

Unraveling the complex G protein interaction signatures of Free Fatty Acid Receptors using BRET ER/K biosensors

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Introduction: It is clear that many GPCRs interact with multiple different G α proteins. Intermolecular BRET biosensors have therefore been used to assess the full range of G proteins activated by a given receptor. A key limitation is that expression levels of receptors and signaling proteins involved will impact sensor readouts. We have generated a range of expression level independent BRET sensors based on described FRET sensors (1) and used these to assess G protein coupling to free fatty acid (FFA) family of GPCRs (2).

Methods: Sensors were generating by cloning the GPCR fused at its C terminus to mNeonGreen, a 10nm ER/K linker (1), Nanoluciferase (NLUC), and peptide corresponding to the final 28 a.a. specific G α 's. Sensors for 10 different G α 's, and one control (lacking any G α) were generated for each GPCR. Sensors were expressed in HEK293 cells and BRET monitored after NLUC substrate addition, before and for 2 min after ligand addition. Area under the ligand response curve (AUC) was calculated to determine responses for each GPCR-G α combination.

Results: The biosensor approach was validated using GPCRs classically associated with Gs (β_2 -AR), Gi/o (D₂), Gq/11 (FFA1), and G12/13 (GPR35) (each n \geq 3). In each case, sensors reported significant increases (p<0.05) in AUC for members of the classically associated G α family. Extending these studies to two short chain fatty acid receptors, FFA2 (n=3) and FFA3 (n=3), demonstrated that activation by the endogenous ligand, propionate (C3), resulted in significant increases (p<0.05) in AUC for all G α 's other than Gs for FFA2 for each member of the Gi/o family as well as for G12 for FFA3. An FFA2 allosteric agonist, AZ1729 (3), produced significant (p<0.05) AUC responses only for Gi1/2, Gi3 and G12 sensors. Co-addition of C3 and AZ1729, resulted in a synergistic increase to Gi1/2, Gi3, and G12 AUC compared with C3 (10.3-fold, 7.2-fold and 10.30-fold) or AZ1729 (7.3-fold, 9.6-fold, 5.7-fold) alone. Activation of the long chain fatty acid receptor, FFA4, by endogenous ligand, α -linolenic acid, or by various synthetic agonists (each n \geq 3) (2), resulted in significant AUC increases (p<0.05) for all G α 's except G12 and G13.

Conclusion: These studies demonstrate the complex GPCR-G protein coupling profiles of the FFA family of GPCRs using novel BRET ER/K biosensors.

References

1. Malik RU et al. *J Biol Chem* (2013) 288: 17167-17178.
2. Milligan G. et al. *Chem Rev* (2017) **117**: 67-110.
3. Bolognini D. et al. *J Biol Chem* (2016) **291**: 18915-18931.