nm. The generation of blue fluorescent cells could be attributed to the activation of G-protein-coupled receptors (Figure 2A, top). The agonist effect could be inhibited by the β 2AR antagonist, propranolol. When pre-incubated with propranolol, the percentage of cells that emitted blue fluorescence after agonist stimulation was similar to untreated cells (Figure 2A, middle and bottom). Dose response curves for isoproterenol and propranolol were also calculated using GraphPad Prism 5 software. As shown in Figure 2B, HEK/293- β 2a2 cells responded to the known β 2AR agonist, isoproterenol, with a dose-dependent increase in β -lactamase activity. The EC_{50} value of isoproterenol was found to be 15.5 nmol/L, which is not only similar to previously reported data for HEK-293 cells (11 nmol/L) using an isoproterenol-stimulated β 2AR sequestration assay^[25] but also data obtained in C2C12 cells (27 nmol/L) using a β -galactosidase based EFC assay^[7]. Dose-dependent inhibition was demonstrated under the treatment of the β 2AR antagonist, propranolol, in the presence of 1 μ mol/L isoproterenol (Figure 2C). The IC_{50} of 51.9 nmol/L obtained for propranolol using β -lactamase EFC assay was also consistent with reported data^[26].

Screening of β 2AR agonists from natural compound library

To see if this cell-based assay was suitable for high-throughput screening, the Z' factor value was calculated as described previously by Zhang *et al*^[24]. Half of the plate was treated with 1 μ mol/L isoproterenol, and half of the plate was untreated. Four wells (A1, B1, G12, and H12) were designed for reagent control to calculate the blue/green ratio for the demonstration of β -lactamase activity. The Z' factor was 0.68 in our experiment (Figure 3A). To further assess the value of this cell model for searching agonists targeting β 2-adrenoceptor, we screened 352 HPLC-fractionated samples from the ethanol extracts of Chinese medicinal herbs in a 96-well format. Isoproterenol was used as the positive control, and 1% DMSO was used as the negative control. Cells were incubated with

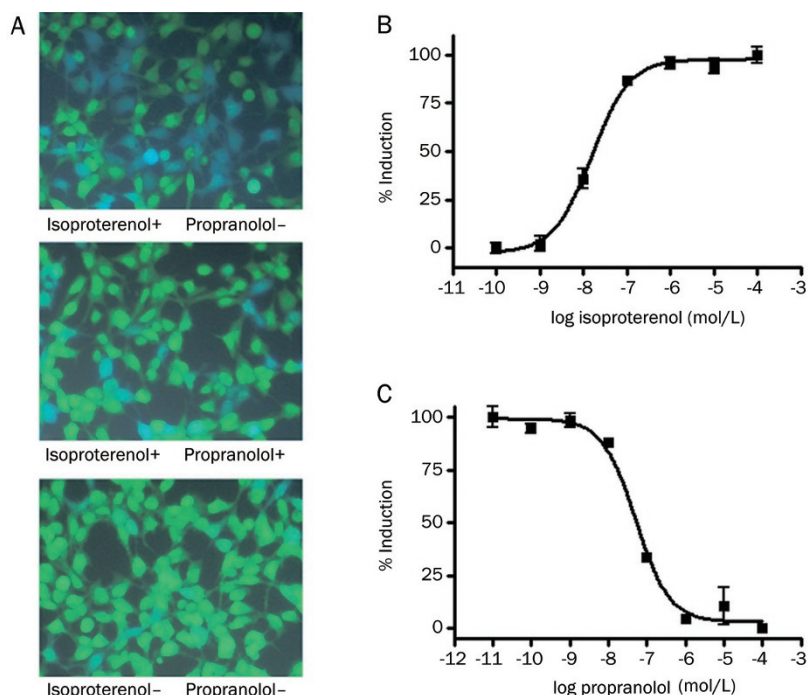


Figure 2. Characteristics of stable cell line HEK/293-β2a2. (A) Immunofluorescence assay of β-lactamase activity. Stable cell line HEK/293-β2a2 treated with 1 μmol/L isoproterenol (top); pre-incubated with 10 μmol/L propranolol and then treated with 1 μmol/L isoproterenol (middle); untreated (bottom) shown clear difference in blue cells percentage imaged by fluorescence microscope (magnification×100). (B) and (C) Dose response curves for isoproterenol and propranolol. Data represent means±SD for triplicate samples.

fractionated samples for 5 min in a serum-free medium. All wells were treated identically, and all contained 1% DMSO, a concentration which has been determined to have no adverse effect on the cells or the assay (data not shown). The fraction activity was expressed as the percentage of maximal β-lactamase induction by 1 μmol/L isoproterenol. As shown in Figure 3B, three fractions each exhibited a significant effect on the induction of β-lactamase activity (>40%). The effect of the other fractions on the induction of β-lactamase complementation is also shown in Figure 3. These fractions, which had activities below three times of standard deviation, were considered insignificant.

Hits validation

To confirm the activity of the primary screen hits, we repeated the receptor/β-arrestin interaction assay in triplicate (Figure 4A). Fractions 10-B11 and 10-C11 were identified to have >50% of maximum induction while fraction 10-D11 showed >40% of maximum induction. Three positive hits were further validated with the melanocortin receptor 4 (MC4R) cell line, which was also generated based on the β-lactamase complementation assay. As expected, they all had no agonist effect on the MC4R cell line (Figure S1). β2-adrenoceptor is a classic G-protein-coupled receptor used for investigation. Once the ligand binds to a β2-adrenoceptor, the receptor will change the conformation and couple to a heterotrimeric G-protein, resulting in an elevation of cAMP levels. Thus, hits obtained from the primary screening were also validated with another functional assay to measure the induction of cAMP expression. As shown in Figure 4B, three fractions had nearly the same intensity of cAMP induction as the positive control, isoproterenol. These two experiments confirm the effect of the three fractions on β2-adrenoceptor activation. Three positive fractions

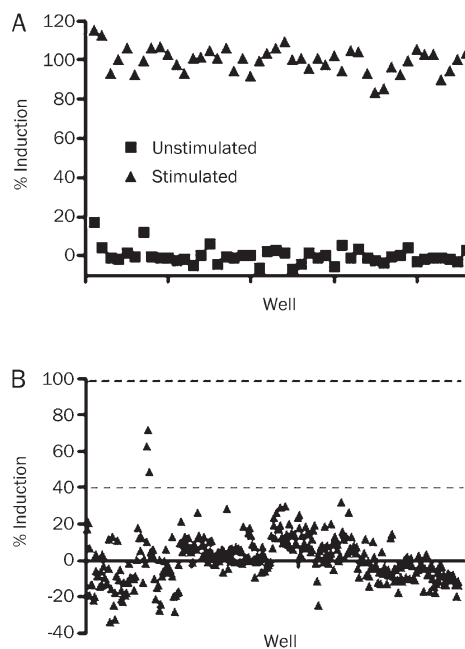


Figure 3. Z' factor determination and high-throughput screening. (A) Z' factor determination. This experiment was representative of three experiments. (B) High-throughput screening from natural compound library. Each triangle represents the reading from one well. The induction of β-lactamase activity was expressed as a percentage of isoproterenol induction (performed in the same plate).

were consecutively obtained from *Fructus Aurantii Immaturus*. HPLC chromatograms of fractions 10-B11, 10-C11, and 10-D11 are shown in Figure 5. In addition, the methyl thiazolyl tetrazolium (MTT) test showed that fraction treatment at test-

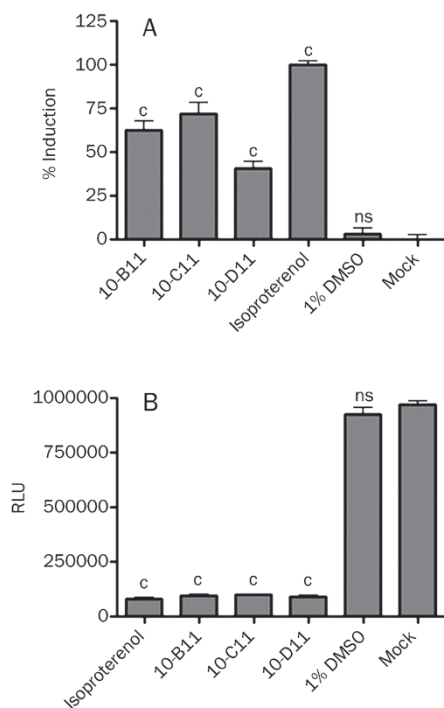


Figure 4. Positive hits validation. (A) Repeat of the receptor/ β -arrestin interaction assay. Data represent means \pm SD for triplicate samples. (B) Positive hits validation by cAMP quantitation. Three hits have similar potential compared with positive control isoproterenol in elevating cAMP expression. Data represent means \pm SD for triplicate samples. $^cP < 0.01$ vs Mock.

ing condition did not decrease cell viability (data not shown). All these results suggest that a cell-based functional assay system based on β -lactamase complementation technology has high selectivity for agonist identification. It is a useful tool for high-throughput screening and monitoring GPCR activation.

Discussion

We established a cell-based high-throughput screening assay system for monitoring G-protein-coupled receptor activation using β -lactamase complementation technology. Detection of β -lactamase activity recovery with CCF4/AM could be observed and imaged in intact cells by a fluorescence microscope. This system is also adapted for sorting or analyzing by flow cytometry. The two fragments of β -lactamase are only about 10 kDa and 19 kDa. They are relatively smaller than many other tag proteins, and their size decreases the possibility of altering the function of chimeric fusion proteins. The small size, monomeric characteristic, and availability of a cell-permeable fluorescent substrate suggest that a system based on the β -lactamase enzyme complementation has wide potential in biological studies and drug discovery.

GPCR activation leads to signal transduction through two mediators: G-proteins and β -arrestins. Agonists bind to GPCR and activate heterotrimeric G-proteins. This activation promotes second-messenger signaling and subsequently regulates different physiological functions. Another pathway

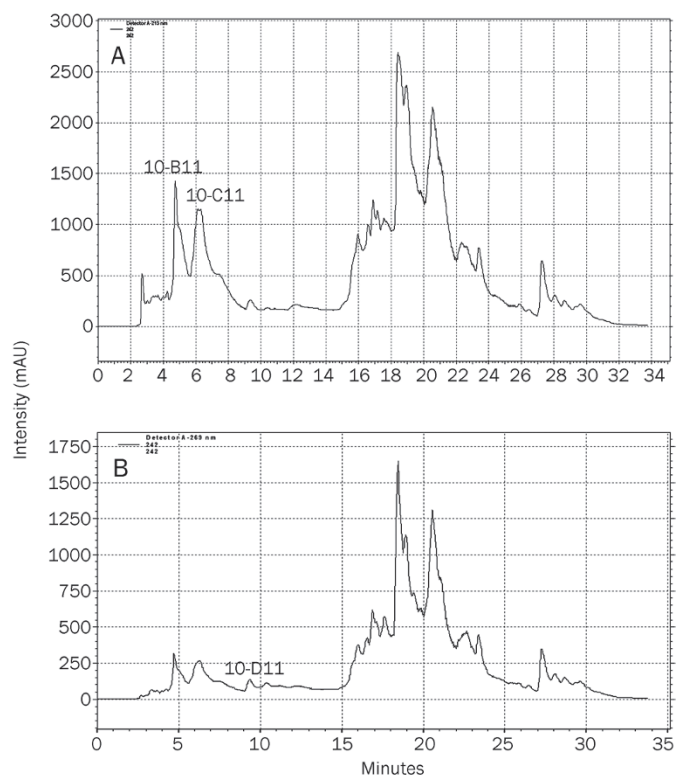


Figure 5. HPLC chromatograms of three positive fractions. (A) Fraction 10-B11 and 10-C11 (Detector A-215 nm). (B) Fraction 10-D11 (Detector A-269 nm). The arrows indicated the positions of active fractions.

for GPCR signal transduction is mediated by β -arrestins. The activated receptors can be phosphorylated by G-protein-coupled receptor kinases and recruit β -arrestins, which terminate signal transduction through G-proteins. Receptors move into clathrin-coated pits with β -arrestins from cell membranes, which cause receptor internalization and desensitization. Meanwhile, β -arrestins also serve as a scaffold for numerous signaling molecules that are stimulated by receptor agonists, which initiate a second parallel set of signals^[27]. Traditionally, it was thought that ligand binding stimulated or inhibited all receptor functions to an equal extent, but in recent years, researchers have found “biased ligands” in many GPCRs^[28–30]. Perfect biased ligands stimulate one receptor activity without stimulating other known receptor activity. The development of β -arrestin-biased ligands might provide opportunities for novel therapies^[31]. Our cell-based model provided a tool for the measurement of agonist-promoted interactions between GPCRs and β -arrestins. Along with assays that measure second-messenger production (for example, Ca^{2+} or cAMP), our model remains promising in the search for a biased ligand specific to the GPCR signal pathway that can activate only G-protein- or β -arrestins-mediated signal transduction.

Traditional Chinese medicinal herbs are effective in the treatment of many diseases. In this study, we screened a traditional Chinese medicinal herb library for hits that could activate β_2 -adrenoceptor, recruit β -arrestin and cause the

complementation of a β -lactamase fragment. Three hits were found from 352 fraction samples. Validated with the measurement of cAMP production, the three herbal extracts are each possible agonists of β 2-adrenoceptor.

In summary, our achievement demonstrated that a cell-based assay using β -lactamase complementation technology was suitable for ligand screening. This assay was also validated on two other GPCRs: melanocortin receptor 4 and dopamine receptor 5 (data not shown). Our approach may provide an alternative strategy for scientists interested in this area.

Acknowledgements

This research was supported by the National Natural Science Foundation of China (grant No 30873103). We thank Xiao-yu LIU of HD Biosciences for assistance in the preparation of Chinese medicinal herb extracts.

Author contribution

Chuan-ke ZHAO designed and performed the research, analyzed the data and wrote the paper; Qi YIN discussed and revised the paper; and Shi-you LI revised the paper.

References

- Pierce KL, Premont RT, Lefkowitz RJ. Seven-transmembrane receptors. *Nat Rev Mol Cell Biol* 2002; 3: 639–50.
- Wise A, Jupe SC, Rees S. The identification of ligands at orphan G-protein coupled receptors. *Annu Rev Pharmacol Toxicol* 2004; 44: 43–66.
- Wise A, Gearing K, Rees S. Target validation of G-protein coupled receptors. *Drug Discov Today* 2002; 7: 235–46.
- Barak LS, Ferguson SS, Zhang J, Caron MG. A beta-arrestin/green fluorescent protein biosensor for detecting G protein-coupled receptor activation. *J Biol Chem* 1997; 272: 27497–500.
- Zhang J, Barak LS, Anborgh PH, Laporte SA, Caron MG, Ferguson SS. Cellular trafficking of G protein-coupled receptor/beta-arrestin endocytic complexes. *J Biol Chem* 1999; 274: 10999–1006.
- Angers S, Salahpour A, Joly E, Hilairet S, Chelsky D, Dennis M, et al. Detection of beta 2-adrenergic receptor dimerization in living cells using bioluminescence resonance energy transfer (BRET). *Proc Natl Acad Sci USA* 2000; 97: 3684–9.
- Yan YX, Boldt-Houle DM, Tillotson BP, Gee MA, D'Eon BJ, Chang XJ, et al. Cell-based high-throughput screening assay system for monitoring G protein-coupled receptor activation using beta-galactosidase enzyme complementation technology. *J Biomol Screen* 2002; 7: 451–9.
- Barnea G, Strapps W, Herrada G, Berman Y, Ong J, Kloss B, et al. The genetic design of signaling cascades to record receptor activation. *Proc Natl Acad Sci USA* 2008; 105: 64–9.
- van Der Lee MM, Bras M, van Koppen CJ, Zaman GJ. Beta-arrestin recruitment assay for the identification of agonists of the sphingosine 1-phosphate receptor EDG1. *J Biomol Screen* 2008; 13: 986–98.
- Zhao X, Jones A, Olson KR, Peng K, Wehrman T, Park A, et al. A homogeneous enzyme fragment complementation-based beta-arrestin translocation assay for high-throughput screening of G-protein-coupled receptors. *J Biomol Screen* 2008; 13: 737–47.
- Doucette C, Vedvik K, Koepnick E, Bergsma A, Thomson B, Turek-Etienne TC. Kappa opioid receptor screen with the Tango beta-arrestin recruitment technology and characterization of hits with second-messenger assays. *J Biomol Screen* 2009; 14: 381–94.
- Ghosh RN, DeBiasio R, Hudson CC, Ramer ER, Cowan CL, Oakley RH. Quantitative cell-based high-content screening for vasopressin receptor agonists using transfluor technology. *J Biomol Screen* 2005; 10: 476–84.
- Garippa RJ, Hoffman AF, Gradl G, Kirsch A. High-throughput confocal microscopy for beta-arrestin-green fluorescent protein translocation G protein-coupled receptor assays using the Evotec Opera. *Methods Enzymol* 2006; 414: 99–120.
- Bertrand L, Parent S, Caron M, Legault M, Joly E, Angers S, et al. The BRET2/arrestin assay in stable recombinant cells: a platform to screen for compounds that interact with G protein-coupled receptors (GPCRs). *J Recept Signal Transduct Res* 2002; 22: 533–41.
- Vrecl M, Jorgensen R, Pogacnik A, Heding A. Development of a BRET2 screening assay using beta-arrestin 2 mutants. *J Biomol Screen* 2004; 9: 322–33.
- Galarneau A, Primeau M, Trudeau LE, Michnick SW. Beta-lactamase protein fragment complementation assays as *in vivo* and *in vitro* sensors of protein protein interactions. *Nat Biotechnol* 2002; 20: 619–22.
- Wehrman T, Kleaveland B, Her JH, Balint RF, Blau HM. Protein-protein interactions monitored in mammalian cells via complementation of beta-lactamase enzyme fragments. *Proc Natl Acad Sci USA* 2002; 99: 3469–74.
- Philippon A, Dusart J, Joris B, Frere JM. The diversity, structure and regulation of beta-lactamases. *Cell Mol Life Sci* 1998; 54: 341–6.
- Moore JT, Davis ST, Dev IK. The development of beta-lactamase as a highly versatile genetic reporter for eukaryotic cells. *Anal Biochem* 1997; 247: 203–9.
- Zlokarnik G, Negulescu PA, Knapp TE, Mere L, Burres N, Feng L, et al. Quantitation of transcription and clonal selection of single living cells with beta-lactamase as reporter. *Science* 1998; 279: 84–8.
- Milic M, Bao X, Rizos D, Liu F, Ziegler MG. Literature review and pilot studies of the effect of QT correction formulas on reported beta2-agonist-induced QTc prolongation. *Clin Ther* 2006; 28: 582–90.
- Li SY, Chen C, Zhang HQ, Guo HY, Wang H, Wang L, et al. Identification of natural compounds with antiviral activities against SARS-associated coronavirus. *Antiviral Res* 2005; 67: 18–23.
- Zhang YL, Zhang HQ, Liu XY, Hua SN, Zhou LB, Yu J, et al. Identification of human dopamine receptors agonists from Chinese herbs. *Acta Pharmacol Sin* 2007; 28: 132–9.
- Zhang JH, Chung TD, Oldenburg KR. A simple statistical parameter for use in evaluation and validation of high throughput screening assays. *J Biomol Screen* 1999; 4: 67–73.
- Ferguson SS, Menard L, Barak LS, Koch WJ, Colapietro AM, Caron MG. Role of phosphorylation in agonist-promoted beta 2-adrenergic receptor sequestration. Rescue of a sequestration-defective mutant receptor by beta ARK1. *J Biol Chem* 1995; 270: 24782–9.
- Allen M, Hall D, Collins B, Moore K. A homogeneous high throughput nonradioactive method for measurement of functional activity of Gs-coupled receptors in membranes. *J Biomol Screen* 2002; 7: 35–44.
- Lefkowitz RJ, Shenoy SK. Transduction of receptor signals by beta-arrestins. *Science* 2005; 308: 512–7.
- Bisello A, Chorev M, Rosenblatt M, Monticelli L, Mierke DF, Ferrari SL. Selective ligand-induced stabilization of active and desensitized parathyroid hormone type 1 receptor conformations. *J Biol Chem* 2002; 277: 38524–30.
- Holloway AC, Qian H, Pipolo L, Ziogas J, Miura S, Karnik S, et al. Side-chain substitutions within angiotensin II reveal different requirements

- for signaling, internalization, and phosphorylation of type 1A angiotensin receptors. *Mol Pharmacol* 2002; 61: 768–77.
- 30 Groer CE, Tidgewell K, Moyer RA, Harding WW, Rothman RB, Prisinzano TE, *et al*. An opioid agonist that does not induce micro-opioid receptor – arrestin interactions or receptor internalization. *Mol Pharmacol* 2007; 71: 549–57.
- 31 Violin JD, Lefkowitz RJ. Beta-arrestin-biased ligands at seven-transmembrane receptors. *Trends Pharmacol Sci* 2007; 28: 416–22.
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Erratum

A major error was found in the title of our recent publication in *Acta Pharmacol Sin* 2010 Sep; 31 (9): 1026–30. Epub 2010 Aug 2.

The false title is “Non-peptidic **glucose-like peptide-1** receptor agonists: aftermath of a serendipitous discovery”.

The correct title is “Non-peptidic **glucagon-like peptide-1** receptor agonists: aftermath of a serendipitous discovery”.

The authors felt very sorry for this oversight and an unforgivable mistake.

The editorial office and the authors appreciated Prof George HOLZ very much for his great help.