

1 **Molecular and functional insights into Gonadotrophin Hormone Receptor**
2 **Dimerisation and Oligomerisation**

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4 Uche C Agwuegbo¹ and Kim C Jonas^{1,2,*}

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6 1. Vascular Biology Research Centre, St George's University of London, Cranmer
7 Terrace, London, SW17 0RE, UK.

8 2. Institute of Medical and Biomedical Education, St George's University of London,
9 Cranmer Terrace, London, SW17 0RE, UK.

10 * Corresponding author, kjonas@sgul.ac.uk; +44 2087255953

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12 **Key words:** Gonadotrophin hormone receptors, Gonadotrophins, G protein-coupled
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23 **Abstract**

24 The gonadotrophin hormones, follicle stimulating hormone and luteinising hormone, are
25 essential for reproduction. They work in concert to control multiple aspects of gonadal function
26 to ultimately produce meiotically competent and fertilisable gametes, provide the optimal
27 endometrial environment and support for implantation and maintain pregnancy via
28 progesterone production throughout the first trimester of pregnancy. These complex and
29 multidimensional functions are mediated via the gonadotrophin hormone receptors, luteinising
30 hormone receptor (LHR) and follicle stimulating hormone receptor (FSHR), Class A G protein-
31 coupled receptors (GPCR), which couple to multiple G protein-dependent and independent
32 signal pathways to control these physiological processes. Over the last two decades, a
33 plethora of experimental evidence has shown that GPCRs can associate to form dimers and
34 oligomers. This association provides a means of mediating the diverse functional
35 requirements of a single receptor subtype and for the gonadotrophin hormone receptors, has
36 been shown to alter the pharmacology and signal activation profile of these receptors. This
37 review will detail the historical and current evidence detailing the formation of gonadotrophin
38 hormone receptor homomers and heteromers. We will discuss the functional insight gained
39 from *in vitro* and *in vivo* studies, and the potential impact in modulating reproductive health
40 and disease.

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49 **1. Introduction**

50 The coordinated actions of receptor-mediated processes ensure the correct functioning of all
51 physiological and endocrine processes. This is particularly pertinent for the gonadotrophin
52 hormone receptors (GpHRs), follicle-stimulating hormone receptor (FSHR) and luteinising
53 hormone receptor (LHR), whose actions are essential for reproduction and fertility(1-3).
54 Localised to specialised cellular compartments of the testes and ovaries, the GpHRs control
55 and regulate gonadal functions; in males, regulating testosterone production and the
56 proliferation and maturation of sperm(1), and in females, regulating gonadal steroidogenesis,
57 follicular growth and maturation, follicle recruitment and dominant follicle selection, ovulation,
58 and corpus luteum function(2,3). Due to the importance of GpHRs in initiating and maintaining
59 fertility, they are key targets of assisted reproductive technologies, particularly *in vitro*
60 fertilisation (IVF). Additionally, in recent years, the extragonadal expression of GpHRs has
61 been reported, with proposed roles in prostate cancer(4), placental function(5,6), osteoclast
62 activity(7,8), thermogenesis(9) and the development of Alzheimer's disease(10). Thus,
63 understanding the mechanisms underpinning how GpHRs function is imperative for the
64 generation of more efficacious, targeted, effective and potentially personalised
65 pharmacological-based therapeutic strategies for improvements in reproductive health, and
66 also non-reproductive health and disease. This review will discuss a concept that has emerged
67 over the last two decades as an important modality for regulating GpHR function, namely, the
68 formation of GpHRs dimers and oligomers. We will first briefly appraise the functional roles of
69 di/oligomerisation for the wider G protein-coupled receptors superfamily, before delving into
70 the evidence presented for GpHR di/oligomerisation, and the impact on GpHR function. We will
71 finish with discussing pertinent and outstanding questions in our GpHR di/oligomerisation and
72 future perspectives for this important area of research.

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75 **2. Why do GpHR di/oligomerisation?**

76 For many years, the accepted central dogma for how GPCRs functioned, was as a single
77 receptor unit, binding hormone and mediating activation of a linear, single G protein-
78 dependent signal pathway. However, an explosion of literature over the last 20-30 years has
79 questioned this idea and presented evidence that this view of GPCR signalling was perhaps,
80 unsurprisingly, too simplistic. Indeed, most GPCRs, including the GpHRs, can couple to more
81 than 1 G protein-dependent signal pathway, may have more than 1 endogenous ligand, and
82 can activate non-G protein-dependent pathway activation via recruitment of the molecular
83 scaffolding protein, β -arrestin (recently reviewed in (11-13)). In addition, more recent studies
84 have added additional layers to this complexity with internalised GPCRs, including the
85 luteinising hormone receptor(14), and closely related the thyroid stimulating hormone
86 receptor(15,16), able to sustain cAMP-dependent signalling from the endosome(17,18). The
87 complex requirements of a single GPCR subtype show the diverse modalities that GPCRs
88 need to exploit to mediate their physiological effects. A concept that has emerged as
89 increasingly important mode of regulating GPCRs functionality, is receptor dimerisation and
90 oligomerisation. The association of GPCRs, including the GpHRs, with self (homomerisation)
91 or with other GPCRs (heteromerisation) has been shown to occur, providing a platform to
92 regulate different aspects of a GPCRs lifecycle(19), and afford the ability to diversify functional
93 responses, regulate the magnitude of signal response, the specificity of signal produced and
94 ligand directed biased signalling(11,12,20).

95

96 Although studies have shown that GPCRs, including FSHR and LHR can di/oligomerise, most
97 information has been derived from heterologous cell systems over-expressing receptors,
98 therefore, the significance in health and disease remains largely unclear. Innovative
99 approaches to study the GpHR di/oligomerisation have pushed forward our understanding of

100 the physiological relevance of Class A GPCR di/oligomerisation, showing its role and
101 relevance to regulating GpHR function, *in vivo*.

102 **3. Biochemical and biophysical evidence of gonadotrophin receptor dimerisation**

103 The first suggestion or inference that GpHR could associate, originated from the electron
104 microscopy studies of the 1970's and 1980's. Using ovarian theca and granulosa cells and
105 binding of labelled hCG to LHR, showed that LHR existed as 'clusters'(21,22). The proposed
106 clustering of LHR on hCG binding was suggested to stagnate the receptors, thereby
107 minimising the lateral diffusion of receptors, to negatively regulate G protein-coupling(22). In
108 more recent years, this concept has been further explored using time-resolved
109 phosphorescence anisotropy, a biophysical technique that can track the movement of LHR
110 within the plasma membrane. Utilising endogenous LHR expression in the Leydig tumour cell
111 line, MA-10 cells, and ovine luteal cells, binding of hCG to LHR, was shown to rotationally
112 immobilise LHR, the result of increased LHR-LHR interactions induced by ligand binding. This
113 suggested the presence of a ligand-induced increase in LHR clustering into specialised micro-
114 domains, and potentially association to form homomers(23,24). This 'clustering' could of
115 course be attributed to the initial steps in receptor desensitisation processes, and clustering
116 into clathrin coated pits for receptor internalisation, in line with classical Class A GPCR
117 endocytosis on ligand binding(17,18). More recent studies of LHR suggest that internalisation
118 kinetics and routing fate of LHR (and FSHR) are different to more typical Class A GPCRs,
119 such as β 2 adrenergic receptor, (25,26), with LHR (and FSHR) trafficking to a newly identified
120 and smaller endosomal compartment- the very early endosome. However, how
121 di/oligomerisation of LHR (and FSHR) links with internalisation to endosomal compartments
122 remains to be determined. Interestingly, the rotational diffusion rate of LHR was different when
123 bound to LH and hCG, suggesting potential differential regulation of LHR clustering and
124 potentially homomerisation by these two hormones(23).

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126 Co-immunoprecipitation of differentially epitope-tagged LHR and FSHR provided some of the
127 first biochemical evidence supporting the formation of LHR and FSHR dimers and oligomers.
128 Initially Western blotting of single tagged LHR showed varying molecular weight LHR species,
129 suggesting the presence of LHR monomers, dimers and oligomers in whole cell lysates(27).
130 To confirm that the larger dimeric and oligomeric bands were formed from interacting LHR
131 protomers, co-immunoprecipitation of two differentially tagged-LHR was carried out, showing
132 the presence of higher molecular forms of LHR and supporting the hypothesis that LHR does
133 form dimeric and higher order oligomeric LHR. Interestingly, hCG dose dependently increased
134 the number of LHR dimers and oligomers observed, but only when the LHR was stably
135 expressed, and therefore efficiently trafficked to the membrane in its mature post-
136 translationally modified form(27). As co-immunoprecipitations analyse whole cell fractions, it
137 is likely that this reflects differences in plasma membrane versus intracellular endoplasmic
138 reticulum localised LHR, and the differences in cellular localisation of LHR that stable versus
139 transient transfection of LHR results in(27). Latter studies utilising the proximity-based
140 resonance energy transfer technique of bioluminescence energy transfer (BRET), confirmed
141 the association of LHR into dimers/oligomers, and demonstrated the specificity of this
142 interaction. Interestingly, ligand treatment was shown to have no effect on LHR association
143 via this method(28).

144

145 Direct biochemical evidence for the formation of FSHR homomers was also achieved using a
146 combination of co-immunoprecipitation, FRET and BRET techniques. Similarly, to findings
147 with the LHR, FSHR was shown to reside as monomers, dimers and oligomeric
148 complexes(29,30), with FSH treatment having little effect on the relative ratios of FSHR
149 monomers, dimers and oligomers, formed, in concordance with later BRET studies and single
150 molecule imaging studies of LHR dimers and oligomers(31).

151

152 The subcellular location and timing of GpHR homomer formation has been identified. Using a
153 combination of co-immunoprecipitation with subcellular fractionation and BRET techniques
154 LHR and FSHR dimers and oligomers were shown to be localised to the plasma membrane
155 and endoplasmic reticulum(29). Moreover, utilisation of a misfolded mutant LHR that was
156 shown to be retained in the ER, revealed association of wild type LHR with the misfolded ER
157 retained mutant LHR, showing that LHR dimers and oligomers were formed during the post-
158 translational processing and modification of LHR within the ER. Analogous studies to
159 investigate this question with FSHR, showed FSHR dimers and oligomers were also formed
160 in the ER following translation.

161

162 **4. GpHR di/oligomerisation and functional asymmetry**

163 A debated and remaining question in GPCR di/oligomer field is how ligand binding within a
164 dimer or oligomer impacts on subsequent ligand binding to additional receptor protomers
165 within a complex. An elegant study by Urizar et al(32), provided the link between glycoprotein
166 hormone receptor di/oligomerisation, and negative cooperativity, via BRET and competition
167 binding assays. To explore the effect of ligand binding within a dimer on subsequent ligand
168 bindings to additional receptor protomers within a complex, a chimeric receptor of the LHR
169 extracellular domain fused to the transmembrane region of the TSHR was generated. When
170 the full length, wild type TSHR was expressed alone and binding assays conducted, as
171 expected, hCG failed to compete with TSH for binding to TSHR. However, when TSHR was
172 co-expressed with the chimera LT receptor, hCG competed with TSH for binding, and
173 desorption studies using radiolabelled TSH with 'cold' hCG suggested the presence of two
174 binding sites per dimer, as opposed to a single binding pocket formed by the dimeric
175 interaction of the two receptor protomers(32). This evidence suggests that GpHR dimers, and
176 potentially oligomers are linked by strong negative cooperativity, with binding of ligand to one
177 receptor protomer within the dimer, decreasing the binding affinity of ligand to the unbound
178 receptor protomer. This important series of experiments demonstrate that GpHR display

179 strong functional asymmetry via allosteric communication between protomers within a dimer
180 and potentially oligomer(32).

181

182 In subsequent follow-up studies the mode of communication between receptors within a dimer
183 was explored. The transmembrane membrane was shown to mediate the allosteric
184 communication between GpHR protomers in dimers(33), in line with previous reports and now
185 accepted mode of allosteric communication between Class C GPCR dimers(12). Interestingly,
186 homomers comprised of constitutively active receptors failed to display any evidence of
187 negative cooperativity with symmetry in ligand binding observed(33). A question that remains
188 is the link between negative cooperativity and preference of signal pathway activated and/or
189 magnitude of signal response observed. For the related TSHR, it has been proposed that
190 single receptor occupancy within a dimer is sufficient for Gs activation, but occupation of both
191 receptor protomers is required for Gq activation (and can also mediate Gs activation(34). How
192 negative cooperativity regulates ligand binding and subsequent G protein-signalling within an
193 oligomer remains to be determined.

194

195 **5. Functional complementation of GpHR - roles *in vitro* and *in vivo***

196 To explore the functional significance of GpHR dimers and oligomers, the phenomenon of
197 functional complementation (also known as transactivation or intermolecular cooperation) has
198 been exploited. This experimental approach utilises the relatively compartmentalised nature
199 of GPCR ligand binding and signal activation and has used to explore the impact of
200 di/oligomerisation on the functions of number of GPCRs, including the GpHRs (reviewed by
201 (35)). The structural properties of GpHRs make them a particularly good experimental model
202 for utilising functional complementation to study di/oligomerisation, as ligand binding is largely
203 mediated by the large extracellular N terminus domain of the receptors, and G protein coupling
204 for signal propagation, by transmembrane domains 5-7, as evidenced by the naturally and

205 experimentally induced activating and inactivating mutations of the GpHRs. Using these
206 discrete functional properties, mutant GpHRs have been generated that are either 'binding
207 defective'- that cannot not bind to ligand but are theoretically still able to couple of G proteins,
208 or 'signal defective' that can bind ligand but cannot couple to G protein(s) to generate
209 intracellular signals, that if expressed by themselves are functionally inactive. Remarkably,
210 when co-expressed, these mutant receptors undergo functional complementation, or trans-
211 activation, with the 'signal defective' receptor binding ligand, and the 'binding defective'
212 receptor coupling to G protein. The functioning of these receptors can only occur via
213 interaction of at least 1 signal- and 1 binding- defective receptor protomers within a complex,
214 and thus restoration of functional response is via the formation of dimers and oligomers(35).
215 This experimental paradigm has therefore proved a valuable and highly utilised model for
216 studying the functional consequences of GpHR di/oligomerisation.

217

218 *In vitro aspects*

219 The ability of GpHRs to undergo functional complementation was first reported by Osuga et
220 al(36). Using a mutant LHR that had previously been identified from a patient with Leydig cell
221 hypoplasia that was shown to possess a premature stop codon at transmembrane domain 5,
222 producing a truncated LHR, they showed that expression of this TM6/7 deleted LHR could
223 bind hCG, but couldn't activate cAMP production. To explore aspects of functional
224 complementation, a chimeric FSHR/LHR, termed FLR, was generated, comprised of the
225 FSHR extracellular domain and TM region of LHR, which generated cAMP in response to FSH
226 (and not hCG)(36). Remarkably, when the FLR was co-expressed with the truncated mutant
227 LHR and treated with hCG, cAMP production was observed, showing that the mutant LHR,
228 that could bind hCG, but not activate cAMP production, had trans-activated the FLR. This
229 showed that GpHRs could undergo functional complementation, via intermolecular
230 communication and cross talk between these two receptor species(36), and most likely via the
231 formation of dimers/oligomers.

232

233 To explore the minimum TM domain requirements for transactivation to occur and how
234 receptor protomers engaged in transactivation communicate, further deletions of the LHR
235 were undertaken. It was found that co-expression of the LHR extracellular region fused to
236 transmembrane domain 1 (TM1), with FLR was sufficient to mediate hCG-dependent
237 cAMP(36). Interestingly, fusion of the LHR extracellular domain to the single membrane
238 spanning CD8 peptide failed to mediate hCG dependent cAMP production, suggesting the
239 requirement of transmembrane mediated inter-protomer communication for transactivation to
240 occur(36). In contrast to this, other studies have shown that the extracellular domains of the
241 GpHRs fused to CD8 or GPI to anchor to the extracellular region of the receptor to the plasma
242 membrane were sufficient to mediate transactivation(37,38), and suggest a 'kiss and run' type
243 interaction of the extracellular domain of one protomer directly interacting and activating
244 neighbouring receptors. The latter view contrasts with our knowledge on how other GPCRs
245 undergoing transactivation communicate and with more recent evidence in the GpHR field,
246 whereby activation of neighbouring receptor protomers occurs via intermolecular
247 communication via the TM bundles(12,28,30,33,39). Our own studies that have utilised a
248 combination of single molecule imaging and molecular modelling would also suggest that
249 dimerisation, and indeed oligomerisation of LHR undergoing functional complementation is via
250 intermolecular communication between the TM bundles(31), showing the importance of inter
251 protomer TM bundle cross talk in directing the functions of GpHR dimers and oligomers.

252

253 Studies have shown that not all binding and signal defective mutant LHR and FSHR can
254 undergo functional complementation(38,40), suggesting a specificity in pairing, and structural
255 requirements for inter-protomer communication and ultimately transactivation to occur. The
256 leucine rich repeats (LRR) of the extracellular region of LHR are essential for mediating ligand
257 binding and appear to be a key factor in the facilitation of transactivation. It was found that
258 binding defective mutant LHR with mutations that are localised to the LRR regions 1-3 were

259 able to undergo functional complementation when co-expressed with signal defective LHR.
260 However, if binding defective mutations were located to LRR regions 4-8, they were unable to
261 undergo functional complementation(38,40). This most likely reflects the close proximity of
262 LRR 4-8 to the hinge region of the LHR, which is crucial for transducing ligand binding to the
263 TM region for signal activation. Additionally, as the hinge region also contains a suppressor of
264 TM activation to constrain the unbound receptor in an inactive conformation, which on ligand
265 binding is relaxed, mutations in LRR4-8, may interfere with the conformational changes that
266 occur to allow TM activation, and thus transactivation fails to occur.

267

268 Signal specificity of both LHR and FSHR has also been demonstrated using functional
269 complementation. Co-expression of CD8 or GPI-anchored FSHR with differential signal
270 defective mutant FSHR were shown to preferentially activate both cAMP and IP3, only cAMP,
271 or only IP3(41), suggesting differential activational states and receptor conformations that
272 mediate the specificity of G protein-coupling and subsequent intracellular pathway activation
273 within dimers and oligomers. Our *in vivo* and *in vitro* studies studies utilising LHR
274 transactivation mutations also support this idea(31), further details of which will be further
275 discussed in latter sections.

276

277 We would be remiss in admitting more atypical examples of GpHR transactivation within
278 heteromers that has been observed. LHR has been shown to transactivate the epidermal
279 growth factor receptor (EGFR), leading to the downregulation of ovarian aromatase
280 expression during the LH surge during the ovarian cycle(42). In two different mouse models
281 with inactivating mutations of the EGFR, it was shown that the down-regulation of the Cyp19a1
282 via hCG (used to mimic the LH surge), was markedly impaired but not completely abolished,
283 as it requires efficient signalling via transactivation of the EGFR. The *in vivo* of LHR-mediated
284 transactivation of the EGFR in ovulation has also been determined(43-45).

285

286 *In vivo aspects*

287 Most studies investigating the functional consequences of GPCR di/oligomerisation to date
288 have primarily utilised *in vitro* approaches, which rely upon the over-expression of GPCRs in
289 heterologous cell lines. As a result, the physiological consequence of GPCR di/oligomerisation
290 remains poorly understood and our understanding is lacking in the physiological necessity and
291 functional relevance of GPCR di/oligomerisation *in vivo*. A study that made significant inroads
292 into proving the physiological relevance of Class A GPCR di/oligomerisation, utilised a
293 functional complementation approach and the LHR as a model GPCR, to determine if LHR
294 di/oligomerisation was sufficient to mediate LHR functions *in vivo*. Using the LHR knockout
295 (LuRKO) mice, and a BAC transgenic approach, targeted co-expression of ligand binding
296 defective LHR (C22A mutation, LHR^{B-}) with signal defective mutant LHR (a deletion of
297 transmembrane domains 6 and 7, LHR^{S-}) could rescue the hypogonadal phenotype of male
298 LuRKO mice(46). Testes size and serum testosterone levels were equivalent to that of wild
299 type litter mates. Importantly, the infertile phenotype of LuRKO animals was also reversed,
300 with litter frequency and sizes from mating studies in LuRKO mice co-expressing LHR^{B-} and
301 LHR^{S-} equivalent to wild type littermates(46). Serum LH was slightly raised in comparison to
302 wild type littermates, showing the increased hypothalamic-pituitary drive to the testes to initiate
303 and maintain LH-dependent testosterone production. This seminal study showed the first *in*
304 *vivo*, physiological evidence for GPCR di/oligomerisation, and importantly, that
305 di/oligomerisation was a functionally relevant mode of LHR activation/signal propagation *in*
306 *vivo*(46). Although there has been a subsequent *in vitro* study that critically debated whether
307 the observations of this *in vivo* study were due to idiosyncrasies of the BAC transgenic method
308 for introducing the LHR^{B-} and LHR^{S-} mutants into the LuRKO mouse background(47), the
309 control experiments from this study conclusively showed that the expression of single LHR^{B-}
310 or LHR^{S-} mutants failed to rescue the infertile and hypogonadal phenotype of these mice(46).
311 Thus, confirming the specificity of the BAC approach, and confidence in LHR functional

312 complementation occurring in male mice *in vivo*. Interestingly, co-expression of LHR^{B-} and
313 LHR^{S-} in female LuRKO mice, failed to rescue the infertile and hypogonadal phenotype of
314 these animals. This may reflect the low levels of LHR^{B-} and LHR^{S-} expression female mice,
315 and the inability to induce sufficient LHR expression during the ovarian cycle. It may also
316 additional reflect the inability of functional complementation to mediate the multiple signalling
317 and functional requirements of LHR in females, as previously discussed. Further insights as
318 to why the latter proposition may be the case, will be discussed in the next section.

319

320 **6. Single molecule imaging of gonadotrophin receptor dimerisheteroation**

321 The advances in technology development have resulted in an explosion of single molecule
322 imaging and single particle tracking techniques. These advances have given unprecedented
323 insight into the nature and composition of GpHR homomers and heteromers, and the link with
324 signal activation.

325

326 Early insights into the membrane organisation of LHR were provided by single particle tracking
327 studies, which presented evidence of rat LHR localising to small specialised membrane
328 microdomains, or lipid rafts, following binding of hCG(48). These microdomains were
329 approximately 3 times smaller than unbound receptor, and importantly, when disrupted using
330 the cholesterol depleting agent, methyl- β -cyclodextrin, showed a decreased hCG-dependent
331 cAMP production(48). As a total abrogation of hCG-dependent cAMP production was not
332 observed, it suggests that localisation of hCG-bound LHR to lipid rafts is not essential for Gs
333 coupling. In support of this, although constitutively active mutant LHR also localised to lipid
334 rafts, and these LHR microdomains were approximately the same size as hCG-bound LHR,
335 cAMP production was not affected by raft disruption using methyl- β -cyclodextrin(49). That
336 said, the relationship between raft location and coupling of LHR to alternate G protein-
337 dependent and independent pathways remains unknown. Additionally, whether ligand-specific

338 differences in raft localisation and/or size of these microdomains exist. This may be
339 particularly pertinent given the changing lipid environment, and LH/hCG/LHR signal
340 requirements of the ovarian follicle during folliculogenesis and corpus luteum function.

341

342 A recent study utilising a combination of precision FRET and fluorescent correlation
343 spectroscopy with photon counting histogram methods, has explored the nature and
344 composition of FSHR homomers within the plasma membrane. Using a fusion of a C tail-
345 truncated human FSHR, the C terminus tail of the LHR and fluorescent proteins compatible
346 with FRET to the proximal end of the C-tail, FSHR was shown to traffick to and reside in the
347 plasma membrane almost exclusively as homodimers(50). These findings contrast with
348 previous crystal structure analysis of the FSHR ectodomain and hinge region, which proposed
349 that FSHR exists as trimers(51). However, the crystal structures lack the vital interactions of
350 the extracellular and transmembrane domain, due to complexities of generating stable crystals
351 with intact transmembrane domains. Additionally, the functional and physiological significance
352 of different FSHR complexes remains to be determined.

353

354 Our recent studies have provided significant insight into how the composition and functional
355 role of each protomer within an LHR oligomer can regulate signal amplitude. Utilising a
356 functional complementation approach, and the same functional complementation LHR^{B-} and
357 LHR^{S-} from our previously discussed *in vivo* study, we employed the super resolution imaging
358 approach of photoactivated dye, localisation microscopy (PD-PALM), to quantitate the number
359 of LHR monomer, dimers and oligomers at the plasma membrane, and determine the nature
360 of these complexes. Using heterologous cell lines that stably expressed either the wild type
361 mouse LHR, or co-expressing LHR^{B-} and LHR^{S-} mutant receptors, we observed that
362 approximately 40% of LHR formed homomers, with remaining 60% residing as monomers(31).
363 Sub-analysis of the types of associated LHR complexes that were observed revealed that the

364 predominant associated form of wild type LHR was dimers, with a small number of lower order
365 and higher order wild type LHR complexes. In contrast, in cells co-expressing LHR^{B-} and
366 LHR^S, approximately half the number of dimers were observed, with a concomitant increase
367 in lower order trimers and tetramers(31). Interestingly, treatment with LH or hCG had no effect
368 on the number of associated wild type or functional complementation LHR, nor the types of
369 oligomeric complexes formed, in line with previous findings from BRET analysis(28).
370 Assessment of LH and hCG-dependent Gs and Gq-association in wild type LHR and functional
371 complementation models, revealed that LH-dependent Gq activation was impaired in the
372 functional complementation model, suggesting that for full LH-dependent Gq activation, an
373 element of cis or unidirectional activation of LHR is required(31). This may shed light as to
374 why the female functional complementation mutant mice were infertile, as previously studies
375 have shown LH-dependent Gq activation is required for ovulation(52). Generation of cell lines
376 with varying cell surface ratios of LHR^{B-}:LHR^{S-} revealed that cells with an excess of LHR^{S-}
377 :LHR^{B-} resulted in amplification of Gs and Gq-dependent signals. Interestingly, the difference
378 in Gs and Gq signal observed, corresponded with an enrichment of LHR^{B-} receptor protomers
379 in both trimers and tetramers, suggesting, that modulation of specific the composition and
380 functional role adopted by a protomer engaged in an oligomeric complex, can fine tune the
381 amplitude of signal response generated.

382

383 **7. Gonadotrophin receptor heteromers**

384 The formation of LHR/FSHR heteromers has long been a debated topic. With specific
385 relevance to female reproductive physiology, LHR and FSHR are co-expressed within
386 granulosa cells of the dominant pre-ovulatory follicle, suggesting a functional role for LHR-
387 FSHR crosstalk in mediating/facilitating ovulation. As such, heteromerisation of LHR/FSHR
388 was first demonstrated by BRET(53), and subsequently by fluorescent correlation
389 spectroscopy(50), using heterologous cell models, co-expressing LHR and FSHR.
390 Importantly, heteromerisation of LHR/FSHR was shown to alter the pharmacology of FSHR

391 and LHR, enhancing dissociation of bound ligands, and negatively impacting on Gs-
392 dependent signal pathway activation(53). As LHR is known to signal via both Gs and Gq-
393 dependent pathways, with the latter particularly important for mediating LH-dependent
394 ovulatory events(52), studies by our group further investigated the impact of LHR/FSHR on
395 LH-dependent Gq activation. We found a change from a transient to more sustained calcium
396 signal, that was both dependent on Gq activation and influx of extracellular calcium(54).
397 Importantly, the presence of a sustained calcium response was also confirmed in human
398 granulosa lutein cells, which endogenously co-express LHR and FSHR, which was also found
399 to be sensitive to extracellular calcium channel blockers. PD-PALM studies revealed an LH-
400 dependent increase in LHR/FSHR heteromerisation, with specific enrichment in
401 heterotetramers, suggesting that modulation of LHR/FSHR heteromers mediated the switch
402 from transient to sustained LH-dependent calcium signalling(54). Investigation of cross-talk
403 between LH/hCG and FSH has shown that co-treatment of FSH with either LH or hCG results
404 in potentiation of their respective effects on apoptosis and steroidogenesis(55), giving an
405 insight into the potential physiological roles of LHR/FSHR heteromers. However,
406 understanding the exact physiological role of LHR/FSHR heteromers *in vivo*, remains to be
407 determined.

408

409 **8. Perspectives and conclusions**

410 It is unquestionable that GpHRs can associate to form homomers and heteromers. The
411 formation of homomers and heteromers has been shown to impact on aspects of GpHR
412 pharmacology, including ligand binding, signal specificity and signal magnitude. Despite the
413 overwhelming experimental evidence detailing the functional relevance of GpHR
414 di/oligomerisation, limitations in probing FSHR and LHR *in vivo* coupled with the technical
415 complexities, e.g., the lack of specific FSHR and LHR antibodies, mean that our knowledge of
416 the physiological roles of these receptor complexes remains limited. However, with the relative
417 ease and decreasing cost of gene modification techniques such as CRISPR, future research

418 programs in this area will enhance our knowledge to gain insight into the physiological role(s)
419 of GpHR homomers and heteromers.

420

421 With technological advances, come questions. The intriguing finding from the single molecule
422 imaging studies of LHR that 60% of LHR at the plasma membrane appears to be
423 monomers(31), suggests a distinct functional role for monomers in mediating/regulating LHR
424 functions. Indeed, deciphering how different ‘flavours’ of GpHR monomers, homomers and
425 heteromers regulate gonadotrophin hormone functions will provide much needed insight into
426 how these distinct receptor complexes regulate and fine-tune GpHR functions *in vitro* and
427 importantly, *in vivo*. Identifying the unique ‘signatures’ of GpHR complexes at the cell surface,
428 the drivers which control the formation of these complexes, and the resulting cellular
429 responses, will provide invaluable insight into how GpHR di/oligomerisation impacts
430 reproductive health, and potential dysregulation in disease. Such advances will provide the
431 opportunity for novel and potentially personalised pharmacological treatment strategies for
432 treatment reproductive pathologies, and may lead to advances for the improvement in the
433 success rate of assisted reproductive technologies.

434

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