



# Toward identifying specific roles for G-protein $\beta$ and $\gamma$ subunit variants in olfactory reception

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## A commentary on

### Expression profile of G-protein $\beta\gamma$ subunit gene transcripts in the mouse olfactory sensory epithelia

by Sathyanesan, A., Feijoo, A. A., Mehta, S. T., Nimarko, A. F., and Lin, W. (2013). *Front. Cell. Neurosci.* 7:84. doi: 10.3389/fncel.2013.00084

G-proteins mediate many cellular signaling processes; some are restricted to certain tissues or cell types, whereas others are involved in more general activities. For example, information regarding a change in the concentration of peptides, hormones, lipids, neurotransmitters, ions, odorants and tastants or an influx of photons to the eye can be transmitted to a cell via G-proteins.

Heterotrimeric G proteins are composed of three subunits:  $G\alpha$ ,  $G\beta$  and  $G\gamma$ . When activated, the  $G\alpha$  subunit binding GTP, and the  $G\beta\gamma$  heterodimer act on their effectors. In both vertebrates and invertebrates, many genes encode different variants of each subunit. In mammals, 20 genes encode the  $G\alpha$ , 5  $G\beta$ , and 12  $G\gamma$  subunits (Malbon, 2005; Dupre et al., 2009), and there is also a considerable amount of variability in more simple organisms, such as *Drosophila melanogaster*, with 6 genes for  $G\alpha$ , 3 for  $G\beta$  and 2 for  $G\gamma$  (Wolfgang et al., 1990; Yarfitz et al., 1991; Schulz et al., 1999; Boto et al., 2010).

Expression studies generally offer basic information on the possible biological function of gene products and have been extensively applied to G-proteins in many species. Previous reports have confirmed that gene expression is cell specific in some cases. For example, in *Drosophila*, the  $G\alpha_q$ -1 isoform (Lee et al., 1994) and

the  $G\beta 76C$  subunit (Yarfitz et al., 1991) were found to be specifically expressed in photoreceptor cells, highlighting their role in phototransduction.

The possibility of relating specific expression of certain genes (specially for the less known  $G\beta$  and  $G\gamma$  subunits) with particular functions in a comprehensive way is both a very interesting and hot issue for very different fields (see for example, O'Neill et al., 2012; El-Haibi et al., 2013). The attempt to relate particular  $G\beta$  and  $G\gamma$  variants with olfactory reception tissues in mice is in the basis of the article by Sathyanesan et al. recently published in *Frontiers in Cellular Neuroscience* (2013, 7, 84).

In many vertebrates, olfactory reception is mediated by odorant receptors that belong to the G-protein-coupled receptor (GPCR) family (Mombaerts, 1999).

The expression pattern and functional roles for  $G\alpha$  proteins in olfactory reception have been deeply studied. Golf was found to be highly and almost exclusively expressed in olfactory receptor neurons (ORNs) of the main olfactory epithelium (MOE) (Jones and Reed, 1987), and a lack of Go in the vomeronasal organ (VNO) of mice elicits behavioral deficits (Chamero et al., 2011). However, the  $G\beta\gamma$  subunits are not well-studied in mice, and only a few reports refer to the gene or protein expression of particular variants (Kulaga et al., 2004; Lin et al., 2007; Kerr et al., 2008; Li et al., 2013).

The  $G\beta\gamma$  heterodimer is a functional structure that, unlike  $G\alpha$ , does not change its conformational state when it dissociates from the heterotrimer. *In vitro* studies show a high variability of possible  $G\beta$  and  $G\gamma$  combinations, though the possibilities are more restricted in the

native situation (Milligan and Kostenis, 2006).

In their paper in *Frontiers in Cellular Neuroscience*, Sathyanesan et al. (2013) performed a comprehensive study of the expression pattern of all  $G\beta$ - and  $G\gamma$ -encoding genes (17) in mice olfactory receptor epithelia, MOE and VNO, in adult animals and also at different postnatal stages.

To this end, the researchers analyzed gene expression by RT-PCR and quantitative PCR using RNA extracted from both organs and designing specific primers for each  $G\beta$  and  $G\gamma$  subunit based on the 3' UTR region in an attempt to overcome possible homology. The authors reported strong expression of the  $\beta 1$ ,  $\gamma 8$ , and  $\gamma 13$  genes in MOE, confirming previous results from other studies (Lin et al., 2007; Kerr et al., 2008), and also detected for the first time the expression of  $G\beta 2,4$  and 5 and  $G\gamma 2,3,5,10,11$ , and 12 in this tissue. A quantitative analysis confirmed that  $\beta 1$ ,  $\gamma 8$  and  $\gamma 13$  are the most abundant transcripts in the main olfactory epithelium of the mouse. Sathyanesan et al. similarly analyzed the expression of the  $G\beta$  and  $G\gamma$  subunits in VNO, and their results showed the expression of only  $G\beta 1$  among the  $G\beta$  group (and perhaps a very weak signal for  $G\beta 2$ ), and the strong presence of  $G\gamma 2,3,8$ , and 13.

These data are based on the total RNA present in the organs. Thus, for further detail on the presence of distinct G proteins in different cell types in these olfactory organs, the authors performed *in situ* RNA hybridization experiments (RISH) to localize the  $G\beta$  and  $G\gamma$  transcripts to specific cells.

Although the RISH results did not consistently agree with the data from

PCR experiments, as explained in the manuscript, such a situation can be due to the different sensitivities of the techniques or to the inherent technical difficulties of each. Nevertheless, the RISH data are reliable, considering that they show the most restrictive results. G $\beta 1$  appears to be the only variant expressed in MOE and VNO olfactory receptor neurons. With regard to G $\gamma$  subunits, some expression specificity was detected, as G $\gamma 2$  and G $\gamma 12$  were only localized to supporting cells. The authors performed double-labeled experiments to show that, in a considerable proportion of neurons in MOE, Golf and G $\gamma 13$  are expressed in the same cells, as are G $\beta 1$  and G $\gamma 13$ . Therefore, Sathyanesan et al. propose that these three subunits may be part of the same heterotrimer.

The authors also show convincing results regarding specific expression, depending on the cell type, of the G $\gamma$  subunits in VNO. VNO sensory neurons differentially express two types of G $\alpha$  proteins: Gi2 in the apical layer and Go in the basal layer (Jia and Halpern, 1996). Sathyanesan et al. found 4 types of G $\gamma$  in VNO: G $\gamma 2,3,8$ , and 13. Although all are expressed in the Gi2 layer, only was found to G $\gamma 8$  localize to the basal Go layer of neurons.

Further experiments testing the cellular location and protein interactions will be necessary to confirm these data, but the finding of gene expression specificity for some G $\beta$  and G $\gamma$  subunits is an important step toward unraveling olfactory transduction in mammals and the role of G-proteins in the development of olfactory reception tissues.

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