

## An Interleukin 5 Mutant Distinguishes between Two Functional Responses in Human Eosinophils

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### Summary

Interleukin 5 (IL-5) is the key cytokine involved in regulating the production and many of the specialized functions of mature eosinophils including priming, adhesion, and survival. We have generated a point mutant of human IL-5, IL-5 (E12K), which is devoid of agonist activity in both a TF-1 cell proliferation assay and a human eosinophil adhesion assay. However, IL-5 (E12K) is a potent and specific antagonist of both these IL-5-dependent functional responses. In both receptor binding and cross-linking studies the wild-type and IL-5 (E12K) mutant exhibit virtually identical properties. This mutant protein was unable to stimulate tyrosine phosphorylation in human eosinophils, and blocked the phosphorylation stimulated by IL-5. In contrast, IL-5 (E12K) is a full agonist in a human eosinophil survival assay, although with reduced potency compared to the wild-type protein. This IL-5 mutant enables us to clearly distinguish between two IL-5-dependent functional responses and reveals distinct mechanisms of receptor/cellular activation.

IL-5 is a cytokine secreted predominantly by T cells, but also by mast cells and eosinophils (1–3), that has multiple effects on cells of the eosinophil and basophil lineage. IL-5 induces the differentiation and expansion of eosinophil precursors in the bone marrow (4) and plays a key role in regulating many of the specialized functions of mature, terminally differentiated eosinophils, including adhesion (5), priming of both degranulation and chemotaxis (6, 7), and the promotion of cell survival (8).

While eosinophils appear to play a protective role during the host defense response to parasitic infection, accumulating clinical and experimental evidence has also implicated the eosinophil in the pathophysiology of a number of allergic diseases such as bronchial asthma, allergic rhinitis, and atopic dermatitis (9–11). Administration of neutralizing anti-IL-5 monoclonal antibodies in mouse, guinea pig, or primate models of allergic asthma inhibits the development of airway eosinophilia and bronchial hyperreactivity (12–16). This highlights the eosinophil, and more specifically IL-5, as an attractive target for therapeutic intervention in allergic diseases such as asthma.

The effects of IL-5 are mediated through a heterodimeric receptor complex composed of a specific ligand binding  $\alpha$  subunit and a  $\beta$  subunit that is shared by the specific receptor  $\alpha$  chains for IL-3 and GM-CSF (17). Although this common  $\beta$  chain ( $\beta_c$ ) alone has no inherent ligand

binding capacity, it confers high-affinity ligand binding to the receptor  $\alpha/\beta$  chain complex. This phenomenon of increased binding affinity in the presence of the  $\beta_c$  chain has been termed affinity conversion, and is most profound for IL-3 (1,000-fold), with smaller effects on GM-CSF binding (20–100-fold) and IL-5 binding (2–4-fold) (18–20). The roles of the respective  $\alpha$  and  $\beta_c$  chains in signal transduction are unclear. Analysis of chimeric receptors composed of the extracellular domain of the IL-5 receptor  $\alpha$  chain and the intracellular domain of the  $\beta_c$  chain suggests that ligand-induced dimerization of the cytoplasmic domains of  $\beta_c$  may be sufficient for receptor activation and the transduction of a proliferative signal (21). In contrast, deletion and mutational analysis has defined specific regions within the cytoplasmic domains of both the IL-5 receptor  $\alpha$  and  $\beta_c$  chains that are essential for the coupling of IL-5 receptors to a proliferative signal (21–24). Thus, although  $\beta_c$  is indispensable for the signals leading to proliferation, it is possible that the receptor  $\alpha$  chain may contribute to cytokine signal transduction either directly, or by promoting receptor oligomerization. Furthermore, the relative contributions of the two receptor chains to signal transduction leading to functional responses other than proliferation are uncharacterized. Although the precise molecular mechanisms by which the active IL-5 receptor complex directs these diverse cellular responses in the eosinophil are also

poorly understood, one of the earliest measurable signaling events is the phosphorylation of a variety of cellular proteins on tyrosine residues (25).

IL-5 is a disulphide-linked homodimeric glycoprotein with 115 amino acids in each chain (26). Elucidation of the IL-5 crystal structure revealed a novel two domain structure in which each domain adopts a four  $\alpha$ -helix bundle structure similar to the cytokine fold in IL-2, IL-4, growth hormone, and GM-CSF (27). Among these related structures, IL-5 is unique in that each bundle is composed of three helices from one monomer and a fourth helix which is contributed by the second monomer. Recently, the receptor binding sites on IL-5 have been defined by site-directed mutagenesis. Residues E88, R90, and E109, within the carboxy terminus of IL-5, define the IL-5 receptor  $\alpha$  chain binding site (28, 29). In addition, the IL-5 mutant E12A, IL-5 (E12A), exhibits partial agonist activity in a TF-1 cell proliferation assay, suggesting that the highly conserved E12 in the amino-terminal helix defines a point of contact with the receptor  $\beta_c$  chain which is important for  $\beta_c$  chain activation and subsequent signal transduction (28). Furthermore, in a TF-1 proliferation assay the IL-5 mutant E13Q (Due to a single amino acid difference at the amino terminus, IL-5 [E12] as described here is equivalent to IL-5 [E13] as described by Tavernier [29].), IL-5 (E13Q), was completely inactive and exhibited antagonist properties (29).

In this study we have produced an IL-5 mutant containing a charge reversal at position 12, IL-5 (E12K). It has been characterized in a variety of functional and receptor binding assays, and we have used this protein as a tool to reveal distinct mechanisms of receptor/cellular activation in human eosinophils.

## Materials and Methods

**Reagents.** Recombinant IL-3, TNF- $\alpha$ , and GM-CSF were obtained from R&D systems (Abingdon, UK). BS<sup>3</sup> (bis[sulfosuccinimidyl] suberate)<sup>1</sup> cross-linking reagent was from Pierce and Warriner (Chester, UK). Na<sup>125</sup>I for protein iodination, and <sup>125</sup>I-labeled interleukin 5 (<sup>125</sup>I-IL-5) were obtained from Amersham International (Amersham, Buckinghamshire, UK). Anti-phosphotyrosine antibody (4G10) was obtained from TCS Biologicals (Buckingham, UK). All other chemicals were obtained from Sigma (Poole, Dorset, UK).

**Site-directed Mutagenesis and Protein Expression.** Recombinant human IL-5 was expressed in *E. coli* and purified to homogeneity as previously described (30). IL-5 (E12K) was generated by site-directed mutagenesis, expressed in *E. coli* and purified as previously described (28) with the following modifications. Renaturation was carried out by dropwise dilution of the protein in 6 M urea, into 0.1 M ethanolamine, pH 9.8, containing 2.5 M urea, 10% ammonium sulphate, 1 mM reduced glutathione, and 0.1 mM oxidized glutathione to a final protein concentration of 10  $\mu$ g/ml,

<sup>1</sup>Abbreviations used in this paper: BS<sup>3</sup>, bis(sulfosuccinimidyl) suberate; ED<sub>80</sub>, concentration required to give 80% of the maximum biological response; IL-5R-zz, extracellular domain of the IL-5 receptor  $\alpha$  chain, fused with the IgG binding domain of protein A; MOPS, 3-[N-morpholino] propanesulfonic acid; SPA, scintillation proximity assay.

and stirred overnight at 4°C. The solution was dialyzed against three changes of 0.1 M ethanolamine, and concentrated 10-fold by ultrafiltration before gel filtration. Circular dichroism spectral analysis confirmed that the secondary and tertiary structure was identical to wild-type IL-5 (data not shown).

**Receptor Binding Assays.** In competition binding studies IL-5 and IL-5 (E12K) were assayed for their relative ability to bind to recombinant IL-5 receptor  $\alpha$  chain alone, or the receptor  $\alpha/\beta$  chain complex as expressed on TF-1 cells, as previously described (31).

In brief, for low-affinity binding studies the extracellular domain of the IL-5 receptor  $\alpha$  chain, fused with the IgG binding domain of protein A (IL-5R $\alpha$ -ZZ), was incubated with rabbit IgG and anti-rabbit-coated fluoromicrosphere beads (Amersham International, UK) for 2 h at 4°C. The resultant complex was incubated with 100 pM <sup>125</sup>I-IL-5 in the presence of increasing concentrations of unlabeled cytokines as indicated. After incubation for 4 h at room temperature, bound radioligand was measured in a Wallac 1450 Microbeta scintillation counter set up in scintillation proximity assay (SPA) mode. Nonspecific binding was measured in the presence of a 500-fold molar excess of unlabeled IL-5.

For cell-based binding studies TF-1 cells ( $1 \times 10^6$ ) were incubated with 200 pM <sup>125</sup>I-IL-5 for 2 h at room temperature in the presence of increasing concentrations of unlabeled cytokines as indicated. After separation of bound and free ligand by centrifugation through oil (16% paraffin oil, 84% silicone oil; BDH, Dorset, UK) the cell-associated radioligand was quantified in a gamma counter. Nonspecific binding was measured in the presence of a 300-fold molar excess of unlabeled IL-5.

**TF-1 Cell Proliferation Assay.** Cytokine-induced proliferation of the human erythroleukemia cell line TF-1 was measured as previously described (28). In brief, assays were performed in 96-well microtiter plates containing  $5 \times 10^3$  cells/well with the indicated cytokines. After incubation for 60–72 h at 37°C the induction of proliferation was measured using the Cell Titer 96™ nonradioactive cell proliferation assay (Promega, Southampton, UK), according to the manufacturer's instructions.

**Eosinophil Purification.** Human peripheral blood eosinophils were isolated from healthy donors with mildly elevated eosinophil count, by a CD16-negative selection protocol as previously described (32). Eosinophil purity was always >95%.

**Eosinophil Activation Assay.** Cytokine-induced activation of purified eosinophils, as measured by adhesion to immobilized IgG, was assayed as previously described (32). In brief,  $5 \times 10^3$  eosinophils were incubated with the indicated cytokines for 30 min at 37°C, in a 96-well microtiter plate precoated with human IgG. After a washing step, the adherent eosinophils were lysed and the endogenous peroxidase activity measured in a colorimetric assay using *o*-phenylenediamine as a substrate.

**Eosinophil Survival Assay.** Purified eosinophils were suspended at  $10^6$  cells/ml in DMEM containing 10% FCS, 50 U/ml penicillin, and 50  $\mu$ g/ml streptomycin (GIBCO, Paisley, Scotland) in microtiter plates, in the presence or absence of wild-type IL-5 or IL-5 (E12K). After incubation at 37°C for 72 h, cell viability was assessed by trypan blue exclusion, counting a minimum of 200 cells. For antibody blocking experiments, human eosinophils were incubated with the indicated concentrations of cytokines in the presence or absence of 250  $\mu$ g/ml anti-IL-5 neutralizing antibody, TRFK-5 (33). For polymyxin B experiments agonists were preincubated with 500 U/ml polymyxin B for 1 h at 37°C before addition of purified human eosinophils.

**Cross-linking of Radiolabeled IL-5 or IL-5 (E12K) to IL-5 Receptor  $\alpha$  and  $\beta$  Chains Expressed on COS Cells.** For cross-linking studies recombinant human IL-5 and IL-5 (E12K) were iodinated by

a modified chloramine-T method essentially as described previously (28).

COS cells were plated at 10% confluence in 175-cm<sup>2</sup> flasks and grown overnight in DMEM containing 10% FCS, 50 U/ml penicillin, and 50 µg/ml streptomycin. Cells were then transiently transfected with human IL-5 receptor  $\alpha$  and  $\beta_c$  chain expression plasmids (10 µg each) by a calcium phosphate precipitation method, using the Mammalian Transfection Kit (Stratagene, Cambridge, UK), according to the manufacturers instructions. Cells were harvested 48 h after transfection in PBS containing 5 mM EDTA and counted.

Transfected cells ( $5 \times 10^5$ ) were incubated for 2 h at 4°C in 200 µl binding medium (RPMI 1640 containing 0.5% wt/vol bovine serum albumin and 20 mM Hepes, pH 7.4) with 70 nM <sup>125</sup>I-IL-5 or <sup>125</sup>I-IL-5 (E12K) in the presence or absence of a 100-fold molar excess of unlabeled IL-5. Cells were then washed twice, resuspended in 1 ml of ice-cold PBS, and 50 mM BS<sup>3</sup> cross-linker was added to a final concentration of 1 mM. After incubation for 30 min at 4°C, cells were washed twice in PBS and resuspended in 150 µl lysis buffer (50 mM Hepes pH 7.4, containing 2 mM EDTA, 2 mM EGTA, 1 mM 4-[2-aminoethyl] benzenesulfonyl fluoride, 5 µg/ml leupeptin, 5 µg/ml pepstatin, 5 µg/ml antipain, and 1% vol/vol Triton X-100). The cell lysate was incubated for 10 min on ice, centrifuged for 10 min at 12,000 rpm in a microfuge, and the supernatant mixed with 4× SDS sample buffer containing 10% 2-ME. Extracts were analyzed on a 4–12% NuPAGE SDS-polyacrylamide gel (Novex, San Diego, CA) run in MOPS buffer (3-[N-Morpholino] propane-sulfonic acid). After electrophoresis gels were fixed and dried and the radiolabeled bands visualized using a PhosphoImager (Molecular Dynamics, Buckinghamshire, UK).

**Detection of Phosphotyrosine Containing Proteins by Western Blotting.** Human eosinophils ( $5 \times 10^5$ ) were incubated in the presence of indicated concentrations of cytokines for 5 min at 37°C, in PBS containing 0.4% human serum albumin. The cells were then pelleted at 4°C, resuspended in 1× NuPAGE sample buffer containing 2.5% 2-ME and 1 mM sodium orthovanadate and boiled for 5 min. Extracts were analyzed on a 4–12% NuPAGE SDS-polyacrylamide gel run in MOPS buffer and transferred to hybond-ECL membrane (Amersham International, Amersham, UK). Blots were probed with the anti-phosphotyrosine antibody 4G10, and proteins visualized using a horseradish peroxidase-conjugated goat anti-mouse secondary antibody (Sigma, Poole, Dorset, UK) with enhanced chemiluminescence (ECL) detection (Amersham International, UK).

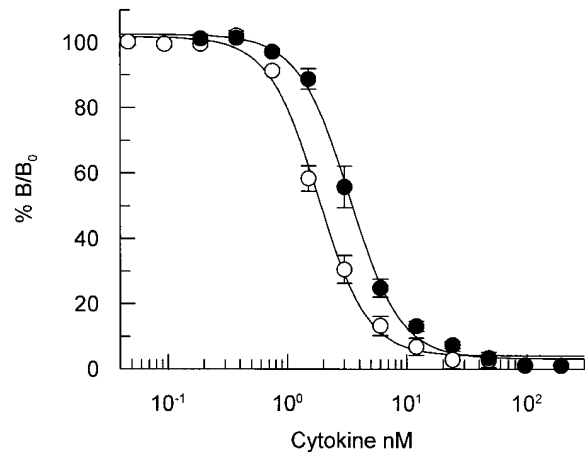
**Data Analysis.** Receptor binding and biological data were analyzed using Graft 3.01 as previously described (28).

## Results

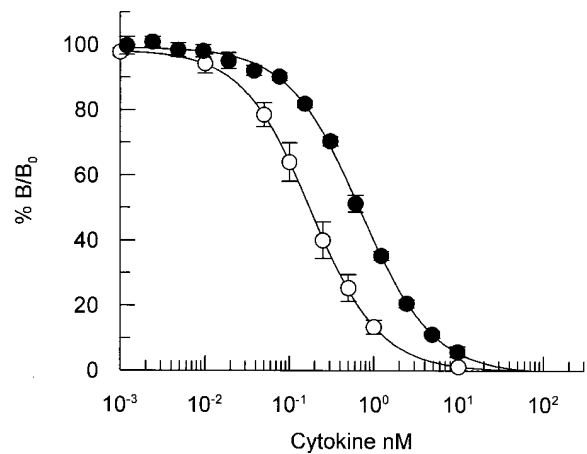
### Relative Receptor Binding Properties of IL-5 and IL-5 (E12K).

Wild-type human IL-5 and a mutant IL-5 containing a charge reversal mutation at position 12, IL-5 (E12K), were expressed in *E. coli* and purified to homogeneity. As a means of comparing the receptor binding properties of wild-type IL-5 and IL-5 (E12K), both proteins were assayed for their relative ability to bind to the IL-5 receptor  $\alpha$  chain alone, or to the high-affinity  $\alpha/\beta$  receptor complex. In competition binding experiments, using the recombinant extracellular domain of the IL-5 receptor  $\alpha$  chain in an SPA assay format, IL-5 (E12K) exhibited a  $1.7 \pm 0.2$ -fold

A

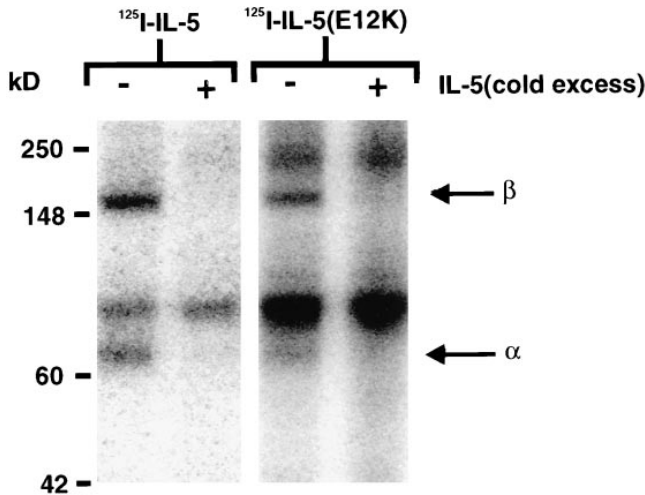


B



**Figure 1.** Comparative binding of wild-type IL-5 and IL-5 (E12K) to recombinant IL-5 receptor  $\alpha$  chain (A) or to the receptor  $\alpha/\beta$  complex on TF-1 cells (B). (A) Recombinant IL-5 receptor  $\alpha$ -chain, immobilized on SPA beads, were incubated with 100 pM <sup>125</sup>I-IL-5 in the presence of increasing concentrations of unlabeled wild-type IL-5 (open circles) or IL-5 (E12K) (closed circles). After incubation for 4 h at room temperature samples were counted in a Wallac 1450 microbeta counter set up in SPA mode. The results show competition as percentage of maximum binding (% B/B<sub>0</sub>). Each value represents the mean  $\pm$  SEM of four independent experiments for wild-type and six independent experiments for IL-5 (E12K). (B) TF-1 cells were incubated at room temperature for 2 h in the presence of 200 pM <sup>125</sup>I-IL-5 and increasing concentrations of unlabeled wild-type (open circles) or IL-5 (E12K) (closed circles). After separation of bound and free ligand the cell-associated radioligand was quantified in a gamma counter. The results show competition as percentage of maximum binding (% B/B<sub>0</sub>). Each value represents the mean  $\pm$  SEM of five independent experiments.

( $n = 6$ )-fold reduction in binding affinity to the receptor  $\alpha$  chain relative to wild-type IL-5 (Fig. 1 A). The interaction with the high-affinity receptor complex was assessed in a cell based competition binding assay using TF-1 cells which express both the IL-5 receptor  $\alpha$  and  $\beta_c$  chains. In this cell based system IL-5 (E12K) exhibited a  $4.5 \pm 1.2$ -fold ( $n = 5$ ) reduction in binding affinity relative to wild-type IL-5 (Fig. 1 B). The shift in relative binding of IL-5 (E12K) and wild-type IL-5 to the high-affinity  $\alpha/\beta$  complex, compared



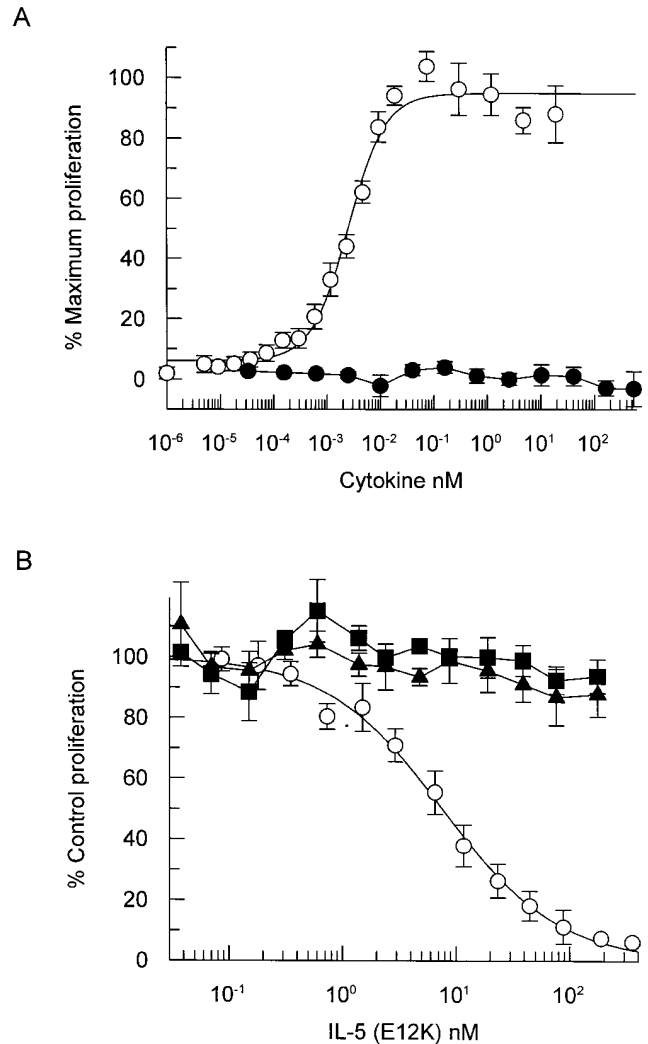
**Figure 2.** Chemical cross-linking of wild-type IL-5 and IL-5 (E12K) to IL-5 receptors. COS cells transfected with human IL-5 receptor  $\alpha$  and  $\beta_c$  chain cDNAs were incubated with  $\sim 70$  nM  $^{125}\text{I}$ -IL-5 or  $^{125}\text{I}$ -IL-5 (E12K) for 2 h at  $4^\circ\text{C}$  in the presence or absence of a 100-fold molar excess of unlabeled IL-5, then cross-linked with 1 mM BS $^3$ . After detergent lysis of cells, soluble extracts were analyzed by SDS-PAGE and radiolabeled bands visualized by a PhosphoImager.

to the  $\alpha$  chain alone, was found to be statistically nonsignificant ( $P = 0.084$ ), as assessed by an analysis of variance technique.

**Chemical Cross-linking of Radiolabeled IL-5 and IL-5 (E12K) to IL-5 Receptors.** To determine whether IL-5 (E12K) is able to interact with the IL-5 receptor  $\beta_c$  chain, we employed receptor cross-linking studies. The chemical cross-linking of  $^{125}\text{I}$ -IL-5 or  $^{125}\text{I}$ -IL-5 (E12K) to COS cells transfected with IL-5 receptor  $\alpha$  and  $\beta_c$  chains revealed identical patterns of cross-linked species of  $\sim 70$  and 150 kD under reducing conditions (Fig. 2), which correspond to the predicted molecular mass of the receptor  $\alpha$  and  $\beta_c$  subunits, respectively, bound to a monomer of IL-5. Furthermore, the appearance of these bands was blocked by addition of a 100-fold molar excess of unlabeled IL-5 during the incubation, confirming that both  $^{125}\text{I}$ -IL-5 and  $^{125}\text{I}$ -IL-5 (E12K) can be specifically cross-linked to both IL-5 receptor  $\alpha$  and  $\beta_c$  chains.

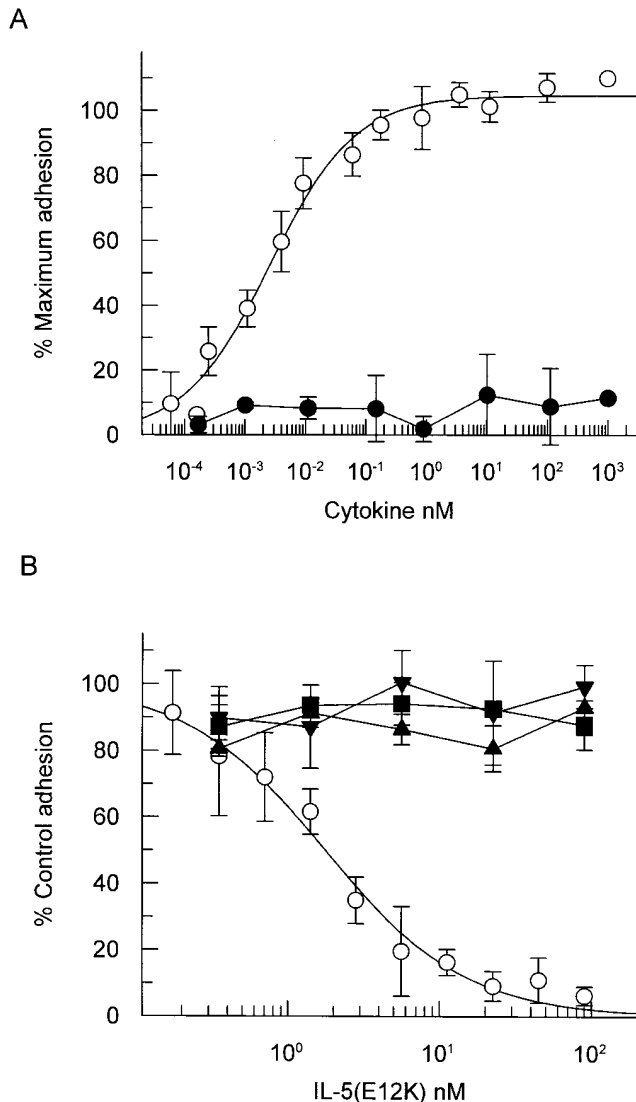
**Biological Properties of IL-5 (E12K) in a TF-1 Proliferation Assay.** Wild-type IL-5 induced the proliferation of the human erythroleukemia cell line TF-1 in a concentration-dependent fashion ( $\text{ED}_{50} = 2.6 \pm 0.6$  pM,  $n = 4$ ), while IL-5 (E12K) was unable to stimulate proliferation even at concentrations up to 400 nM, a concentration 160,000-fold higher than the concentrations of wild-type IL-5 required to elicit half maximal proliferation of TF-1 cells (Fig. 3 A).

Having determined that IL-5 (E12K) was able to bind to IL-5 receptors with almost wild-type affinity, yet was unable to elicit a proliferative response in TF-1 cells, we tested this mutant for its ability to antagonize the effects of IL-5 in a TF-1 proliferation assay. As seen in Fig. 3 B, IL-5 (E12K) antagonized the effect of IL-5 in a concentration-



**Figure 3.** IL-5 (E12K) exhibits no agonist activity (A) and is a specific IL-5 antagonist (B) in a TF-1 cell proliferation assay. (A) TF-1 cells were incubated with increasing concentrations of either wild-type (open circles) or IL-5 (E12K) (closed circles) for 60–72 h and the induction of proliferation was measured using a nonradioactive cell proliferation assay. Each value represents the mean  $\pm$  SEM of four independent experiments. (B) TF-1 cell proliferation was assayed at either 77 pM IL-5 (open circles), 133 pM IL-3 (closed squares) or 14 pM GM-CSF (closed triangles) in the presence of increasing concentrations of IL-5 (E12K). The concentrations of wild-type cytokines represent  $\sim \text{ED}_{80}$ 's (concentration of cytokine required to give 80% of the maximum biological response) for proliferation in our TF-1 cell line. Each value represents the mean  $\pm$  SEM of six independent experiments. 100% values for IL-5-, IL-3-, and GM-CSF-induced proliferation were equivalent to  $A_{550}$  of  $0.68 \pm 0.02$ ,  $0.73 \pm 0.18$ , and  $0.64 \pm 0.04$ , respectively. Unstimulated levels were  $0.03 \pm 0.02$ .

dependent manner. Proliferation of TF-1 cells, in the presence of 77 pM wild-type IL-5, was totally inhibited by 100 nM IL-5 (E12K) and 50% inhibition was achieved at  $10.3 \pm 2.8$  nM ( $n = 6$ ), which represents a 130-fold molar excess of IL-5 (E12K) over wild-type. Furthermore, in the same functional assay IL-5 (E12K) did not significantly inhibit TF-1 proliferation stimulated by IL-3 or GM-CSF (Fig. 3 B). This demonstrates not only that IL-5 (E12K) is a



**Figure 4.** IL-5 (E12K) exhibits no agonist activity (A) and is a specific IL-5 antagonist (B) in a cytokine induced eosinophil activation assay. (A) Human eosinophils were incubated with increasing concentrations of either wild-type (open circles) or IL-5 (E12K) (closed circles) for 30 min at 37°C, in a 96-well microtiter plate precoated with human IgG. After a washing step, the adherent eosinophils were lysed and the endogenous peroxidase activity measured in a colorimetric assay. (B) In antagonist experiments eosinophils were incubated with either 20 pM IL-5 (open circles), 180 pM IL-3 (closed squares), 6 pM GM-CSF (closed triangles), or 1,000 pM TNF- $\alpha$  (inverted triangles) in the presence of increasing concentrations of IL-5 (E12K). The concentrations of wild-type cytokines represent  $\sim$ ED<sub>80</sub>'s in the eosinophil adhesion assay. In both panels each value represents the mean  $\pm$  SEM of three independent experiments using different blood donors. 100% values for IL-5-, IL-3-, GM-CSF-, and TNF- $\alpha$ -induced adhesion were  $0.56 \pm 0.07$ ,  $0.76 \pm 0.05$ ,  $0.41 \pm 0.07$ , and  $0.49 \pm 0.06$ , respectively. Unstimulated levels were  $0.15 \pm 0.05$ .

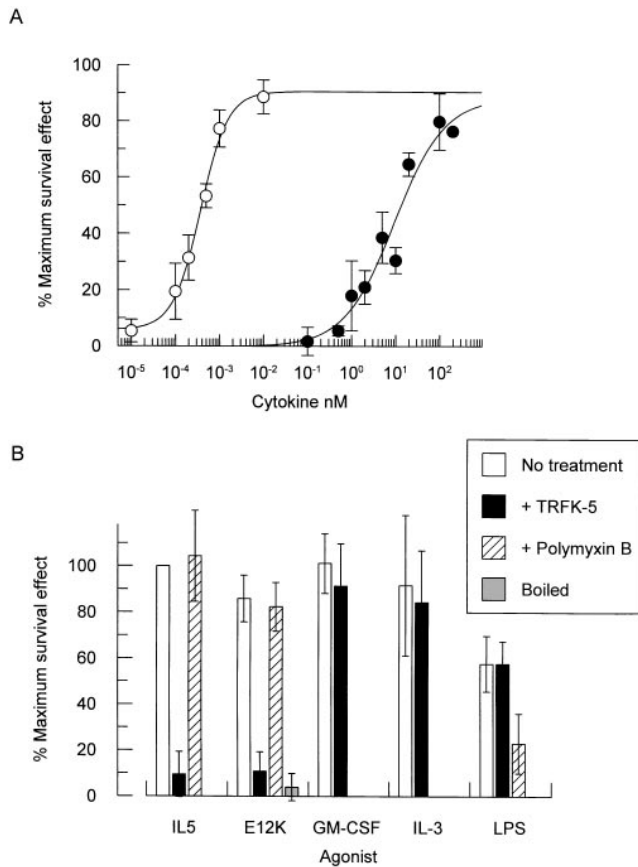
specific IL-5 receptor antagonist, but also excludes the possibility of IL-5 (E12K)-dependent cellular toxicity.

**Biological Properties of IL-5 (E12K) in an Eosinophil Activation Assay.** To confirm and extend our observations to additional IL-5-induced cellular responses, we switched

our studies to human peripheral blood eosinophils, a terminally differentiated cell type in which IL-5 stimulates multiple functional responses. We have established a sensitive and reliable in vitro assay for analysis of cytokine-induced eosinophil activation based on adhesion to immobilized IgG. In this assay, wild-type IL-5 induced a concentration-dependent increase in eosinophil adhesion to IgG (ED<sub>50</sub> =  $1.3 \pm 0.1$  pM,  $n = 3$ ), while the mutant E12K was incapable of inducing eosinophil adhesion even up to concentrations of 1  $\mu$ M, a concentration 400,000-fold higher than the ED<sub>50</sub> for wild-type IL-5 (Fig. 4 A). In the same functional assay IL-5 (E12K) completely antagonized the effects of IL-5 in a concentration-dependent fashion. Eosinophil adhesion in the presence of 20 pM wild-type IL-5 was completely inhibited by 50 nM IL-5 (E12K) with 50% inhibition achieved at  $1.9 \pm 0.1$  nM ( $n = 3$ ), which represents a 95-fold molar excess of IL-5 (E12K) over wild-type (Fig. 4 B). In addition, the inhibitory effect of IL-5 (E12K) was specific for IL-5 induced adhesion with no significant inhibition of IL-3, GM-CSF, or TNF- $\alpha$  across the range of IL-5 (E12K) concentrations found to inhibit IL-5 (Fig. 4 B).

**Biological Properties of IL-5 (E12K) in an Eosinophil Survival Assay.** Mature eosinophils that have been separated from peripheral blood do not survive more than 4 d in vitro without the addition of cytokines. The eosinophilopoietic cytokines IL-5, IL-3, and GM-CSF have all been reported to promote eosinophil survival, and so maintain cell viability (8, 34, 35). We therefore assessed the relative ability of wild-type IL-5 and IL-5 (E12K) to promote eosinophil survival. Purified human peripheral blood eosinophils were incubated in the presence of increasing concentrations of wild-type IL-5 or IL-5 (E12K) and eosinophil viability measured by trypan blue exclusion, after a period of 72 h. Both IL-5 and IL-5 (E12K) were able to promote eosinophil survival in a concentration-dependent manner with ED<sub>50</sub> values of  $0.4 \pm 0.1$  pM and  $20.6 \pm 8.7$  nM, respectively ( $n = 5$ ) (Fig. 5 A). Although IL-5 (E12K) promoted eosinophil survival with a 50,000-fold reduction in biological potency with respect to wild-type IL-5, it was still a full agonist capable of eliciting a maximal biological response not significantly different from wild-type IL-5.

One trivial explanation for the agonist activity of IL-5 (E12K) in the eosinophil survival assay is the contamination of this *E. coli* expressed cytokine with bacterial LPS, which has been reported to promote eosinophil survival through the autocrine production of GM-CSF (36). To exclude this possibility a series of control experiments were performed (Fig. 5 B). First, the agonist activity of IL-5 (E12K) was completely abolished after boiling for 15 min. Furthermore, pretreatment with polymyxin B, an LPS inhibitor (36), blocked LPS- but not IL-5 or IL-5 (E12K)-induced eosinophil survival. In addition, the anti-IL-5 neutralizing antibody TRFK-5 blocked the agonist activity of IL-5 (E12K) and wild-type IL-5, but not IL-3, GM-CSF or LPS in the eosinophil survival assay (Fig. 5 B). This confirms that the eosinophil survival activity is specifically mediated via IL-5 (E12K) and excludes the possibility of an LPS-dependent mechanism.



**Figure 5.** Both wild-type IL-5 and IL-5 (E12K) promote eosinophil survival (A). The agonist activity of E12K can be abolished by boiling, or pretreatment with a neutralizing anti-IL-5 antibody but not by polymyxin B (B). (A) Increasing concentrations of either wild-type (open circles) or IL-5 (E12K) (closed circles) were incubated with purified human eosinophils for 72 h at 37°C and eosinophil viability measured by trypan blue exclusion. Data expressed as a percentage of the maximum survival effect obtained with wild-type IL-5 in each experiment (66% ± 5.6 viable cells, compared to 2.7% ± 1.7 in the absence of cytokine). Each value represents the mean ± SEM of six independent experiments using different blood donors. (B) Human eosinophils were incubated with either 1 pM IL-5, 50 nM IL-5 (E12K), 40 pM IL-3, 20 pM GM-CSF, or 1 ng/ml LPS alone, or in the presence of 250 μg/ml anti-IL-5 neutralizing antibody, TRFK-5, or after pretreatment with 500 U/ml polymyxin B for 1 h at 37°C. For boiling experiments, IL-5 (E12K) was boiled for 15 min before incubation with eosinophils. Eosinophil viability was measured after 72 h by trypan blue exclusion. The concentrations of cytokines represent ~ED<sub>80</sub>'s in the eosinophil survival assay. Data expressed as a percentage of the maximum survival effect obtained with wild-type IL-5 in each experiment (61% ± 8.0 viable cells, compared to 17% ± 3.7 in the absence of cytokine). Each value represents the mean ± SEM of at least three independent experiments.

**Relative Effects of IL-5 and IL-5 (E12K) on Tyrosine Phosphorylation in Eosinophils.** Stimulation of eosinophils with IL-5 induced an increase in the phosphotyrosine content of a number of proteins, in a concentration-dependent manner, with the predominant phosphorylated species at 150-kD (Fig. 6). In contrast to wild-type IL-5, IL-5 (E12K) did not stimulate tyrosine phosphorylation even at concentrations up to 5 μM. Furthermore, 500 nM IL-5 (E12K) completely

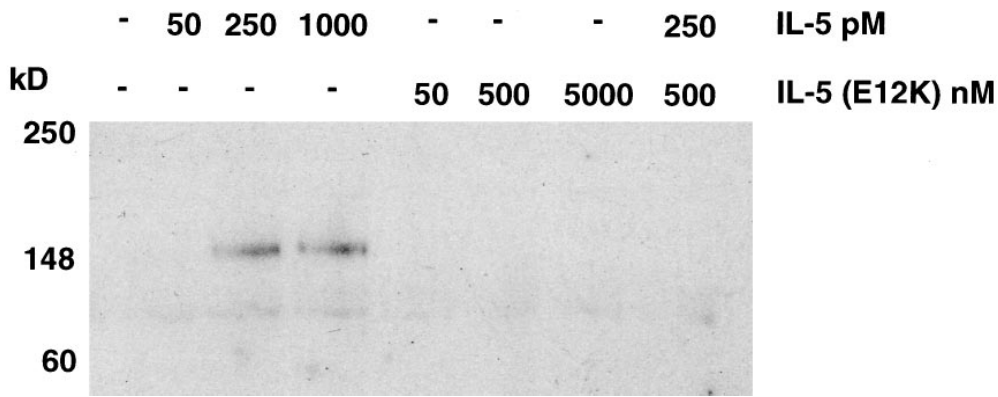
inhibited the tyrosine phosphorylation of this 150-kD protein stimulated by 250 pM IL-5 (Fig. 6).

## Discussion

In a systematic program of alanine scanning mutagenesis we previously demonstrated that IL-5 (E12A) exhibited partial agonist activity in a TF-1 cell proliferation assay and proposed that E12 defined a contact point for the IL-5 receptor β chain. We based this conclusion not only on biological activity but on the reduction in affinity of E12A for the IL-5 receptor α/β chain compared to the α chain alone (28). Generating a charge reversal at this putative β<sub>c</sub> chain binding site residue, IL-5 (E12K), has allowed us to further investigate the effect of this mutation.

Glutamic acid-12 resides on the amino-terminal helix of IL-5 at an analogous position to E21 in GM-CSF and E22 in IL-3. Charge reversal mutations at these positions in IL-3 and GM-CSF resulted in proteins which were unaltered in binding their respective α chains, but demonstrated a reduction in high-affinity binding (37, 38). For IL-5 (E12K) we also see essentially wild-type binding to the IL-5 receptor α chain, but a reduction in high affinity binding of ~2.6-fold. Although this result is entirely consistent with the reduction in high affinity we previously observed with IL-5 (E12A) (28), the apparent reduction in the affinity of IL-5 (E12K) for the α/β complex is not statistically significant ( $P = 0.084$ ). Since the affinity conversion conferred by β<sub>c</sub> on the IL-5/IL-5 receptor-α chain complex is small (two- to fourfold), compared to the much larger shifts in affinity seen with both GM-CSF and IL-3, this lack of statistical significance is probably due to the difficulty of measuring such small changes. The contribution of the β<sub>c</sub> chain to IL-5 (E12K) high-affinity binding is difficult to measure clearly through competition binding studies. However we demonstrated by chemical cross-linking studies that, as with wild-type IL-5, IL-5 (E12K) is still capable of contacting not only the receptor α chain but also the β<sub>c</sub> chain. This result is consistent with the pattern of cross-linking obtained with IL-5 (E13Q), which has a similar antagonist biological profile in TF-1 cells (29). Such cross-linking experiments have not been published with human GM-CSF (E22R) or IL-3 (E21R) and it is not clear whether these mutants can similarly cross-link to the β<sub>c</sub> chain in the absence of high affinity binding. However, a biologically active murine GM-CSF mutant (E21A) has been described which binds the α/β complex with low affinity only, yet is able to cross-link to both the α and β<sub>c</sub> chains of the receptor (39). As suggested by Tavernier et al. (29), this may indicate that affinity conversion and receptor activation can be uncoupled.

In contrast to IL-5 (E12A), the IL-5 (E12K) mutant was completely devoid of agonist activity in a TF-1 cell proliferation assay and was a full and selective antagonist of IL-5 in this functional assay. While IL-3 (E21R) retains agonist activity for TF-1 proliferation at 20,000-fold reduced potency relative to wild-type IL-3 (37), GM-CSF (E22R) lacks agonist activity and is a specific GM-CSF antagonist



**Figure 6.** Wild-type IL-5 but not IL-5 (E12K) stimulates tyrosine phosphorylation of a 150-kD protein in human eosinophils. Human eosinophils were stimulated with the indicated concentrations of cytokines for 5 min at 37°C. Cells were pelleted, resuspended in SDS sample buffer and lysates were electrophoresed and immunoblotted with anti-phosphotyrosine antibody.

(40). Furthermore, the antagonistic properties of IL-5 (E12K) are not restricted to inhibition of IL-5-induced cell proliferation. In human eosinophils IL-5 (E12K) was a selective antagonist of IL-5 induced eosinophil activation, as assessed by adhesion to immobilized IgG, exhibiting no effect on eosinophil activation induced by the related cytokines IL-3 and GM-CSF or the unrelated cytokine TNF- $\alpha$ .

Taken together these mutagenesis studies suggest that the structural conservation at both the cytokine and receptor level for IL-5, IL-3, and GM-CSF underlie a common mechanism of receptor binding and activation, and that the acidic residues within the amino terminal helix of all three cytokines form a contact point for the  $\beta_c$  chain which contributes to receptor activation and, for the latter two cytokines at least, high-affinity binding. Somewhat surprisingly however, IL-5 (E12K) was clearly a full agonist in an eosinophil survival assay, albeit with a 50,000-fold reduction in potency relative to wild-type IL-5. This survival effect of IL-5 (E12K) was not due to the presence of contaminating LPS and was consistently seen in the same eosinophils in which IL-5 (E12K) antagonized IL-5-induced adhesion to IgG. Clearly this single point mutation in IL-5 is able to distinguish between two IL-5-dependent functional responses in the same cell type. This apparent functional dichotomy in the actions of IL-5 (E12K) implies that different functional responses within the same cell, activation versus survival, are mediated through distinct mechanisms of cellular activation.

In the context of IL-5, E12 appears to be necessary for activating the IL-5 receptor  $\beta_c$  leading to an eosinophil activation signal. However, the fact that IL-5 (E12K) promotes eosinophil survival suggests that residues on IL-5 other than E12 are involved in receptor activation leading to a survival signal. The reduction in potency of IL-5 (E12K) with respect to wild-type IL-5 further suggests that, although not required for full agonist activity, E12 must contribute to the efficiency of receptor activation. This would indicate that multiple points of contact between IL-5

and its receptor complex are required for complete receptor activation. Since IL-5 (E12K) can be cross-linked to the receptor  $\beta_c$ , residues other than E12 may directly contact  $\beta_c$  contributing to its complete activation. Such a model may also account for the ability of human IL-3 (E22R) and murine GM-CSF (E21A) to exhibit biological activity despite the lack of high-affinity binding (37,39). However, a direct role for the IL-5 receptor  $\alpha$  chain in signal transduction has been inferred from both deletion and point mutation studies in the intracellular domain of the  $\alpha$  chain, resulting in the loss of IL-5-induced cell proliferation (21, 22). Furthermore, it has been proposed that the ligand-specific  $\alpha$  subunits of the IL-5, IL-3, and GM-CSF receptors may mediate cytokine-specific signals leading to different cellular responses (41). Since mutations at E12 appear to be unaffected in  $\alpha$ -chain binding, it is also possible that elements of the survival signal emanate from the  $\alpha$  chain.

These properties of IL-5 (E12K) highlight its potential use as a tool for dissecting signaling pathways from the activated IL-5 receptor complex leading to distinct IL-5-dependent functional responses, in particular eosinophil survival. Initial experiments in eosinophils measuring one of the earliest detectable signaling events, tyrosine phosphorylation, highlighted opposing effects of IL-5 and IL-5 (E12K). Wild-type IL-5, but not IL-5 (E12K), was able to activate tyrosine phosphorylation of a major 150-kD protein species and furthermore, the IL-5-dependent phosphorylation of this protein was inhibited by IL-5 (E12K).

In summary, we have identified an IL-5 point mutant, IL-5 (E12K), which lacks agonist activity and is a potent antagonist in both a TF-1 cell proliferation assay and an eosinophil activation assay, yet exhibits agonist activity in promoting eosinophil survival. The dual agonist and antagonist properties of IL-5 (E12K) can be explained via distinct mechanisms of cellular/receptor activation. We are currently attempting to elucidate the precise mechanism by which this IL-5 mutant exerts its agonist effect.

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