

## Transdominant inhibition of tyrosine kinase activity in mutant insulin/insulin-like growth factor I hybrid receptors

(*in vitro* assembly/heterologous receptors/substrate phosphorylation/autophosphorylation)

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Communicated by Donald F. Steiner, October 15, 1990

**ABSTRACT** Classical insulin and insulin-like growth factor I (IGF-I) receptors exist as well defined  $\alpha_2\beta_2$  heterotetrameric complexes that are assembled from two identical  $\alpha\beta$  heterodimeric half-receptor precursors. Recent evidence suggests that insulin and IGF-I half-receptors can heterologously assemble to form  $\alpha_2\beta_2$  insulin/IGF-I hybrid receptor complexes *in vivo* and *in vitro*. We have utilized hybrid receptor complexes to examine ligand-stimulated transmembrane signaling of wild-type insulin ( $\alpha\beta_{\text{INS.WT}}$ ) or IGF-I ( $\alpha\beta_{\text{IGF.WT}}$ ) half-receptors assembled with a kinase-defective insulin half-receptor mutant ( $\alpha\beta_{\text{INS.A/K}}$ ). *In vitro* assembly of either ( $\alpha\beta_{\text{IGF.WT}}/(\alpha\beta_{\text{INS.A/K}}$ ) or ( $\alpha\beta_{\text{INS.WT}}/(\alpha\beta_{\text{INS.A/K}}$ ) hybrid receptors resulted in decreased substrate protein kinase activity. The degree of protein kinase inactivation directly correlated with the amount of immunologically cross-reactive hybrid receptors formed. In contrast to substrate kinase activity, insulin-stimulated autophosphorylation of the ( $\alpha\beta_{\text{INS.WT}}/(\alpha\beta_{\text{INS.A/K}}$ ) hybrid receptor complex was completely unaffected in comparison to the wild-type ( $\alpha\beta_{\text{INS.WT}}/(\alpha\beta_{\text{INS.WT}}$ ) receptor. To assess a molecular basis for this difference, autophosphorylation of a hybrid receptor composed of a truncated  $\beta$ -subunit insulin half-receptor with the kinase-defective half-receptor, ( $\alpha\beta_{\text{INS.ACT}}/(\alpha\beta_{\text{INS.A/K}}$ ), demonstrated the exclusive autophosphorylation of the ( $\alpha\beta_{\text{INS.A/K}}$ ) half-receptor  $\beta$  subunit. These results demonstrate that ligand-dependent substrate phosphorylation by insulin and IGF-I holoreceptors requires interactions between two functional  $\beta$  subunits within the  $\alpha_2\beta_2$  heterotetrameric complex and occurs through an intramolecular trans-phosphorylation reaction.

Insulin and insulin-like growth factor I (IGF-I) holoreceptors share a large degree of structural and functional similarity (1, 2). The mature  $\alpha_2\beta_2$  holoreceptors are synthesized from  $\alpha\beta$  fusion proreceptor precursors that are cotranslationally acylated and glycosylated and then transported to the Golgi apparatus where additional glycosylation, proteolytic cleavage, and assembly into an  $\alpha_2\beta_2$  complex occurs (3–9). The mature  $\alpha_2\beta_2$  receptor state is required for ligand-stimulated transmembrane signaling, since lower oligomeric forms such as isolated  $\alpha\beta$  heterodimers (10–14) and truncated  $\alpha_2\beta\beta'$  receptors (15) are kinase-inactive species. Immunologically cross-reactive  $\alpha_2\beta_2$  insulin/IGF-I hybrid receptors have been described that appear to result from the heterologous assembly of individual  $\alpha\beta$  insulin and IGF-I receptor precursor proteins *in vivo* (16, 17). In parallel, we have observed that insulin/IGF-I hybrid receptor complexes can be assembled from purified insulin and IGF-I  $\alpha\beta$  half-receptors *in vitro* (18).

The identification of hybrid receptor complexes has raised the question whether functional alterations in transmembrane signaling could result from the assembly of wild-type receptor precursors with dysfunctional receptor subtypes. For example, it has been observed that heterozygote individuals that express both wild-type and mutant tyrosine kinase-defective insulin receptor precursors have severe insulin resistance and diabetes (19–22). In these patients, the degree of insulin resistance is significantly greater than would be predicted from a simple loss of half the normal complement of insulin receptors. Similarly, cell lines that coexpress both endogenous wild-type and transfected kinase-defective insulin receptors display a marked decrease in insulin sensitivity and/or responsiveness (23–26). To determine whether defective insulin signaling through hybrid receptors could contribute to the effects observed, we have examined the kinase activity of  $\alpha_2\beta_2$  heterotetrameric hybrid receptors composed of kinase-defective A/K<sub>1018</sub> mutant insulin half-receptors (24) assembled *in vitro* with a wild-type insulin or IGF-I half-receptor.

### MATERIALS AND METHODS

**Isolation of  $\alpha\beta$  Heterodimeric Insulin and IGF-I Receptor Complexes.** Human placenta membranes (20 mg/ml) were treated with 2 mM dithiothreitol for 5 min at pH 8.5 to reduce and dissociate the  $\alpha_2\beta_2$  heterotetrameric receptors, followed by detergent solubilization and partial purification by Bio-Gel A-1.5m gel filtration or wheat germ agglutinin-Sepharose chromatography (27). The ( $\alpha\beta_{\text{IGF.WT}}$ ) half-receptors were isolated by immunoaffinity purification by using the anti-IGF-I receptor monoclonal antibody  $\alpha\text{IR-3}$  or by immunodepletion of ( $\alpha\beta_{\text{INS.WT}}$ ) half-receptors by using the anti-insulin receptor monoclonal antibody 83-7 (28–31). The ( $\alpha\beta_{\text{INS.WT}}$ ), ( $\alpha\beta_{\text{INS.A/K}}$ ), and ( $\alpha\beta_{\text{INS.ACT}}$ ) half-receptors were obtained from cDNA-transfected cell lines expressing the wild-type human insulin receptor (32), the A/K<sub>1018</sub> mutant insulin receptor (24), and a deletion-mutated insulin receptor lacking 43 COOH-terminal amino acid residues (33), respectively. Cell membranes (3 mg/ml) were prepared (27) and treated with alkaline pH plus dithiothreitol as described above for the human placenta membranes followed by partial purification by Bio-Gel A-1.5m gel filtration chromatography or polysilyne-Sepharose affinity chromatography (34).

**Bio-Gel A-1.5m Column Gel Filtration Chromatography.** Isolated ( $\alpha\beta_{\text{IGF.WT}}$ ) and ( $\alpha\beta_{\text{INS.A/K}}$ ) half-receptors (2 pmol/ml) were mixed and incubated in 50 mM Hepes (pH 7.8) containing 100 nM insulin, 100 nM IGF-I, or 100 nM insulin/100 nM IGF-I for 1 hr at 23°C. Samples were then applied to

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Abbreviation: IGF-I, insulin-like growth factor I.  
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Bio-Gel A-1.5m gel filtration columns (1.6 × 50 cm) equilibrated in 50 mM Tris·HCl, pH 7.6/0.1% Triton X-100/150 mM NaCl/0.02% NaN<sub>3</sub> (Bio-Gel buffer) at 4°C. Fractions containing the α<sub>2</sub>β<sub>2</sub> heterotetrameric and αβ heterodimeric insulin and/or IGF-I receptors were identified by <sup>125</sup>I-labeled IGF-I and <sup>125</sup>I-labeled insulin binding.

**Immunoabsorption of Insulin and IGF-I Receptor Complexes.** Bio-Gel A-1.5m gel filtration column fractions containing the α<sub>2</sub>β<sub>2</sub> heterotetrameric or αβ heterodimeric insulin and IGF-I receptors were incubated (1:500 dilution) with Sepharose-coupled anti-insulin receptor monoclonal antibody 83-7 and anti-IGF-I receptor monoclonal antibody αIR-3 for 16 hr at 4°C. The antibody-receptor complexes were collected by centrifugation and the supernatants were assayed for <sup>125</sup>I-labeled insulin and <sup>125</sup>I-labeled IGF-I binding.

**Kinase Assays.** Isolated (αβ)<sub>INS.WT</sub>, (αβ)<sub>INS.ΔCT</sub>, and (αβ)<sub>INS.A/K</sub> half-receptors (0.1–5 pmol/ml) were mixed and incubated in 50 mM Hepes, pH 7.8/100 nM insulin for 1 hr at 22°C. The samples were immunoabsorbed with Sepharose-coupled monoclonal antibody 83-7 and resuspended in 50 mM Hepes, pH 7.8/100 nM insulin/10 mM MnCl<sub>2</sub>/10 mM MgCl<sub>2</sub>. Substrate phosphorylation was initiated by the addition of poly(Glu-Tyr) (2 mg/ml) and [γ-<sup>32</sup>P]ATP (100 μM, 3 μCi/nmol; 1 Ci = 37 GBq), and the reaction was terminated (20 min) by precipitation onto Whatman 3MM filter paper with 10% (wt/vol) trichloroacetic acid. Autophosphorylation was initiated by the addition of [γ-<sup>32</sup>P]ATP (100 μM, 3 μCi/nmol) and terminated (5 min) by addition of 5 mM ATP/5 mM EDTA/100 mM sodium fluoride/10 mM sodium pyrophosphate. Samples were centrifuged at 12,000 × g, resuspended in Laemmli sample buffer (35) containing 300 mM dithiothreitol, heated for 5 min at 100°C, and resolved on 7.5% polyacrylamide gels containing SDS as described (33).

The isolated (αβ)<sub>IGF.WT</sub> and (αβ)<sub>INS.A/K</sub> half-receptors (0.3–3 pmol/ml) were mixed and incubated for 1 hr at 22°C with 100 nM insulin or 100 nM insulin/100 nM IGF-I, followed by the addition of 10 mM MnCl<sub>2</sub>/10 mM MgCl<sub>2</sub>/poly(Glu-Tyr) (2 mg/ml). Substrate phosphorylation was then assayed directly in solution by the addition of [γ-<sup>32</sup>P]ATP (100 μM, 3 μCi/nmol) and terminated as described above.

**RESULTS AND DISCUSSION**

**Heterologous Assembly of (αβ)<sub>INS.A/K</sub> and (αβ)<sub>IGF.WT</sub> Half-Receptors.** To determine whether mutant/wild-type hybrid receptors could be formed *in vitro*, the highly related but immunologically distinct (αβ)<sub>IGF.WT</sub> and (αβ)<sub>INS.A/K</sub> half-receptor species were examined for heterologous assembly into (αβ)<sub>IGF.WT</sub>/(αβ)<sub>INS.A/K</sub> complexes (Fig. 1). Equal amounts of (αβ)<sub>IGF.WT</sub> and (αβ)<sub>INS.A/K</sub> were incubated with insulin alone, IGF-I alone, or insulin/IGF-I and then subjected to Bio-Gel A-1.5m gel filtration chromatography to determine the receptor association state. Insulin treatment of the mixed αβ half-receptors resulted in a characteristic mobility shift of insulin binding activity (Fig. 1A), consistent with an insulin-induced *in vitro* assembly of α<sub>2</sub>β<sub>2</sub> heterotetrameric complexes (12). The formation of an (αβ)<sub>INS.A/K</sub>/(αβ)<sub>INS.A/K</sub> heterotetrameric receptor was specific for insulin, since IGF-I treatment had no effect on the association state of the (αβ)<sub>INS.A/K</sub> half-receptor. Similarly, IGF-I incubation of the mixed half-receptors resulted in a specific mobility shift of IGF-I binding activity, consistent with the formation of an (αβ)<sub>IGF.WT</sub>/(αβ)<sub>IGF.WT</sub> heterotetrameric IGF-I receptor complex, whereas insulin was without effect (Fig. 1B). The simultaneous treatment of the mixed half-receptors with a combination of insulin plus IGF-I resulted in a mobility shift of both insulin and IGF-I binding activity to

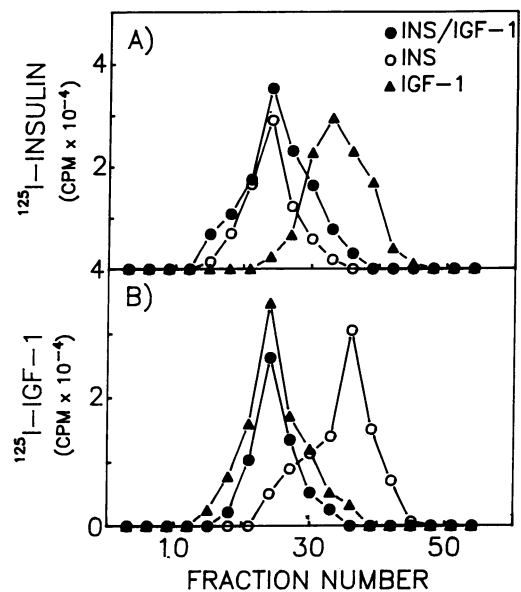


FIG. 1. Ligand-dependent association of (αβ)<sub>IGF.WT</sub> and (αβ)<sub>INS.A/K</sub> heterodimeric half-receptors into an α<sub>2</sub>β<sub>2</sub> heterotetrameric state. (αβ)<sub>IGF.WT</sub> half-receptor and the (αβ)<sub>INS.A/K</sub> half-receptor were obtained from human placenta membranes and transfected Rat-1 fibroblast membranes. Equal amounts of (αβ)<sub>IGF.WT</sub> and (αβ)<sub>INS.A/K</sub> half-receptors (2 pmol/ml) were mixed and incubated with 100 nM insulin (○), 100 nM IGF-I (▲), or 100 nM insulin/100 nM IGF-I (●) for 1 hr at 23°C. The samples were then applied to Bio-Gel A-1.5m gel filtration columns (1.6 × 50 cm) equilibrated in 50 mM Tris·HCl, pH 7.6/150 mM NaCl/0.1% Triton X-100. Fractions (0.45 ml) were collected after voiding 20 ml and assayed (200 μl) for <sup>125</sup>I-labeled insulin (<sup>125</sup>I-insulin) (A) and <sup>125</sup>I-labeled IGF-I (<sup>125</sup>I-IGF-I) binding (B). In the absence of ligand treatment, the mixed αβ heterodimeric half-receptors displayed peak insulin and IGF-I binding at fractions 33–36 (data not shown).

the expected position of an α<sub>2</sub>β<sub>2</sub> heterotetrameric receptor (Fig. 1).

Although these data demonstrate that both the (αβ)<sub>INS.A/K</sub> and (αβ)<sub>IGF.WT</sub> half-receptors associate, in a ligand-specific manner, to a heterotetrameric state, such association could have occurred in either a homologous or heterologous fashion. To determine the degree of insulin/IGF-I hybrid receptor formation, the peak fractions in Fig. 1 were immunoabsorbed with the anti-insulin receptor specific monoclonal antibody 83-7 and the anti-IGF-I receptor monoclonal antibody αIR-3. In the absence of ligand, the mixed αβ heterodimeric half-receptors displayed absolute antibody specificity without any detectable cross-reactivity (e.g., 98–100% of binding precipitated only by the cognizant monoclonal antibody) (Table 1). Antibody specificity (90–100%) was also maintained subse-

Table 1. Insulin and IGF-I induced heterologous assembly of (αβ)<sub>IGF.WT</sub> and (αβ)<sub>INS.A/K</sub> half-receptors into an immunological cross-reactive α<sub>2</sub>β<sub>2</sub> heterotetrameric hybrid receptor complex

Treatment	Receptor species	<sup>125</sup> I-labeled insulin precipitated, %		<sup>125</sup> I-labeled IGF-I precipitated, %	
		83-7	αIR-3	83-7	αIR-3
Untreated	αβ	98	0	1	100
Insulin	α <sub>2</sub> β <sub>2</sub> /αβ	95	3	10	100
IGF-I	α <sub>2</sub> β <sub>2</sub> /αβ	93	0	9	100
Insulin/IGF-I	α <sub>2</sub> β <sub>2</sub>	95	55	54	100

Bio-Gel A-1.5m gel filtration column fractions containing the α<sub>2</sub>β<sub>2</sub> heterotetrameric or αβ heterodimeric insulin and IGF-I receptors (Fig. 1) were immunoabsorbed with 83-7 or αIR-3 followed by <sup>125</sup>I-labeled insulin and <sup>125</sup>I-labeled IGF-I binding.

quent to separate insulin or IGF-I treatment of the mixed  $\alpha\beta$  heterodimeric half-receptors. However, incubation of the mixed  $\alpha\beta$  heterodimeric half-receptors with the combination of insulin plus IGF-I resulted in  $\alpha_2\beta_2$  heterotetrameric receptors that displayed partial (54–55%) cross-reactivity of ligand binding precipitation by both 83-7 and  $\alpha\text{IR-3}$  (Table 1). Thus, incubation of approximately equal amounts of  $(\alpha\beta)_{\text{IGF.WT}}$  and  $(\alpha\beta)_{\text{INS.A/K}}$  half-receptors with a combination of insulin plus IGF-I resulted in the random formation of  $\alpha_2\beta_2$  heterotetrameric hybrid receptors. This is in accord with our previous observations of random heterologous association between the wild-type insulin and IGF-I half-receptors (18).

**Inhibition of  $(\alpha\beta)_{\text{IGF.WT}}$  Kinase Activity by  $(\alpha\beta)_{\text{INS.A/K}}$  Half-Receptor.** We next examined the effect of hybrid formation between the  $(\alpha\beta)_{\text{INS.A/K}}$  and  $(\alpha\beta)_{\text{IGF.WT}}$  half-receptors on IGF-I-stimulated substrate protein kinase activity (Fig. 2). A fixed concentration of  $(\alpha\beta)_{\text{IGF.WT}}$  was incubated with various amounts of  $(\alpha\beta)_{\text{INS.A/K}}$  prior to the simultaneous addition of insulin plus IGF-I. The addition of increasing amounts of  $(\alpha\beta)_{\text{INS.A/K}}$  resulted in a progressive decrease in IGF-I-stimulated substrate kinase activity (Fig. 2A), which directly correlated ( $r^2 = 0.91$ ) with an increased *in vitro* assembly of  $(\alpha\beta)_{\text{IGF.WT}}/(\alpha\beta)_{\text{INS.A/K}}$  heterotetrameric hybrid receptors. Kinase inactivation did not simply result from heterologous hybrid formation between the insulin and IGF-I  $\alpha\beta$  half-receptors *per se*, since  $(\alpha\beta)_{\text{INS.WT}}/(\alpha\beta)_{\text{IGF.WT}}$  hybrid receptor complexes displayed ligand-stimulated kinase activity in response to both insulin and IGF-I (data not shown). In addition, the  $(\alpha\beta)_{\text{IGF.WT}}$  displayed a linear 4.5-fold increase in substrate protein kinase activity as the concentration of  $(\alpha\beta)_{\text{IGF.WT}}$  was increased 6-fold in the presence of IGF-I alone (data not shown) or insulin plus IGF-I (Fig. 2B).

In these experiments it was necessary to drive the formation of the insulin/IGF-I hybrid receptors by a combination of insulin plus IGF-I (Fig. 1). To confirm the ligand specificity of substrate kinase activation of the  $(\alpha\beta)_{\text{IGF.WT}}$  half-receptor,

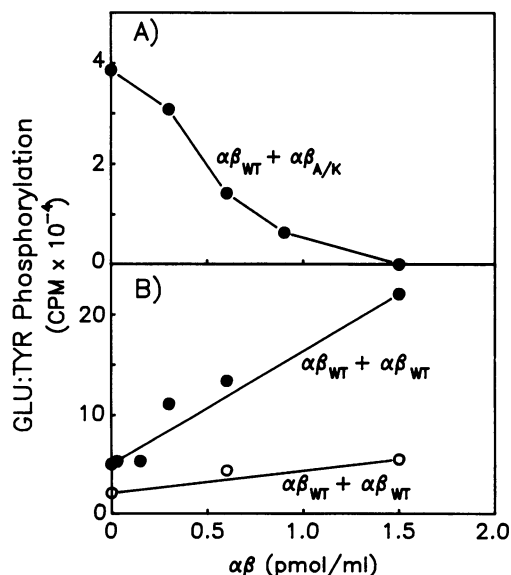


FIG. 2. Transdominant inhibition of  $(\alpha\beta)_{\text{IGF.WT}}$  half-receptor substrate kinase activity by *in vitro* assembly with the  $(\alpha\beta)_{\text{INS.A/K}}$  half-receptor. (A) A fixed amount of  $(\alpha\beta)_{\text{IGF.WT}}$  half-receptor (0.3 pmol/ml) was mixed with increasing relative amounts of  $(\alpha\beta)_{\text{INS.A/K}}$  half-receptor and incubated with 100 nM insulin/100 nM IGF-I for 1 hr at 22°C and substrate phosphorylation was determined. (B) The  $(\alpha\beta)_{\text{IGF.WT}}$  half-receptor (0.3 pmol/ml) was mixed with increasing relative amounts of homologous  $(\alpha\beta)_{\text{IGF.WT}}$  half-receptors in the presence of 100 nM insulin (○) or 100 nM insulin/100 nM IGF-I (●) for 1 hr at 22°C. Poly(Glu-Tyr) substrate phosphorylation was then determined as described in A.

a comparison between insulin and IGF-I stimulation was determined (Fig. 2B). As observed in Fig. 1, insulin did not induce the self-association of the  $(\alpha\beta)_{\text{IGF.WT}}$  into an  $\alpha_2\beta_2$  heterotetrameric state and was relatively ineffective in activating the IGF-I receptor kinase (Fig. 2B). In the presence of IGF-I alone or insulin plus IGF-I, the linear relationship between  $\alpha\beta$  half-receptor concentration and kinase activity reflected the complete formation of  $\alpha_2\beta_2$  IGF-I heterotetrameric receptor complexes under these conditions (27, 29).

**Inhibition of  $(\alpha\beta)_{\text{INS.WT}}$  Kinase Activity by the  $(\alpha\beta)_{\text{INS.A/K}}$  Half-Receptor.** In a similar paradigm, we examined the effect of the  $(\alpha\beta)_{\text{INS.A/K}}$  half-receptor on insulin-stimulated substrate kinase activity by  $(\alpha\beta)_{\text{INS.WT}}$ . As observed for  $(\alpha\beta)_{\text{IGF.WT}}$  (Fig. 2), the addition of increasing amounts of  $(\alpha\beta)_{\text{INS.A/K}}$  to a fixed concentration of  $(\alpha\beta)_{\text{INS.WT}}$  resulted in a dose-dependent inhibition of  $(\alpha\beta)_{\text{INS.WT}}$  substrate protein tyrosine kinase activity (Fig. 3A). Half-maximal inhibition of the  $(\alpha\beta)_{\text{INS.WT}}$  kinase activity occurred at approximately equal molar amounts of  $(\alpha\beta)_{\text{INS.A/K}}$  whereas maximal inhibition required a 10-fold excess. As previously reported (23–25), the kinase-defective  $(\alpha\beta)_{\text{INS.A/K}}$  was substrate-kinase-inactive (Fig. 3B). In contrast, the isolated  $(\alpha\beta)_{\text{INS.WT}}$  displayed a linear 6-fold increase in insulin-stimulated substrate phosphorylation over an 8-fold  $(\alpha\beta)_{\text{INS.WT}}$  concentration range (Fig. 3B). In addition, we have observed that  $(\alpha\beta)_{\text{INS.A/K}}$  inhibition of substrate kinase activity is specific for the  $(\alpha\beta)_{\text{INS.WT}}$  half-receptor, since addition of  $(\alpha\beta)_{\text{INS.A/K}}$  to  $\alpha_2\beta_2$  wild-type insulin receptors had no effect on substrate kinase activity (data not shown). Furthermore, *in vitro* assembly of the  $(\alpha\beta)_{\text{INS.A/K}}$  half-receptor with prephosphorylated and autoactivated  $(\alpha\beta)_{\text{INS.WT}}$  half-receptors (10) did not result in diminished substrate kinase activity (data not shown). These data demonstrate that the transdominant inhibition of  $(\alpha\beta)_{\text{INS.WT}}$  substrate kinase activity by

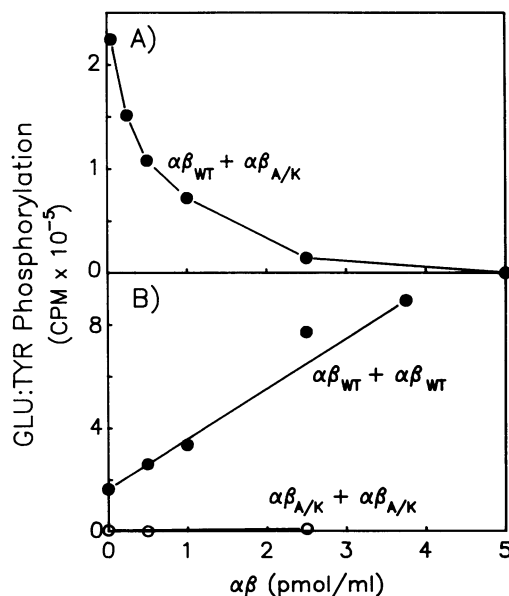


FIG. 3.  $(\alpha\beta)_{\text{INS.A/K}}$  inhibition of insulin-stimulated protein kinase activity of the  $(\alpha\beta)_{\text{INS.WT}}$  half-receptor. (A) A fixed amount of  $(\alpha\beta)_{\text{INS.WT}}$  half-receptor (0.5 pmol/ml) was mixed with increasing amounts of  $(\alpha\beta)_{\text{INS.A/K}}$  half-receptor prior to the addition of 100 nM insulin for 1 hr at 22°C. The samples were immunoabsorbed to anti-insulin receptor monoclonal antibody 83-7 and substrate phosphorylation was initiated. (B)  $(\alpha\beta)_{\text{INS.WT}}$  (●) and  $(\alpha\beta)_{\text{INS.A/K}}$  (○) insulin half-receptors (0.5 pmol/ml) were mixed with increasing amounts of homologous half-receptor in the presence of 100 nM insulin for 1 hr at 22°C. Samples were then immunoabsorbed to monoclonal antibody 83-7 and assayed for poly(Glu-Tyr) substrate phosphorylation as described in A.

$(\alpha\beta)_{\text{INS.A/K}}$  results from impaired insulin-signaling within a wild-type/mutant hybrid receptor complex.

**Autophosphorylation of Wild-Type/Mutant Hybrid Receptor Complexes.** To further investigate the insulin-signaling defect in the wild-type/mutant hybrid receptor, insulin-stimulated  $\beta$ -subunit autophosphorylation was examined (Fig. 4). In contrast to the complete inhibition of substrate kinase activity (Fig. 3), the  $(\alpha\beta)_{\text{INS.WT}}/(\alpha\beta)_{\text{INS.A/K}}$  hybrid receptor (Fig. 4A, lane 2) displayed  $\beta$ -subunit autophosphorylation that was essentially identical compared to the  $(\alpha\beta)_{\text{INS.WT}}/(\alpha\beta)_{\text{INS.WT}}$  holoreceptor complex (Fig. 4A, lane 1). As expected, the *in vitro*-assembled  $(\alpha\beta)_{\text{INS.A/K}}/(\alpha\beta)_{\text{INS.A/K}}$  holoreceptors alone were completely devoid of insulin-stimulated  $\beta$ -subunit autophosphorylation (Fig. 4A, lane 3).

One possible explanation for apparently normal  $\beta$ -subunit autophosphorylation but defective substrate kinase activity in the  $(\alpha\beta)_{\text{INS.WT}}/(\alpha\beta)_{\text{INS.A/K}}$  hybrid receptor would be an altered intramolecular autophosphorylation cascade. To address this issue, autophosphorylation was reexamined in hybrid receptors formed from the heterologous assembly of the  $(\alpha\beta)_{\text{INS.A/K}}$  half-receptor with the  $\beta$ -subunit C-terminal-truncated  $\alpha\beta$  half-receptor,  $(\alpha\beta)_{\text{INS.}\Delta\text{CT}}$ . Insulin-stimulated autophosphorylation of the  $(\alpha\beta)_{\text{INS.}\Delta\text{CT}}/(\alpha\beta)_{\text{INS.A/K}}$  hybrid receptor complex demonstrated the specific labeling of the  $M_r$  95,000  $(\alpha\beta)_{\text{INS.A/K}}$   $\beta$  subunit, without significant autophosphorylation of the truncated  $M_r$  90,000  $(\alpha\beta)_{\text{INS.}\Delta\text{CT}}$   $\beta$  subunit (Fig. 4B, lane 2). As reported (33), autophosphorylation of the  $(\alpha\beta)_{\text{INS.}\Delta\text{CT}}/(\alpha\beta)_{\text{INS.}\Delta\text{CT}}$  receptors exclusively identified the truncated  $M_r$  90,000  $\beta$  subunit species (Fig. 4B, lane 1). These data directly demonstrate an intramolecular trans-phosphorylation of the  $(\alpha\beta)_{\text{INS.A/K}}$   $\beta$  subunit by the kinase-active  $(\alpha\beta)_{\text{INS.}\Delta\text{CT}}$   $\beta$  subunit within the  $\alpha_2\beta_2$  heterotetrameric  $(\alpha\beta)_{\text{INS.}\Delta\text{CT}}/(\alpha\beta)_{\text{INS.A/K}}$  hybrid receptor complex.

In summary, we have identified a transdominant inhibition of ligand-stimulated substrate kinase activity in hybrid insulin

and IGF-I holoreceptors composed of  $\alpha\beta$  wild-type and  $\alpha\beta$  kinase-defective half-receptors. However, these hybrid receptors display essentially normal  $\beta$ -subunit autophosphorylation activity that occurs by an intramolecular trans-phosphorylation mechanism. Thus, insulin binding must necessarily stimulate the kinase activity of one of the  $\alpha\beta$  half-receptors that subsequently utilizes the other  $\alpha\beta$  half-receptor as a phosphotyrosine acceptor substrate. Since autophosphorylation has been established (36–39) to result in the activation of substrate kinase activity, we hypothesize that the presence of a kinase-defective half-receptor within an  $\alpha_2\beta_2$  hybrid receptor complex results in premature termination of the activating signal by preventing a secondary back phosphorylation of the wild-type half-receptor species.

We thank Dr. Steve Jacobs for providing the IGF-I receptor-specific monoclonal antibody  $\alpha\text{IR-3}$ , Dr. Jonathon Whittaker for providing the NIH 3T3 HIR3.5 cell line, and Corrine Conner for excellent technical assistance. This research was supported by grants from the National Institutes of Health, The March of Dimes Birth Defects Foundation, Wellcome Trust, and Serono Diagnostics.

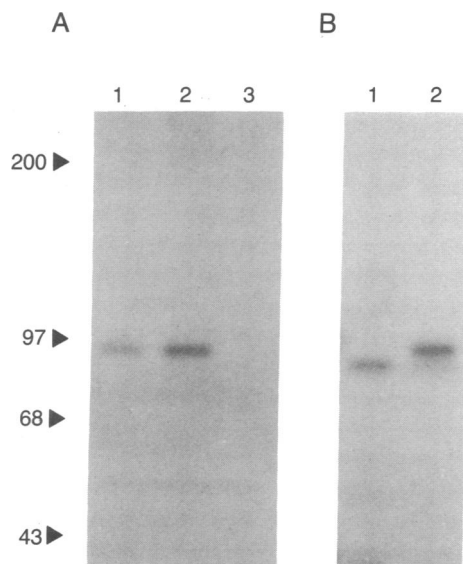


FIG. 4. Insulin-stimulated autophosphorylation of wild-type/mutant hybrid receptor complexes. (A) The  $(\alpha\beta)_{\text{INS.WT}}$  (0.2 pmol/ml; lane 1),  $(\alpha\beta)_{\text{INS.WT}}$  plus  $(\alpha\beta)_{\text{INS.A/K}}$  (0.2 pmol/ml + 1.0 pmol/ml; lane 2), and  $(\alpha\beta)_{\text{INS.A/K}}$  (1.0 pmol/ml; lane 3) half-receptors were incubated with 100 nM insulin for 1 hr at 22°C, then immunoabsorbed with the anti-insulin receptor monoclonal antibody 83-7, and autophosphorylated. (B) The  $(\alpha\beta)_{\text{INS.}\Delta\text{CT}}$  (0.1 pmol/ml; lane 1) and  $(\alpha\beta)_{\text{INS.}\Delta\text{CT}}$  plus  $(\alpha\beta)_{\text{INS.A/K}}$  (0.1 pmol/ml + 0.6 pmol/ml; lane 2) half-receptors were incubated with 100 nM insulin for 1 hr at 22°C, then immunoabsorbed with the monoclonal antibody 83-7, and autophosphorylated as described in A. Positions of molecular weight markers are indicated ( $\times 10^{-3}$ ).

1. Rechler, M. M. & Nissley, S. P. (1985) *Annu. Rev. Physiol.* **47**, 425–442.
2. Goldfine, I. D. (1987) *Endocr. Rev.* **8**, 235–255.
3. Van Obberghen, E., Kasuga, M., LeCam, M., Hedo, J. A., Itin, A. & Harrison, L. C. (1981) *Proc. Natl. Acad. Sci. USA* **78**, 1052–1056.
4. Forsayeth, J., Maddux, B. & Goldfine, I. D. (1986) *Diabetes* **35**, 837–846.
5. Hedo, J. A., Kasuga, M., Van Obberghen, E., Roth, J. & Kahn, C. R. (1981) *Proc. Natl. Acad. Sci. USA* **78**, 4791–4795.
6. Jacobs, S., Kull, F. C. & Cuatrecasas, P. (1983) *Proc. Natl. Acad. Sci. USA* **80**, 1228–1231.
7. Deutsch, P. J., Wan, C. F., Rosen, O. M. & Rubin, C. S. (1983) *Proc. Natl. Acad. Sci. USA* **80**, 133–136.
8. Ronnett, G. V., Knutson, P., Kohanski, R. A., Simpson, T. L. & Lane, M. D. (1984) *J. Biol. Chem.* **259**, 4566–4575.
9. Hedo, J. A., Collier, E. & Watkinson, A. (1987) *J. Biol. Chem.* **262**, 954–957.
10. Boni-Schnetzler, M., Rubin, J. B. & Pilch, P. F. (1986) *J. Biol. Chem.* **261**, 15281–15287.
11. Boni-Schnetzler, M., Kaligian, A., DelVecchio, R. & Pilch, P. F. (1988) *J. Biol. Chem.* **263**, 6822–6828.
12. Morrison, B. D., Swanson, M. L., Sweet, L. J. & Pessin, J. E. (1988) *J. Biol. Chem.* **263**, 7806–7813.
13. Sweet, L. J., Morrison, B. D. & Pessin, J. E. (1987) *J. Biol. Chem.* **262**, 6939–6942.
14. Wilden, P. A., Morrison, B. D. & Pessin, J. E. (1989) *Biochemistry* **28**, 785–792.
15. O'Hare, T. & Pilch, P. F. (1988) *Biochemistry* **27**, 5693–5700.
16. Moxham, C. D., Duronio, V. & Jacobs, S. (1989) *J. Biol. Chem.* **264**, 13238–13244.
17. Soos, M. A. & Siddle, K. (1989) *Biochem. J.* **263**, 553–563.
18. Treadway, J. L., Morrison, B. D., Goldfine, I. D. & Pessin, J. E. (1989) *J. Biol. Chem.* **264**, 21450–21453.
19. Grunberger, G., Zick, Y. & Gorden, P. (1984) *Science* **223**, 932–934.
20. Moller, D. E. & Flier, J. S. (1988) *N. Engl. J. Med.* **319**, 1526–1529.
21. Odawara, M., Kadowaki, T., Yamamoto, R., Shibasaki, Y., Tobe, K., Accili, D., Bevins, C., Mikami, Y., Matura, N., Akanuma, Y., Takaku, F., Taylor, S. I. & Kasuga, M. (1989) *Science* **245**, 66–68.
22. Taira, M., Taira, M., Hasimoto, N., Shimada, F., Suzuki, Y., Kanatsuka, A., Nakamura, F., Ebina, Y., Tatibana, M., Makino, H. & Yoshida, S. (1989) *Science* **245**, 63–66.
23. Chou, C. K., Dull, T. J., Russell, D. S., Gherzi, R., Lebowitz, D., Ullrich, A. & Rosen, O. M. (1987) *J. Biol. Chem.* **262**, 1842–1847.
24. McClain, D. A., Maegawa, H., Lee, J., Dull, T. J., Ullrich, A. & Olefsky, J. (1987) *J. Biol. Chem.* **262**, 14663–14671.
25. Russell, D. S., Gherzi, R., Johnson, E. L., Chou, C. K. & Rosen, O. M. (1987) *J. Biol. Chem.* **262**, 11833–11840.

26. McClain, D. A., Maegawa, H., Thies, R. S. & Olefsky, J. (1990) *J. Biol. Chem.* **265**, 1678–1682.
27. Feltz, S. M., Swanson, M. L., Wemmie, J. A. & Pessin, J. E. (1988) *Biochemistry* **27**, 3234–3242.
28. Kull, F. C., Jacobs, S., Su, Y. F., Svoboda, M. E., Van Wyk, J. J. & Cuatrecasas, P. (1983) *J. Biol. Chem.* **258**, 6561–6566.
29. Wilden, P. A., Treadway, J. L., Morrison, B. D. & Pessin, J. E. (1989) *Biochemistry* **28**, 785–792.
30. Soos, M. A., Siddle, K., Baron, M. D., Heward, J. M., Luzio, J. F., Bellatin, J. & Lennox, E. S. (1986) *Biochem. J.* **235**, 199–208.
31. Soos, M. A., O'Brien, R. M., Brindle, N. P. J., Stigter, J. M., Okamoto, A. K., Whittaker, J. & Siddle, K. (1989) *Proc. Natl. Acad. Sci. USA* **86**, 5217–5221.
32. Whittaker, J., Okamoto, A. K., Thys, R., Bell, G. I., Steiner, D. F. & Hofmann, C. A. (1987) *Proc. Natl. Acad. Sci. USA* **84**, 5237–5241.
33. McClain, D. A., Maegawa, H., Levy, J., Huecksteadt, T., Dull, T. J., Lee, J., Ullrich, A. & Olefsky, J. M. (1988) *J. Biol. Chem.* **263**, 8904–8911.
34. Morrison, B. D., Feltz, S. M. & Pessin, J. E. (1989) *J. Biol. Chem.* **264**, 9994–10001.
35. Laemmli, U. K. (1970) *Nature (London)* **227**, 680–685.
36. Rosen, O. M., Herrera, R., Olowe, Y., Petruzzelli, L. M. & Cobb, M. H. (1983) *Proc. Natl. Acad. Sci. USA* **80**, 3237–3240.
37. Yu, K.-T. & Czech, M. P. (1984) *J. Biol. Chem.* **259**, 5277–5286.
38. Klein, H. H., Freidenberg, G. R., Kladdde, M. & Olefsky, J. M. (1986) *J. Biol. Chem.* **261**, 4691–4697.
39. Yu, K.-T. & Czech, M. P. (1986) *J. Biol. Chem.* **261**, 4715–4722.