

Roles of G Proteins in Coupling of Receptors to Ionic Channels and Other Effector Systems

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ABSTRACT

Guanine nucleotide binding (G) proteins are heterotrimers that couple a wide range of receptors to ionic channels. The coupling may be indirect, via cytoplasmic agents, or direct, as has been shown for two K⁺ channels and two Ca²⁺ channels. One example of direct G protein gating is the atrial muscarinic K⁺ channel K⁺[ACh], an inwardly rectifying K⁺ channel with a slope conductance of 40 pS in symmetrical isotonic K⁺ solutions and a mean open lifetime of 1.4 ms at potentials between -40 and -100 mV. Another is the clonal GH₃ muscarinic or somatostatin K⁺ channel, also inwardly rectifying but with a slope conductance of 55 pS. A G protein, G_K, purified from human red blood cells (hRBC) activates K⁺[ACh] channels at subpicomolar concentrations; its α subunit is equipotent. Except for being irreversible, their effects on gating precisely mimic physiological gating produced by muscarinic agonists. The α_K effects are general and are similar in atria from adult guinea pig, neonatal rat, and chick embryo. The hydrophilic βγ from transducin has no effect while hydrophobic βγ from brain, hRBCs, or retina has effects at nanomolar concentrations which in our hands cannot be dissociated from detergent effects. An anti-α_K monoclonal antibody blocks muscarinic activation, supporting the concept that the physiological mediator is the α subunit not the βγ dimer. The techniques of molecular biology are now being used to specify G protein gating. A "bacterial" α_{i-3} expressed in *Escherichia coli* using a pT7 expression system mimics the gating produced by hRBC α_K.

I. INTRODUCTION

G proteins play a central role in coupling receptors to effector systems (Figure 1). By the latest count about 85 distinct receptors, identified by pharmacological and/or molecular biological means, are coupled to effector functions by G proteins (e.g., Table 1). The number of G proteins carrying out this task is rather small in comparison, comprising 12 known and possibly 4 to 5 as yet to be discovered distinct Gα (Table 2). However, the recognition of their complexity is increasing with the discovery of up to 4 distinct β and at least 3 γ subunits that associate with various αs (Table 3). Like receptors, which are increasing in number rapidly, effectors affected by the activated forms of G proteins are also increasing (Table 4),

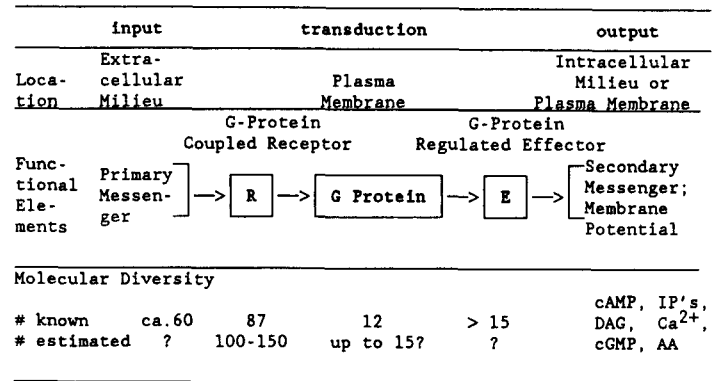


FIGURE 1. Flow of information through G protein-dependent signal transduction systems as found in vertebrates.

most notably through the discovery in 1986—1987 that ionic channels form part of the family of molecules regulated by G proteins. Using cell-free systems, such as provided by excision of membrane patches from cells and incorporation of plasma membrane vesicles into lipid bilayers, it was shown that ionic channels are indeed regulated by activated G proteins. Some of these channels had long before been predicted to be under the control of G protein-coupled receptors by means other than soluble second messengers. The mechanism by which G proteins regulate some of these channels is at the very least "mem-

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Table 1
Examples of Receptors Acting on Cells via G proteins

Type of receptor	Membrane function/system affected	Effect	Coupling protein involved	Examples of target cells(s)/organs
Neurotransmitters				
Adrenergic				
beta-1	AC	Stimulation	G _s	Heart (68), fat (62)
	Ca channel	Stimulation	G _s	Heart (4), skeletal muscle (5)
beta-2	AC	Stimulation	G _s	Liver (69), lung (70), lymphoid cells (71)
alpha-1	PhL C	Stimulation	G _{plc}	Smooth muscle (72), liver (73, 74)
	PhL A ₂	Stimulation	G _{pta}	FRTL-5 cells (75)
	PhL D	Stimulation	G _{pld}	Liver (76)
alpha-2A, -2B	AC	Inhibition	G _i	Platelet (77, 78), fat (79, 80) liver (81)
	Ca channel	Closing	G _o (G _p ?)	Symp. presynapse (24, 82)
Dopamine				
D-1	AC	Stimulation	G _s	Caudate Nucleus (83), parathyroid (84)
D-2	AC	Inhibition	G _i	Pituitary lactotrophs (85) and melanotrophs (86)
Acetylcholine				
Muscarinic M ₁ -type (M ₁ , M ₃ , M ₅)	PhL C	Stimulation	G _{plc}	Pancreatic acinar cell (87), parotid (88), CNS (88)
	K channel (M)	Closing	?	CNS, Symp. ganglia (89)
Muscarinic M ₂ -type (M ₂ , M ₄)	AC	Inhibition	G _s	Heart (68, 90)
	K channel	Opening	G _k (G _i)	Heart (2), CNS (91), lactotroph (3)
GABA _B	Ca channel	Closing	G _o (G _p ?)	Dorsal root ganglia (83, 25)
	K channel	Opening	G _k (G _o ?)	Pyramidal cells (92)
	AC	Inhibition	G _i	CNS (93)
Purinergic P1				
Adenosine A-1 or Ri	AC	Inhibition	G _i	CNS (94), fat (95)
	K channel	Opening	G _k (G _i)	heart (96)
Adenosine A-2 or Ra	AC	Stimulation	G _s	Fat (97), kidney (98), CNS (99)
Purinergic P _{2X} and P _{2Y}	PhL C (PIP ₂)	Stimulation	G _{plc}	Turkey erythrocytes (100)
	PhL C (PC)	Stimulation	G-(?)	Liver (101)
	PhL D	Stimulation	G-(?)	Liver (76)
Serotonin (5HT)				
S-1a (5HT-1a)	AC	Inhibition	G _i	Pyramidal cells, CNS (102)
	K channel	Opening	G _k (G _i ?, G _o ?)	Pyramidal cells (103)
S-1c (5HT-1c)	PhL C	Stimulation	G _{plc}	Blowfly salivary gland (88), smooth muscle (103)
S-2 (5HT-2)	AC	Stimulation	G _s	Skeletal muscle (104), CNS (105)
	Ca channel	Closing	G _o (G _p ?)	Dorsal root ganglia (82)
Histamine				
H-1	PhL C	Stimulation	G _{plc}	CNS (106), chromaffin cells (107)
H-2	AC	Stimulation	G _s	Heart (108), CNS (109), gastric mucosa (110)
Peptide Hormones				
Pituitary				
Adrenocorticotropin (ACTH)	AC	Stimulation	G _s	Adrenal cortex (111)
	Ca channel	Opening	G _k (?)	Adrenal glomerulosa (112)
	PhL A ₂	Stimulation	G-(?)	Adrenal cortex (113)
Opioid (mu, kappa, delta)	AC	inhibition	G _i	NG-108 (114, 115), luteal (116)
	Ca Channel	Closing	G _o (G _p ?)	NG-108 (29), dorsal root ganglia (117, 118)
	K channel	Closing	G _k (G _i ?)	CNS (119)
Luteinizing hormone (LH)	AC	Stimulation	G _s	Granulosa (120), luteal (121), Leydig (122)
	PhL C	Stimulation	G _p	Ovary (123)
Follicle stimulating hormone (FSH)	AC	Stimulation	G _s	Granulosa (120), Sertoli (122)
Thyrotropin (TSH)	AC	Stimulation	G _s	Thyroid (124)
Melanocyte stimulating hormone (MSH)	AC	Stimulation	G _s	Melanocytes (125)
Hypothalamic				
Corticotropin releasing hormone (CRF)	AC	Stimulation	G _s	Corticotroph (126)

Table 1 (continued)
Examples of Receptors Acting on Cells via G proteins

Type of receptor	Membrane function/system affected	Effect	Coupling protein involved	Examples of target cells(s)/organs
Growth hormone releasing hormone (GRF)	AC	Stimulation	G _s	Somatotroph (127, 128)
Gonadotropin releasing hormone (GnRH)	PhL A ₂	Stimulation	G _{pta}	Gonadotroph (129), granulosa (130), Leydig (131)
	PhL C	Stimulation	G _{plc}	Gonadotroph (132), granulosa (33), luteal (134), Leydig (131)
Thyrotropin releasing hormone (TRH)	Ca channel	Opening	G _i -type	GH ₃ (31)
	PhLC	Stimulation	G _{plc}	Lactotroph (135, 136), thyrotroph (137)
Somatostatin (SST or SRIF)	AC	Inhibition	G _i	GH ₄ C ₁ (138)
	AC	Inhibition	G _i	Lactotrophs (139), somatotrophs (140), corticotrophs (141) lymphoid cells (142), liver (143), heart (143), renal cortex (143)
Vasopressin	K channel	Opening	G (G _i ?)	Lactotrophs (3)
	Ca channel	Closing	?	Corticotrophs (28)
V-1a (vasopressor, glycogenolytic)	PhL C	Stimulation	G _{plc}	Smooth muscle, liver (73), sympath. ganglia (144)
V-1b (pituitary)	PhL D	Stimulation	G-(?)	Liver (76)
	PhL A ₂	Stimulation	G-(?)	Renal mesangial cells (145)
V-2 (antidiuretic)	AC	Inhibition	G _i	liver (146)
Oxytocin	PhL C	Stimulation	G _{plc}	Pituitary (147)
Other hormones	AC	Stimulation	G _s	Distal and collecting tubule (148)
	PhL C	Stimulation	G _{plc}	Uterus (149)
Chorionic gonadotropin	AC	Stimulation	G _s	luteal (150), Leydig (151)
Glucagon	AC	Stimulation	G _s	Liver (152), fat (62), heart (153, 154), pancreatic <i>beta</i> -cells (155)
	Ca pump	Inhibition	G _s (?)	Liver (156)
Cholecystokinin (CCK)	PhL C	Stimulation	?	Liver (157)
	AC	Stimulation	G _s	Pancreatic acini (158)
Secretin	PhL C	Stimulation	G _{plc}	Pancreatic acini (159)
	AC	Stimulation	G _s	Fat (62) pancreatic acini (160)
Vasoactive intestinal peptide (VIP)	AC	Stimulation	G _s	CNS (161), pancreatic acini (160), intestinal mucosa (162)
Parathyroid hormone (PTH)	PhL C	Stimulation	G _{plc}	Sensory ganglia (163)
	AC	Stimulation	G _s	Renal cortex (164)
Angiotensin II	PhL C	Stimulation	G _{plc}	Renal cortex (165)
	PhL C	Stimulation	G _{plc}	Glomerulosa cells (166), smooth muscle (167)
Calcitonin	AC	Inhibition	G _i	Liver (163), glomerulosa cells (168), Leydig cells (169), renal cortex (170)
	Ca channel	Stimulation	G _i -type	Y1 adrenal cells (30)
Calcitonin gene-related peptide	PhL A ₂	Stimulation	G-(?)	Renal medulla (171), fibroblasts (172)
	PhL D	Stimulation	G-(?)	Liver (76)
Calcitonin	AC	Stimulation	G _s	Renal cortex (173)
	AC	Stimulation	G _s	Skeletal muscle (174)
Calcitonin gene-related peptide	PhL C	Stimulation	G _{plc}	Skeletal muscle (175)
Other Regulatory Factors				
Chemoattractant (fMet-Leu-Phe or fMLP)	PhL C	Stimulation	G _{plc}	Neutrophils (176)
Thrombin	PhL A ₂	Stimulation	G-(?)	Neutrophils (177)
	PhL C	Stimulation	G _{plc}	Platelets (178), fibroblasts (172)
	AC	Inhibition	G _i	Fibroblasts (172)
	PhL A ₂	Stimulation	G-(?)	Platelets (179)

Table 1 (continued)
Examples of Receptors Acting on Cells via G proteins

Type of receptor	Membrane function/system affected	Effect	Coupling protein involved	Examples of target cells(s)/organs
Bombesin	PhL C	Stimulation	G _{plc}	Fibroblasts (180)
Gastrin releasing peptide	PhL C	Stimulation	G _{plc}	Fibroblasts (180)
Platelet-derived growth factor	PhL C	Stimulation	G _{plc}	Fibroblasts (181)
IgE	PhL C	Stimulation	G _{plc}	Mast cells (182)
	PhL A ₂	Stimulation	G-(?)	Mast cells (183)
Bradykinin	PhL C	Stimulation	G _{plc}	Fibroblasts (180), NG-108 (184), endothelial cells (185) renal papilla (186)
	PhL A ₂	Stimulation	G _{pla}	Fibroblasts (172), endothel. cells (187), kidney (188)
	K channel	Stimulation	G _k (G _i ?)	NG-108 (189)
	AC	Inhibition	G _i	NG-108 (184), fibroblasts (172)
Neurokinin/tachykinin NK1 (substance P)	PhL C	Stimulation	G _{plc}	CNS (190), smooth muscle (190) salivary gland (191)
	PhL A ₂	Stimulation	G-(?)	Smooth muscle (192)
Neuropeptide Y	AC	Inhibition	G _i	Heart (193)
	K channels	Stimulation	G _k	Heart (261)
	Ca channels	Inhibition	G _o	Sensory ganglia (26)
Peptide YY	?	?	?	CNS (194)
Tumor necrosis factor (TNF)	?	?	?	Monocytes (195)
Colony stimulating factor (CSF-1)	?	?	?	Monocytes (196)
Interleukin-1	PhL C (PC)	Stimulation	G(?)	T-cells (197)
Neurotensin	PhL C	Stimulation	G _{plc}	CNS (198)
	AC	Inhibition	G _i	Neuroblastoma (199)
Atrial natriuretic factor	AC	Inhibition	G _i	Aorta (200)
Epidermal growth factor	PhL D	Stimulation	G-(?)	Liver (76)
Phosphatidic acid	PhL C	Stimulation	G _{plc}	A431 carcinoma cells (201)
Platelet activating factor (PAF)	PhL C	Stimulation	G _{plc}	Platelets (202), liver (203)
	PhL A ₂	Stimulation	G-(?)	Fibroblasts (172)
	AC	Inhibition	G _i	Fibroblasts (172)
Galanin	AC	Inhibition	G _i	Pancreatic <i>beta</i> -cells (204)
Kyotorphin	PhL C	Stimulation	G _{plc} , G _o ?	CNS (205)
Prostanoids				
Prostaglandin E ₁ , E ₂	AC	Inhibition	G _i	Fat (206), kidney (207)
Prostaglandin F _{2 alpha}	PhL C	Stimulation	G _{plc}	Luteal cells (208)
Prostacyclin (PGI ₂) PGE ₁ , PGE ₂	AC	Stimulation	G _s	Luteal cells (209, 210), kidney (211)
Leukotriene D ₄ , C ₄	PhL A ₂	Stimulation	G _{pla}	Endothelial cells (212, 213)
	PhL C	Stimulation	G _{plc}	RBL-1 cells (214)
Sensory				
Light (Rhodopsins)	cGMP-PDE	Stimulation	Tr(G _{tc})	Retinal rod cells (night) (215)
	cGMP-PDE	Stimulation	Tc(G _{tc})	Retinal cone cells (color) (216a)
Olfactory signals	AC	Stimulation	G _{olf}	Olfactory cilia (216a)
	PhL C	Stimulation	G _p ?	Olfactory cilia (217)

Note: AC: adenylyl cyclase; PhL C: unless denoted otherwise, phospholipase C with specificity for phosphatidylinositol bisphosphate; PhL A₂: phospholipase A₂ (substrate specificity unknown); PIP₂: phosphatidylinositol bisphosphate; PC: phosphatidylcholine.

brane delimited" and independent of any phosphorylation event or of changes in cytoplasmic levels of second messengers such as cAMP, Ca²⁺, or IP₃ and is very likely due to direct interaction of the G protein α subunit and the channel proper (for

review of initial findings, see Reference 1). Our group was prominent in providing some of the initial as well as subsequent supporting data for these conclusions.²⁻¹² The issue whether $\beta\gamma$ dimers stimulate K⁺ channel activity in inside-out mem-

Table 2
Diversity of Mammalian G Protein α Subunits

Name	Purified ^a	Cloned ^a	Function(s) ^a Identified
α_s (1 gene, 4 splice variants)	Yes	Yes	Yes: 3 ^b
3 α_s s (3 genes: α_{s1} , α_{s2} , α_{s3}) ^c	Yes	Yes	Yes
α_{o1} (1 gene: α_{o1})	Yes	Yes	Yes (?) ^d
α_{o2} (splice variant of α_{o1} gene ?)	Yes	No	No
α_i -rod	Yes	Yes	Yes
α_i -cone	No	Yes	Inferred
α_{of}	No	Yes	Inferred
α_{qx}	Yes	Yes	No
" α_p "-PhLC ^e	?, No ^f	?, No ^f	Yes
" α_p "-PhLA ^e	?, No ^f	?, No ^f	Yes

^a For detailed referencing see Birnbaumer et al.²¹⁸

^b See Table 3.

^c Named in chronologic order of cloning (Suki et al.²¹⁹).

^d Stimulatory roles in both K⁺ channel (Yatani et al.⁷) and phospholipase C regulation (Kikuchi et al.²²⁰) have been reported.

^e PTX sensitive and PTX insensitive activities have been reported, heterotrimeric nature of the G protein involved is inferred from PTX sensitivity and from inhibition of "G_p"-mediated activities by $\beta\gamma$ dimers (Moriarty et al.²²¹). For detailed discussion see Birnbaumer et al.²²²

^f The PTX sensitive form may have been purified/cloned, but the PTX-insensitive forms are unknown.

Table 3
Diversity in Subunit Composition of G Protein $\beta\gamma$ Dimers^a

	Refs.
β Subunits: β_1 (β_{36}), β_2 (β_{35}), β_3 (migration unknown) 4 cloned	
β_1	223-225
β_2	226-228
β_3	229
β_4	259
2 seen on SDS-PAGE	227,230,231,260
γ Subunits	
γ_T -1 (10 kDa); γ_T -2 (6kDa)	232-234
One cloned	235-236
2 seen on SDS-PAGE	237
Antigenicity distinct from that of γ_{GS}	238
Silver staining distinct from that of γ_{GS}	260
γ_G -a (6 kDa), γ_G -b (10 kDa):	239
Peptide map of 6 kDa form distinct from that of the 6 kDa γ_T	240
Antigenicity distinct from that of γ_T s	238
One cloned	241
At least two identified on SDS-PAGE	260
γ_G -c (7 kDa):	231
not cloned (?)	"
Migration on SDS-PAGE distinct from that of other γ_{GS}	260
Antigenicity distinct from that of other γ_{GS}	231

^a β and γ subunits are purified as dimers composed of mixtures of the subunits described in this table.

^b The protein of this cloned γ subunit has not yet been identified.

brane patches is not yet settled and, as shown later, is the object of intense research.¹³⁻¹⁵ The recognition that G proteins may affect ionic channels under cell-free conditions led us, as well as others to investigate this possibility further. By the most recent count (August 1989), six classes of ionic channels, comprising at least 12 separate molecular species defined by distinct kinetic and pharmacological properties, have been shown to be stimulated or inhibited under cell-free conditions by exogenously added G protein α subunits or by activation of a nearby G protein by GTP γ S, as seen in inside-out membrane patches or after incorporation of membrane vesicles into planar lipid bilayers. These ionic channels include channels that are essentially silent unless a G protein is activating them, referred to as G protein-gated ion channels,^{2,12} as well as channels that are merely regulated by the G proteins, such as an ATP-sensitive, K⁺ channel,¹⁶⁻¹⁸ an amiloride sensitive monovalent cation channel,¹⁹ a Ca²⁺-activated, charybdotoxin-sensitive, K⁺ channel,²⁰ and various voltage-gated ion channels,^{4,5,22-24} (summarized in Table 5).

In addition to the approximately 12 channels thus far shown to be influenced by G proteins in a manner that appears to be direct, i.e., not involving protein phosphorylation or changes in levels of second messengers such as cyclic nucleotides, diacylglycerol, or Ca²⁺, there may still be several more. This is suggested by reports on effects of PTX treatment or of GTP γ S and G protein subunit injection on whole cell currents. These effects include the inhibition by GTP γ S or pertussis toxin (PTX)-sensitive, G_o- and G_i-type G proteins of voltage-gated Ca²⁺ channels in chick²⁴ and rat²⁵⁻²⁷ dorsal root ganglion cells and of what seem to be similar channels in AtT-20²⁸ and NG108-15²⁹ cells, and the stimulation by PTX-sensitive, G_i-type G protein of voltage-gated Ca²⁺ channels in adrenal Y1 and GH₃ cells^{30,31} (summarized in Table 6). However, in these cases the involvement of soluble second messengers in the mediation of the effects of the G protein has yet to be ruled out.

The case of cell-free regulation of dihydropyridine (DHP)-sensitive, Ca²⁺ channels by G_s is of special interest. It was unexpected for two reasons: one because DHP-sensitive Ca²⁺ channels had been shown to be stimulated upon phosphorylation by the catalytic unit of cAMP-dependent protein kinase,³² indicating that nature uses dual pathways to regulate a single function, one fast and membrane delimited, the other slower with a longer life span; the other, because it had been thought that G proteins might be "monogamous", i.e., specific for single effector functions, and here we were faced with proof for multifunctionality in G protein actions.

These advances were all the result of a multidisciplinary approach to the problem of signal transduction by G proteins which brought together classical biochemistry, sophisticated single channel recordings, and modern molecular biology. The background experiments, especially those of Nargeot et al.,³³ Soejima and Noma,³⁴ Pfaffinger et al.,³⁵ and Breitwieser and Szabo,³⁶ which led to the discovery that ionic channels are effector systems of G proteins akin to adenylyl cyclase and

Table 4
Classes G Protein Effector Functions

Effector class ^a	G Protein(s) ^b	Effect	Refs./Comment
Proven in Cell-Free Assays			
Adenylyl cyclases	G _i / ^c "G _i "	Stimulation/inhibition	Table 3
Retinal cGMP-specific PDE	G _i	Stimulation	Table 3
Phospholipase C	"G _p " (G _o ?)	Stimulation	222,242,243
Phospholipase A ₂	"G _p "	Stimulation	75,172,222
Ionic Channels (six types)	G _k = G _i s/G _o s; G _s	Stimulation/inhibition	Table 6
Suggested by Indirect or Intact Cell Studies^c			
Na/H Antiport	"G _p " (?)	Stimulation	244 ^d
Voltage-gated Ca ²⁺ channels	G _s , G _o s (?)	Inhibition	Table 7
Insulin-sensitive glucose transporter	G _s (?)	Stimulation	245 ^e
Liver Ca ²⁺ Pump	G _s (?)	Inhibition	246 ^f
Renal Na/K ATPase	?	Inhibition	247 ^g
More ionic channels (?)	?	Stimulation/inhibition	Table 7
Phospholipase D	"G _p " (?)	Stimulation	76 ^h
Nonretinal phosphodiesterases (PDEs)	?	Stimulation	248 ⁱ

^a The number of distinct gene products in each class may be high. For example, work from the laboratories of Gilman and Randall Reed shows that there are at least three adenylyl cyclases; the atrial, GH₃, and hippocampal neuron G_k-gated K⁺ channels differ in conductance and G protein specificity; the heart and skeletal muscle dihydropyridine sensitive Ca²⁺ channels stimulated by G_s are products of separate genes.

^b Except in the cases of adenylyl cyclase inhibition ("G_i") and phospholipase C activation ("G_p"), the G protein involved is known. In the cases of "G_i" and "G_p" *in vitro* experiments with activated purified α subunits have failed. Although some preliminary data indicate that the PTX sensitive "G_p" may be one of the G_o variants, confirmation is needed.

^c These are systems in which mediation by phosphorylation has not been ruled out, but where indications are strong that a direct G protein regulation is possible.

^d Inferred from persistent PTX-sensitive thrombin response in TPA-desensitized fibroblasts (Paris and Pouyssegur²⁴⁴).

^e Inferred from kinetics and toxin sensitivity of agonist-mediated regulation of glucose transport in fat cells (Kuroda et al.²⁴⁵).

^f Inferred from effect of glucagon peptide 19-29 on liver C pump (Mallat et al.²⁴⁶).

^g Inferred from dopamine mediated regulation of Na-K ATPase (Bertorello and Aperia²⁴⁷).

^h Inferred from the fact that receptors that increase phospholipase D activity are coupled by G proteins and that GTP γ S stimulates phospholipase D activity in isolated membranes (Bocchino et al.⁷⁶).

ⁱ Inferred from agonist-induced PTX-insensitive increased rates of cAMP hydrolysis in Hughes et al.²⁴⁸

Table 5
Ionic Channels Regulated Under Cell-Free Conditions by Pure G Protein and/or GTP γ S

Channel type	G protein	Effect	Tissue/cell	Refs.
K _{CK} 40 pS	G _i (1,2,3)	Stim.	Heart	2,7,8,10,96
K _{CK} 50 pS	G _{i3}	Stim.	GH ₃	3,8,43
K _{CK} 4 types	G _{o1}	Stim.	Hippocampal neurons	12
K _{ATP}	G _{i3}	Stim.	RIN, heart, skeletal muscle	16-18
Na(K) _(Aml)	G _{i3}	Stim.	Renal med. collect. tubule	19
Ca _(DHP) L-type	G _s (all 4)	Stim.	Heart, skeletal muscle	4,5,11,21
Na _(TTX)	G _s	Inh.	Heart	22
K _{Ca(Charyb)}	? (Iso/GTP)	Stim.	Uterine smooth muscle	20
Ca _(Ni) T-type	? (GTP γ S)	Inh.	Rat DRGs	23

Table 6
Ionic Channels That May Be Under Direct Regulation by G Proteins as Inferred from Whole Cell Recordings

Channel type	Agonist	G protein ^a	Effect	Cell	Mimicked by TPA	Refs.
Ca (N-type?)	Opioid	G _o >G _i	Inh.	NG108-15	?	29
Ca (L-type)	Ang II	G _i	Stim.	Y1 Adrenal	?	30
Ca (L-type)	GnRH	G _i /G _o	Stim.	GH ₃	?	31
Ca (L-type)	SST	? GTP-γS	Inh.	AtT-20	Yes	28
Ca (?)	GABA _(B)	? GTP-γS	Inh.	Chick DRG	Yes (total)	24,249
	NE (αAR)	? GTP-γS	Inh.	Chick DRG	Yes	24,249
Ca (?)	GABA _(B)	? GTP-γS	Inh.	Rat DRG	No	25,250,251
Ca (?)	NPY	G _o >G _i	Inh.	Rat DRG	Yes (partial)	26,252
	Bradyk.	G _o = G _i	Inh.	Rat DRG	Yes	27

^a All the G proteins involved are PTX-sensitive.

cGMP-phosphodiesterase, and the initial experiments showing effects of purified G proteins and protein subunits on channels in excised membrane patches, were reviewed in Brown and Birnbaumer.¹ The present article focuses on some of the more recent results from our laboratories dealing with a PTX-sensitive, G_i-type family of G proteins, their effects of ionic channels, our efforts in assigning defined functions to individual G proteins as they are known from biochemical and molecular cloning studies, and some speculations that follow from the results obtained as to why G proteins dissociate and how transduction pathways are set up.

II. PRIMARY STRUCTURE OF G PROTEINS

The primary αβγ structure of G proteins is summarized in Figure 2 and Tables 2 and 3, and has been reviewed by Lochrie and Simon,³⁷ as well as by us.³⁸ Key to the understanding of their functioning is of course both their subunit dissociation cycle superimposed on their GTPase cycle, and their transient — or perhaps not so transient — interactions with receptors and effectors. In our view it is clear that the receptor signal is carried to the effector by the α subunit as exemplified in the experiment of Figure 3. At the time of this writing, 12 α subunits, encoded in 9 genes, 4 β subunits, and at least 3 γ subunits (all mammalian) are known (Tables 2 and 3). β and γ subunits form dimers which may be of several types, β₃₆γ and β₃₅γ, if a cell expresses only one type of γ subunit, or of more types if it expresses more β or γ subunits. To our knowledge, α subunits, when combining with βγ dimers to form holo-G proteins, do not distinguish among βγ dimers (Figure 4). This is not to say that all tissues have the same complement of βγ dimers. Quite the contrary, as shown in Figure 5, β₃₅/β₃₆ ratios differ in human placenta, human erythrocytes, bovine brain, and bovine retinal rod cells. α Subunits bind GTP, hydrolyze GTP, dissociate from the βγ dimer on activation by

GTP analogs, and/or NaAlF₄/GDP, and, with few exceptions, are substrates for ADP-ribosylation by cholera and/or pertussis toxin (CTX and PTX). Studies with transducin-α identified an arginine at the approximate center of the molecule as the amino acid ADP ribosylated by CTX³⁹ and a cysteine at position -4 from the carboxyl terminal end as the site of ADP-ribosylation by PTX.⁴⁰ It is worth mentioning that not all G proteins have been either purified or cloned, and that not all the known G proteins, such as G_{o1} and G_{o2}^{41,42} or some of the G_is, have unequivocally assigned functions, or that all functionally recognized G proteins, such as the inhibitory G_i of adenylyl cyclase or the stimulatory G_p of phospholipases, have been identified.

III. FUNCTIONAL STUDIES

A. Combined Use of Natural and Recombinant α Subunits Made in Bacteria to Define Their Functions

The abundance at which different α_s and α_i molecules are expressed varies from tissue to tissue, raising the question as to whether functional differences are associated with the structural differences or whether the G_ss and, respectively, the G_is should be thought of merely as isoforms. Although the final word on these questions is not yet in, we have during this last year developed the method(s) required to answer them. Thus, we have so far succeeded in purifying two types of G_i from human erythrocytes (hRBCs) and a third from bovine brain, from which we also purified the two forms of G_o. This allowed us to test for potential differential biological functions. We also expressed biologically active forms of the α subunits in bacteria, designated as recombinant α subunits, so that we could predict/confirm biological functions of various cloned and/or purified G protein α subunits.

We have not yet carried out all of the studies. However, we were able to determine first of all that both the natural purified and the recombinant forms of α₁₃, α₁₁, and α₁₂ all stimulate K⁺

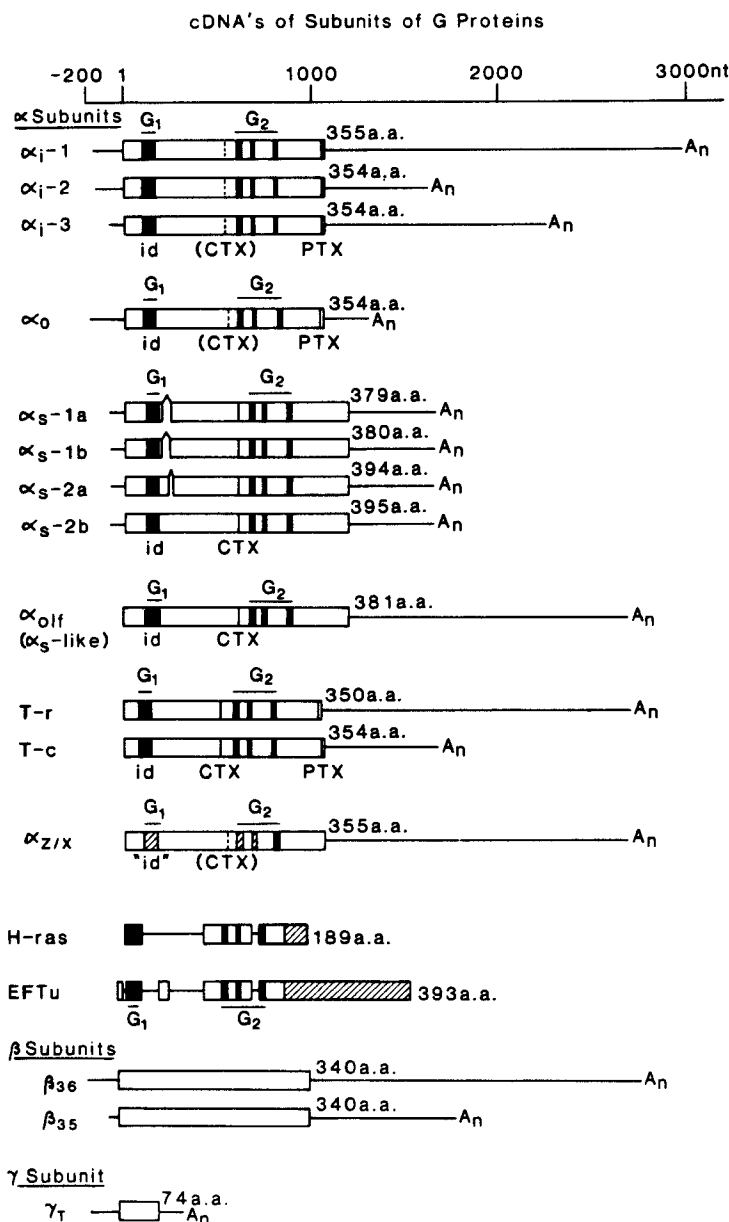


FIGURE 2. Schematic representation of vertebrate α subunit mRNA molecules as deduced from cDNA cloning. Open boxes represent the open reading frames or coding sequences and lines represent 5' and 3' untranslated sequences which may be incomplete. Black boxes within the open reading frames of α subunits denote sequences highly homologous to those known in bacterial elongation factor TU to be involved in GTP binding and hydrolysis. Sequences homologous to these are present also in the *ras* molecules. The mRNA molecules encoding the β_{36} , β_{35} , and γ_T are shown for comparison. The position of amino acids ADP-ribosylated by CTX and PTX are indicated. i.d., location of the identity box. The scale is in nucleotides.

channels (Figure 6; References 7 and 43) We failed to observe significant differences in the potency with which type 1, 2, and 3 α_i molecules activated K^+ channels, even though the recombinant forms all had potencies between 30- and 50-fold lower than their natural counterparts.⁷ Thus, with respect to

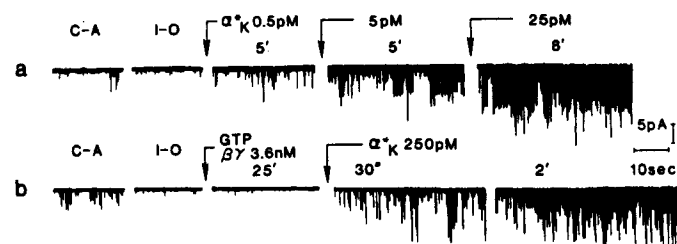


FIGURE 3. (a) Stimulation of single channel K^+ currents in GH₃ cell membrane patches by increasing concentrations of native GTP γ S-activated human erythrocyte α_{13} (α_K). (b) Lack of intrinsic stimulatory effects of human erythrocyte $\beta\gamma$ dimer added in the presence of GTP in a responsive membrane patch. Lubrol PX (500 nM) was present throughout. This and similar figures shown next present segments of records obtained in the cell-attached mode (C-A) and in the inside-out mode after patch excision and before additions (I-O). Additions are denoted by arrows and time elapsed between addition of test substances are given above the records.⁹

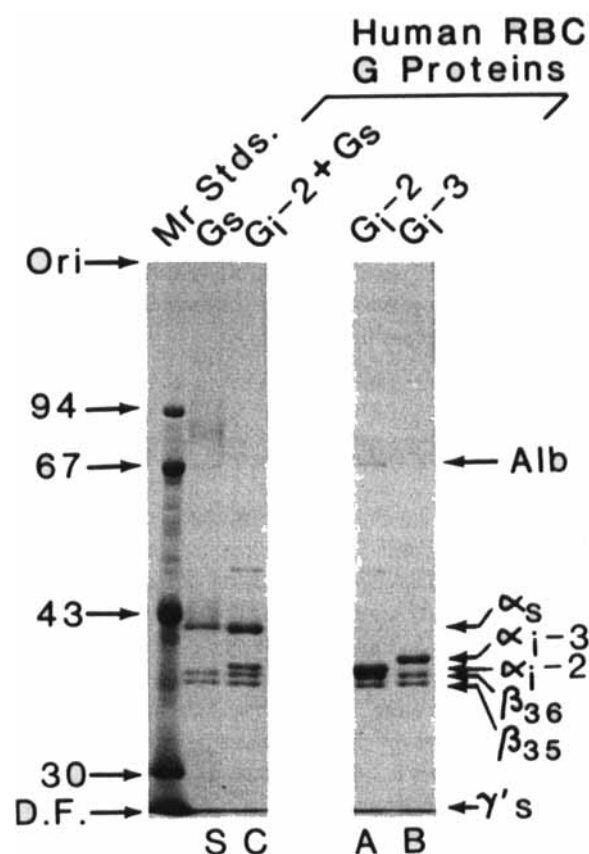


FIGURE 4. Urea gradient/SDS-PAGE analysis of G_α , G_{i-2} , and G_{i-3} purified from human erythrocyte membranes. Shown is the photograph of a Coomassie blue stained gel. Note: (1) differential migration of α subunits as well as of the two β subunits, (2) that each of the proteins has its share of $\beta\gamma$ dimers, and (3) that this does not differ from one-to-another.

atrial muscarinic K^+ channels, G_{i1} , G_{i2} , and G_{i3} must be considered iso-G proteins. Studies are currently in progress in atrial membrane patches in which the endogenous G_K has been

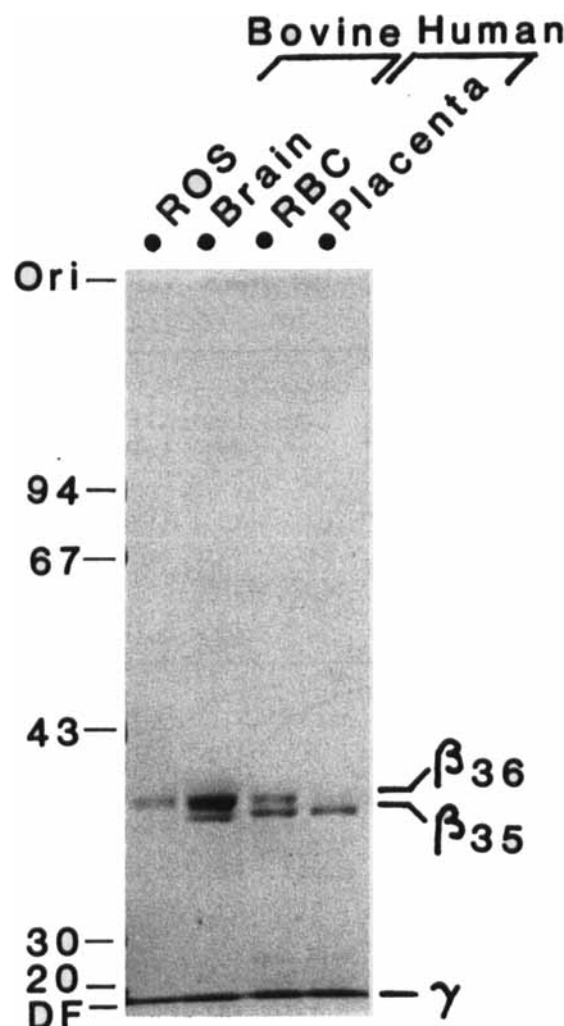


FIGURE 5. Urea gradient/SDS-PAGE analysis of four preparations of $\beta\gamma$ dimers. Shown is the photograph of a Coomassie blue stained gel. Note that different tissues express β_{35} and β_{36} at different ratios.

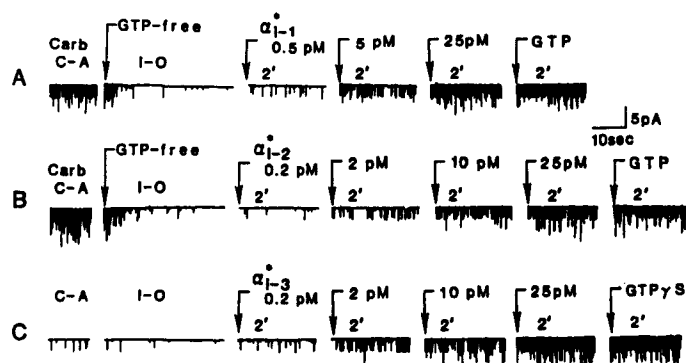


FIGURE 6. Lack of significant difference in stimulation of atrial G_i -gated K^+ channels obtained with the activated α subunits of G_{i-1} , G_{i-2} , and G_{i-3} . Presence or absence carbachol (Carb) is denoted at the beginning of each train of records.⁷

uncoupled from receptors by treatment with PTX,^{1,3} to determine whether muscarinic and/or P_2 purinergic receptors exhibit selectivity for interaction with one or another of the G_i proteins.

Second, by testing the effect of recombinant α_s , we were able to determine that the stimulation of Ca^{2+} channels obtained with purified hRBC G_s ^{4,5,11} is indeed mediated by G_s as opposed to being due to a contaminant. Furthermore, in collaboration with Michael Graziano and Al Gilman we tested the recombinant forms of three of the possible four splice variants of α_s for their Ca^{2+} channel stimulatory activity and found that they all do so with indistinguishable potency and efficacy.²¹ As was the case with recombinant α_i subunits, the recombinant α_s subunits also displayed a 20-fold reduced potency with respect to that of native human erythrocyte α_s . This applied not only to Ca^{2+} channel stimulation, but equally to adenylyl cyclase stimulation. Very likely, bacteria fail to carry out a critical posttranslational modification that exists only in eukaryotic cells, or alternatively, modify the α subunit in a manner that eukaryotic cells do not.

B. Gating of Ionic Channels as a Tool to Discover New Roles for G Proteins: Effects of G_o on Neuronal K^+ Channels

One of the properties of the "muscarinic" K^+ channels is that they are essentially silent in the absence of stimulation by an activated G protein (G_k). That is, in the absence of activated G protein their P_o is close to zero. The possibility existed that not only G_i proteins regulate K^+ channels but also the structurally closely related G_o . Since nervous tissue is rich in G_o , central nervous system neurons, specifically hippocampal pyramidal cells, were placed into culture and studied for potential presence of both G_i - and G_o -gated K^+ channels. Although these studies are still in progress,²⁵⁸ the initial findings with highly purified bovine brain GTP γ S-activated G_{o1} (G_{o1}^*) were of interest.¹² They identified the existence of several novel G-protein gated, more precisely G_o -gated K^+ channels, that are distinct from G_i -gated K^+ channels. Thus, application of purified bovine brain G_{o1}^* to the cytoplasmic aspect of inside-out membrane patches of cultured hippocampal pyramidal cells resulted in appearance of three new types of single channel K^+ currents consistent with the existence of the three nonrectifying of K^+ channels having sizes of 13, 40, and 55 pS, respectively, plus an inwardly rectifying K^+ channel with a slope conductance of 40 pS. No such channel activities were observed with hRBC G_i^*-3 or hRBC α_i^*-3 . G_{o2}^* or α_{o2}^* have not been tested in this system as yet. In contrast to earlier observations with G_o^* added to guinea pig atrial membrane patches, which showed only marginal effects of GTP γ S-activated G_{o1} at 2 nM², the hippocampal K^+ channel is highly sensitive to G_{o1}^* ,¹² and significant activation was obtained at 1 pM and half maximal effects were obtained at about 10 pM. The identity of the active G protein in the G_{o1} preparations used was confirmed with recombinant GTP γ S-activated α_{o1} . The G_o -gated channels were

stimulated in the absence of Ca^{2+} or ATP, in the presence or absence of AMP-P(NH)P, added routinely to inhibit ATP-sensitive, 70 pS K^+ channels. Furthermore, EGTA did not interfere with the actions of $\text{G}_{\text{o}1}$ or recombinant $\alpha_{\text{o}1}$. Thus, in hippocampal pyramidal cells of the rat, $\text{G}_{\text{o}1}$ is a G_k , and the K^+ channels gated by it are several and differ from those present in atrial cells in various aspects including G protein specificity. These findings raise the question which if any of the G proteins that gate K^+ channels regulate the other known PTX sensitive effector systems.

IV. $\beta\gamma$ DIMERS INHIBIT K^+ CHANNEL GATING BY G PROTEIN

Logothetis et al.^{13,14} reported twice that $\beta\gamma$ dimers stimulate atrial K^+ channel activity. Their finding is not reproduced in our hands. Quite the contrary, when we add $\beta\gamma$ dimers to inside-out membrane patches in which K^+ channels have been stimulated either by GTP only (baseline activity) or by carbachol plus GTP (agonist-stimulated activity), we find consistently inhibition of activity (Figure 7). On their own, i.e., when added to silent patches in the absence of GTP, $\beta\gamma$ dimers have no effect under our assay conditions (Figure 3; References 8 to 10).

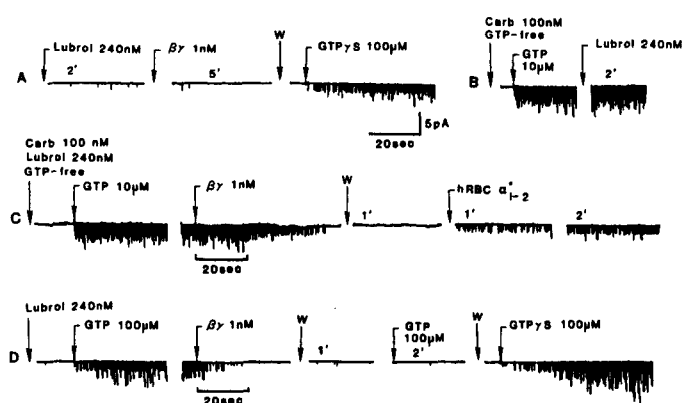


FIGURE 7. Inhibition of G protein-gated K^+ channel activity by $\beta\gamma$ dimers. Inside-out membrane patches from adult guinea pig atrial cells were exposed to the bathing solution (140 mM KCl, 2 mM MgCl_2 , 5 mM EGTA, 10 mM HEPES-K, pH 7.4) containing the additives shown on the figure. The pipette solution was identical to the bathing solution and contained 100 nM carbachol when indicated. The composition of bathing solution was changed by a concentration clamp method. Note that Lubrol PX and/or bovine serum albumin, used to maintain $\beta\gamma$ dimers in suspension do not interfere with stimulation of activity by GTP (D) or GTP plus agonist (B, C), and that $\beta\gamma$ dimers have no effect on their own (A) but inhibit atrial K^+ channel stimulation by the membrane G_k (C, D). Inhibition is faster and is elicited with lower concentrations of $\beta\gamma$ dimers when K^+ channels are operating under baseline conditions (GTP only, experiment D: cessation of activity after 16 s) than when they are stimulated by agonist (Carbachol plus GTP: experiment C: cessation of activity after 50 s). Numbers above records denote time elapsed in min between solution change and the beginning of the record shown.

Concentration effect studies showed clearly that $\beta\gamma$ dimers are more potent in inhibiting agonist-independent than agonist-stimulated activity and that this phenomenon applies not only to $\beta\gamma$ dimers suspended in Lubrol-PX, such as those from human placenta, human erythrocytes, and bovine brain, but also to $\beta\gamma$ dimers presented to the patches in aqueous media, such as transducin $\beta\gamma$ (Figure 8).

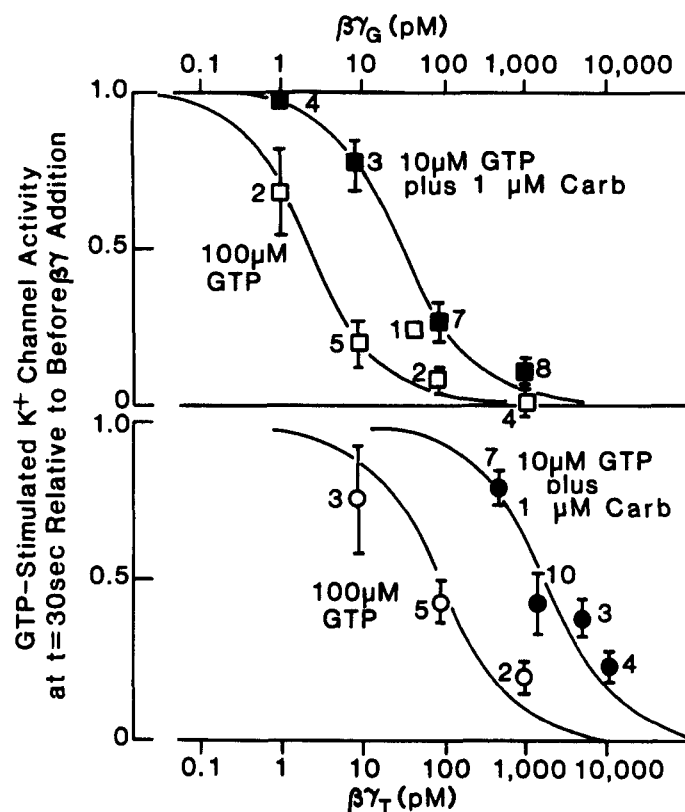


FIGURE 8. Effect of agonist (carbachol) on the dose-dependent inhibition by $\beta\gamma$ dimers of GTP-dependent K^+ channel activities in inside-out guinea pig atrial membrane patches by $\beta\gamma$ dimers. When 1 μM carbachol was present in the pipette, GTP was 10 μM ; in the absence of carbachol GTP was 100 μM . $\beta\gamma_{\text{G}}$: data obtained with $\beta\gamma$ dimers derived from either human erythrocyte, human placenta, or bovine brain were pooled. $\beta\gamma_{\text{T}}$: data obtained with $\beta\gamma$ derived from bovine rod outer segments. (We thank Dr. Tony Evans for the gift of human placental $\beta\gamma$ dimers, and Dr. J.-K. Ho, for the gift of bovine rod outer segment $\beta\gamma$ dimers.)

The fact that $\beta\gamma$ dimers inhibit GTP-dependent activity in the absence of agonist at lower concentrations as compared to inhibition in the presence of agonist was observed previously in purified components reconstituted into phospholipid vesicles (Figure 9; References 44 and 45) and serves to support our thesis that $\beta\gamma$ dimers act in intact membranes as suppressor of "noise" generated by agonist-unoccupied receptors.

The inhibitory effects of $\beta\gamma$ dimers obtained by us need to be contrasted to stimulatory effects obtained by Clapham, Neer, and their collaborators.^{13-15,46} We do not understand exactly

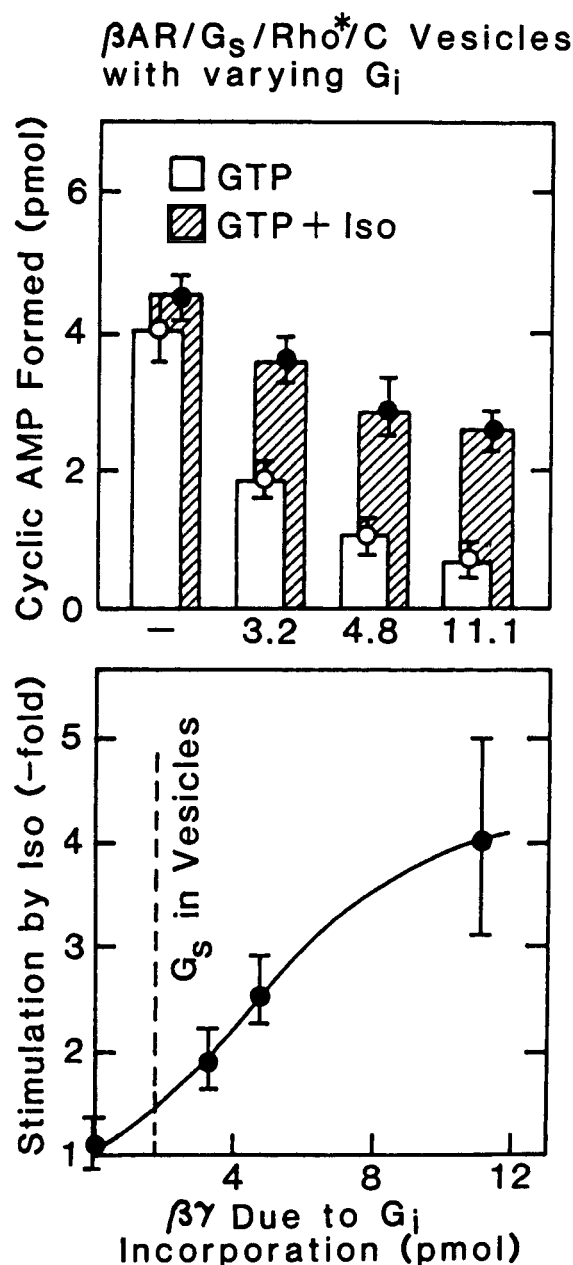


FIGURE 9. $\beta\gamma$ Dimers formed upon activation of G_i by rhodopsin, are potent inhibitors of agonist-independent but not of agonist-stimulated G_s activity. β -Adrenoceptors, G_s , and resolved bovine brain catalytic unit of adenylyl cyclase were incorporated into phospholipid vesicles in the presence of photoactivated rhodopsin and the indicated amounts of G_i . Vesicles were resuspended in buffer with GTP and assayed for adenylyl cyclase activity in the absence and the presence of the β -adrenoceptor agonist isoproterenol. Top panel: absolute activities; Bottom panel: relative stimulation by isoproterenol. Note that the selective inhibition of receptor plus GTP induced activation of G_s by $\beta\gamma$ dimer is an essential feature for functional expression of the agonist effect.⁴⁴

the reasons for the discrepancy since we obtain inhibition also in the absence of Lubrol PX using $\beta\gamma$ dimers from transducin which are water soluble. The claim that $\beta\gamma$ dimers may be acting by stimulation of arachidonic acid formation and acting through a metabolite⁴⁶ is suspect. In an adjacent report, Kurachi et al.⁴⁷ describe that arachidonic acid and its metabolites require for their action the presence of GTP and are blocked by GDP β S. Logothetis et al.^{13,15} and presumably also Kim et al.⁴⁶ obtain stimulation with $\beta\gamma$ dimers in the absence of GTP.

V. CURRENT VIEWS ON HOW SIGNAL TRANSDUCTION BY GTP AND RECEPTORS COMES ABOUT AND WHICH RECEPTOR ACTS ON WHICH G PROTEIN TO REGULATE WHICH EFFECTOR SYSTEM

Taken together the results discussed in the previous sections lead to several conclusions: (1) ionic channels are targets of direct regulation by G proteins as are adenylyl cyclase and the cGMP-specific phosphodiesterase; (2) α subunits and not $\beta\gamma$ dimers are the specificity determinants of signal transduction pathways; (3) several G proteins may have the same function, e.g., stimulation of K^+ channels by three G_s ; and (4) a single G protein may have more than one function, i.e., be multifunctional (e.g., stimulation of adenylyl cyclase and the dihydropyridine sensitive Ca^{2+} channel by a single G_s).

Two independent sets of questions emerge from these findings. The first deals with the subunit dissociation reaction and asks what type of advantage it confers onto the system by its existence. An answer to this can be found upon analyzing in detail the G protein regulatory cycle and the mechanism by which receptors promote G protein activation by GTP.

The second set of questions deals with crosstalk between signal transduction pathways, i.e., whether receptors act on more than one G protein, and if so, which and whether G proteins interact with more than one receptor as well as with more than one effector, and if so, how frequent this is.

A. Role of Subunit Dissociation: Requirement for Catalytic Action of Receptors

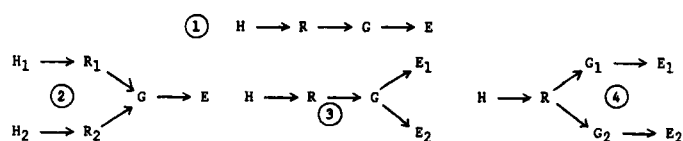
In the second half of the 1970s, it was demonstrated that receptors act catalytically rather than stoichiometrically to activate adenylyl cyclase,⁴⁸ and in 1980 it was found that receptors in addition to promoting GDP/GTP exchange^{49,50} also participate in promoting the activation reaction proper of adenylyl cyclase by GTP and its analogs,⁵¹⁻⁵³ reviewed in Reference 54. These two findings, catalytic action and stabilization of the nucleotide-activated form of adenylyl cyclase, now known to be in fact G_s , are thermodynamically impossible, unless the stabilized form of the G protein undergoes some type of additional spontaneous change that releases it from

microscopic reversibility constraints, which happens in all enzymatically catalyzed reactions where the product is chemically different from the starting substrate. Dissociation of $\beta\gamma$ from the activated and stable receptor-G protein complex is such a reaction change. We therefore propose that the role of the dissociation reaction is a requirement without which receptors cannot act catalytically. Both the catalytic nature of receptor-mediated activation of G_i and the fact that receptor affects not only GDP release but also the rate at which inactive guanine nucleotide-occupied G protein isomerizes from an inactive to an active state have been confirmed in reconstitution experiments with purified receptor and purified G_i .^{55,56} Since receptors do not interact with the α subunits except in the context of $\beta\gamma$,⁵⁷⁻⁶¹ reassociation of the GTPase-deactivated α with $\beta\gamma$ is essential for restimulation by receptors. This then leads to a description of the G protein regulatory cycle under influence of receptor as depicted in Figure 10.

Taken together, it is thus possible to ascribe three generic functions to $\beta\gamma$ dimers (Table 7): (1) activation of α subunit by receptor, for without $\beta\gamma$, receptors do not interact with α ; (2) amplification of the receptor signal, for without dissociation, receptors cannot act catalytically; and (3) noise reduction, for unoccupied receptors are not silent.

B. Specificities in Receptor-G Protein-Effector Interaction

Ever since the discoveries in the late 1960s that up to five different hormone receptors can activate a single adenylyl cyclase system in an isolated membrane^{62,63} and, in the early and mid-1970s, that receptors can be transferred from one cell to another⁶⁴ and that there are no species and/or tissue specificity restrictions as to the source of G_i for reconstitution of a hormonally stimutable adenylyl cyclase system in *cyc*⁻ membranes,^{65,66} it has been clear that single G proteins are designed to interact with classes of receptors as opposed to single receptor subtypes. The discovery that the same splice variant of G_i that activates adenylyl cyclase is also able to regulate Ca^{2+} channel activity²¹ indicates that one G protein can interact with more than one effector. The discovery that three different G_i proteins all activate the same K^+ channel,⁷ indicates that several G proteins may regulate a single effector. Ashkenazi et al.⁶⁷ showed that single receptors may affect more than one G protein. It follows that transmembrane wiring diagrams are in fact combinations of the following four basic configurations:



The complexity that may exist in the wiring of transmembrane signal transmission was well illustrated by the findings of Ewald

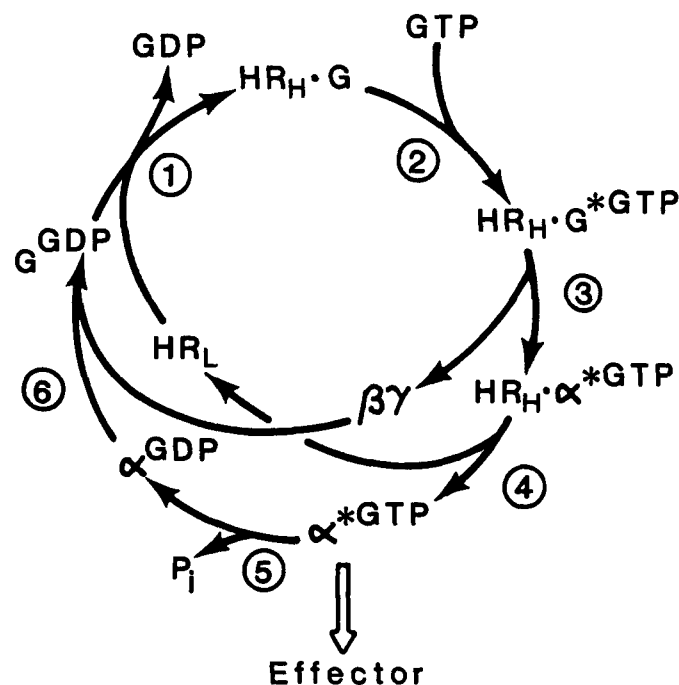


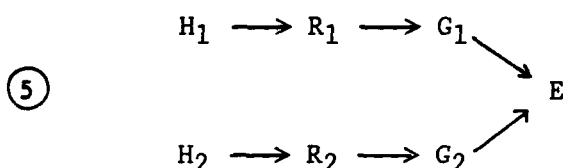
FIGURE 10. Integrated view of receptor-mediated catalytic activation of a G protein in the context of the dual subunit dissociation and GTPase cycles of G proteins. The role of receptors is to promote nucleotide exchange and to stabilize a GTP-dependent "activated" form of the G protein and the G protein undergoes a cyclical dissociation-reassociation reaction and oscillates between GDP, nucleotide free, and GTP states. The cycles are driven energetically forward by the capacity of the G protein to hydrolyze GTP²⁵³⁻²⁵⁷ and kinetically by the dissociation of $\beta\gamma$ dimer from the activated receptor-G proteins complex. The receptor has high affinity for agonist (R_H) when associated with the nucleotide-free trimeric $\alpha\beta\gamma$ form of the G protein, and has low affinity for the agonist (R_L) when it is free. Furthermore, the receptor has higher affinity for the trimeric $\alpha\beta\gamma$ form of G than the G-GDP thus accounting for the finding that GDP and GDP analogs promote the R_H to R_L transition. The $\beta\gamma$ dimers are required for the interaction of α with R, and after formation of G-GTP, no G^* forms unless it is "aided" by receptor. Thus, receptor has an even higher affinity for the G^*GTP state than the nucleotide free state of G. As a consequence receptor dissociation is absolutely dependent on reaction 2 (subunit dissociation). Thermodynamic reasons do not allow R both to stabilize the G^* state and to dissociate from it. Reaction 3 states further that the α^*GTP loses its ability to stay associated with receptor and decomposes further into free activated α^*GTP plus free receptor, thus accounting for the fact that under "working" conditions (saturation by both GTP and hormone, and hence sustained regulation of effector) only a small proportion of receptors are found in their high affinity, G-protein associated state. It follows that the G protein cycle is driven forward not only by the GTPase but also, and obligatorily so, by the subunit dissociation reaction. The scheme accounts for the experimental findings: (a) that receptors act catalytically and therefore need to dissociate from the G protein at one time or another;⁴⁸ and (b) that receptors accelerate the transition from inactive G-GTP γ S to active G * -GTP γ S transition and therefore must have higher intrinsic affinity for the activated than the inactive state.⁵¹⁻⁵³

et al.²⁷ in PTX-treated rat sensory neurons. On studying the efficacy with which brain G_i and G_o reconstitute Ca^{2+} current regulation by neuropeptide Y (NPY) and bradykinin, they discovered that the effect of NPY could be fully reconstituted with G_o , with G_i being much less potent, while the effect of

Table 7
Intramembrane Roles of G Protein $\beta\gamma$ Dimers

Reaction	Product	Role
Reassociation with α^{GDP}	G^{GDP}	Activation by R
Dissociation from $HR.G^{GTP}$	$HR.\alpha^{*GTP}$	Signal amplification
Reassociation with $R.\alpha^{*GTP}$	$R.G^{GTP}$	Noise reduction

bradykinin could only be partially reconstituted by G_o , requiring G_i to achieve full reconstitution (Diagram 5).



The important notion that emerges from these findings is that the wiring diagrams describing signal transduction by G proteins needs to be determined individually and separately for each cell or tissue of interest. This includes the determination not only of which receptors are present but also which G proteins and effectors process the receptor signals.

VI. CONCLUSION

Signal transduction by G proteins is a fundamental and widespread mechanism used by a wide variety of hormones, neurotransmitters, and auto- and paracrine factors to regulate cellular functions. G proteins modulate not only cAMP formation, but also intracellular Ca^{2+} mobilization, arachidonic acid release, and, very importantly, membrane potential, with the latter not only being a trigger for neurotransmitter release, but also a conductor of nerve impulses. In tissues such as secretory cells, it is the main regulator of Ca^{2+} entry. In heart, action potentials play the dual role of determining the frequency of contraction, and through modulation of the duration of the depolarized state, membrane potential determines Ca^{2+} entry and the force of contraction. More subtle changes in resting membrane potential alter the cell's predisposition to be stimulated by other factors and hormones. It is easy to imagine that persistent changes in membrane potential may affect acutely and chronically the cell's proliferative properties.

The mechanism by which G proteins are activated provides for amplification, reversal of action, and continued monitoring of hormone: for an amplification because few receptor molecules may act catalytically to activate many G proteins molecules; for reversal of action because they have an internal turnoff mechanism whereby the $G\alpha$ subunit hydrolyzes GTP to GDP; and for continued monitoring of the primary messenger level because each activation cycle requires not only GTP but also occupied receptor.

Not all G proteins are known and some are known of which their functions are still unknown. However, more G proteins and more effector functions affected by them will surely be found. Work is in progress to unravel a complicated network of interactions between receptors, G proteins, and effector systems that not only affect regulation of metabolic activities of organs such as liver, heart, and fat, but also of the integrative functions of the central nervous system.

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