

Comprehensive Assessment of the Association between *FCGRs* polymorphisms and the risk of systemic lupus erythematosus: Evidence from a Meta-Analysis

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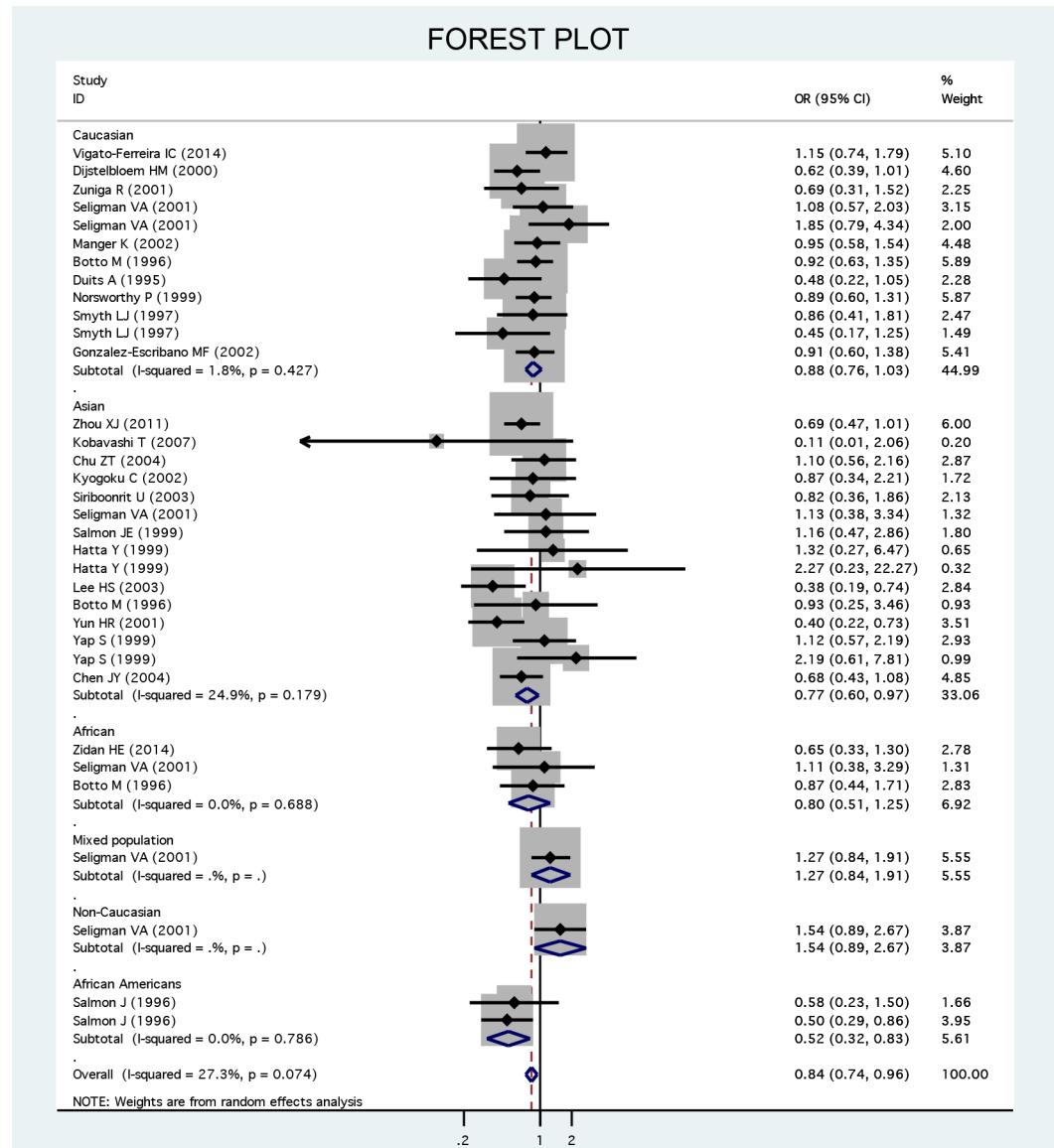
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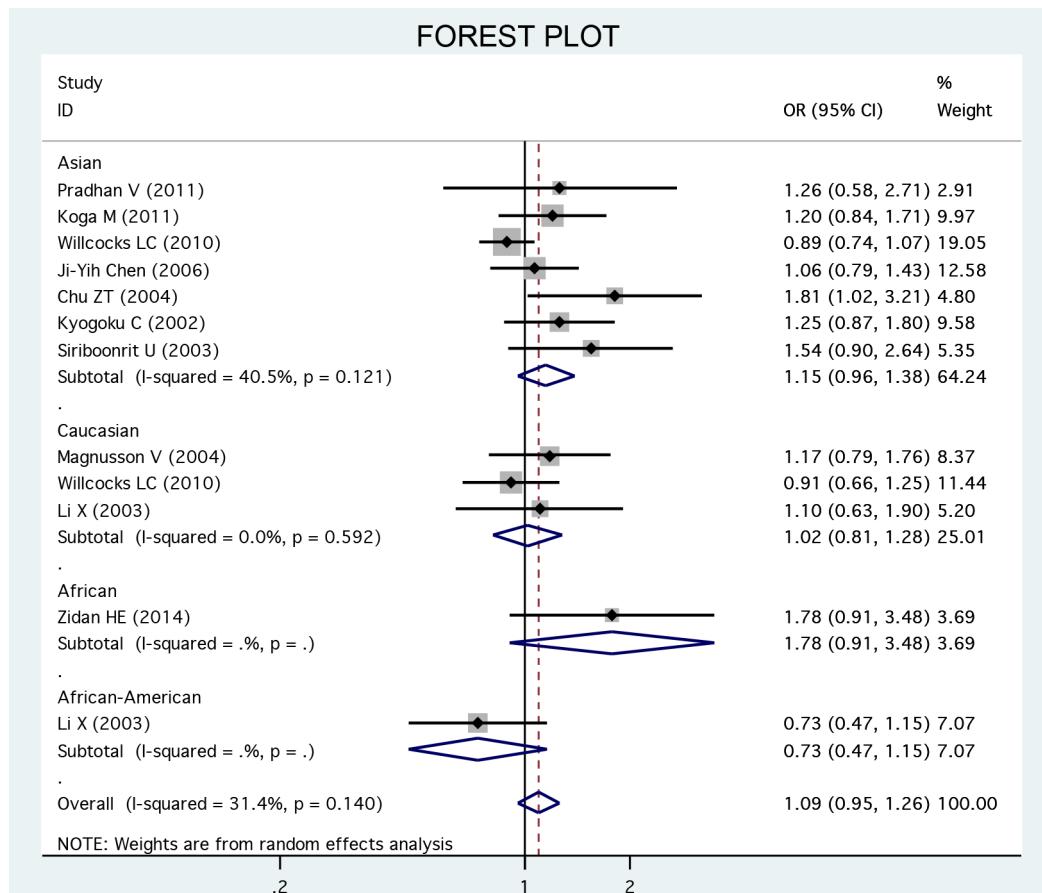
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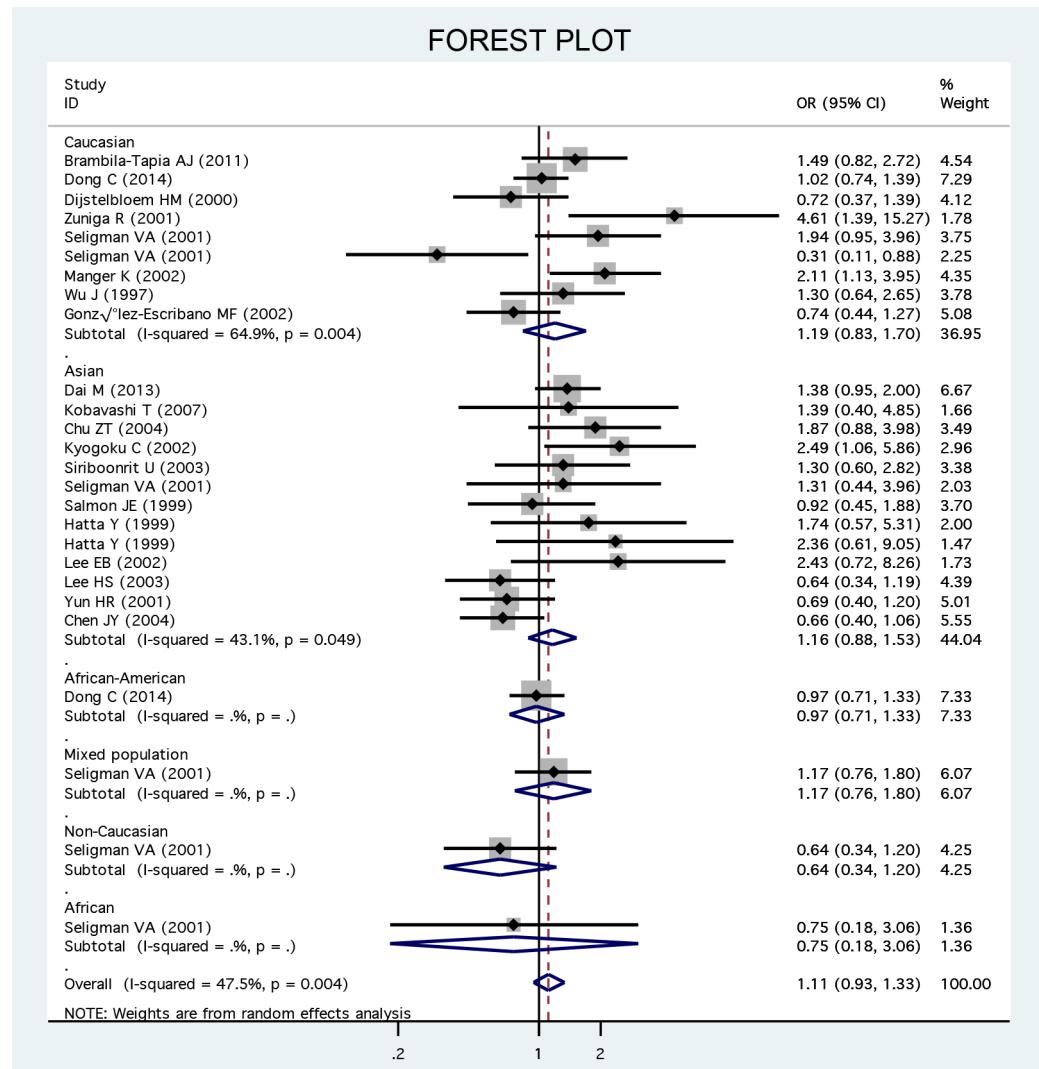
Supplementary Fig S1a. Forest plot for the meta-analysis of the association between FCGRs polymorphisms and SLE. (FCGR2A rs1801274 and SLE (AA/AG vs GG)



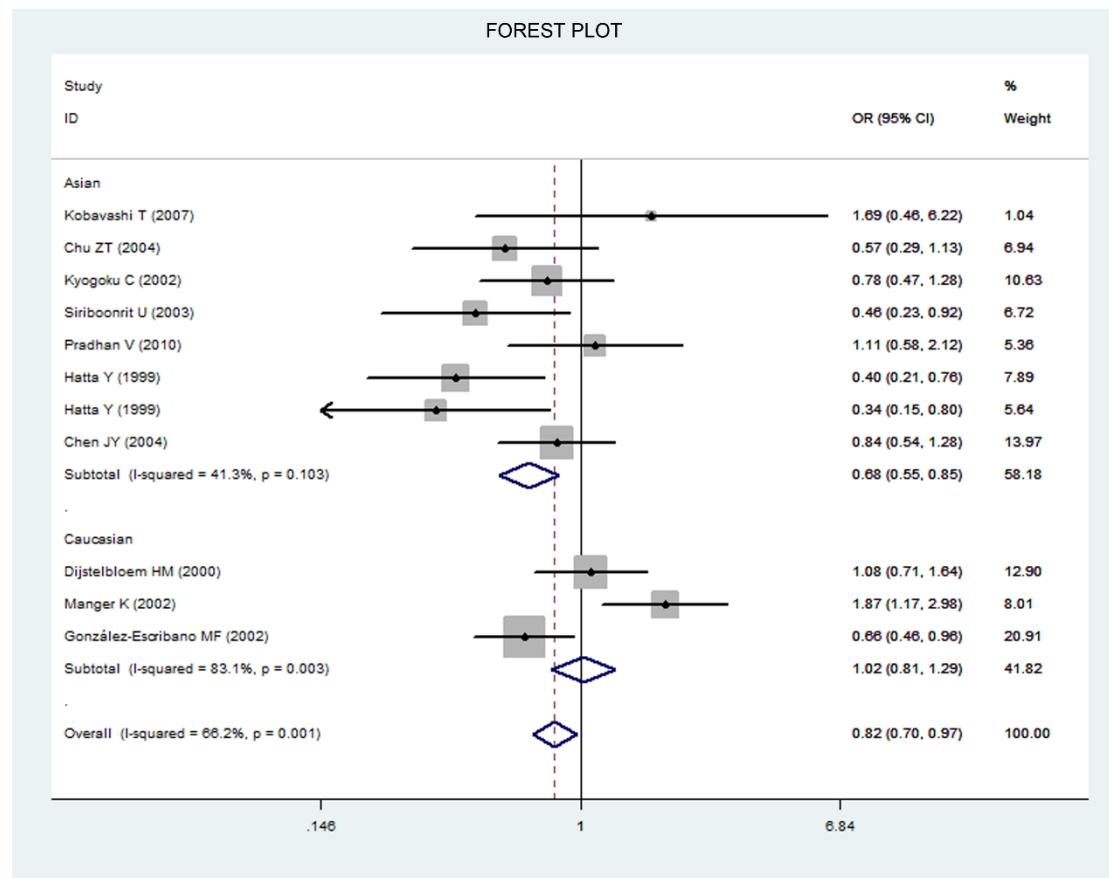
Supplementary Fig S1b. Forest plot for the meta-analysis of the association between FCGRs polymorphisms and SLE. (FCGR2B rs1050501 and SLE (CC/CT vs TT)



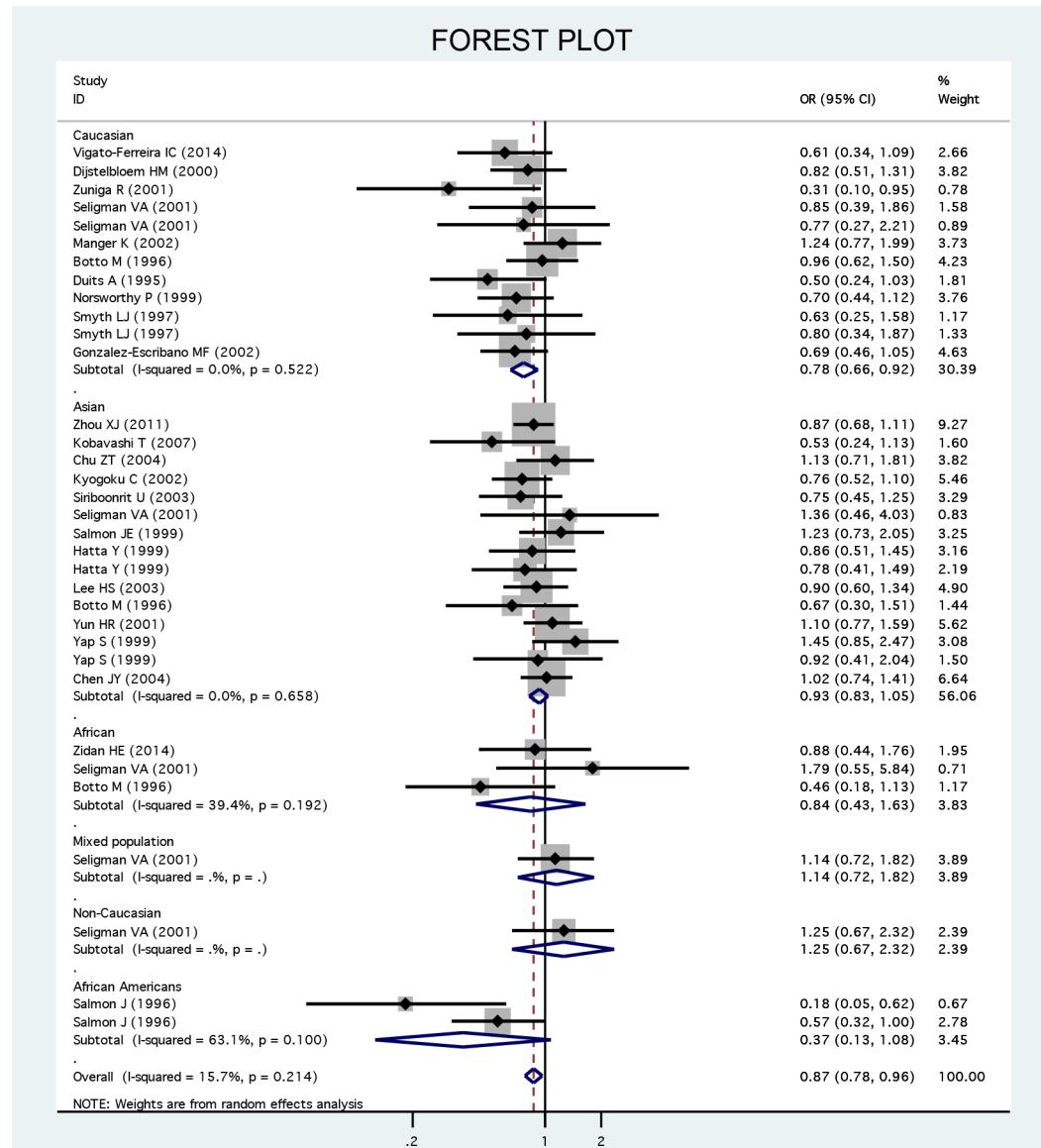
Supplementary Fig S1c. Forest plot for the meta-analysis of the association between FCGRs polymorphisms and SLE. (FCGR3A rs396991 and SLE (TT/TG vs GG)



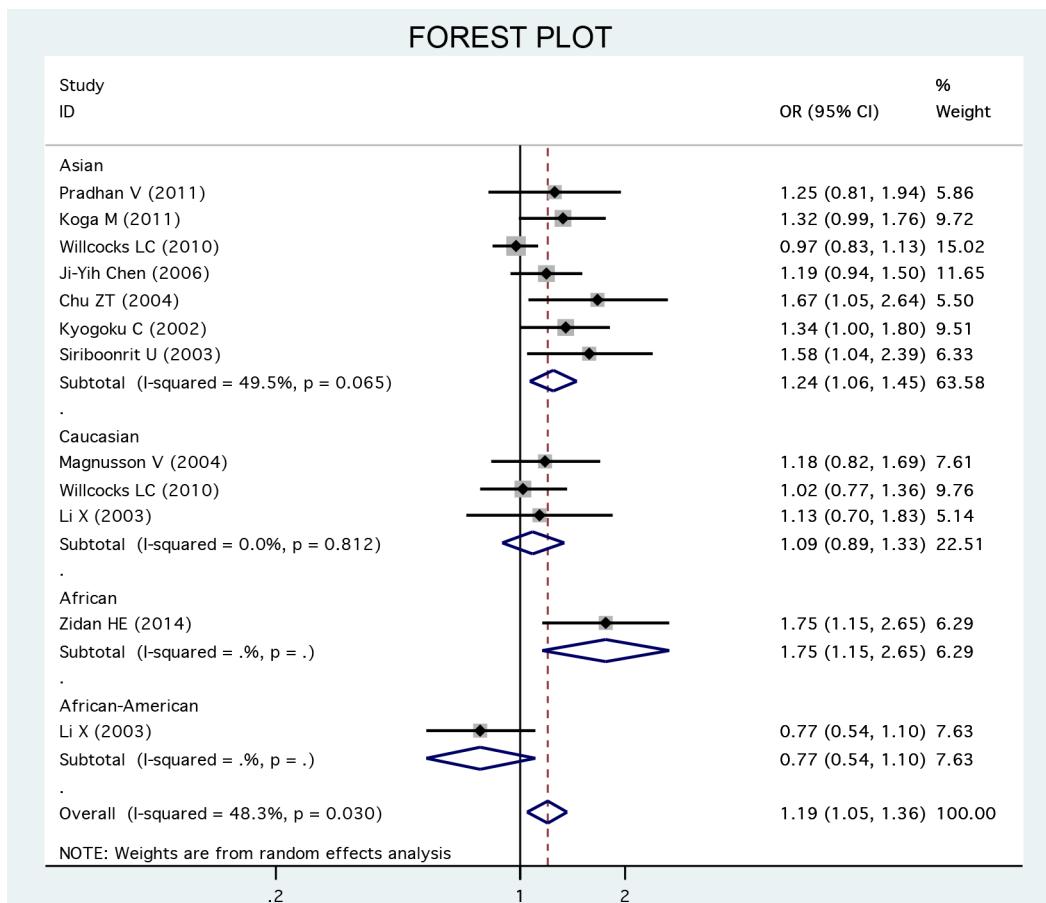
Supplementary Fig S1d. Forest plot for the meta-analysis of the association between FCGRs polymorphisms and SLE. (FCGR3B NA1•NA2 and SLE (NA1 NA1/ NA1 NA2 vs NA2 NA2)



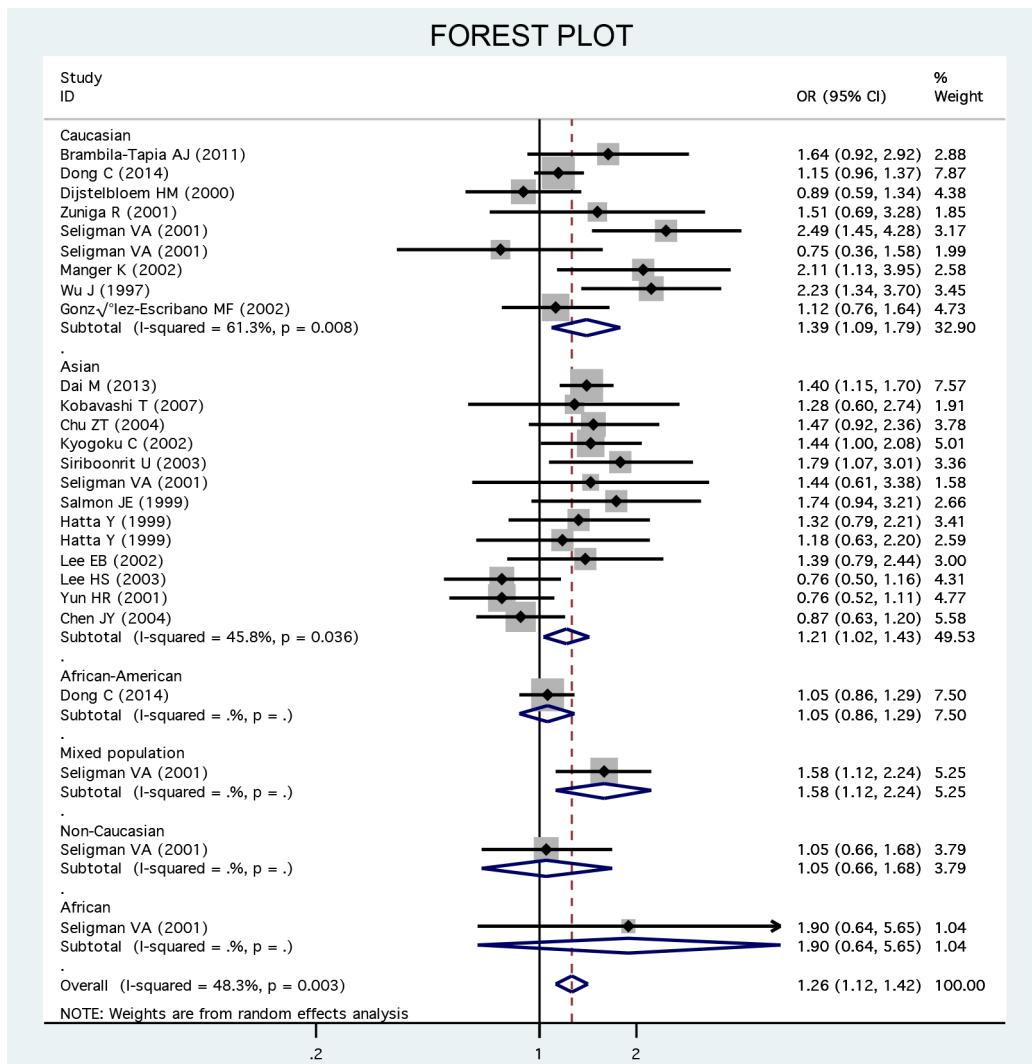
Supplementary Fig S2a. Forest plot for the meta-analysis of the association between FCGRs polymorphisms and SLE. (FCGR2A rs1801274 and SLE (AA vs AG/GG)



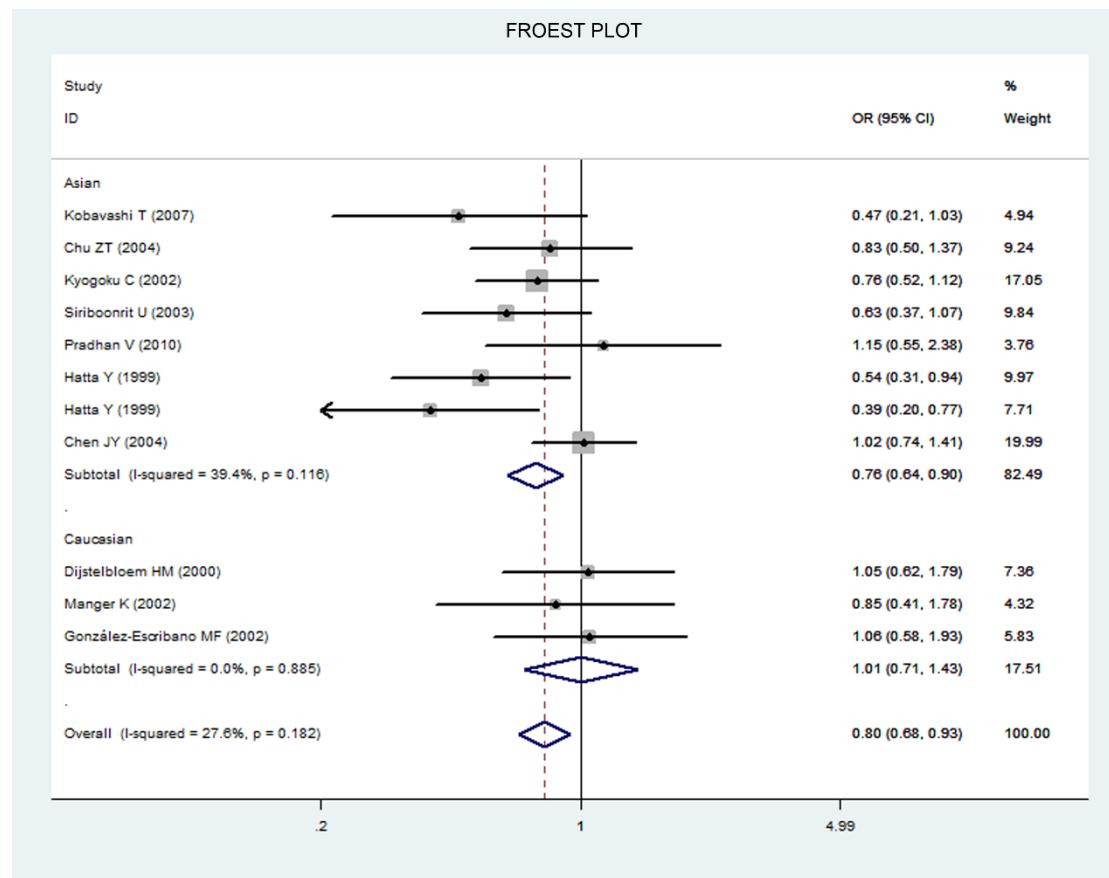
Supplementary Fig S2b. Forest plot for the meta-analysis of the association between FCGRs polymorphisms and SLE. (FCGR2B rs1050501 and SLE (C vs T)



Supplementary Fig S2c. Forest plot for the meta-analysis of the association between FCGRs polymorphisms and SLE. (FCGR3A rs396991 and SLE (T vs G)



Supplementary Fig S2d. Forest plot for the meta-analysis of the association between FCGRs polymorphisms and SLE. (FCGR3B NA1•NA2 and SLE (NA1NA1 vs NA1NA2/NA2NA2)



Supplementary Table S1 Begg's Test of publication under three models

	allele model		Recessive model		Dominant model	
	Z	P	Z	P	Z	P
FCGR2A(rs1801274)	0.53	0.594	1.16	0.248	0.77	0.441
FCGR2B(rs1050501)	1.58	0.115	0.21	0.837	1.85	0.064
FCGR3A(rs396991)	1.15	0.252	1.54	0.123	2.07	0.038
FCGR3B(NA1•NA2)	1.25	0.213	0.93	0.350	0.93	0.350