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The Coagulation Profile Monitoring of COVID-19 Patients with Standard and Viscoelastic Point of Care Hemostasis Tests

Lazarevic M^{1,2*}, Vukovic N³, Milic D^{1,2}, Kamenov A^{1,2}, Peric V^{1,2}, Golubovic M^{1,2}, Stosic M^{1,2}, Spasic D², Stojiljkovic V^{1,2} and Stokanovic D¹

¹Medical School of Nis, University of Nis, Serbia

²Clinic of Cardiovascular Surgery, University Clinical Center Nis, Serbia

³Clinic for Anesthesiology and Intensive Therapy, University Clinical Center Nis, Serbia

*Corresponding author:

Milan Lazarevic,

Clinic of Cardiovascular Surgery, University Clinical Center Nis, Medical School of Nis, University of Nis, Serbia Received: 15 Apr 2023 Accepted: 20 May 2023 Published: 30 May 2023 J Short Name: JCMI

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1. Abstract

1.1. Introduction: Coagulation disorders during COVID-19 infection are associated with poor prognosis and disease severity, because two processes that interfere each other are thrombosis and inflammation. Very important issue for clinicians is timely and adequate hemostasis and inflammation monitoring in order to prevent and treat potentially letal consequences.

1.2. Subjects and Methods: The study was approved by the Ethics Committee of the Clinical center Nis, Serbia. One hundered fourty two patients presented with COVID-19 ARDS were admitted to the ICU in Clinic for anesthesiology Clinical Center Nis, from 14th April 2020 to 25 th May 2020. On admission blood was collected for biochemical and coagulation testing. The data obtained was analyzed using Statistical Package for Social Sciences (SPSS v. 25, Chicago, IL, USA).

1.3. Results: Among all parameters assessed, mortality was associated with higher age (p<0.05), higher factor I (p<0.05), INR (p<0.001), D-Dimer (p<0.001), ADP (p<0.001), ASPI (p<0.001), TRAP (p<0.001), PSEP (p<0.001), A5extest (p<0.01), A10ex test (p<0.01), A5 fib (p<0.001), A10fib (p<0.001) and MCF fib (p<0.001), but lower CT extest (p<0.05). Mortality was associated with extreme values of D-Dimer above 1000 (p<0.001), ADP above 590 (p<0.001), ASPI above 800 (p<0.001), TRAP above 1500 (p<0.001) and PSEP above 1000 (p<0.05).

1.4. Discussion: In our study three variables resulted with extraordinary discriminating capacity with AUC>0.9, those calculated cut-of values of D-dimer, thrombin activating peptide receptor (TRAP) test and A10 in FIBTEM. Predictive ability of D-dimer grows over the time. In our study, A10 FIBTEM stood out as the best predictive marker for mortality outcome among the data obtained by thromboelastometry. Not only does it correlate with the MCF value of the same test, A10 FIBTEM has proven to be more useful in predicting different clinical outcomes. It belongs to the group of markers that depict the strength of the clot and showed the strongest predictive potential among other FIBTEM elements that are generally elevated in hypercoagulable states such as CO-VID19 infection.

1.5. Conclusion: The key to success in the treatment of Covid 19 infection is timely and adequate therapy and patient monitoring, which is impossible without early risk stratification and mortality prediction. Sophisticated hemostasis parameters can contribute to early risk assessment, which was initially performed only on the basis of the patient's clinical status.

2. Introduction

The pandemic of COVID-19 was challenge for healthcare systems around the world, with a development of acute respiratory distress syndrome (ARDS) and the need for admission to an intensive care unit (ICU) or death. A lot of different symptoms are present but the most important are severe lung dysfunction, a need for ventilation, shock and multiple organ failure [1].

Coagulation disorders during COVID-19 infection are associated with poor prognosis and disease severity, because two processes that interfere each other are thrombosis and inflammation [2]. Due to viral infection, pathogens initiate complex systemic inflammatory responses as part of innate immunity. Activation of host immune systems results in activation of coagulation and thrombin generation and that process is called immunothrombosis [3].

Inflammation is present in patients with SARS-CoV-2 infection, levels of IL-6 are elevated, C reactive protein and procalcitonin, and also fibrinogen [4]. Endothelial cell activation and damage results in disruption of the natural antithrombotic state [5]. This inflammation and activation of coagulation is the cause for the elevated D-dimer levels, as increased levels have been associated with thromboembolism[6]. Some patients have systemic inflammatory response syndrome (SIRS) or cytokine storm, which may explain more dramatic changes in coagulation tests, including significantly elevated D-dimer and changes in other hemostasis tests, especially as the disease progresses[7,8].

The receptor for virus to adhere is an angiotensin-converting enzyme 2 receptor on endothelial cells and viral replication causes inflammatory cell infiltration, endothelial cells death, and microvascular thrombosis [9]. As a result, microcirculatory dysfunction contributes to the clinical symptoms in patients with COVID-19.

Very important issue for clinicians is timely and adequate hemostasis and inflammation monitoring in order to prevent and treat potentially letal consequences.

3. Subjects and Methods

The study was approved by the Ethics Committee of the Clinical center Nis, Serbia. One hundered fourty two patients presented with COVID-19 ARDS were admitted to the ICU in Clinic for anesthesiology Clinical Center Nis, from 14th April 2020 to 25 th May 2020. All patients, aged from 36 to 84 years, female and male, were under tracheal intubation and mechanical ventilation and they were all in rolled in study. On admission blood was collected for biochemical and coagulation testing.

Blood samples were taken from the antecubital vein and stored in serum vacutainer tubes without additives for c reactive protein (CRP), using the immunoturbidimetry method on a Beckman Coulter AU 680 analyzer (Beckman Coulter Inc., Brea, CA, USA). Presepsin (pg/mL) levels were measured from the whole blood specimens using chemiluminescence enzyme immunoassay technology and Magtration® technology on a PATHFAST Immunoanalyzer (Mitsubishi Chemical Europe GmbH, Düsseldorf, Germany).

For coagulation profile samples testing (D-dimer, prothrombin time, partial thromboplastin time, fibrinogen concentration and anti-Xa) we used 4ml whole blood citrated tubes, and tests were performed on ACL TOP 350 coagulometer (Instrumentation Laboclinandmedimages.com ratory, USA).

Viscoelastic test (Clot Pro, Enicor, Germany) was also performed from 4ml citrated whole blood. The output of the instrument consists, among all of 1. coagulation time (CT, seconds) 2. clot amplitude after 5 and 10 minuts(A5, A10) 3.maximum clot firmness (MCF, mm) 4. clot formation time (CFT, seconds).

For impedance aggregometry-platelet function testing (Multiplate, Roch, Germany) we took blood in litium-heparinised 4ml tubes, and we used different platelet agonists in three separate tests to measure platelet aggregation 1. adenosine diphosphate in ADP test (aggregation units per minute-AU/min), 2.arachidonic acid in ASPI test (aggregation units per minute-AU/min) and 3.thrombin in TRAP test (aggregation units per minute-AU/min).

For whole blood count measure we used Horiba ABX 200 (Horiba Medical, France) counter and blood was drown in 4ml tubes with ethylenediaminetetraacetic acid (EDTA).

4. Statistical Analyses

The data obtained was analyzed using Statistical Package for Social Sciences (SPSS v. 25, Chicago, IL, USA). According to the normality of distribution, continuous variables were presented as means with standard deviation, or as median with interquartile range. Categorical variables were presented as absolute and relative numbers. The differences between two tested groups were tested by parametric Student's t-test, or non-parametric Mann-Whitney U-test and Fischer's exact test. The correlation between continuous variables was assessed according to Pearson's correlation coefficient. Univariate and multivariate binary logistic regression was performed to determine statistically significant predictors of dependent variables. We evaluated the discriminatory power of various laboratory parameters and determined the optimal cut-off values by the receiver operating characteristic (ROC) curve analyses. ROC curves for multiple variables were constructed based on probabilities obtained by binary logistic regression modeling and compared with DeLong test using MedCalc (v. 19.0; MedCalc Software Ltd, Ostend, Belgium). A p-value less than 0.05 was a measure of statistical significance.

5. Results

Among all parameters assessed, mortality was associated with higher age (p<0.05), higher factor I (p<0.05), INR (p<0.001), D-Dimer (p<0.001), ADP (p<0.001), ASPI (p<0.001), TRAP (p<0.001), PSEP (p<0.001), A5extest (p<0.01), A10ex test (p<0.01), A5 fib (p<0.001), A10fib (p<0.001) and MCF fib (p<0.001), but lower CT extest (p<0.05) (Table 1).

When we analyzed all these parameters in the view of their normal ranges, mortality was associated with normal range factor I, rather than values above it (p<0.05). Values below normal range were associated with survival in cases of ADP (p<0.001), ASPI (p<0.001) and TRAP (p<0.001). On the contrary, death occurrence was more frequent in patients with D-Dimer (0<0.05), PSEP (p<0.001), A10

fib (p<0.001) and MCFfib above normal range (p<0.01). Good clinical outcome, in terms of survival, was associated with higher ranges (normal or above normal) of CT extest (p<0.05), but lower ranges (normal or below normal) of A10 extest (p<0.05). As for CFT extest, death outcome was the most common in patients in normal range values, while the survivers had this parameter above or below normal range (p<0.05).

Mortality was associated with extreme values of D-Dimer above 1000 (p<0.001), ADP above 590 (p<0.001), ASPI above 800 (p<0.001), TRAP above 1500 (p<0.001) and PSEP above 1000 (p<0.05).

Using ROC analysis, we identified optimal cut-off values for a number of inflammation and coagulation parameters, with the highest sensitivity and specificity in discriminating patients with later exitus (Graph 1.). Good discriminatory ability (AUC>0.7) was shown for the following parameters: Factor I (\geq 9.14), INR (\geq 1.38), PSEP (\geq 335), A5 fib (\geq 28) and MCF fib (\geq 36). Cut-off values for ADP (\geq 591) and ASPI (\geq 728) were excellent in discriminating patients with exitus. The best discriminators, with AUC>0.9, were D-Dimer (\geq 860), TRAP (\geq 1180) and A10 fib (\geq 30), all significantly better than the other parameters, but without statistically significant difference among them. After making combinations of 2 or 3 of these last parameters, we have found that both D-Dimer and TRAP were worse than the combination of other 2 parameters (p<0.05), but not A10 fib which was equally good as all the 2 or 3-parameter combinations tested (Table 2).

We have found various degree of correlation between analyzed variables (Table 3). There was a strong positive correlation among ADP, ASPI and TRAP (p<0.001), as well as among A5extest, A10 extest and MCFextest (p<0.001), and between A5fib and MCF

fib (p<0.001). Strong negative correlation existed between CFT ex test and A5extest (p<0.001), A10 ex test (p<0.001) and MCF extest (p<0.001). Presepsin concentration was in weak positive correlation with A10fib (p<0.001). Factor I moderately directly correlatied with A5 fib (p<0.001) and MCFfib (p<0.001), besides, its weak positive correlation was noted with A5 extest (p<0.001), A10 extest (p<0.001), MCF extest (p<0.001) and CT fib (p<0.001).

After performing univariate binary logistic regression (Table 4), in order to find predictors of mortality, a number of parameters stood out. In cases where the same parameter has been found significant in various forms (countinous variable, according to normal range, or according to previously found cut-off value), the variable with the highest predictive value was chosen to be included in the multivariate model. The following variables cut-off values, previously determined, were more valuable than absolute values or standard cut-offs: age, Factor I, INR, DDimer, ASPI, PSEP, A5 extest, A10 extest and A5 fib.

Interestingly, TRAP and A10 fib cut-off values, previously shown to have high discriminatory ability, were less valuable in the logistic regression modelling of mortality. Due to the total patients number and high co-linearity between some variable, the number of predictors in the multivariate model had to be reduced. The most fitted multivariate model (χ 2=141.007, p<0.001) explains 63.0-91.6% in the death occurrence variance. All three variables included in the model were found to be independent predictors of mortality. DDimer above 860 increases the risk of death 24 times (p<0.01). TRAP values (binned according to normal value range), with each higher value bin, the risk is 22 times higher (p<0.01). A10fib values above normal range brings 290 times greater risk of death (p<0.05).

		No mortality (N=104)	Mortality (N=38)	t* or Z** or $\chi 2^{***}$ (p)		
Age (years)		62.67±12.10	66.71±8.44	2.229 (0.028)*		
Factor I		7.62±1.54	8.56±2.29	2.344 (0.023)*		
Factor I (>4)		103 (99.0%)	34 (89.5%)	4.944 (0.018)***		
AntiXa		0.38±0.14	0.37±0.15	0.323 (0.747)*		
INR		1.26±0.16	1.48±0.23	5.500 (0.000)*		
INR (>1)		102 (98.1%)	38 (100.0%)	0.003 (1.000)***		
DDimer		444.0 (407.0-737.5)	1285.0 (970.0-1542.0)	8.393 (0.000)**		
DDimer (>230)		89 (85.6%)	38 (100.0%)	4.697 (0.011)***		
DDimer (>10	00)	2 (1.9%)	27 (71.1%)	77.640 (0.000)***		
aPTT		37.53±8.92	36.38±7.14	0.712 (0.478)*		
	<34	45 (43.3%)	20 (52.6%)			
aPTT	34-38	14 (13.5%)	3 (7.9%)	1.348 (0.510)***		
	>38	45 (43.3%)	15 (39.5%)			
ADP		365.14±150.88	670.21±224.77	7.753 (0.000)*		

 Table 1: Inflammation and coagulation parameters according to mortality

-2023				
	<406	61 (58.7%)	3 (7.9%)	
ADP	406-992	43 (41.3%)	43 (41.3%) 30 (78.9%)	
	>992	0 (0.0%)	5 (13.2%)	
ADP (>590)	1	5 (4.8%)	24 (63.2%)	54.771 (0.000)***
ASPI		486.61±229.59	843.84±217.04	8.326 (0.000)*
ASPI (<790)		103 (99.0%)	15 (39.5%)	66.129 (0.000)***
ASPI (>800)		0 (0.0%)	22 (57.9%)	66.896 (0.000)***
TRAP		548.25±293.04	1375.90±367.61	13.884 (0.000)*
	<923	923 94 (90.4%) 7 (18.4 23-1509 10 (9.6%) 14 (36.5)	7 (18.4%)	
TRAP	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	14 (36.8%)	78.997 (0.000)***	
	>1509	0 (0.0%)	14 (36.8%) 78.997 (0.000)*** 17 (44.7%)	
TRAP (>150	0)	0 (0.0%)	20 (52.6%)	59.435 (0.000)***
PSEP		293.0 (239.0-395.5)	593.0 (446.2-743.8)	5.074 (0.000)**
PSEP (>337)		27 (26.0%)	33 (86.8%)	39.818 (0.000)***
PSEP (>1000))	0 (0.0%)	3 (7.9%)	5.005 (0.018)***
CTextest		65.38±16.38	57.58±15.77	2.535 (0.012)*
	<38	1 (1.0%)	3 (7.9%)	
CTextest	38-65	47 (45.2%) 23 (60.5%)		8.958 (0.011)***
	>65	56 (53.8%)	12 (31.6%)	
A5extest		53.0 (47.0-56.0)	56.5 (51.0-60.0)	2.628 (0.009)**
A5extest (<3	8)	7 (6.7%)	0 (0.0%)	1.446 (0.190)***
A10extest		59.5 (55.0-63.0)	63.5 (58.5-66.0)	3.174 (0.002)**
	<38	10 (9.6%)	0 (0.0%)	
A10extest	38-67	89 (85.6%)	33 (86.8%)	6.415 (0.040)***
	>67	5 (4.8%)	5 (13.2%)	•
MCFextest	1	61.0 (58.0-64.0)	63.0 (60.0-67.2)	2.405 (0.016)**
	<38	4 (3.8%)	0 (0.0%)	
MCFextest	38-68	89 (85.6%)	32 (84.2%)	2.099 (0.350)***
	>68	11 (10.6%)	6 (15.8%)	
CFTextest	1	53.0 (41.2-63.0)	50.5 (44.0-53.2)	1.005 (0.315)**
	<38	20 (19.2%)	2 (5.3%)	
CFTextest	38-69	63 (60.6%)	31 (81.6%)	6.111 (0.047)***
	>69	21 (20.2%)	5 (13.2%)	
CTfib	1	67.5 (45.2-86.5)	63.0 (47.8-73.5)	0.332 (0.740)**
CTfib (>70)		46 (44.2%)	12 (31.6%)	1.357 (0.185)***
A5fib		22.62±6.43	27.45±6.58	3.939 (0.000)*
A5fib (>9)		103 (99.0%)	38 (100.0%)	0.000 (1.000)***
A10fib		17.48±4.67	36.26±4.46	21.472 (0.000)*
A10fib (>23)		13 (12.5%)	37 (97.4%)	84.189 (0.000)***
MCFfib		24.62±7.63	30.90±7.91	4.294 (0.000)*
MCFfib (>25)	48 (46.2%)	28 (73.7%)	7.409 (0.004)***

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	AUC (95%CI for AUC)	р	cut-off	sensitivity (%)	specificity (%)
Age	0.637 (0.536-737)	0.013	68	71.1	64.4
Factor I	0.702 (0.591-0813)	0	9.14	71.1	74
Anti-Xa	0.487 (0.378-0.596)	0.816	0.4	55.3	47.1
INR	0.790 (0.696-0.884)	0	1.38	76.3	75
D-Dimer	0.961 (0.930-0.991)	0	860	86.8	95.2
aPTT	0.474 (0.373-0.575)	0.637	31.2	73.7	37.5
ADP	0.878 (0.816-0.940)	0	591	63.2	95.2
ASPI	0.848 (0.772-0.924)	0	728	65.8	97.1
TRAP	0.955 (0.923-0.987)	0	1180	76.3	99
PSEP	0.779 (0.701-0.856)	0	335	89.5	74
CT extest	0.364 (0.261-0.468)	0.013	37	100	1
A5 extest	0.644 (0.544-0.744)	0.009	57	50	77.9
A10 extest	0.674 (0.574-0.774)	0.002	64	50	82.7
MCF extest	0.632 (0.533-0.730)	0.016	59	97.4	27.9
CFT extest	0.445 (0.349-0.540)	0.315	41	92.1	25
CT fib	0.482 (0.381-0.583)	0.74	43	97.4	22.1
A5 fib	0.704 (0.597-0.810)	0	28	68.4	78.8
A10 fib	0.989 (0.968-1.000)	0	30	97.4	100
MCF fib	0.703 (0.607-0.798)	0	36	42.1	94.2

Table 2: Optimal cut-off values for a number of inflammation and coagulation parameters, with the highest sensitivity and specificity in discriminating patients with increased mortality

Table 3: Correlation between various coagulation and inflammation parameters

	FI	AntiXa	INR	DD	aPTT	ADP	ASPI	TRAP	PSEP	СТе	A5e	A10e	MCFe	CFTe	CTf	A5f	A10f	MCFf
	-0.058	0.226	0.284	0.176	0.274	-0.01	-0.104	0.092	0.114	0.273	0.066	0.098	0.117	-0.029	0.39	0.208	0.338	0.152
Age	0.497	0.007	0.001	0.036	0.001	0.91	0.219	0.276	0.175	0.001	0.438	0.244	0.165	0.731	0	0.013	0	0.071
EI		0.046	0.011	0.261	0.296	0.078	0.156	0.181	0.073	-0.274	0.432	0.405	0.396	-0.177	-0.39	0.68	0.11	0.613
FI		0.584	0.899	0.002	0	0.354	0.064	0.031	0.391	0.001	0	0	0	0.035	0	0	0.195	0
AntiVo			0.042	-0.026	0.29	0.038	-0.043	-0.006	-0.033	0.062	0.006	0.011	-0.018	0.042	0.076	0.143	0.029	0.089
Ашла			0.622	0.757	0	0.653	0.612	0.94	0.696	0.467	0.942	0.897	0.834	0.621	0.366	0.089	0.731	0.29
INID				0.351	0.093	0.249	0.192	0.218	0.144	-0.158	0.184	0.145	0.09	-0.167	-0.107	0.305	0.5	0.334
INK				0	0.272	0.003	0.022	0.009	0.088	0.061	0.029	0.086	0.288	0.047	0.207	0	0	0
חח					0.146	0.487	0.513	0.555	0.196	0.037	0.341	0.362	0.319	-0.15	-0.007	0.357	0.572	0.371
DD					0.083	0	0	0	0.019	0.66	0	0	0	0.075	0.933	0	0	0
aPTT						-0.031	0.087	-0.012	-0.09	0.283	0.157	0.16	0.227	-0.09	0.069	0.421	-0.048	0.461
ai i i						0.711	0.305	0.883	0.288	0.001	0.062	0.057	0.006	0.286	0.411	0	0.572	0
ADP							0.838	0.768	0.175	-0.135	0.392	0.444	0.346	-0.291	-0.178	0.147	0.532	0.312
ADI							0	0	0.038	0.106-9	0	0	0	0	0.034	0.081	0	0
ASPI								0.794	0.159	-0.266	0.49	0.574	0.49	-0.359	-0.362	0.306	0.424	0.414
ASIT								0	0.058	0.001	0	0	0	0	0	0	0	0
ΤΡΑΡ									0.184	-0.247	0.426	0.422	0.44	-0.286	0.206	0.259	0.627	0.313
									0.029	0.003	0	0	0	0.001	0.014	0.002	0	0
PSEP										-0.08	-0.022	-0.001	-0.021	0.024	0.069	0.042	0.351	0.034
1.511										0.344	0.793	0.992	0.801	0.778	0.415	0.62	0	0.69
CTe											-0.399	-0.355	-0.277	0.315	0.722	-0.351	-0.127	-0.319
											0	0	0.001	0	0	0	0.131	0

45.						0.907	0.933	-0.77	-0.562	0.663	0.265	0.703
Abe						0	0	0	0	0	0.001	0
A 10a							0.883	-0.728	-0.558	0.624	0.269	0.691
Alle							0	0	0	0	0	0
MCEa								-0.772	-0.463	0.633	0.279	0.665
WICTE								0	0	0	0.001	0
CETa									0.484	-0.433	-0.213	-0.464
Crie									0	0	0.011	0
CTF										-0.429	0.089	-0.421
CII										0	0.294	0.142
A 5 F											0.294	0.917
ASI											0	0
A 10£												-0.329
AIUI												0

Table 4: Binary logistic regression of mortality predictors

	Univariate		Multivariate	
	OR (95%CI for OR)	n	OR (95% CI for OR)	n
Age (>68)	4 445 (1 981-9 970)	0		P
Factor I	1 392 (1 093-1 773)	0.007		
Factor I (>4)	0.083 (0.009-0.764)	0.007		
Factor I (>9.14)	7,000 (3,062-16,003)	0.028		
	<u> </u>	0		
	480.244 (41.870-3040.800)	0		_
INK (21.58)	9.007 (4.051-23.005)	0		
DDimer	1.006 (1.004-1.008)	0		
DDimer (>1000)	125.182 (26.168-598.831)	0		
DDimer (≥860)	130.680 (35.590-479.835)	0	23.735 (2.824-199.461)	0.004
ADP	1.011 (1.007-1.016)	0		
ADP (<normal range="">)</normal>	17.007 (5.085-56.873)	0		
ADP (>590)	33.943 (11.139-103.433)	0		
ASPI	1.011 (1.007-1.016)	0		
ASPI (<normal range="">)</normal>	157.933 (19.847-1256.730)	0		
ASPI (≥728)	64.744 (17.131-244.686)	0		
TRAP	1.006 (1.004-1.009)	0		
TRAP (<normal range="">)</normal>	23.421 (8.669-63.278)	0	21.983 (2.365-204.311)	0.001
TRAP (≥1180)	331.889 (40.372-2728.392)	0		
PSEP	1.003 (1.001-1.004)	0		
PSEP (>337)	18.822 (6.668-53.131)	0		
PSEP (≥335)	24.241 (7.870-74.663)	0		
CTextest	0.970 (0.947-0.994)	0.014		
CTextest (<normal range="">)</normal>	0.376 (0.186-0.763)	0.007		
A5extest	1.100 (1.032-1.173)	0.003		
A5extest (≥57)	3.522 (1.604-7.734)	0.002		
A10extest	1.121 (1.035-1.215)	0.005		
A10extest (<normal range="">)</normal>	3.985 (1.300-12.211)	0.016		
A10extest (≥64)	4.778 (2.117-10.782)	0		
MCFextest	1.107 (1.026-1.195)	0.009		

MCFextest (≥59)	14.307 (1.875-109.149)	0.01		
A5fib	1.128 (1.056-1.204)	0		
A5fib (≥28)	8.076 (3.521-18.525)	0		
A10fib	1.895 (1.410-2.546)	0		
A10fib (>23)	259.000 (32.697-2051.584)	0	289.509 (3.438-24378.575)	0.012
			· · · · · · · · · · · · · · · · · · ·	
MCFfib	1.111 (1.053-1.172)	0		
MCFfib MCFfib (>25)	1.111 (1.053-1.172) 3.267 (1.441-7.406)	0 0.005		



Graph 1: ROC analysis, sensitivity and specificity of inflammation and coagulation parameters

6. Discussion

Extensive activation of the coagulation cascade leading to disseminated intravascular coagulation and thrombosis and inevitable progression of Covid 19 and mortality. In the background of these two extremes are hypercoagulability and aberrant fibrinolysis when extreme values of PT, aPTT, platelet count, fibrinogen and fibrin can be expected and associated with COVID 19 mortality [10]. Our idea was to find some parameters that would be markers of the mentioned pathophysiological mechanisms and find predictive cut-off values for them that would enable sufficiently early detection and stratification of the most at-risk Covid-19 patients.

Endothelial injury and consequently tissue factor genesis, and inhibited fibrinolysis due to changes in concentrations of urokinase-type plasminogen activator and plasminogen activator inhibitor-1 are main pathophysiologic mechanisms of focal or disseminated intravascular coagulation [11]. Severe endothelial injury followed with vascular thrombosis and angiogenesis are three principal morphological findings in Covid-19 related acute respiratory distress syndrome [12].

Vascular injury reflects extensive D-dimer elevations [13]. Therefore, this marker is recognized by the International Society of clinandmedimages.com Thrombosis and Haemostasis (ISTH) as the most important among the data we receive from routine initial analysis in patient risk stratification [14].

Concordance (C) statistic with value of area under the ROC (Receiver Operating Characteristic) curve (AUC) is a gold standard for outcome prediction for different predictive models [15]. Generally accepted value of it for excellent predictive ability of some diagnostic test is 0.8 [16].

In our study three variables resulted with extraordinary discriminating capacity with AUC>0.9, those calculated cut-of values of D-dimer, thrombin activating peptide receptor (TRAP) test and A10 in FIBTEM. These parameters were the basis for creating predictive models for estimating mortality in the most severe Covid-19 patients. Due to the hypofibrinolytic profile in thromboelasatometry, there was concern about the predictive ability of the D-dimer [17]. Cut-off value of D-dimer, which is roughly four times larger than normal, reflects hypercoagulability over hypofibrinolysis in critical ill Covid-19 patients. Predictive ability of D-dimer grows over the time. This fact should be kept in mind when analyzing the results, since our research includes patients with already developed ARDS [17].

Although the characteristic hypercoagulable profile, decreased CFT and increased MCF, was thromboelastometrically confirmed, there are no data on the correlation between thromboelastometric parameters and clinical outcomes [18]. In our study, A10 FIBTEM stood out as the best predictive marker for mortality outcome among the data obtained by thromboelastometry. Not only does it correlate with the MCF value of the same test, A10 FIBTEM has proven to be more useful in predicting different clinical outcomes. It belongs to the group of markers that depict the strength of the clot and showed the strongest predictive potential among other FIBTEM elements that are generally elevated in hypercoagulable states such as COVID19 infection. The clinical advantage of this marker would be its rapid detection, both in relation to MCF and in relation to conventional laboratory tests. Given these advantages, A10 Fibtem could be a useful parameter when admitting patients to the ICU [19, 20].

An elevated level of TRAP has a great predictive potential for intrahospital mortality in COVID 19. This result can be very useful in daily clinical practice, considering that the TRAP test represents baseline platet aggregation and is independent of the influence of acetylsalicylic acid derivatives and P2Y12 inhibitors. High TRAP test levels we interpret as overactivated phenotype of platelets which may be associated with a hypercoagulable state, disease progression and mortality.

Presepsin is a biomarker that has been studied in relation to sepsis, a potentially life-threatening condition that occurs when the body's response to infection causes damage to its own tissues and organs. There is some research that suggests that presepsin levels may be elevated in individuals with severe COVID-19, particularly those who develop sepsis as a complication of the disease.

Bacteremic co-infection is a leading cause of ICU admission, mechanical ventilation and mortality in individuals with CO-VID-19, and studies have shown that patients with severe CO-VID-19 who develop sepsis have a higher risk of death compared to those who do not. Due to the similar immunological and pathophysiological background of sepsis and covid 19, and frequent bacterial co-infection, it was logical to test the predictive ability of presepsin. High values of it may be a useful tool in predicting which individuals with COVID-19 are at higher risk of developing sepsis and potentially dying from the disease [21]. However, the cut-off value of presepsin that we obtained is slightly above the upper limit and is significantly lower compared to other studies [22]. Relationship between presepsin and COVID-19 mortality is still being studied, and more research is needed to fully understand the extent of this connection. Other factors, such as age, underlying health conditions, and access to medical care, may also play a role in determining an individual risk of mortality from COVID-19 [23].

7. Conclusion

The key to success in the treatment of Covid 19 infection is timely and adequate therapy and patient monitoring, which is impossible without early risk stratification and mortality prediction. Sophisticated hemostasis parameters can contribute to early risk assessment, which was initially performed only on the basis of the patient's clinical picture. Hypercoagulability is the main hemocoagulation disorder in COVID 19 infection. Of all the tested pointof-care parameters, D-dimer, A10 FIBTEM and TRAP stood out and their combinations are characterized by an outstanding predictive potential for the detection of in-hospital mortality of COVID 19 infection. These parameters are easy to interpret, easily available, are elements of different hemostasis tests and are markers of different effects on hypercoagulability.

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