Supplementary Figures

Figure S1.A. Manhattan plot of the FTND meta-analysis. The red line represents the genome-wide significance threshold \( (P < 5 \times 10^{-8}) \), while the blue line represents suggestive association threshold \( (P < 5 \times 10^{-5}) \).
Figure S1.B. QQ plot of the FTND meta-analysis.
Figure S1.C. Manhattan plot of the TTFC meta-analysis. The red line represents the genome-wide significance threshold ($P < 5 \times 10^{-8}$), while the blue line represents suggestive association threshold ($P < 5 \times 10^{-5}$).
Figure S1.D. QQ plot of the TTFC meta-analysis.
Figure S2. Chromosomal regions with markers reaching genome-wide significance in FTND and TTFC meta-analyses. A. CHRNA5-CHRNA3-CHRNB4 region on chromosome 15 for FTND; B. CHRNA5-CHRNA3-CHRNB4 region on chromosome 15 for TTFC; C. AA333164 region on chromosome 14 for TTFC; D. SORB2 region on chromosome 4 for TTFC; E. BG182718 region on chromosome 14 for TTFC.