# Alternate Splicing of Interleukin-1 Receptor Type II (IL1R2) In Vitro Correlates with Clinical Glucocorticoid Responsiveness in Patients with AIED

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Alternate Splicing of Interleukin-1 Receptor Type II (IL1R2) In Vitro Correlates with Clinical Glucocorticoid Responsiveness in Patients with AIED

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Abstract
Autoimmune Inner Ear Disease (AIED) is poorly characterized clinically, with no definitive laboratory test. All patients suspected of having AIED are given glucocorticoids during periods of acute hearing loss, however, only half initially respond, and still fewer respond over time. We hypothesized that AIED is a systemic autoimmune disease characterized by dysfunctional peripheral blood mononuclear cells (PBMC) responses to a unique cochlear antigen(s). To test this hypothesis, we examined end-stage AIED patients undergoing cochlear implant surgery and compared autologous perilymph from AIED patients to controls. We determined that autologous perilymph from AIED patients was unable to induce expression of a long membrane-bound Interleukin-1 Receptor Type II (mIL1R2) transcript in PBMC as compared with controls, despite similar expression of the short soluble IL1R2 (sIL1R2) transcript (p<0.05). IL1R2 is a molecular decoy that traps interleukin-1β (IL-1β) and does not initiate subsequent signaling events, thereby suppressing an inflammatory response. IL1R2 transcript length is regulated by alternate splicing, and the major inhibitory function is attributed to the full-length mIL1R2. In addition, IL1R2 expression is induced by dexamethasone. Separately, we prospectively examined patients with newer onset glucocorticoid-responsive AIED. Immediately prior to clinical treatment for acute deterioration of hearing thresholds, their PBMC demonstrated a robust induction of mIL1R2 in PBMC in response to dexamethasone in vitro that correlated with a clinical response to prednisone in vivo (p<0.0001) as measured by hearing restoration. In contrast, clinically steroid unresponsive patients demonstrated high basal levels of mIL1R2 in their PBMC and only minimally augmented expression in response to dexamethasone. Thus, induced expression of mIL1R2 appears to be a protective mechanism in hearing homeostasis and warrants further investigation in a large prospective clinical trial to determine if IL1R2 can be used as a specific biomarker for AIED.

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Introduction
Sensorineural hearing loss (SNHL) is a common otologic problem that typically is not amenable to medical intervention. Potentially reversible SNHL can be divided into two sub-groups: (1) Autoimmune Inner Ear Disease (AIED) and (2) sudden SNHL (SSNHL). Timely administration of glucocorticoids in these two conditions may result in hearing preservation. While SSNHL is usually an isolated, unilateral and single event, patients with AIED usually experience multiple episodes of rapid hearing loss in both ears. This disorder can be an isolated process affecting the ear only, or it may be part of a systemic autoimmune disorder in approximately 30% of those afflicted [1]. Further characterization of AIED has been hampered by limited access to the inner ear, and by the absence of a reliable biomarker that distinguishes AIED from patients with other forms of SNHL [2]. Thus, the incidence and prevalence of AIED is unknown, however an initial response to glucocorticoids may assist to define this clinical disorder [1,3,4,5]. In patients with AIED, hearing preservation with glucocorticoids is often only maintained short term [3,6]. Of the 60 [3] to 70% [6] of patients who are initially steroid responsive, only 14% remain so after 34 months [6]. It is unknown whether these patients who initially respond, and then become refractory to steroid therapy, have undergone a change in immune response that precludes continued response to corticosteroids. Moreover, on a molecular level, it is unknown how steroids induce a clinical response.

Medical therapy for AIED patients refractory to corticosteroids is limited. Methotrexate has been shown to be beneficial in several small series; however, in a large prospective trial, methotrexate was comparable to placebo in maintaining hearing in AIED patients [5]. In animals, suppression of TNFa by Etanercept was helpful in treating AIED, however, a recent human clinical trial failed to confirm this [7]. When medical therapy fails to restore hearing, and hearing aids are no longer beneficial, cochlear implantation is often used to restore hearing in patients with AIED who develop bilateral, severe to profound SNHL. Identification of a more effective medical therapy for AIED is contingent on
developing a better understanding of the mechanism(s) that cause this disorder.

A much more common entity treated with corticosteroid therapy is sudden sensorineural hearing loss (SSNHL). A fraction of patients with SSNHL are considered to have AIED [8], as the majority of these patients are thought to have either viral or vascular pathophysiologic causes of their disease. Nonetheless, therapy for all patients with a sudden decline in hearing is similar and usually treatment entails the prompt use of corticosteroids. Evidence of an autoimmune mechanisms being important in SSNHL has been reported, and includes an increased prevalence of antibodies specific for a HSP70 (a heat shock protein), commonly found in the blood of a cohort of SSNHL patients [9], and an increased frequency of the HLA-DRB1*04 allele in SSNHL, which is associated with a poor response to glucocorticoids [10]. HLA-DRB1*04 has also been associated with other autoimmune disorders such as autoimmune thyroiditis and rheumatoid arthritis [11,12].

Many autoantibodies to cochlear and other self-antigens in human serum have been shown to be enriched in AIED patients. Presumptive markers for AIED that has been identified include autoantibodies to: ANA (18–43%), HSP70 (22–89%) and others [recently reviewed [13]]. However, no single antibody is present in all patients with this disease [2]. An increased prevalence of several autoantibodies have correlated with glucocorticoid-responsive AIED, including antibodies to both HSP70 [14] and CTCL12, recently suggested to be a biomarker for this disease [15]. HSP 70 garnered a great deal of attention because of its high specificity for AIED [14]; however its pathogenic role is controversial [16], since mice immunized with bovine HSP 70 failed to experience hearing loss [17].

Animal models of AIED have demonstrated that the inner ear is accessible from the peripheral circulation and no strict anatomic barrier exists within the cochlea. Immune cells have been shown to traffic into the inner ear from the peripheral circulation [2,18], and radiolabeled lymphocytes have been shown to enter the scala tympani of the cochlea [2,18]. Furthermore, T cells from the systemic circulation have been shown to proliferate in the endolymphatic sac, and perilymph proteins of the cochlea can activate immunocytes in the endolymphatic sac [19]. These passive transfer experiments suggest that there is no anatomic barrier to trafficking lymphocytes from the systemic circulation into the inner ear.

Recent animal studies suggest that macrophages (MΦ), which are part of the innate immune response, can augment adaptive T-cell responses in AIED [20]. In these experiments, re-exposure to KLLH antigen in the presence of systemic lipopolysaccharide (LPS) caused upregulation of Interleukin-1β (IL-1β) expression in the cochlea, leukocyte entry into the cochlea, and subsequent hearing loss. As anticipated, LPS induced Toll Like Receptor 4 (TLR4) signaling and augmented IL-1β expression and was associated with hearing loss. These experiments suggest that the innate immune system may participate in the development of AIED [20]. In addition, the production of IL-1β by MΦ, likely causes T-cell polarization to Th1-like response that might perpetuate AIED. A critical role of pro-inflammatory CD4+, Th1-like T-cells has been shown in the cochlin peptide–murine model [21]. Interferon-gamma (IFN-γ)-producing, Th1-like T cells have been identified in the peripheral blood of AIED patients [22]. Modulation of interleukin-1β levels has resulted in IFN-γ expression by T-cells in other autoimmune disorders. In murine autoimmune encephalomyelitis, blockade of the IL-1 receptor antagonist (IL-1RA) permits soluble IL-1β to bind the IL-1 receptor type 1 (IL-1R1), augmenting the development of clinical encephalomyelitis and T-cells expression of IFN-γ [23]. Thus, the absence of IL-RA expression, or other molecules that oppose the IL-1β inflammatory cascade, during an immune response, can promote the development of autoimmune disease, including AIED.

**Results**

We hypothesized that patients with AIED have a systemic immunologic disease manifested in the inner ear initiated by a unique cochlear antigen. Given that PBMC are capable of trafficking to the inner ear during inflammation [2], we anticipated that immunocytes that had previously encountered a cochlear antigen would respond differently to cochlear proteins compared with those that had not been previously exposed. For this reason, we compared the effect of adding autologous perilymph to peripheral blood mononuclear cells (PBMC) in patients with AIED who underwent cochlear implantation (end stage disease) compared to PBMC from patients undergoing implantation for longstanding, non-immunologic, stable, SNHL (hereafter called controls) using RNA expression microarray analysis. Patient data can be seen in table 1. Results for each patient were compared to unstimulated autologous PBMC (negative control) and pneumococcal-stimulated autologous PBMC. The rationale for the pneumococcal stimulation is that the standard care for all patients undergoing cochlear implantation is prior immunization with pneumococcal vaccine to prevent meningitis [24]. Thus, all patients were primed by vaccination with a 23-valent-pneumococcal vaccine (Pneumovax®). The majority of those with AIED also had systemic autoimmune disease. Interestingly, none of these subjects had the same autoimmune disorder, reducing the possibility of a single II HLA allele as the mediating factor. Microarray results for AIED and control subjects were analyzed by comparing unstimulated PBMC, with autologous perilymph stimulated, and pneumococcal stimulated, PBMC (figure 1A). Analysis of the microarray data was performed by normalizing data sets by RNA and an ANOVA analysis was performed on grouped arrays by condition with a Benjamini and Hochberg correction, and a threshold of 2.0 fold change (Genesifter, VixXlabs). Of note, only 10 genes were differentially expressed in these experiments when the threshold was set at 2 and a Benjamini & Hochberg correction applied (p<0.05). Of those 10 genes, only one, the interleukin-1 receptor type 2 (IL1R2) (affymetrix ID 205403, nt 1104–1485 (long membrane bound form of IL1R2 (mIL1R2)), was differentially expressed when autologous perilymph was added to PBMC of control subjects, as compared with AIED subjects (Figure 1A, 1B). Analysis of IL-1RI, IL-1β, IL-1RA, heat shock proteins (HSP) and TNF-α all failed to show differential expression (not shown).

**IL1R2 Expression**

To confirm our microarray results, we used a quantitative real time polymerase chain reaction (Q-RT-PCR). IL1R2 exists in two forms: membrane-bound (mIL1R2) and soluble (sIL1R2). The RNA primers used to detect the membrane bound, long form of the IL1R2 message were derived from the coding region of the IL1R2 gene present on the Affymetrix gene array chip (figure 2A). The shorter soluble form is made by alternative splicing [25], or by enzymatic cleavage of the membrane bound protein by either a metalloproteinase, aminopeptidase or elastase [26,27,28]. However, the shed IL1R2 size varies with a metalloproteinase producing a 45 kD product, and aminopeptidase and elastase producing a 60 kD product [26]. Interestingly, both AIED patients and controls make transcripts for the shorter form, although at slightly lower levels in AIED subjects compared with controls (affymetrix ID 211372, nt 635–1084) (figure 2B). RNA for the longer,
Table 1. Clinical History of AIED and control patients undergoing cochlear implantation.

<table>
<thead>
<tr>
<th>Patient</th>
<th>Status</th>
<th>Autoimmune History</th>
<th>Hearing History</th>
<th>PTA &amp; HINT</th>
<th>Serology</th>
</tr>
</thead>
<tbody>
<tr>
<td>40 y/o female(^1,2)</td>
<td>Control (birth anoxia?)</td>
<td>None</td>
<td>Congenital loss – non progressive</td>
<td>PTA 100 dB; HINT 60 dB quiet: 0%</td>
<td>NP*</td>
</tr>
<tr>
<td>49 y/o male(^1,2)</td>
<td>Control</td>
<td>None</td>
<td>40+ years loss</td>
<td>PTA 100 dB; HINT 50 dB quiet: 10%</td>
<td>NP*</td>
</tr>
<tr>
<td>79 y/o female(^1,3)</td>
<td>Control (LVA)</td>
<td>None</td>
<td>70+ years LVA</td>
<td>PTA 110; HINT 55 dB quiet: 0%;</td>
<td>NP*</td>
</tr>
<tr>
<td>1.5 y/o female(^2)</td>
<td>Control (Cx 26)</td>
<td>None</td>
<td>Congenital – stable</td>
<td>PTA: NR at 102 dB by ABR; IT-MAIS: 12%</td>
<td>NP*</td>
</tr>
<tr>
<td>36 y/o female(^1,2)</td>
<td>AIED, progressive loss</td>
<td>Rheumatoid Arthritis</td>
<td>29 years, periods of rapid loss</td>
<td>PTA 120 dB; HINT: 55 dB quiet: 0%</td>
<td>P0, Rheumatoid factor IgM (14.2) &amp; IgG (25.3), immune complex, ANA 1:1280 (Hep-2), IgM phospholipids (10.9), immune complex (26.1)</td>
</tr>
<tr>
<td>56 y/o male(^1,2)</td>
<td>AIED, rapid loss</td>
<td>Type I IDDM</td>
<td>2 years, rapid loss</td>
<td>PTA 120; HINT 70 dB quiet: 0%</td>
<td>ANA 1:1280 (mouse kidney), ANA 1:320 (Hep-2), weakly positive IgM phospholipid Ab (12.1)</td>
</tr>
<tr>
<td>58 y/o female(^1,2,3)</td>
<td>AIED, rapid loss</td>
<td>Hashimoto’s thyroiditis</td>
<td>4 years, rapid loss</td>
<td>PTA 110 dB; HINT at 50 dB quiet: 0%</td>
<td>Borderline for immune complex (21.4), weakly positive IgM phospholipids (11.7), borderline rheumatoid factor IgG (20.8)</td>
</tr>
<tr>
<td>51 y/o female(^1,2)</td>
<td>AIED, progressive loss</td>
<td>None</td>
<td>Sudden loss left ear 10 years prior, sudden loss right ear &lt;6 months</td>
<td>PTA 80 dB; HINT 65 dB quiet: 19%</td>
<td>68 kD positive</td>
</tr>
<tr>
<td>3 y/o male(^2)</td>
<td>AIED, b/l rapid SNHL</td>
<td>None</td>
<td>6 months, rapid loss</td>
<td>PTA 120 dB; IT-MAIS 8%</td>
<td>Negative</td>
</tr>
</tbody>
</table>

PTA: Pure tone average of the audiogram at 500, 1000, 2000 and 4000 Hz, (although 4000 Hz is not traditionally included, this information is important to speech perception for cochlear implantees and therefore PTA represents these four frequencies for these patients). NP: Not Performed: control subjects did not have serology for AIED performed as there was no clinical indication for such studies, HINT: Hearing in Noise Test; IT-MAIS: Infant-Toddler Meaningful Auditory Integration Scale, LVA: Large Vestibular Aqueduct; Cx26: connexin 26 gene mutation. Superscript numbers in column 1 refer to inclusion of the patient in data sets for the below figures: 1: included in analysis for figure 1; 2b & c: included in analysis for figure 2b or 2c. In one patient, all RNA was used for microarray analysis (79 y/o female, control).

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Figure 1. Differential expression of IL1R2 by microarray in response to autologous cochlear perilymph in patients undergoing cochlear implantation. 1A Comparison of RNA expression using the Affymetrix U133A 2.0 chip to compare cultured PBMC of controls (CTRL) and AIED patients for the following conditions: either unstimulated (UNTX), stimulated with 23-valent pneumococcal vaccine (PNEUMO) or autologous cochlear perilymph (C). Only 10 genes (panel A) were significantly different (p < 0.05) when a threshold of 2 and a Benjamini & Hochberg correction was applied (Genesifter, VixXlabs). Only one of them unique to the cochlear fluid stimulated condition IL1R2 (affy ID 205403), p < 0.05. Array data in 1A can be viewed at www.ncbi.nlm.nih.gov/geo/ using the user “Vambutas” with a password “daniella” to view series “GSE4277” containing 18 arrays from the six patients with “1” next to their description in table 1. 1B A comparative analysis of IL1R2 expression in these arrays is seen in panel 1B, with standard deviations shown.
doi:10.1371/journal.pone.0005293.g001
membrane bound form, is not expressed in perilymph stimulated PBMC from AIED patients (no transcript detected at 45 cycles); however, in perilymph stimulated control PBMC, mIL1R2 RNA is strongly expressed (figure 2B). The difference was statistically significant using a Mann Whitney test \( (p = 0.03) \) (in stat). Notably, the inhibitory function of IL1R2 is selectively attributed to this long, membrane bound form [29].

Soluble IL1R2 levels, which reflect of the shorter IL1R2 message (affymetrix 211372), were reduced in AIED patients in culture supernatants of PBMC stimulated with perilymph measured by ELISA (figure 2C). This finding parallels our microarray results for Affymetrix probe 211372 (not shown) and our Q-R-PCR expression of sIL1R2 in AIED and control subjects (figure 2B). Interestingly, at least for the AIED cochlear implant recipients, there did not appear to be increased sIL1R2 as the mechanism for IL1R2 reduction compared to controls.

**IL1R2 splicing correlates with clinical steroid responsiveness**

Only 60% of patients studied by the AIED study group were steroid responsive [3]. To date, it is unknown why some AIED patients respond while others do not, although presence of certain antibodies have been correlated with steroid responsiveness [14,15]. We hypothesized that robust expression of mIL1R2, in response to corticosteroids may be indicative of a clinical response. To test the hypothesis, we enrolled patients with acute deterioration of their hearing that met the same criteria that the AIED study group used [3]. Blood for PBMC isolation was drawn immediately prior to oral steroid therapy (60 mg of prednisone daily for a minimum of 7 days with a variable taper thereafter). A repeat audiogram was performed between 7–10 days post initiation of treatment, and in a subset of patients, blood for PBMC isolation was again drawn. Table 2 describes the eighteen patients enrolled in this study. The average hearing recovery at all frequencies tested is shown in Table 2. All responders returned to their prior baseline hearing thresholds at the end of the glucocorticoid therapy. Non-responders were defined as those patients that showed less than 5 dB improvement at all frequencies tested, or further decline in pure tone thresholds. In these studies, we observed that pre-treatment expression of mIL1R2 mRNA by PBMC in response to dexamethasone stimulation \( \text{in vitro} \), correlated with clinical responsiveness to oral steroid therapy. Corticosteroid responders had absent pre-treatment, basal mIL1R2 expression; however, they dramatically augmented...
Table 2. Clinical History of patients treated with prednisone.

<table>
<thead>
<tr>
<th>PATIENT</th>
<th>PRIOR TX?</th>
<th>PMH</th>
<th>SEROLOGY</th>
<th>MRI</th>
<th>AFFECT</th>
<th>EAR</th>
<th>250 Hz</th>
<th>500 Hz</th>
<th>1000 Hz</th>
<th>2000 Hz</th>
<th>4000 Hz</th>
<th>AVG</th>
<th>Improve</th>
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<tbody>
<tr>
<td>56 WM (NR1)</td>
<td>YES</td>
<td>NC</td>
<td>NA</td>
<td>NEG</td>
<td>PRETX</td>
<td>35</td>
<td>45</td>
<td>60</td>
<td>105</td>
<td>120</td>
<td>−1 dB</td>
<td></td>
<td></td>
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<tr>
<td>42 WF (NR2)</td>
<td>YES</td>
<td>Hypo-thyroid</td>
<td>ANA 1:320, Immune complex +</td>
<td>NEG</td>
<td>PRETX</td>
<td>50</td>
<td>75</td>
<td>90</td>
<td>95</td>
<td>95</td>
<td>−1 dB</td>
<td></td>
<td></td>
</tr>
<tr>
<td>61 HF (NR3)</td>
<td>YES</td>
<td>NC</td>
<td>NA</td>
<td>NEG</td>
<td>PRETX</td>
<td>65</td>
<td>60</td>
<td>40</td>
<td>35</td>
<td>45</td>
<td>−3 dB</td>
<td></td>
<td></td>
</tr>
<tr>
<td>57 WM (NR4)</td>
<td>YES</td>
<td>Brother W/Crohns</td>
<td>ANA1:2560, P0 pos</td>
<td>NEG</td>
<td>PRETX</td>
<td>55</td>
<td>60</td>
<td>75</td>
<td>65</td>
<td>80</td>
<td>−19 dB</td>
<td></td>
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<tr>
<td>45 WM (NR 5)</td>
<td>YES</td>
<td>NC</td>
<td>RF borderline +</td>
<td>NEG</td>
<td>PRETX</td>
<td>60</td>
<td>60</td>
<td>65</td>
<td>65</td>
<td>70</td>
<td>+2 dB (NS)</td>
<td></td>
<td></td>
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<tr>
<td>62 WF (NR6)</td>
<td>YES</td>
<td>NC</td>
<td>Weak+68 kD Borderline typell collagen</td>
<td>NEG</td>
<td>PRETX</td>
<td>10</td>
<td>10</td>
<td>10</td>
<td>10</td>
<td>40</td>
<td>−9 dB</td>
<td></td>
<td></td>
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<tr>
<td>60 WF (NR7)</td>
<td>YES</td>
<td>Pulm collag vasc disease</td>
<td>P0 pos</td>
<td>NEG</td>
<td>PRETX</td>
<td>60</td>
<td>60</td>
<td>60</td>
<td>55</td>
<td>75</td>
<td>−1 dB</td>
<td></td>
<td></td>
</tr>
<tr>
<td>WM (NR8)</td>
<td>YES</td>
<td>Father w/Auto-immune disease req. steroids</td>
<td>Neg</td>
<td>NEG</td>
<td>PRETX</td>
<td>50</td>
<td>55</td>
<td>65</td>
<td>65</td>
<td>60</td>
<td>0 dB</td>
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<tr>
<td>46 WF (NR9)</td>
<td>NO</td>
<td>Multiple Sclerosis</td>
<td>NA</td>
<td>NEG</td>
<td>PRETX</td>
<td>95</td>
<td>90</td>
<td>95</td>
<td>120</td>
<td>120</td>
<td>−9 dB</td>
<td></td>
<td></td>
</tr>
<tr>
<td>56 WF (R1)</td>
<td>YES</td>
<td>Daughter W/IDDMM</td>
<td>ANA 1:560, Phospholipid IgG &amp; IgM pos</td>
<td>NEG</td>
<td>PRETX</td>
<td>55</td>
<td>60</td>
<td>60</td>
<td>40</td>
<td>40</td>
<td>+37 dB</td>
<td></td>
<td></td>
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<tr>
<td>54 WF (R2)</td>
<td>YES</td>
<td>Hashimotos Thyroiditis</td>
<td>ANA 1:160, 68 kD pos</td>
<td>NEG</td>
<td>PRETX</td>
<td>105</td>
<td>115</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>+39 dB</td>
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<tr>
<td>39 WM (R3)</td>
<td>YES</td>
<td>NC</td>
<td>NA</td>
<td>NEG</td>
<td>PRETX</td>
<td>40</td>
<td>45</td>
<td>35</td>
<td>15</td>
<td>30</td>
<td>+15 dB</td>
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<tr>
<td>69 WF (R4)</td>
<td>YES</td>
<td>Crohn's Disease</td>
<td>NA</td>
<td>NEG</td>
<td>PRETX</td>
<td>40</td>
<td>35</td>
<td>30</td>
<td>30</td>
<td>20</td>
<td>+6 dB</td>
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<tr>
<td>59 WF (R5)</td>
<td>YES</td>
<td>Fibromyalgia</td>
<td>NA</td>
<td>NEG</td>
<td>PRETX</td>
<td>105</td>
<td>105</td>
<td>105</td>
<td>100</td>
<td>95</td>
<td>+9 dB</td>
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<tr>
<td>53 WM (R6)</td>
<td>YES</td>
<td>Psoriasis</td>
<td>NA</td>
<td>NEG</td>
<td>PRETX</td>
<td>50</td>
<td>55</td>
<td>45</td>
<td>25</td>
<td>25</td>
<td>+17 dB</td>
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<td>46 WM (R7)</td>
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<td>NC</td>
<td>NA</td>
<td>NEG</td>
<td>PRETX</td>
<td>55</td>
<td>60</td>
<td>30</td>
<td>10</td>
<td>60</td>
<td>+11 dB</td>
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<td></td>
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<tr>
<td>79 WF (R8)</td>
<td>YES</td>
<td>NC</td>
<td>NA</td>
<td>NEG</td>
<td>PRETX</td>
<td>50</td>
<td>40</td>
<td>40</td>
<td>55</td>
<td>80</td>
<td>+11 dB</td>
<td></td>
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</tr>
<tr>
<td>37 WF (R9)</td>
<td>YES</td>
<td>NC</td>
<td>Neg</td>
<td>NEG</td>
<td>PRETX</td>
<td>20</td>
<td>30</td>
<td>45</td>
<td>60</td>
<td>75</td>
<td>+19 dB</td>
<td></td>
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</tr>
</tbody>
</table>

"Prior tx?": although all had prior rapid declines in hearing requiring prednisone therapy, none were treated in a 3 month period prior to enrollment. Several patients had occasional vestibular symptoms, however, these symptoms did not coincide with periods of hearing fluctuation. Serology was performed at Immco Diagnostics. NA = not available; serologic testing at Immco could not be performed as these tests were not covered by the patient’s insurance carrier. Audiometric change shown represents the change between the average of the post-treatment pure tone thresholds from the pre-treatment pure tone average threshold. NS = not significant: average post-treatment hearing changes of less than 5 dB are not felt to represent significant changes. All declines in hearing are reported as 0 dB. NR = clinical non-responder; R = clinical responder.

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mIL1R2 expression in response to dexamethasone stimulation, p<0.0001 (figure 3A), using a Mann Whitney test. Although the mean and standard deviations are large in the groups largely due to the steroid non-responsive patient that clustered with the steroid responsive group, the median fold change of the non-responsive group was 47 as compared to the median of the responsive group was 2.37×10^6. If we set a threshold for clinical steroid responsiveness to be a greater than 500 fold induction of mIL1R2 in vitro, then the sensitivity of this test is 100% with a confidence interval of 71.7 to 100%. The specificity would be 88.9% with a confidence interval of 31.8–99.7%. The dramatic difference in the pre-treatment fold change of the corticosteroid responders was inversely proportional to basal mIL1R2 expression by unstimulated PBMC (45 cycles, S.D. 0.0 in replicate samples), (denoted “unstim” in figure 3B), compared to corticosteroid non-responders (28.48 cycles, S.D. ±3.34) (Figure 3B). Post-treatment basal expression of mIL1R2 in steroid responders was detected at a much lower cycle number by Q-RT-PCR than pre-treatment (figure 3B). Pre-treatment basal mIL1R2 expression in glucocorticoid-responders dramatically increased from no expression at 45 cycles to detectable expression at 30 cycles post-treatment, compared with 30 cycles to 27 cycles post-treatment in non-responders suggesting that the glucocorticoid responsiveness correlated with induction of mIL1R2 RNA expression (figure 3B). Furthermore, as seen with the cochlear implant patients, despite absent mIL1R2 expression, sIL1R2 expression was observed at reasonable levels (figure 3C). Interestingly, the steroid refractory patients demonstrated high mIL1R2 expression in absence of stimulation prior to treatment, with no major change in expression in response to dexamethasone (figure 3B, denoted “dex stim”). This expression pattern suggests some endogenous stimulation of mIL1R2 expression.

It is paradoxical that AIED cochlear implant recipients demonstrated absent basal expression of mIL1R2, yet steroid refractory AIED patients exhibit high levels of mIL1R2 in absence of in vitro stimulation. Steroid-refractory patients with high basal

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**Figure 3. Pre-treatment induced expression of mIL1R2 in response to dexamethasone correlates with post-treatment clinical hearing restoration.** 3A 18 patients with presumptive AIED and sudden declines in hearing were enrolled, blood drawn prior to treatment, and treated with 60 mg prednisone/day for 7 days and then tapered. Pre-treatment PBMC were divided and incubated with either no stimulus or dexamethasone (8 μg/ml is shown). IL1R2 mRNA was measured by Q-RT-PCR. Post-treatment audiograms were obtained between 7–14 days later. Audiometric improvement (∆) was measured as the average of 250, 500, 1,2,& 4 kHz. Responders (R) all returned to prior baseline hearing. Pre-treatment levels of mIL-1R2 Q-RT-PCR (expressed as fold change over baseline) predicted steroid responsiveness in AIED (P<0.0001, Mann Whitney test). 3B Pre and post treatment mIL1R2 levels in a subset of clinical responders and non-responders. Pre-treatment responders have no basal mIL1R2 expression in PBMC (unstim) at 45 cycles (in replicate samples), compared with non-responders that have substantially higher basal levels. Although PBMC from both responders and non-responders demonstrate increased mIL1R2 RNA expression in response to in vitro dexamethasone stimulation (dex stim), responders are exquisitely sensitive. The concentration of dexamethasone shown for the PBMC stimulation experiments was 20 μg/ml, however, 4 μg/ml and 8 μg/ml produced identical results, suggesting maximal stimulation even at 4 μg/ml. 3C Despite clear differences in mIL1R2 expression patterns, sIL1R2 expression is differences between responders and non-responders are minimal suggesting alternate splicing in clinical responders.

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expression of mIL1R2 may represent a different disorder. Alternatively, steroid-refractory patients have a different inflammatory microenvironment that overweights the effects of mIL1R2-induced expression. The absent basal expression of mIL1R2 in AIED cochlear implant recipients may reflect the absence of acute inflammation at the time of implantation. Our results demonstrate that low basal and induced expression of mIL1R2 in AIED during periods of inflammation and rapid hearing decline correlated with the clinical response to glucocorticoids.

**Discussion**

IL1R2 is known to be a molecular decoy that sequesters IL-1β, and blocks the initiation of downstream signaling, thereby preventing inflammation [as reviewed [30,31]]. The IL1R2 protein is expressed on B cells, monocytes/macrophages and neutrophils [30], and may be expressed in response to many stimuli, including oxygen radicals, IL-4, IL-13, and glucocorticoids [30], possibly explaining the ability of steroids to ameliorate AIED. Release of the soluble IL1R2 (sIL1R2) may also be induced by various stimuli, although the major inhibitory function of IL1R2 has been ascribed to the membrane bound form [29]. The induction of IL1R2 by IL-4 [32] certainly suggests that the maintenance of a T\(_p\)2-like phenotype is important in controlling inner ear disease, whereas expression of IFN\(_γ\), that fosters a T\(_H\)1-like response, has been observed in AIED patients [22], and blocks the expression of IL1R2 [33]. IL1R2 is an important mechanism that prevents IL-1β-mediated inflammation at sites where inflammation is poorly tolerated. In the brain, IL1R2 is preferentially expressed compared to IL1R1, the cognate IL-1β receptor [34]. Restoring IL1R2 expression in IL1R2 deficient cells resulted in significant cellular protection from IL-1β mediated damage [35]. Treatment of mice with IL1R2 expressing cells reduced inflammation in collagen-induced arthritis [36], demonstrating that IL1R2 may be beneficial in diseases where IL-1β is dominant and endogenous counteracting antagonists are absent.

In steroid responsive patients, sIL1R2 RNA was detected in unstimulated PBMC, however, the long mIL1R2 transcript could not be detected. In the presence of dexamethasone, the long transcript could readily be detected. LPS stimulation of monocytes results in reduced mRNA synthesis of IL1R2 [28]. Although transcriptional induction of IL1R2 from dexamethasone stimulation of monocytes has been studied extensively by Northern analysis [37,38,39], alternate splicing has not been reported, perhaps because these studies were performed in a limited number of control subjects or small molecular weight shifts induced by alternate splicing near the 3’ end may not have been detected in these studies. Dexamethasone has however, been reported to induce differential splicing in other genes [40]. Furthermore, alternate splicing of critical regulatory genes has been identified to associate with risk of other autoimmune diseases, such as alternate splicing of BANK1 in Systemic Lupus Erythematosus [41]. Investigation into whether genetic polymorphisms affect the splicing of the IL1R2 message is underway in our laboratory.

Although it has been shown that no anatomic barrier exists preventing leukocyte trafficking to the inner ear [2,18], the robust expression of mIL1R2 in control cochlear implant subjects, and AIED patients who responded to glucocorticoids suggests that the inner ear likely is at least a partially functionally immunoprivileged site. We hypothesize that AIED patients who experience a break in tolerance and fail to express IL1R2 in response to antigenic stimuli, have unopposed IL-1β expression that leads to hearing loss. Unopposed expression of IL-1β has been shown to induce sensorineural deafness in patients with Muckle-Wells syndrome, an autoimmune disorder. In these patients, restoration of hearing was observed following anakinra therapy, an IL-1 receptor antagonist [42,43], with normalization of IL-1β serum levels [43].

We further hypothesized that the expression of IL1R2, in response to various stimuli, likely represents an innate immune response by macrophages. Macrophages are the predominant producer of IL1R2 [37,39]. Macrophage ingress into the cochlea has been demonstrated in traumatic injury: perhaps best characterized during periods of acoustic-induced trauma [44]. Furthermore, during acoustic trauma, pro-inflammatory cytokines, including IL-1β are expressed [45]. Similarly, IL-1β expression was observed in response to LPS in an animal model of AIED [20]. Reduction of mononuclear cell secretion of IL-1β was observed following anakinra therapy and correlated with hearing restoration in a patient with Muckle-Wells syndrome [43]. We hypothesize that as the AIED progresses, an increased production of IFN-γ blocks expression of IL1R2 in the cochlea. Preliminary flow cytometry experiments demonstrate induction of IL1R2 surface expression in response to dexamethasone occurs in blood monocytes in these patients (not shown). We are also examining the perilymph of these patients to ascertain whether it is the cytokine microenvironment, or the presence of a unique protein or autoantigen found in perilymph that is inducing differential IL1R2 expression in perilymph from AIED and controls.

To date, ability to identify glucocorticoid responsive AIED patients prior to treatment has been limited to a correlation with seropositivity for either HSP70 or CTCL2 [14,15]. Thus, identifying mIL1R2 levels that predict steroid responsiveness early on in disease may help better phenotype those with AIED, and those who will benefit from this therapy thereby avoiding the complications associated with this treatment in patients not expected to respond.

The observations we describe suggest that mIL1R2 expression may be an important mechanism involved in glucocorticoid responsiveness in AIED; however, given that steroids control inflammation by multiple mechanisms, it is probable that other pathways contribute to the observed response. Clinical therapies that alter IL-1β expression have not been used in AIED patients.

**Methods**

**Patient Recruitment**

**Ethics Statement.** These studies were approved by the North Shore-LIJ Health System IRB. All patients signed written consent for inclusion into these studies. In the case of the two children, their parents provided written consent. Assent was not obtained from the children due to their young age.

**Cochlear Implant Subjects.** All patients met current criteria for cochlear implantation and had a bilateral profound SNHL. All patients with AIED had periods of rapid hearing loss and were no longer steroid responsive. All adult controls had stable SNHL for over 30 years of clear, non-immunologic origin. A total of 9 patients were recruited (7 adults 2 children) representing 4 controls and 5 with AIED, and their clinical demographics shown in table 1.

At the time of cochlear implant surgery; blood was drawn to isolate PBMC. Perilymph fluid was harvested from the cochleostomy in a bloodless field. In order to control for a cochlear perilymph specific response, we compared unstimulated PBMC, to cochlear perilymph stimulated PBMC, and pneumococcal stimulated PBMC, from both AIED and control patients. Three microarrays were performed per study subject: 1) perilymph+autologous PBMC, 2) pneumococcal antigen+autologous PBMC, and 3) PBMC alone and administration of pneumococcal vaccine...
2 weeks prior to cochlear implant surgery, considered to be standard of care prior to cochlear implantation, served as a positive control.

**Prospectively Enrolled, Steroid Treated Subjects**

Eighteen patients with AIED who had not undergone cochlear implantation were studied in a separate IRB approved protocol (Table 2). They all had a clinical history suggestive of AIED, and met criteria defined for AIED trials [3]. At the time of enrollment, they all demonstrated sudden declines in their hearing and were treated with 60 mg of prednisone daily for 7 days with a variable taper thereafter. The responders were defined as a 5 dB or greater average improvement at 250, 500, 1000, 2000 and 4000 Hz. All patients recruited must have had prior audiograms demonstrating their baseline-hearing threshold. Furthermore, SNHL of greater than 30 dB at one or more frequencies in both ears with evidence of active deterioration (elevated threshold) in at least one ear of 15 dB at one frequency (excluding 250 Hz as a sole indicator), or 10 dB at 24-frequencies developing in >3 days but <90 days [3,16]. If the hearing loss evolved in less than 3 days, prior similar hearing declines must have occurred or the patient must have a systemic autoimmune disorder AND the patient must meet the audiology criteria outlined above. All patients with retrocochlear (vestibular schwannoma or other internal auditory canal) pathology, patients who received prednisone or other immunosuppressive therapy within 3 months, or patients with retrocochlear symptoms coinciding with periods of hearing fluctuation [3] were all excluded from this study. All patients received a minimum of 60 mg of prednisone daily for 7 days with a variable taper thereafter.

**Microarray Analysis.** PBMC were isolated over a Ficoll-hypaque gradient, divided into 5 ml cultures (2 x 10^6 cells/ml) and cultured in RPMI+10%FCS for 16 hours at 37°C with 5% CO₂ with one of 3 stimuli (1) without antigenic stimulus (untx), (2) or (3) with 150 nM and 60 ng of total RNA. The DNA sequences were compared to a T7 DNA polymerase promoter. The DNA was stranded, ethanol precipitated and biotinylated cRNA generated by in vitro transcription (ENZO). cRNA was purified on a spin column (Qiagen), quantitated, and 20 μg fragmented and hybridized to the Affymetrix HG U133A 2.0 array (Affymetrix). Data sets were normalized using RNA and an ANOVA analysis was performed on grouped arrays by condition with a Benjamini and Hochberg correction, and a threshold of 2.0 fold change (Genesifter, VixXlabs). Array data in 1A can be viewed at www.ncbi.nlm.nih.gov/geo/ using the user “Vambutas” with a password “daniella” to view series “GSE4277” containing 18 arrays from the six patients with “1” next to their description in table 1. The array data is MIAME compliant. Upon publication, this data will be publicly released, with the password removed.

**Q-RT-PCR.** Quantitative Real Time RT-PCR (Q-RT-PCR) was performed on PBMC from patients using TaqMan chemistry. The relative abundance of IL1R2 mRNA for 2 membrane bound (mIL1R2) and the soluble (sIL1R2) coding regions associated with the 2 affymetrix probes sets 211372 (sIL1R2) and 205403 (mIL1R2) were compared to β-actin was determined using the Eurogentec RTqPCR mastermix (Eurogentec, Belgium) and ABI PRISM 7700 Sequence Detection System. Membrane bound IL1R2, as reflected by affymetrix probe set 205403 was detected by patients at 129–125, 1295–1314, and taqman probe 1271–1278, whereas the shorter soluble form (sIL1R2) reflected by the affymetrix probe 211372 and primers nt 790–830–849, and taqman probe 820–827. The alternative splice site described by Liu changing GAA to TAA, introducing a stop codon, occurs at nt 1118 [25]. These primers were added at final concentration of 200 nM and 100 nM respectively to 50 ng of total RNA. The conditions were 48°C for 30’, 95°C for 10’ and 45 cycles of 95°C for 15’ and 60°C for 1’. Data was analyzed using Sequence Detection System software version 1.9.1. Results were expressed as Ct (Threshold cycle) values, which is inversely proportional to the starting template copy number. Relative abundance of IL1R2 in cochlear fluid stimulated cells was calculated compared to untreated control samples using delta delta Ct method (User Bulletin #2, ABI). In all prospectively enrolled, steroid treated patients, the unstimulated condition is the average of 2 independent PBMC cultures (SD for responders was 0.0, and SD for non-responders was 3.34 cycles).

**ELISA Analysis.** ELISA for soluble IL-1R2 (sIL-1R2). Supernatants from 16-hour PBMC cultures from 3 AIED patients and 2 controls were used to determine the level of soluble receptor (See table 1 for inclusion). ELISA was performed according to manufacturers’ instructions (R&D Systems). Stimulus conditions were either pneumococcal stimulus as used above or autologous cochlear perilymph and compared to unstimulated PBMC. A large number of duplicate samples were run to ensure accuracy.

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**Author Contributions**

Conceived and designed the experiments: AV. Performed the experiments: AV JD. Analyzed the data: AV ML VB. Contributed reagents/materials/analysis tools: JD EG MG. Wrote the paper: AV.

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