



The Role of C-Reactive Protein and Fibrinogen in the Development of Intracerebral Hemorrhage: A Mendelian Randomization Study in European Population

Biyang Wang^{1†}, Xiaoyu Zhang^{1,2†}, Di Liu¹, Jie Zhang¹, Mingyang Cao¹, Xin Tian¹, Isinta Elijah Maranga¹, Xiaoni Meng¹, Qiuyue Tian¹, Feifei Tian¹, Weijie Cao¹, Wei Wang², Manshu Song^{2*} and Youxin Wang^{1*}

OPEN ACCESS

Edited by:

Lei Zhang,
Soochow University, China

Reviewed by:

Wei-Min Chen,
University of Virginia, United States
Ping Zeng,
Xuzhou Medical University, China

*Correspondence:

Manshu Song
m.song@ecu.edu.au
Youxin Wang
wangy@cmmu.edu.cn

[†]These authors have contributed
equally to this work

Specialty section:

This article was submitted to
Statistical Genetics and Methodology,
a section of the journal
Frontiers in Genetics

Received: 02 October 2020

Accepted: 06 January 2021

Published: 04 February 2021

Citation:

Wang B, Zhang X, Liu D, Zhang J,
Cao M, Tian X, Maranga IE, Meng X,
Tian Q, Tian F, Cao W, Wang W,
Song M and Wang Y (2021) The Role
of C-Reactive Protein and Fibrinogen
in the Development of Intracerebral
Hemorrhage: A Mendelian
Randomization Study in
European Population.
Front. Genet. 12:608714.
doi: 10.3389/fgene.2021.608714

¹Beijing Key Laboratory of Clinical Epidemiology, School of Public Health, Capital Medical University, Beijing, China,

²School of Medical and Health Sciences, Edith Cowan University, Joondalup, WA, Australia

Background: The causal association of C-reactive protein (CRP) and fibrinogen on intracerebral hemorrhage (ICH) remains uncertain. We investigated the causal associations of CRP and fibrinogen with ICH using two-sample Mendelian randomization.

Method: We used single-nucleotide polymorphisms associated with CRP and fibrinogen as instrumental variables. The summary data on ICH were obtained from the International Stroke Genetics Consortium (1,545 cases and 1,481 controls). Two-sample Mendelian randomization estimates were performed to assess with inverse-variance weighted and sensitive analyses methods including the weighted median, the penalized weighted median, pleiotropy residual sum and outlier (MR-PRESSO) approaches. MR-Egger regression was used to explore the pleiotropy.

Results: The MR analyses indicated that genetically predicted CRP concentration was not associated with ICH, with an odds ratio (OR) of 1.263 (95% CI = 0.935–1.704, $p = 0.127$). Besides, genetically predicted fibrinogen concentration was not associated with an increased risk of ICH, with an OR of 0.879 (95% CI = 0.060–18.281; $p = 0.933$). No evidence of pleiotropic bias was detected by MR-Egger. The findings were overall robust in sensitivity analyses.

Conclusions: Our findings did not support that CRP and fibrinogen are causally associated with the risk of ICH.

Keywords: C-reactive protein, fibrinogen, single-nucleotide polymorphisms, Mendelian randomization, intracerebral hemorrhage

INTRODUCTION

Globally, stroke is a leading cause of death with a high societal burden in most regions (GBD 2015 Mortality and Causes of Death Collaborators, 2016). Among adults, the risk of stroke from the age of 25 years is approximately 25% (Feigin et al., 2018). Hemorrhagic stroke (HS) as a subtype of stroke carries high morbidity and mortality rates (Stokum et al., 2015),

and intracerebral hemorrhage (ICH) is by far the most common type of HS (Qureshi et al., 2009). Inflammation plays an important part in pathogenesis of stroke by influencing the development of atherosclerosis and plaque instability (Barone and Feuerstein, 1999; Scirica and Morrow, 2006).

C-reactive protein (CRP) and fibrinogen, considered as well-proven clinical markers of systemic inflammation, are acute-phase protein synthesized by hepatocytes against inflammation (Dalmon et al., 1993; Pepys and Hirschfield, 2003) and can increase the risk of cardiovascular disease (Scirica and Morrow, 2006; Zhang et al., 2014) and stroke (Coull et al., 1991; Cao et al., 2007; Jiménez et al., 2015). Significantly increased levels of fibrinogen are commonly found in patients with stroke, suggesting that fibrinogen is elevated before thrombotic incidents occur and is a risk factor for stroke (Coull et al., 1991). However, Roudbary et al. (2011) revealed that CRP concentration was not improved in patients with HS. No associations of CRP and fibrinogen with ICH were identified in a nested case-control study (Karim et al., 2020). The observational epidemiologic studies on the associations of CRP and fibrinogen with ICH showed inconsistent results (Coull et al., 1991; Roudbary et al., 2011). Furthermore, potential unmeasured confounders and reverse causation bias in observational studies limit the ability to ascertain causal inferences.

Mendelian randomization (MR) is a genetic epidemiological method to explore the association between the exposure and outcome, using genetic variants as instrumental variables (IVs) for the exposure (Smith and Ebrahim, 2003). Because of the independent segregation and randomized assignment of alleles at meiosis, MR approach can control potential confounders and reverse causation, making stronger causal inference (Lawlor et al., 2008). Therefore, we conducted two-sample MR analysis to assess the causal relationships of CRP and fibrinogen in the development of ICH in European population.

MATERIALS AND METHODS

Study Design and Data Sources

A two-sample MR approach was used to investigate the causal effects of CRP and fibrinogen on the risk of ICH. The study design is under the assumption that the genetic variants are associated with CRP and fibrinogen, but not with confounders. Besides, the genetic variants affect risk of ICH only through exposure and not through any alternative pathways.

Information on genetic variants associated with level of CRP was collected from a meta-analysis of genome-wide association study (GWAS), which is currently the largest study attempted to identify genetic variants in relation to CRP concentration involving 204,402 individuals from 88 previous population-based cohort studies (Ligthart et al., 2018). In genetic variants associated with fibrinogen, we used previously published genetic variants of a GWAS meta-analysis involving more than 100,000 subjects (Sabater-Lleal et al., 2013).

Summary statistics data on associations of genetic variants with ICH were obtained from the published GWAS meta-analysis by the International Stroke Genetics Consortium (ISGC)

of 3,026 participants (1,545 cases and 1,481 controls; Woo et al., 2014). All data in our MR analyses were restricted to individuals of European ancestry only.

Genetic Variants

We used single-nucleotide polymorphisms (SNPs) published previously, which reached genome-wide significance ($p < 5 \times 10^{-8}$) for CRP and fibrinogen concentrations as MR IVs. The selected SNPs were independent, namely, not in linkage disequilibrium ($r^2 < 0.2$). Nineteen SNPs (11 for CRP and 8 for fibrinogen) were not presented in ISGC datasets. For the unavailable SNPs in outcome datasets, we replaced them with proxy SNPs. The proxy SNPs in linkage disequilibrium ($r^2 > 0.8$) were identified for two SNPs. Accordingly, 42 SNPs for CRP and 16 SNPs for fibrinogen were included in the analysis of ICH. The summary genetic association data are reported in **Supplementary Table S1**.

Mendelian Randomization Analysis

We performed two-sample MR analyses to estimate the associations of CRP and fibrinogen with ICH using summarized data. Causal effects on ICH of CRP and fibrinogen concentrations were estimated using the conventional inverse-variance weighted (IVW) method (Burgess et al., 2013). We also conducted sensitivity analyses using the weighted median (WM), the penalized weighted median (PWM), and pleiotropy residual sum and outlier (MR-PRESSO) methods (Bowden et al., 2015, 2016; Verbanck et al., 2018). For MR-Egger regression analysis, we assessed directional pleiotropy based on its intercepts (Burgess and Thompson, 2017). A leave-one-out analysis (omitted one SNP in turn) was performed to test the influence of outlying values (Burgess and Thompson, 2017). Heterogeneity of individual genetic variants was evaluated by Cochran's Q test. All results are presented as an odds ratio (OR) with 95% confidence interval (CI) of the outcomes per predicted increase in CRP and fibrinogen concentrations. The associations of each SNP with CRP and fibrinogen concentrations are further plotted compared to their effects for the outcomes. All analyses were performed by the TwoSampleMR and MR-PRESSO packages with R version 4.0.2.

RESULTS

Causal Association of CRP With ICH

The results of associations between genetically determined CRP and fibrinogen and the risk of ICH were presented in **Table 1**. Genetic predisposition to CRP levels were not observed to be statistically significantly associated with ICH by performing IVW method (OR = 1.263, 95% CI = 0.935–1.704, $p = 0.127$). The lack of causal association remained in all sensitivity analyses (all $p > 0.05$; **Table 1**).

The MR-Egger method showed no evidence of directional pleiotropy for the association of CRP with ICH [odds (intercept), -0.010 ; $p = 0.480$; **Table 2**]. For IVs, MR-PRESSO did not detect any potential outliers. Likewise, no heterogeneity was observed among individual SNPs of CRP for ICH ($Q = 36.775$,

$p = 0.616$, **Table 2**). We calculated the individual and pooled MR estimates between each CRP-related SNP and the risk for ICH shown as forest plots and scatter plots in **Figure 1**. The result of leave-one-out sensitivity analysis showed that the association between CRP and ICH was not substantially driven by any individual SNP (**Supplementary Figure S1**).

Causal Association of Fibrinogen With ICH

Regarding fibrinogen, we found no causal effect of genetically instrumented fibrinogen on ICH (OR = 0.879, 95% CI = 0.042–18.281, $p = 0.933$). No significant association was observed for ICH in sensitivity analyses that were performed by WM, PWM, MR-PRESSO, and MR-Egger methods (**Table 1**).

The MR-Egger method showed no evidence of directional pleiotropy for the association of fibrinogen with ICH (**Table 2**). For IVs, MR-PRESSO did not detect any potential outliers. We calculated the individual and pooled MR estimates between each fibrinogen-related SNP and risk for ICH shown as forest plots and scatter plots in **Figure 2**. Furthermore, analysis on

leaving out each SNP revealed that the inverse association between fibrinogen concentrations and ICH was not substantially driven by any individual SNP (**Supplementary Figure S2**). However, the Cochran Q statistic was 28.028 with an associated $p < 0.05$, suggesting some heterogeneity in the effect estimates of fibrinogen and ICH (**Table 2**), but there was no clear evidence of directional pleiotropy (p for intercept > 0.05 , **Table 2**).

DISCUSSION

In the present study, we assessed whether high circulating levels of CRP and fibrinogen are causally associated with ICH using two-sample MR analysis in European population. In the present study using publicly available summary statistics data, we did not find CRP and fibrinogen levels might increase ICH risk. The findings were overall robust in sensitivity analyses.

Apart from being markers of systemic inflammation, CRP and fibrinogen are acute-phase protein induced by proinflammatory cytokine contributing to host defense against infection (Dalmon et al., 1993; Pepys and Hirschfield, 2003). Previous studies investigated associations between CRP and fibrinogen and ICH but reported inconsistent results. A large-scale cohort study found that CRP and fibrinogen were not associated with a significantly greater risk of HS (Jiménez et al., 2016), while a retrospective cohort study suggested that increased CRP was a significant risk factor for in-hospital mortality among patients with cardiovascular disease including ICH (Yoshinaga et al., 2017).

In our analysis, we did not observe the relationships of CRP and fibrinogen with ICH. These findings suggested that the role of CRP and fibrinogen may be less important in causing the risk of ICH. A previous MR study indicated that CRP concentration itself was unlikely to be even a modest causal factor in coronary heart disease (Wensley et al., 2011). Our findings corroborate earlier studies that showed CRP had no clear effect on ICH risk (Liu et al., 2014). Similar results were also found in a meta-analysis consisting of six population-based prospective studies (Georgakis et al., 2019). Another meta-analysis has also suggested that elevated baseline CRP levels exhibited no clear effect on HS (Zhou et al., 2016). However, evidence from a few prospective studies showed that CRP level in HS patients was significantly elevated (Das et al., 2014; Xue et al., 2017).

Fibrinogen participates in platelet aggregation, thrombogenic activity, atherogenesis, and inflammation, and the role of fibrinogen is probably various in the different subtypes of stroke. Our findings were supported by previous studies, which also reported

TABLE 1 | Mendelian randomization (MR) estimates of exposure with intracerebral hemorrhage from the inverse-variance weighted (IVW) and sensitivity analysis.

Phenotype	IVs (SNPs)	OR (95% CI)	p
CRP			
IVW	42	1.263 (0.935–1.704)	0.127
Weighted median	42	1.458 (0.977–2.175)	0.065
Penalized weighted median	42	1.466 (0.957–2.247)	0.079
MR_Egger	42	1.432 (0.906–2.266)	0.133
MR-PRESSO	42	1.236 (0.950–1.522)	0.154
Fibrinogen			
IVW	16	0.879 (0.042–18.281)	0.933
Weighted median	16	0.438 (0.016–11.771)	0.623
Penalized weighted median	16	0.438 (0.014–13.561)	0.637
MR_Egger	16	1.663 (0.004–746.651)	0.872
MR-PRESSO	16	1.221 (–1.893 to 4.335)	0.901

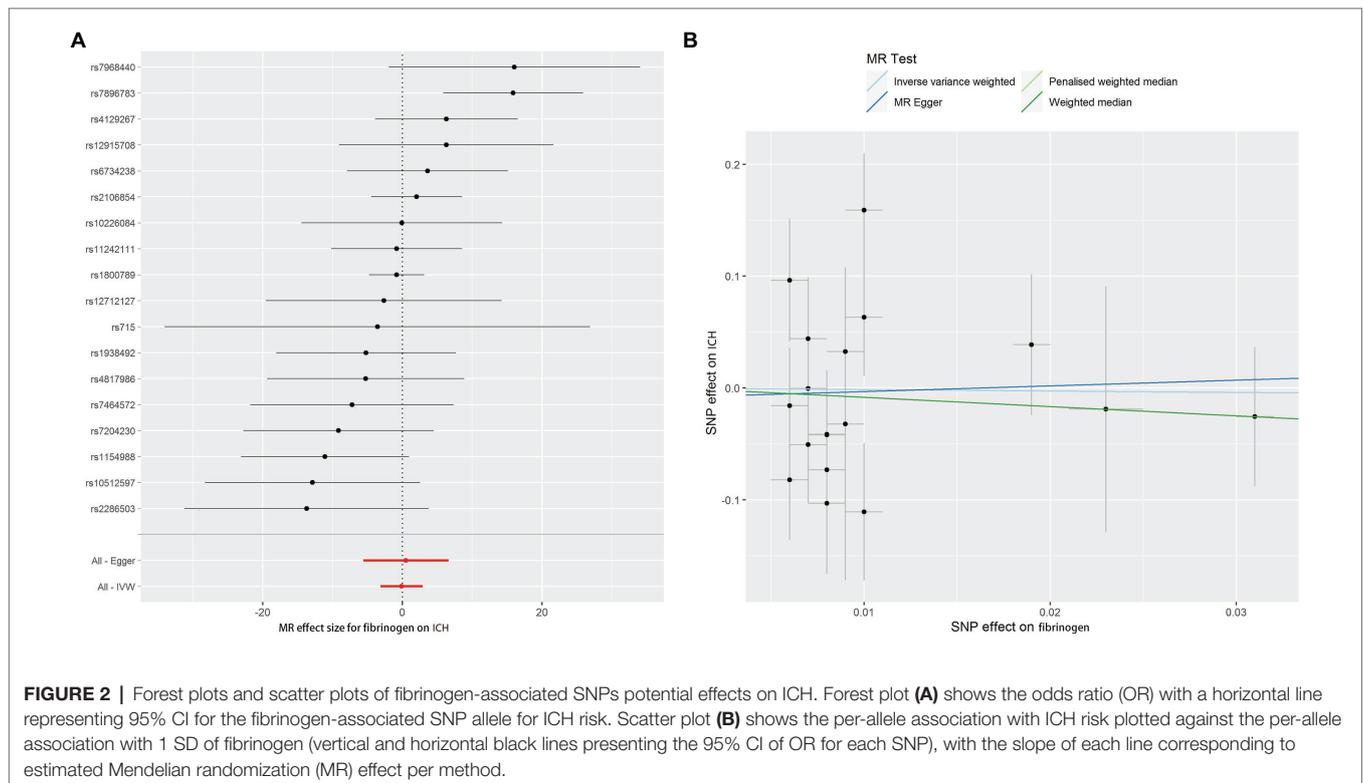
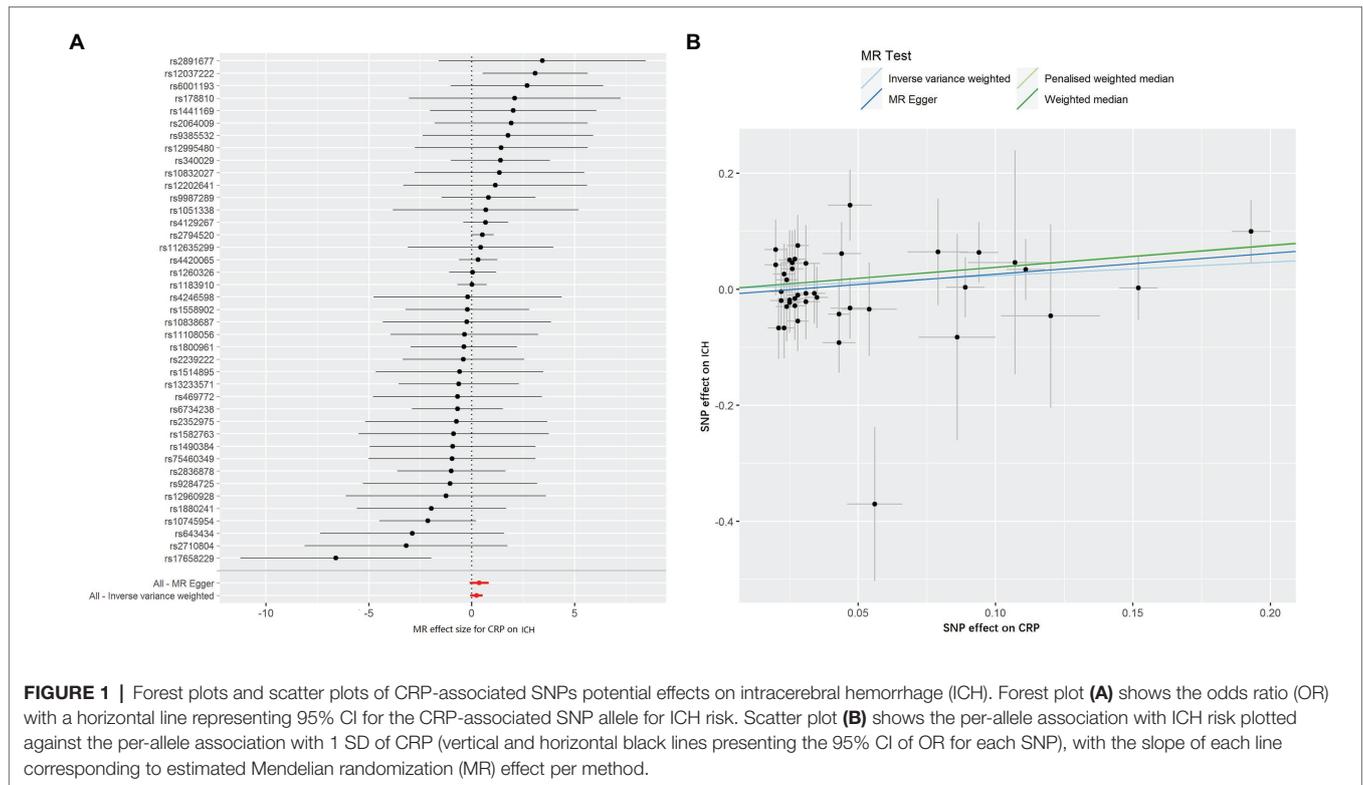
CRP, C-reactive protein; IVW, inverse-variance weighted; MR, Mendelian randomization; WM, weighted median; PWM, penalized weighted median; OR, odds ratio; MR-PRESSO, pleiotropy residual sum and outlier; CI, confidence interval.

TABLE 2 | Heterogeneity tests and MR-Egger intercept of CRP and fibrinogen causally linked to ICH.

Outcome	Exposure	Intercept	p^a	Cochran's Q	Q_df	p^b
ICH	CRP	–0.010	0.480	36.775	40	0.616
ICH	Fibrinogen	–0.008	0.815	28.028	15	0.045

^aValue of p for MR-Egger intercept.

^bValue of p for heterogeneity tests by performing inverse-variance weighted method. CRP, C-reactive protein; ICH, intracerebral hemorrhage.



that elevated levels of fibrinogen did not exhibit suggestive evidence of association with HS (Alvarez-Perez et al., 2011). In line with our results, no significant association between

fibrinogen and ICH was observed in observational studies (Woodward et al., 2005; Welsh et al., 2008; Folsom et al., 2016). However, greater plasma fibrinogen concentration was associated

with increased risk of ICH in these prospective studies (Sato et al., 2006; Sturgeon et al., 2008). These findings should be interpreted cautiously as higher CRP and fibrinogen levels may reflect subclinical infection, chronic infectious diseases, preexisting disease, and socioeconomic or lifestyle characteristics. Besides, these opposite results may be due to different study populations and ethnic groups (Iso et al., 2012; Shi et al., 2016).

The pathogenesis of the associations of CRP and fibrinogen with the risk of ICH is unclear. CRP plays a direct role in the pathogenesis of atherosclerosis and is upregulated significantly in atheromatous plaques, where it may promote low-density lipoprotein cholesterol uptake by macrophages (Torzewski et al., 2000). Moreover, these inconsistent previous results may be due to reverse causal bias or confounders from atherosclerosis (Libby et al., 2011) or inflammation (Hartwig et al., 2017). One possible explanation is that the previous finding was a false-positive outcome because the effect of confounding was not controlled for, whereas in our studies, the genetic variants associated with exposure explained a larger proportion of variance, showing the true relationship of CRP and fibrinogen with ICH. Another possible explanation is that a mass of variants resulted in greater pleiotropy potential, which may have diluted the association in our analysis.

The major strengths of this study are using data from large-scale GWAS studies and ISGC collaboration. We used a two-sample MR approach assessing CRP and fibrinogen levels in relation to the risk of ICH in European-descent individuals, which reduces bias of population stratification. Moreover, in terms of the MR analysis, we performed conventional IVW, WM, PWM, MR-PRESSO, and MR-Egger methods to avoid reverse causation and to reduce other confounding factors. Lastly, there is no strong evidence of pleiotropic effects for the genetic instruments, suggesting there was less likelihood of CRP and fibrinogen-related SNPs are associated with other phenotypes.

The present study also has some limitations. Interpreting the magnitude of estimates for the effect of CRP and fibrinogen on ICH risk requires caution. First, stratified analyses or analyses adjusted for other covariates were not possible on the account of using the available summary statistics datasets. In addition, the genetic IVs accounted for approximately 7.0% of the total variation in CRP and 3.7% of plasma fibrinogen variation (Sabater-Lleal et al., 2013; Ligthart et al., 2018), which might be low for the use as IVs, and any bias from weak instruments was in the direction of the null (Pierce and Burgess, 2013). Nevertheless, MR analysis likely reflects lifelong exposure to elevated CRP and fibrinogen levels. However, it is possible that only exposure in a specific window of time (e.g., early life) affects ICH risk. Lastly, we used a relatively small sample size to explore the causal relationship between CRP, fibrinogen, and ICH with the

power of less than 0.90. Thus, the nonsignificant but still suggestive associations between CRP and fibrinogen levels and ICH risk should be further validated in future studies with larger independent populations and larger datasets offering greater statistical power.

In conclusion, these MR analyses did not find evidence to support the causal relationship between CRP and fibrinogen with ICH. The results add to the burgeoning evidence that refutes the harmful role of CRP and fibrinogen in ICH. Further research is required to clarify this finding, using larger samples for undertaking “adjusted” MR analyses.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding authors.

AUTHOR CONTRIBUTIONS

BW and XZ drafted the manuscript. YW, MS, DL, JZ, MC, XT, IM, XM, QT, FT, WC, and WW critically revised the manuscript. All authors contributed to the article and approved the submitted version.

FUNDING

The study was supported by grants from the National Natural Science Foundation of China (81673247, 81872682, and 81773527) and the China-Australian Collaborative Grant (NSFC 81561128020-NHMRC APP1112767).

ACKNOWLEDGMENTS

Data on CRP and fibrinogen associated single-nucleotide polymorphisms have been derived from published articles (Sabater-Lleal et al., 2013; Ligthart et al., 2018). Data on intracerebral hemorrhage were accessed through the ISGC Cerebrovascular Disease Knowledge Portal (Woo et al., 2014).

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fgene.2021.608714/full#supplementary-material>

REFERENCES

- Alvarez-Perez, F. J., Castelo-Branco, M., and Alvarez-Sabin, J. (2011). Usefulness of measurement of fibrinogen, D-dimer, D-dimer/fibrinogen ratio, C reactive protein and erythrocyte sedimentation rate to assess the pathophysiology and mechanism of ischaemic stroke. *J. Neurol. Neurosurg. Psychiatry* 82, 986–992. doi: 10.1136/jnnp.2010.230870
- Barone, F. C., and Feuerstein, G. Z. (1999). Inflammatory mediators and stroke: new opportunities for novel therapeutics. *J. Cereb. Blood Flow Metab.* 19, 819–834. doi: 10.1097/00004647-199908000-00001
- Bowden, J., Davey Smith, G., and Burgess, S. (2015). Mendelian randomization with invalid instruments: effect estimation and bias detection through egger regression. *Int. J. Epidemiol.* 44, 512–525. doi: 10.1093/ije/dyv080

- Bowden, J., Davey Smith, G., Haycock, P. C., and Burgess, S. (2016). Consistent estimation in Mendelian randomization with some invalid instruments using a weighted median estimator. *Genet. Epidemiol.* 40, 304–314. doi: 10.1002/gepi.21965
- Burgess, S., Butterworth, A., and Thompson, S. G. (2013). Mendelian randomization analysis with multiple genetic variants using summarized data. *Genet. Epidemiol.* 37, 658–665. doi: 10.1002/gepi.21758
- Burgess, S., and Thompson, S. G. (2017). Interpreting findings from Mendelian randomization using the MR-egger method. *Eur. J. Epidemiol.* 32, 377–389. doi: 10.1007/s10654-017-0255-x
- Cao, J. J., Arnold, A. M., Manolio, T. A., Polak, J. F., Psaty, B. M., Hirsch, C. H., et al. (2007). Association of carotid artery intima-media thickness, plaques, and C-reactive protein with future cardiovascular disease and all-cause mortality: the cardiovascular health study. *Circulation* 116, 32–38. doi: 10.1161/circulationaha.106.645606
- Coull, B. M., Beamer, N., de Garmo, P., Sexton, G., Nordt, F., Knox, R., et al. (1991). Chronic blood hyperviscosity in subjects with acute stroke, transient ischemic attack, and risk factors for stroke. *Stroke* 22, 162–168. doi: 10.1161/01.str.22.2.162
- Dalmon, J., Laurent, M., and Courtois, G. (1993). The human beta fibrinogen promoter contains a hepatocyte nuclear factor 1-dependent interleukin-6-responsive element. *Mol. Cell. Biol.* 13, 1183–1193. doi: 10.1128/mcb.13.2.1183
- Das, S., Roy, S., Kaul, S., Jyothi, A., and Munshi, A. (2014). CRP gene (1059G>C) polymorphism and its plasma levels in ischemic stroke and hemorrhagic stroke in a south Indian population. *Inflammation* 37, 1683–1688. doi: 10.1007/s10753-014-9897-y
- Feigin, V. L., Nguyen, G., Cercy, K., Johnson, C. O., Alam, T., Parmar, P. G., et al. (2018). Global, regional, and country-specific lifetime risks of stroke, 1990 and 2016. *N. Engl. J. Med.* 379, 2429–2437. doi: 10.1056/NEJMoa1804492
- Folsom, A. R., Gottesman, R. F., Appiah, D., Shahar, E., and Mosley, T. H. (2016). Plasma d-dimer and incident ischemic stroke and coronary heart disease: the atherosclerosis risk in communities study. *Stroke* 47, 18–23. doi: 10.1161/STROKEAHA.115.011035
- GBD 2015 Mortality and Causes of Death Collaborators (2016). Global, regional, and national life expectancy, all-cause mortality, and cause-specific mortality for 249 causes of death, 1980–2015: a systematic analysis for the global burden of disease study 2015. *Lancet* 388, 1459–1544. doi: 10.1016/s0140-6736(16)31012-1
- Georgakaki, M. K., Malik, R., Björkbacka, H., Pana, T. A., Demissie, S., Ayers, C., et al. (2019). Circulating monocyte chemoattractant protein-1 and risk of stroke: meta-analysis of population-based studies involving 17 180 individuals. *Circ. Res.* 125, 773–782. doi: 10.1161/CIRCRESAHA.119.315380
- Hartwig, F. P., Borges, M. C., Horta, B. L., Bowden, J., and Davey Smith, G. (2017). Inflammatory biomarkers and risk of schizophrenia: a 2-sample Mendelian randomization study. *JAMA Psychiatry* 74, 1226–1233. doi: 10.1001/jamapsychiatry.2017.3191
- Iso, H., Noda, H., Ikeda, A., Yamagishi, K., Inoue, M., Iwasaki, M., et al. (2012). The impact of C-reactive protein on risk of stroke, stroke subtypes, and ischemic heart disease in middle-aged Japanese: the Japan public health center-based study. *J. Atheroscler. Thromb.* 19, 756–766. doi: 10.5551/jat.11999
- Jiménez, M. C., Rexrode, K. M., Glynn, R. J., Ridker, P. M., Gaziano, J. M., and Sesso, H. D. (2015). Association between high-sensitivity C-reactive protein and total stroke by hypertensive status among men. *J. Am. Heart Assoc.* 4:e002073. doi: 10.1161/jaha.115.002073
- Jiménez, M. C., Rexrode, K. M., Kotler, G., Everett, B. M., Glynn, R. J., Lee, I. M., et al. (2016). Association between markers of inflammation and total stroke by hypertensive status among women. *Am. J. Hypertens.* 29, 1117–1124. doi: 10.1093/ajh/hpw050
- Karim, M. A., Kartsonaki, C., Bennett, D. A., Millwood, I. Y., Hill, M. R., Avery, D., et al. (2020). Systemic inflammation is associated with incident stroke and heart disease in east Asians. *Sci. Rep.* 10:5605. doi: 10.1038/s41598-020-62391-3
- Lawlor, D. A., Harbord, R. M., Sterne, J. A., Timpson, N., and Davey Smith, G. (2008). Mendelian randomization: using genes as instruments for making causal inferences in epidemiology. *Stat. Med.* 27, 1133–1163. doi: 10.1002/sim.3034
- Libby, P., Ridker, P. M., and Hansson, G. K. (2011). Progress and challenges in translating the biology of atherosclerosis. *Nature* 473, 317–325. doi: 10.1038/nature10146
- Ligthart, S., Vaez, A., Vösa, U., Stathopoulou, M. G., de Vries, P. S., Prins, B. P., et al. (2018). Genome analyses of >200,000 individuals identify 58 loci for chronic inflammation and highlight pathways that link inflammation and complex disorders. *Am. J. Hum. Genet.* 103, 691–706. doi: 10.1016/j.ajhg.2018.09.009
- Liu, Y., Wang, J., Zhang, L., Wang, C., Wu, J., Zhou, Y., et al. (2014). Relationship between C-reactive protein and stroke: a large prospective community based study. *PLoS One* 9:e107017. doi: 10.1371/journal.pone.0107017
- Pepys, M. B., and Hirschfield, G. M. (2003). C-reactive protein: a critical update. *J. Clin. Invest.* 111, 1805–1812. doi: 10.1172/JCI18921
- Pierce, B. L., and Burgess, S. (2013). Efficient design for Mendelian randomization studies: subsample and 2-sample instrumental variable estimators. *Am. J. Epidemiol.* 178, 1177–1184. doi: 10.1093/aje/kwt084
- Qureshi, A. I., Mendelow, A. D., and Hanley, D. F. (2009). Intracerebral haemorrhage. *Lancet* 373, 1632–1644. doi: 10.1016/S0140-6736(09)60371-8
- Roudbary, S. A., Saadat, F., Forghanparast, K., and Sohrabnejad, R. (2011). Serum C-reactive protein level as a biomarker for differentiation of ischemic from hemorrhagic stroke. *Acta Med. Iran.* 49, 149–152.
- Sabater-Lleal, M., Huang, J., Chasman, D., Naitza, S., Dehghan, A., Johnson, A. D., et al. (2013). Multiethnic meta-analysis of genome-wide association studies in >100 000 subjects identifies 23 fibrinogen-associated loci but no strong evidence of a causal association between circulating fibrinogen and cardiovascular disease. *Circulation* 128, 1310–1324. doi: 10.1161/CIRCULATIONAHA.113.002251
- Sato, S., Iso, H., Noda, H., Kitamura, A., Imano, H., Kiyama, M., et al. (2006). Plasma fibrinogen concentrations and risk of stroke and its subtypes among Japanese men and women. *Stroke* 37, 2488–2492. doi: 10.1161/01.STR.0000242473.13884.8e
- Scirica, B. M., and Morrow, D. A. (2006). Is C-reactive protein an innocent bystander or proatherogenic culprit? The verdict is still out. *Circulation* 113, 2128–2151. doi: 10.1161/circulationaha.105.611350
- Shi, H., Leng, S., Liang, H., Zheng, Y., and Chen, L. (2016). Association study of C-reactive protein associated gene HNF1A with ischemic stroke in Chinese population. *BMC Med. Genet.* 17:51. doi: 10.1186/s12881-016-0313-3
- Smith, G. D., and Ebrahim, S. (2003). 'Mendelian randomization': can genetic epidemiology contribute to understanding environmental determinants of disease? *Int. J. Epidemiol.* 32, 1–22. doi: 10.1093/ije/dyg070
- Stokum, J. A., Kurland, D. B., Gerzanich, V., and Simard, J. M. (2015). Mechanisms of astrocyte-mediated cerebral edema. *Neurochem. Res.* 40, 317–328. doi: 10.1007/s11064-014-1374-3
- Sturgeon, J. D., Folsom, A. R., Longstreth, W. T. Jr., Shahar, E., Rosamond, W. D., and Cushman, M. (2008). Hemostatic and inflammatory risk factors for intracerebral hemorrhage in a pooled cohort. *Stroke* 39, 2268–2273. doi: 10.1161/STROKEAHA.107.505800
- Torzewski, M., Rist, C., Mortensen, R. F., Zwaka, T. P., Bienek, M., Waltenberger, J., et al. (2000). C-reactive protein in the arterial intima: role of C-reactive protein receptor-dependent monocyte recruitment in atherosclerosis. *Thromb. Vasc. Biol.* 20, 2094–2099. doi: 10.1161/01.atv.20.9.2094
- Verbanck, M., Chen, C. Y., Neale, B., and Do, R. (2018). Detection of widespread horizontal pleiotropy in causal relationships inferred from Mendelian randomization between complex traits and diseases. *Nat. Genet.* 50, 693–698. doi: 10.1038/s41588-018-0099-7
- Welsh, P., Lowe, G. D., Chalmers, J., Campbell, D. J., Rumley, A., Neal, B. C., et al. (2008). Associations of proinflammatory cytokines with the risk of recurrent stroke. *Stroke* 39, 2226–2230. doi: 10.1161/STROKEAHA.107.504498
- Wensley, F., Gao, P., Burgess, S., Kaptoge, S., Di Angelantonio, E., Shah, T., et al. (2011). Association between C reactive protein and coronary heart disease: Mendelian randomisation analysis based on individual participant data. *BMJ* 342:d548. doi: 10.1136/bmj.d548
- Woo, D., Falcone, G. J., Devan, W. J., Brown, W. M., Biffi, A., Howard, T. D., et al. (2014). Meta-analysis of genome-wide association studies identifies 1q22 as a susceptibility locus for intracerebral hemorrhage. *Am. J. Hum. Genet.* 94, 511–521. doi: 10.1016/j.ajhg.2014.02.012
- Woodward, M., Lowe, G. D., Campbell, D. J., Colman, S., Rumley, A., Chalmers, J., et al. (2005). Associations of inflammatory and hemostatic variables with the risk of recurrent stroke. *Stroke* 36, 2143–2147. doi: 10.1161/01.STR.0000181754.38408.4c
- Xue, Y., Zhang, L., Fan, Y., Li, Q., Jiang, Y. A. -O., and Shen, C. (2017). C-reactive protein gene contributes to the genetic susceptibility of hemorrhagic

- stroke in men: a case-control study in Chinese Han population. *J. Mol. Neurosci.* 62, 395–401. doi: 10.1007/s12031-017-0945-6
- Yoshinaga, R., Doi, Y., Ayukawa, K., and Ishikawa, S. (2017). High-sensitivity C reactive protein as a predictor of inhospital mortality in patients with cardiovascular disease at an emergency department: a retrospective cohort study. *BMJ Open* 7:e015112. doi: 10.1136/bmjopen-2016-015112
- Zhang, Y., Zhu, C. -G., Guo, Y. -L., Xu, R. -X., Li, S., Dong, Q., et al. (2014). Higher fibrinogen level is independently linked with the presence and severity of new-onset coronary atherosclerosis among Han Chinese population. *PLoS One* 9:e113460. doi: 10.1371/journal.pone.0113460
- Zhou, Y., Han, W., Gong, D., Man, C., and Fan, Y. (2016). Hs-CRP in stroke: a meta-analysis. *Clin. Chim. Acta* 453, 21–27. doi: 10.1016/j.cca.2015.11.027

Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Copyright © 2021 Wang, Zhang, Liu, Zhang, Cao, Tian, Maranga, Meng, Tian, Tian, Cao, Wang, Song and Wang. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.