

**Association between paraoxonase 1 -108C/T polymorphism and coronary heart disease: an updated meta-analysis**

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**ADMINISTRATIVE INFORMATION****Support** - No funding.**Review Stage at time of this submission** - Preliminary searches.**Conflicts of interest** - None declared.**INPLASY registration number:** INPLASY202430117**Amendments** - This protocol was registered with the International Platform of Registered Systematic Review and Meta-Analysis Protocols (INPLASY) on 27 March 2024 and was last updated on 27 March 2024.**INTRODUCTION**

**Review question / Objective** At present, no consensus is reached among articles that investigate the relationship of paraoxonase 1 (PON1) -108C/T polymorphism with susceptibility of coronary heart disease (CHD) so far.

**Condition being studied** In this regard, the present meta-analysis was conducted to comprehensively review existing articles related to the relationship of PON1 -108C/T polymorphism with CHD susceptibility.

**METHODS**

**Participant or population** PON1 -108C>T polymorphism and CHD.

**Intervention** To evaluate the impact of PON1 -108C>T polymorphism on CHD clinical outcomes.

**Comparator** The PON1 -108T allele is the risk factor of CHD.

**Study designs to be included** Meta analysis.

**Eligibility criteria** We selected studies based on the following criteria: (a) case-control articles evaluating the relationship of PON1 -108C>T with CHD susceptibility; (b) studies conducted by enrolling irrelevant subjects; (c) studies with enough data on ORs and 95% CIs. At the same time, reports, reviews, letters and comments were eliminated from this study. For duplicates, the one that had the greatest sample size was adopted for the present meta-analysis.

**Information sources** Web of Science, PubMed, CNKI and Embase were systemically retrieved to enroll eligible articles published before June 2020.

**Main outcome(s)** The meta-analysis was conducted to found the relationship of PON1 -108C/T polymorphism with CHD susceptibility.

**Quality assessment / Risk of bias analysis** Begg's funnel plot was employed to investigate the bias of publication, with  $P < 0.05$  indicating the presence of significant bias of publication.

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**Strategy of data synthesis** STATA12.0(Stata Corporation, College Station, Texas) was utilized for meta-analysis.

**Subgroup analysis** Subgroup analyses is based on HWE,sample size and race.

**Sensitivity analysis** We also conducted sensitivity for evaluating result stability through eliminating a study each time for determining its impact on combined ORs.

**Country(ies) involved** China.

**Keywords** Coronary heart disease, -108C/T, Risk.

**Contributions of each author**

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