

Clinical Significance of Quantitative D-Dimer Testing IN Suspected Deep Vein Thrombosis

Thesis

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LIST OF ABBREVIATIONS

AAD	Acute aortic dissection
AP	Emergency department.
APAs	Antiphospholipid antibodies
APC	Activated protein C
APTT	Activated partial thromboplastin time
ATIII	Antithrombin III
BMI	Body mass index
CBC	Complete blood count
CCDS	Colour coded duplex sonography
CT	Computed tomography
CTPA	Computed tomographic pulmonary angiography
CTV	Computed tomographic venography
CUS	Compression ultra sonography
DIC	Dissamminated intravascular coagulation
DVT	Deep venous thrombosis
ELISA	Enzyme-linked immuno-sorbent assay
FII	Prothrombin
FRM	Fibrin related marker
HIT	Heparin-induced thrombocytopenia
K⁺-EDTA	Potassium ethylene diamine tetra-acetic
MRI	Magnetic resonance imaging
MTHFR	Methylenetetrahydrofolate reductase
NPV	Negative predictive value
OCPs	Oral Contraceptives pills
PC	Protein C
POC	Point of care
PPP	Platelet Poor Plasma
PPV	Positive predictive value
PS	Protein S

LIST OF ABBREVIATIONS (CONT.)

PSGL-1	P-selectin glycoprotein ligand 1
PT	Prothrombin time
PTT	Partial thromboplastin time
RCL	Reactive centre loop
ROC	Receiver operating characteristic
S	Sensitivity
SD	Standard deviation
SLE	Systemic lupus erythematosus
SNP	Single nucleotide polymorphism
SP	Specificity
T	Thrombin
TLC	Total leucocytic count
TM	Thrombomodulin
TM	Thrombomodulin
t-PA	Tissue plasminogen activator
U-PA	Urokinase-type plasminogen activator
US	Ultra sonography
V/Q Scan	Ventilation-perfusion scintigraphy
Va	Activated factor V
VIIIa	Activated factor VIII
Vitamin B	Pyridoxal phosphate
Vitamin B¹² ...	Cobalamin
VTE	Venous thromboembolism
X²	Chi-square test
A^{20210G}	Prothrombin gene mutation

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INTRODUCTION

Deep venous thrombosis (DVT) is a common condition with significant morbidity and mortality if not diagnosed and treated in a timely manner. The clinical signs and symptoms of DVT are non-specific and objective testing is required for diagnosis (*Heim et al., ۲۰۰۴*).

Accurate diagnosis of deep-vein thrombosis minimizes the risk of thromboembolic complication and averts the exposure of patients without thrombosis to the risk of anticoagulant therapy (*Wells et al., ۲۰۰۳*).

Diagnostic imaging with venography was previously the gold standard for DVT evaluation and may still be utilized if other techniques are inconclusive. Although effective, the technique is invasive, expensive, time consuming, requires specialized personnel and may introduce a variety of complications (*Fox and Bertoglio, ۲۰۱۱*).

Compression ultra sonography (CUS) is a highly accurate method for the detection of deep vein thrombosis in symptomatic outpatients and has replaced venography and other diagnostic methods in common practice (*Bernardi et al., ۲۰۰۸*). However ultrasonography is expensive and time consuming (*Chen et al., ۲۰۰۸*).

Moreover, a significant proportion of out patients may be lost at follow-up and has poor sensitivity for detection of calf vein

thrombosis. In addition, three fourths or more of the patients with suspected DVT have negative ultra sound findings and require repeated imaging to identify the further 2% to 6% in whom occlusive proximal DVT becomes apparent in a week (*Kelly et al., 2007*).

Therefore, diagnostic strategies have been developed aiming at reduction of the need for imaging techniques and instead including D-dimer testing (*Jennersjo et al., 2009*). Plasma D-dimers are generated when the endogenous fibrinolytic system degrades fibrin as in venous thromboembolism, and they consist of 2 identical subunits derived from 2 fibrin molecules. Unlike fibrinogen degradation product, which are derived from fibrinogen and fibrin (*Kelly et al., 2007*). It's a degradation product of cross linked fibrin. D-dimer assays are sensitive but non-specific marker for venous thromboembolism (*Wells, 2007*).

D-dimer testing has become rapid simple and inexpensive and it has the potential to detect thrombosis in any part of the venous system. If the sensitivity of the D-dimer test for deep venous thrombosis consistently very high, its negative predictive value will also be high and reliably exclude the presence of disease. Therefore D-dimer assays has been suggested as an initial test to rule out DVT to reduce the number of patients requiring diagnostic imaging (*Heim et al., 2008*).

Normal result from a highly sensitive D-dimer test effectively ruled out deep vein thrombosis among patients with

either low or moderate clinical. This makes ultrasound testing unnecessary (*Fancher et al., ۲۰۰۴*).

When D-dimer testing is positive, CUS or colour coded duplex sonography (CCDS) should initially be performed to accurately exclude or confirm DVT. In patients with suspected DVT, serial CUS can be avoided if the D-dimer test is negative. Whereas in cases of negative sonographic findings, an additional negative D-dimer test allows exclusion of acute proximal DVT (*Melly et al., ۲۰۰۶*).

AIM OF THE STUDY

The aim of this study is to assess the role of D-Dimer and its effectiveness in diagnosis and ruling out DVT in clinically suspected patients, as well as the possibility of reducing the need of ultrasoungraphy.

THROMBOPHILIA

Hypercoagulable states can be defined as a group of inherited or acquired conditions associated with a predisposition to venous thrombosis, arterial thrombosis or both. Although most inherited conditions appear to increase only the risk of venous thromboembolic events (VTEs), some of the acquired conditions have been associated with both VTEs and arterial thrombosis. These include cancer, myeloproliferative syndromes, antiphospholipid antibodies (APAs), hyperhomocysteinemia, and heparin-induced thrombocytopenia (*Deitcher, 2010*).

Thrombosis becomes more common as age increases and its occurrence is frequently associated with risk factors such as trauma (accidental or surgical), pregnancy, malignant disease, immobility or oral contraceptives. Thrombosis, however, may develop at a younger age and sometimes in the absence of an easily identifiable risk factor. Recently it has become increasingly recognised that patients who have defects or abnormalities which alter the physiological haemostatic balance in favour of fibrin formation or persistence are at increased risk of clinical thrombosis. These patients may be considered to have thrombophilia. It must, however, be realised that many patients with laboratory evidence of a thrombophilic abnormality remain clinically asymptomatic (*Dahlbäck, 2004*).

History of thrombophilia:

Egeberg in 1960 was the first to describe a thrombophilia caused by a hereditary deficiency of antithrombin. Members of the family described in the report suffered from recurrent venous thrombosis, and the disorder was inherited in an autosomal dominant pattern. The deficiency of this naturally occurring anticoagulant protein remained the only identified cause of inherited thrombophilia for many years (*Greer, 2008*).

Since the early 1980s there has been an explosion of new knowledge, with the identification of protein C (PC) deficiency, and three years later, protein S (PS) deficiency was described as additional causes of inherited thrombophilia. However, altogether these three defects only account for less than 10% of selected cases of juvenile and/or recurrent venous thrombosis and for less than 10% of unselected cases, and this was disappointing at that time.

This situation changed dramatically in 1993 when Dahlback and coworkers reported that venous thrombosis often is associated with hereditary resistance to activated protein C (APC) the protease generated by the thrombomodulin-PC anticoagulant pathway to inactivate activated factor V and VIII (Va and VIIIa).

APC resistance is associated with factor V Leiden which is the most frequent cause of inherited thrombophilia, accounting for 20% to 50% of cases (*Greer, 2007*).

Mild hyperhomocysteinemia was found in 19% of patients with juvenile venous thrombosis and family studies showed that in most cases the abnormality was inherited (*Oger et al., 2007*). While the genetic lesions for deficiencies of Antithrombin III deficiency, Protein C deficiency, Protein S deficiency, and activated protein C resistance can be found in single genes encoding the defective proteins, inherited hyperhomocysteinemia may be caused by defects in several genes encoding different enzymes involved in the metabolism of the amino acid (*Oger et al., 2007*).

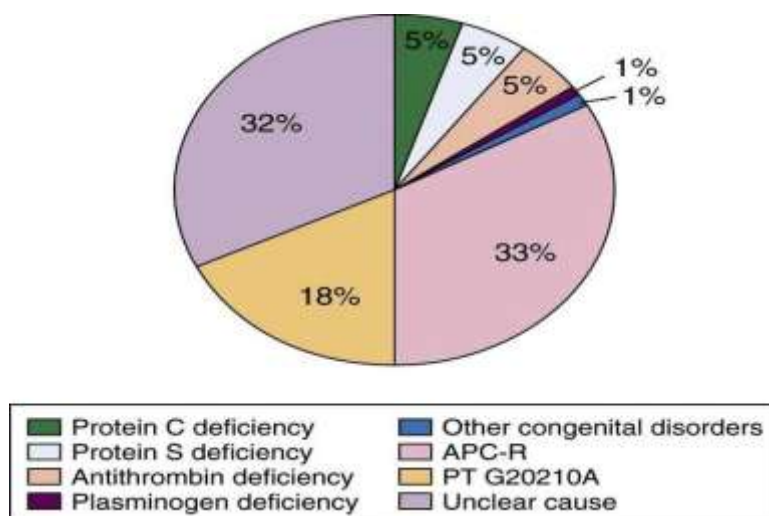


Figure (1): Results of testing for congenital hypercoagulable states. APC-R, activated protein C resistance; PT G20210A, prothrombin G20210A mutation (*Deitcher et al., 2000*).