Aberration in monocyte

count as anticipator of adverse terminal events in COVID-19: case-control study

Aberración en el recuento de monocitos como anticipador de eventos terminales adversos en COVID-19:

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Abstract

Background: Aberrancy in monocytes account is customarily recognized in patients with COVID-19 strikingly in gravely affected patients, the exacting interrelation with detrimental consequences still in the gap. The study was carried out to found any significant monocyte level changes in COVID-19 and its relation to diagnostic usefulness and expectation of adverse outcomes. A sample of 100 patients was recruited into the analysis; all with diagnosis compatible with pneumonia underwent care inwards of Al-Hussein Hospital for isolation; 50 participators with COVID-19 as cases and 50 participators with influenza A elected control. The diagnosis was suggested by clinical and radiological attributes and verified by polymerase chain reaction. Monocytes total and laboratory analysis of inflammatory indices and organ vitality were achieved for all participants. The mean age was 64.4, including 56 males and 44 females. 66% of cases were monocytopenic, 34% were non-monocytopenic. 12% of control were monocytopenic while 88% were non-monocytopenic with significant difference (P value=0.001). Forty-five participants were critical while 55 were non-critical, 37 COVID-19, and eight controls were critical and admitted to the ICU. Fortyone participants died, 35 from the COVID-19 category and six from controls. 30 of COVID-19 who were critical in ICU were monocytopenic (all died), three were non-monocytic (only one died) and 4 had monocytosis (all died), 6 of control who were critical in ICU were monocytopenic (all died), 2 had normal count (all survived) and no one had monocytopenia (Odd ratio=1.200). Deviation in monocyte count from the norm is a valuable discriminator for the diagnosis of CO-VID-19 and a suitable anticipator across all spectrums for its adverse consequences.

Keywords: COVID-19, Monocytes, Influenza A, ICU

Resumen

Antecedentes: la aberrancia en los monocitos se reconoce habitualmente en pacientes con COVID-19 de manera sorprendente en pacientes gravemente afectados, la interrelación exigente con consecuencias perjudiciales aún está en la brecha. El estudio se llevó a cabo para encontrar cambios significativos en el nivel de monocitos en COVID-19 y su relación con la utilidad diagnóstica y la expectativa de resultados adversos. Se reclutó para el análisis una muestra de 100 pacientes, todos con diagnóstico compatible de neumonía atendidos en las salas del Hospital Al-Hussein para aislamiento; 50 participantes con COVID-19 como casos y 50 participantes con influenza A elegidos como control. El diagnóstico fue sugerido por atributos clínicos y radiológicos y verificado por reacción en cadena de la polimerasa. Se logró el análisis total de monocitos y de laboratorio de los índices inflamatorios junto con la vitalidad del órgano para la suma total de todos los participantes. La edad media fue de 64,4 años, incluidos 56 varones y 44 mujeres. El 66% de los casos fueron monocitopénicos, el 34% no monocitopénicos. El 12% de los controles eran monocitopénicos mientras que el 88% eran no monocitopénicos con una diferencia significativa (valor de p = 0,001). 45 participantes fueron críticos mientras que 55 no lo fueron, 37 COVID-19 y 8 controles fueron críticos y fueron admitidos en UCI. 41 de los participantes murieron, 35 de la categoría COVID-19 y 6 de los controles. 30 de los COVID-19 que fueron críticos en la UCI fueron monocitopénicos (todos murieron), 3 no monocitopénicos (solo 1 murió) y 4 tuvieron monocitosis (todos murieron), 6 de los controles que fueron críticos en la UCI fueron monocitopénicos (todos murieron), 2 tenían un recuento normal (todos sobrevivieron) y nadie tenía monocitopenia (Odd ratio = 1.200). La desviación en el recuento de monocitos de la norma es un discriminador valioso para el diagnóstico de COVID-19 y un anticipador adecuado en todos los espectros por sus consecuencias adversas.

Palabras clave: COVID-19, Monocitos, Influenza A, UCI

Introduction

COVID-19 pandemic is the biggest terrible fulmination to humankind in the new century that retrogresses life's standardness. COVID-19 caused by SARS-CoV-2 spread intercontinentally impulse world health organization (WHO) to certify it as pandemic¹. COVID-19 sustains a significant health, social, and economic strain with measurable morbidness and lethality^{2,3}. It is fundamentally acute unspecified pulmonary sickness but eventually incorporates a multitude of systems, its manifestations are intermingled with a diversity of respiratory disorders of varied causations endorsing a secure analysis for discrimination, particularly when the cases are overwhelmed⁴.

Despite the fact that the verificatory diagnostic investigation for COVID-19 is polymerase chain reaction, its employment is impeded by irrefutable shortcomings encompassing protracted overturn time with elevated false negative yields uniquely in poverty-stricken nations⁵. Instantaneous authentication of the diagnosis of COVID-19 with sooner discerning those who are in neediness for vigorous interventions is pivotal in COVID-19, so mandating conceptualization a parameter that is effortlessly approachable and promptly obtainable⁶. Monocytes and other innate immune cells execute an orchestrated duty in the pathogenesis of a multitude of viral infections, infiltrating regionally and spread out distantly with antagonistic viral defensive influences7. Monocytes are thought-about the front-line defender among other immune cells, and they are the cardinal originator of the inflammatory milieu in CO-VID-19; they serve bi-directionally as inflammatory instigators and tissue repairers thus contributing to the diversity of end results encompassing resolution⁸. There are several affirmations for monocyte dysregulation in predisposing to hyperinflammatory upset and thence instigating to build up the seriousness and fatality of COVID-199. Monocytes, among other defensive cells, carry out an influential participant in the advancement of varied complexities and grim residual jeopardy in COVID-19 substantially thrombotic perturbations, organ dysfunctions, and hyperinflammatory distress predisposing to the upraised incidence of deadly consequences of this potentially lethal infection¹⁰. Various hematological parameters submit to aberrancy in COVID-19 incorporating aberration in monocytes count and its interconnection with the aggressiveness of outcome that deserves methodical exploration¹¹. Plenty of studies theorize utilization of hematological indicators that seems to be linked to heightened seriousness and lethality of COVID-19¹².

We hypothesized that the remarkable alterations of monocytes inspected recurrently in COVID-19 are noteworthy and plausibly have impactful diagnostic and prognostic implications. We intend to devise an inventive, easily memorized, affordable, reputable, and simply practicable test with diagnostic and expectation magnitude for unfavorable implications of COVID-19.

Materials and methods

Type of study: A case-control study had been preceded from the beginning of first week of February 2020 to the end of the last week of April 2020.

Study population: The participators constituted patients looking for medical facilities in isolation departments of AL-Hussein teaching hospital in Al-Nasiriya city, the diagnosis was assured by any feasible details in the case sheets of patients, whole numbers of patients accounted for in the analysis was 100 patients, with 50 cases affirmed to be COVID-19 pneumonia and 50 candidate verified to be influenza A pneumonia elected as control.

Inclusion criteria: whatever patients who were exceedingly suspicious to harbor pneumonia from a radiological and clinical point of view and proved by positive polymerase chain reaction for COVID-19 or influenza A.

Exclusion criteria: 1-Whatever patient had receiving medications currently or remotely known to affect monocytes level; 2-Whatever patients are known to have blood dyscrasia or malignancy; 3-Whatever patient with a known background of autoimmune disorders.

Sampling size: Sampling size was acceptable, restrained by the approachability of patients and time scale of analysis, but sampling task for control, systematized strategy for random sampling was accomplished to construct the control.

Ethical concern: An ethical endorsement was derived from Al-Hussein teaching hospital authority; an enlightened acceptance was adopted from all participators.

Study tools:The questionnaire: Unique design of the questionnaire was fabricated for harvesting the information and was inspected and gauged by three official persons (of community medicine and physician) for checking the validness and prospering of the questionnaire. The questionnaire delineated by 2 subsections:

First subsection: Inclusive of questioning relevant to identity (age, gender, marital status, residency, and occupation).

Second subsection: Inclusive of questioning relevant to manifestations of present complaints, history of preexisting co-morbidities, duration of manifestations, existing and precedent medications consumption.

Third subdivision: Assigning complexities evolved for the time of accessing care in isolation division of the hospital that direct referral of patients to intensive care department inclusive of recovery and death.

Diagnostic procedures: Every patient was appraised by focus clinical examination inclusive of vital signs, oxygen saturation was determined for all participators to assign for grading of severity, the patients were rendered hypoxic when oxygen saturation is lower than 93% ¹³. The radiological valuation was carried out for all participators by CT-scan of the chest that skillful radiologist interpreted to prescribe abnor-

malcy. Polymerase chain reaction handled on nasopharyngeal swab sample was practiced upon entrance to the medical ward. All candidates submitted to laboratory assessment by complete blood count to count on monocytes; the average count of monocytes in the laboratory of our hospital is 200-800 cell/microliter. The participators were partitioned into monocytopenic when monocytes lower than 200 cells/microliter and non-monocytopenic, non-monocytopenic is further partitioned into normomonocytic in those possessing average monocytes to count and those with monocytosis when monocytes count greater than 800 cells/microliter. Laboratory indicant of inflammatory reactants was explored in sort of ferritin, C-reactive protein titer (CRP), ESR, lactate dehydrogenase (LDH), d-dimer, creatinine phosphokinase (CPK), and interleukin-6. An estimation of organ vitality was implemented by blood urea, serum creatinine, liver function, coagulation profile, and arterial blood gas analysis. Cardiac function was gauged for all participants by troponin and electrocardiogram. The participators were categorized into critical and non-critical categories in agreement with the grading of severity. Noncritical-categories are of moderate and severe COVID-19 pneumonia stayed in the common isolation ward, cases of critical status transposed to ICU. Critical severe type, meet any of the following: (a) respiratory failure occurred and mechanical ventilation was required; (b) shock happened; (c) patients complicated with organ failure required ICU admission¹⁴. Any patient match WHO criteria for discharge was appointed to be recovered¹⁵.

Statistical analysis: statistical package of social sciences (SPSS) version 25 was employed for data assay, descriptive statistics, frequencies, percentages, associations, the test of significance (Chi-square, Fischer exact test, T-test, and ANOVA test) was employed for interpretation for categorical variables, means, and standard deviation were employed to present data of continuous variables. A p-value of less than (0.05) was specified as statistically significant.

Results

A sum of 100 participators engaged within the analysis, dispersed evenly in two categories (COVID-19) as cases and (influenza A) as control, mean age of the overall sample was 64.4, mean age of case group was 63.54 simultaneously for control was 65.26 with no significant statistical association (P-value: 0.599). Come on, bio-demographic attributes mostly they displayed no significant statistical association where P values greater than 0.05. Whilst gender was statistically significant amongst two categories (P-value: 0.001) as illustrated in Table 1. Pertaining to co-morbidities, hypertension and pulmonary disease display significant statistical difference among cases and control where P values: 0.0001 and 0.002 respectively, whilst no significant statistical difference between cases and control in respect to diabetes mellitus and heart disease where P values: 0.834 and 0.545 respectively as shown in table 2. 66%, 26% and 8% of COVID -19 category had monocytopenia, normal count, and monocytosis respectively whilst 12%,54%, and 34% of influenza A category had monocytopenia, normal count, and monocytosis respectively with a significant statistical difference where P value = 0.001 as displayed in Figure 1.

74% of COVID-19 were critical whilst 16% of control were critical, 26% of COVID-19 group were non-critical whilst 84% of the control group were non-critical with a significant difference where (p value=0.001), as illustrated in figure 2. There is the significant statistical difference amongst cases and controls in all outcomes (ICU admission, death, and recovery) where (P-value=0.001 for all) as displayed in table 3.

37 participators with COVID-19 were critically ill, 35 died and 2 recovered, 30 dead were monocytopenic, 1 was normomonocytic whilst 4 have had monocytosis, 2 who were recovered were normomonocytic with significant statistical significance where P value=0.0001 for both death and recovery in critical patients in relation to monocytes count as exhibited in table 5.

Of 33 COVID-19 participators, 30 people with COVID-19 who were monocytopenic died, 4 with monocytosis died whilst just 1 normomonocytic died, 12 with normomonocytic recovered, whilst no participator with monocytosis recovered, with significant statistical significance between recovery and death where (P value =0.001) for both as displayed in table 6.

		Р	CR		Chi-	
		COVID-19	Influenza A	Total	Square P value	
	Male	36	20	56		
Conder	Iviale	64.3%	35.7%	100%	10.390	
Gender	Famala	14	30	44	0.001	
	Female	31.8%	68.2%	100%		
		18	12	30		
	Governmental	60.0%	40.0%	100%		
Occupation	o	5	6	11	1.731	
Total	Self-employer	45.5%	54.5%	100%	0.630	
		8	9	17		
	Housewife	47.1%	52.9%	100%		
	Retired	19	23	42		
	Retired	45.2%	54.8%	100%		
	Single	1	0	1		
Marital	Single	100%	0.0%	100%	1.010	
status	Married	49	50	99	0.315	
	Married	49.5%	50.5%	100%		
	Magiriya	15	14	29		
	Nasiriya.	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	100%			
	Suq –	14	13	27		
	Alsyookh	51.9%	48.1%	100%		
Desidence	Chatra	7	9	16	0.471	
Residence	Shatra	43.8%	56.3%	100%	0.631	
		8	9	17		
	Chibayesh	47.1%	52.9%	100%		
	Difess	6	5	11		
	Rifaee	54.5%	45.5%	100%		
Total		50	50	100		
Total		50.0%	50.0%	100%		

Table 2: Distribution according to comorbid diseases								
			PCR	Total	Chi- square			
		COVID	Influenza A	Total	P value			
	No	8	29	37				
hyportoncion	NO	21.6%	78.4%	100%	18.919ª			
hypertension	Yes	42	21	63	.0001			
	res	66.7%	33.3%	100%				
	No	30	27	57				
Heart disease		52.6%	47.4%	100%	.367ª			
neart uisease	Yes	20	23	43	.545			
	res	46.5%	53.5%	100%				
	No	17	18	35				
Diabetes	NO	48.6%	51.4%	100%	.044ª			
mellitus	Yes	33	32	65	.834			
	res	50.8%	49.2%	100%				
	No	42	17	59				
Pulmonary	NU	71.2%	28.8%	100%	25.837ª			
diseases	Vaa	8	33	41	0.002			
	Yes	19.5%	80.5%	100%				
Total		50	50	100				
iotai		50.0%	50.0%	100%				

Figure 1: Distribution of cases and control according to monocyte count. Chi-square=31.640, P value=0.001.

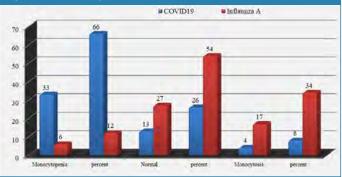
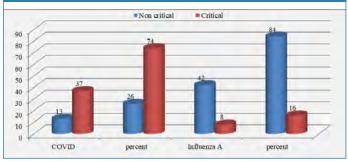


Table 3: Difference in distribution of main point outcomes of CO-VID-19 and influenza

		T	otal		
		COVID-19	Influenza A	Total	X ² , P value
Admission	Isolation ward	13	42	55	
		23.6%	76.4%	100%	33.980ª, 0.001
	ICU	37	8	45	33.700, 0.001
		82.2%	17.8%	100%	
Recovery	No	35	6	41	
			14.6%	100%	34.766ª,
	Yes	15	44	59	0.001
		25.4%	74.6%	100%	
Death	No	15	44	59	
		25.4%	74.6%	100%	34.766ª, 0.001
	Yes	35	6	41	34.700-, 0.001
		85.4%	14.6%	100%	
Total		50	50	100	
		50.0%	50.0%	100%	





	Outcome		Monocytop	penia		Normal monocyte Monocytosis							
		COVID- 19	Influenza A	Total	X ² , P, odd's ratio	COVID- 19	Influenza A	Total	X ² , P, odd's ratio	COVID- 19	Influenza A	Total	X ^{2,} P, odd's ratio
		3	0	3	.591 ^d	10	25	35	1.970 ^e	0	17	17	21.000
Severity	Non critical	100.%	0.0%	100%	0.442	28.6%	71.4%	100%	0.160	0.0%	100%	100%	0.001
erit,	Oritical	30	6	36	1 000	3	2	5	272	4	0	4	0.067
`	Critical	83.3%	16.7%	100%	1.200	60.0%	40.0%	100%	.267 -	100%	0.0%	100%	
Þ	Isolation word	3	0	3	.591 ^d	10	25	35	1.970 ^e	0	17	17	
dmi	ISUIALIUTI WULU	100%	0.0%	100%	.442	28.6%	71.4%	100%	.160	0.0%	100%	100%	21.000 ^r 0.001 a
Admission	1011	30	6	36	1 200	3	2	5	2/7	4	0	4	
ă	ICU	83.3%	16.7%	100%	1.200	60.0%	40.0%	100%	.267	100%	0.0%	100%	
-	Abcont	30	6	36	.591 ^d	1	0	1	2.130 ^e	4	0	4	
Recovery	Absent	83.3%	16.7%	100%	.442	100%	0.0%	100%	.144	100%	0.0%	100%	21.000
over	Dresent	3	0	3	022	12	27	39	2.250	0	17	17	0.001a
<	Present	100%	0.0%	100%	.833	30.8%	69.2%	100%	3.250	0.0%	100%	100%	
	Abcont	3	0	3	.591 ^d	12	27	39	2.130 ^e	0	17	17	21.000 ^f
Death	Absent	100%	0.0%	100%	.442	30.8%	69.2%	100%	.591ª	0.0%	100%	100%	0.001
ath	Dracant	30	6	36	1.200	1	0	1	200	4	0	4	.058
	Present	83.3%	16.7%	100%	1.200	100%	0.0%	100%	.308	100%	0.0%	100%	
	Total	33	6	39		13	27	40		4	17	21	
	IUIdI	84.6%	15.4%	100%		32.5%	67.5%	100%		19.0%	81.0%	100%	

						1
Critical- Severity			Recov	very	Total	FE, p value
	erniour eevenity		No	Yes	Total	TE, p value
	Monocytopenia	Count	30	0	30	
	Monocytopenia	%	100.0%	0.0%	100.0%]
	Normal	Count	1	2	3	23.962
Recovery	Normai	$\begin{array}{c c c c c c c c c c c c c c c c c c c $	100.0%			
, , , , , , , , , , , , , , , , , , , ,	Monocytosis	Count	4	0	4	0.0001
				0.0%		
	Total			2		
	TOLAI	%	94.6%	5.4%	100.0%	
			No	Yes		
	Monocytopenia	Count	0	30	30	
		%	0.0%	100.0%	100.0%	1
Death	Normal	Count		1	3	1
Douin		%	66.7%	33.3%	100.0%	23.962
	Monocytosis	Count	0	4	4	0.0001
		%	0.0%	100.0%	100.0%	
	Total	Count	2	35	37	
	, otai	%	5.4%	94.6%	100.0%	1

Table 6: Comparison of monocyte between recovered patients and died among COVID-19

			Death	Recovery	Total	Chi-square, p value
	Monocytopenia	Count	30	3	33	
	Monocytopenia	%	90.9%	9.1%	100.0%	
Monoovto	Normal	Count	1	12	13	32.617
Monocyte		%	7.7%	92.3%	100.0%	0.001
	Monocytosis	Count	4	0	4	
		%	100.0%	0.0%	100.0%	
Total		Count	35	15	50	
TOLAI		%	70.0%	30.0%	100.0%	

Discussion

In our analysis, we are approaching substantive perceptions: two-thirds of COVID-19 presented with reduced monocytes count, and the bulk of influenza A participators have had more elevated monocytes than counterpart COVID-19 with significant statistical difference implying that monocyte is a vigorous diagnostic indicant that is useful for discrimination, we are nearly in line with Chen et al.¹⁶. Monocytes were much lower in passed away COVID-19 participators than those who remained alive with significant difference between fatality and recovery patients according to monocytes (P value: 0.001), and monocytes were anticipant for death in COVID-19 by 20 folds (Odd ratio=1.200). Our study determinations are concordant with Pakos et al.¹⁷ that perceive dead persons had lower monocytes and had an inverse relation with fatality. Our study 30 monocytopenic who were critically ill with COVID-19 died (P value=0.0001), concluding that monocytopenia is intimately interrelated with lethality in critically COVID-19. A study done by Blomme et al.¹⁸ extrapolating that passed away patients with COVID-19 had lower monocytes than alive patients. Our analysis's Pivotal determination is that monocytes are lower in critical than non-critical grouping and it anticipating severity by 17 folds (Odd ratio=1.200) and anticipates admittance to ICU (odd ratio=1.200), a recent meta-analysis by Bao et al.19 surmised that monocytes is much reduced in severely affected clients. In contrast to our study, Bastug et al. 20 did not set up dissimilarity between critical and non-critical patients in regard to monocytes in COVID-19, this is owing to the fact that there might be a difference in study population or design of the study. But Anurag et al.²¹ declare comparable findings to our discernment in that monocytes discordantly reciprocating with severity. Peculiarly bulk of recovered patients 12 out of 13 with COVID-19 had normal monocytes count. Two out of three of critical normomonocytic clients were saved, substantiate impactful evidence that normalizing monocytes at any grade of severity might harbinger the onset of recovery. Fathi et al.22 ascertained that COVID-19 patients had appreciably higher monocytes sooner in recovered patients with COVID-19. Strikingly all critically ill patients in the COVID-19 grouping who undergone admittance to ICU with monocytosis died with high statistical significance disclosing that monocytosis is an expectant for lethality and most facets of dreadful terminal events of COVID-19. Mei et al.23 revealed the hospitalized patients with COVID-19 exhibiting monocytosis in the midst of other laboratory variables go through all spans of unfavorable terminal events with exposing the worse prognostication properties. Looking at a bio-demographic trait of our analysis, cases, and control were in a nearly exact fitting situation to elude from selection bias, so most traits displayed no significant statistical relation. Come to comorbidities in our study; COVID-19 grouping demonstrates a sizable number

of underlying pre-existing conditions, hypertension was the dominating then diabetes mellitus, heart disease and the slightest is pulmonary illnesses, our discerning in agreement with Guan et al.²⁴. There is significant statistical interrelation joining cases and control regarding hypertension and pulmonary disorders where P values=0.001 and 0.002 respectively in accordance with Shen et al.^{25,} but there was no significant interrelation regarding diabetes and cardiac diseases where P values=0.834, 0.545, respectively, this is nearly compatible with what was attained by virtue of Zayet et al.²⁶

Conclusion

Deviation in monocytes count from the norm is a valuable discriminator for diagnosis of COVID-19 and suitable anticipator overall spectrum of adverse consequences.

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