Objective: To study the Ala53Thr and Ala30Pro mutations of the α-synuclein gene in a large number of Chinese patients with Parkinson disease (PD) as well as controls.

Methods: We recruited 183 Chinese patients with sporadic PD, 17 with younger-onset PD (onset age < 50 years), and 7 with PD and a positive family history as well as 227 unaffected Chinese control subjects from the outpatient departments of 2 major hospitals in Hong Kong. All subjects were assessed for the diagnosis of PD by a consultant neurologist or geriatrician. Subjects were interviewed with a standard questionnaire that also questioned for family history. Venous blood samples were obtained from the subjects and genomic DNA was extracted and studied for the presence of Ala53Thr mutation in exon 4 and Ala30Pro mutation in exon 3 of the α-synuclein gene using a polymerase chain reaction restriction fragment length polymorphism method.

Results: None of the Chinese PD patients or controls had either the Ala53Thr (exon 4) or Ala30Pro (exon 3) mutation of the α-synuclein gene.

Conclusion: We failed to discover Ala53Thr or Ala30Pro mutations in a large number of Chinese patients with PD and control subjects, adding to the emerging consensus that variations in the α-synuclein gene are associated with PD in few families worldwide.

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SUBJECTS AND METHODS

Patients and controls were recruited from the outpatient departments of the United Christian and Prince of Wales hospitals in Hong Kong. All subjects were assessed by a consultant neurologist or geriatrician for the diagnosis of PD. The criteria for diagnosis is in accordance with those described by Maraganore et al. In the selection of controls, 10-year age group, sex, and locality of residence were taken into consideration to make the control group as comparable to the PD group as possible. (For more details of the study, refer to Chan et al.)

The subjects were interviewed with a standard questionnaire and were questioned for family history of PD. If there was a family history, further details regarding which relatives were affected (ie, parents, siblings, uncles, aunts, nephews, or nieces) were obtained. Venous blood samples were taken from the subjects, and genomic DNA was extracted by the salting out method. The patients and controls were screened for the presence of the Ala53Thr mutation in exon 4 of the α-synuclein gene (the Contursi mutation) using the polymerase chain reaction (PCR) Ysp45I restriction fragment length polymorphism (RFLP) method described by Polymeropoulos et al. DNA that was obtained from an affected Contursi kindred member and contained the Ala53Thr mutation was kindly donated by Polymeropoulos et al. and was used as a positive control. The primer sequences used for the DNA amplification were 5'-GCTAATCAGCAATTTAGCTAG-3' (forward) and 5'-GATAATTTCTTTAGTGCTACG-3' (reverse). The amplification reaction was performed in a 50-µl volume that contained 100 ng of template DNA, 300 ng of each primer, 2 mmol/L magnesium chloride, 0.24 mmol/L deoxynucleotidetriphosphates, and 2 U of DNA polymerase (Red Hot DNA polymerase; Advanced Biotechnologies, Surrey, England). The samples were subjected to an initial denaturation at 94°C for 5 minutes followed by 17 cycles of denaturation at 94°C for 15 seconds, annealing at 50°C for 30 seconds, and extension at 72°C for 30 seconds. The second phase of the amplification reaction involved 18 cycles of denaturation (94°C for 15 seconds), annealing (50°C for 30 seconds), and extension (72°C for 1 minute 30 seconds). The 216-base pair (bp) PCR product was digested with the restriction enzyme Ysp45I (New England Bio Labs, Beverly, Mass) according to the supplier's protocol, resolved on a 3% agarose gel (agarose 1000; Life Technologies, Melbourne, Australia), and visualized using ethidium bromide staining. Mutant alleles yield restriction fragments of 128 and 88 bp using this technique (Figure).

The Ala30Pro mutation in exon 3 of the α-synuclein gene was examined using the MvaI-based PCR-RFLP method previously described by Kruger et al. In this case, the forward and reverse primers were 5'-AAGTGATTTATGTTTCC-3' and 5'-AATGACAATTGGGGTTACCC-3', respectively. The PCR reactions (25 µL) contained 100 ng of template DNA, 20 pmol/L of each primer, 2.1 mmol/L magnesium chloride, 0.225 mmol/L deoxynucleotidetriphosphates, and 1 U of DNA polymerase. The samples were subjected to an initial denaturation at 95°C for 5 minutes followed by 35 cycles of denaturation at 95°C for 1 minute, annealing at 54°C for 1 minute, and extension at 72°C for 1 minute. A final extension period of 72°C for 5 minutes completed the reaction. The 192-bp PCR products were digested with MvaI restriction enzyme and resolved using 8% polyacrylamide gel electrophoresis followed by silver staining. Alleles containing the Ala30Pro mutation yield 136- and 56-bp fragments following MvaI digestion.

COMMENT

The failure to discover the Ala53Thr or Ala30Pro mutations in a large number of Chinese patients with PD and control subjects adds to the emerging consensus that variations in the α-synuclein gene are associated with PD in few families worldwide. Moreover, studies have thus far failed to reveal other mutations in the coding region of the α-synuclein gene in large groups of patients with familial and sporadic PD. Studies are yet to determine whether any variations in the regulatory or intronic regions of the α-synuclein gene are associated with PD. Efforts may be better spent in elucidating the mechanism by which the Ala53Thr and Ala30Pro variants of α-synuclein lead to neurodegeneration in individuals possessing these mutations. Such mechanistic information may suggest other candidate genes with causative variants potentially more common in patients with PD worldwide than the Ala53Thr or Ala30Pro variants of α-synuclein.

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Corresponding author: Daniel K. Y. Chan, MBBS, FRACP, Department of Geriatric Medicine, Prince of Wales Hospital, Level 3, High Street Building, High Street, Randwick NSW 2031, Sydney, Australia.
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