# High Affinity IgE-Fc Receptor $\alpha$ and $\gamma$ Subunit Interactions

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# ABSTRACT

**Objective:** To explore the relationships between the subunits ( $\alpha$ ,  $\beta$  and  $\gamma$ ) of the high affinity IgE receptor (Fc $\epsilon$ RI) and its ability to mediate transmembrane signaling.

Study Design: Experimental study.

Place and Duration of Study: Department of Molecular Biology and Biotechnology, University of Sheffield, UK, from 2008 to 2009.

**Methodology:** The approach employed was to create a chimera (human  $\alpha\gamma\gamma$ ) using the extracellular (EC) domain of the human high affinity IgE receptor. The alpha subunit (huFccRI $\alpha$ ) of IgE receptor was spliced onto the rodent gamma TM and cytoplasmic domain (CD). This was transfected into the Rat Basophilic Leukemia cell line in order to assess the possibility of selectively activating cells transfected with this single pass construct for antigen induced mediator release.

**Results:** The RBLs cell lines transfected with the huFccRI $\alpha/\gamma/\gamma$  cDNA constructs were assessed for the cell surface expression of the huFccRI $\alpha$  subunit and the response to the antigenic stimulus by looking for degranulation and intracellular Ca<sup>2+</sup> mobilisation. The results obtained showed the absence of huFccRI $\alpha$  subunit expression on the surface of transfected cells as seen by flowcytometric studies,  $\beta$ -hexosaminidase assays and intracellular calcium mobilisation studies.

**Conclusion:** In the present study the grounds for non-expression of huFc $\epsilon$ RI $\alpha$ / $\gamma$ / $\gamma$  cDNA remains elusive but may be due to the fact that the human-rodent chimeric receptors are assembled differently than the endogenous rodent receptors as seen in study in which COS 7 cells were transfected with human/rat chimeric complexes.

Key Words: The high affinity IgE receptor (FceRI). Rat basophilic leukemia cells (RBLs). Immunoglobulin E (IgE).

#### **INTRODUCTION**

The High Affinity IgE Receptor, FcERI has the highest affinity of all immunoglobulin receptors with a binding constant in the 10<sup>-9</sup> to 10<sup>-10</sup> M range for its ligand. IgE.<sup>1,2</sup> FccRI exists as a tetramer on the surface of human mast cells and basophils consisting of a  $\alpha$ -chain, a  $\beta$ -chain and a disulphide-linked dimer of  $\gamma$ -chains, while on Langerhans cells, dendritic cells and monocytes the FccRI is present as a trimer.<sup>3</sup> The subunits ( $\alpha$ ,  $\beta$  and  $\gamma$ ) of FcERI are composed of three domains: extracellular (EC), transmembrane (TM) and cytoplasmic (CT) domains.<sup>4,5</sup> Our group is exploring the relationships between the structure of the FcERI and its ability to mediate transmembrane signaling.6,7 The approach employed was to create a chimera (human $\alpha\gamma\gamma$ ) using the extracellular (EC) domain of the human high affinity IgE receptor alpha subunit (huFceRIa) spliced onto the rodent gamma TM and cytoplasmic domain (CD) and transfect them into the rat basophilic leukemia cell line in

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order to assess the possibility of selectively activating cells transfected with this single pass construct for antigen induced mediator release. This strategy might facilitate assessment of the surface expression of the huFc $\epsilon$ RI $\alpha$  ligand binding domain independent of the endogenous rodent huFc $\epsilon$ RI $\alpha$ .

## METHODOLOGY

The study was carried out from 2008 to 2009 at the Department of Molecular Biology and Biotechnology, Sheffield, United Kingdom. Splice Over Extension (SOE) PCR strategy was used to construct the  $\alpha\gamma\gamma$  chimaeric receptor, where the  $\alpha$  chain is of human origin<sup>8</sup> and the  $\gamma$  chain is of rat.<sup>9</sup>

Primer A - Forward Primer for Human FcεRIα:

5'-TCCTGCTAGCATGAAGAAGATGGCTCCT-'3 with a Nhel restriction site.

Primer B - Reverse Primer for Human Fc $\epsilon$ RI $\alpha$ :

5'-AGGCGTCGACGTACTTCTCACGCGGA-'3 with a Sall restriction site.

Primer C - Forward Rat FccRIy Primer:

5'-CTCACTCGAGGGAGAGCCGCAGCTC-'3 with a Xhol restriction site.

Primer D - Reverse Rat FcεRlγ Primer:

5'-AAGACTCGAGCTATTGGGGTGGTTT-'3 with a Xhol restriction site.

Primer E - Forward Human FccRI $\alpha$  Rat FccRI $\gamma$  Linker Primer:

#### 5'-CCGCGTGAGAAGTACGGAGAGCCGCAGCTC-'3.

Primer F - Reverse Human FccRl $\alpha$  Rat FccRl $\gamma$  Linker Primer:

5'-GAGCTGCGGCTCTCCGTACTTCTCACGCGG-'3.

Primer G - Reverse Human TM Region Internal Primer:

5'-TCCTGTCGACTGAGATAAATAATCCT-'3 with a Sall restriction site.

Primer H - Forward Rat CD Region Internal Primer:

5'-GGACGTCGACCGACTCAAGATCCAGG-'3 with a Sall restriction site.

The huFc $\in$ RI $\alpha$  gene cDNA was provided in the vector pGEM3Zf, (kindly provided by U. Blank and J-P. Kinet), and was used as the PCR template. The pUC18 plasmid was used as the cloning vector, and the pEE6 plasmid (Celltech) was used as the transfection vector. Miniprep plasmid DNA samples were sent for sequencing to the company Cogenics, using the M13-48REV and M13For-40 primers, after which the results were analysed using ALIGN, CHROMA and CLONE computer programs. The confirmed, non-mutated, clones were used to subclone the cDNA construct (huFc $\epsilon$ RI $\alpha/\gamma/\gamma$ ) into expression vector pEE6 for subsequent transfection into mammalian cells (RBL-2H3.1). Vectors pEE6 and pUC18 with huFccRI $\alpha/\gamma/\gamma$  were digested using the EcoRI and Hind III restriction enzymes (Figure 1). The desired bands were excised from the agarose gel and purified using the Quantum Prep<sup>™</sup> Freeze N Squeeze DNA gel extraction spin columns kit. The purified bands were ligated together and a negative (water) control was also set up to eliminate self-ligations. The ligation reactions were transformed into XLI-Blue supercompetent cells. Qiagen/Wizard DNA miniprep kit was employed for small scale DNA extraction. Test digestion of 10µl miniprep plasmid DNA was carried out using the EcoRI and Hind III restriction enzymes followed by resolving on TAE gel to confirm the presence of the desired fragment. The miniprep plasmid DNA was transformed into XLI-Blue competent cells. QIAGEN Plasmid Midi Kit was employed for large scale DNA extraction.

Expression of the  $\alpha\gamma\gamma$  chimeric receptor: The RBL-2H3.1 cell line<sup>10</sup> was employed as the host for the huFccRI $\alpha/\gamma/\gamma$  construct. Stable transfections were carried out using electroporation. In each case, a negative control using water in place of DNA was setup. The transfected cells were then used to carry out flow cytometric studies to ascertain the presence of the desired huFccRI $\alpha$  receptors on the cell surface. Following flowcytometric studies stocks of the transfected cells were prepared by cryogenic preservation and employed for release assays and intracellular calcium mobilisation studies. The RBL-2H3.1 transfected cell line with the huFccRI $\alpha/\gamma/\gamma$  cDNA

constructs was monitored for  $\beta$ -hexosaminidase release through endogenous and huFc $\epsilon$ RI $\alpha$  transfected receptors by challenging them with mouse IgE with DNP-HSA specificity and human IgE with NIP-HSA specificity.<sup>11,12</sup> The RBL-2H3.1 cell lines transfected with the huFc $\epsilon$ RI $\alpha$ / $\gamma$ / $\gamma$  cDNA constructs were assessed for surface expression of huFc $\epsilon$ RI $\alpha$  by employing the flow cytometry technique. Cells were labelled in series with a combination of human IgE (JW8), biotinylated antihuman IgE and streptavidin R-phycoerythrin antibodies. Intracellular Ca<sup>2+</sup> mobilization in RBL-2H3.1 transfected with huFc $\epsilon$ RI $\alpha/\gamma/\gamma$  cDNA constructs brought about by antigenic stimulus (NIP/DNP) was assessed using the Ca<sup>2+</sup> indicator Fluo-3AM.<sup>11,12</sup> IgE sensitized cells loaded with Fluo-3AM were analysed using a FACSort flow cytometer preset for Fluo-3AM studies. Data was recorded in the form of a density plot of mean fluorescence against time. After an initial background reading of 30-50 seconds, samples were activated with the appropriate cross-linking agent (NIP/DNP 100 ng/ml) and in the event of no calcium mobilization, the samples were challenged with a calcium ionophore (ionomycin) as a positive control to ensure that the cells have been loaded with Fluo-3AM.



**Figure 1:** Test Digestion of Miniprep and Midiprep Plasmid DNA was carried out using the EcoRI and Hind III restriction enzymes followed by resolving on TAE gel to confirm the presence of the desired fragment.

In Lane 1 (Figure A) bands visible are 2686bp (pUC18), 6200bp (pEE6) and Lane 1 (Figure B) 1117bp (huFccRl $\alpha/\gamma/\gamma$  construct) while lane M is the 1Kbp DNA marker.



Figure 2: Assessment of cell surface expression of huFc $\epsilon$ RI $\alpha$  transfected receptors in RBL-2H3 cells transfected with huFc $\epsilon$ RI $\alpha$ / $\gamma$ / $\gamma$  construct by flow cytometry.

Figure 2a shows the control histogram while Figure 2b shows the cell population histograms of sample of huFccRla/ $\gamma/\gamma$  transfected cell lines which were assessed for huFccRla receptor expression.

## RESULTS

Data from the FACS analysis of the transfected RBLs is shown in Figure 2. The data demonstrates the absence of huFccRla receptor subunit on the cell surface of transfected RBLs (Figure 2b). The RBLs transfected with huFccRla/ $\gamma/\gamma$  cDNA constructs when sensitised with human IgE and subsequently challenged with human IgE specific NIP-HSA model antigen failed to release above background readings (Figure 3). RBL-2H3 trans-



**Figure 3:** Release of  $\beta$ -hexosaminidase through rodent FccRI $\alpha$  and huFccRI $\alpha$  transfected receptors in RBL-2H3 transfected with human huFccRI $\alpha/\gamma/\gamma$  construct in response to antigenic stimulus.

Cells were cultured, harvested, resuspended at 0.5 x 10<sup>6</sup>/ml in appropriate media with NIP-specific hulgE (SPE, 1/500)/ DNP-specific mIgE and plated into 96 well plates. Next days, cells were washed and activated with DNP-HSA/NIP-HSA cross-linking agent (0.1-1000 ng/ml) for 20 minutes prior to incubation with  $\beta$ -hexosaminidase substrate for 2 hours.  $\beta$ -hexosaminidase release was assessed as described in material and methods. Data is presented as mean  $\pm$  S.D. from three separate experiments performed in triplicate. Released  $\beta$ -hexosaminidase is expressed as a percentage of total  $\beta$ -hexosaminidase.



Figure 4: Assessment of Intracellular calcium level of RBLs transfected with huFccRIa/ $\gamma/\gamma$  cDNA constructs following activation in the absence of extracellular calcium.

RBL-2H3 cells transfected with the huFccRla/ $\eta/\eta$  cDNA constructs were sensitised with (Figure A) DNP-specific mIgE (SPE-7, 1/500) or (Figure B) NIP-specific hulgE (JW8, 1/500) for 16 hours. After an initial background reading the cells were activated with appropriate cross-linking agent DNP (Figure A) and NIP (Figure B) at a concentration of 100 ng/ml. Following excitation at 488 nm emitted fluorescence was recorded at 525 nm using a FACSort flow cytometer. The cells were then activated with ionomycin (10  $\mu$ M, Figure B).

fected with huFcɛRIα/ $\gamma/\gamma$  cDNA constructs sensitized with mIgE exhibited an increase in mean fluorescence peaking at 26 seconds after being activated by mIgEspecific DNP-HSA (Figure 4a) and no calcium mobilization was observed on being sensitized with huIgE and activated by huIgE-specific NIP-HSA (Figure 4b). The same transfected RBLs on failure to elicit calcium mobilization with huIgE-specific NIP-HSA were activated by ionomycin as a positive control and exhibited an immediate increase in mean fluorescence which was maintained until a slow decrease occurred relating to a depletion of intracellular Ca<sup>2+</sup> (Figure 4b).

#### DISCUSSION

The technique of constructing novel chimera has provided insight into the understanding the phenomena of allergy.8 In addition, research is being done in treatment of cancers by constructing chimeric antigen receptor-engineered T-cells.9 Various studies have been conducted using chimera made up of the CD of the  $\gamma$ subunit FccRI or TCR $\zeta$  and the EC domain of CD4<sup>10</sup> or Tac, the interleukin 2 receptor.<sup>11</sup> In the study by Letourneur et al. (1991) a chimeric receptor consisting of EC domain of  $\alpha$  chain of the interleukin receptor (Tac) and CT domain of either  $\zeta$  or  $\gamma$ . When they are expressed in T-cells or RBLs could be activated leading to release of interleukin 2 in T-cells and serotonin in RBLs. stably transfected into RBL-2H3 cells. The two constructs prepared by Repetto et al. (1996) using the EC and cytoplasmic (CT) domains of the FcERI and the IL-2R p55 subunit<sup>1</sup> were  $\alpha/\gamma/\gamma$  and  $1/\gamma/\gamma$  and results demonstrated that  $Fc \in RI\alpha$  EC and  $Fc \in R\gamma$  CT domains are mandatory for signalling process leading to the response characteristically seen after the receptor aggregation.<sup>12</sup> The RBLs cell lines transfected with the huFc $\epsilon$ RI $\alpha/\gamma/\gamma$  cDNA constructs were assessed for the cell surface expression of the huFc $\epsilon$ RI $\alpha$  subunit and the response to the antigenic stimulus by looking for degranulation and intracellular Ca<sup>2+</sup> mobilisation. The results obtained showed the absence of huFcERIa subunit expression on the surface of transfected cells as seen by flowcytometric studies,  $\beta$ -hexosaminidase assays and intracellular calcium mobilisation studies. Similar results were obtained previously by Higginbottom in 1996 (PhD thesis). In contrast, a study by Repetto et al. a  $\alpha/\gamma/\gamma$  construct when stably transfected into RBL-2H3 cells lead to expression of the receptor which was able to exhibit all the signalling events similar to the native rat FcERI thus demonstrating the mandatory role of both the  $Fc \in Rl\alpha$  EC domain and FcR-y CT domains.12 It was expected that similar expression if achieved in the present study could lead to mutations being introduced in TM domains of the  $\gamma$  that would provide insight into the intersubunit interactions. The huFccRIa/ $\gamma/\gamma$  cDNA construct had the problem of presence of spurious point mutations, which needed

correction. Similar problems were encountered in the generation and transfection of  $\alpha/\gamma/\gamma$  cDNA constructs into RBL cells in a study by Higginbottom (1996). It is, therefore, surprising that the results published by Repetto and collaborators (1996) could not be reproduced in the present and previous study conducted by Higginbottom (1996). In the study by Repetto et al. VIS expression vector (unavailable), which possesses a visna virus promoter for constitutive expression, was used for transfection into COS and RBL cell lines,12 in addition in the study by Higginbottom pMAMneo vector, which is an inducible vector, was employed for transfection of  $\alpha/\gamma/\gamma$  cDNA constructs into RBL cells.<sup>13</sup> In the present study, constitutive expression pEE6 vector, which has been successfully used for the expression of wild-type and mutant huFc $\in$ RI $\alpha$  constructs into RBL-2H3 cell line, was used and the reason for the nonexpression of huFc $\epsilon$ RI $\alpha/\gamma/\gamma$  cDNA constructs remains uncertain. The human-rodent chimeric receptors are assembled differently than the endogenous rodent receptors as seen in study in which COS 7 cells were transfected with human/rat chimeric complexes.14

#### CONCLUSION

In the present study, the grounds for non-expression of huFccRIa/ $\gamma/\gamma$  cDNA remains elusive but may be due to the fact that the human-rodent chimeric receptors are assembled differently than the endogenous rodent receptors as seen in study in which COS 7 cells were transfected with human/rat chimeric complexes.

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