Effect of CD14 polymorphisms on the Risk of Cardiovascular Disease: Evidence from a Meta-Analysis

Running head: CD14 polymorphisms and Cardiovascular

Jin-Jian Xu^{1,2#}, Ke-Qi Liu^{3#}, Zhi-Min Ying^{4#}, Xiao-Wei Zhu¹, Xue-Jin Xu², Pian-Pian Zhao¹, Wei-Yang Bai¹, Mo-Chang Qiu³, Pei-Kuan Cong², Xing-Wei Zhang², Hou-Feng Zheng^{1,2*}

- 1, Diseases & Population (DaP) Geninfo Lab, School of Life Sciences, Westlake University and Westlake Institute for Advanced Study, Hangzhou, Zhejiang, China.
- 2, Institute of Ageing Research and the Affiliated Hospital, School of Medicine, Hangzhou Normal University, Hangzhou, Zhejiang, China
- 3, Jiangxi Medical College, Shangrao, Jiangxi, China
- 4, The Second Affiliated Hospital, Zhejiang University School of Medicine, Hangzhou, Zhejiang, China

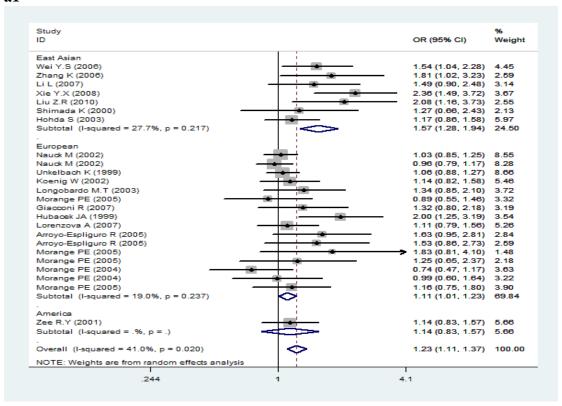
*Corresponding to: Hou-Feng Zheng, MD, PhD.

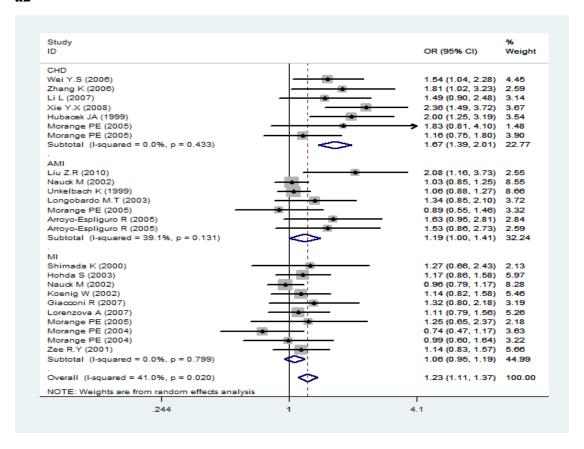
E-mail: zhenghoufeng@wias.org.cn

Table S1. The information of publication bias in all Polymorphisms

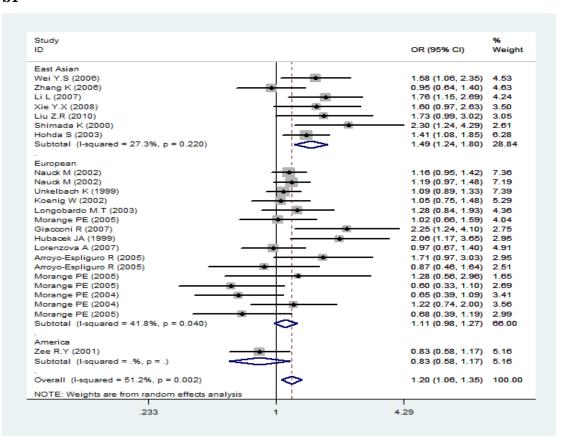
	Allele model		Dominant model		Recessive model	
Polymorphisms	$\overline{\mathbf{Z}}$	P	$\overline{\mathbf{Z}}$	P	$\overline{\mathbf{Z}}$	P
C-159T	1.77	0.077	1.4	0.161	1.4	0.161
C-260T	1.56	0.118	2.51	0.012	1.12	0.264

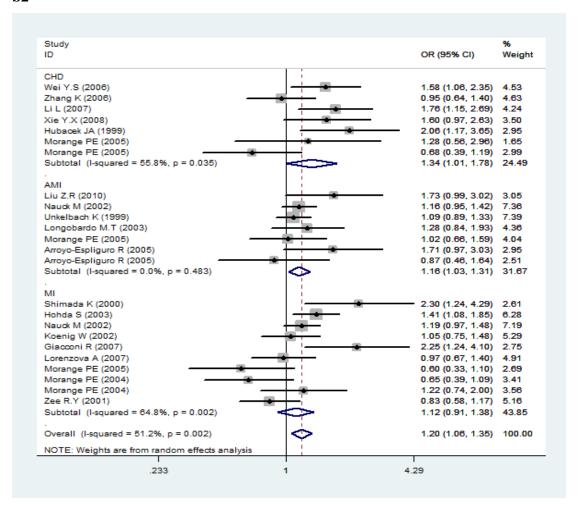
a1



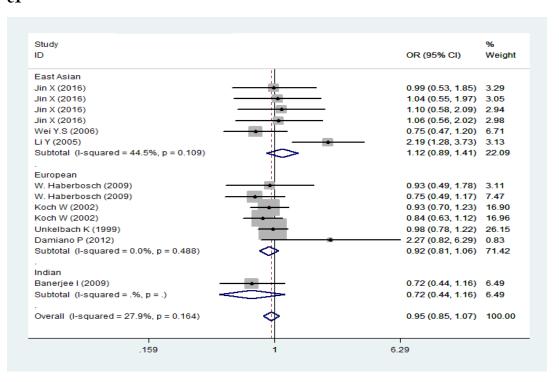


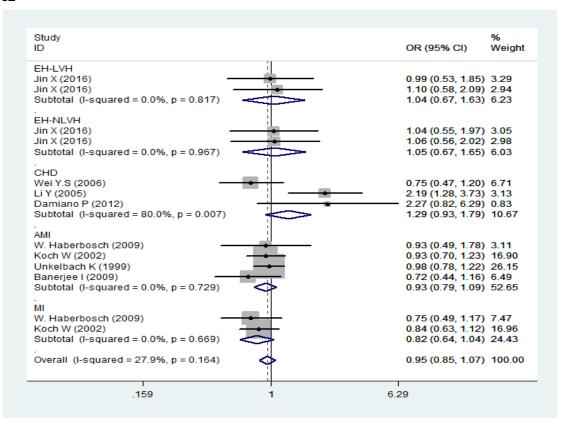
b1



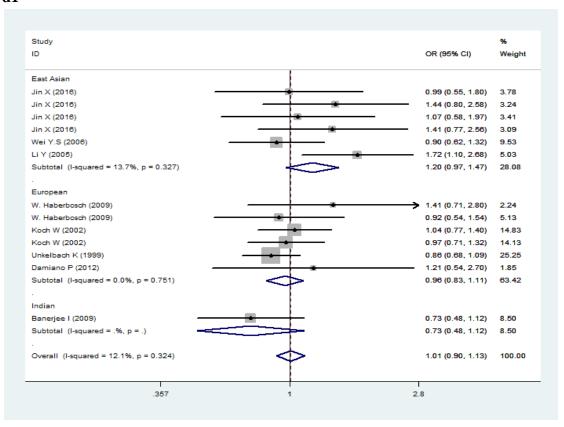


c1





d1



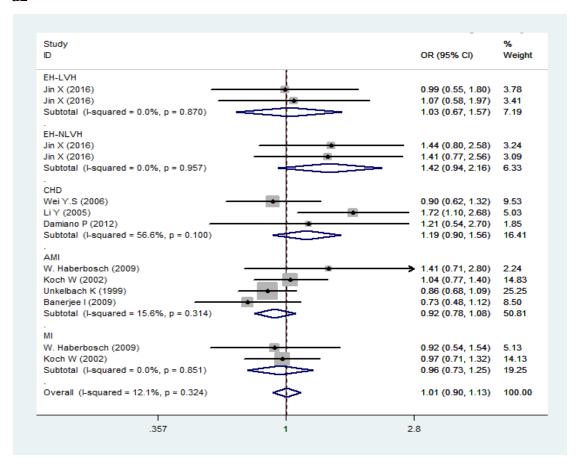
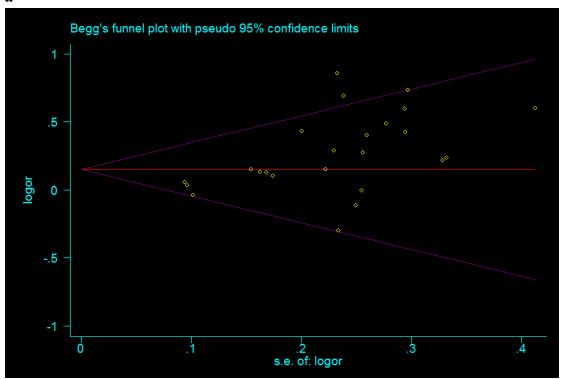
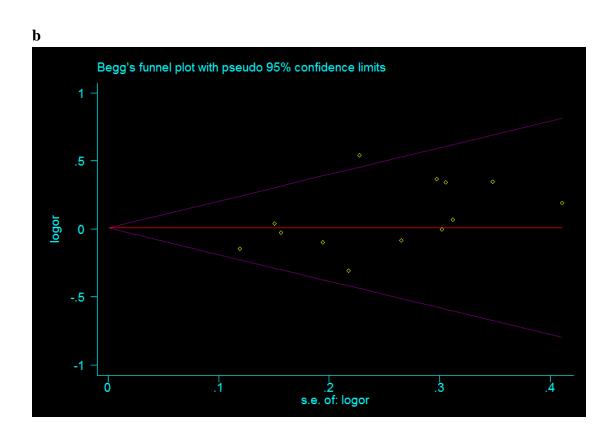
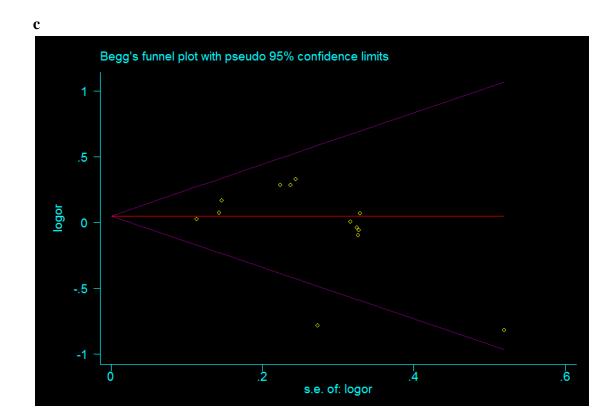


Figure S1. Forest plot for the meta-analysis of the association between CD14 gene polymorphisms and CVD under the Dominant and Recessive models

- (a1) rs2569190 and CVD. (CC VS.TT+CT), stratification by ethnicity
- (a2) rs2569190 and CVD. (CC VS.TT+CT), stratification by endpoints
- (b1) rs2569190 and CVD. (TT VS.CC+CT), stratification by ethnicity
- (b2) rs2569190 and CVD. (TT VS.CC+CT), stratification by endpoints
- (c1) C-159T and CVD. (CC VS.TT+CT), stratification by ethnicity
- (c2) C-159T and CVD. (CC VS.TT+CT), stratification by endpoints
- (d1) C-159T and CVD. (TT VS.CC+CT), stratification by ethnicity
- (d2) C-159T and CVD. (TT VS.CC+CT), stratification by endpoints







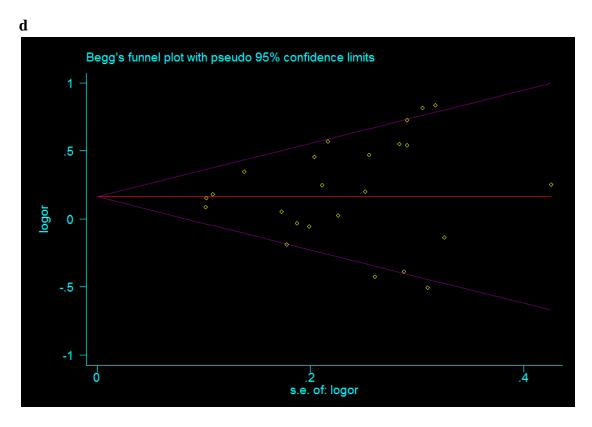
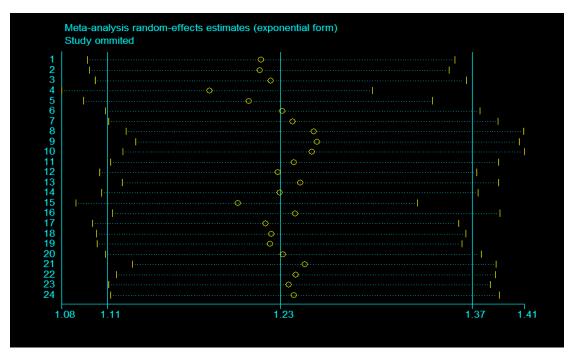


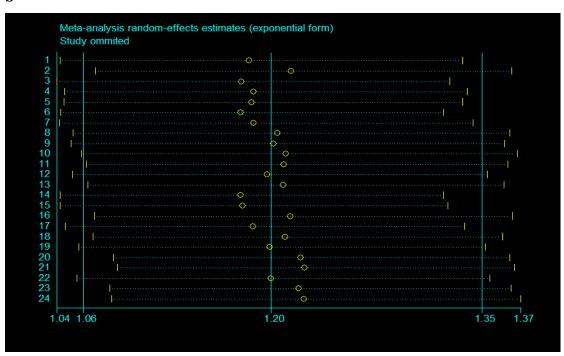
Figure S2. Begg's funnel plot of publication bias in the meta-analysis of the association of CD14 polymorphisms with CVD risk

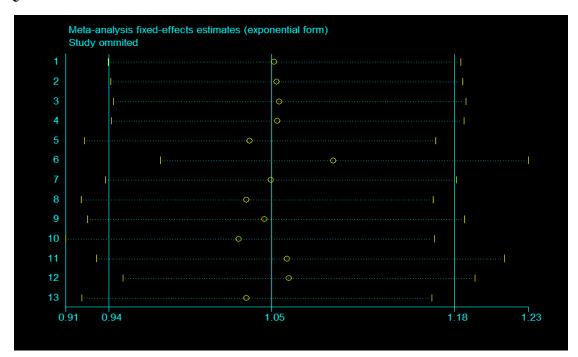
- (a) rs2569190 and CVD. (CC VS.TT+CT)
- (b) rs2569190 and CVD. (TT VS.CC+CT)
- (c) C-159T and CVD. (CC VS.TT+CT)
- (d) C-159T and CVD. (TT VS.CC+CT)

a



b





 \mathbf{d}

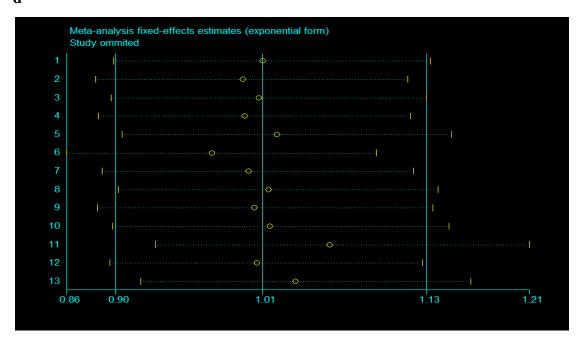


Figure S3. Sensitivity analysis to assess the stability of the meta-analysis

- (a) rs2569190 and CVD. (CC VS.TT+CT)
- (b) rs2569190 and CVD. (TT VS.CC+CT)
- (c) C-159T and CVD. (CC VS.TT+CT)
- (d) C-159T and CVD. (TT VS.CC+CT)