

Supplementary Material

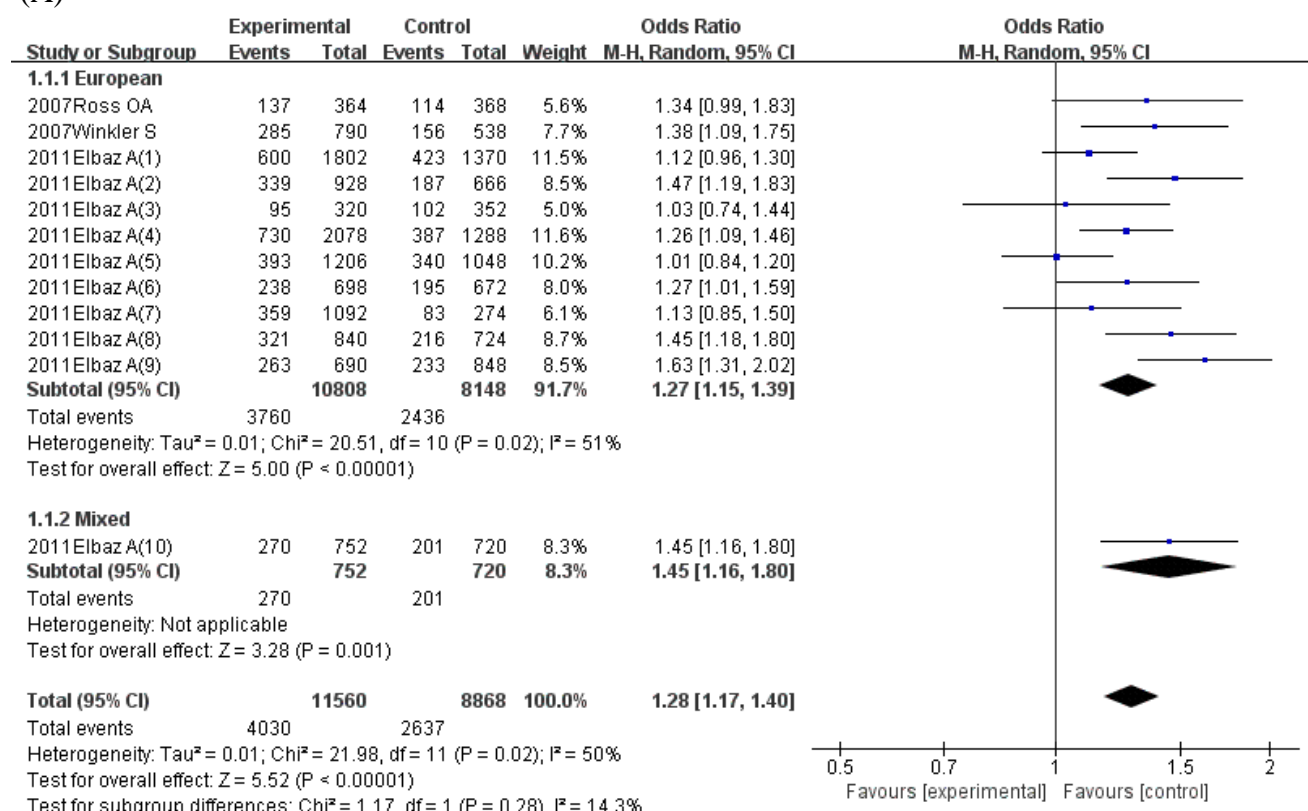
A comprehensive analysis of the association between *SNCA* polymorphisms with the risk of Parkinson's disease

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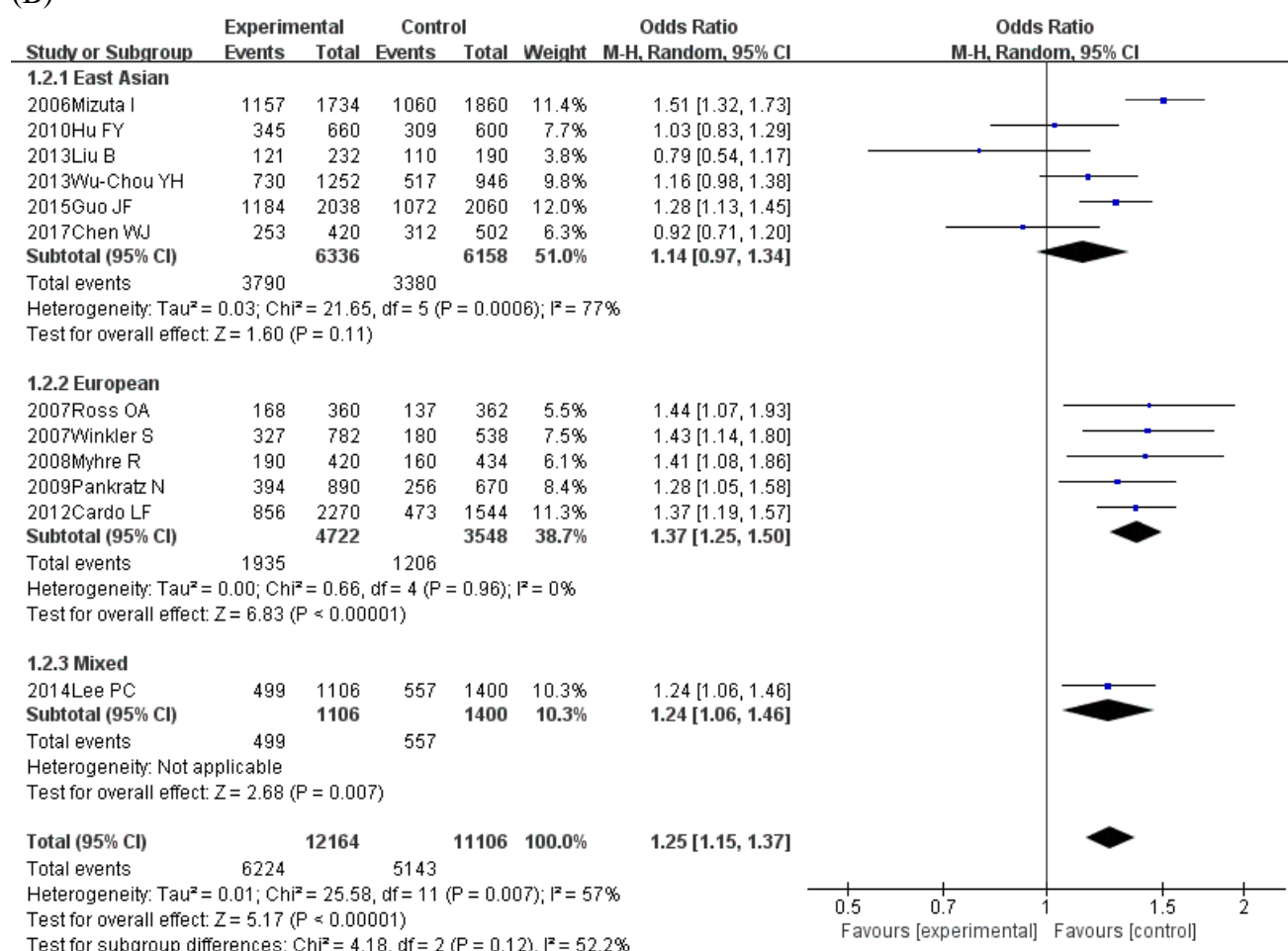
[†] These authors have contributed equally to this work and are co-first authors.

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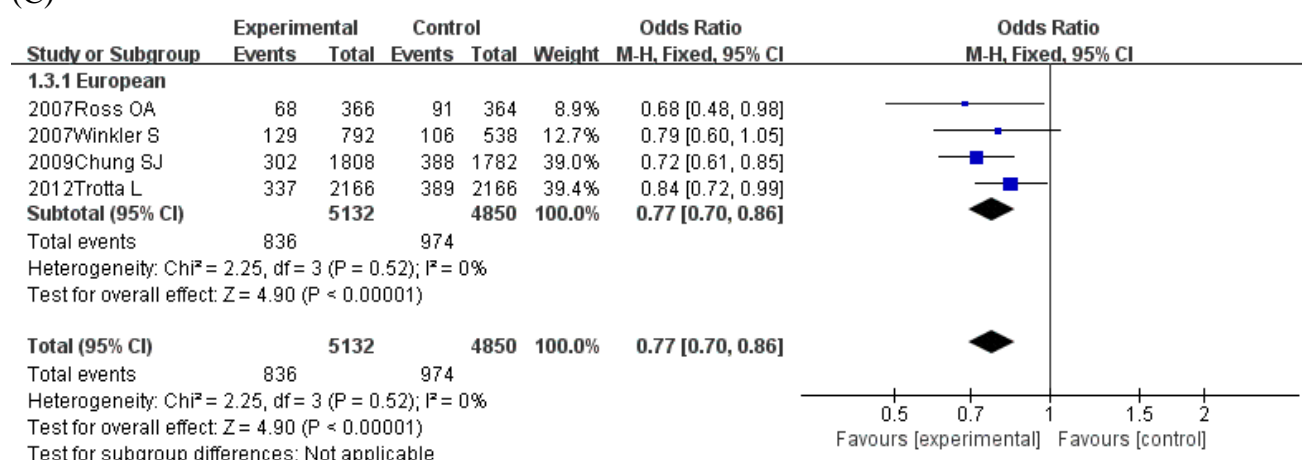
(A)



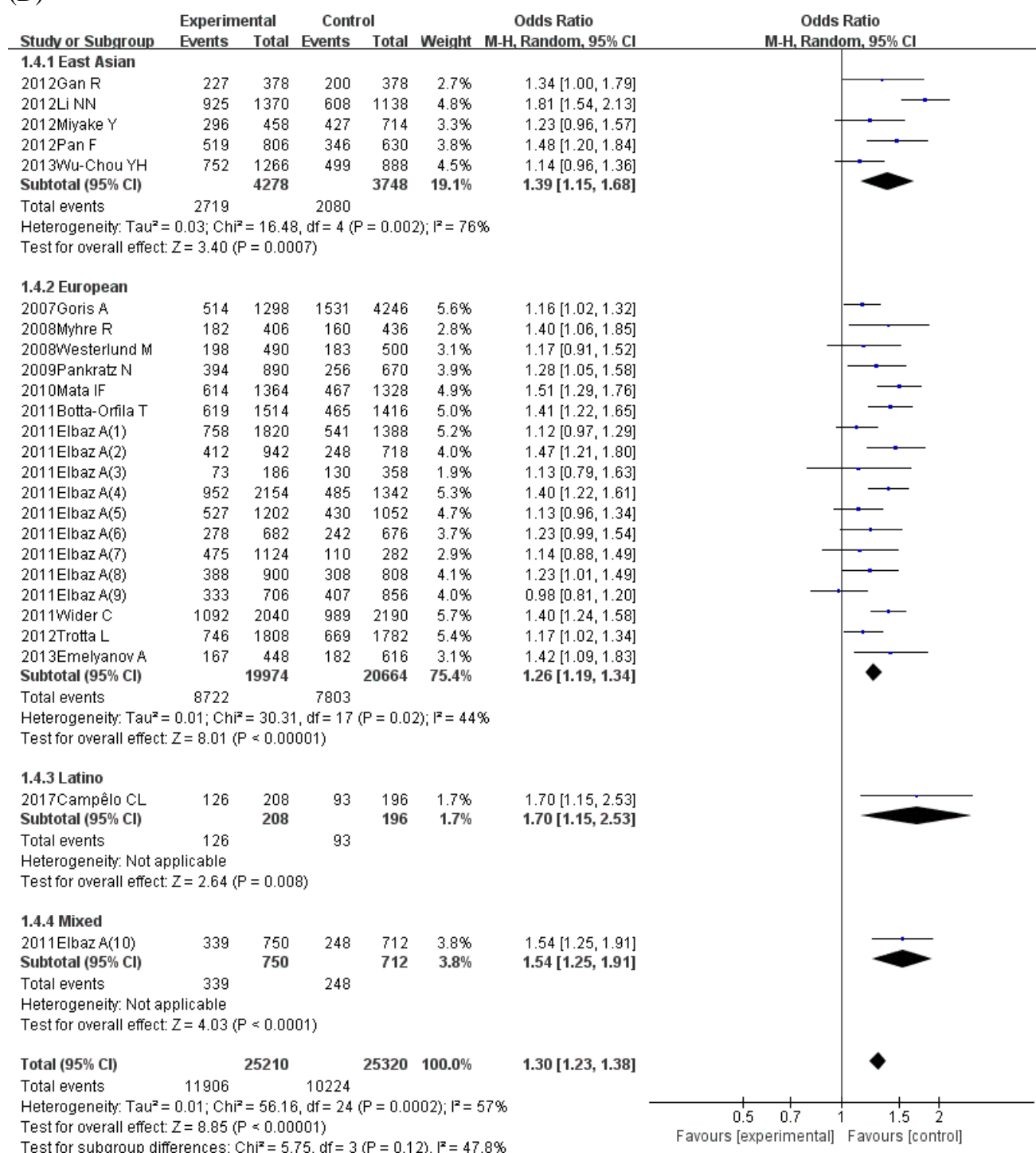
(B)



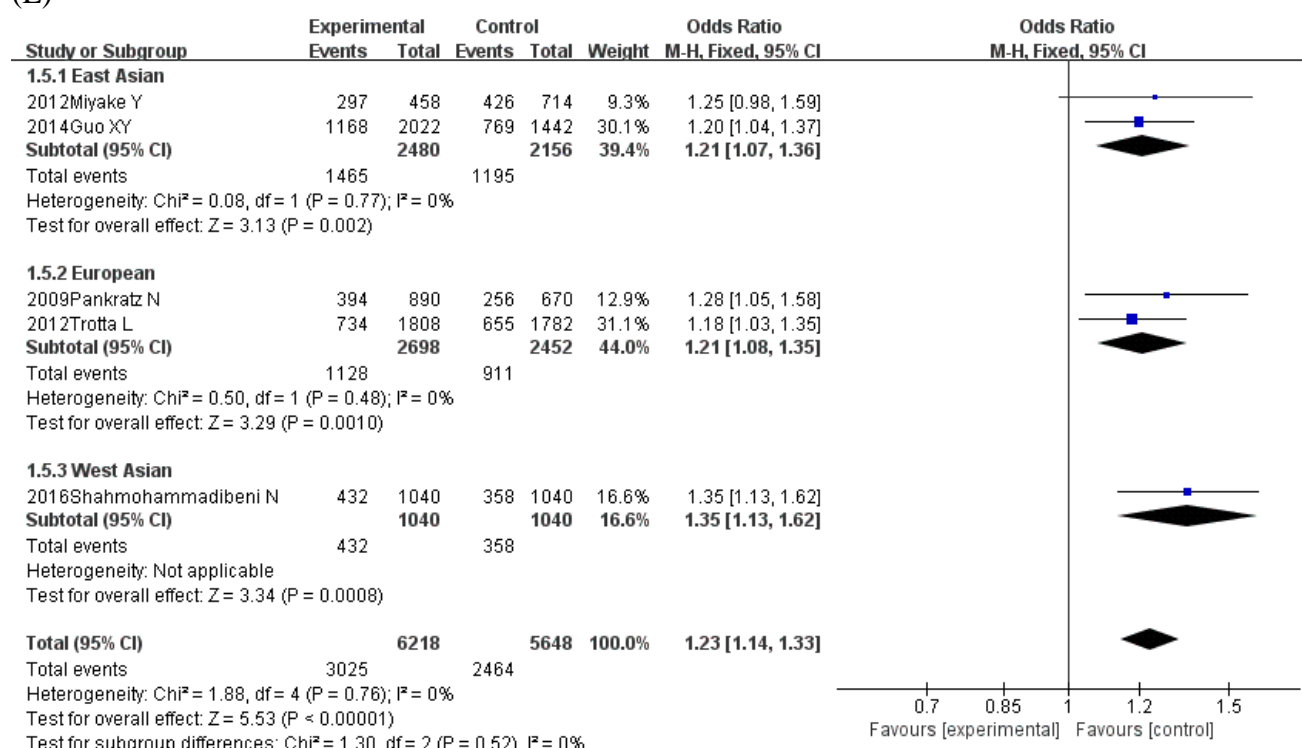
(C)



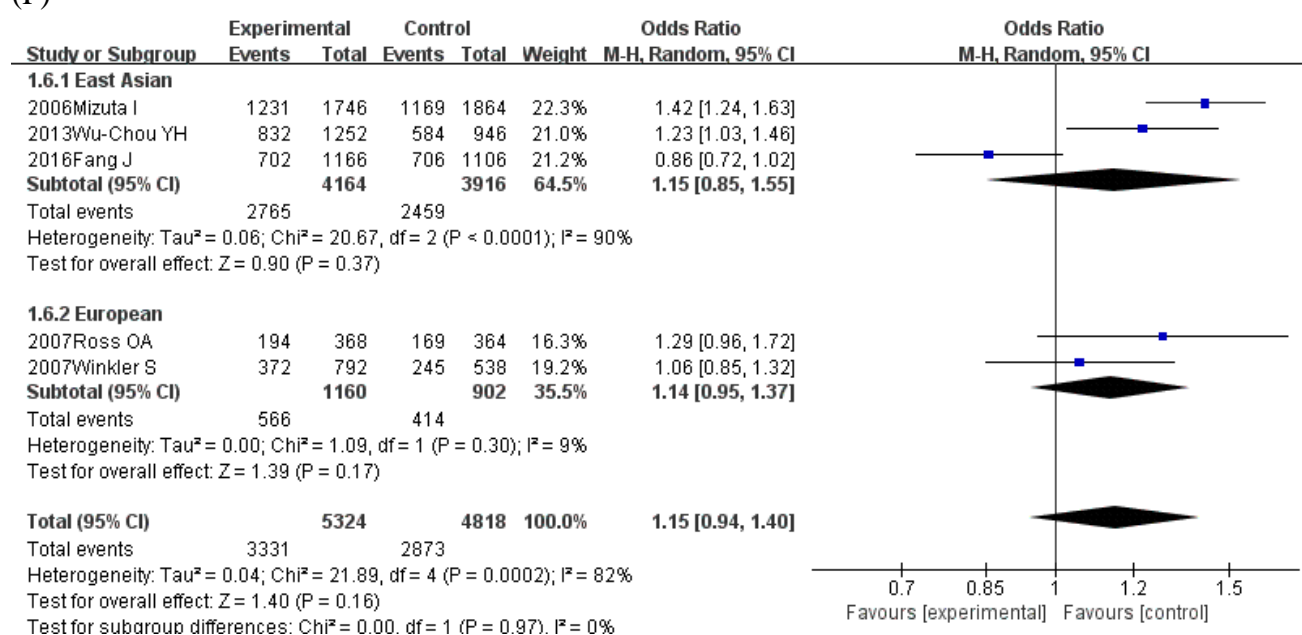
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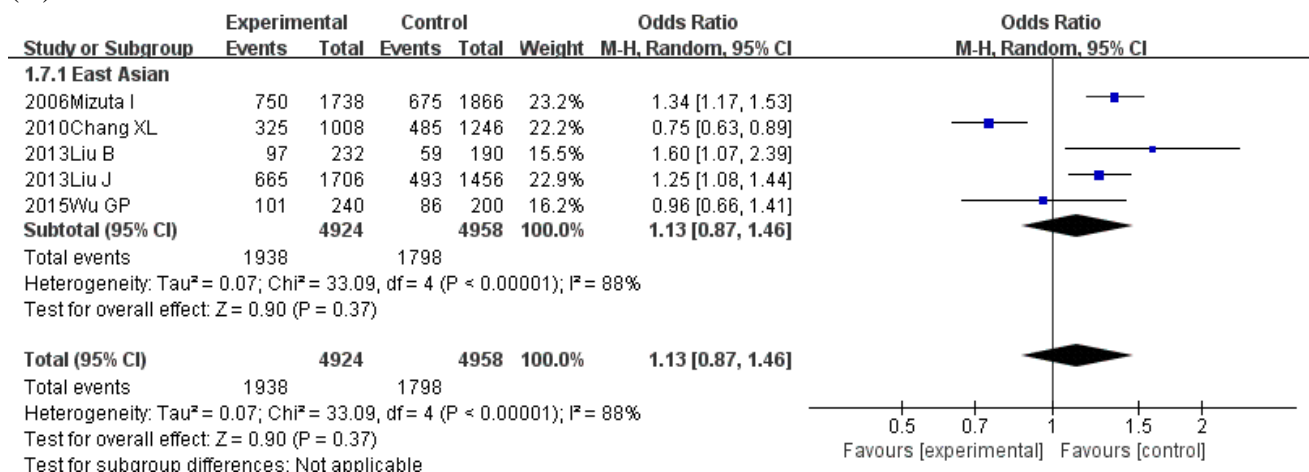
(E)



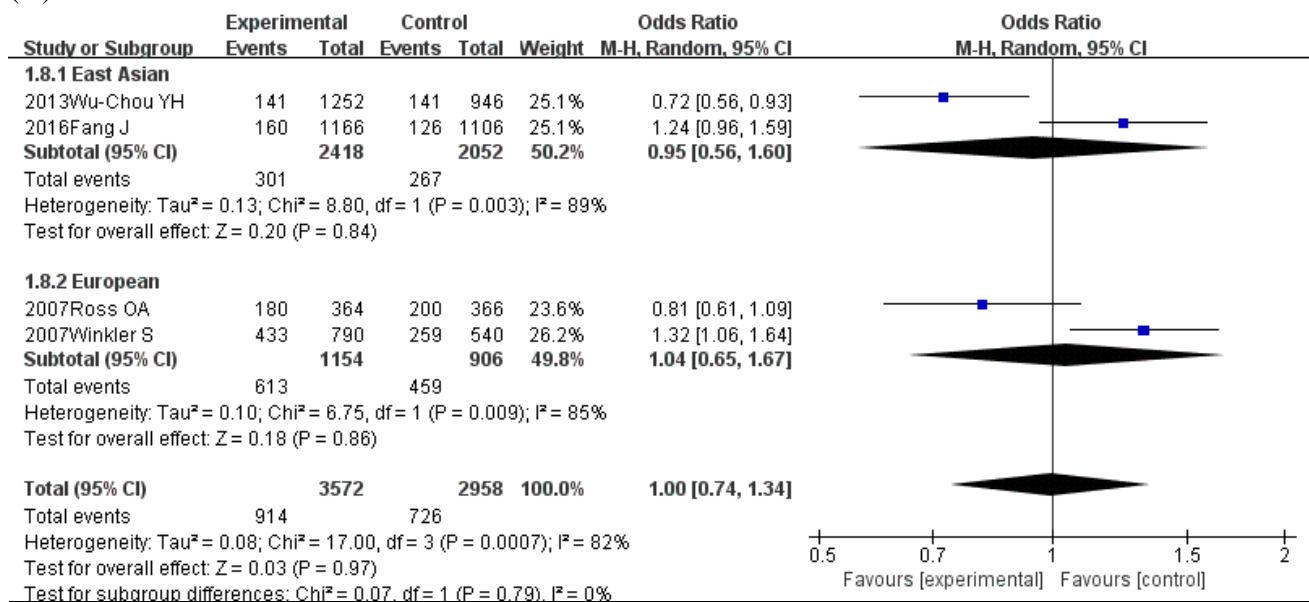
(F)



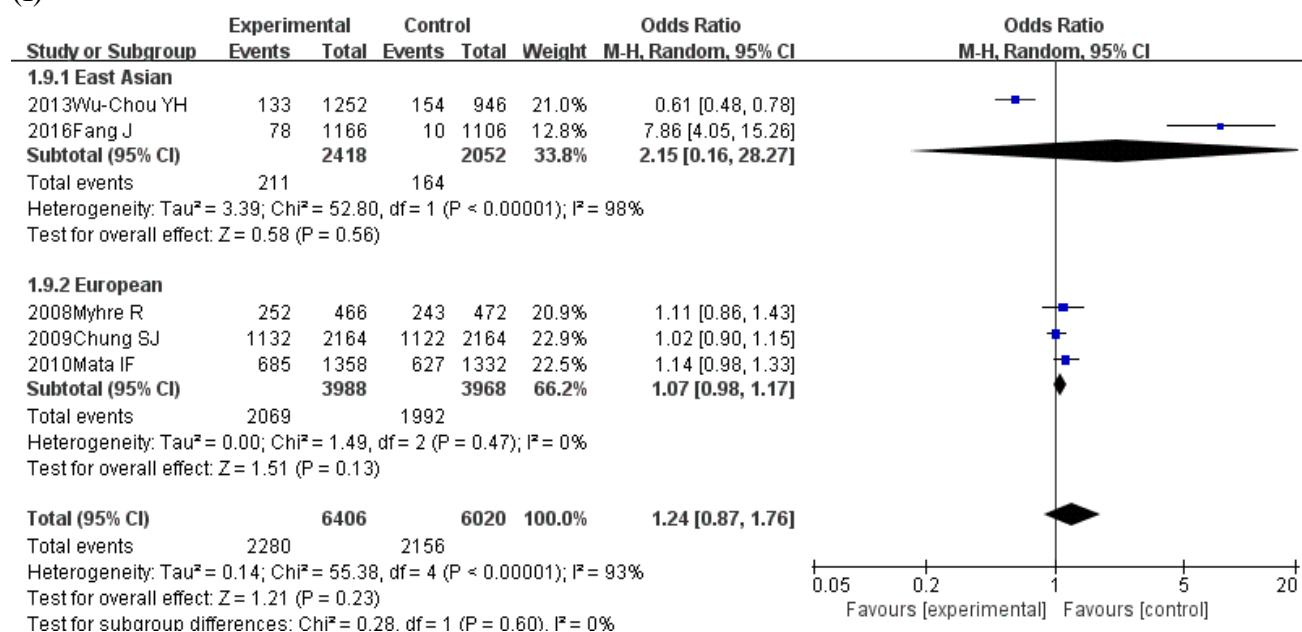
(G)



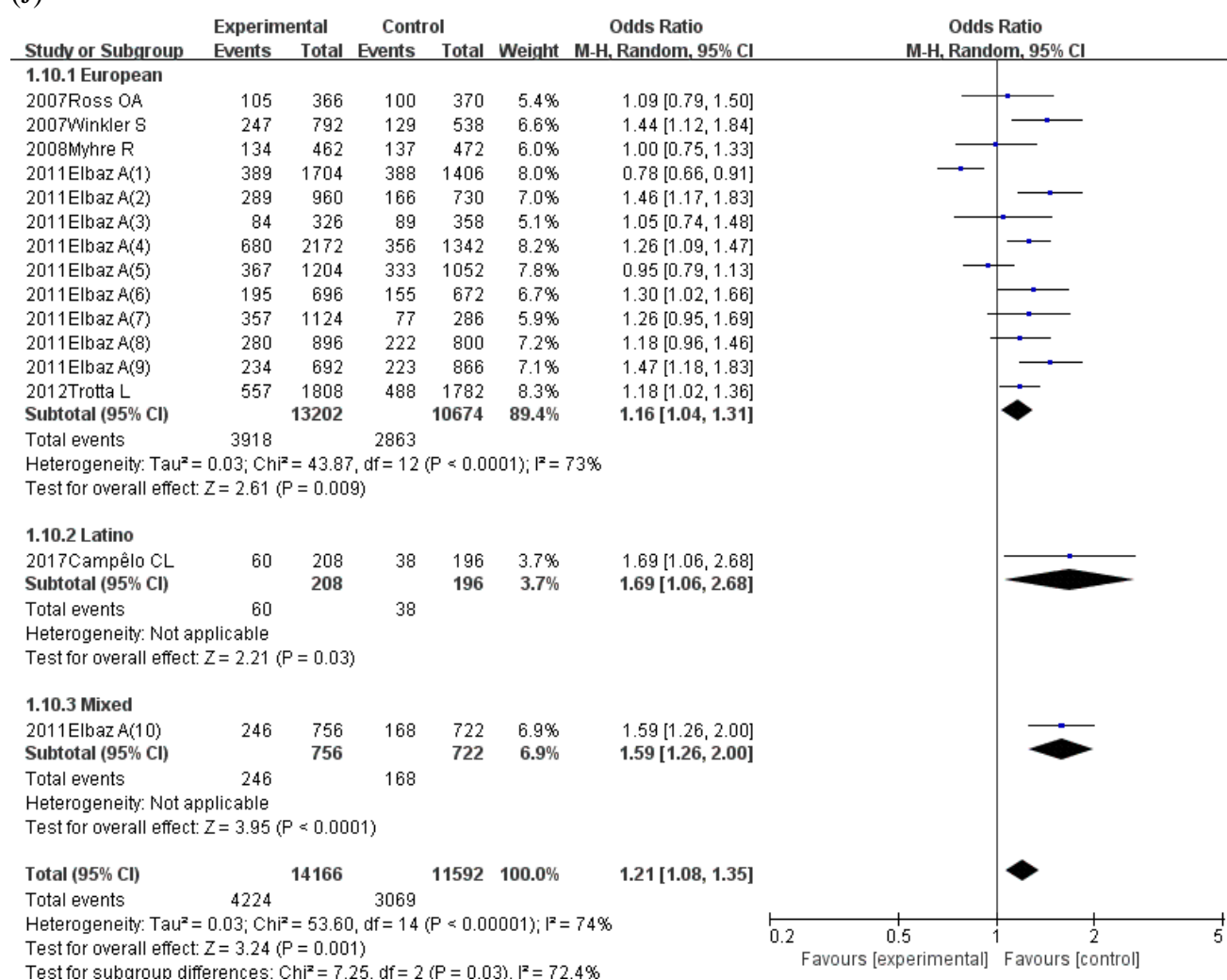
(H)



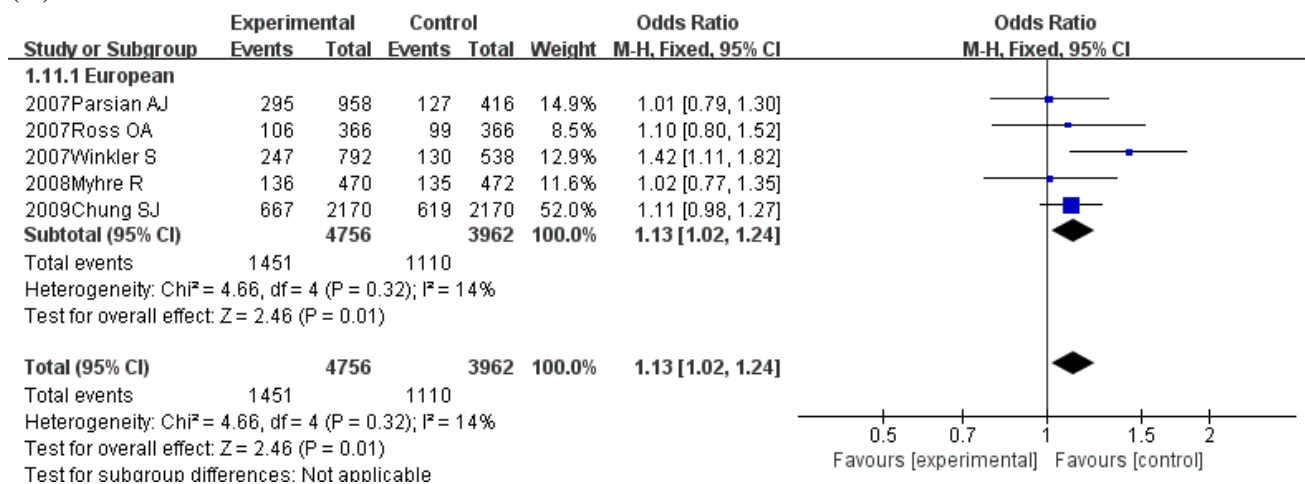
(I)



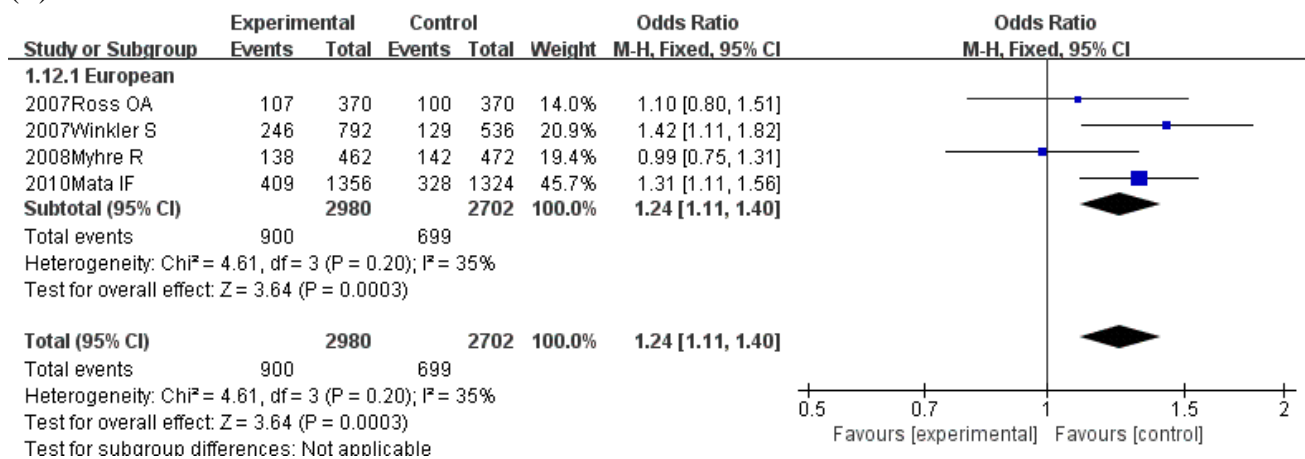
(J)



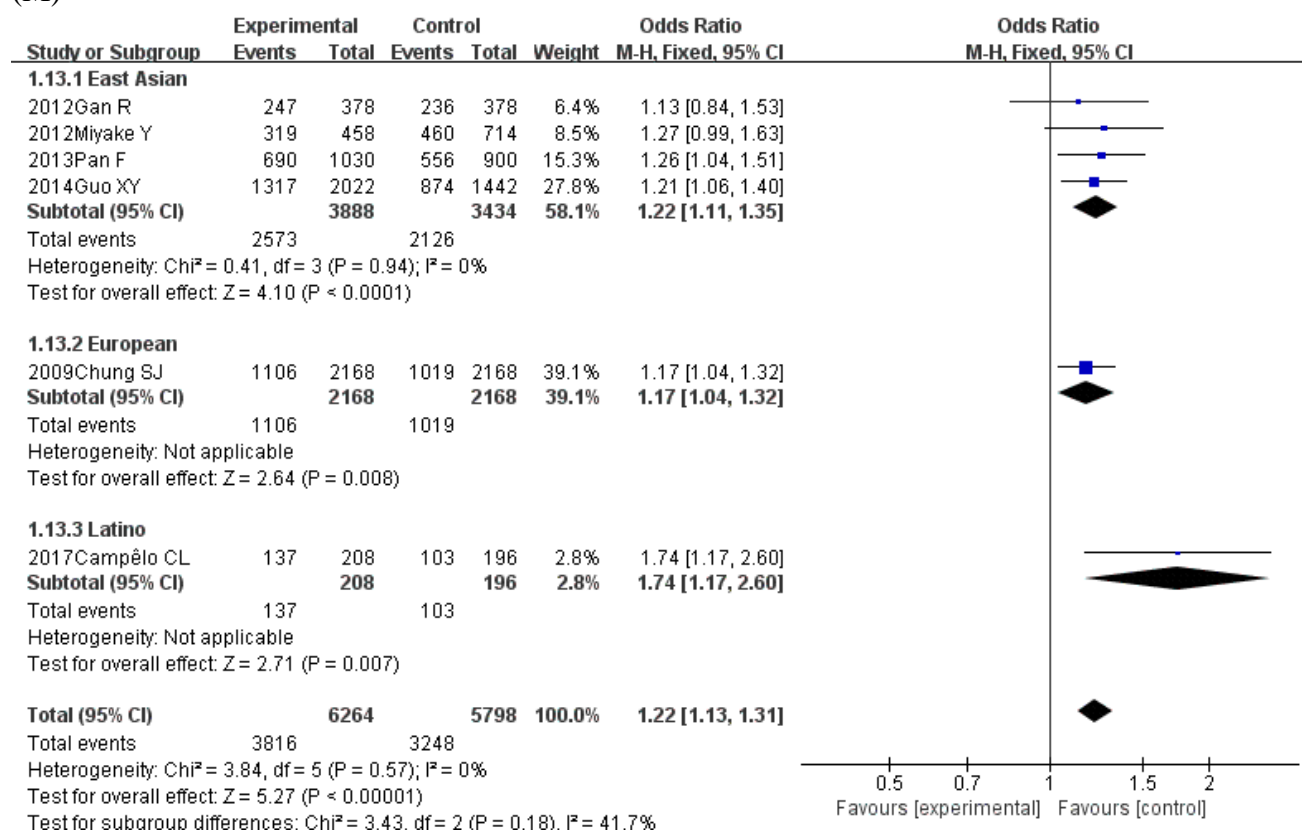
(K)



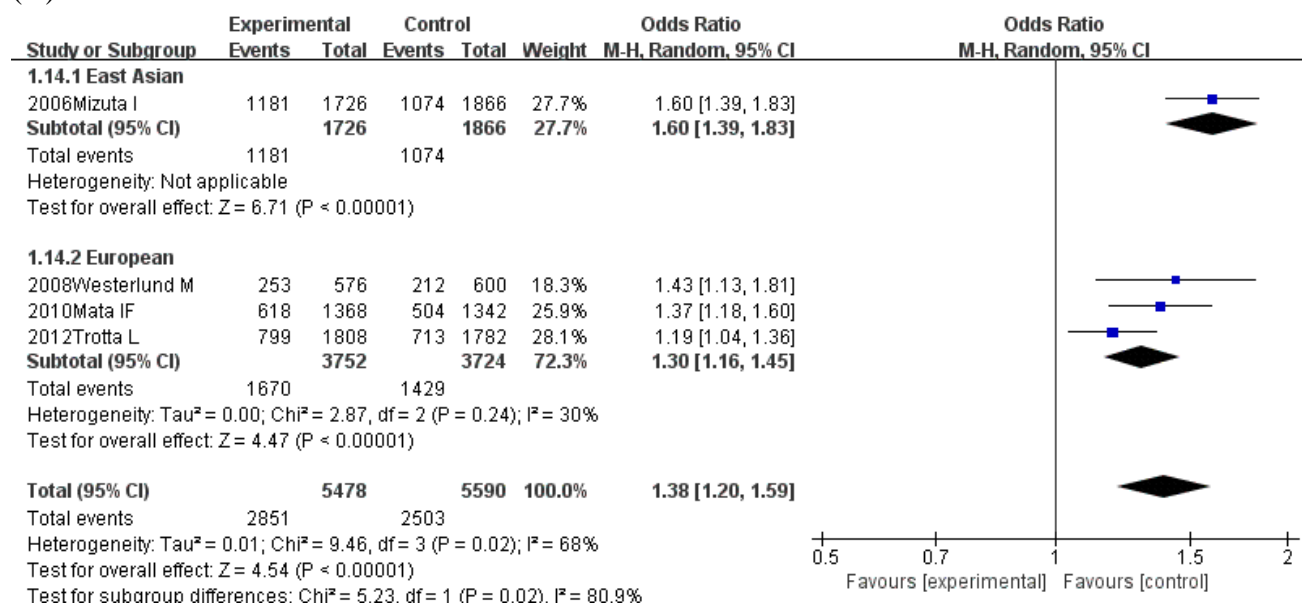
(L)



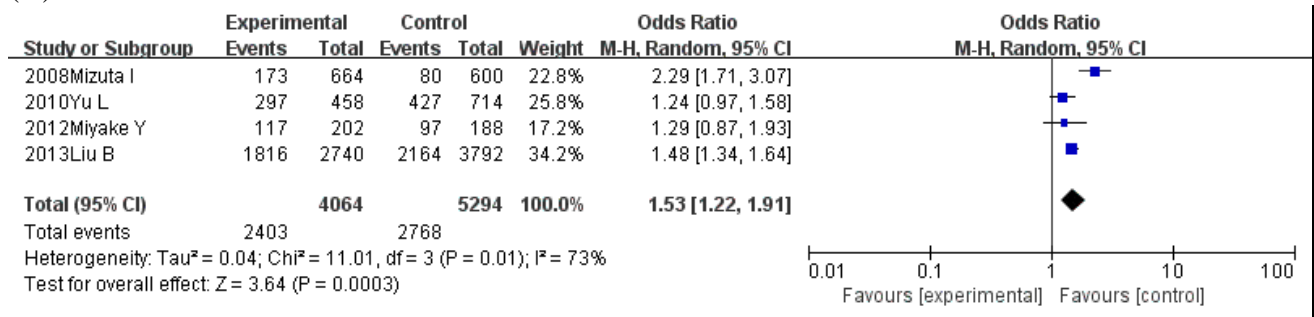
(M)



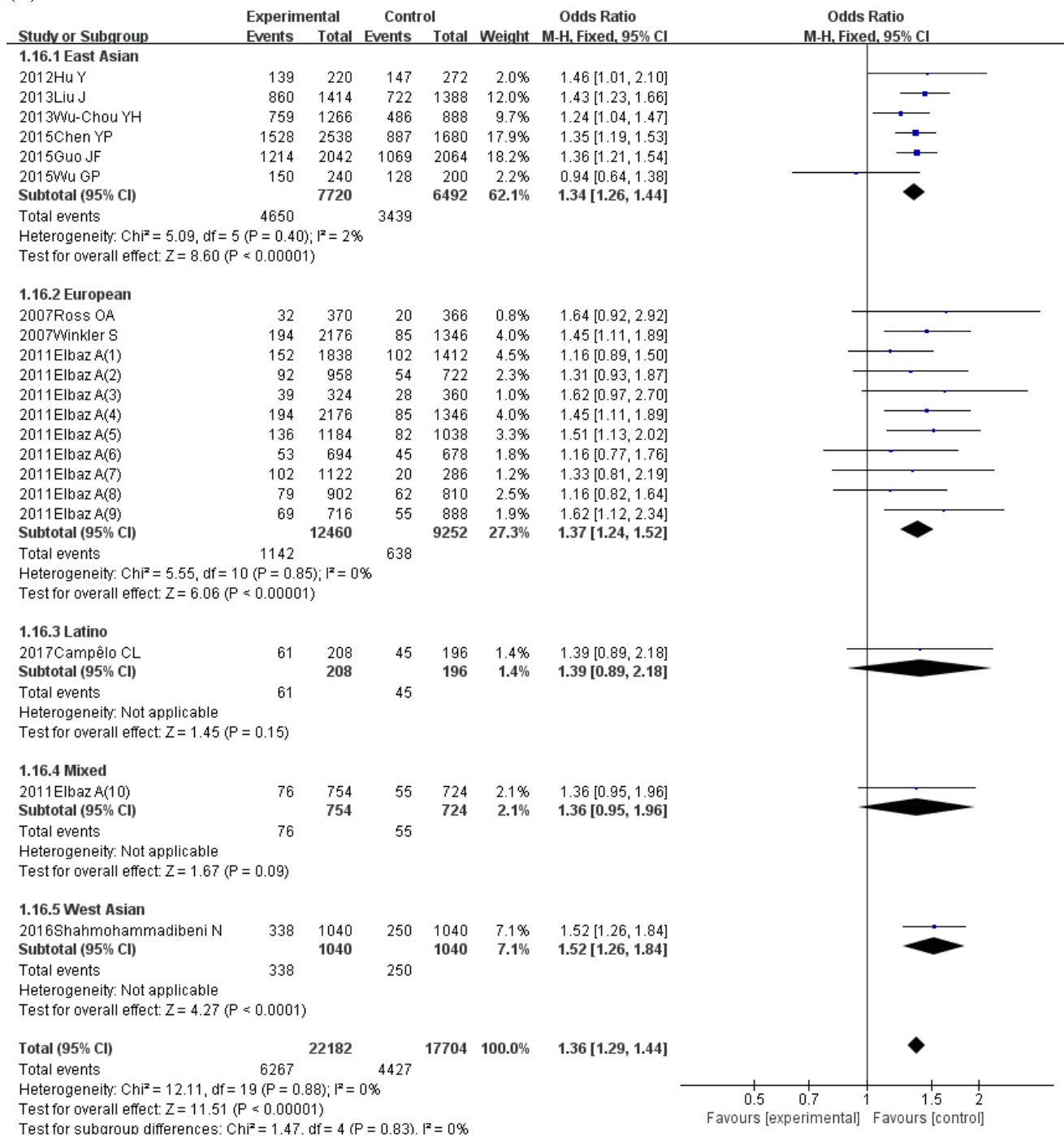
(N)



(O)



(P)



Supplementary Figure 1: Forest plots of the association between each *SNCA* variant (allele) included and PD risks in total and by ethnicity. (A)-(P) were individually responsive to the variants rs181489(T), rs356165(G), rs356186(A), rs356219(G), rs356220(T), rs356221(A), rs894278(G), rs2301134(A), rs2301135(G), rs2583988(T), rs2619363(T), rs2619364(G), rs2736990(G), rs2737029(G), rs7684318(C), rs11931074(T).