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# The orphan G protein-coupled receptor 25 (GPR25) is activated by Apelin and Apela in non-mammalian vertebrates

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#### A R T I C L E I N F O

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#### ABSTRACT

G protein-coupled receptor 25 (GPR25) is an orphan G protein-coupled receptor in vertebrates, that has been implicated to be associated with autoimmune diseases and regulate blood pressure in humans. However, the endogenous ligand of GPR25 remains unknown in vertebrates. Here, we reported that in non-mammalian vertebrates (zebrafish, spotted gars, and pigeons), GPR25 could be activated by Apelin and Apela peptides, which are also the two endogenous ligands of vertebrate Apelin receptor (APLNR). Using the pGL3-CRE-luciferase reporter assay and confocal microscopy, we first demonstrated that like APLNR, zebrafish GPR25 expressing in HEK293 cells could be effectively activated by zebrafish Apelin and Apela peptides, leading to the inhibition of forskolin-stimulated cAMP production and receptor internalization. Like zebrafish GPR25, pigeon and spotted gar GPR25 could also be activated by Apelin and Apela, and their activation could inhibit forskolin-induced cAMP accumulation. Interestingly, unlike zebrafish (/spotted gar/pigeon) GPR25, human GPR25 could not be activated by Apelin and Apela under the same experimental conditions. RNA-seq analysis further revealed that GPR25 is expressed in a variety of tissues, including the testes and intestine of zebrafish/spotted gars/humans, implying the potential roles of GPR25 signaling in many physiological processes in vertebrates. Taken together, our data not only provides the first proof that the orphan receptor GPR25 possesses two potential ligands 'Apelin and Apela' and its activation decreases intracellular cAMP levels in non-mammalian vertebrates, but also facilitates to unravel the physiological roles of GPR25 signaling in vertebrates.

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#### 1. Introduction

Up to date, nearly 87 class A G protein-coupled receptors (GPCRs) are still classified as orphan GPCRs and their ligands have not yet been identified [1]. The G protein—coupled receptor 25 (GPR25) is an example of orphan receptors in the class A GPCR family. GPR25 was first cloned in 1997 and mapped on human chromosome 1 [2]. GPR25 is expressed in human memory T-cells and NK-cells and identified as a primary causal gene associated with autoimmune diseases revealed by *cis*-eQTL mapping based on a genome-wide associated with arterial stiffness, and presumably, it may be capable of binding an unknown ligand to regulate blood pressure (BP) [4].

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Despite the progress on understanding the potential roles of GPR25, the identity of its endogenous ligand remains unknown. Within the class A GPCR family, GPR25 shares a relatively high degree of amino acid sequence identity (29–34%) with vertebrate Apelin receptor (APLNR) [5]. It is clear that APLNR is activated by two distinct peptides, named Apelin and Apela (also known as Elabela and Toddler), which are encoded by two separate genes (*APELIN* and *APELA*), in humans [6], rats [7] and zebrafish [8,9].

Apelin is a peptide of 36 or 13 amino acids (designated as Apelin-36 and Apelin-13 respectively) and identified as the first endogenous ligand of APLNR [10]. Both Apelin and APLNR are reported to be widely expressed in the central nervous system (CNS) and peripheral tissues and involved in the regulation of many physiological/pathological processes, including the cardiovascular functions, angiogenesis, water balance and stress-induced disorders in vertebrates [11–13]. Apela is an alternative endogenous ligand of APLNR which was identified in zebrafish and mammals







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just recently. Apela is a peptide of 36 (Apela-36) or 22 (Apela-22) amino acids. Despite the low degree of amino acid sequence identity (25%) shared between Apelin and Apela peptides [14], both peptides can equipotently activate APLNR, which is functionally coupled to Gi protein and its activation decreases intracellular cAMP levels. In zebrafish, Apela is reported to be essential for cell movement during gastrulation and heart development [8,9]. In mammals, Apela is highly expressed in the kidneys of humans [6] and rats [7] and plays important roles in angiogenesis [6], fluid homeostasis [7], placental development [15] and cardiovascular development [16].

The structural similarity shared between APLNR and GPR25 led us to hypothesize that like APLNR, GPR25 could be activated by Apelin and/or Apela in vertebrates. To test this hypothesis, in this study, we cloned GPR25 from several representative vertebrate species, including zebrafish, pigeons, spotted gars and humans, and tested whether this receptor could be activated by Apelin and Apela *in vitro*. Our data, for the first time, demonstrated that the orphan receptor GPR25 can be activated by Apelin and Apela in nonmammalian vertebrates, and its activation can decrease intracellular cAMP levels. Our findings will help to unravel the physiological roles of GPR25 signaling in vertebrates.

#### 2. Materials and methods

#### 2.1. Chemicals and peptides

Chicken Apelin-36, Apela-32 (without a disulfide bond) and zebrafish Apelin-36, Apela-36 and Apela-22 (without a disulfide bond) peptides were synthesized (>95% purity) by GL Biochem (Shanghai, China) and their structures were verified by mass spectrometry. Dual-luciferase reporter assay system was purchased from Promega (Madison, Wisconsin). Primers used in this study were listed in Supplementary Table 1. All other experimental reagents were purchased from Sigma (Sigma-Aldrich, China).

#### 2.2. Animal experiments

Zebrafish, pigeons and spotted gars purchased from local suppliers were sacrificed and various tissues were collected. All these animal experiments were conducted in accordance with the Guidelines for Experimental Animals issued by the Ministry of Science and Technology of the People's Republic of China. The experimental protocol used in this study was approved by the Animal Ethics Committee of College of Life Sciences, Sichuan University.

# 2.3. Cloning of zebrafish, spotted gar, pigeon and human GPR25 and construction of expression plasmids

Genomic DNA was extracted from human embryonic kidney 293 (HEK293) cells or from heart tissues of zebrafish, pigeon and spotted gar using a genome DNA extraction kit (Tiangen Biotech, China) following the manufacturer's instructions. According to the predicted (or reported) cDNA sequences of zebrafish spotted (XM\_021479994), pigeon (XP\_005510493), gar (XM\_015342099), and human (NM\_005298) GPR25 deposited in the GenBank, gene-specific primers were designed to obtain the complete open reading frame (ORF) of GPR25 from these species. The ORF of GPR25 was amplified with the high-fidelity KOD DNA polymerase (Toyobo, Japan) using genomic DNA as the template (note: the coding regions of GPR25 in these species are intronless). Then, PCR products were cloned into pcDNA3.1 (+) expression vector (Invitrogen, Carlsbad, CA) and sequenced by Beijing Genome Institute (BGI).

Using the same approach, the expression plasmids encoding zebrafish APLNR (also named APLNRa, NM\_001075105) and human APLNR (NM\_005161) were also constructed in this study. For internalization assays, three pcDNA3.1 (+) expression plasmids encoding zebrafish GPR25-EGFP, human GPR25-EGFP and APLNR-EGFP were constructed by tagging EGFP to the C-terminus of GPR25 or APLNR.

#### 2.4. Functional characterization of GPR25 in cultured HEK293 cells

According to the method established in our previous studies [17,18], the functionality of various cloned receptors was examined in HEK293 cells by a system of co-transfected pGL3-CRE-luciferase reporter construct and receptor expression plasmids. In brief, HEK293 cells transiently expressing GPR25/APLNR were treated with Apelin/Apela peptide for 6 h in the presence of forskolin (5  $\mu$ M). Then, receptor-inhibited cAMP levels were monitored by a pGL3-CRE-Luciferase reporter assay [17,18].

#### 2.5. Internalization assay by confocal microscopy

HEK293 cells cultured on six-well plates were transfected with 1  $\mu$ g/well expression plasmid encoding zebrafish/human GPR25-EGFP (or human APLNR-EGFP). Then, cells were subcultured on poly-D-lysine coated glass coverslips in 24-well plates for 24 h. Cells were treated with serum-free DMEM medium or ligands (10  $\mu$ M Apelin or Apela) for 30 min, and then fixed with 4% paraformaldehyde for 10 min. After fixation, cells were washed 3 times in cold PBS, and a subsequent 1-min incubation with 2  $\mu$ g/ml DAPI in PBS was used to stain the nuclei. EGFP signals were examined by a Leica TCS SP5 II confocal microscope (Leica, Germany).

#### 2.6. Statistical analysis

The luciferase activities in each treatment group were expressed as percentage compared with the control group. The data were analyzed by nonlinear regression followed by one-way ANOVA using GraphPad Prism 5 (GraphPad Software, San Diego, CA). To validate the results, all these experiments were repeated at least 3 times.

#### 3. Results

#### 3.1. Cloning of GPR25 from zebrafish, pigeons and spotted gars

To identify the potential ligands for the orphan receptor GPR25 in vertebrates, we first cloned the coding region of *GPR25* from several representative vertebrate species, including zebrafish (MG732937), spotted gars (MG732939) and pigeons (MG732938) (Supplementary Figs. 1–3). The coding region of zebrafish, spotted gar and pigeon *GPR25* are predicted to encode a receptor of 377, 376 and 371 amino acids respectively, which are similar in length to human GPR25 (361 amino acids, NM\_005298). Amino acid sequence alignment revealed that zebrafish GPR25 shows high amino acid identity with that of spotted gars (52%), pigeons (38%), coelacanth (45%) and humans (39%), and the highest identity was noted in the seven transmembrane domains (TM1-7). In addition, the conserved D/ERY motif known for G protein-coupling and two cysteine residues for a disulfide bond formation are also present in GPR25 of all vertebrate species examined (Fig. 1).

Interestingly, we also found that vertebrate GPR25 shares some degree of amino acid sequence identity with zebrafish/human APLNR (34%/29%) (Fig. 1).



**Fig. 1.** Alignment of zebrafish GPR25 (zfGPR25) with that of human GPR25 (hGPR25), pigeon GPR25 (piGPR25), coelacanth GPR25 (coGPR25, XP\_005988535), spotted gar GPR25 (sgGPR25), zebrafish APLNR (zfAPLNR, NM\_001075105) and human APLNR (hAPLNR, NM\_005161). Identical residues are shaded in black, and similar residues are shaded in gray. Asterisks indicate the 'D/ERY motif' and 'NPxxY motif'. The two cysteine residues used for disulfide bond formation are marked by arrowheads. Horizontal lines mark the seven transmembrane domains (TM1-7).

# 3.2. Functional analysis of GPR25 in zebrafish, pigeons, spotted gars and humans

To test whether like APLNR, GPR25 could be activated by Apelin and Apela peptides, zebrafish, spotted gar, pigeon and human GPR25 were transiently expressed in HEK293 cells and treated by zebrafish or chicken Apelin and Apela peptides (Fig. 2A). The receptor-inhibited cAMP levels were then monitored by a pGL3-CRE-luciferase reporter system established in our laboratory [17].

As shown in Fig. 2B, zebrafish Apelin-36, Apela-36 and Apela-22 could inhibit forskolin (5  $\mu$ M)-induced luciferase activity of HEK293 cells expressing zebrafish GPR25 dose-dependently, and their EC<sub>50</sub> values are: 0.39  $\pm$  0.14  $\mu$ M, 0.32  $\pm$  0.09  $\mu$ M and 0.52  $\pm$  0.16  $\mu$ M, respectively. Likewise, zebrafish GPR25 could also be activated by chicken Apelin-36 (EC<sub>50</sub>: 0.66  $\pm$  0.16  $\mu$ M) and Apela-32 (EC<sub>50</sub>: 0.74  $\pm$  0.17  $\mu$ M) (Fig. 2C). As a positive control, zebrafish APLNR could be potently activated by zebrafish Apelin-36 (EC<sub>50</sub>: 1.14  $\pm$  0.78 nM), Apela-36 (EC<sub>50</sub>: 1.41  $\pm$  0.36 nM) and Apela-22 (EC<sub>50</sub>: 1.87  $\pm$  0.68 nM) (Fig. 2D) [8,9]. All these findings clearly demonstrated that like APLNR, GPR25 is a functional receptor coupled to Gi-cAMP signaling pathway and can be activated by Apelin and Apela in zebrafish, despite a relatively high concentration of each peptide is required for its activation.

Similarly, we also demonstrated that both chicken (and/or zebrafish) Apelin and Apela could activate pigeon or spotted gar GPR25 expressed in HEK293 cells (Fig. 3A-D), with Apelin being slightly more potent than Apela. Strikingly, human GPR25 expressed in HEK293 cells showed no response to chicken Apelin/ Apela treatment (Fig. 3E), though both peptides could potently

activate human APLNR under the same condition (EC<sub>50</sub> of Apelin-36:  $0.64 \pm 0.07$  nM; Apela-32: 0.64 nM  $\pm 0.07$ ) (Fig. 3F). These findings further support the hypothesis that like APLNR, GPR25 can be activated by Apelin and Apela in non-mammalian vertebrates.

#### 3.3. GPR25 internalization induced by Apelin and Apela

To investigate whether like APLNR, GPR25 could be induced to internalize following agonist treatment, HEK293 cells expressing zebrafish GPR25-EGFP were treated by 10  $\mu$ M of zebrafish Apelin-36 or Apela-36 for 30 min, and the receptor internalization was examined by confocal microscopy. As shown in Fig. 4A, the strong signal of zebrafish GPR25-EGFP was mainly localized at the cell membrane in the absence of peptide. However, when treated by Apelin-36/Apela-36, GPR25-EGFP signal was rapidly internalized to form large and hollow intracellular vesicles, as shown in Fig. 4B and C. As a positive control, human APLNR could also internalize after Apelin and Apela treatment and formed the large and hollow intracellular vesicles in the cytoplasm (Fig. 4H and I). These findings further indicate that like APLNR, zebrafish GPR25 could be activated by both Apelin and Apela, thus resulting in agonist-induced receptor internalization (Fig. 4).

Interestingly, unlike zebrafish GPR25-EGFP or human APLNR-EGFP, human GPR25-EGFP appears unable to internalize with peptide treatment (Fig. 4), although its signal could be observed at the cell membrane and cytoplasm (surrounding nuclei) in the absence of peptide (Fig. 4D). This finding suggests that human GPR25 may not be activated by Apelin/Apela *in vitro*, as shown in our functional assays (Fig. 3E).



**Fig. 2.** Functional analysis of zebrafish GPR25. **(A)** Amino acid sequences of zebrafish (zfApelin-36, zfApela-36 and zfApela-22) and chicken (cApelin-36, cApela-32) peptides used (note: chicken Apelin/Apela peptides were deduced from their cDNA sequences, KX017222/KX017223). **(B, C)** Effects of zebrafish Apelin-36/Apela-36/Apela-22 **(B)** or chicken Apelin-36/Apela-32 **(C)** on forskolin ( $5 \mu$ M)-stimulated luciferase activity in HEK293 cells expressing zebrafish GPR25 (zfGPR25), monitored by a pGL3-CRE-luciferase reporter system. **(D)** Effects of zebrafish Apelin-36/Apela-36/Apela-36/Apela-32 con forskolin-stimulated luciferase activity in HEK293 cells expressing zebrafish APLNR (zfAPLNR). **(E)** No effect on forskolin-stimulated luciferase activity was noted when HEK293 cells transfected with an empty pcDNA3.1 (+) vector were treated with Apelin/Apela peptides. Each data point represents mean  $\pm$  SEM of three replicates (N = 3).

#### 4. Discussion

In this study, the orphan receptor *GPR25* was cloned from several representative vertebrate species including zebrafish. Although the overall amino acid sequence identity among these species is not high (38–52%), our synteny analysis clearly indicates that the cloned zebrafish, spotted gar and pigeon *GPR25* are orthologous to human *GPR25* (Supplementary Fig. 4). In addition, we noted that the seven transmembrane domains of vertebrate GPR25 share higher sequence identities with those of human GPR25, ranging from 48% to 53%. Like other members of class A

GPCR, the conserved D/ERY motif for G protein-coupling and NPxxY motif are also observed in these cloned GPR25. All these conserved structural features imply that GPR25 may be a functional receptor in vertebrates. Moreover, we noted that vertebrate GPR25 shows a relatively high degree of amino acid sequence identity with that of zebrafish APLNR (34%) and human APLNR (29%), particularly in their transmembrane domains. This finding implies that GPR25 may have a close evolutionary relationship with APLNR. This speculation was supported by the phylogenetic analysis shown in Supplementary Fig. 5, in which vertebrate GPR25 shows a much closer evolutionary relationship to vertebrate APLNR, than to other



**Fig. 3.** Functional analysis of GPR25 in pigeons, spotted gars and humans. (**A**, **B**) Effects of chicken Apelin/Apela (**A**) or zebrafish Apelin/Apela (**B**) on forskolin (5 μM)-stimulated luciferase activity of HEK293 cells expressing pigeon GPR25 (piGPR25). (**C**, **D**) Effects of zebrafish Apelin/Apela (**C**) or chicken Apelin/Apela (**D**) on forskolin-stimulated luciferase activity of HEK293 cells expressing spotted gar GPR25 (sgGPR25). (**E**) Effects of chicken Apelin/Apela on forskolin-stimulated luciferase activity of HEK293 cells expressing spotted gar GPR25 (sgGPR25). (**E**) Effects of chicken Apelin/Apela on forskolin-stimulated luciferase activity of HEK293 cells expressing human GPR25 (hGPR25). (**F**) Effects of chicken Apelin/Apela on forskolin-stimulated luciferase activity of HEK293 cells expressing human APLNR (hAPLNR). Each data point represents mean ± SEM of three replicates (*N* = 3).

members of class A GPCR family, such as angiotensin II receptor type 1/2 (AGTR1/2).

The structural similarity, together with the close phylogenetic relationship, between GPR25 and APLNR led us to hypothesize that GPR25 and APLNR may share functional similarity in vertebrates, and GPR25 can also be activated by the two known endogenous ligands of APLNR: Apelin and Apela. In agreement with this hypothesis, we proved that like zebrafish/human APLNR, zebrafish GPR25 could be activated by both Apelin and Apela, and receptor activation could inhibit forskolin-stimulated cAMP production, monitored by a pGL3-CRE-luciferase reporter system [17,18]. Our finding, for the first time, indicates that zebrafish GPR25 possesses two potential ligands 'Apelin and Apela' and is functionally coupled to Gi protein(s), whose activation inhibits adenylate cyclase (AC) activity, similar to APLNR. Like zebrafish GPR25, pigeon and spotted gar GPR25 could also be activated by Apelin and Apela (Fig. 3A-D). Strikingly, we found that chicken Apelin-36 and Apela-32 seem unable to activate human GPR25 (Fig. 3E), though both peptides could activate human APLNR potently (Fig. 3F). Since confocal microscopy indicates human GPR25 could localize to the cell membrane (Fig. 4D), the inability of Apelin/Apela in activating human GPR25 is likely due to their extremely low (or no) binding affinity to GPR25 in vitro. This is partially congruent with the finding in a recent study, in which the recombinant alkaline phosphataseconjugated Apela cannot bind to rat GPR25 in vitro [7]. Taken together, our data suggest that like APLNR, GPR25 could be activated by Apelin and Apela in vertebrates, or, at the very least, in non-mammalian vertebrates.

Ligand-induced receptor endocytosis is a common feature of many GPCRs in response to ligand stimulation [19]. In this study, we

found that zebrafish GPR25 is localized at the cell surface. Nevertheless, human GPR25 and APLNR could also show some degree of intracellular retention (Fig. 4), as suggested in previous studies reporting intracellular retention of human APLNR expressed in HEK293 cells [6,20]. Like human APLNR (Fig. 4) [6,20], zebrafish GPR25 is rapidly internalized and forms the large, hollow vesicles upon Apelin and Apela stimulation (Fig. 4), whereas human GPR25 seems unable to undergo internalization. This finding further supports the notion that both Apelin and Apela could activate GPR25 in non-mammalian vertebrates.

The activation of GPR25 by Apelin and Apela in non-mammalian vertebrates, and not in mammals, suggests that mammalian GPR25 may have undergone a dramatic structural change after the divergence of birds and mammals, thus leading to its functional difference between non-mammalian and mammalian species (Figs. 2–4). Similar to the findings in the present study, another orphan receptor Bombesin receptor subtype-3 (BRS3) was demonstrated to be potently activated by gastrin-releasing peptide (GRP) and neuromedin B (NMB) in non-mammalian vertebrates (chickens and spotted gars), but not in mammals (mice) in our recent study [21]. These studies from our laboratory may provide an alternative key approach to de-orphanize other orphan GPCRs from an evolutionary perspective in future. In this study, we also noted that activation of GPR25 in zebrafish, spotted gars and pigeons requires high concentrations of Apelin or Apela ( $\geq$ 100 nM), therefore, we still cannot exclude the possibility that other potential 'high-affinity' ligand(s) for GPR25 may exist in vertebrates.

The functional similarity between GPR25 and APLNR, ie both can be activated by Apelin and Apela and are coupled to Gi-cAMP signaling pathway (Figs. 2–4), strongly suggests that GPR25



Fig. 4. Confocal microscopy showing the internalization of GPR25/APLNR. (A-C) Zebrafish Apelin-36 (B) and Apela-36 (C) could induce the internalization of zebrafish GPR25-EGFP expressed in HEK293 cells, while in the absence of ligand (A: control), GPR25-EGFP was mainly localized at cell membrane. (D-F) Chicken Apelin (E)/Apela (F) seems unable to induce the internalization of human GPR25-EGFP. (G-I) Human APLNR expressed at cell membrane (G: control) was rapidly internalized when treated by chicken Apelin-36 (H) or Apela-32 (I). Arrowheads mark the internalized GPR25/APLNR signals in the large, hollow vesicles.

signaling may play important physiological and pathological roles in vertebrates more or less similar to those of APLNR signaling, such as regulation of blood pressure [4,11]. Despite the failure of Apelin and Apela in activating human GPR25 in vitro, the activation of GPR25 by Apelin or Apela in vivo should not be ruled out, as this may be possible in the presence of some unknown accessory proteins in vivo. In addition, there has been increasing evidence showing that many GPCRs possess constitutive activity in the absence of their endogenous ligands, such as melanocortin receptor 4 (MC4R) in the absence of  $\alpha$ -melanocyte-stimulating hormone ( $\alpha$ -MSH) [22]. Therefore, there is also a possibility that GPR25 may possess constitutive activity to regulate physiological/pathological processes, such as blood pressure and arterial stiffness in humans [4]. It is reported that APLNR is capable of forming hetero-dimers with other GPCRs, such as kappa opioid receptor [23], thus, we should not exclude the possibility that GPR25 may hetero-dimerize with other GPCR(s) (e.g. APLNR) to exert its functions. Our future studies on these issues will help to define the functionality of vertebrate GPR25 more precisely.

To understand the physiological roles of GPR25 in vertebrates, RNA-Seq data was used to analyze the distribution of *GPR25* in zebrafish, spotted gars and humans (Supplementary Fig. 6). *GPR25* mRNA was detected to be widely expressed in most tissues examined, including the brain, heart, gills, kidneys, bones, intestines and testes of zebrafish and spotted gars. Although zebrafish *APLNR* was previously reported to be expressed in abundance during embryonic development [8,9], *GPR25* mRNA was hardly detected in zebrafish embryos. It implies that unlike APLNR, GPR25 signaling may not play an active role in zebrafish embryogenesis [8]. Like zebrafish and spotted gar *GPR25*, human *GPR25* mRNA is also expressed in a variety of tissues, including the testes and intestines [24]. The wide tissue distribution of *GPR25* across vertebrate species hints that GPR25 may play a wide range of roles in vertebrates, which is worthy of further investigation.

In summary, this study represents the first to report that the orphan receptor GPR25 can be activated by Apelin and Apela in non-mammalian vertebrates, and its activation can decrease intracellular cAMP levels. Our data suggests that GPR25 possesses two potential ligands 'Apelin and Apela' and is coupled to Gi-cAMP signaling pathway. These findings imply that GPR25 signaling may act in a way more or less similar to APLNR signaling in vertebrates. Although many questions regarding GPR25 have yet been answered, our study undeniably sets up a critical starting point to explore the physiological roles of GPR25 signaling in vertebrates, including zebrafish and humans.

#### **Competing financial interests**

The authors declare no competing financial interests.

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#### Appendix A. Supplementary data

Supplementary data related to this article can be found at https://doi.org/10.1016/j.bbrc.2018.04.229.

#### **Transparency document**

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## **Supplementary Materials**

## Title:

The orphan G protein-coupled receptor 25 (GPR25) is activated by Apelin and Apela in non-mammalian vertebrates

### Authors:

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1	ATG	GCA	AGC	AGC	'ACA	GAG	ATG	GCA	CAC	TCT	GGA	ATT	ACA	ATG	AGC	TTG	ACA	AGT	GAA	TAT	GAC	TAT	GAC	TAT	'CCA	ATT	AAT	TCC	ACG	GAC	90
1	М	A	S	S	Т	Е	М	A	Η	S	G	I	Т	М	S	L	Т	S	Е	Y	D	Y	D	Y	Ρ	I	Ν	S	Т	D	30
91	GAG	AAT	CCA	ATC	TAC	ACC	CTT	CCC	GAC	GCI	GAA	CTG	CTO	GCCC	ATG	TCT	'AAC	ATC	TAC	ATT	CCT	GTG	CTG	TAC	ATC	ATT	ATG	TTC	CTC.	ACC	180
31	Ε	Ν	Ρ	I	Y	Т	L	Ρ	D	A	Ε	L	L	Ρ	М	S	Ν	I	Y	I	Ρ	V	L	Y	I	I	М	F	L	Т	60
181	GGC	TCT	TTG	GGA	AAC	CTG	TTT	GTG	ATT	GTO	GTC	ATT	GGA	AAG	CGG	CGC	AAG	JAAA	AGT	GGA	CGC	CTG	GTA	GAC	ACC	TTC	GTG	CTG	AAC	CTG	270
61	G	S	L	G	Ν	L	F	V	Ι	V	V	I	G	K	R	R	K	K	S	G	R	L	V	D	Т	F	V	L	Ν	L	90
271	GCA	CTA	GCC	GAC	CTC	GTG	TTT	GTT	CTC	ACG	CTG	CCA	ATO	TGG	GCA	ATT	TCG	ACC	CGT	TAT	GAC	GAG	TGG	CCC	TTT	GGT	GAG	GTG	TTA	TGC	360
91	A	L	A	D	L	V	F	V	L	Т	L	Ρ	Μ	W	A	I	S	Т	R	Y	D	Е	W	Ρ	F	G	Ε	V	L	С	120
361	AAG	ATT	AGC	AGC	'TTT	ATC	'ATT	GCG	GTT	AAC	CGC	TTT	TCC	CAAC	ATC	TTC	TTC	CTC	ACC	TGC	ATG	AGC	GTG	GAC	CGC	TAC	стg	GCA	GTG	GTG	450
121	К	I	S	S	F	I	I	A	V	Ν	R	F	S	Ν	I	F	F	L	Т	С	М	S	V	D	R	Y	L	A	V	V	150
451	CGC	CTA	ATG	GAC	TCT	CGG	TTT	CTC	CGC	'AGC	CAGC	'AAT	TGI	GCA	CAA	ATT	ACC	TGT	GGC	ATC	GTG	TGG	GTG	GTT	TCC	TTC	TTT	СТА	GGA'	TCT	540
151	R	L	М	D	S	R	F	L	R	S	S	Ν	С	A	Q	I	Т	С	G	I	V	W	V	V	S	F	F	L	G	S	180
541	CCA	TCT	CTG	GCG	TAC	CGT	CAC	TTG	ATC	AAC	'AAT	TCG	GTO	GTGC	TCT	GAA	GAT	TCC	AAA	TCC	TCT	TTT	GTT	CAA	GGA	ATG	AAT	CTT	CTG.	ACT	630
181	Ρ	S	L	A	Y	R	Η	L	I	Ν	Ν	S	V	С	S	Е	D	S	K	S	S	F	V	Q	G	М	Ν	L	L	Т	210
631	ATA	TTG	CTA	ACC	TTC	TTG	CTC	CCG	GTA	CTC	ATT	CTA	GGC	CTC	TGT	TAT	GGG	TCG	ATT	TTG	GTC	AAC	CTG	CGA	CGT	CAC	TGT	CAC	AAT	ССТ	720
211	I	L	L	Т	F	L	L	Ρ	V	L	I	L	G	L	C	Y	G	S	I	L	V	Ν	L	R	R	Н	С	Η	Ν	Ρ	240
721	GCC	AAC	ACA	CGC	ACT	GAT	GCC	AGA	CGA	AGG	CAT	TCG	GTI	AAG	ATT	GTG	TTT	GCT	ATC	ATT	TCA	GCC	TTT	CTG	ATC	TCC	TGG	CTT	CCA'	TTC	810
241	A	Ν	Т	R	Т	D	A	R	R	R	Н	S	V	K	I	v	F	A	I	I	S	A	F	L	I	S	W	L	Ρ	F	270
811	AAC	TGC	TTC	AAG	GCC	ATC	CAT	GTT	GCG	TTA	CTG	ATC	ATC	CAAT	'GGA	GAT	CTT	TAAT	GAG	GAC	ACC	TAT	GTT	GTC	ATA	CAT	AGA	GGA	CTC.	ATG	900
271	Ν	С	F	К	A	I	Н	V	A	L	L	I	I	Ν	G	D	L	Ν	Е	D	Т	Y	V	V	I	Н	R	G	L	М	300
901	CTC	TCC	TGC	TGT	CTG	GCA	TTT	CTC	AAC	AGC	TGC	GTC	'AA1	CCG	GCT	ATC	TAC	TTC	TTT	СТС	GAT	CAA	CAT	TTC	AGA	CGC	AGA	GCT	TCC.	ATG	990
301	L	S	С	С	L	A	F	L	Ν	S	С	V	Ν	Ρ	A	I	Y	F	F	L	D	Q	Н	F	R	R	R	A	S	М	330
991	TTG	TGT	CTG	AGC	TGT	CTG	AGT	CAA	AAT	GAC	CAA	GCA	CAC	CAG	AGT	TAC	ATC	ACC	TCG	AAT	TCT	TAC	TCC	AAC	GGC	ACA	тст	GAG	ACC	TGC	1080
331	L	С	L	S	С	L	S	Q	Ν	D	Q	A	Η	Q	S	Y	I	Т	S	Ν	S	Y	S	Ν	G	Т	S	Е	Т	С	360
1081	TCT	GGA	AAT	ACA	TCA	ACC	CGC	GGG	CGG	CTI	TTC	TCA	CTI	ACT	CAA	AAG	GCA	TGA	11	34											
361	S	G	Ν	т	S	т	R	G	R	L	F	S	L	Т	0	К	А	*	37	8											

**Supplementary Fig. 1** The nucleotide and deduced amino acid sequence of zebrafish *GPR25* (accession no. MG732937).

1 ATGGCGGCGAATACCGTGGACACGACGGACTACGAGTACTCTGAAAATGACTTCAACGGAAGCGAGAACTACACATACCTGTATCCTGAC 90 1 M A A N T V D T T D Y E Y S E N D F N G S E N Y T Y L Y P D 30 31 L H E L L D C P Q K S L P W Q S V I I P T L Y F L I F F T G 60 181 TTCATTGGGAATCTGTGCGTCATCGTCATCGTCCATGGCCTCCAAGCACAAGAACAGGAGGCTGGTGGACACCTTTGTCATTAACCTGGCCGTG 270 61 F I G N L C V I V I M A S K H K N R R L V D T F V I N L A V 90 271 GCGGACTTGGTCTTCGTGTGCACTTTACCGTTCTGGGGGGATTTCGGCGGCGAAGGAAAACCAATGGGACTTCGGCGATTTCCTCTGCAAG 360 91 A D L V F V C T L P F W G I S A A K E N Q W D F G D F L C K 120 361 TTTAGCAGCTACGTCATCTCCGTGAACCGCTTCTCCAACATTTTTTTCCTCACCTGCATGAGCGTGGACAGGTACTTGGCCGTCGTGCGG 450 121 F S S Y V I S V N R F S N I F F L T C M S V D R Y L A V V R 150 451 ATGCTGGACTCCAGATACCTTCGCAGCAACCAATGCGTACGCTTGACCTGTTGCGTCATCTGGTTGGCGTCCATCGCGCCTAGGAGTGCCC 540 151 M L D S R Y L R S N O C V R L T C C V I W L A S I A L G V P 180 541 TCTCTTATCTACCGGAGGGTGGAGGAAAATGGCGCGGCCCGTTTGTGCACCGAAATCGCCGACTCAGCTGTTTTCAACGGGATCAGCCTT 630 181 S L I Y R R V E E N G A A R L C T E I A D S A V F N G I S L 210 211 V A L F L A F V L P V V I I V A C Y C S I L V K L R N P P G 240 721 CTTCAAAAACCCCAAAAACCGAACAGCGCCGCAGACACTCGCTCCAAGATTGTGCTGGTCATCATCGTGGCCTTCGTCGTCTCCTGGCTGCCG 810 241 L Q N P K T E Q R R R H S L K I V L V I I V A F V V S W L P 270 811 TTCAACGTCTTCAAAACCATCCTCATCGCCTGCCAGCTCCAGTTCGCGTCGCTGTCCTGCGGCCTCTCAAACGACCCTGGCGAGGGGGCCTG 900 271 F N V F K T I L I A C Q L Q F A S L S C A S Q T T L A R G L 300 901 ACCCTGTCCTCTTGCCTTCCTAAACAGCTGCGCTAACCCCGGCCATTTACATCTTCCTGGACCACCACCTTCAGGCATCGCGCTCTG 990 301 T L S S C F A F L N S C A N P A I Y I F L D H H F R H R A L 330 991 CGCTTCTGTCTCGGCTGGTTCGGCCCACAGATCAAGCGGCGGAGTGGAGCTCCAGCCGGGTCCTCTCCCGGGCTCACCATTGACAGTTAC 1080 331 R F C L G W F G P Q I K R R S G A P A G S S S G L T I D S Y 360 1081 TCGGTGTCCAGTTACCCTCGCTGCCGCATGTCTTCAGTTCACTTGCAGTGA 1131 361 S V S S Y P R C R M S S V H L Q \* 377

**Supplementary Fig. 2** The nucleotide and deduced amino acid sequence of spotted gar *GPR25* (accession no. MG732939).

1 ATGGCTCCCGAGGAGCTCGTCACCTCGGATGACTACGAGTACCCACCGGTGACCAGCGCGATGGAGAACCAGTGGGTGTCCTCCAGGGGG 90 1 M A P E E L V T S D D Y E Y P P V T S A M E N Q W V S S R G 30 31 E L F F T T I F I P I L Y S L I F L L G L V G N L F V I V L 181 ATGGCCAAGAAGAGGGGGCAAGAGGATGGTGGACACGTTCGTGCTGAACCTGGCGGTGGCCGATGTCGTGTTTGTCTGCACGTTGCCC 270 61 M A K K S R G K R M V D T F V L N L A V A D V V F V C T L P 90 91 F W V A A G A Q G N R W L L G E G L C K L S S Y V I A V N R 120 361 TGCTCCAGCATCCTCTTCCTCACCGCCCTCAGCGTGGAGCGCTACCTGGTCATCAGGAAGGTGCTGGACACCAAGGTGATGGGTTCCCAG 450 121 C S S I L F L T A L S V E R Y L V I R K V L D T K V M G S Q 150 451 AGACACGTGCGCGTCACCTGCGGCGTCATCTGGGCCGTGTCCCTCCTCCTGGGCGCCCCCACCCTGCTGTACCGGCGGCGGCGGGGGGGC 540 151 R H V R V T C G V I W A V S L L L G A P T L L Y R R L D G D 180 541 GACTGCTGCGGATGAAGATGGAGAGGAGTTCAGCCTGGCCATGGTCTTCCTCACCTTCCTCCTCCTGCCCTTGGGCGTCATCTCCTCTCTGCTAC 630 181 D C W D E D G E D F S L A M V F L T F L L P L G V I S F C Y 210 631 TGCTCCATCTACTGCCGGGCTCCAGCGCCACGTCCGGCCTGGGCAGGGGTGTCCGACGGGCCCCCCGCGCCATCGTCACCATCGTCACCGCCCCC720 211 C S I Y C R L Q R H V R L G R G V R R S H R A I V T I V T A 240 721 TTCCTCTGCTCCTGGCTGCCCCTGAACGCCTGCAAGGTGCTGCTCTTCTTCCTCGCCAAGGGGATGCTGGTCCTGTCTCAGGGCCAGGAG 810 241 F L C S W L P L N A C K V L L F F L A K G M L V L S Q G Q E 270 811 GTGGCCCTCAGGTGGCTGGTGGCCACCAGCACCTGCCTGGCATTCGTCAACAGCTGCGTCAACCCTCTCATCTATGCCCTGATGGACGGA 900 271 V A L R W L V A T S T C L A F V N S C V N P L I Y A L M D G 300 901 CACTGCCGGCCCCGATGTCCCCTAGGTCCCCGTGCGCCGGGGGCACAGGTCCCAGAGCAGCCGCCCCCTCCTCCACCAGCGACTCCAGCCTC 990 301 H C R P R C P L G P R A P G T G P R A A A P S S T S D S S L 330 331 L F G V W T R S R C R T R H Q P A P G G T P L S P P A S S P 360 1081 GGTGCCACCGCTGTCCCCCTGGTCCCCAGCTGTGA 1116 361 G A T A V P P G P O L \* 372

**Supplementary Fig. 3** The nucleotide and deduced amino acid sequence of pigeon *GPR25* (accession no. MG732938).



**Supplementary Fig. 4**. Synteny analysis of *GPR25* in vertebrates. The orphan receptor GPR25 is localized in a syntenic region conserved in all vertebrate species examined including humans, pigeons, zebrafish, spotter gars and coelacanths. The orthologous gene was marked in the matching colors. The number below each gene indicates the gene location (Megabyte, Mb) in corresponding chromosome (chr) or scaffold shown in brackets on the right.



**Supplementary Fig. 5** Phylogenetic analysis of GPR25 among vertebrates. Phylogenetic analysis was performed using the amino acid sequences containing GPR25, APLNR, AGTR1, AGTR2, BDKRB1 and BDKRB2 from vertebrate species, including human and other vertebrate species indicated. Phylogenetic tree was constructed using the Maximum Likelihood method based on the JTT matrix-based model in MEGA7 [1]. Numbers near branch points indicate the bootstrap values.



**Supplementary Fig. 6** Expression of *GPR25* in zebrafish, spotted gars and humans. (**A**, **B**) The tissue distribution of *GPR25* mRNA transcript was examined in zebrafish (**A**) and spotted gars (**B**) using RNA-Seq data (Accessions: SRP044781 and SRP044782) [2], including 11 different tissues: brain, gill, heart, muscle, liver, kidneys, bone, intestines, embryos, ovaries and testes. The quantification of reads was performed with Salmon v0.8.2 [3] against the Ensembl database. (**C**) The tissue distribution of human *GPR25* mRNA transcripts was examined by analyzing the RNA-Seq data obtained from the GTEx Consortium (https://www.gtexportal.org/home/gene/GPR25) [4]. The transcripts per million (TPM) values were used to estimate the abundance of *GPR25* mRNA transcript.

Gene/construct name	Sense/ antisense	Primer sequence (5'-3')	Size (bp)	
<sup>b</sup> Primers for construction	on of the expression	ion plasmids		
Zebrafish gpr25	Sense	CGG <u>GGTACC</u> GCACAGATGGCAAGCAGCAC	1187	
	Antisense	CGG <u>GGTACC</u> AGAAGCGAAAGTTGGCAGTC		
Human GPR25	Sense	CGG <u>GGTACC</u> AGGCCATGGCCCCCACAGAG	1321	
	Antisense	CGG <u>GGTACC</u> GGAACAGCTAGTCACAGGAG		
Pigeon GPR25	Sense	CCG <u>GAATTC</u> GCCCCCATGGCTCCCGAGGAG	1202	
	Antisense	CCG <u>GAATTC</u> TCAAACCGGGTTCCTGTTACC		
Spotted Gar gpr25	Sense	CCG <u>GAATTC</u> GCGACATGGCGGCGAATACC	1149	
	Antisense	CCG <u>GAATTC</u> GAGAGTAGGTCCTTCACTGC		
Zebrafish aplnra	Sense	CCG <u>GAATTC</u> CAAATGGAGCCAACGTC	1242	
	Antisense	CCG <u>GAATTC</u> ATACTCGCATCCACTCATC		
Human APLNR	Sense	CCG <u>GAATTC</u> AGCATGGAGGAAGGTGGTG	1148	
	Antisense	CCG <u>GAATTC</u> CTAGTCAACCACAAGGGTC		
Zebrafish Gpr25-EGFP	Sense	GTGTGGTGGAATTCGCCACCATGGCAAGCAGCACAGAGAT	1171	
	Antisense	TCCTCGCCCTTGCTCACCATTGCCTTTTGAGTAAGTGAGA		
Human GPR25-EGFP	Sense	TAGTCCAGTGTGGGGGGAATTCGCCACCATGGCCCCCACAG	1128	
	Antisense	CTCGCCCTTGCTCACCATCCAGGAGGCCGAGGCAGTGT		
Human APLNR-EGFP	Sense	GTGTGGTGGAATTCGCCACCATGGAGGAAGGTGGTGAT	1180	
	Antisense	TCCTCGCCCTTGCTCACCATGTCAACCACAAGGGTCTCCT		
pcDNA3.1-EGFP	Sense	ATGGTGAGCAAGGGCGAGGA	6159	
	Antisense	GGTGGCAATTCCACCACACT		

# Supplementary Table 1 Primers used in this study <sup>a</sup>

<sup>a</sup> All primers were synthesized by Tsingke (Beijing, China).

<sup>b</sup>Restriction sites added at the 5'-end of the primers are underlined.

### **References in supplementary materials**

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