

La diagnostica di laboratorio delle trombofilie

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Quali alterazioni trombofiliche ricercare

- Difetto di Antitrombina
- Difetto di Proteina C
- Difetto di Proteina S
- Mutazione Fattore V Leiden
- Mutazione G20210A della protrombina
- Lupus Anticoagulant e anticorpi antifosfolipidi
- Disfibrinogenemia
- Aumento Fattore VIII (Fattori IX, XI e altri)
- Iperomocisteinemia

Quando NON eseguire i test

- Trombosi acuta (eseguire la diagnostica per la HIT, se presente sospetto)
- Durante le terapie anticoagulanti
- Gravidanza o terapia ormonale
- In tutte le condizioni associate ad aumento delle proteine della fase acuta (post-chirurgia, mal. infiammatorie)

Se indispensabile, test genetici sempre eseguibili, ma in genere non sono consigliabili screening parziali

Proteina C e gravidanza

- Donna di 36 anni esaminata alla 25° settimana di gravidanza
- Il padre in trattamento anticoagulante per 2 pregressi episodi di TEV (1 TVP prossimale a 30 anni dopo gesso e recidiva idiopatica di TVP a 62 anni)
- Livello della Proteina C durante la gravidanza = 66% (n.v. > 68%)
- 2 anni dopo il parto:
Proteina C attività = 47%
Proteina C antigene = 51%
- Conferma del difetto di Proteina C nello zio paterno

Table 2 Distribution of different mutations within the genes for antithrombin, protein C and protein S

Mutation types	<i>SERPINC1</i>	<i>PROC</i>	<i>PROS1</i>
Missense/nonsense	155 (56%)	231 (74.5%)	171 (64%)
Splicing	17 (6%)	25 (8%)	27 (10%)
Regulatory	0 (0%)	12 (4%)	3 (1%)
Small deletions	52 (19%)	24 (8%)	28 (10%)
Small insertions	23 (8%)	13 (4%)	15 (6%)
Small indels	2 (0.8%)	3 (1%)	4 (1.5%)
Gross deletions	24 (9%)	2 (0.5%)	14 (5%)
Gross insertions/ duplications	1 (0.4%)	0 (0%)	4 (1.5%)
Complex rearrangements	2 (0.8%)	0 (0%)	2 (1%)
Total	276	310	268

According to HGMD database (The Human Gene Mutation Database <http://www.hgmd.cf.ac.uk/ac/search.php>), accessed on 14th December 2015.

Difetti eredofamiliari di Antitrombina, Proteina C e Proteina S

- Ereditarietà: autosomica dominante
- Livelli attesi: eterozigote \Rightarrow 50-60%
omozigote \Rightarrow 0-15%
o doppio eterozigote
- Tipo I \Rightarrow ridotto dosaggio funzionale e immunologico
- Tipo II \Rightarrow ridotto dosaggio funzionale
normale dosaggio immunologico

Classificazione dei difetti eredofamiliari di Antitrombina

Tipo	Alterazione
I	Ridotta sintesi e/o aumentato turnover
II PE	Alterata funzionalità sito reattivo e sito di legame per eparina (Pleiotropic Effect)
II RS	Alterata funzione del solo sito reattivo (Reactive site)
II HBS	Alterata funzione del solo sito di legame per eparina (Heparin Binding Site)

Dosaggio dell'Antitrombina

Metodi funzionali

Test cromogenici che possono essere eseguiti:

- in presenza di eparina = attività di cofattore dell'eparina
- in assenza di eparina = attività antitrombinica progressiva
- con trombina (umana o bovina) o FXa

Metodi immunologici

- immunodiffusione radiale (RID), laser nefelometria, elettroimmunodiffusione (sec. Laurell)

Test di laboratorio e difetti eredofamiliari di Antitrombina

Tipo	Dosaggio funzionale (con trombina o FXa)		Dosaggio immunologico
	Cofattore eparinico	Attività progressiva	
I	↓	↓	↓
II PE	↓	↓	↔
II RS	↓	↓	↔
II HBS	↓	↔	↔

AT levels in different kindreds (Demers et al, 1993, modified)

Pts	AT Def. type	AT anti-IIa UI/ml	AT anti-Xa UI/ml	AT:Ag UI/ml
1	I	0.64	0.45	0.42
2	I	0.70	0.53	0.50
3	I	0.55	0.40	0.36
4	II	0.65	0.57	0.70

Minore probabilità di falsi negativi (valori normali in soggetti portatori del difetto) con l'uso FXa

Influenza del Cofattore Eparinico II

- Il Cofattore Eparinico II inibisce la trombina umana, ha un debole effetto sulla trombina bovina e non inibisce il FXa
- Se il livello di Cofattore Eparinico II è particolarmente alto, l'uso di trombina umana nel test determina una sovrastima dei livelli di Antitrombina del 20-30%
- L'interferenza del Cofattore Eparinico II si può quindi evitare usando FXa e in parte usando trombina bovina

Antitrombina Cambridge II (Ala384Ser) (Perry et al, 1998)

- Difetto di tipo II RS (dosaggio immunologico normale)
- Dosaggio cromogenico con FXa e trombina umana normale
- Dosaggio cromogenico con trombina bovina moderatamente ridotto

FXa (n=105)	Trombina umana (n=25)	Trombina bovina (n=59)
83%	76%	71%

Antithrombin Cambridge II (A384S): an underestimated genetic risk factor for venous thrombosis

Javier Corral,¹ David Hernandez-Espinosa,¹ Jose Manuel Soria,² Rocio Gonzalez-Conejero,¹ Adriana Ordonez,¹ Jose Ramon Gonzalez-Porras,³ Elena Perez-Ceballos,¹ Ramon Lecumberri,⁴ Ignacio Sanchez,¹ Vanessa Roldan,¹ Jose Mateo,² Antonia Minano,¹ Marcos Gonzalez,³ Ignacio Alberca,³ Jordi Fontcuberta,² and Vicente Vicente¹

Table 1. Characteristics of patients with venous thrombosis and control subjects

Characteristic	Patients	Controls	Multivariate analysis	
			P	OR (95% CI)
AT 1246 (A384S), no.				
G/G	1001 1.7%	1016 0.2%	—	—
G/T	17	2	.002	9.75 (2.2-42.5)

Prevalenza altri difetti di AT nei soggetti con precedente tromboembolismo venoso = 0.6%

4/17 pazienti = forma omozigote

Antithrombin Cambridge II (A384S) supports a role for antithrombin deficiency in arterial thrombosis

Vanessa Roldán¹; Adriana Ordoñez¹; Francisco Marín²; Esther Zorio³; José M. Soria⁴; Antonia Miñano¹; Francisco España³; Rocio González-Conejero¹; Javier Pineda⁵; Amparo Estellés³; Jordi Fontcuberta⁴; Vicente Vicente¹; Javier Corral¹

Thromb Haemost 2009; 101: 483–486

	All Patients (n= 1224)	Controls (n= 1649)	Crude OR (95%CI) p value	Adjusted OR (95%CI) p value
Age	51 ± 13	47 ± 16	p<0.001	
AT A384S	1214 (99.2)	1644 (99.7)	2.71	5.66
G/G (%)	10 (0.8)	5 (0.3)	(0.85–9.10)	(1.53–20.88)
G/T (%)			0.059	0.009
		Young Patients (n= 533)	Crude OR (95%CI) p value	Adjusted OR (95%CI) p value
Age		40 ± 5	p<0.001	
AT A384S		528 (99.1)	3.11	9.98
G/G (%)		5 (0.9)	(0.78–12.41)	(1.60–62.24)
G/T (%)			0.059	0.009

The phenotypic and genetic assessment of antithrombin deficiency

P. C. COOPER*, F. COATH[†], M. E. DALY[†], M. MAKRIS*,[†]

Int. Jnl. Lab. Hem. 2011, **33**, 227–237

AT defect	AT level: bovine thrombin-based assay (IU/dl)	AT level: bovine FXa-based assay (IU/dl)	AT antigen level (IU/dl)
Cambridge II	67	93	87
Denver	63	90	88
Reference range	86–132	83–135	83–124

Discrepancy between antithrombin activity methods revealed in Antithrombin Stockholm: do factor Xa-based methods overestimate antithrombin activity in some patients?

BLOOD, 15 MARCH 2002 • VOLUME 99, NUMBER 6

Table 1. Antithrombin levels with antigen and activity methods

	Antigen (Lia test), %	Thrombin-based activity test, %	Factor Xa-based activity test, %
Proband (our results)	93	60-64	110
Proband (Canadian results)	NA	72-74	53-56
Sister (our results)	86	62	92
Brother (our results)	92	55	89

Great discrepancy in antithrombin activity measured using five commercially available functional assays

Kaija Javela ^{*}, Sari Engelbarth, Leena Hiltunen, Pirjo Mustonen, Marja Puurunen

Thrombosis Research 132 (2013) 132–137

These patients belonged to 94 different families.
100 patients were type I and 104 patients type II.

Conclusions: There was great inter-assay variability especially in type II deficient patients, but also in patients with type I deficiency. However, most of the patients defined as having normal AT activity by some methods had thrombotic symptoms. Most tested assays find type I AT deficient patients accurately. In our study population only methods A1 and C could find most patients with type II AT deficiency, whereas methods A2, B and D misdiagnosed the majority of patients as non-deficient.

	Method	Principle	Incubation time (s)
A1	STA-Stachrom AT III®	Thrombin-based	60 [*]
A2	STA-Stachrom AT III®	Thrombin-based	300
B	Berichrom Antithrombin®	Thrombin-based	180 [*]
C	Innovance Antithrombin®	FXa-based	180 [*]
D	HemosIL®	FXa-based	120 [*]

Antithrombin heparin binding site deficiency: A challenging diagnosis of a not so benign thrombophilia

Christelle Orlando ^{a,*}, Olivier Heylen ^a, Willy Lissens ^b, Kristin Jochmans ^a

Thrombosis Research 135 (2015) 1179–1185

Objectives and Methods: The study population consisted of 82 genetically confirmed HBS deficient patients sharing six different mutations. Plasma samples of 35 of them, including one homozygous patient, were used for the evaluation of 4 commercial activity assays in their ability to diagnose HBS deficiency. We assessed mutation-specific prevalence of venous and arterial thrombosis and the contribution of additional thrombophilic risk factors.

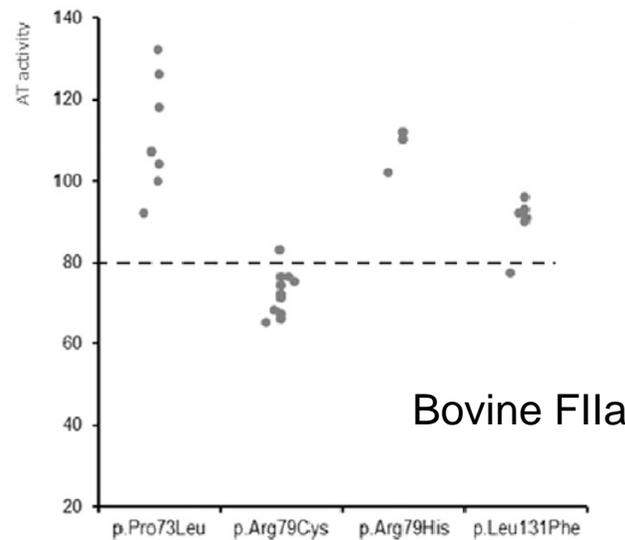
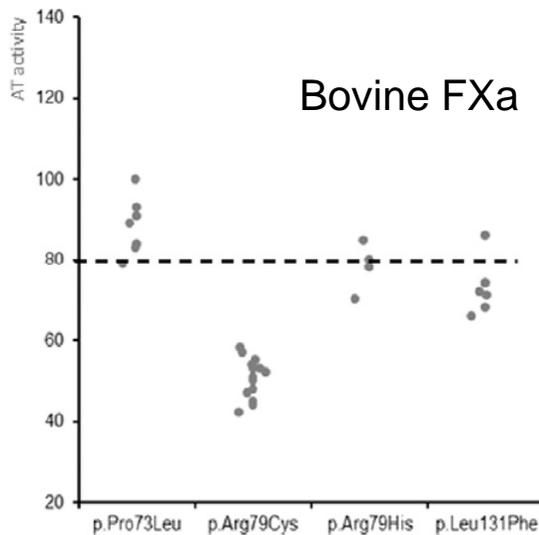
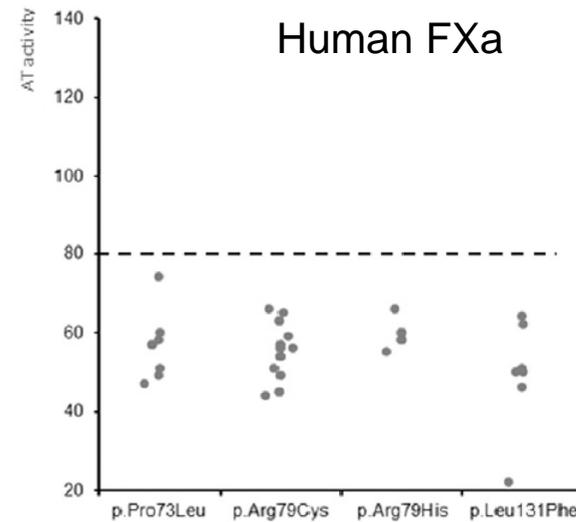
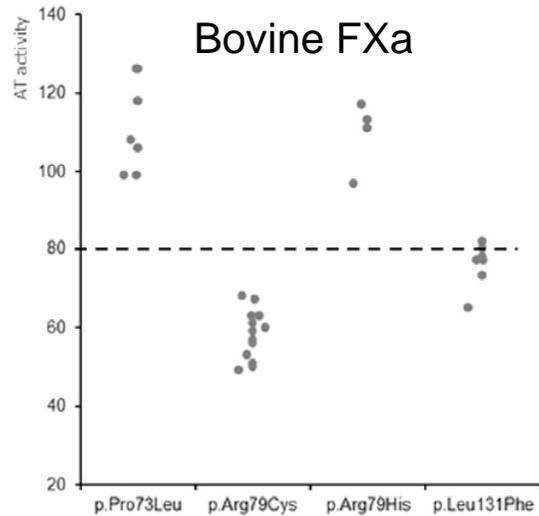
	LIQ	COA	INN	BIO
Substrate source	Bovine FXa	Bovine FXa	Human FXa	Bovine FIIa
Incubation time	100-140 s	100-140 s	180-190 s	55-65 s
Heparin concentration	3000U/mL	3000U/mL	1500U/mL	5000U/mL

LIQ = HemosIL®Liquid AT, COA = Coamatic® AT, INN = Innovance®, BIO = Biophen® AT.

Antithrombin heparin binding site deficiency: A challenging diagnosis of a not so benign thrombophilia

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The phenotypic and genetic assessment of antithrombin deficiency

P. C. COOPER*, F. COATH†, M. E. DALY†, M. MAKRIS*,†

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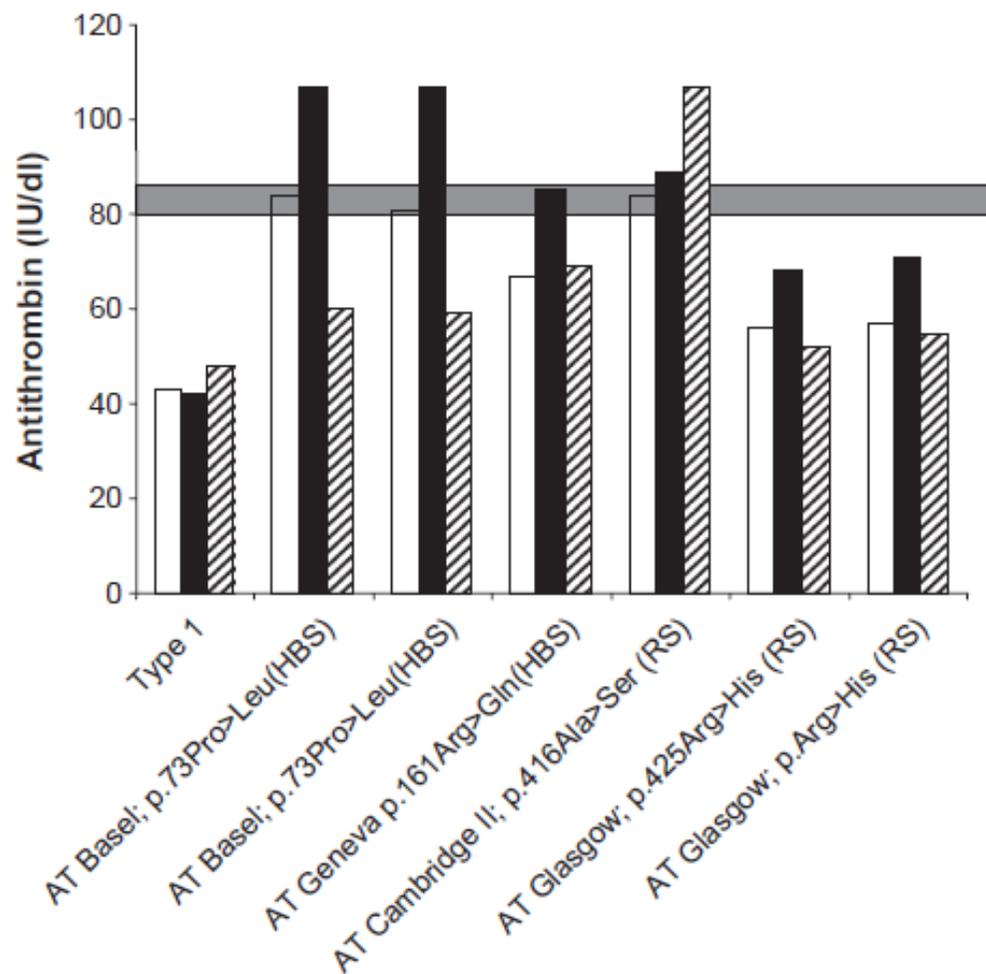


Figure 4. Antithrombin (AT) activity measured using a thrombin-based assay with 30-s incubation of sample dilution with bovine thrombin (□) or with a 180-s incubation with bovine thrombin (■) and assayed using a new human FXa-based assay (Innovance, Siemens) (▨). The broad horizontal bar indicates the lower limit of the normal range for AT in these assays.

This suggests that not only the source of enzyme (FIIa or FXa) is responsible for the difference in performance of AT functional tests. Table 1 demonstrates the assay characteristics of the most commonly used hc-anti-FXa AT assays. All but one reagent contains bovine FXa as enzyme in the reaction. The different tests differ in plasma dilution, in the added heparin concentration, in the source of chromogenic substrate and also in the incubation time. It can be concluded that clarification of the situation concerning heparin cofactor AT functional assays warrants future research to establish improved recommendation for AT testing.



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Editorial

Puzzling questions on antithrombin: Diagnostic limitations and real incidence in venous and arterial thrombosis



In conclusion, despite of being an old molecule, there are still many interesting issues concerning antithrombin. We must improve the methods to diagnose antithrombin deficiency. And we should know the real incidence of antithrombin defects in atherothrombotic disorders.

Principali domain funzionali della Proteina C

- Sito di attivazione
- Sito attivo
- Siti che legano i substrati (FVIII e FV)
- Sito di legame con Proteina S
- Siti di interazione con superfici fosfolipidiche, Ca^{2+} , EPCR

Dosaggio della Proteina C

Metodi funzionali

Attivazione della PC con Protac e diluizione con plasma carente o tampone

- Metodi coagulativi
- Metodi cromogenici

Metodi immunologici

- ELISA
- Elettroimmunodiffusione (sec. Laurell)

Dosaggio della Proteina C: metodi funzionali coagulativi

- Sensibili a tutte le possibili alterazioni della PC (interazione PC/Ca²⁺, fosfolipidi, PS, FV e FVIII)
- Considerati a lungo i metodi di prima scelta, in quanto identificano tutti i possibili difetti
- Numerose interferenze

Dosaggio della Proteina C: metodi funzionali cromogenici

- Evidenziano solo alterazioni del sito reattivo e del sito coinvolto nell'attivazione con Protac
- Nessuna interferenza

Dosaggio della Proteina C: metodi funzionali coagulativi vs cromogenici

Metodi coagulativi

- Falsi valori ridotti per aumentati livelli FVIII, mutazione FV Leiden
- Falsi valori normali in presenza di LAC e eparina#

Metodi cromogenici

- Falsi valori normali in un raro difetto di tipo II (indicato come IIb; prevalenza 5% di tutti i difetti di tipo II) caratterizzato da una alterazione del sito di interazione con Ca²⁺ e fosfolipidi

disponibili metodi insensibili fino a 1-2 U/ml

Classificazione dei difetti eredofamiliari di Proteina C

Tipo	Dosaggio funzionale		Dosaggio immunologico
	coagulativo	cromogenico	
I	↓	↓	↓
II a	↓	↓	↔
II b	↓	↔	↔

Altri fattori Vit-K dipendenti nella norma

Dosaggio della Proteina C: scelta del metodo funzionale

- Le possibili interferenze sul test coagulativo sono numerose e si possono presentare una certa frequenza
- Il difetto di tipo IIb (alterato test coagulativo e normale test cromogenico) è molto raro
- L'uso di un metodo cromogenico è raccomandabile (pochi soggetti saranno non diagnosticati se si adotta questo metodo)
- L'uso di un metodo coagulativo è comunque indicato in casi selezionati in cui si sospetta la presenza di un difetto di tipo IIb

Diagnostica dei difetti eredofamiliari di Proteina S

Complicata da:

- Presenza nel plasma di 2 forme di PS:
libera (40%) = attiva come cofattore della PC
legata al C4bBP (60%) = inattiva
- C4bBP è una proteina della fase acuta
aumento del C4bBP = aumento della forma legata e
riduzione della forma libera
- Nel plasma possono essere misurate attività,
proteina totale e forma libera

Dosaggio della Proteina S

Metodi funzionali

Per il dosaggio dell'attività

- Coagulativi (PT, aPTT)

Metodi immunologici

Per il dosaggio della proteina S totale e della frazione libera

- ELISA, nefelometrici, elettroimmunodiffusione (sec. Laurell)

Classificazione dei difetti eredofamiliari di Proteina S

Tipo	Dosaggio funzionale	Dosaggio immunologico	
		Libera	Totale
I	↓	↓	↓
II	↓	↔	↔
III	↓	↓	↔
Altri fattori Vit-K dipendenti nella norma			

Dosaggio funzionale della Proteina S

Metodi coagulativi sensibili a:

- Mutazione FV Leiden
- Presenza LAC e aumento di fattore VIII (metodi basati su PTT)
- Presenza di fattore VIIa (metodi basati su PT)

Dosaggio della Proteina S: test di primo livello

- Raccomandabile utilizzare come test di primo livello un metodo immunologico per il dosaggio della frazione libera in sostituzione del metodo funzionale coagulativo
- Il dosaggio della frazione libera correla bene con il dosaggio funzionale nei difetti di tipo I e III, ma non in quelli di tipo II

TEST OF THE MONTH

AJH Educational Material

Am. J. Hematol. 89:1147–1150, 2014.

Activated protein C resistance testing for factor V Leiden

■ Practical Considerations for Laboratories and Clinicians

The presence of FV_{Leiden} can also be determined by DNA testing, which is not affected by any of the interferences described above.

However, APC-R assays are advantageous as they are easily automated, cost effective [37,38], and may detect rare causes of APC resistance other than FV_{Leiden} . They are also necessary for detecting pseudohomozygous FV_{Leiden} [11] and for assessing phenotypic FV_{Leiden} thrombophilia in bone marrow or liver transplant patients [39].

APC-R assays that dilute patient plasma into factor V-deficient plasma (or the relevant equivalent for the assay) are much more accurate for detecting FV_{Leiden} than are assays that do not. Although concordance of APC-R assays with DNA analysis is extremely high when the APC-R assay includes a dilution step, it is reasonable for laboratories to perform DNA analysis on patients with abnormal APC-R results, as this practice helps confirm that there has been no specimen mix-up during testing [39]. With most APC-R assays, lupus anticoagulants can cause falsely normal results (Table I). Therefore, if DNA analysis is not available, any patient with an abnormal result should be tested for a lupus anticoagulant.

SPECIAL ARTICLE

International consensus statement on an update of the classification criteria for definite antiphospholipid syndrome (APS)

S. MIYAKIS,* M. D. LOCKSHIN,† T. ATSUMI,‡ D. W. BRANCH,§ R. L. BREY,¶ R. CERVERA,**
R. H. W. M. DERKSEN,†† P. G. DE GROOT,†† T. KOIKE,‡ P. L. MERONI,‡‡ G. REBER,§§
Y. SHOENFELD,¶¶ A. TINCANI,*** P. G. VLACHOYIANNOPOULOS††† and S. A. KRILIS*

**APS diagnosis is possible only if one clinical
and one laboratory criterion are present**

Laboratory criteria

- Lupus Anticoagulant (LA) present in plasma, detected according to the guidelines of the ISTH (SSC on LA/phospholipid dependent antibodies)
- Anti phospholipid antibodies
 - Anticardiolipin antibodies (IgG and/or IgM) and
 - Anti β 2 glycoprotein I antibodies (IgG and/or IgM) in serum or plasma, present in medium or high titer (> 40 U or $>$ the 99th percentile), measured by a standardized ELISA

Test positivity must be confirmed on two or more occasions at least 12 weeks apart

- LA and solid-phase aPL (Ab aCL and Ab anti β 2GPI) coexist in a limited number of patients with the APS syndrome
- Diagnosis must be based on both LA and solid-phase aPL detection

Arthritis & Rheumatism

An Official Journal of the American College of Rheumatology
www.arthritisrheum.org and wileyonlinelibrary.com

SPECIAL ARTICLE

International Consensus Guidelines on Anticardiolipin and Anti- β_2 -Glycoprotein I Testing

Report From the 13th International Congress on Antiphospholipid Antibodies

Gabriella Lakos,¹ Emmanuel J. Favaloro,² E. Nigel Harris,³ Pier Luigi Meroni,⁴
Angela Tincani,⁵ Richard C. Wong,⁶ and Silvia S. Pierangeli⁷

Vol. 64, No. 1, January 2012, pp 1–10

OFFICIAL COMMUNICATION OF THE SSC

Update of the guidelines for lupus anticoagulant detection

V. PENGO,* A. TRIPODI,† G. REBER,‡ J. H. RAND,§ T. L. ORTEL,¶ M. GALLI** and P. G. DE GROOT††

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LA laboratory detection

- Patients selection
- Blood collection
- Choice of tests
- Diagnostic steps
- Transmission results
- When testing

Diagnostic steps

1. Screening tests
2. Mixing tests
3. Confirmatory tests

Screening tests (aPTT, DRVVT)

Thrombin time
to exclude UH

Altered

Mixing tests

Normal

Altered

Factor deficiency

Confirmatory tests

Altered

Normal

Factor inhibitor

LA diagnosis

**LA
diagnostic
steps**



Laboratory methods to detect antiphospholipid antibodies

Steven A. Krilis^{1,3} and Bill Giannakopoulos^{1,2,3}

Hematology 2014

Summary of commonalities and contrasts between recent ISTH, BCSH, and CLSI guidelines for LAC detection

Area of recommendation	ISTH 2009	CLSI 2014
Sample preparation	Double centrifugation	Double centrifugation
Assays to use	dRVVT and aPTT	dRVVT and aPTT and/or others
Testing order	Screen-mix-confirm	Screen-confirm-mix
Ratio derivation	NPP denominator	RI mean denominator
RI/cutoffs	99th percentile	97.5th percentile (if Gaussian)
Calculations for phospholipid dependence	Percent correction of screen by confirm or LAC ratio (screen/confirm)	Percent correction of screen by confirm, or LAC ratio (screen/confirm)
Mixing test	Perform on 1:1 mixture with NPP; interpret with ICA or mixing test-specific cutoff	Perform on 1:1 mixture with NPP; interpret with ICA or mixing test-specific cutoff
Testing patients on VKAs	Undiluted plasma if INR < 1.5; mix with NPP if INR > 1.5 but < 3.0	Screen and confirm on 1:1 mixture with NPP; TSVT and ET or PNP
Testing patients on UFH	Interpret with caution	Can detect LAC in some cases where heparin neutralizer is effective
Interpretive reporting	Recommended	Recommended

ET indicates Ecarin time; ICA, index of circulating anticoagulant; PNP, platelet neutralization procedure; RI, reference interval; TSVT, Taipan snake venom time; and UFH, unfractionated heparin.

To Mix or Not to Mix in Lupus Anticoagulant Testing? That is the Question

Armando Tripodi, Ph.D.¹

Semin Thromb Hemost 2012;38:385–389.

ISTH guidelines on Lupus Anticoagulant testing

V. Pengo

Thrombosis Research 130 (2012) S76–S77

Diagnostic steps are:

a) screening step; b) mixing test; c) confirmatory test.

All these steps are needed and integrated tests where screening plasma is tested in the presence of low and high phospholipid concentration without a mixing test should be discouraged.

More on: laboratory investigation of lupus anticoagulants: mixing studies are sometimes required

A. TRIPODI* and V. PENGO†

- Reporting a negative LA without having performed mixing tests may be dangerous
- The LA activity of some positive plasma samples is fully unveiled only when normal plasma is added, because of the so-called “lupus cofactor” phenomenon
- This unknown cofactor (prothrombin? β 2-glycoprotein I?) which potentiates the inhibitory activity of LA is responsible for the paradoxical prolongation of the clotting time observed in the mixing test

J Thromb Haemost 2011; 9: 2126–7.

Optimisation of lupus anticoagulant tests: should test samples always be mixed with normal plasma?

Maarten T. T. Pennings¹; Philip G. de Groot¹; Joost C. M. Meijers^{2,3}; Albert Huisman¹; Ronald H. W. M. Derksen⁴; Rolf T. Urbanus¹

Thromb Haemost 2014; 112: 736–742

We conclude that mixing studies do not have an additional value in the exclusion of a single coagulation factor deficiency as the cause of a false positive screening assay. As a consequence, the merits of mixing during LA-assessment are questionable. Moreover, mixing tests might induce false negative LA results due to dilution effects. Since LA is readily detected with the combination of screening test and phospholipid confirm, we suggest to omit the mixing test during routine LA-assessment.