

No association between the tumor necrosis factor- α gene promoter polymorphisms and schizophrenia in a Japanese population

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Abstract

Tumor necrosis factor- α (TNF- α) is a pleiotrophic cytokine and exerts neuroprotective and neurodegenerative effects in brain. Several studies have indicated that TNF- α is likely related to the pathogenesis of schizophrenia. Recent genetic investigations have revealed that a TNF- α gene promoter polymorphism (–G308A) is associated with schizophrenia, although negative findings have also been reported. To assess whether the TNF- α gene promoter variants including –G308A could be implicated in vulnerability to schizophrenia, we conducted a case-control association analysis (265 cases and 424 controls) and the transmission disequilibrium test (TDT) analysis (83 trios) for four polymorphisms (–G238A, –G308A, –C857T and –T1031C) in Japanese subjects. In a case-control analysis, there was no significant association between the promoter polymorphisms or haplotypes in the TNF- α gene and schizophrenia. In the TDT analysis, we also did not observe transmission distortion. Our results suggest that the above four polymorphisms in the promoter region of the TNF- α gene appear not to confer increased susceptibility for schizophrenia in a Japanese population.

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1. Introduction

Schizophrenia is a complex genetic disorder and affects approximately 1% of the population worldwide. The pathogenesis of schizophrenia is still unclear; however, cytokines might be implicated in the etiology or

pathology of schizophrenia (for review, Nawa et al., 2000). Tumor necrosis factor- α (TNF- α), a pleiotrophic cytokine, exerts neuroprotective and neurodegenerative effects in brain (for review, Venters et al., 2001). Several studies have shown that blood concentrations and in vitro production of TNF- α were significantly higher in patients with schizophrenia than in healthy controls (Monteleone et al., 1997; Kowalski et al., 2001; Theodoropoulou et al., 2001), whereas some studies failed to find this increase (Haack et al., 1999; Erbağci et al., 2001). Buka et al. (2001) have reported that elevated TNF- α levels of maternal serum at the time

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Table 1
Primers and probes used in genotyping

| SNP | Primers | Probes |
|---------|---|---|
| –G238A | 5'-CAGTCAGTGGCCCAGAAGAC-3' 5'-CCCTCACACTCCCCATCCT-3' | 5'-VIC-CTCGGAATCGGAGCAG-NFQ-3' 5'-FAM-CTCGGAATCAGAGCAG-NFQ-3' |
| –G308A | 5'-CCAAAAGAAATGGAGGCAATAGGTT-3' 5'-GGACCCTGGAGGCTGAAC-3' | 5'-VIC-CCCGTCCCCATGCC-NFQ-3' 5'-FAM-CCCGTCCCTCATGCC-NFQ-3' |
| –C857T | 5'-GGGCTATGGAAGTCGAGTATGG-3' 5'-GTCCTGGAGGCTCTTTCACT-3' | 5'-VIC-CCCTGTCTTCGTTAAGG-NFQ-3' 5'-FAM-CCTGTCTTCATTAAGG-NFQ-3' |
| –T1031C | 5'-GTGAGGCCGCCAGACT-3' 5'-GCCCCCTCCAGACCCTGA-3' | 5'-VIC-CTTTTCCTTCGTCTTCTC-NFQ-3' 5'-FAM-TTTTCCTTCATCTTCTC-NFQ-3' |

of birth were associated with schizophrenia and related psychotic disorders in offspring, although Brown et al. (2004) found a significant association between maternal interleukin-8 but not TNF- α levels during the second trimester and the risk of schizophrenia spectrum disorders in offspring. Interestingly, Skurkovich et al. (2003) reported a case of schizophrenia whose negative symptoms improved with antibodies to TNF- α and to interferon- γ . Thus, these findings suggest that cytokines, including TNF- α , are likely related to the pathogenesis of schizophrenia.

Wilson et al. (1997) demonstrated that the minor A-allele of a polymorphism at the –308 position in the promoter region of the TNF- α gene, located on chro-

mosome 6p21.1–21.3, is a much more powerful transcriptional activator than the major G-allele. Recently, several studies have shown that the –G308A polymorphism of the TNF- α gene is associated with schizophrenia (Boin et al., 2001; Meira-Lima et al., 2003; Schwab et al., 2003; Tan et al., 2003). However, other studies have failed to confirm this association (Riedel et al., 2002; Handoko et al., 2003; Pae et al., 2003; Tsai et al., 2003; Duan et al., 2004; Hashimoto et al., 2004; Hänninen et al., 2005; Kampman et al., 2005; Pae et al., 2006). This inconsistency requires further investigations. Therefore, we performed a case-control association study and the transmission disequilibrium test (TDT) analysis in Japanese subjects to assess whether

Table 2
Genotype and allele frequencies of the TNF- α gene promoter region polymorphisms in cases and controls

| SNP | Genotype (%) | | | P^a | Allele (%) | | P^a |
|------------------------|--------------|------------|----------|-------|------------|------------|-------|
| –G238A | G/G | G/A | A/A | | G | A | |
| Cases | 259 (97.7) | 6 (2.3) | 0 (0.0) | 1.00 | 524 (98.9) | 6 (1.1) | 0.64 |
| With family history | 110 (99.1) | 1 (0.9) | 0 (0.0) | 0.58 | 221 (99.5) | 1 (0.5) | 0.32 |
| Without family history | 149 (96.8) | 5 (3.2) | 0 (0.0) | 0.83 | 303 (98.4) | 5 (1.6) | 1.00 |
| Controls | 412 (97.2) | 11 (2.6) | 1 (0.2) | | 835 (98.5) | 13 (1.5) | |
| –G308A | G/G | G/A | A/A | | G | A | |
| Cases | 258 (97.4) | 7 (2.6) | 0 (0.0) | 0.66 | 523 (98.7) | 7 (1.3) | 0.66 |
| With family history | 107 (96.4) | 4 (3.6) | 0 (0.0) | 1.00 | 218 (98.2) | 4 (1.8) | 1.00 |
| Without family history | 151 (98.0) | 3 (2.0) | 0 (0.0) | 0.42 | 305 (99.0) | 3 (1.0) | 0.43 |
| Controls | 409 (96.5) | 15 (3.5) | 0 (0.0) | | 833 (98.2) | 15 (1.8) | |
| –C857T | C/C | C/T | T/T | | C | T | |
| Cases | 184 (69.4) | 73 (27.6) | 8 (3.0) | 0.20 | 441 (83.2) | 89 (16.8) | 0.08 |
| With family history | 78 (70.3) | 28 (25.2) | 5 (4.5) | 0.50 | 184 (82.9) | 38 (17.1) | 0.26 |
| Without family history | 106 (68.8) | 45 (29.2) | 3 (2.0) | 0.17 | 257 (83.4) | 51 (16.6) | 0.13 |
| Controls | 271 (63.9) | 130 (30.7) | 23 (5.4) | | 672 (79.2) | 176 (20.8) | |
| –T1031C | T/T | T/C | C/C | | T | C | |
| Cases | 188 (71.0) | 74 (27.9) | 3 (1.1) | 0.43 | 450 (84.9) | 80 (15.1) | 1.00 |
| With family history | 79 (71.2) | 31 (27.9) | 1 (0.9) | 0.67 | 189 (85.1) | 33 (14.9) | 1.00 |
| Without family history | 109 (70.8) | 43 (27.9) | 2 (2.3) | 0.67 | 261 (84.7) | 47 (15.3) | 0.93 |
| Controls | 306 (72.2) | 108 (25.5) | 10 (2.3) | | 720 (84.9) | 128 (15.1) | |

^a Compared with control group.

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