

**Application of a chemiluminescent methodology for detection of minimal residual disease in childhood acute lymphoblastic leukemia**

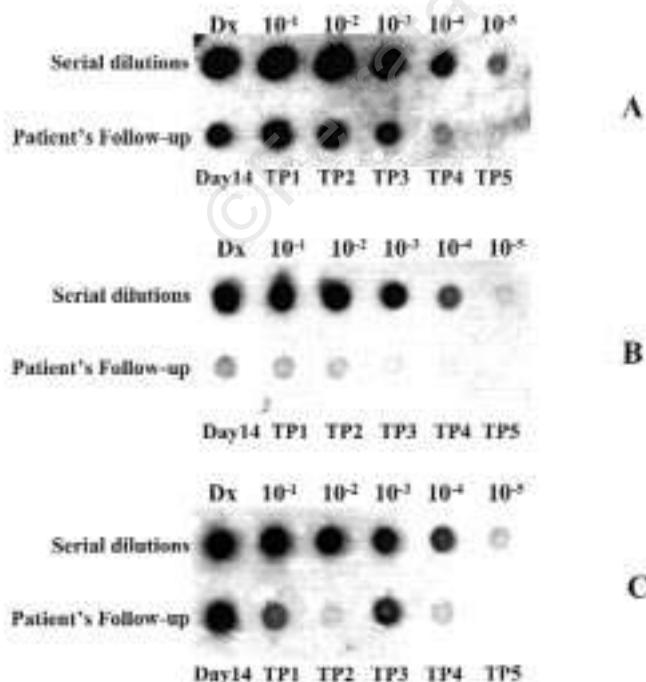
Analysis of minimal residual disease (MRD) can predict outcome in childhood acute lymphoblastic leukemia (ALL). We applied a chemiluminescent methodology in 20 children with ALL. We detected MRD at different time-points throughout the follow-up of our patients, concluding that chemiluminescent detection of MRD is a reliable, safe and sensitive method.

Recent prospective studies clearly demonstrate the prognostic value of MRD in children with ALL.<sup>1,2</sup> Several methodologies are available for MRD analyses in ALL patients.<sup>1-5</sup> In order to test a safe, sensitive and reliable method for those laboratories in which neither radioactive analysis nor the TaqMan strategy can be carried out,<sup>3</sup> we present here a report on application of digoxigenin (DIG)-labeled patient-specific probes for a chemiluminescent detection of MRD in children with ALL.

Twenty children with ALL diagnosed at our institution and treated according to the ongoing protocol of the Associazione Italiana di Ematologia ed Oncologia Pediatrica (AIEOP-ALL 95) were included in this study. The children's characteristics are listed in Table 1. We collected samples at 5 time-points (TP): after 43 days (TP1), after three months (TP2), five (TP3), seven (TP4) and 24 months (TP5) of therapy. T-cell receptor (TCR)  $\gamma$  and TCR  $\delta$  gene rearrangements were identified and characterized by performing diagnostic polymerase chain reactions (PCR), heteroduplex and sequencing analyses using standardized techniques.<sup>6,7</sup> PCR of follow-up were performed using different protocols in order to amplify TCR $\gamma$  and TCR $\delta$  rearrangements.<sup>7</sup> Sev-

en microliters of the products were spotted onto positively charged nylon membranes (NYTRAN N+, Roche Boehringer). Next, 200 pmol of each oligonucleotide were 3'-end labeled with DIG-ddUTP using a DIG oligonucleotide 3'-end labeling kit (Roche Molecular Biochemicals, Mannheim, Germany) according to manufacturer's instructions. The membranes were prehybridized at 68°C for 2 hours using 25 mL hybridization solution (Roche Molecular Biochemicals) and then hybridized in a sealed plastic bag with 3 mL hybridization solution and 20  $\mu$ L of the labeling probe (200 pmol). The membranes were incubated overnight at 54°C and then washed twice in 2  $\times$  SSC, 0.1% SDS at room temperature for 5 min. The membranes were soaked with blocking solution (Roche Molecular Biochemicals) for 30 min at room temperature. Twenty-four milliliters of blocking solution containing 2.4  $\mu$ L of anti-DIG-alkaline phosphatase Fab fragments (Roche Molecular Biochemicals) were added, followed by incubation for 30 min. Finally, the membranes were washed twice with washing buffer (Roche Molecular Biochemicals) for 15 min and incubated with detection buffer (Roche Molecular Biochemicals) for 5 min. A chemiluminescent substrate (CSPD, Roche Molecular Biochemicals) was added to the surface of the membrane. The hybridization bags were sealed and incubated at room temperature for 5 min in the dark. Finally the membranes were air-dried, placed in a plastic hybridization bag and incubated at 37°C for 15 min. We exposed each membrane to an X-OMAT AR film (Kodak) for a period ranging from 15 min to 4 hours, in order to increase signal detection.

The sensitivities for respective PCR targets using the chemiluminescent method are summarized in Table 1. Among our cases we designed probes which reached different sensitivities ranging from 10<sup>-3</sup> to 10<sup>-6</sup>. Figure 1 shows representative results regarding three paradigmatic situations of MRD analysis which might influence clinical decisions. Figure 1A shows persistence of a leukemic clone in a child with T-cell acute lymphoblastic



**Figure 1.** Chemiluminescent dot-blot hybridization of follow-up samples for cases LG (1A), SR (1B) and FD (1C), respectively; the patient-specific probes of all three cases resulted in a sensitivity of 10<sup>-4</sup>, as shown in serial dilutions prepared with leukemic DNA into normal healthy donor peripheral blood DNA. Minimal residual disease was detected at different time-points in the three cases: a persistence of leukemic clone was noted in LG (1A) until the patient underwent allogeneic bone marrow transplantation; there was a progressive eradication in patient SR (1B), as well as in the majority of the other cases; the clone re-emerged at the beginning of the re-induction phase (TP3) in case FD (1C).

**Table 1. Characteristics of children with ALL regarding rearrangements used for the chemiluminescent methodology.**

Pts	Sex/Age at Dx	Immuno-phenotype	Rearrangement at Dx	Junctional region	Patient-Specific Probes (5'-3')	Probe Sensitivity
AG	M/3ys	Common	Vδ2/Dδ3	-3/6/0	ctgtgacTCCCGTactgggg	10 <sup>3</sup>
AD	M/4ys	Common	Vδ2/Dδ3	-3/4/-2	cctgtgactTCTTTgggggata	10 <sup>3</sup>
CA	M/7ys	Common	Vδ2/Dδ3	-5/11/0	tgccctgtTTCACAGGGGAactg	10 <sup>4</sup>
FD	F/3ys	Pre-B	Vδ2/Dδ3	-1/del 80bp/Hept	tgtgacacTCTCTacagtgtctac	10 <sup>4</sup>
GA	F/11ys	Common	Vδ2/Dδ3	0/3/0	ctgtgacaccTTTactggggg	10 <sup>3</sup>
MM	F/4ys	Pre-B	Vδ2/Dδ3	-8/6/0	tactgtgcctCTTTCactgggg	10 <sup>4</sup>
SS	M/3ys	Common	Vδ2/Dδ3	-5/3/-2	tgtgcctgtTTTgggggat	10 <sup>3</sup>
CS	F/10ys	Common	Vδ2/Dδ3	-3/del 100bp	tgtgcctgtacCCTTCAGGCCCTGT	10 <sup>4</sup>
CM	M/4ys	Common	Vδ2/Dδ3	-5/4/0	tgtgcctgtGCTactGGGGG	10 <sup>4</sup>
FL	F/2ys	Common	Vγ2	-4/7/-1	cctgggaCAGCCGAaattatt	10 <sup>4</sup>
SR	F/3ys	Common	Vδ2/δd3	0/4/-3	cgtgacaccCCGAggggaata	10 <sup>4</sup>
SA	M/2ys	Common	Vδ2/δd3	-1/6/0	cctgtgacacGGCGTactggg	10 <sup>4</sup>
MG	M/12ys	T-ALL	Vγ4	-2/18/-6	ggatgCCCCGAAGTAGATTATTctt	10 <sup>4</sup>
BS	M/7ys	T-ALL	Vγ4	-4/8/-6	ccaccgggaTCTTCTGgtataaga	10 <sup>4</sup>
DSE	F/13ys	T-ALL	Vδ1/Jδ1	-2/25*/0	tttGAgggatCCCACCCATacac	10 <sup>6</sup>
LG	M/2ys	T-ALL	Vδ1/Jδ1	-1/37^/-1	CCGCTcctcctGGAAtggg	10 <sup>4</sup>
SE	F/7ys	T-ALL	Vδ1/Jδ1	-4/14*/0	gggTcttTAGgggggTacc	10 <sup>4</sup>
SR	F/3ys	T-ALL	Vδ1/Jδ1	-6/44*/-5	CCAagtGGTGGGAAACACGT	10 <sup>4</sup>
TD	M/3ys	T-ALL	Vδ1/Jδ1	-2/18*/0	gggTAGGTATTATacac	10 <sup>4</sup>
LG	M/13ys	T-ALL	Vδ2/δd3	-3/13/0	gacACCGGAGAAATGact	10 <sup>4</sup>

Pts: patents; Dx: diagnosis; ys: years; del: deletion; bp: base pair; hept: heptamer; \*D<sub>1</sub>/D<sub>2</sub>/D<sub>3</sub> gene segments are included; ^:D<sub>2</sub>/D<sub>3</sub> gene segments are included; capital letters represent N-region inserted nucleotides.

leukemia (case LG) who subsequently received an allogeneic bone marrow transplantation. Figure 1B (case SR) demonstrates evident progressive decrement of the chemiluminescent signal indicating eradication of the leukemic clone, whereas Figure 1C (case FD) shows how reappearance of a leukemic clone at the beginning of the reinduction phase (TP3) has not been followed by a subsequent relapse (TP4).

There are several advantages in using the chemiluminescent method for detection of MRD.<sup>5</sup> Firstly, the procedure is not radioactive and can, therefore, be performed in any laboratory where radioactive products are not allowed or the TaqMan strategy is too expensive or not yet applied. Secondly, because of the short exposure time (2-10 min), the whole procedure can be shortened in the future. However, end labeling of the oligonucleotide can be done at any time and the labeled probes can be stored at -20°C and used up to four times. In addition, the hybridization solution of DIG-labeled probes can be reused several times in order to analyze all time-points. We decided to use TCRδ and TCRγ gene junctional regions as patient-specific targets, because these rearrangements represent stable markers of disease throughout the duration of treatment and between diagnosis and relapse, as previously demonstrated.<sup>8,9</sup> Our data confirm that the chemiluminescent methodology can be performed in any laboratory and provides information important for clinical and therapeutic decisions. This procedure is faster and safer than the radioactive dot-blot technique, reducing biological hazards for the operator.

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