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RESEARCH ARTICLE

EVALUATION OF T- REGULATORY CELLS IN BREAST CANCER.

Ola Sayed Mohamed Ali¹, *Maha A. El-Taweel², Iman Attia Abdel El-Gawad² and Eman Ahmed Gouda³.

1. Professor of Biochemistry, Faculty of Pharmacy for girls, Al-Azhar University.
2. Associate professor of clinical pathology, National Cancer Institute, Cairo University.
3. B.Sc. of pharmaceutical sciences, Faculty of Pharmacy, Cairo University.

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Abstract

Treg cells have an important role in cancer, enabling tumors to elude host anti-tumor immune response. CD4+CD25+ Treg cells constitute 5–10% of peripheral blood CD4+ lymphocytes. Also, CD8+CD25+ cells have been shown a suppressive effect and share phenotypic and functional features with CD4+CD25+ Treg cells.

Aim of the work: This work was done to evaluate the frequency of expression of different Treg cells in the peripheral blood of breast cancer patients and to elucidate their impact on clinical and laboratory parameters.

Patients and Methods: This study included forty-seven Egyptian breast cancer patients (30 de novo patients and 17 after surgery patients) and 13 healthy controls and 18 benign breast patients. They were tested by Flow Cytometry

Results: CD4+CD25+ Treg cells were significantly higher in the de novo compared to the healthy controls and in the after-surgery group compared to the healthy controls and benign breast groups ($P < 0.0001$), while CD4+CD25hi were significantly higher in both de novo and after surgery patients as compared to healthy controls ($P < 0.0001$).

Conclusion: some Treg cells as CD4+CD25+ cells are predominant in breast cancer patients.

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Introduction: -

Breast cancer is the most frequently diagnosed cancer and the leading cause of cancer deaths among females worldwide, accounting for 23% of the total new cancer cases and 14% of the total cancer deaths in 2008 (Jamal et al. 2011). Breast cancer patients have immune system functional deviations, that affect immune cells that infiltrate the tumor, but also cells from the blood compartment, even in the absence of a metastatic disease (Pinho et al., 2016). Regulatory T-cells (Treg cells) are a variety of cells that display the capacity to suppress T-cell proliferation in vitro and prevent immune pathology in vivo Treg cells have an important role in cancer, enabling tumors to elude host anti-tumor immune response (Maggiet al., 2005).

CD4+CD25+ Treg cells constitute 5–10% of peripheral blood CD4+ lymphocytes (Maggiet al., 2005). They express CD25 (the alpha chain of the IL-2 receptor), cytotoxic T lymphocyte-associated antigen-4 (CTLA-4), a glucocorticoid induced TNF receptor (GITR), and a novel member of the fork head transcription factors (Foxp3) (Zhang et al., 2007).

Corresponding Author:- Maha A El-Taweel.

Address:- Associate professor of clinical pathology, National Cancer Institute, Cairo University.

CD8+CD25+ cells have been recently shown a suppressive effect. CD8+CD25+Treg cells share phenotypic and functional features with CD4+CD25+ Treg cells (Cosmi et al., 2003).

The aim of this study is to evaluate the frequency of expression of different Treg cells in the peripheral blood of Egyptian breast cancer patients and compare their expression with healthy controls and benign breast patients. In addition, to study the relationship between different Treg cells and clinical and laboratory findings and response to treatment in such patients.

Patients and Methods: -

Forty-seven breast cancer patients, 30 de novo patients (before surgery) and 17 after surgery patients, 13 benign breast disease patients, and 18 apparently healthy volunteering females as a control group were included in this study. The patients were presented to the outpatient's clinic at the National Cancer Institute (NCI), Cairo University over a period of consecutive 11 months. Malignancy was confirmed by histopathological examination. Written informed consent was obtained before enrollment into the study according to ethical committee. Investigations performed included routine hematological and biochemical investigations, and CEA and CA15.3 evaluations.

Methods: -

Flow cytometric analysis of Treg cells: -

Isolation of mononuclear cells and cell culture: -

Three to four ml of heparinized blood from each sample was added to 15 ml centrifuge tube and diluted 1:1 with sterile PBS without Magnesium or Calcium salts (*Biowest, France catalog no: L0615*) under sterile conditions. Ficoll-Hypaque media of density 1.077 gm/ml (*Lymphosep, Lymphocyte Separation Media- Biowest, France Catalog no: L0560*) was used to separate the PBMCs. The mononuclear cells were washed and re-suspended under sterile conditions in 3 ml RPMI 1640 with L-Glutamine medium (*Biowest, France catalog no: L050*) containing 10% Fetal Bovine Serum (*Biowest, South America catalog no: S1810*) and 2 % Antibiotic-Antimycotic 100x (*Biowest, France catalog no: L0010*) which contains Penicillin G, Streptomycin Sulfate & Amphotericin B. Two ng/ml IL2 (*Sigma, catalog no: 151342*) were added for T cell activation. The cells were incubated at 37°C, 5% CO₂ for 48 hours.

Monoclonal Antibodies: -

Fluorescent labeled mouse monoclonal antibodies anti-Human CD4-FITC (*Catalog no: F0766, Clone MT310*), CD8-PE (*Catalog no: R0806, Clone DK25*), CD25-FITC (*Catalog no: F0801, Clone ACT-1*), CD25-PE (*Catalog no: R0811, Clone ACT-1*) and CD3-RPE-Cy5 (*Catalog no: C7067, Clone UCHT1*) were obtained from DakoCytomation, Denmark.

Flowcytometric analysis: -

CD4+CD25+ Treg cells were studied with triple staining of anti-CD4 FITC, anti-CD25 PE and anti-CD3 RPE-Cy5, and CD8+CD25+ Treg cells were studied with anti-CD25 FITC, anti-CD8 PE and anti-CD3 RPE-Cy5 using COULTER® EPICS® XL™ Flow Cytometer instrument. To determine the percentage of CD4+Treg cells, lymphocytes were gated by plotting forward versus side scatter followed by gating on CD3+CD4+ T cells. These cells were then analyzed for CD25 expression; CD4+CD25hi were also identified as a tail from the major population of CD4+CD25 cells (Baumgartner and McCarter, 2009). CD8+ Treg cells were detected as percentage of CD8+CD25+ population gated from CD3+ lymphocytes (Fig 1).

Statistical analysis: -

Data management and analysis were performed using the Statistical Analysis System (SPSS) version 17 software. Comparison of groups with respect to numerical variables was done using the Mann-Whitney test. The Kruskal Wallis test was used to compare medians of 3 or more independent groups. Spearman's rank correlation was used to measure the strength of association between 2 numerical variables. All p-values were two sided, P-values ≤ 0.05 were considered significant.

Results: -

Characteristics of the de novo, after surgery & benign patients' groups are shown in (Table 1.a & 1.b).

CD4+ lymphocytes % were higher in normal controls and de novo patients compared to the after-surgery patients (P=0.04) (Table 2).

CD4+CD25+ Treg cells were significantly higher in the de novo compared to the healthy controls and in the after-surgery group compared to the healthy controls and benign breast groups ($P < 0.0001$), while CD4+CD25hi were significantly higher in both de novo and after surgery patients as compared to healthy controls ($P < 0.0001$) (Table 2).

A significant positive correlation was detected between CD4+CD25+ and CD4+CD25hi Treg cells in both de novo and benign patients' groups, while a significant negative correlation was detected between CD4+CD25 and CD8+CD25+ Treg cells in the after-surgery group (Table 3).

CD4+CD25+ Treg cells showed significant negative correlation with age in both de novo and after surgery groups ($P = 0.038$, $r = -0.388$ & $P = 0.012$, $r = -0.612$ respectively), and significant positive correlation with CEA in the benign group ($P = 0.005$, $r = 0.722$). CD8+CD25+ Treg cells showed significant positive correlation with age in the after-surgery group ($P = 0.048$, $r = 0.487$) (Table 4).

CD4+CD25+ and CD4+CD25hi Treg cells were higher in premenopausal compared to postmenopausal de novo breast cancer patients ($P = 0.017$ and 0.051 respectively). CD4+CD25hi Treg cells were significantly higher in the non-metastatic compared to the metastatic patients in the de novo group ($P = 0.041$), while CD4+CD25hi Treg cells were higher in grade III compared to grade II patients in the after-surgery group ($P = 0.011$) (Data not shown)

Discussion: -

In Egypt, breast cancer ranked the first, constituting 17.50% of total malignancies in females (Meloni et al., 2006).

Although T cells represent the most important immunological response in tumor growth, they become suppressive CD4⁺ and CD8⁺Treg cells after chronic stimulation and interactions with tumor cells, thus promoting rather than inhibiting cancer development and progression (Mokhtar et al., 2007).

In this study, CD4+ lymphocytes were predominant in peripheral blood of de novo breast cancer patients and healthy controls groups compared to the after-surgery group ($P = 0.04$).

Mozaffari et al. (2007) reported the same predominance of absolute number of CD4+ lymphocytes although it was statistically non-significant. Leong et al. (2006) showed that CD4+ lymphocyte % was the 2nd predominant subset (after CD8+) in the tumor microenvironment. Different sample analyses and different sample sizes may be the contributing factors for this discrepancy.

Tumor cells commonly participate in the generation of Treg cells, which provides an explanation for the observation that elevated numbers of Treg cells have been found in many types of cancer. It appears that TGF- β , secreted by the tumor itself or tumor-stimulated myeloid cells, plays a central role in tumor-mediated development of Treg cells by converting naïve T cells into Treg cells (Perez et al., 2007).

As regards CD4+CD25+ and CD4+CD25hi, they were significantly higher in the de novo and the after-surgery cancer patients compared to the healthy controls and the benign patients' group.

Our result is consistent with Wang et al. (2011) who found the same result in their study on peripheral blood of de novo breast cancer patients.

In our study, CD4+CD25hi subset was higher in the after-surgery patients compared to the de novo patients ($P = 0.04$) and this was in accordance with the finding of Mozaffari et al (2007).

CD8+CD25+Treg cells share phenotypic and functional features with CD4+CD25+ Treg cells (Cosmi et al., 2006). We detected that the benign patients group had the highest percentage of CD8+CD25+% Treg cells subset, followed by the after-surgery group. The de novo breast cancer patients and healthy controls group were almost the same, the differences were non-significant. Mozaffari et al. (2007) reported a significant increase in CD8+CD25hi Treg cells in breast cancer patients after receiving adjuvant therapy compared to the de novo patients. Consistently, Shao et al. (2017) reported that the proportion of regulatory T cells (Treg) after chemotherapy in the disease control patients decreased significantly compared with that of the progressive patients and this difference affected

significantly the overall survival. Pinho et al (2016) found that breast cancer patients have higher percentage of effector CD8+ T cells compared to healthy donors.

T-reg cells can offer a new insight into improving cancer therapies. Combined chemotherapeutic and immunomodulatory agents that showed their effect on T-reg can lead to marked suppression of tumor growth and prolongation of survival time (Xia et al., 2017 and Hekim et al., 2017).

A significant positive correlation was detected between the CD4+CD25+ % and the CD4+CD25hi % in the de novo breast cancer patients group ($P < 0.0001$, $r = 0.673$) and in the benign group ($P = 0.002$, $r = 0.79$), while there was a significant negative correlation between the CD4+CD25+ % and CD8+CD25+ % in the after-surgery patients group ($P = 0.018$, $r = -0.583$).

In agreement with our results, Zhu et al. (2015) and Leong et al. (2006) have demonstrated an insignificant negative correlation between the CD4+CD25+ cells and CD8+CD28+CD25+ obtained from TILs of breast cancer patients. They also found a significant positive correlation between CD4+CD25+ and CD8+CD28- subsets. In that study, CD8+CD28+CD25+ were defined as effector T cells and CD8+CD28- subset as Treg cells.

In our study, the observed negative relationship may be explained by the finding that not all CD8 + CD25+ population are Treg cells (Bisikirska et al., 2005).

Correlating CD4+CD25+ % and CD8+CD25+% with some of the prognostic factors of breast cancer revealed a significant negative correlation between the CD4+CD25+ % and the age in the de novo and after surgery groups. However, a significant positive correlation was found between CD8+CD25+% and age in the after-surgery group.

Similarly, Chui et al. (2004) reported that CD4+ Treg cells were higher in the blood of breast cancer patients < 65 years in age. Tsaknaridis et al. (2003) also demonstrated that the suppressive activity of human CD4+CD25+ T cells declines with age in his study done on normal subjects. In contrast, Leong et al. (2006) found a significant positive correlation with age (> 50 years old).

This diversity of results may be due to differences in sample size or ethnic diversity.

Regarding tumor markers, a significant positive correlation was detected in the benign patients between CD4+CD25+ % and CEA ($P = 0.005$, $r = 0.722$), while CA15.3 didn't show any significant results. In contrast, Abo-Elainen et al. (2008) reported a significant positive correlation between CD4+CD25+% in the peripheral blood of 30 female breast cancer patients and CA15.3 ($P = 0.880$, $r = 0.001$). This may be explained by the different stages of breast cancer patients enrolled in both studies.

CD4+CD25hi % was significantly higher in the non-metastatic group compared to the metastatic group in de novo patients ($P = 0.041$). Many studies have demonstrated that a higher Treg cells number within the tumor microenvironment or in peripheral blood would imply a worse prognosis and reduced survival (Wilke et al., 2010).

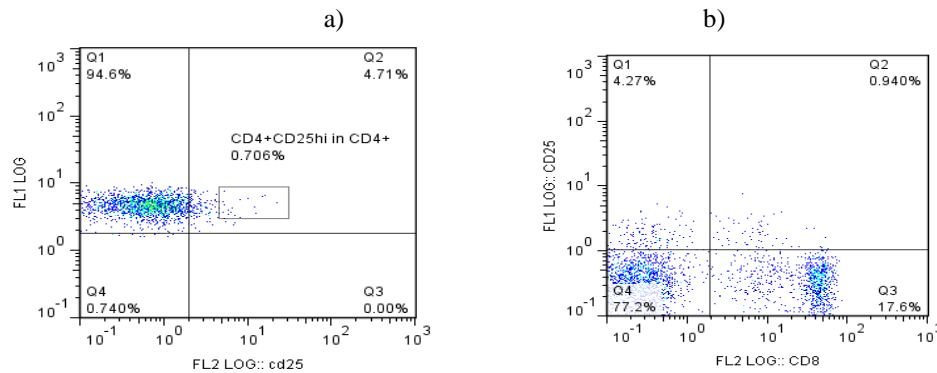
Significant higher percentage of CD4+CD25hi in the after-surgery group was detected in grade III compared to grade II ($P = 0.011$). This was inconsistent with Leong et al. (2006).

We didn't find any significant relationship between CD4+CD25+ % or CD4+CD25hi % and the tumor stage, ER, PR or Her2neu status in both the de novo and the after-surgery patients groups. Our results are consistent with Song et al (2015) but different from Perez et al. (2007). Also, Mahmoud et al. (2011) and Bates et al. (2006) found significant correlations between Treg cells infiltration and ER negativity in breast cancer patients. Liu et al (2014) found that CD4+CD25+FOXP3+ regulatory TILs are a poor prognostic indicator in ER+ breast cancer, but a favorable prognostic factor in the HER2+/ER-ve subtype. Leong et al. (2006) found a significant positive correlation between CD4+CD25- T subset of TILs in breast cancer patients and ER. Estrogen has been reported to play a role in regulating the activation of human T cells particularly CD4+ and CD8+ T cells. The importance of ER- α in regulating the immune system is further supported by an in vivo murine model, where a high frequency of immature double CD4+CD8+ thymocytes was found in ER- α negative mice as compared to the ER- α positive mice (Erlandsson et al., 2001).

This discrepancy between results may be attributed to the predominance of ER positivity in our studied population, different ethnic groups or menopausal status of the studied population.

In conclusion, some Treg cells as CD4+CD25+ cells are predominant in breast cancer patients, and show negative correlation with age and with CD8+CD25+ lymphocytes, which are predominant in the benign patients and after surgery patients.

Treg cells may play a role in modulation of effector T cell responses against breast tumors. Some Treg cells such as CD4+CD25+ cells maybe increased in cases of malignancy, while others, as CD8+CD25+ lymphocytes seem to play a protective role against malignancy. Further studies and larger sample sizes are needed to clarify the diagnostic and prognostic role of Treg cells in breast cancer progression. Treg can act as an immune-modulator in the therapeutic treatment of breast cancer patients weather alone or in combination with therapeutic vaccines and cytokines.



(Fig 1) a):- Positive expression of CD4+CD25+ and CD4+CD25hi subsets gated from CD4+ lymphocytes in a de novo patient. b) CD8+CD25+ subsets histogram from CD3+ lymphocytes gate in an after-surgery patient.

Table (1.a):- Clinical data of the de novo, after surgery and benign patients.

		De novo patients		After-surgery patients		Benign patients		Total	
		n = 30	Percentage %	n= 17	Percentage %	n= 13	Percentage %	n= 60	Percentage %
Age	< 50 years	14	47	7	41	7	54	28	47
	≥ 50 years	16	53	10	59	6	46	32	53
Family history	Positive	4	13	0	0	0	0	4	7
	Negative	26	87	17	100	13	100	56	93
Marital status	Married	29	97	17	100	9	69	55	92
	Single	1	3	0	0	4	31	5	8
Menarche status	Pre-menopause	12	40	7	41	9	69	28	47
	Post-menopause	18	60	10	59	4	31	32	53

n: number of patients.

Table (1.b):- Clinical and pathological data of the de novo and after surgery patients.

		De novo patients		After surgery patients		Total	
		n = 30	Percentage	n= 17	Percentage	n= 47	Percentage
Tumor size	Tumor size \leq 5 cm	24	80	12	71	36	77
	Tumor size > 5 cm	6	20	5	29	11	23
Lymph node status	Positive	29	97	13	76	42	89
	Negative	1	3	4	24	5	11
Distant metastasis	Present	14	47	5	29	19	40
	Absent	16	53	12	71	28	60
Tumor stage *	Stage I	1	3	0	0	1	2
	Stage II	9	30	7	41	16	34
	Stage III	5	17	4	24	9	19
	Stage IV	15	50	6	35	21	45
Grade	II	26	87	12	71	38	81
	III	4	13	5	29	9	19
Tumor focality	Unifocal	26	87	14	82	40	85
	Multifocal	4	13	3	18	7	15
ER	Positive	24	80	10	59	34	72
	Negative	6	20	7	41	13	28
PR	Positive	11	37	14	82	25	53
	Negative	19	63	3	18	22	47
HER2/neu	Positive	4	13	7	41	11	23
	Negative	26	87	10	59	36	77

* Stage according to TNM classification for breast cancer (AJCC).

n: number of patients, **Grade:** Pathological grade, **ER:** Estrogen receptor, **PR:** Progesterone receptor, **HER2/neu:** Human Epidermal growth factor Receptor 2.

Table (2):- Comparison of different lymphocyte populations in the different studied groups (de novo, after surgery, benign patients and healthy controls) using Krusal-Wallis test:

	Healthy controls	De novo patients	After surgery patients	Benign patients	P-value
	Median & IR (Q1 – Q3)	Median & IR (Q1 – Q3)	Median & IR (Q1 – Q3)	Median & IR (Q1 – Q3)	
Total lymphocytes %	84.7 (82 – 87.8)	88.65 (86 – 92.3)	89.2 (81 – 91)	90 (88 – 91)	0.15
CD3+ lymphocytes %	76.95 (67.6 – 88.1)	69.8 (58.2 – 85.1)	67.15 (47.8 – 82.6)	77.8 (64.6 – 85.4)	0.27
CD4+ lymphocytes %	65.25 a (61.9 – 72.4)	65.6 b (57.5 – 75)	55.7 a,b (46.5 – 67)	52.7 (51 – 71.9)	0.04
CD8+ lymphocytes %	17 (12.1 – 20.5)	16.5 (13.5 – 20.9)	19.7 (12.7 – 33.7)	16.6 (9.4 – 20.9)	0.57
CD4+/CD8+ ratio	3.57 (3.4 – 5.9)	3.79 (3.3 – 4.6)	2.83 (1.45 – 4.1)	3.46 (3.2 – 5.1)	0.13
CD4+CD25+ %	2.1 c,d (1.7 – 2.5)	3.78 c (3.1 – 4.6)	4.46 d,e (3.95 – 4.9)	3.11 e (1.9 – 3.6)	<0.0001
CD4+CD25hi %	0.27 f,g (0.15 – 0.35)	0.7 f (0.47 – 0.85)	0.8 g (0.7 – 1.1)	0.43 (0.22 – 0.57)	<0.0001
CD8+CD25+ %	0.81 (0.54 – 0.85)	0.89 (0.5 – 1.12)	0.94 (0.7 – 1.2)	0.96 (0.56 – 1.7)	0.24

IR: Interquartile range

* Groups with the same letter show statistically significant difference

Table (3):- Correlation analysis between the different Treg cells subsets % in the different patients' groups

			CD4+CD25hi %	CD8+CD25+ %
De novo patients	CD4+CD25+ %	r	0.673	-0.159
		P	< 0.0001**	0.438
	CD4+CD25hi %	r		-0.229
		P		0.261
After surgery patients	CD4+CD25+ %	r	0.33	-0.583
		P	0.212	0.018*
	CD4+CD25hi %	r		-0.007
		P		0.978
Benign patients	CD4+CD25+ %	r	0.79	-0.116
		P	0.002**	0.721
	CD4+CD25hi %	r		0.068
		P		0.842
* Significant.				
**Highly significant				

Table (4):- Correlation analysis between different Treg cells subsets % with the clinical, histopathological and laboratory data in the different patients' groups (de novo, after surgery and benign patients):

			CD4+CD25+ %	CD4+CD25hi %	CD8+CD25+ %	
Age	De novo	r	-0.388	-0.214	0.119	
		P	0.038*	0.266	0.554	
	After surgery	r	-0.612	0.267	0.487	
		P	0.012*	0.301	0.048*	
	Benign	r	0.29	0.035	0.248	
		P	0.336	0.914	0.437	
Tumor size	De novo	r	-0.075	-0.201	0.21	
		P	0.701	0.296	0.294	
	After surgery	r	0.025	-0.051	0.467	
		P	0.926	0.846	0.058	
	CEA	De novo	r	-0.279	0.034	-0.131
			P	0.143	0.859	0.515
After surgery		r	0.148	0.221	0.108	
		P	0.584	0.393	0.679	
Benign	r	0.722	0.493	0.155		
	P	0.005**	0.103	0.631		
CA 15-3	De novo	r	-0.025	-0.065	-0.002	
		P	0.896	0.737	0.993	
	After surgery	r	-0.091	-0.422	0.174	
		P	0.737	0.092	0.504	
	Benign	r	0.544	0.371	-0.315	
		P	0.055	0.236	0.318	

* Significant. ** Highly significant.

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