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High-sensitivity C-reactive protein in stroke patients – The importance in consideration of influence of multiple factors in the predictability for disease severity and death

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Abstract

High sensitivity C-reactive protein (hsCRP) has been evaluated as a biomarker in stroke and relevant pathological diseases. While its predictive values in several pathological phenotypes have been confirmed, controversy exists among different studies. This review summarizes reports of the predictive values of hsCRP for the diagnosis, etiology, prognosis and mortality of stroke diseases. The current literature suggests that CRP expression is influenced by multiple factors, such as polymorphisms, the genomic backgrounds and gender. However, few reported studies analyzed data based on all these multiple factors. Future studies should focus on comprehensive analysis based on multiple factors.

Keywords

Atherosclerosis; High sensitivity C-reactive protein (hsCRP); Ischemic; Polymorphism; Predictability; Stroke

1. Introduction

Stroke is a major cerebrovascular disease threatening human health and life with high morbidity, disability and mortality [1]. According to the data from Global Burden of Disease

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Study [2], worldwide in 2010 there were an estimated 11,569,538 incident ischemic strokes and 5,324,997 events of incident hemorrhagic stroke; furthermore, 2,835,419 individuals died from ischemic stroke and 3,038,763 from hemorrhagic stroke. Stroke is the number one cause of the death in several countries such as China [3]. Stroke was defined as a sudden onset of loss of global or focal cerebral function persisting for more than 24 h [4]. Biomarkers that predict the occurrence and outcome from ischemic stroke are critical for prevention and treatment. Serum biomarkers are considered to be the most valuable adjunct to routine clinical examination and imaging data [5]. Inflammation has an important role in the development of atherosclerosis and during the ischemic event. Inflammatory markers such as fibrinogen and hsCRP have been reported as a predictable marker for the stroke severity and outcome [6]. It has been reported that it is possible to use the increase in the concentration of acute phase reactants especially the hsCRP to help predict cerebrovascular mortality [7].

Human CRP (Pentraxin 1, Ptx1) is an acute phase reactant that is rapidly produced by the liver after tissue injury or infection [5,7]. CRP is a sensitive indicator of inflammation [8,9]. Although many inflammatory biomarkers have been reported to be useful in predicting clinical outcome after stroke, hsCRP remains one of the most widely used in clinical practice [8–12]. Increased hsCRP has been associated with the development of atherosclerosis, ischaemic attacks, hemorrhagic stroke, as well as disease outcomes [13–17]. Studies suggest that post-stroke inflammatory responses may exacerbate tissue damage after cerebral infarction and affect clinical outcomes [18,19]. Recent studies have shown that elevated hsCRP values independently predict the risk of future cardiovascular diseases and ischemic cerebrovascular diseases, including transient ischemic attack in the elderly [20–22]. After acute ischemic stroke (AIS), a sustained inflammatory response indicated by increased level of CRP has been reported in about 75% of patients. A strong and persistent inflammatory response is associated with a worse outcome. Moreover, CRP at discharge has been shown to be related to a 1-year outcome [23–25]. Recently, a meta-analysis suggested that elevated baseline hsCRP levels are independently associated with excessive ischemic stroke risk.

However, the physiologic role of CRP is not well understood; it potentially has anti-inflammatory properties as well as proinflammatory effects [26]. Inflammation may not only be the consequence of brain infarction, it may also contribute to ischemic damage. In addition, the role of inflammatory markers in predicting functional outcome in stroke remains controversial [27]. This review summarizes the data on the predictive value of hsCRP for ischemic stroke, analyzes the potential influence of CRP on functional outcomes, and identifies the critical issues that are important for the future studies.

2. CRP

CRP (protein CRP) is named because of its precipitation reaction with phosphocholine moiety of pneumococcal cell wall C-polysaccharide. According to the information on Ensembl data base (http://useast.ensembl.org/Homo_sapiens/Gene/Summary?db=core;g=ENSG00000132693;r=1:159712289-159714589), the CRP gene is located on human chromosome 1, between 159,712,289 and 159,714,589 bps. It has 7

transcripts, of which 5 encode proteins. Although the detailed functions of these protein variants are not fully known, the protein with 224 aa produced by the CRP-001 transcript is commonly known as the CRP protein. The protein is a member of the pentraxin family of proteins. The pentraxins are an ancient family of proteins with a unique architecture found as far back in evolution as the Horseshoe crab. CRP structure is strong phylogenetic conservation and high resistance to proteolysis [28]. Therefore, it is relatively stable and easy to be measured.

Expression of CRP is regulated mainly at the transcriptional level by cytokines such as interleukin-1, interleukin-6, and tumor necrosis factor; among them, interleukin-6 is the primary circulating physiologic mediator [26]. Recent studies have examined the effects of IL-1 β , the IL-1 receptor antagonist anakinra and the IL-6 receptor blocker tocilizumab on the regulation of expression of CRP [29]. An estimation suggests that inter-individual variation in blood CRP level is 35–40% heritable [30] and it is affected by gene polymorphisms. The expression of CRP is also influenced by environmental factors such as bacterial infection and injury. Plasma CRP concentrations increase rapidly and dramatically (100-fold or more) in response to tissue injury or inflammation. Therefore, high levels of CRP (10–1000 mg/L), are associated with serious infections or inflammatory diseases such as arthritis. The hsCRP, on the other hand, measures CRP in the low range, from 0.5 to 10 mg/L. At such range, the hsCRP test is used to identify low but persistent levels of inflammation. The hsCRP is more precise than standard CRP when measuring baseline concentrations. Induction of an hsCRP is rapid and its half-life is long enough in the blood for a steady time course. Therefore, plasma hsCRP is very useful for the diagnostic workup of inflammatory and infectious diseases [26].

3. HsCRP level in early diagnosis of stroke

At present, the diagnosis of stroke depends on clinical examination and neuro-imaging techniques [31]. It is critically important to identify effective biomarkers for the early diagnosis of acute stroke and to establish biological procedures for early diagnosis [5]. Early studies of different populations suggested that modest elevation of hsCRP is a predictor of future vascular events in apparently healthy individuals [32,33]. However, there are few data on the value of hsCRP for the early diagnosis of ischemic stroke.

In 2012, Youn et al. reported in the Keran population that higher hsCRP levels were associated with larger infarct volumes in AIS (Table 1) [26]. Their results suggest that elevated hsCRP levels may serve as a helpful serologic marker in the evaluation of the severity of AIS [34]. Kanai et al. investigated the reciprocal association between serum levels of hsCRP (Table 1) and intercellular adhesion molecule-1 (sICAM-1) and the risk of brain infarction in type 2 diabetic patients [18]. Their study suggests that both measurements of hsCRP and sICAM-1 levels are useful as a predictor of future stroke in diabetic subjects [34]. However, these studies are in Asian population. These findings have not been confirmed in the other regions such as US and European countries.

4. HsCRP in the prediction of atherosclerosis—artery stenosis or occlusion

Ischaemic stroke accounts for 70–80% of all cases of stroke [35]. Ischaemic stroke is mainly caused by atherosclerosis and thrombotic obstruction of cerebral blood flow (Table 1) [36,37]. Atherosclerosis, as a complex and systemic disease, may unequally induce intracranial stenosis, which may in turn impose the limitation of particular inflammatory mediator in describing atherosclerosis [38,39]. Middle cerebral artery stenosis is also associated with advanced age, hypertension, diabetes mellitus and hyperlipidaemia. The risk of stroke increases with the prevalence of atherosclerosis. Therefore, identification of biomarkers for atherosclerosis—artery stenosis or occlusion is essential for the early prevention of stroke. Considerable studies have already shown that an elevated serum CRP level reflects an increased tendency for plaque rupture and a high atherosclerotic burden [13,32,33,40–42].

In 2011, Whiteley et al. showed that higher levels of hsCRP, along with IL-6 and fibrinogen measured after stroke onset in 817 patients are significantly associated with increased incidence of occlusive vascular events and vascular death, as well as nonvascular causes of death after stroke [43]. In a prospectively study of intracranial large artery atherosclerosis progression in 25 of 75 patients using transcranial Doppler follow-up during a median follow-up time of 23 months, Arenillas et al. reported that hsCRP and plasminogen activator inhibitor-1 measured 3 months after onset of ischemic stroke are highly predictive of intracranial large artery atherosclerosis progression [44]. Similarly, Yoshida et al. showed that CRP, along with IL-6 and protein-conjugated acrolein (PC-Acro) is the risk factor for carotid atherosclerosis [45]. In a study on the transcription levels of ABCA1, ABCG1 and SR-BI and plasma CRP in Chinese populations with various risk factors for atherosclerosis, Li et al. reported that circulating CRP was increased almost in all the risk populations except in males [46]. Similarly, hsCRP and Lp-PLA2 was found to improve ischemic stroke risk prediction in the Atherosclerosis Risk in Communities study [47].

At the same time, there are also negative reports. In a study of the association between inflammatory biomarkers and progression of intracranial large artery stenosis after ischemic stroke, Shimizu and colleagues reported that the Cox proportional hazard analysis did not show a significant association between hsCRP and intracranial large artery atherosclerosis progression. The authors concluded that, because the effects of inflammatory factors may continuously affect the vessel walls during the post stroke period, the biomarkers may be more predictive in the chronic rather than the acute stage of stroke and may be important for ILA progression [41]. In a study of the influence of polyvascular atherothrombotic disease on stroke patient prognosis, its relation with inflammatory markers and the progression of atherothrombotic disease, Blanco et al. found a non-significant trend to higher hsCRP levels in patients with vascular recurrences [19]. The authors explained that this is possibly because the calculation of the sample size was based on incorrect assumptions regarding IL-6 levels [42].

The above data suggest that the hsCRP is associated with the progression of atherosclerosis and possibly with artery stenosis or occlusion. The failure to detect significant relationship between hsCRP and the progression of atherosclerosis may be due to limited sample size.

5. hsCRP and death risk

An early risk assessment of the severity of disease and prognosis using biomarkers is pivotal for optimized care and the allocation of healthcare resources [48]. The predictability of death risk for the stroke patients by hsCRP has been evaluated extensively (Table 1). Current research indicated that expression of hsCRP can be used to predict mortality from as short as three months to as long as 5 year.

At least two studies have evaluated the value of hsCRP for predicting the risk of early death of stroke patients. In a prospective study of 189 patients with AIS who were admitted within 24 h after the onset of symptoms, Tu et al. measured short-term functional outcome by the modified Rankin scale (mRS), 90 days after admission. They concluded that hsCRP was independent predictor of short-term outcome and mortality (Table 1) after AIS [49]. Huang et al. (Table 1) found that elevated plasma hsCRP independently predicted risk of all-cause death within 3 months after acute stroke in Chinese patients. Compared with low hsCRP, patients with high hsCRP (>3 mg/L) (Table 1) had a significantly higher rate of all-cause death (0.71% vs. 10.00%; $p < 0.001$) at 3 months after stroke onset [50]. Their study suggests that the level of hsCRP is a better predictor of risk of death than for disease relapse. These studies suggest that shortly after the onset of AIS, increased CRP levels may reflect an accompanying inflammatory reaction. Elevated CRP levels may reflect the extent of brain injury. Therefore, the elevated level of hsCRP can predict the risk of early death of stroke patients.

Li and Liu prospectively studied one year outcomes in 374 patients with ischemic stroke who were admitted within 24 h after the onset of symptoms. They measured serum levels of PCT, hsCRP and NIH stroke scale of the patients at the time of admission. Clinical follow-up was performed at 1 year. Their data showed that hsCRP on admission was independent predictor of 1 year mortality with an adjusted hazard ratio of 15.37 (95% confidence interval [CI], 3.25–41.08) [48].

Several prospective studies have demonstrated that high hsCRP was an independent predictor of long-term mortality in patients with stroke. Idicula et al. (Table 1) prospectively studied 498 patients with ischemic stroke who were admitted within 24 h after the onset of symptoms [51]. Their investigation indicated that elevated hsCRP concentration was an independent predictor of long-term mortality over a period of 2.5 years after ischemic stroke ($p = 0.002$). Their results also suggested that hsCRP concentration was also an independent predictor of short-term mortality after supratentorial intracerebral hemorrhage [51]. From a prospective cohort study, Makita et al. (Table 1) found that baseline serum hsCRP level was an independent predictor for future ischemic stroke and all-cause mortality in apparently healthy Japanese men. In their study, the mean follow-up period was 2.7 years [12].

The 5-year is the longest reported follow-up period. Chan et al. [52] investigated biomarkers of various pathophysiologic pathways including hsCRP with mortality in stroke patients (Table 1). During the 5-year follow-up, the mortality was 37.0% (34 out of 92). Authors concluded that hsCRP was the most significant independent predictor of both short- and long-term mortality after stroke. Mitchell et al. [53] admitted 467 patients with first ischemic stroke; at 5-year follow-up, they concluded that hsCRP was associated with risk of death. Similarly, several other reports have suggested that increased hsCRP level is associated with high risk of death stroke patients with variety of mortality rate. These studies included Hartmann et al. (41.0% mortality rate) [54], Whiting et al. (60.0% mortality rate) [55], Kammersgaard et al. (58.4%) [56], and Putaala et al. (10.7%) [57].

At present, hsCRP is probably the only independent prognostic factors in predicting both short- and long-term mortality.

6. Application in etiological subtypes of ischemics stroke

Like many known disease, ischemic stroke is a group of diseases with many different etiologies. At present, precise etiological classification depends heavily on the results of auxiliary examination. However, most of these examinations are difficult to complete at the early onset of a stroke because of the emergency nature of the disease. Based on the TOAST etiology classification system, ischemic stroke is divided into five categories (Table 1): largeartery atherosclerosis (LAA), cardioembolism (CE), small-artery occlusion (SAA), stroke of other determined etiology (SOE), and stroke of undetermined etiology (SUE) [58,59]. As hsCRP levels vary in apparently healthy population and in stroke patients, it might be a useful marker for the ischemic stroke classification. Liu et al. found that hsCRP and D-dimer levels were significantly different among the subtypes and were the highest in CE (cardioembolism). Using the data, hsCRP > 6.96 mg/L was classified as the CE subtype, with a sensitivity of 41% and a specificity of 74% [58]. The authors suggested that using a combination of hsCRP and DD may improve the diagnostic accuracy of CE in the acute phase [58]. Future studies on this aspect will improve the classification of the of stroke subtypes.

7. hsCRP and prognosis

CRP levels strongly correlate with stroke severity and independently predict mortality and stroke recurrence in AIS patients (Table 1) [60], but only hsCRP is an independent predictor for unfavorable outcome [61]. The level of hsCRP can predict the severity of AIS patients. By examining the level of hsCRP in 316 patients with AIS, Luo et al. found that elevated CRP level can reflect the severity of AIS [61].

The different time points of blood sampling for the measurement of hsCRP levels are also studied. Yeh et al. studies the hsCRP levels during both acute and convalescent phases of 160 patients [27]. They found that Hs-CRP levels during both acute phase and convalescent phase of IS were significantly and independently predictive of 90-day major adverse neurological event (MANE) [27]. Authors also found that circulating level of hsCRP at 48 h was better than the level at day 21 after acute IS for predictive of 90-day MANE

[27]. Their finding supports the validity of using serum hsCRP level in predicting clinical outcome in patients with and without IS regardless of the timing for blood sampling [27].

During the subacute phase of stroke, hsCRP values are associated with the level of functional disability. Song et al. examined the hsCRP level in a total of 309 patients with first-ever stroke within 24 h after symptom onset [62]. The hsCRP measurements were made at admission and on the seventh hospital day. Their study showed that both hsCRP values on admission and on the seventh hospital day were significantly correlated with modified Rankin Scale (mRS) scores obtained at 4 times after the onset of stroke. Authors suggest that hsCRP values on the subacute phase have sufficient value as predictors of the prognosis of functional disability after first-ever stroke [62].

However, the prognostic value of hsCRP on stroke has its limitations. Elkind et al. found that, after adjusting for confounders, hsCRP was not associated with risk of recurrent stroke or recurrent stroke, myocardial infarction, or vascular death but was associated with risk of death [53]. In a study to investigate the association between the progression of silent cerebral infarction (SCI) and levels of soluble adhesion molecules and high-sensitivity C-reactive protein (hsCRP) in type 2 diabetic patients, Kawamura et al. reported that hsCRP was not associated with the progression of SCI. By examining the hsCRP and ferritin levels of 62 patients prospectively within 48 h after onset of ischemic stroke, Ozkan et al. (Table 1) reported that neither hsCRP nor ferritin levels could predict functional disability 3 months after stroke onset [27].

8. CRP and hemorrhagic stroke

Studies so far indicate that hsCRP is not a predictive marker for hemorrhagic stroke. Roudbary et al. assessed the concentration of hsCRP in patients with documented ischemic and hemorrhagic stroke in the first 24 h of the onset of symptoms [17]. Their study revealed that types of stroke were related to elevated hsCRP levels [17]. Serum level of hsCRP (Table 1) in ischemic patients was 18.92 ± 11.28 and in hemorrhagic group was 2.65 ± 1.70 . This difference was statistically significant ($p < 0.0001$) [17]. They concluded that hsCRP level is increased in patients with ischemic but not in hemorrhagic stroke, which may make it a useful adjunctive method for determining type of stroke. At present other markers, but not hsCRP, are potentially used in the diagnosis of hemorrhagic stroke.

9. Other neurological and related conditions

In addition its prediction value in the stroke, hsCRP has also been investigated for its prediction values in stroke relevant diseases. Wersching et al. examined 447 community-dwelling and stroke-free individuals from the Systematic Evaluation and Alteration of Risk Factors for Cognitive Health (SEARCH) Study. They found that serum CRP is linked to cerebral microstructural integrity and cognitive function [63]. Song et al. examined 212 patients with de novo Parkinson's disease (PD), 253 patients with acute ischemic cerebrovascular disease and 119 healthy subjects and investigated the differences in hsCRP among these 3 groups. Their results suggested that elevated hsCRP might have a clinical value as a risk factor for Parkinson's disease [15]. In a study of CRP and lipid levels in

women Everett et al. found that hsCRP associated more closely with ischemic stroke than with CHD. They suggested that concomitant evaluation of lipid levels and hsCRP may improve risk assessment for stroke as well as CHD [11]. These and many other studies have indicated that hsCRP is associated with large number of diseases and disorders. Currently, the mechanistic linkages between CRP and these diseases are not completely understood.

10. CRP polymorphism influence the level of hsCRP

Like many other genes, the expression level of CRP is influenced by its polymorphism [64] (Fig. 1). Polymorphism of CRP has been investigated in different ethnic groups and different diseases. Sheu et al. [65] measured CRP concentrations and investigated 6 polymorphisms of CRP gene in 369 Chinese men. They found that CRP 1009A>G genotypes and associated haplotypes were associated with lower fasting serum hsCRP concentrations in a group of elderly Chinese men. Liu et al. [66] then investigated the potential association of the CRP gene + 1444C/T polymorphism with symptomatic carotid artery stenosis. They found that CRP + 1444 CC genotype is a risk factor for >70% carotid artery stenosis. The serum CRP level was associated with the presence of carotid stenosis.

Kettunen et al. investigated 5 polymorphisms and measured CRP concentration of hsCRP in men in a Finnish middle-aged study population. They found that CRP gene allelic variation is associated with CRP levels in both sexes and with one carotid artery elasticity parameter (SI) in men in a population-based Finnish cohort of middle-aged subjects [64]. Assayag genotyped the -757T>C CRP gene SNP and determined the concentration of serum CRP, the intima-media thickness (IMT) of the common carotid artery and the existence of plaque/s in 612 apparently healthy men and women aged 66 ± 10 years in Israel [67]. Their study provided evidence that CRP-757C allele of CRP gene is associated with carotid atherosclerosis, independently of traditional vascular risk factors.

By studying the C1444T polymorphism of CRP 75 in consecutive patients with a first-ever cerebral ischemic event attributable to symptomatic intracranial atherosclerotic disease (ICAD) [68], Arenillas et al. suggested that presence of the T allele within the CRP gene C1444T polymorphism is associated with a higher risk of further ischemic events in symptomatic ICAD patients.

Polymorphisms in other locations in the genome are also found to influence the level of hsCRP. Bernardo et al. investigated the Influence of the CD14 C260T promoter polymorphism on C-reactive protein levels in patients with coronary artery disease [69]. Their finding indicated that homozygosity status of the T allele was independently associated with hsCRP levels and suggested that T homozygotes of this functional polymorphism are at increased ischemic risk.

However, there are negative reports suggest there is no association of polymorphisms of CRP with disease. Ghattaset al reported that CRP 1059G/C gene variation influences plasma CRP levels. Conversely, this polymorphism was not associated with the risk for acute myocardial infarction in an Egyptian population [70]. Huang assessed the potential genetic determinants for serum hsCRP levels in a cohort with well-controlled nondiabetic

hypertension. They reported that CRP genotype had a limited effect on serum hsCRP levels in subjects with well-controlled hypertension; suggesting that the impacts of clinical rather than genetic determinants influence serum hsCRP levels for cardiovascular risk stratification in this intermediate-risk Chinese Taiwanese cohort [71].

These results above probably are the results of variation of function of the polymorphism, different ethnic genetic backgrounds, and potential gene-environmental interaction.

11. Ethnic difference on the prediction values of hsCRP

The predictive value of hsCRP may vary among different ethnic groups (Fig. 1). At present, data on the early diagnosis based on the expression level of hsCRP are all from Asian population [26,34]. The reports on prediction values on diagnosis of disease recurrence and major adverse neurological events are also from Asian population [27,61,62]. Although the prediction of death rate is well recognized among all ethnic groups, two studies that reported the prediction values of hsCRP on the risk of early death of stroke patients are all from Asian [49,50]. It is not clear whether it is true that some polymorphism are only associated with men [64,65] while others are associated with both sexes, and some of the polymorphism do not have prediction values. The values of hsCRP expression level for the prediction of atherosclerosis—artery stenosis or occlusion have been recognized in a variety of ethnic groups. However, there is a possibility that it predicts different phenotypes in different populations [35–47]. Therefore, it is most likely that the level of hsCRP and its prediction values on variety of stroke phenotypes are influenced by the ethnic genome background.

12. Summary and future direction

It is clear that hsCRP has an important role in the prognosis and risk factor prediction in stroke diseases, including long term and short time death risk, early diagnosis, prediction of etiological subtypes, and prognosis (Table 2). CRP is potentially valuable for the prediction of atherosclerosis (Table 2). Its value on the prediction is unknown for on idiopathic Parkinson's disease, cerebral microstructural integrity and cognitive function, and Hemorrhage. Reports on its predictive value on functional disability, Cognitive decline in late midlife, and genetic influence are controversial (Table 2).

CRP expression and its predictive value are influenced by gene polymorphisms [65], the genomic backgrounds of different racial groups [26,27,35,46,50], sex and the environment (Fig. 1). Therefore it is important to consider these factors when evaluating the predictive value of hsCRP on stroke phenotypes in the future studies. These factors may also contribute to the controversial differences between studies in the value of hsCRP. Based on the current report, hsCRP can be used in early diagnosis of stroke, prediction of atherosclerosis—artery stenosis or occlusion, death risk, and etiological subtypes of ischemic stroke.

In addition, hsCRP may be a valuable marker for some cerebrovascular relevant diseases, but further investigation is necessary. For example, the value of hsCRP levels as risk factors for the initiation of Parkinson's disease [15]. Interestingly, in the prediction of

stroke prognosis, lesion volume and all of the evaluated biochemical parameters are not deterministic factors [72].

In many studies, hsCRP has been investigated together with other biomarkers. In order to increase the sensitivity and specificity of diagnosis, the combination of multi-biomarkers may be necessary [5,27,34,41,47–49,53,58,62,73–75]. These markers may together with hsCRP predict risk in different stages of the stroke and/or phenotypes. For example, recent observations suggest that measurement of serum MMP-9, TIMP-1, and hsCRP levels may be useful in the diagnosis and management of ischemic stroke patients providing a more feasible blood biomarker index [74]. Recently, the association between cerebrovascular disease and adhesion molecules has received much attention. A study suggests that sICAM-1 levels may be a potential marker for silent cerebral infarction, which may lead to future stroke and vascular dementia, and that this marker could be useful in monitoring disease progression and as a surrogate marker in treatment studies [75,76].

Lowering CRP levels may be of value in stroke patients or those with other relevant diseases. It has been reported that statin therapy significantly alters serum hsCRP level in patient after acute ischemic stroke [14]. While more studies may be necessary to determine whether statins improve the clinical outcome among patients with AIS with elevated CRP levels [26], statins should be given as early as possible in patients with acute ischemic stroke, especially those with high hsCRP concentrations [14].

Quick and accurate detection methods for CRP levels should be developed. For example, a real-time, on-line and low-cost widerange -CRP assay is a reasonable alternative to the Behring Dade hsCRP method in acute stroke/transient ischemic attack patients [77].

There are several important questions regarding the role of CRP polymorphisms. It is important to understand whether polymorphisms affect the expression level or function of the CRP, whether the altered level of CRP reflect the risk level in stroke disease, and whether single plays a key role or multiple polymorphisms act together.

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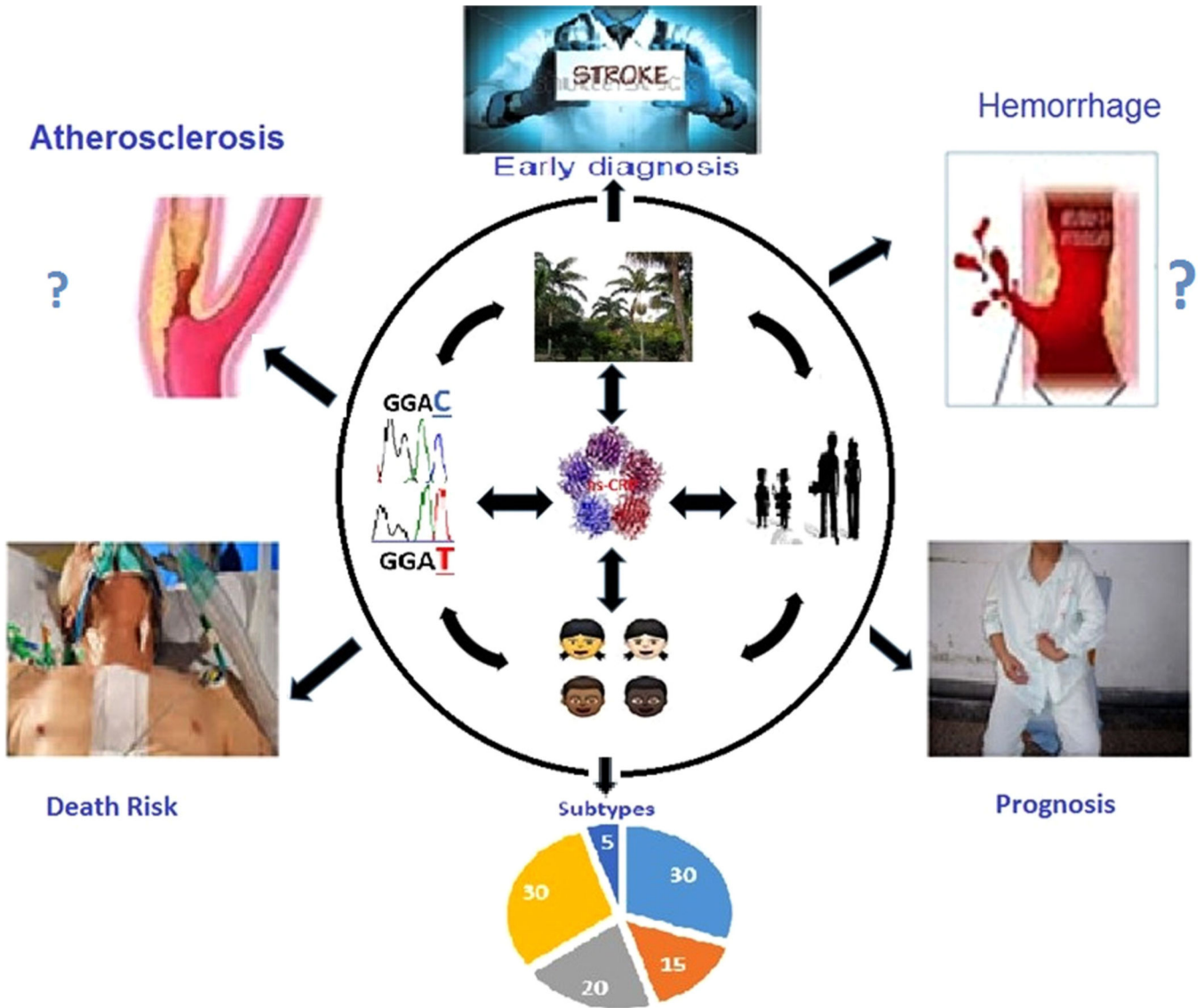


Fig. 1. Multiple factors determine the prediction function of hsCRP on stroke phenotypes. Level of hsCRP is influenced by its gene polymorphism, genomic background of the ethnic groups, gender and it interacts with environment. These multiple factors interact each other to eventually determine the level of hsCRP. Accordingly, the level of hsCRP achieves its values in whether it predicts different disease phenotypes.

Table 1

Potential clinical applications of hsCRP level reported in literature.

Studies	Phenotype	CRP changes	Effect and/or description
Xie et al. [13]	Atherosclerosis	Significantly high	Patients with ICAO (internal carotid artery occlusion), MCAO (middle cerebral artery occlusion) and ICAO + MCAO had significantly higher serum hsCRP concentrations
Youn et al. [26]	Early diagnosis	High	Serve as a helpful serologic marker in the evaluation of the severity of AIS
Kanai et al. [34]		>0.83 mg/L	Measurement the risk of brain infarction in type 2 diabetic patients
Li et al. [48]	Death risk	High	Independent predictor of long-term mortality after ischemic stroke in a Chinese sample
Chan et al. [52]		>3.4 mg/L	Most significant independent predictor of both short- and long-term mortality after stroke
Huang et al. [50]		>3.0 mg/L	Elevated plasma hsCRP independently predicted risk of all-cause death within 3 months after acute IS in Chinese patients
Makita et al. [12]		High	CRP levels can be used to predict future ischemic stroke and mortality in Japanese men from the general population, independently from traditional cardiovascular risk factors
Tu et al. [49]		High	Independent predictors of short-term outcome and mortality after AIS
Elkind et al. [53]		High	Increase with stroke severity and may be associated with mortality to a greater degree than recurrence
Liu et al. [58]	Etiological subtypes of IS	>6.96 mg/L	Different among the subtypes and were the highest in CE(cardioembolism), followed by LAA(large-artery atherosclerosis) and SAA (small-artery occlusion)
Ozkan et al. [27]	Prognosis	>0.5 mg/dL	Predict functional disability 3 months after stroke onset
Gong et al. [60]		>3.0 mg/L	Elevated hsCRP independently predicts 1-year poor outcome in acute stroke
Park et al. [74]		High	hsCRP levels, along with MMP-9, TIMP-1 may be useful in the diagnosis of ischemic stroke patients
Yeh et al. [40]		High	Serum level of hsCRP during both acute and convalescent phases was significantly and independently predictive of 90-day MANE (major adverse neurological event)
Youn et al. [26]		Higher	Reflecting a large volume of infarct, may serve as a helpful serologic marker in the evaluation of severity of acute ischemic stroke
Roudbary et al. [17]	Hemorrhagic stroke	Average About 10.7906 mg/L	hsCRP level is increase in patients with Ischemic infarct dramatically but not in hemorrhagic stroke which might be considered as a usefully adjunct method for determining type of stroke in patients with cerebrovascular problems

Predictive values of CRP on different phenotypes.

Table 2

Valuable	Potentially valuable	Unknown value	Controversial
1. Death risk [12,48-50,52-57]	Atherosclerosis [13,46]	1. Idiopathic Parkinson's disease [15]	1. Cognitive decline in late midlife [78]
2. Etiological subtypes [58,59]		2. Cerebral microstructural integrity and cognitive function [63]	2. Genetic determinants influence [71]
3. Early diagnosis [26,34]		3. Hemorrhage [17]	3. Functional disability [62,79-81]
4. Prognosis [26,27,40,60,74]			