Relationship between asthma severity and progression of Alzheimer's disease

Severity of asthma is occasionally modulated by neuropsychiatric conditions. However, little is known about the impact of cognitive decline on asthma severity. Cognitive decline is a core symptom in patients with Alzheimer's disease (AD). AD is a disease characterized by progressive cholinergic failure that could possibly reduce airway hyperresponsiveness to cholinergic stimulation and thus symptoms of asthma. Furthermore, the functions of T lymphocytes—which play a crucial role in the development of chronic asthma—are partially impaired in patients with AD related diseases.

We hypothesised that declining cognitive function might result in an improvement in asthma, and prospectively studied the contribution of the progression of AD to the clinical course of asthma.

Eight patients with asthma of mean (SE) duration 15.3 (0.9) years with concomitant AD were identified and prospectively followed for 5 years from 1995 to 2000. All subjects were treated with oral theophylline (200 mg twice daily) and a 200 µg dose of fenoterol given by a metered-dose inhaler as needed. Family members of the patients completed a diary card that recorded asthma symptoms, of the patients.

During the study period, n=8, p=0.26). No other precipitating factors for asthma were identified during the study period in any subject. Both overall attack frequency and severity of asthma symptoms decreased significantly during the progression of cognitive impairment in asthma patients with AD. However, peripheral cholinergic function might not be impaired in the airways in patients with AD despite an extensive loss of central cholinergic neurons. It has been reported that the nervous system may modulate immunological and inflammatory responses. Our results suggest that progression of AD might provide an ameliorating effect on the clinical course of asthma, probably due to alterations in the immunological responses including eosinophilic inflammation in the airway.

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References


IL-1 haplotypes and lung function decline

We read with interest the paper by Joos et al. on the association of IL-1 gene haplotypes with decline in lung function in smokers and share their view on a possible role of IL-1 genetics in inflammatory respiratory diseases. We have analysed the same polymorphism by

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The editors will decide as before whether to also publish it in a future paper issue.

Table 1: Assessment of asthma severity and change in cognitive function at study entry (baseline) and 5 year follow up (end point) in asthma patients with Alzheimer’s disease

<table>
<thead>
<tr>
<th>Case</th>
<th>Age (y)</th>
<th>Sex</th>
<th>MMSE score</th>
<th>Asthma symptom score</th>
<th>Daily inhaler puffs</th>
<th>Number of hospital admissions for asthma</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>Baseline</td>
<td>End point</td>
<td>Baseline</td>
<td>End point</td>
</tr>
<tr>
<td>1</td>
<td>67</td>
<td>M</td>
<td>23</td>
<td>18</td>
<td>6.4</td>
<td>1.2</td>
</tr>
<tr>
<td>2</td>
<td>66</td>
<td>M</td>
<td>21</td>
<td>16</td>
<td>8.6</td>
<td>2.2</td>
</tr>
<tr>
<td>3</td>
<td>70</td>
<td>F</td>
<td>23</td>
<td>17</td>
<td>10.2</td>
<td>1.6</td>
</tr>
<tr>
<td>4</td>
<td>65</td>
<td>M</td>
<td>22</td>
<td>15</td>
<td>7.8</td>
<td>1.2</td>
</tr>
<tr>
<td>5</td>
<td>65</td>
<td>M</td>
<td>21</td>
<td>16</td>
<td>7.5</td>
<td>0.4</td>
</tr>
<tr>
<td>6</td>
<td>69</td>
<td>F</td>
<td>23</td>
<td>17</td>
<td>9.4</td>
<td>3.6</td>
</tr>
<tr>
<td>7</td>
<td>66</td>
<td>F</td>
<td>22</td>
<td>16</td>
<td>9.2</td>
<td>2.4</td>
</tr>
<tr>
<td>8</td>
<td>68</td>
<td>M</td>
<td>23</td>
<td>21</td>
<td>7.6</td>
<td>7.4</td>
</tr>
<tr>
<td>Mean (SE)</td>
<td>67.0 (0.7)</td>
<td>22.3 (0.3)</td>
<td>17.0 (0.7)*</td>
<td>8.4 (0.5)</td>
<td>1.8 (0.4)</td>
<td>2.2 (0.3)</td>
</tr>
</tbody>
</table>

MMSE=Mini-Mental State Examination; SE=standard error.
*p<0.0001 (Wilcoxon rank test) compared with baseline data in all asthma patients with Alzheimer’s disease; †p<0.0001 compared with baseline data in seven asthma patients with Alzheimer’s disease (cases 1–7).

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The same methods in adult incident non-smoking asthmatic patients and non-smoking controls. Our results indicate that the association of IL-1 genetics with rate of decline in lung function is not limited to smokers.

New adult asthma cases and controls were selected from a cohort of the Mini-Finland Health Survey (MFHS) and later re-evaluated. A more detailed description of the methods used in MFHS has been published elsewhere. The accuracy of the method of asthma case ascertainment has also recently been described. 1 IL-1 haplotypes were found to be significantly associated with the rate of decline of lung function in non-smoking incident cases of asthma (new asthma during follow-up) but not in controls (Table 1). Of the individual haplotypes, Joos et al found that \( \text{IL1RN A2/IL1B} -511T \) was associated with a rapid decline of lung function in smokers and \( \text{IL1RN A2/IL1B} -511T \) with a slow decline. In our control group the observed differences were not significant. Surprisingly, in the asthma group the haplotypes had the opposite effects from those in smokers: \( \text{IL1RN A2/IL1B} -511T \) was associated with a slower decline in lung function and \( \text{IL1RN A2/IL1B} -511T \) with a more rapid decline. \( \text{IL1RN A2/IL1B} -511T \) has previously been found to be associated with many inflammatory diseases. The function of these haplotypes would therefore appear to be disease specific.

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References

Authors’ reply
Karjalainen and colleagues present interesting data on the relationship of IL-1β and IL-1 receptor antagonist haplotypes and the rate of decline of lung function in incident asthmatic subjects in a Finnish cohort. We reported that the \( \text{IL1RN A1/IL1B} -511T \) haplotype was associated with a more rapid decline in lung function in smokers in the Lung Health Study; in contrast, they found that this same haplotype was associated with a slower rate of decline in lung function in patients with asthma. The authors suggest that this apparent contradiction may be because the function of these haplotypes is disease specific. We agree that a different effect of the same haplotype could occur because of fundamental differences in the pathophysiological processes which cause airflow obstruction in asthma and chronic obstructive pulmonary disease (COPD). In asthma, CD4+ Th2 cells underlie persistent eosinophilic inflammation and remodelling in medium sized and larger airways. In COPD, neutrophils and CD8+ cells appear to play an important role in the airflow limitation by causing proteolytic destruction of peripheral lung parenchyma and fibrous scarring of the small blood vessels and respiratory bronchioles. Although inflammation appears to be central to both processes, the roles of IL-1 and IL-1 receptor antagonist in these conditions is unknown and it is possible that the different polymorphisms which are typed to establish haplotypes could have opposite effects.

Alternatively, these apparently contradictory results could be due to different genetic histories of the two study groups. Our study group was taken from the white population in the United States whereas Karjalainen et al studied Finnish individuals. It may be that the polymorphisms which are typed to establish these haplotypes do not, by themselves, change the function or level of expression of the IL-1 proteins but are in linkage disequilibrium with a causal polymorphism(s). In this case, the IL-1 allelic associations could be different in different populations. The bottleneck in the genetic history of the Finnish people could have established a founder effect and resulted in the function altering allele being found on a different genetic background from that in the white population of the United States.

Whatever the correct explanation, these results support the growing evidence that genetic variation at the IL-1 locus is important in modulating the severity and/or functional consequences of a number of inflammatory conditions.

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Molecular analysis of drug resistant TB
Since the mid 1980s the number of notified cases of TB in the UK has continued to rise with the largest increases noted in London and inner city areas. King George Hospital in Goodmayes, Essex provides clinical services to a population of approximately 230 000: 17% are non-white subjects including immigrants from countries with high rates of \( M \) tuberculosis infection and drug resistance. From September 1996 to July 1997 47 adult cases of culture proven TB were identified including seven with drug resistant isolates. None was identified by contact tracing. A previous TB audit of African born patients revealed a high rate of drug resistance (6/24 (25%)) and delays in obtaining drug sensitivities which could have been detrimental to patient management.

Under these circumstances the rapid identification of drug resistance in \( M \) tuberculosis isolates would have been helpful. The aim of this study was to determine retrospectively the usefulness of PCR-reverse hybridisation methods for screening for mutations within or adjacent to \( M \) tuberculosis genes associated with rifampicin ( \( \text{rpoB} \) ) and isoniazid ( \( \text{inhA} \) ) resistance. We also determined whether resistance genotyping combined with IS6110 typing could help to identify clusters of drug resistant cases not previously identified by contact tracing.

Seven consecutive drug resistant \( M \) tuberculosis culture isolates were analysed for rifampicin and isoniazid resistance and the results were compared with conventional susceptibility testing. The commercially available

<table>
<thead>
<tr>
<th>Drug resistance genotype</th>
<th>Resistance genotype</th>
</tr>
</thead>
<tbody>
<tr>
<td>Wild type</td>
<td>( rpoB ) mutation</td>
</tr>
<tr>
<td>Wild type</td>
<td>( \text{inhA} ) mutation</td>
</tr>
<tr>
<td>Wild type</td>
<td>Wild type</td>
</tr>
<tr>
<td>Wild type</td>
<td>Wild type</td>
</tr>
</tbody>
</table>

Table 1
Demographic data, site, phenotypic and genotypic resistance of the seven resistant study isolates

<table>
<thead>
<tr>
<th>Isolate</th>
<th>Age</th>
<th>Country of birth</th>
<th>Site of TB</th>
<th>Drug resistance</th>
<th>Resistance genotype</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>24</td>
<td>Nigeria</td>
<td>Pulmonary</td>
<td>INH/RIF</td>
<td>Wild type</td>
</tr>
<tr>
<td>2</td>
<td>21</td>
<td>Somalia</td>
<td>Pulmonary</td>
<td>INH/RIF</td>
<td>Wild type</td>
</tr>
<tr>
<td>3</td>
<td>40</td>
<td>Zaire</td>
<td>Pulmonary</td>
<td>INH/RIF</td>
<td>Wild type</td>
</tr>
<tr>
<td>4</td>
<td>17</td>
<td>Zaire</td>
<td>Pulmonary</td>
<td>INH/RIF</td>
<td>Wild type</td>
</tr>
<tr>
<td>5</td>
<td>20</td>
<td>Zaire</td>
<td>Pulmonary</td>
<td>INH/RIF</td>
<td>Wild type</td>
</tr>
<tr>
<td>6</td>
<td>42</td>
<td>UK</td>
<td>Pulmonary</td>
<td>INH/RIF</td>
<td>Wild type</td>
</tr>
<tr>
<td>7</td>
<td>44</td>
<td>Somalia</td>
<td>Sternum</td>
<td>PZ</td>
<td>Wild type</td>
</tr>
</tbody>
</table>

INH=isoniazid; RIF=rifampicin; PZ=pyrazinamide. Isolates 3 and 5 had indistinguishable IS6110 types. Isolates 1 and 4 were not typable due to insufficient culture and the banding pattern of isolate 6 was uninterpretable.

Table 1 Decline in FEV\(_1\) (ml/year) in non-smoking asthmatics and controls in different putative IL-1 haplotypes

<table>
<thead>
<tr>
<th>IL-1 haplotype</th>
<th>(1) IL1RN A2/IL1B -511T</th>
<th>(2) IL1RN A2/IL1B -511C</th>
<th>(3) IL1RN A1/IL1B -511T</th>
<th>(4) IL1RN A1/IL1B -511C</th>
<th>p value*</th>
<th>Post hoc tests between allele groups</th>
</tr>
</thead>
<tbody>
<tr>
<td>Non-smoking controls</td>
<td>41.8 (13.8)</td>
<td>44.8 (12.3)</td>
<td>50.7 (11.2)</td>
<td>45.5 (19.4)</td>
<td>0.27</td>
<td>Not tested</td>
</tr>
<tr>
<td>(n=124)</td>
<td>(n=54)</td>
<td>(n=7)</td>
<td>(n=31)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Non-smoking new asthma</td>
<td>63.2 (24.6)</td>
<td>37.7 (20.7)</td>
<td>30.9 (16.9)</td>
<td>51.0 (24.8)</td>
<td>0.0443</td>
<td>3&lt;1, p=0.02, 2&lt;1, p=0.06</td>
</tr>
<tr>
<td>(n=40)</td>
<td>(n=18)</td>
<td>(n=4)</td>
<td>(n=14)</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

*Two way analysis of variance; T1SD test of means.
assay INNO-LIPA Rif.TB<sup>®</sup> was used to detect rpoB mutations and an in-house PCR-reverse hybridisation line probe was used to detect mutations in or adjacent to the katG, inhA, and 

**References**


7. A previous paper from this hospital described apical lung bullae in four young male marijuana smokers, three West Indian and one Caucasian. We describe further cases (one woman) with large upper lobe bullae. All were Caucasian.

8. **Acknowledgements**

We would like to thank the Steering Group Members of the “Molecular Epidemiology of Tuberculosis in London” for allowing us access to their database and to the Mycobacterial Reference Laboratory (Dulwich) for conventional susceptibility testing.

**Pathophysiology of COPD**

The paper by Dentener et al. is interesting and contributes to the understanding of the pathophysiology of chronic obstructive pulmonary disease (COPD). It is becoming clear that COPD is a systemic syndrome, and this paper suggests some potential mechanisms. However, a number of issues merit further comment.

It is noted that, in healthy controls, there is a wide range of C reactive protein (CRP) values extending well beyond what would be considered to be the normal range. The reason for this is unclear, but it does suggest that these individuals are not as healthy as described. In addition, patients with stable COPD have a range of CRP values that also extend beyond this normal range. This is not consistent with previous studies, which suggests that, in patients with stable COPD, the range of CRP values falls within the normal range. Although patients with bronchiectasis were excluded, it is possible that undiagnosed bronchiectasis may have been present. Previous work has shown that 29% of patients presenting with what appeared to be stable COPD had CT evidence of at least mild bronchiectasis. This could conceivably explain a wider range of CRP levels. In addition, it is interesting that after just 5 days of treatment for an acute exacerbation of COPD the CRP had returned to a level below that of the stable cohort in the study. Since standard treatment for an exacerbation is able to achieve this in just a few days, it suggests that the stable group may have contained individuals that were in fact not so stable.

The authors allude to the potential confounding effect of systemic corticosteroids in the study. The changes in total leucocyte count during the exacerbation are likely to be due to the effect of prednisolone, making it difficult to interpret the changes in leucocyte count. In stable patients the action of corticosteroids may also confound the results. It is possible that, even in patients using inhaled corticosteroids, leucocyte numbers could be affected since there may be significant bioavailability at higher doses. Leucocyte count should therefore not be used as a marker for systemic inflammation in these patients.

3. It would appear that the exacerbations of COPD might have been mild, despite the presence of severe COPD on lung function criteria. Although PaO<sub>2</sub> was slightly higher and PaCO<sub>2</sub> slightly lower than in the stable group, these differences were small in magnitude. The pH was not significantly different and, in fact, the stable group contained individuals with a lower pH range (7.30–7.50) than in the exacerbated group (pH 7.34–7.49). Although the mean CRP level appears higher than in stable patients, the range does not differ significantly. This may therefore have led to a less profound change in inflammatory markers than might have been expected, and a study looking at more severe exacerbations may be more revealing.

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to the discrepancy between the studies by Gompertz et al and ours. No relationship was observed between CRP level and severity of disease, whereas separation of patients into normometabolic and hypermetabolic subgroups revealed enhanced levels of CRP in the latter group. Further research is needed to elucidate the cause for the production of these acute phase proteins in subgroups of clinically stable COPD patients and its involvement in the pathogenesis of the disease. It is of interest to note that, although only some of the clinically stable patients had enhanced CRP levels, almost all patients who had an exacerbation of disease had increased CRP levels on day 1 of the exacerbation which declined during treatment.

We agree with Dr Kelly that treatment with corticosteroids could affect the leucocyte count in patients with clinically stable COPD. As reported in our paper, increased circulating levels of leucocytes were observed in the subgroup of clinically stable COPD patients treated with oral corticosteroids compared with those who did not receive oral corticosteroids. However, comparison of subgroups of patients without corticosteroid treatment (oral or inhaled) with control subjects still revealed significantly increased leucocyte counts (data not shown). This indicates that the enhanced levels of circulating leucocytes are not solely due to corticosteroid use, and could be a marker of the systemic inflammatory process in COPD. In line with this hypothesis, Noguera et al reported enhanced circulating levels of polymorphonuclear cells in patients with stable COPD, none of whom had received steroids before the start of the study (oral or inhaled).

As discussed in our paper, administration of prednisolone during treatment of an exacerbation is most probably the cause for the observed rise in leucocyte counts. In order to determine the effect of exacerbation on leucocyte counts, studies are currently being performed in our hospital in which blood samples are collected before the start of treatment. COPD comprises a heterogeneous group of conditions characterised by chronic airflow limitation and destruction of lung parenchyma with clinical manifestations of dyspnoea, cough, sputum production, and impaired exercise tolerance. The definition of an acute exacerbation of COPD is still imprecise, and is generally based on varying combinations of symptoms. Rodriguez-Roisin et al suggested staging COPD exacerbations based on use of health care. They defined three levels of severity: mild, moderate and severe. During a mild exacerbation the patient has an increased need for medication which he/she can manage in their own environment; patients with a moderate exacerbation have an increased need for medication and need to seek additional medical assistance; the patient with a severe exacerbation recognises obvious and/or rapid deterioration in his/her condition requiring admission to hospital. Based on this definition, the patients included in our study were suffering from severe exacerbations of disease. Only limited information is so far available concerning (changes of) inflammatory markers during exacerbations. In our study the kinetics of pro- and anti-inflammatory markers have been analysed in patients with COPD during the first 7 days in hospital for an exacerbation of the disease. The results showed a significant decline in systemic levels of both CRP (at day 3) and LBP (at day 7) compared with day 1, whereas levels of the anti-inflammatory mediator soluble IL-1 receptor II dramatically increased (until day 5). This change in levels of inflammatory mediators may contribute to the clinical improvement of the patients. Additional studies are required to obtain more insight into the role of the inflammatory processes in the pathogenesis of exacerbations which could contribute to measurable parameters, in order to define the severity or outcome of disease more accurately.

References


Authors’ reply

Dr Kelly has some concerns about the levels of C reactive protein (CRP) in the healthy controls and patients with clinically stable COPD in our study. The control group used consisted of randomly selected subjects over 50 years of age and living in the same area as the patients. These subjects had no evidence of COPD based on questionnaires and lung function testing, did not exhibit any acute or chronic disease, and were not taking medication. Based on these selection criteria, this group was considered a healthy population control group. Although two of the 23 control subjects had enhanced CRP levels, the reason for which is unknown, they were not excluded in order to prevent bias. Non-parametric tests were used to compare the CRP levels between controls and COPD patients, and therefore the results are not affected by these two outliers. Concerning the diagnosis of COPD, all patients in our study underwent high resolution computed tomographic scanning to exclude the presence of bronchiectasis. There is increasing evidence that COPD is characterised by systemic effects which, among other factors, are reflected by enhanced circulating levels of inflammatory mediators in the circulation. In line with the observations of increased levels of the acute phase proteins CRP and lipopolysaccharide binding protein (LBP) in patients in a clinically stable condition, subgroups can be identified before the start of treatment.

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References