Study on the Relativity between Cytogenetics and Cytomorphology and its Prognosis Significance in Children with Acute Myelogenous Leukemia

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ABSTRACT

Objective: The main objective of this study was to retrospectively evaluate that the cytogenetic abnormalities is an important prognostic factor for the cure of acute myeloid leukemia (AML).

Methods: This retrospective study enrolled newly diagnosed 70 cases (37 males and 33 females, aged 10.1 months to 14.5 years) of pediatric patients with AML during 2010 January - 2016 February from the Second Affiliated Hospital of Anhui Medical University. Excluding criteria were cases secondary to treatment-related MDS and AML. Samples were obtained from bone marrow cells in patients after treatment on the anterior superior iliac spine, blood diseases laboratory by direct culture or 24/48 hour short-term culture, G -banding technique for testing. Follow-up of 1 - 60 months, the analysis of treatment response rates of different karyotypes, distribution ratios in various subtypes, normal karyotype and abnormal karyotype.

ISPSS17.0 software statistics was used for statistical analysis. Groups were compared using chi-square test; Survival rate was calculated by method of Kaplan Meier and survival difference between groups were compared with breslow test.

Results: Among 70 cases, 42 cases were detected for chromosomal abnormalities (i.e. 60% of the total number of cases), M3 abnormal karyotype distortion rate of 78.5%, M2 abnormal karyotype aberrations 63.3%, M4 60.0%, M1 50%, M5 lowest 38.9 %, M7 nuclear aberrations highest rate was 100%. Total chromosomal aberration rate was 60%. Acute myeloid leukemia cases, t (8; 21) at most , there are 15 cases, and the presence of abnormal karyotype 86.7% in the original part of differentiated myeloid leukemia (M2); t (15; 17) has 11 cases, exists only in acute promyelocytic cell leukemia (M3). After treatment, the remission rate of t (8; 21) was 80%; the remission rate of t (15; 17) was 90%; the remission rate of other abnormal karyotype abnormalities was 50%; the remission rate of total abnormal karyotype was 71.4%. The event free survival rate was significantly different between normal karyotype, t (8; 21), t (15; 17) and other abnormal karyotype groups (P<0.05).

Conclusions: Acute myeloid leukemia karyotype abnormalities among FAB subtypes are different; M3 is the highest rate of abnormal

Keywords
Abnormal karyotypes,
Acute myeloid leukemia,
Cytogenetics, Prognosis.

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INTRODUCTION

Acute Leukemia belong to the group of hematopoietic stem cell mutation clone malignancy which is characterized by certain blood cells of hematopoietic tissue system hyperplasia, and infiltration to the bloodstream and then to various tissues and organs, which led to a series of clinical manifestations.

In China, pediatric malignancies have the highest incidence of leukemia. With the continuous improvement of chemotherapy, the rapid development of immunology, cytogenetic and molecular genetics, it is no longer considered a fatal disease. Complete remission rate of initial treatment in children with acute myeloid leukemia has reached 80% and five year disease-free survival rate of about 40 - 60%.

Over the past years, blood disease research work caught committed to explore the pathogenesis of acute leukemia from cytogenetic and molecular level that the considerable number of leukemia is chromosomal changes because of genetic abnormalities. Oncogene activation, tumor suppressor gene deletion, causing the proliferation of hematopoietic cells, differentiation and regulation disorders, to malignant transformation.

With the advancement of cultivation method and banding techniques, currently age-related macular degeneration (AMD) chromosomal aberrations detected rate has risen from 50% in the 1960s and 1970s and the current to around 80% - 90%. Although the type of AML chromosomal aberrations up to 100 or more, But it can be summarized into two categories: one is FAB subtype-specific chromosomal rearrangements related, about 60%, including the t (8; 21) (q22; q22), t (15; 17) (q 22; q22 or 21), inv (16) (p13; q22) etc.

Where in t (8; 21) (q22; q22) is closely related to AML-M2, t (15; 17) (q22; q12 or 21) found only in AML-M3, inv (16) (p13; q22) more common in M4EO. Such karyotype on early diagnosis and classification of AML has a guide. FAB subtype and the other is not related to abnormal, most of the number of abnormal, common chromosome number has increased first 8, 21, 19, 11 and 22 there appears karyotype aberrations, M2, M4 medium, M5 minimum. t (15; 17) seen in acute promyelocytic leukemia (APL), prognosis is good; t (8; 21) is more common in M2, prognosis is good, also found in M4 and M5, worse prognosis; +8 Abnormalities found in AML M2, M3, M4, M5 and M6 subtypes, prognosis medium; inv (16) high white blood cells, low platelet poor prognosis, AML patients with normal karyotype prognosis medium.

In this study, 70 cases of hospitalized patients with recently diagnosed AML from January 2009-2015 February were enrolled with karyotype abnormalities, patients with acute myeloid leukemia were involved in the treatment to understand AML patients karyotype distribution, survival follow-up and outcome combined with the clinical diagnosis and treatment. The aim is to analyze the correlation between karyotype and prognosis.

METHODS

2.1 Study

Newly diagnosed 70 cases (37 males and 33 females, aged 10.1 months to 14.5 years) of pediatric patients with AML were enrolled from 2009 January – 2016 February from the Second Affiliated Hospital of Anhui Medical University.
Excluding criteria were cases secondary to treatment related MDS and AML.

By FAB classification, M1 subtype 2 cases, M2 subtype 30 cases, M3 subtype 14 cases, M4 subtype 5 cases, M5 subtype 18 cases, and M7 subtype, one case. Diagnosis and evaluation of the efficacy of all patients were done with reference to "blood disease diagnosis and treatment standards".

2.2 Specimen collection

Patients admitted to hospital after chromosome analysis, samples were obtained from the bone marrow cells before and after treatment in patients from iliac spine. Five ml injection containing two ml broth, culture bottles were immediately taken for inspection.

2.3 Cytogenetic analysis

The samples were submitted to the Second Affiliated Hospital of Anhui Medical University, Laboratory of Hematology line karyotype. Direct method or short-term legal culture piece, karyotype analysis using G banding analysis of 20 metaphase cells, press "Human cytogenetic international nomenclature system (ISCN1995)" were to be carried out.

At least two cells having the same or increase in chromosome structural rearrangements, or three cells having the same chromosome loss, will be confirmed as an exception clone.

2.4 Treatment and efficacy evaluation

Patient with recently diagnosed M3 with all-trans retinoic acid (ATRA) 25mg / (m²•d) and arsenic trioxide (As2O3) 0.2 mg / (kg•d) during induction therapy period, 28 days of oral. In consolidate the treatment period, patient were treated with cytarabine (Ara-C) 100 mg / (m² •q12h) from one to seven days and DNR 40 mg / (m²•d) from one to three days, fowling with ATRA 25 mg / (m²•d) and As2O3 0.2 mg/ (kg• d) for 28 days, and the very next treatment with Ara-C 100 mg/ (m² •q12h) from one to seven days and DNR 40 mg/ (m²•d) from one to three days. After completing the consolidating the treatment, patient entered maintenance treatment with ATRA 25 mg/ (m²•d) for 28 days, Ara-C 75 mg/ (m² •q12h) for five days, harringtonine 3 mg/ (m²•d) for seven days, As2O3 0.2 mg/ (kg• d) for 14 days, Ara-C 75mg/ (m² •q12h) and 6-TG 75 mg/ (m²•d) for 7 days; five times totally. Efficacy criteria: Zhang Zhinan reference "blood disease diagnosis and treatment standards".

Complete remission (CR): it means that the leukemic cells without clinical presentation caused by infiltration, or near normal life values = 1.5 × 10^9/L, platelets = 100 × 10^9/L. Peripheral blood leukocyte no leukemia cells. Bone marrow blasts = 5%, red blood cells and megakaryocytes normal series. Partial remission (PR): Bone marrow myeloblast type I + type II (original ten immature monocytes or ten original immature lymphocytes) equals to >5% and = 20%, or clinically, a standard blood counts by the end of complete remission.

No remission (NR): None of the above criteria. Recurrence was defined by one of the following three criteria including (1) bone marrow blasts >5% and = 20%, after a course of an effective anti-leukemic treatment, bone marrow failed to achieve complete remission. (2) Bone marrow blasts >20%; (3) myeloid leukemia cell infiltration. Overall survival is defined as the time of diagnosis until death or end of follow-up. Disease-free survival is defined as the time to achieve complete remission to relapse during this time, if more than one relapse and remission as per the cumulative basis.

2.5 Experimental Method

2.5.1 Reagent

(1) Medium: containing 20% fetal calf serum and 20 µ/ml heparin RPMI l640.

(2) Mitosis blockers: colchicines, formulated 5 µg/ml were used.

(3) Hypotonic: 0.075 mol/L potassium chloride solution.

(4) Fixative: 1: 3 glacial acetic acid and methanol.

(5) Banding solution: 0.1% trypsin, 0.02% EDTA solution, pH 6.8 phosphate buffer solution, Tris solution 3%.

(6) Giemsa staining solution: phosphate buffer solution concentration of 10%, pH 6.8 ~ 7.4.

2.5.2 Step

(1) Take specimens and inoculation culture: Sterile needle to extract bone marrow fluid 0.5 ~ 2 mL in heparin tube, add broth to 8 mL, washing and percussion, centrifugation (1200 rpm/min, 5 min), repeated 2 - 3 times to wash away the fat and other impurities. Per ml medium 1 ~ 3 × 106 cells were inoculated to a sterile culture medium containing 5 ml in to two flasks, shake gently and placed in the box 37 for 48 hours.

(2) Colchicines treatment: Sample bottle was added with colchicines 50 µL, using liquid 0.05 ml (final concentration 0.05 µg/mL) to continue to foster 50 min.
(3) Preparation of samples:
1) After termination of the culture, the culture was poured into the conical centrifuge tube, 6 minutes centrifugation (1200 rev/min), the supernatant was discarded.
2) The prewarmed 8 ml hypotonic solutions added by repeatedly pipetting at 37°C with bubble method, water bath for 30 minutes.
3) PreFixed: Add fixative 1~5mL, percussion mix placed under room temperature 3~5 min.
4) Centrifugation: 800 to 1000 rev/min.
5) Fixed: The supernatant was discarded, freshly prepared fixative was added to 5~6 mL, mix by pipetting.
6) Centrifugation: 800 to 1000 rev/min, 10 minutes
7) Repeat 5 and 6 steps two times.
8) The supernatant was discarded, adding an appropriate amount of fixative gently pipetting the cell suspension to spare.
9) Drop sheet: Take number of slides that stored in clean ice cold wet sheets, slightly tilted, with a capillary dropper suction cell suspension, dropping from 10 cm height of one to two drops on the slide in the air to be dry.
10) Roasted pieces: slides placed in 70°C oven bake for two hours.

(4) G-banding:
1) The first, vat 25 mL (0.1%) mixed with trypsin 25 mL (0.02%) EDTA solution, Tris was adjusted to pH 7. Second, three vat 50 mL were added at pH 6.8 phosphate buffer. The fourth was added 50 mL (5%) Giemsa dye vat. The three dye vats were kept at 37°C water bath.
2) Slide into the first cylinder 20 - 30 seconds, remove successively rinsed in the second, three-cylinder, followed by the third set dye vat five to 10 minutes.
3) Tap water, dry equipment seized.
4) Each patient observed for at least 20 metaphase cells, abnormal karyotype press "Human Cytogenetic Nomenclature System (ISCN 1995)" are identified and described.

2.6 Statistical Methods

SPSS17.0 software statistics was used for statistical analysis. Groups were compared using chi-square test; Survival rate was calculated by method of Kaplan Meier and survival difference between groups were compared with breslow test.

RESULTS

Among 70 cases, 42 cases were detected for chromosomal abnormalities, i.e. 60% of the total number of cases.

Table 1: Karyotype distribution

<table>
<thead>
<tr>
<th>FAB subtype</th>
<th>Number of cases</th>
<th>Normal karyotype</th>
<th>Abnormal karyotype</th>
<th>Distortion rate (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>M1</td>
<td>2</td>
<td>1</td>
<td>1</td>
<td>50%</td>
</tr>
<tr>
<td>M2</td>
<td>30</td>
<td>11</td>
<td>19</td>
<td>63.3%</td>
</tr>
<tr>
<td>M3</td>
<td>14</td>
<td>3</td>
<td>11</td>
<td>78.5%</td>
</tr>
<tr>
<td>M4</td>
<td>5</td>
<td>2</td>
<td>3</td>
<td>60.0%</td>
</tr>
<tr>
<td>M5</td>
<td>18</td>
<td>11</td>
<td>7</td>
<td>38.9%</td>
</tr>
<tr>
<td>M7</td>
<td>1</td>
<td>0</td>
<td>1</td>
<td>100%</td>
</tr>
<tr>
<td><strong>Total</strong></td>
<td><strong>70</strong></td>
<td><strong>28</strong></td>
<td><strong>42</strong></td>
<td><strong>60%</strong></td>
</tr>
</tbody>
</table>

As can be seen from Table 1, all 70 patients with acute myeloid leukemia, M1 subtype one case of abnormal karyotype; M2 subtype 19 cases of abnormal karyotypes, M3 subtype 11 cases of abnormal karyotypes; M4 subtype three cases of abnormal karyotypes, M5 subtype seven cases of abnormal karyotypes; M7 subtypes one case of abnormal karyotype. M3 abnormal karyotype distortion rate of 78.5%, M2 abnormal karyotype aberrations 63.3%, M4 60.0%, M1 50%, M5 lowest, 38.9%, M7 nuclear aberrations highest rate was 100%. However, due to a number of cases investigated test is too small, it is no clear statistical significance between this groups. So this experiment shows M3 subtype highest distortion, total chromosomal aberration rate 60%.

Table 2: Abnormal karyotype in AML

<table>
<thead>
<tr>
<th>Abnormal karyotype</th>
<th>FAB Subtype</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>M1</td>
<td>M2</td>
</tr>
<tr>
<td>t (8;21)</td>
<td>13</td>
<td>2</td>
</tr>
<tr>
<td>t (5;17)</td>
<td>11</td>
<td>11</td>
</tr>
<tr>
<td>Inv (16) (13; q22)</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>8q-</td>
<td>2</td>
<td>2</td>
</tr>
<tr>
<td>11p+</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>11q-</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>+8</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>+12</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>-7</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>-19</td>
<td>1</td>
<td>1</td>
</tr>
</tbody>
</table>
Table 2 shows that, in acute myeloid leukemia patients, there are 10 kinds of abnormal karyotype and complex chromosomal aberrations. t (8; 21) at most, 15 cases, and the presence of abnormal karyotypes was 86.7% in the original part of differentiated myeloid leukemia (M2); t (5; 17) 11 cases, exists only in acute promyelocytic cell leukemia (M3), inv (16) (p13; q22) in two cases, respectively, in the presence of the M4 and M5, 8q- two cases exist in M2, 11p +1 case and 11q-, are present in M2, the +8 and +12 in one case, all exist in M2, the -19 one case, exist in M2, four patients hyperdiploidy were present in M2 and M5, and the remaining two cases were complicated karyotype, they were present in the M5 and M7.

Table 3: Abnormal karyotype and efficacy in AML

<table>
<thead>
<tr>
<th>Group</th>
<th>CR+PR</th>
<th>NR</th>
<th>Total</th>
<th>Response Rate (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>t (8; 21)</td>
<td>12</td>
<td>3</td>
<td>15</td>
<td>80</td>
</tr>
<tr>
<td>t (5; 17)</td>
<td>10</td>
<td>1</td>
<td>11</td>
<td>90.1</td>
</tr>
<tr>
<td>Other abnormal karyotype</td>
<td>8</td>
<td>8</td>
<td>16</td>
<td>50</td>
</tr>
<tr>
<td>Total</td>
<td>30</td>
<td>12</td>
<td>42</td>
<td>71.4</td>
</tr>
</tbody>
</table>

Fig 1: The event free survival curves of different karyotype in AML

From table 2, table 3 and fig 1 can be seen, M2 subtype in 13 patients with abnormal karyotype t (8; 21), of which 12 cases of remission after treatment, two cases are not alleviated after treatment, remission rate 80%. M3 subtype in 11 patients with abnormal karyotype t (15; 17), 10 patients were in complete remission after treatment, one case was not alleviated, and remission rate was 90.1%. Other abnormal karyotypes total of 16 cases, 50% response rate. There are two cases which inv (16) (p13; q22), one case remission, and one died, two cases of 8 q- were relieved, one case of 11p +, follow-up death, one case of 11q- was remission. One case of +8 was died. One case of +12 was remission. One case of -7 patients was followed up in death. One case of -19 was remission. The event free survival rate was significantly different between normal karyotype, t (8; 21), t (15; 17) and other abnormal karyotype groups (P=0.028 <0.05). Further statistical analysis found that event free survival rate free survival rate between normal karyotype and other abnormal karyotype groups was statistically difference (P=0.004 <0.05), no difference between the rest of the groups.

DISCUSSION

With the continuous improvement of chromosome banding techniques, especially the rapid development of high-resolution technology, such chromosomal abnormalities relationship between certain types of leukemia and has become more and more closely, but also identified a number of subtypes iconic chromosome abnormal characteristics. Meanwhile, with disease progression or improvement of chromosomal abnormality occurs constantly changing, these changes we adjust has a significant role in the treatment and prognosis of the disease. In most patients, the application of cytogenetic methods can be found in clear cell clonal chromosomal abnormalities. The same chromosomal abnormalities are also seen in acute leukemia (AL) of each subtype, which means not only the diagnosis, but also prognostic value, the clinical characteristics of patients; help us to predict the patient’s condition, because the merits of any karyotype changes are not absolute. Chromosomal abnormalities are important prognostic factor, such as t (8; 21) and t (15; 17) or inv (16) of better prognosis, and -5, -7, t (9; 22), and complex chromosomal abnormalities have poor prognosis.

The group of 70 children with acute myeloid leukemia, chromosome aberrations was 60%, suggesting that cytogenetic abnormalities are an important factor in the occurrence of acute myeloid leukemia. Highest karyotype aberration M3, 78.5%, M5 lowest, 38.9%, between the groups, not statistically significant. Considering the small sample volume chromosome aberration rate difference.
between the AML subtypes may vary.

4.1 Acute promyelocytic leukemia (APL) and t (15; 17)

This group included 14 patients with 11 cases of abnormal karyotype APL, where, t (15; 17) 11 cases, accounting for 100%. Alan k Burnett5 and other studies of 218 patients with APL, 187 cases of t (15; 17), accounting for 85.8%, there was no difference between the present studies. Kühnl A and Grimwade D6 studied 1612 cases of AML patients, 932 cases had abnormal karyotype, t (15; 17) 198 cases, accounting for 21.2% response rate was 87%. The patients with t (15; 17) of the total 70 cases of abnormal karyotype AML 15.6% response rate was 100%, more consistent both with David Grimwade and other reports. 10 cases of t (15; 17) have been applied, all-trans retinoic acid treatment, 10 cases of remission, remission rate was 100%. M3 has t (15; 17) translocation PML (promyelocytic leukemia gene) on chromosome 15 and 17, retinoic acid receptor gene (RARα) chromosome formation PML-RARα fusion gene, which is M3 incidence and application of all-trans retinoic acid treatment effective molecular basis. t (15; 17) prognosis is good, has its foundation in molecular biology6-7.

4.2 Acute myeloid leukemia part differentiation type (M2) and t (8; 21)

The patients of inv (16) (p13; q22) there are two cases, one case exists in M4, achieve remission, one case of M5, did not achieve remission. The patients were males, aged four, WBC 56.2 × 10^9/L, platelets 12 × 10^9/L; there are many adverse prognostic factors age, low platelets, and its poor prognosis. Eghtedar A, et al15 summarized 110 cases of inv (16) patients, CR 93%, high white blood cell (>120 × 10^9/L) and low platelets (<30 × 10^9/L) difficult to achieve CR. 65.3%, the prognosis is moderate.

4.3 Clinical relationship of karyotypes inv (16) (p13; q22)

The patients of inv (16) (p13; q22) there are two cases, one case exists in M4, achieve remission, one case of M5, did not achieve remission. The patients were males, aged four, WBC 56.2 × 10^9/L, platelets 12 × 10^9/L; there are many adverse prognostic factors age, low platelets, and its poor prognosis. Eghtedar A, et al15 summarized 110 cases of inv (16) patients, CR 93%, high white blood cell (>120 × 10^9/L) and low platelets (<30 × 10^9 / L) difficult to achieve CR. 65.3%, the prognosis is moderate.

4.4 Clinical relationship of normal karyotype

This group of patients with normal karyotype had 28 patients, seven cases of NR, response rate was 75%, means for survival time is 78.5 ±7.5 month. Torstein Haferlach16 and other reports analyzed 453 cases of newly diagnosed AML patients. It showed normal karyotype and abnormal +8, prognosis medium.

CONCLUSIONS

This article summarizes 70 cases of children with acute myeloid leukemia untreated cases, the application of short-term culture and/ or direct method to do G-banding karyotype analysis, all patients of acute myeloid leukemia were given standard regimen and observed cytogenetic karyotype relations and efficacy, the following conclusions occur:

(1) Acute myeloid leukemia cell has its genetic basis, diagnosis of chromosomal studies in acute myeloid leukemia are important in treatment and prognosis.
(2) Acute myeloid leukemia karyotype abnormalities among FAB subtypes are different; M3 is the highest rate of abnormal karyotype aberrations, M2, M4 medium, M5 minimum.

(3) $t{(15;17)}$ seen in acute promyelocytic leukemia (APL), prognosis is good.

(4) $t{(8;21)}$ is more common in M2, prognosis is good, also found in M4 and M5, worse prognosis;

(5) +8 Abnormalities found in AML M2, M3, M4, M5, M6 subtypes, prognosis medium;

(6) inv (16) high white blood cells, low platelet poor prognosis;

(7) AML patients with normal karyotype prognosis are better than other groups.

This study belongs to the single center study with small sample, only one case of some subtypes, so, we will join with other centers to expand the samples and further improve this study.

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