Butyrylcholinesterase K variant is genetically associated with late onset Alzheimer’s disease in Northern Ireland

S P McIlroy, V L S Crawford, K B Dynan, B M McGleenon, M D Vahidassr, J T Lawson, A P Passmore

Abstract
Alzheimer’s disease (AD) is a progressive neurodegenerative disorder that has been associated, sometimes controversially, with polymorphisms in a number of genes. Recently the butyrylcholinesterase K variant (BCHE K) allele has been shown to act in synergy with the apolipoprotein E e4 (APOE e4) allele to promote risk for AD. Most subsequent replicative studies have been unable to confirm these findings. We have conducted a case-control association study using a clinically well defined group of late onset AD patients (n=175) and age and sex matched control subjects (n=187) from the relatively genetically homogeneous Northern Ireland population to test this association. The BCHE genotypes of patients were found to be significantly different from controls ($\chi^2=23.68$, df=2, $p<0.001$). The frequency of the K variant allele was also found to differ significantly in cases compared to controls ($\chi^2=16.39$, df=1, $p<0.001$) leading to an increased risk of AD in subjects with this allele (OR=3.50, 95% CI 2.20-6.07). This risk increased in subjects 75 years and older (OR=5.50, 95% CI 2.56-11.87). At the same time the APOE e4 associated risk was found to decrease from 6.70 (95% CI 2.40-19.04) in 65-74 year olds to 3.05 (95% CI 1.34-6.95) in those subjects 75 years and older. However, we detected no evidence of synergy between BCHE K and APOE e4. The results from this study suggest that possession of the BCHE K allele constitutes a significant risk for AD in the Northern Ireland population and, furthermore, this risk increases with increasing age.


Keywords: Alzheimer’s disease; BCHE K; APOE

Alzheimer’s disease (AD) is a progressive neurodegenerative disorder, the incidence of which increases sharply in members of the population aged over 65 years. It is characterised by the extracellular accumulation of $\beta$-amyloid (A$\beta$) peptide in the form of amyloid plaques in the brains of affected subjects. It is characterised by the extracellular accumulation of $\beta$-amyloid (A$\beta$) peptide in the form of amyloid plaques in the brains of affected subjects. Although it is clear that a significant proportion of AD has a genetic basis, APOE is the only genetic factor that has been consistently implicated in non-familial late onset disease. However, the possession of the APOE4 allele is neither necessary nor sufficient for disease initiation/progression. This fact has prompted a search for other genetic factors that may confer susceptibility to AD.

One of the candidates that has been identified to date is the K variant of butyrylcholinesterase (BCHE). These authors reported that the BCHE K variant acted in synergy with APOE4 to increase the risk for AD, especially in subjects older than 75 years. This enzyme is an attractive candidate for involvement in AD as it has been hypothesised to participate in the transformation of A$\beta$ from an initially benign form to an eventually malignant form associated with neuritic tissue degeneration and clinical dementia. The activity of this enzyme has been shown to increase with age and be raised in AD. In addition, because BCHe blocks aggregation of A$\beta$ less aggressive long fibrils and possession of the K variant allele is known to result in a 30% reduction in serum cholinesterase activity, the BCHE K variant becomes even more attractive as a candidate risk factor for AD.

Although the findings of Lehmann et al have been replicated or partially replicated in some studies,10–11 a majority have failed to find evidence for association either alone or synergistically with APOE.12–15 Because the frequencies of the BCHE K variant differs in various ethnic populations4–7 and the activity of the enzyme has been shown to be high in an Irish population,9 we decided to test the association of the K variant with AD in a clinically well defined cohort of AD cases and controls from the relatively genetically homogeneous population in Northern Ireland.16

Materials and methods

SUBJECTS
Ethical approval for this study was obtained from the Research Ethics Committee, The Queen’s University of Belfast. Informed written consent was obtained from patients/carers and controls before collection of blood samples. Patients’ blood samples were obtained through The Memory Clinic, Belfast City Hospital and referral from the Alzheimer’s Disease Society, Northern Ireland. Control blood samples were obtained from non-demented healthy spouses and volunteers from retirement clubs throughout Northern Ireland. All patients and controls were white and ascertained to have at least parents and grandparents born in Northern Ireland to ensure ethnicity. Any patient with a family history of
dementia (greater than one first degree relative with any form of dementia) was excluded from the analysis. Diagnosis of probable AD was arrived at using the DSM IV (1994) and NINCDS-ADRDA criteria. Where possible, a CT scan was performed to aid diagnosis.

Age, sex, and ethnically matched volunteers were recruited as control subjects. Assessment of controls included a full medical history and a physical examination. Using the same criteria as was used for the AD patients, a comprehensive family history was taken to exclude those with a history of dementia. Cognitive function was assessed using Mini Mental State Examination (MMSE).

RESULTS

The patient group consisted of 62 males and 113 females with a mean age of 77.7 years (SD 5.94, range 67-94 years). Controls consisted of 58 males and 129 females with a mean age of 77.1 years (SD 6.04, range 67-96 years).

The frequency of the K variant allele was obtained by Cornfield's approximation. A full blood screen was performed as for the patient group. The BCHE K variant was detected by the "salting out" method. The BCHE K variant was detected by the amplification created restriction site method essentially as described and fragments were sized by reference to a sequencing ladder. APOE genotypes were assigned by PCR-RFLP, as previously described.

STATISTICAL ANALYSIS

Genomic DNA was extracted from peripheral blood leukocytes by the "salting out" method. The BCHE K variant was detected by the amplification created restriction site method essentially as described and fragments were sized by reference to a sequencing ladder. APOE genotypes were assigned by PCR-RFLP, as previously described.

Results

Distribution of genotypes was found to differ significantly in cases compared to control subjects (\(\chi^2=23.68, df=2, p<0.001\)) (table 1). The distribution of the genotypes in the control group did not differ from that predicted by Hardy-Weinberg equilibrium; however, the distribution of the genotypes in the cases differed from Hardy-Weinberg. This was not the result of technical error as controls were included in each experiment and in other analyses performed by this group, including APOE in this study, the distribution of genotypes in cases and controls did not differ from that predicted by Hardy-Weinberg equilibrium.

The distribution of BCHE genotypes in cases aged 65 to 74 differed significantly from the distribution in those aged 75 or greater (\(\chi^2=10.85, df=2, p=0.004\)), whereas the genotype distribution in controls did not differ between these age groups (\(\chi^2=2.74, df=2, p=0.25\)) (table 2). Furthermore, while the difference in the distribution of APOE genotypes in these two age groups approached but did not reach significance in the patient group (\(\chi^2=8.40, df=4, p=0.08\)), the distribution of the genotypes in controls between the two age groups was almost identical (\(\chi^2=1.98, df=4, p=0.74\)) (table 3).

The frequency of the K variant allele was also found to be significantly different in cases when compared to controls (\(\chi^2=16.39, df=1, p<0.001\)). Computation of the odds ratio as an estimate of relative risk for disease showed that subjects with the BCHE K variant allele were 3.50 (95% CI 2.20-6.07) times more likely to suffer from AD. Division of the cases and controls into two age groups as in the genotypes and stratifying for APOE \(e4\) showed that the odds ratio associated with the BCHE K variant increased from 1.26 (95% CI NS) in the 65 to 74 year olds (table 4) to 5.50 (95% CI 2.56-11.87) in the group aged 75 years or greater (table 5). At the same time the APOE \(e4\) associated odds ratio decreased from 6.70 (95% CI 2.40-19.04) in the younger group to 3.05 (95% CI 1.34-6.95) in the older group.

The odds ratio for possession of both risk factors increased from 11.26 (95% CI 2.41-68.91) in the under 75 year age group (table 4) to 24.35 (95% CI 6.67 - 88.87) in the subjects of 75 years and older (table 5). However, although 11 cases compared to three controls possessed both risk factors in the younger age group and 26 cases compared to three controls

<table>
<thead>
<tr>
<th>Table 1 Overall distribution of BCHE genotypes</th>
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<tr>
<td></td>
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<td>NN</td>
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<td>NK</td>
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<td>KK</td>
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\(\chi^2=23.68, df=2, p<0.001\).

<table>
<thead>
<tr>
<th>Table 2 Distribution of BCHE genotypes in subjects aged between 65 and 74 years and 75 years and older</th>
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<tbody>
<tr>
<td>Cases</td>
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<td>-------------</td>
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<td>Cases</td>
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</table>

\(\chi^2=1.98, df=4, p=0.74\).
possessed both risk factors in the older age group, we found little evidence of interaction between APOE ε4 and BCHE K either by logistic regression (χ²= 0.13, df=1, p>0.1) or synergy factor analysis (z=0.44, p>0.1).

Discussion

This case-control association study showed a significant difference between the BCHE genotypes of late onset AD cases and control subjects. However, in agreement with two other studies, we noticed that this effect was observed that the APOE ε4 and BCHE K either by logistic regression (χ²= 0.13, df=1, p>0.1) or synergy factor analysis (z=0.44, p>0.1).

Table 4  Carrier odds ratios in the 65 to 74 year subgroup

<table>
<thead>
<tr>
<th>BCHE K</th>
<th>APOEε4</th>
<th>Cases</th>
<th>Controls</th>
<th>OR (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>− −</td>
<td>− −</td>
<td>14</td>
<td>43</td>
<td>Reference</td>
</tr>
<tr>
<td>− +</td>
<td>24</td>
<td>11</td>
<td>6.70 (2.40–19.04)</td>
<td></td>
</tr>
<tr>
<td>+ +</td>
<td>9</td>
<td>22</td>
<td>1.26 (NS)</td>
<td></td>
</tr>
<tr>
<td>+ +</td>
<td>11</td>
<td>3</td>
<td>11.26 (2.41–68.91)</td>
<td></td>
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</tbody>
</table>

Table 5  Carrier odds ratios in the 75 years and older subgroup

<table>
<thead>
<tr>
<th>BCHE K</th>
<th>APOEε4</th>
<th>Cases</th>
<th>Controls</th>
<th>OR (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>− −</td>
<td>− −</td>
<td>21</td>
<td>59</td>
<td>Reference</td>
</tr>
<tr>
<td>− +</td>
<td>5</td>
<td>23</td>
<td>3.05 (1.34–6.98)</td>
<td></td>
</tr>
<tr>
<td>+ +</td>
<td>45</td>
<td>23</td>
<td>5.50 (2.56–11.87)</td>
<td></td>
</tr>
<tr>
<td>+ +</td>
<td>26</td>
<td>3</td>
<td>24.35 (6.67–88.87)</td>
<td></td>
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</table>

maximum effect before the age of 70 years. In addition, Scacchi et al reported no difference between case and control frequencies of subjects older than 80 years even though the APOE ε4 associated risk was present in subjects less than 80 years of age. These reports indicate that it is important to identify a factor that may influence AD risk in the very old and the results of this study suggest that the BCHE K variant is a reasonable candidate.

In conclusion we report that the BCHE K variant is a significant risk factor for late onset AD in the Northern Ireland population. The odds ratio associated with this risk increases in people older than 75 years; however, we were unable to show any synergy with APOE ε4.

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