# **RESEARCH ARTICLE**

# Low Frequency of ETV6-RUNX1 (t 12; 21) in Saudi Arabian Pediatric Acute Lymphoblastic Leukemia Patients: Association with Clinical Parameters and Early Remission

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## Abstract

Background: Pediatric acute lymphoblastic leukemia (pALL) patients at King Abdulaziz Medical City represent a pure Saudi Arabian population. ETV6-RUNX1 positive pALL patients have good prognosis as compared to ETV6-RUNX1 negative counterparts. Therefore, frequencies of these two patient groups have a huge consideration in treatment strategies of pALL in a given population. Different geographical locations have been reported to have different frequencies of ETV6-RUNX1 ranging from 10% in Southeast Asia to 30% in Australia. Aim: Therefore, the objective of this study was to establish the ETV6-RUNX1 status of Saudi Arabian pALL patients and its association with clinical parameters and early remission. Materials and Methods: Clinical parameters and ETV6-RUNX1 status (using FISH technique) of pALL patients attending the Pediatric Oncology Clinic, King Abdulaziz Medical City, Riyadh from 2006 to 2011 were studied. Comparisons between ETV6-RUNX1 positive and negative groups were accomplished using chi-square test or Fisher's exact test. All statistical analyses were performed using SAS version 9.2 (SAS Institute, Inc., Cary, NC). Results: Out of 54 patients, 33 were male and 21 were females (ratio 1.57:1). B- and T-cell lineages were found in 47 (87%) and 7 (13%) patients respectively. Only 5 (9.3%) patients were ETV6-RUNX1 positive while 49(80.7%) were ETV6-RUNX1 negative. All ETV6-RUNX1 patients (100%) were of B-cell lineage and 80% (4/5) were in the 3-7 year age group. None of the ETV6-RUNX11 patients had ≥5% blasts (no remission) at day 14 as compared with 9% in the ETV6-RUNX1 negative group (Figure 1).Conclusions: Frequency of ETV6-RUNX1 positive patients (less than 10%) in our pALL patients is much lower than reported for most European countries, North America, Australia and Japan while it is in accordance with ETV6-RUNX1 frequencies from Egypt (11.6%), Pakistan (10%), Spain (2%) and India (5-7%). This shows ethnic differences in genetics of pALL as well as higher frequencies of ETV6-RUNX1 positive pALL mostly in more industrialized countries, probably due to some industrial pollutants or westernized lifestyle.

Keywords: Acute lymphoblastic leukemia - early remission - ETV6-RUNX1 - translocation - Saudi Arabia

Asian Pac J Cancer Prev, 16 (17), 7523-7527

## Introduction

Acute Lymphoblastic Leukemia (ALL) is the cancer of lymphoid lineage of blood forming pluripotent progenitor cells in the bone marrow (Bhojwani et al., 2015).ALL is characterized by an abnormal huge number of immature lymphocytes or blasts in blood and bone marrow (Ceppi et al., 2015). It is the most common cancer in children worldwide accounting for 25% of all cancers (Pui et al., 2015). National Cancer Institute reported a sharp peak in ALL incidence among children aged 2 to 3 years (>80 cases per million per year), with rates decreasing to 20 cases per million for ages 8 to 10 years. The incidence of ALL among children aged 2 to 3 years is approximately fourfold greater than that for infants and is nearly tenfold greater than that for adolescents aged 16 to 21 years (Ribera & Oriol, 2009). In Saudi Arabia, leukemia has been reported to be third most common cancer in males (Al-Ahmadi & Al-Zahrani, 2013) with ALL being the most common type of leukemia (Alghamdi et al., 2014; Al-Mutlaq et al., 2015). Unlike most of the western populations, Saudi Arabia has 41.7% of the population under 15 years of age, which put a large number of the population at risk of childhood cancer specifically ALL

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(Al-Mutlaq et al., 2015).

Many genetic abnormalities are found in Childhood ALL, which have implications in differential diagnosis/ classification, prognosis, drug selection at different phases of anti-leukemic treatment and in accessing the efficacy of the treatment (Cooper & Brown, 2015). ETV6-RUNX1 ((t (12; 21) (p13; q22)) is one of the most common chromosomal abnormalities in pediatric ALL patients and is associated with favorable prognosis and prolonged survival in majority of the patients (Moorman et al., 2014). In most of the European countries, US, Japan and Australia, ETV6-RUNX1 frequency has been reported to be 20-33% among pediatric ALL (Amor et al., 1998; Iqbal & Tanveer 2007; Awan et al., 2012; Inaba et al., 2013; Iqbal, 2014). Nevertheless, lower frequencies of ETV6-RUNX1 have been reported in some geographical regions (Kwong & Wong., 1997; Eguchi-Ishimae et al., 1998; Garcia-Sanz et al., 1999; Tsang et al., 2001; Rahman et al., 2006; Chung et al., 2010; Faiz & Qazi., 2010; Mazloumi et al., 2012; Iqbal et al. 2014). This shows ethnic differences in occurrence of this prognostically important genetic lesion.

and clinical outcome of childhood ALL from Saudi Arabia, reports regarding the clinical characteristics and clinical outcome of the prognostically important genetic abnormalities are lacking. Therefore, this study was carried out to find out the association of ETV6-RUNX1 positive and ETV6-RUNX1 negative pediatric ALL patients with clinical features and treatment outcome in pediatric ALL patients from King Abdulaziz Medical City, National Guard Health Affairs, Riyadh, Saudi Arabia, which are a true representation of Saudi ethnicity (Wikipedia for Saudi Arabian National Guard, 2015) and therefore best sample to show if there are ethnic differences in genetic epidemiology (e.g., ETV6-RUNX1 frequency) of Saudi pediatric ALL from other ethnic groups in the world.

## **Materials and Methods**

#### Patients

This study was conducted at Division of Pediatric Hematology/Oncology Department of Oncology / King Saud Bin Abdulaziz University for Health Sciences, King Abdulaziz Medical City, National Guard Health Affairs, Riyadh, Saudi Arabia. Inclusion criteria were all

Although there are few reports about the biology

Table 1. Association of Patient Characteristics with day 14 Response in Pediatric Acute Lymphoblastic Leukemia Patients

			Respons	Response Day 14			
Covariate	Statistics	Level	Poor Response N=4	Good Response N=45	P-value*		
gender	N (Row %)	female	1 (5.26)	18 (94.74)	0.555		
Age	N (Row %) N (Row %)	male <= 2 Years	3 (10) 0 (0)	27 (90) 9 (100)	0.471		
	N (Row %)	3 - 7 Years	2 (7.69)	24 (92.31)			
	N (Row %)	8 - 15 Years	2 (14.29)	12 (85.71)			
T-cell	N (Row %)	Negative	2 (4.65)	41 (95.35)	0.016		
	N (Row %)	Positive	100.Q (33.33)	4 (66 <del>.67)</del>			
B-cell WBC	N (Row %)	Negative	2 (33.33) 6.3	<b>10.1</b> <sup>4</sup> (66.67) 41 (95.35)	0.016		12
	N (Row %) N (Row %)	Positive <= 30000	2 (4.65) 7 c 0 <sup>1</sup> (2.7)	41 (95.35) 36 (97.3)			
					0.014		
	N (Row %)	> 30000	<b>75.0</b> <sup>1</sup> <sub>3 (25)</sub>	9 (75)	25.0	30.0	
TEL-AML1	N (Row %)	Negative	4 (9.09)	40 (90.91)	0.482		
	N (Row %)	Positive	0 (0) 56.3	<b>46.8</b> 5 (100)			51
* The p-value is c	alculated by chi-square	test	50.0	54.2	31.3	30.0	
Table 2. Assoc	•	<sup>test</sup>	h day 29 Respo	ed Acu ph		30.0	
	•		h day 29 Respo P 25.0	'ed Acu iph		30.0	
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Table 2. Assoc	•		h day 29 Respo P	'ed Acu iph	c Leukemia 31.3 P-value*		33
Table 2. Assoc Patients	ciation of Patient C	Characteristics with	h day 29 Respo P 	Yed         Acu         ph           ie I         38.0	c Leukemia		33
Table 2. Assoc     Patients     Covariate	ciation of Patient C Statistics	Characteristics with	h day 29 Respo 25.0 Poor Respons 0 0 (0) 1 (2 22)	Ped         Acu         ph           ae I         38.0         espa         23.7         50           21 (100)         20.0% (75)         50         50	<b>c Leukemia</b> <b>31.3</b> P-value* 0.398	30.0	
Table 2. Assoc     Patients     Covariate     gender	ciation of Patient C Statistics N (Row %)	Characteristics with Level female	h day 29 Respo 25.0 Poor Respons 0 0 (0) 1 (2 22)	Ped         Acu         ph           ae I         38.0         espa         23.7         50           21 (100)         20.0% (75)         50         50	<b>c Leukemia</b> <b>31.3</b> P-value* 0.398		
Table 2. Assoc     Patients     Covariate	ciation of Patient C Statistics N (Row %) N (Row %)	Level female <= 2 Years 3 - 7 Years	h day 29 Respo 25.0 Poor Respons 0 0 (0) 1 (2 22)	Ped     Acu     ph       ae I     38.0     esp     23.7     50       21 (100)     29 (96.67%)     9 (100)     100)     100)       trace     26 (100)     100)     100)	<b>c Leukemia</b> <b>31.3</b> P-value* 0.398	30.0	
Table 2. Assoc     Patients     Covariate     gender	ciation of Patient C Statistics N (Row %) N (Row %) N (Row %)	Level female <= 2 Years	h day 29 Respo 25.0 Poor Respons 0 0 (0) 1 (2 22)	Ped     Acu     ph       ae I     38.0     esp     23.7     50       21 (100)     29 (96.67%)     9 (100)     50       29 (96.67%)     9 (100)     51     50       25 (100)     15 (93.73%)     15 (93.73%)     15 (93.73%)	c Leukemia 31.3 P-value* 0.398	30.0	
Table 2. Assoc     Patients     Covariate     gender	ciation of Patient C Statistics N (Row %) N (Row %) N (Row %) N (Row %) N (Row %)	Level female <= 2 Years 3 - 7 Years	h day 29 Respo 25.0 Poor Respons 0 0 (0) 1 (2 22)	Ped     Acu     ph       ae I     38.0     esp     23.7     50       21 (100)     29 (96.67%)     9 (100)     50       29 (96.67%)     9 (100)     51     50       25 (100)     15 (93.73%)     15 (93.73%)     15 (93.73%)	<b>c Leukemia</b> <b>31.3</b> P-value* 0.398	30.0	
Table 2. Assoc         Patients         Covariate         gender         Age	ciation of Patient C Statistics N (Row %) N (Row %) N (Row %) N (Row %) N (Row %) N (Row %)	Level female <= 2 Years 3 - 7 Years 8 - 15 Years	h day 29 Respo 25.0 Poor Respons 0 0 (0) 1 (2 22)	Ped     Acu     ph       ae I     38.0     esp     23.7     50       21 (100)     29 (96.67%)     9 (100)     50       29 (96.67%)     9 (100)     51     50       25 (100)     15 (93.73%)     15 (93.73%)     15 (93.73%)	c Leukemia 31.3 P-value* 0.398 issi issi 0.328	30.0	
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Table 2. Assoc         Patients         Covariate         gender         Age         T-cell	ciation of Patient C Statistics N (Row %) N (Row %)	Level female male <= 2 Years 3 - 7 Years 8 - 15 Years Negative Positive	h day 29 Respo 25.0 Poor Respons 0 0 (0) 1 (2 22)	Ped     Acu     ph       ae I     38.0     esp     23.7     50       21 (100)     29 (96.67%)     9 (100)     50       29 (96.67%)     9 (100)     51     50       25 (100)     15 (93.73%)     15 (93.73%)     15 (93.73%)	c Leukemia 31.3 P-value* 0.398 0.328 0.006	30.0	
Table 2. Assoc         Patients         Covariate         gender         Age	Statistics           N (Row %)	Level female male <= 2 Years 3 - 7 Years 8 - 15 Years Negative Positive Negative	h day 29 Respo 25.0 Poor Respons 0 0 (0) 1 (2 22)	Ped     Acu     ph       ae I     38.0     esp     23.7     50       21 (100)     29 (96.67%)     9 (100)     50       29 (96.67%)     9 (100)     51     50       25 (100)     15 (93.73%)     15 (93.73%)     15 (93.73%)	c Leukemia 31.3 P-value* 0.398 0.328 0.006	30.0	
Table 2. Assoc         Patients         Covariate         gender         Age         T-cell         B-cell	Ciation of Patient C Statistics N (Row %) N (Row %)	Level female male <= 2 Years 3 - 7 Years 8 - 15 Years Negative Positive Negative Positive	h day 29 Respo 25.0 Poor Respons 0 0 (0) 1 (2 22)	Ped     Acu     ph       ae I     38.0     espe     23.7     50       21 (100)     29 (96.67)     50     50       15 (93.73)     15 (93.73)     15 (93.33)       45 (100)     5 (83.33)     5 (83.33)       9 40 (100)     100     100	c Leukemia 31.3 P-value* 0.398 0.328 0.006 0.006	30.0	
Table 2. Assoc         Patients         Covariate         gender         Age         T-cell         B-cell	Statistics           N (Row %)           N (Row %)	Level female male <= 2 Years 3 - 7 Years 8 - 15 Years Negative Positive Negative Positive <=30000	h day 29 Respo 25.0 Poor Respons 0 0 (0) 1 (3.33) 0 (0) 1 (6.25) 1 (16.67) 0 (0) 1 (16.67) 1 (16.67) 0 (0) 0 (0) 0 (0) 0 (0) 1 (16.67) 0 (0) 0 (0) 1 (16.67) 0 (0) 0	Ped     Acu     ph       ae I     38.0     esp     23.7     50       21 (100)     29 (96.67%)     9 (100)     50       29 (96.67%)     9 (100)     51     50       25 (100)     15 (93.73%)     15 (93.73%)     15 (93.73%)	c Leukemia 31.3 P-value* 0.398 0.328 0.006 0.006	30.0	33

\* The p-value is calculated by chi-square test

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pediatric Acute Lymphoblastic Leukemia (ALL) patients between the ages of 2 to 15 years. Study was approved by King Abdullah International Medical Research Centre (KAIMRC), National Guard Health Affairs, Riyadh, Saudi Arabia.

Along with other laboratory parameters for diagnosis and risk stratification, patients were tested for ETV6-RUNX1 using interphase fluorescent in situ hybridization (FISH) as a part of routine laboratory findings.

#### Interphase fluorescent in situ hybridization (FISH)

Selection Of Material: Vysis ETV6/RUNX1 DF FISH Probe Kit (Abbot Laboratories, Illinois, USA) was used to detect ETV6/RUNX1 fusion oncogene resulting from t(12;21)(p13;q22). FISH procedures were carried out according to manufacturer's instructions.

Pre-hybridization, Hybridization and Posthybridization: WBCs were washed with 1X PBS. Cells were fixed in methanol/acetic acid, dropped on slides, and air dried. The slides were pretreated with "0.01% pepsin + 0.02 M HCl" at 37°C for 10 min. The cells and probes were denatured on a heating plate together at 78°C for 10 min. Hybridization was performed overnight at 37°C. Posthybridization washing was done in 23 SSC containing 50% formamide for 7 min at 42°C followed by two washes in 23 SSC (42°C for 7 min). The slides were covered by Vectashield (Vector Laboratories, Burlingame, CA) containing 0.5 g/ml DAPI.

FISH Analysis: Slides prepared by FISH were analyzed using CytoVision 7.0 system (Applied Imaging, Biosciences Centre, Newcastle, UK).

## Treatment protocol & Clinically Follow-up

CCG1991 protocol was used for standard risk patients while CCG1961 protocol was used for high risk patients. Number of blasts at day 14 and day 29 of the treatment were also calculated as a part of routine clinical follow-up.

#### Response criteria

Complete remission (CR or M1) was defined as: Normal bone marrow (with <5% blasts and >25% cellularity), neutrophil counts  $>1.5\times109/1$ , platelet count  $> 100\times109/1$ , and all extramedullary disease resolved. Anything less than CR was considered as incomplete remission (M2 or M3).

## Statistical analysis

The association demographic data, clinical and laboratory parameters and ETV6/RUNX1 status was statistically studied using was performed using SAS version 9.2 (SAS Institute, Inc., Cary, NC). Comparison between ETV6-RUNX1 positive and negative groups done using chi-square test or Fisher's exact test.

# Results

Out of 54 patients, 33 were male and 21 were females (ration1.57:1). B- and T-cell lineage was found in 47 (87%) and 7 (13%) patients respectively. Overall, B-cell lineage and WBC count less than 30,000 were significantly associated with complete remission (M1) at day 14 and

day 29 (Tables 1 & 2).

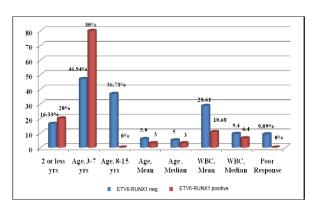
Only 5 (9.3%) patients with ETV6-RUNX1 positive while 49(80.7%) were ETV6-RUNX1 negative. Mean age of ETV6-RUNX1 positive and ETV6-RUNX1 negative patients was  $5.90\pm0.495$  and  $3\pm0.316$ , respectively (<0.001). Moreover, 80% (4/5) of ETV6-RUNX1 positive patients were in 3-7 year age group while 46.94% (23/49) ETV6-RUNX1 negative patients in this age group. Furthermore, though 36.73% of ETV6-RUNX1 negative patients were in 8-15 years age group, no ETV6-RUNX1 positive patients were in this age group (Table 3, Figure 1).

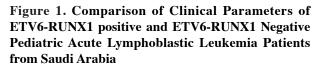
 Table 3. Demographics and Clinical Characteristics of

 ETV6-RUNX1 Positive and ETV6-RUNX1 Negative

 Pediatric Acute Lymphoblastic Leukemia Patients

			P-value*
	Negative (n=49)	Positive (n=5)	
Gender			
Female	19 (38.78)	2 (40)	0.957
Male	30 (61.22)	3 (60)	
Age			
Mean±SEM	5.90±0.495	3±0.316	< 0.001
Median	5	3	0.088
Range	1-13	2-4	
<= 2 Years	8 (16.33)	1 (20)	0.239
3 - 7 Years	23 (46.94)	4 (80)	
8 - 15 Years	18 (36.73)	0 (0)	
T-cell			
Negative	42 (85.71)	5 (100)	0.365
Positive	7 (14.29)	0 (0)	
B-cell			
Negative	7 (14.29)	0 (0)	0.365
Positive	42 (85.71)	5 (100)	
WBC (X103)			
Mean±SEM	28.61±8.64	$10.68 \pm 5.12$	0.083
Median	9.40	6.40	0.63
Range	0-400	3-30	
<=30	38 (77.55)	4 (80)	0.900
> 30	11 (22.45)	1 (20)	
Response @ 14			
Poor Response	4 (9.09)	0 (0)	0.482
Good Response	40 (90.91)	5 (100)	
Response@28			
Poor Response	1 (2.17)	0 (0)	0.739
Good Response	45 (97.83)	5 (100)	





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All ETV6-RUNX1 patients (100%) were of B-cell lineage as compared to 85.71% (42/49) of ETV6-RUNX1 negative patients (p=?). Similarly, none of the ETV6-RUNX1 positive patients was T-cell lineage as compared to 14.29% T-cell lineage ETV6-RUNX1 negative patients. Similarly, none of ETV6-RUNX1 positive patients had  $\geq$ 5% blasts (no remission) at Day 14 as compared to 9% patients from ETV6-RUNX1 negative group (Table 3, Figure 1).

## Discussion

Our study shows a low representation of ETV6-RUNX1 fusion oncogenes among pediatric ALL patients from King Abdulaziz Medical City, National Guard Health Affairs, Riyadh, Saudi Arabia. Most of ETV6-RUNX1 positive (80%) patients belonged to 3-7 years of age group and allETV6-RUNX1 positive showed early remission (day 14) as compared to 9% of ETV6-RUNX1 negative patients not showing day 14 remission.

Frequency of ETV6-RUNX1 in this study is in accordance with previous study from middle-east region reporting 14.7 % ETV6-RUNX1 frequency in pediatric ALL patients (Al-Mulla et al., 2014) although it is much lower than the 20-33% frequency in Europe, US, Japan and Australia (Shurtleff et al., 1995; Harbott et al., 1997; Kobayashi et al., 1997; Amor et al., 1998).

TEL-AML1 frequencies vary greatly in different countries of South East Asia and Southern Asia. A recent study from Pakistani reported the frequency of ETV6-RUNX1 in pediatric ALL patients to be 10.2% (Iqbal, 2014/10). ETV6-RUNX1 frequency has been found to be 17.1% in Korean pediatric ALL population (Chung et al., 2010), Taiwan 17% (Liang et al., 1996) and Malaysia 19% (Gill et al., 2005). In Egypt, ETV6-RUNX1 positivity among pediatric ALL has been reported to be 11.6% (Shaker et al., 2001). Although most of the studies reported higher ETV6-RUNX1 frequencies in Japan, one study from Hiroshima Japan reported it to be 10% (Eguchi-Ishimae et al., 1998). Contrary to other European countries, 2% ETV6-RUNX1 frequency has been reported in Spanish pediatric ALL patients (Garcia-Sanz et al., 1999).All of these studies indicate great ethnic and geographical variations in frequency of this prognostically important genetic abnormality specifically and genetics of pediatric ALL generally, which can have a significant bearing on global pediatric ALL management strategies (Wesołowska-Andersen et al., 2015).

In conclusion, like most of the other developing countries of the region, Saudi Arabia pediatric ALL patients have lower frequencies of ETV6-RUNX1 fusion oncogene than the developed countries like US, Europe, Japan and Canada. Further multicenter studies are required to find out frequencies of other prognostically significant genetic abnormalities in Saudi pediatric ALL patients in order to better plan and manage this highly curable disease in the kingdom.

## References

Al-Ahmadi K, Al-Zahrani A (2013). Spatial autocorrelation of **7526** *Asian Pacific Journal of Cancer Prevention, Vol 16, 2015* 

cancer incidence in Saudi Arabia. *Int J Environ Res Public Health*, **16**, 7207-28.

- Al-Mulla NA, Chandra P, Khattab M, et al (2014). Childhood acute lymphoblastic leukemia in the Middle East and neighboring countries: a prospective multi-institutional international collaborative study (CALLME1) by the Middle East Childhood Cancer Alliance (MECCA). *Pediatr Blood Cancer*, **61**, 1403-10.
- Al-Mutlaq HM, Bawazir AA, Jradi H, et al (2015). A. Patterns of childhood cancer incidence in Saudi Arabia (1999-2008). *Asian Pac J Cancer Prev*, **16**, 431-5.
- Alghamdi IG, Hussain II, Alghamdi MS, et al (2014). The incidence of leukemia in Saudi Arabia. Descriptive epidemiological analysis of data from the Saudi Cancer Registry 2001-2008. Saudi Med J, 35, 674-83.
- Amor DJ, Algar EM, Slater HR, et al (1998). High frequency of t(12;21) in childhood acute lymphoblastic leukemia detected by RT-PCR. *Pathol*, **30**, 381-5.
- Awan T, Iqbal Z, Aleem A, et al (2012). Five most common prognostically important fusion oncogenes are detected in the majority of Pakistani pediatric acute lymphoblastic leukemia patients and are strongly associated with disease biology and treatment outcome. *Asian Pac J Cancer Prev*, 13, 5469-75.
- Bhojwani D, Yang JJ, Pui CH (2015). Biology of childhood acute lymphoblastic leukemia. *Pediatr Clin North Am*, 62, 47-60.
- Ceppi F, Cazzaniga G, Colombini A, et al (2015). Risk factors for relapse in childhood acute lymphoblastic leukemia: prediction and prevention. *Expert Rev Hematol*, **8**, 57-70.
- Chung HY, Kim KH, Jun KR, et al (2010). Prognostic Significance of TEL/AML1 Rearrangement and Its Additional Genetic Changes in Korean Childhood Precursor B-Acute Lymphoblastic Leukemia. *Korean J Lab Med*, **30**, 1-8.
- Cooper SL, Brown PA (2015). Treatment of pediatric acute lymphoblastic leukemia. *Pediatr Clin North Am*, 62, 61-73.
- Eguchi-Ishimae M, Eguchi M, Tanaka K, et al (1998). Fluorescence in situ hybridization analysis of 12;21 translocation in Japanese childhood acute lymphoblastic leukaemia. Jpn J Cancer Res, 89, 783-88.
- Faiz M, Qazi JI (2010). t(12:21) Is underrepresented in childhood B-lineage acute lymphoblastic leukemia in Punjab, Pakistan. J Pediatr Hematol Oncol, 32, 249-51.
- Garcia-Sanz R, Alaejos I, Orfao A, et al (1999). Low frequency of the TEL/AML1 fusion gene in acute lymphoblastic leukemia in Spain. Br J Haematol, 107, 667-9.
- Gill HK, Keoh TS, Dhaliwal JS, et al (2005). TEL-AML1 frequency in multi ethnic Malaysian pediatric acute lymphoblastic leukemia. *Cancer Genet Cytogenet*, **156**, 129-33.
- Harbott J, Viehmann S, Borkhardt A, et al (1997). Incidence of TEL/AML1 fusion gene analyzed consecutively in children with acute lymphoblastic leukemia in relapse. *Blood*, **90**, 4933-7.
- Inaba H, Greaves M, Mullighan CG (2013). Acute lymphoblastic leukaemia. *Lancet*, **381**, 1943-55.
- Iqbal Z (2014). Molecular genetic studies on 167 pediatric ALL patients from different areas of Pakistan confirm a low frequency of the favorable prognosis fusion oncogene TEL-AML1 (t 12; 21) in underdeveloped countries of the region. Asian Pac J Cancer Prev, 15, 3541-6.
- Iqbal Z, Iqbal M, Akhter T (2007). Frequency of BCR-ABL fusion oncogene in Pakistani childhood acute lymphoid leukemia (ALL) patients reflects ethnic differences in molecular genetics of ALL. J Pediatr Hematol Oncol, 29, 585.
- Kobayashi H, Satake N, Kaneko Y (1997). Detection of the Der

(21) t(12;21) chromosome forming the TEL-AML1 fusion gene in childhood acute lymphoblastic leukemia. *Leuk Lymphoma*, **28**, 43-50.

- Kwong YL, Wong KF (1997). Low frequency of TEL-AML1 in adult acute lymphoblastic leukemia. *Cancer Genet Cytoge*, 98, 137-8.
- Liang DC, Chou TB, Chen JS, et al (1996). High incidence of TEL/AML1 fusion resulting from a cryptic t(12;21) in childhood B-lineage acute lymphoblastic leukemia in Taiwan. *Leukemia*, **10**, 991-3.
- Mazloumi SH, Madhumathi DS, Appaji L, et al (2012). Combined study of cytogenetics and fluorescence in situ hybridization (FISH) analysis in childhood acute lymphoblastic leukemia (ALL) in a tertiary cancer centre in South India. *Asian Pac J Cancer Prev*, **13**, 3825-7.
- Moorman AV, Enshaei A, Schwab C, et al (2014). A novel integrated cytogenetic and genomic classification refines risk stratification in pediatric acute lymphoblastic leukemia. *Blood*, **124**, 1434-44.
- Pui CH, Pei D, Coustan-Smith E, et al (2015). Clinical utility of sequential minimal residual disease measurements in the context of risk-based therapy in childhood acute lymphoblastic leukaemia: a prospective study. *Lancet Oncol*, 16, 465-74.
- Rahman SA, Mohadess Ardabili SM, Aghazadeh A, et al (2006). Investigation of TEL-AML1 and BCR-ABL fusion oncogenes in patients affected by acute Lymphoblastic leukemia using interphase in situ hybridization. J Sc Islamic Rep Iran, 17, 17-5.
- Ribera JM, Oriol A (2009). Acute lymphoblastic leukemia in adolescents and young adults. *Hematol Oncol Clin North Am*, **23**, 1033-42.
- Saudi Arabian National Guard (2015). In Wikipedia, The Free Encyclopedia. Retrieved 07:30, May 20, 2015, from http://en.wikipedia.org/w/index.php?title=Saudi\_Arabian\_ National\_Guard&oldid=662239904, May 14.
- Shaker HM, Sidhom IA, El-Attar IA (2001). Frequency and clinical relevance of TEL-AML1 fusion gene in childhood acute lymphoblastic leukemia in Egypt. *J Egyptian Natl Cancer Inst*, **13**, 9-18.
- Shurtleff SA, Buijs A, Behm FG, et al (1995). TEL/AML1 fusion resulting from a cryptic t(12;21) is the most common genetic lesion in pediatric ALL and defines a subgroup of patients with an excellent prognosis. *Leukemia*, **9**, 1985-9.
- Tsang KS, Li CK, Chik KW, et al (2001). TEL/AML1 rearrangement and the prognostic significance in childhood acute lymphoblastic leukemia in Hong Kong. *Am J Hematol*, **68**, 91-8.
- Wesołowska-Andersen A, Borst L, Dalgaard MD, et al (2015). Genomic profiling of thousands of candidate polymorphisms predicts risk of relapse in 778 Danish and German childhood acute lymphoblastic leukemia patients. *Leukemia*, 29, 297-303.