

**Evaluation de la maladie
résiduelle dans les
lymphomes non hodgkiniens**

C Tarella, Torino

**Molecular
techniques
for monitoring
MRD**

Target

Sensitivity

Southern Blot

- **IgH rearrangements**
- **Translocations**

10^{-2}

FISH

- **Translocations**
- **deletions**
- **karyotypic alterations**

5×10^{-2}

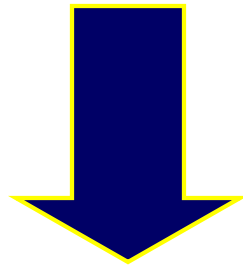
PCR

- **IgH rearrangements**
- **Translocations**

$10^{-4}-10^{-6}$

- ❖ the Molecular Remission (**MR**) achievement is a relevant issue in the management of non-Hodgkin's lymphoma presenting with BM involvement (BM+), in particular follicular lymphoma (**FCL**), mantle-cell lymphoma (**MCL**) and in selected cases of chronic lymphocytic leukemia (**CLL**)
- ❖ a **MR** achievement should always be associated with a Clinical Complete Response
- ❖ the need of a **MR** achievement is strictly related to:
 - disease prognostic presentation
 - type of treatment employed (i.e.: intensive vs. conventional therapeutic approach)

**you no longer need to be a
transplanter to achieve molecular
remission in FCL patients**



**since the use of more effective schedules (i.e.
CHOP, FND) as well as the introduction of
RITUXIMAB, FCL patients might have some
chances of achieving a Molecular Remission
with no need for intensified treatments**

Clinical and molecular response assessed in PB in 194 patients with FL at different time points from the beginning of treatment*

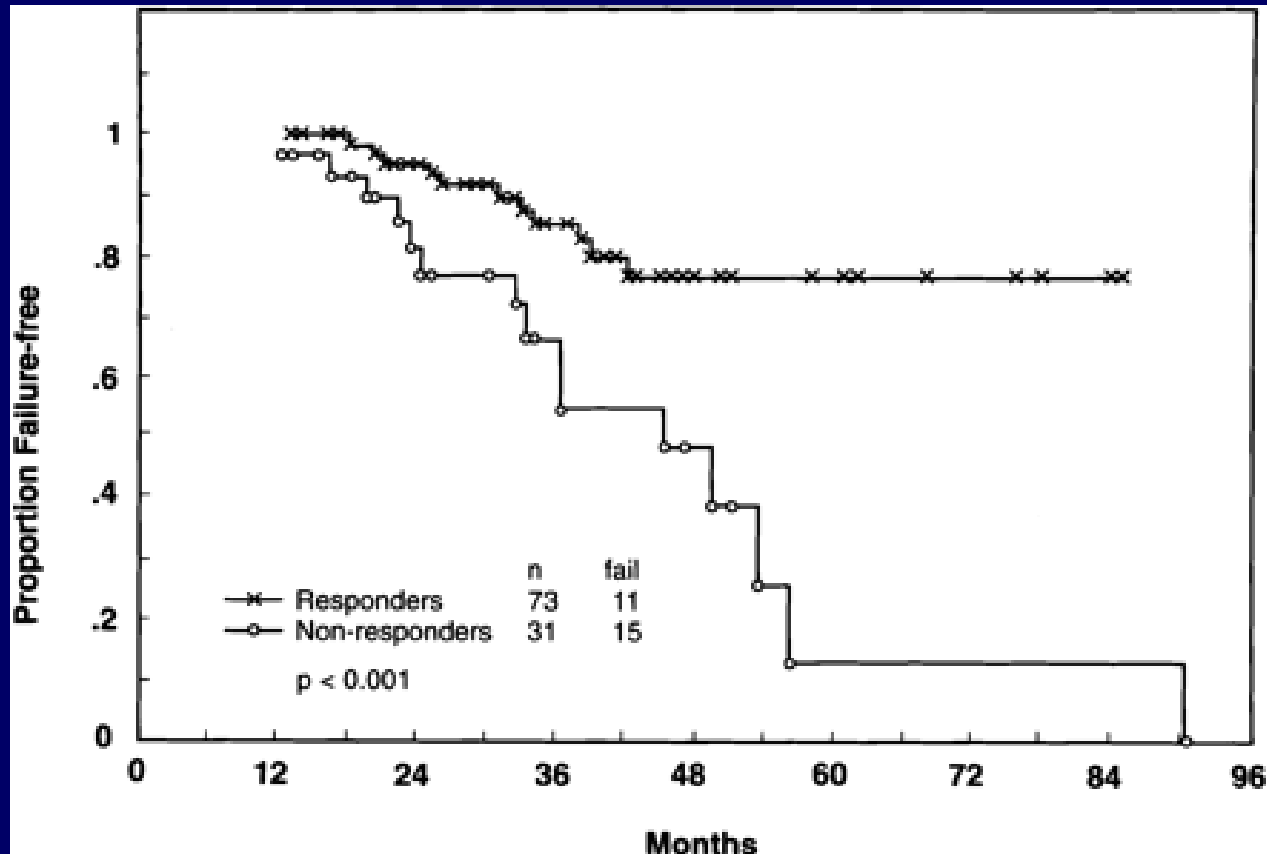
Time point months	No of patients	Clinical CR rate (%)	Molecular (Response rate %)
3-5	118	29	37
6-8	86	63	53
9-14	101	86	56
15-19	74	97	66

*** Therapy was ATT in 87 cases, FND in 24 and CHOP in 12**

Lopez-Guillermo, Blood 91, 2955; 1998

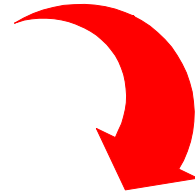
The Clinical Significance of Molecular Response in Indolent Follicular Lymphomas

By López-Guillermo et al.: Blood, Vol. 91: 2955-2960; 1998



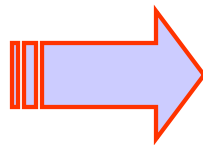
FFS from 1 year after the start of treatment according to the molecular response status within the 1st year (Responders: PCR-negative; Nonresponders: PCR-positive status)

**A Rambaldi et al,
Blood 2002, 99: 856-62**



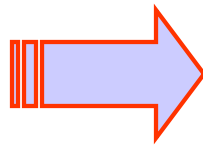
**Molecular monitoring after
sequential CHOP (6 cycles)
and Rituximab administration as
front line treatment in 128 FCL patients**

after CHOP



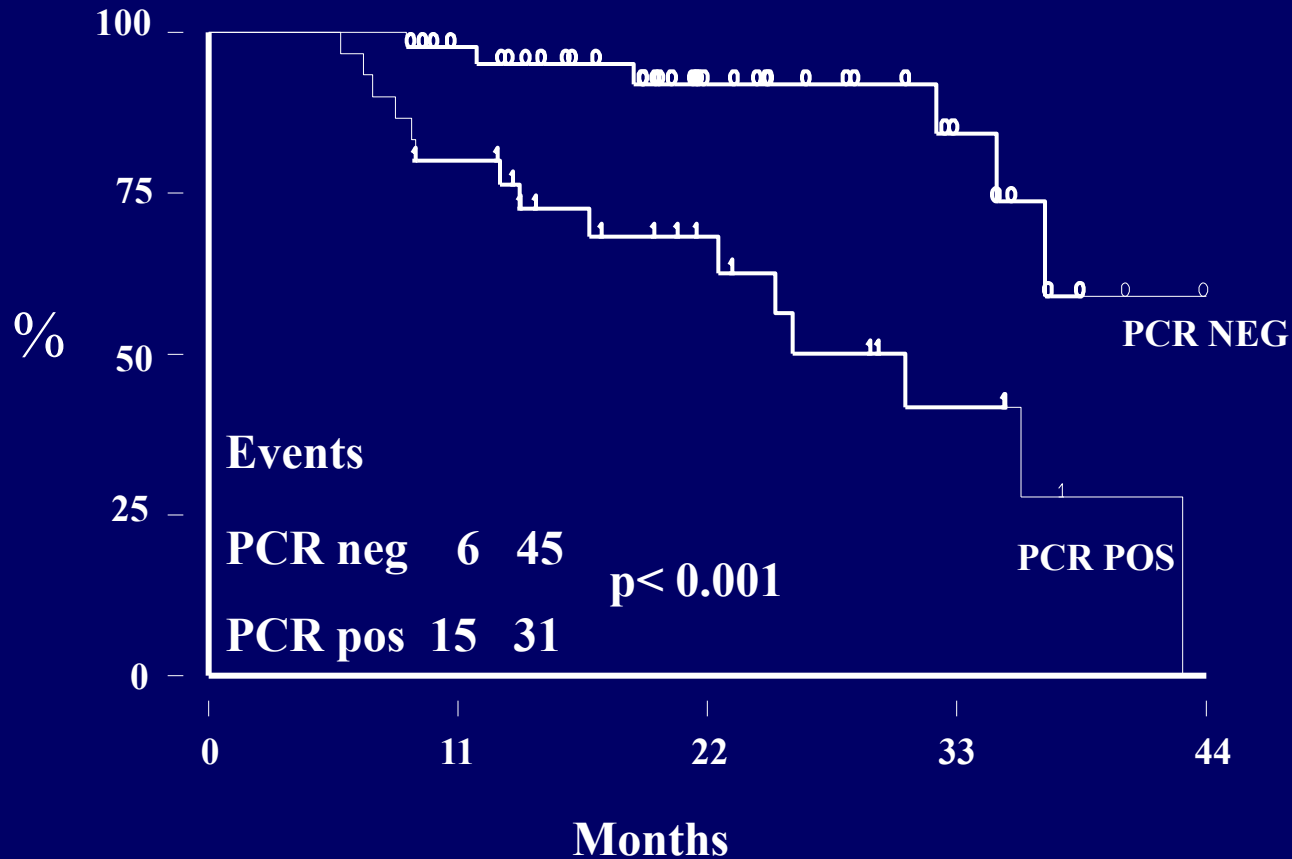
36% BM PCR neg

after Rituximab



74% BM PCR neg

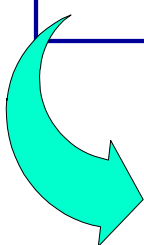
Freedom from recurrence after CHOP-Rituximab according to the PCR status in the BM at week +44



❖ several prospective studies have shown that the **MR** achievement following conventional chemo-immunotherapy (= negative PCR in at least 2 consecutive evaluations, at 6-12 mos. After treatment completion) is a favorable prognostic factor, associated with a prolonged failure-free survival (FFS)

❖ however:

- there is no demonstration of clear advantages in terms of overall survival
- quite a few patients have disease recurrence following MR achievement



At present, **MR** does not appear to be an essential “target” in the management of FCL patients when treated with conventional chemo-immunotherapy

Management of indolent non-Hodgkin's nodal lymphomas

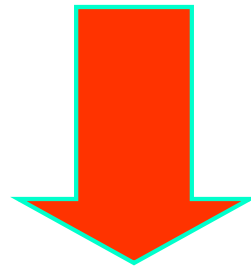
Practice guidelines by SIE, SIES and GITMO

Prof. S. Tura coordinator

❖ ***“ Molecular Response should be checked at the end of first-line therapy in all the patients with an informative probe and a complete clinical remission ”***

❖ ***“ Patients who have achieved a Partial Remission after first-line therapy should be considered for consolidation treatment with one of the following options: Rituximab, autologous SCT, radiotherapy, radioimmunoconjugates (either tositumomab or ibritumomab) “***

FCL e MCL managed with an intensive approach including BM o PBPC autograft



Evidence that achievement of both Clinical and Molecular Remission have a relevant impact on the long-term outcome

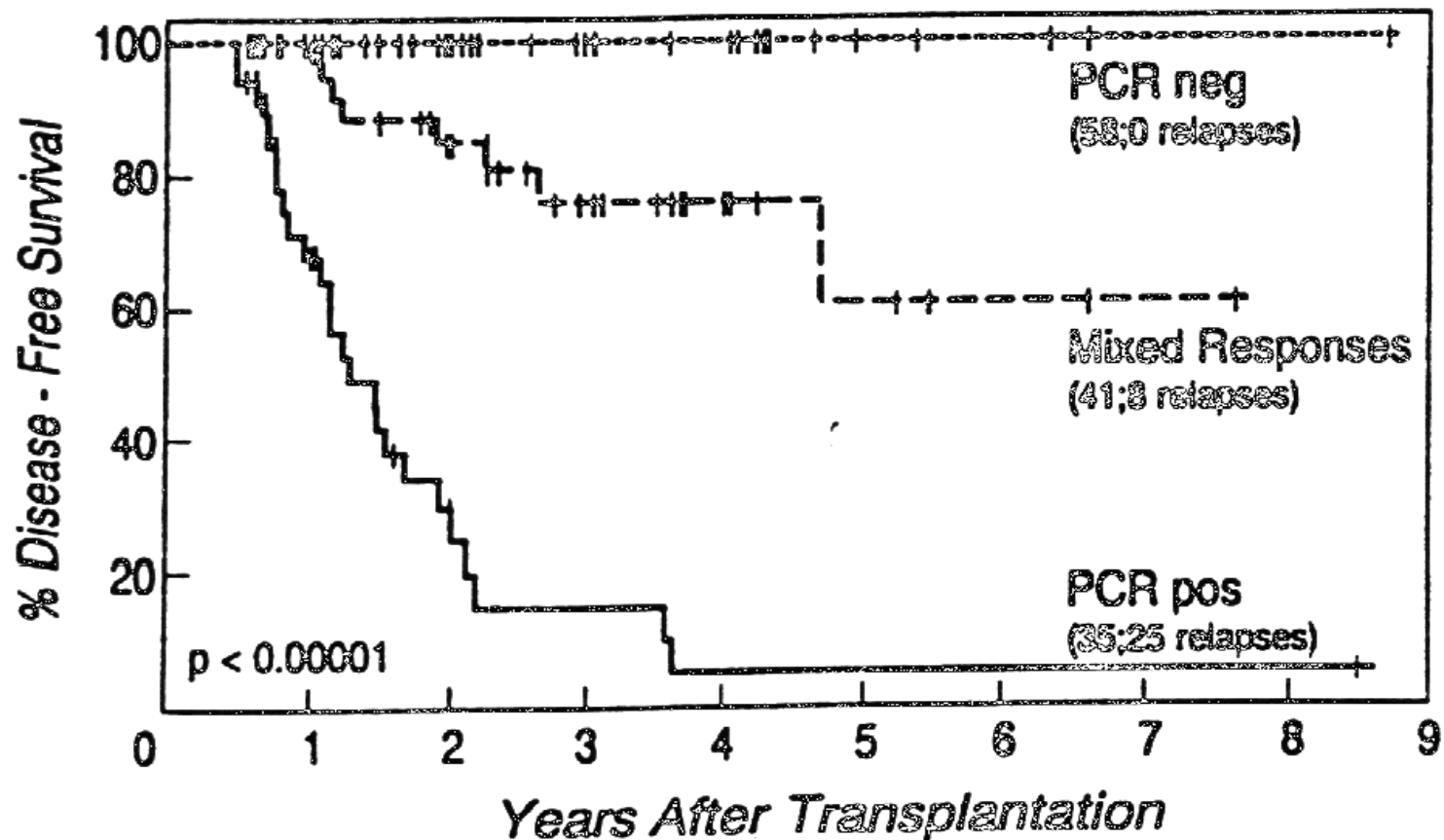
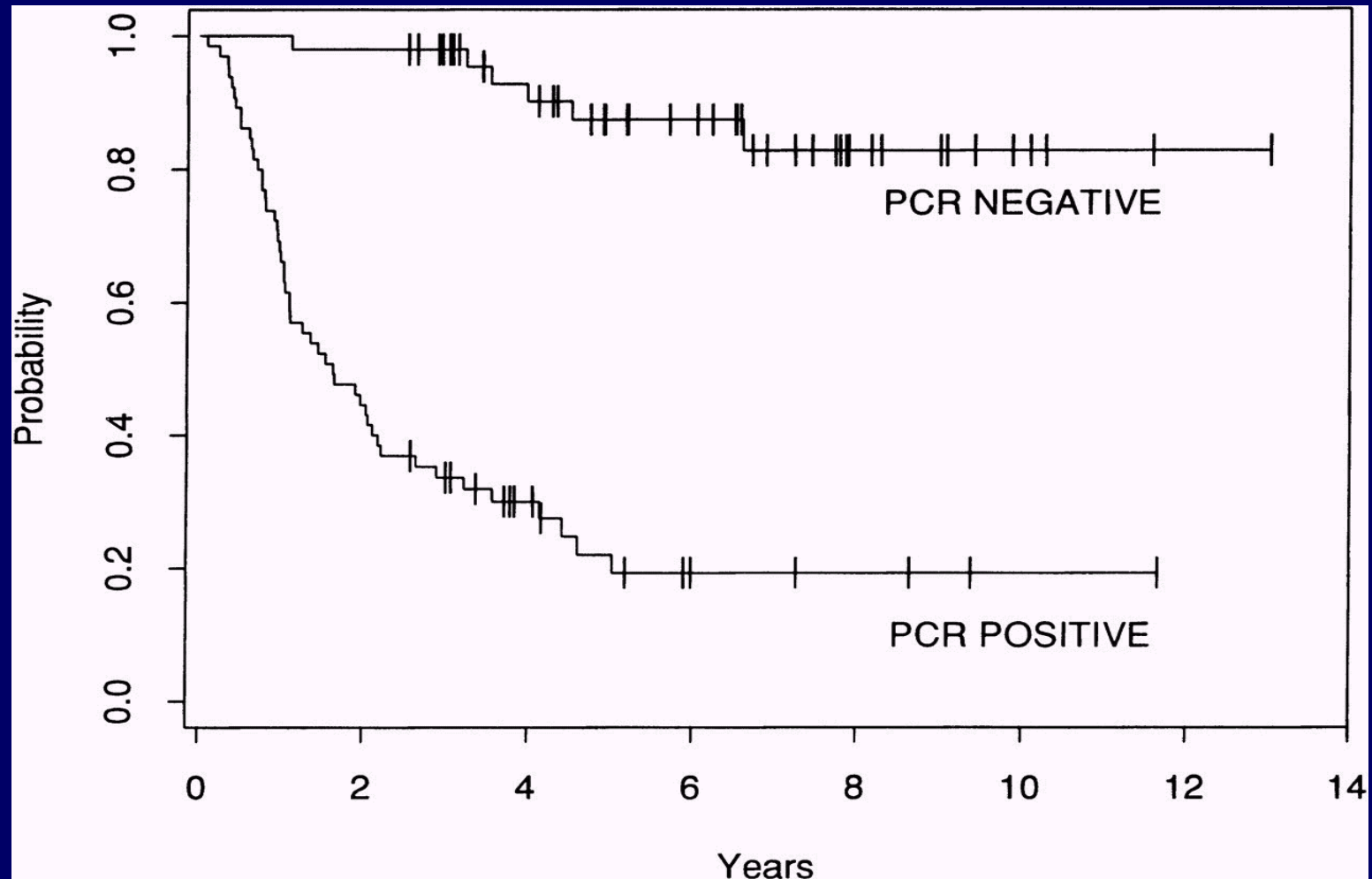


Fig 1. Actuarial probability of disease-free survival after ABMT in 134 patients with B-cell non-Hodgkin's lymphoma.

J.G. Gribben et al Blood 1993

Disease-free survival according to PCR status in FCL patients



Dana Farber study in MCL

Blood 90:4212, 1997

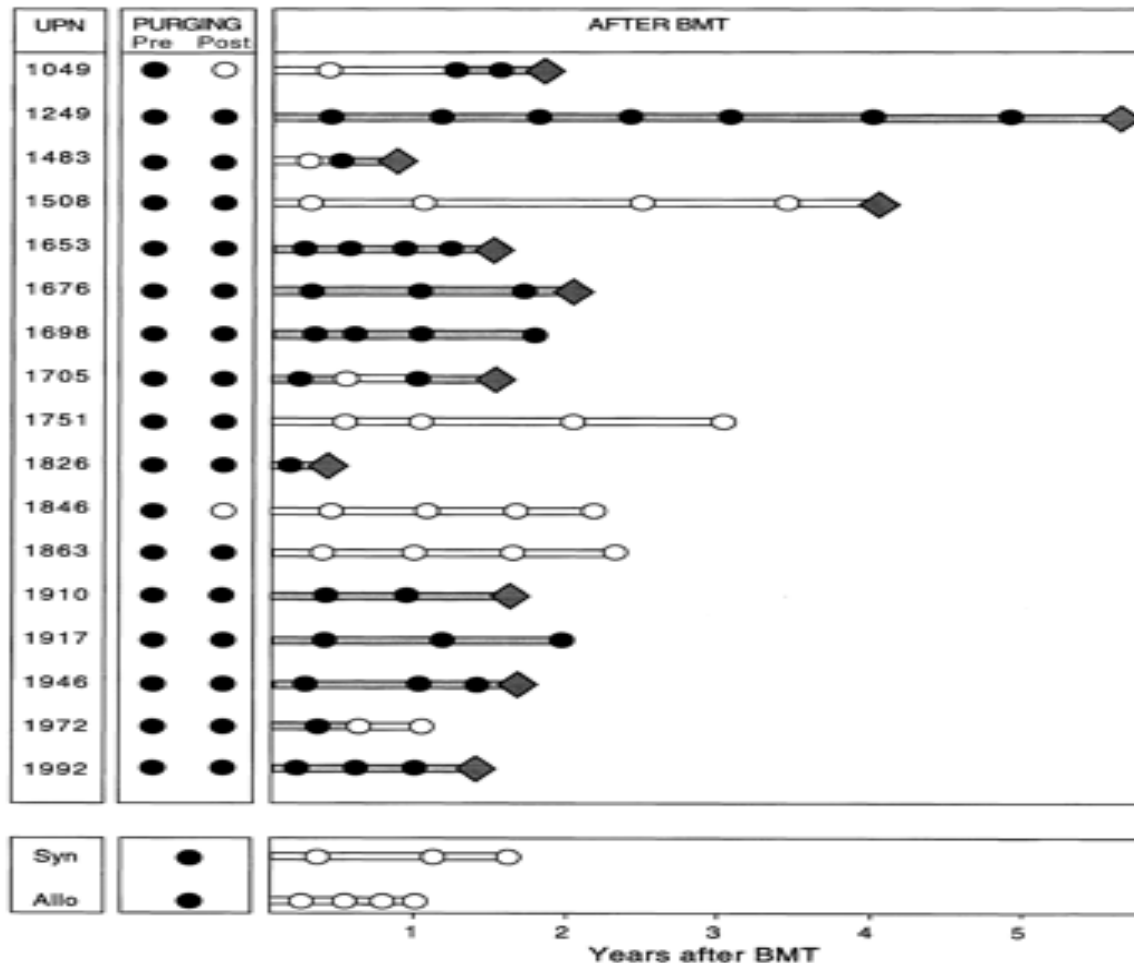
- * complement-mediated in vitro purging of BM cells

- * MRD assessed by PCR on BM harvest post-purging
MRD + in 17/19 pts.

(---> in follicular NHL, post-purging MRD negative in 50% of pts.)

- * more than half of the mantle-cell lymphoma pts.
relapsed within 2 yrs.following autograft

Relapse incidence and PCR status in MCL patients

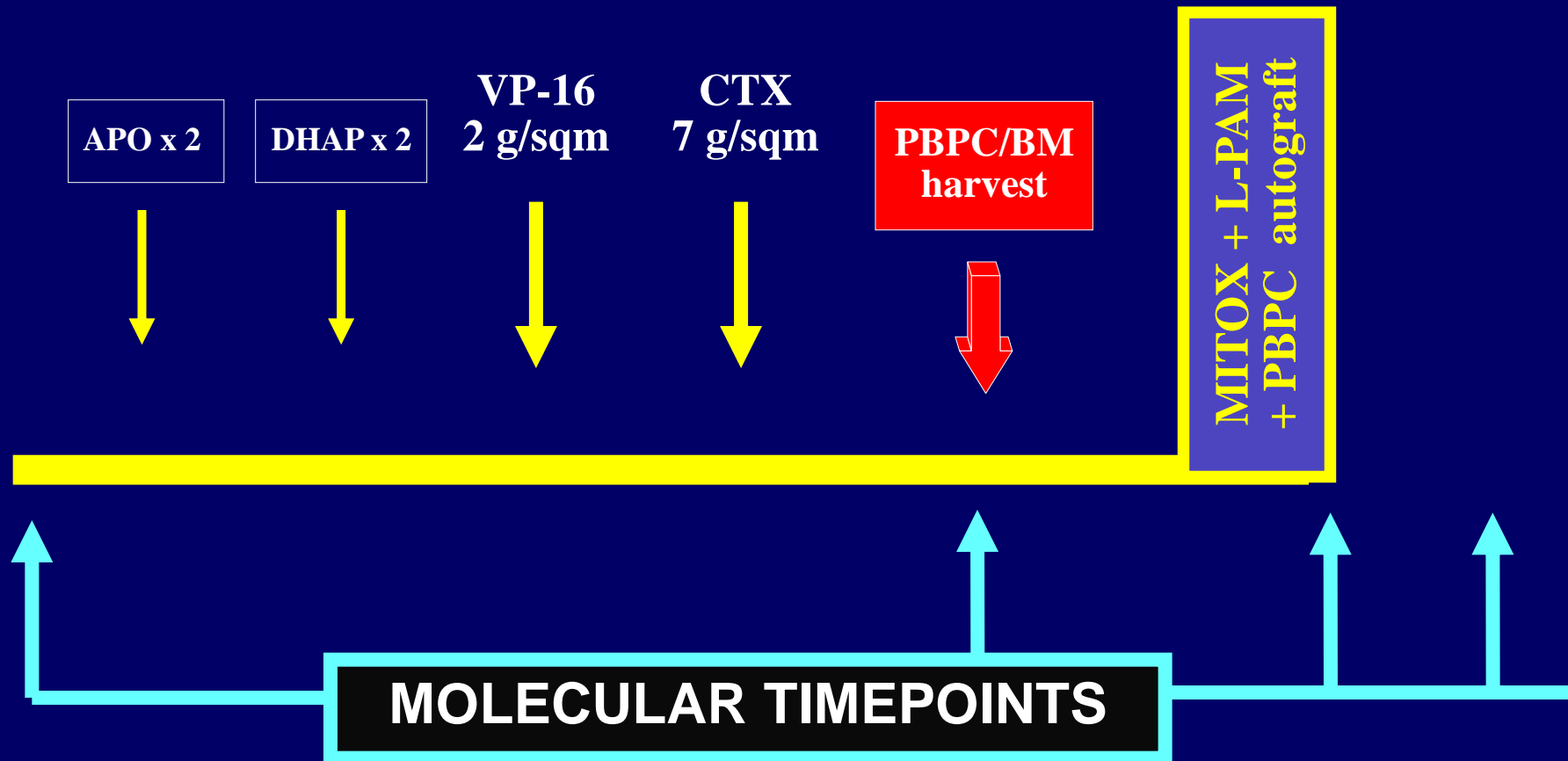


Andersen et al, Blood, 1997: 90,4212-4221

Turin experience

- **when:** since 1990
- **treatment:** high-dose, “purgine-free”, chemotherapy regimen, including intensive debulking and then PBPC autografting
- **patients:** advanced-stage FCL, SLL and MCL, with adverse prognostic features

I-HDS SCHEME FOR BM+, INDOLENT LYMPHOMAS



***Long-term clinical and molecular evaluation
of 70 patients with low-grade lymphoma
(follicular - mantle-cell - lymphocytic)
all treated with the same intensive approach
Including peripheral blood progenitor cell
(PBPC) autograft***

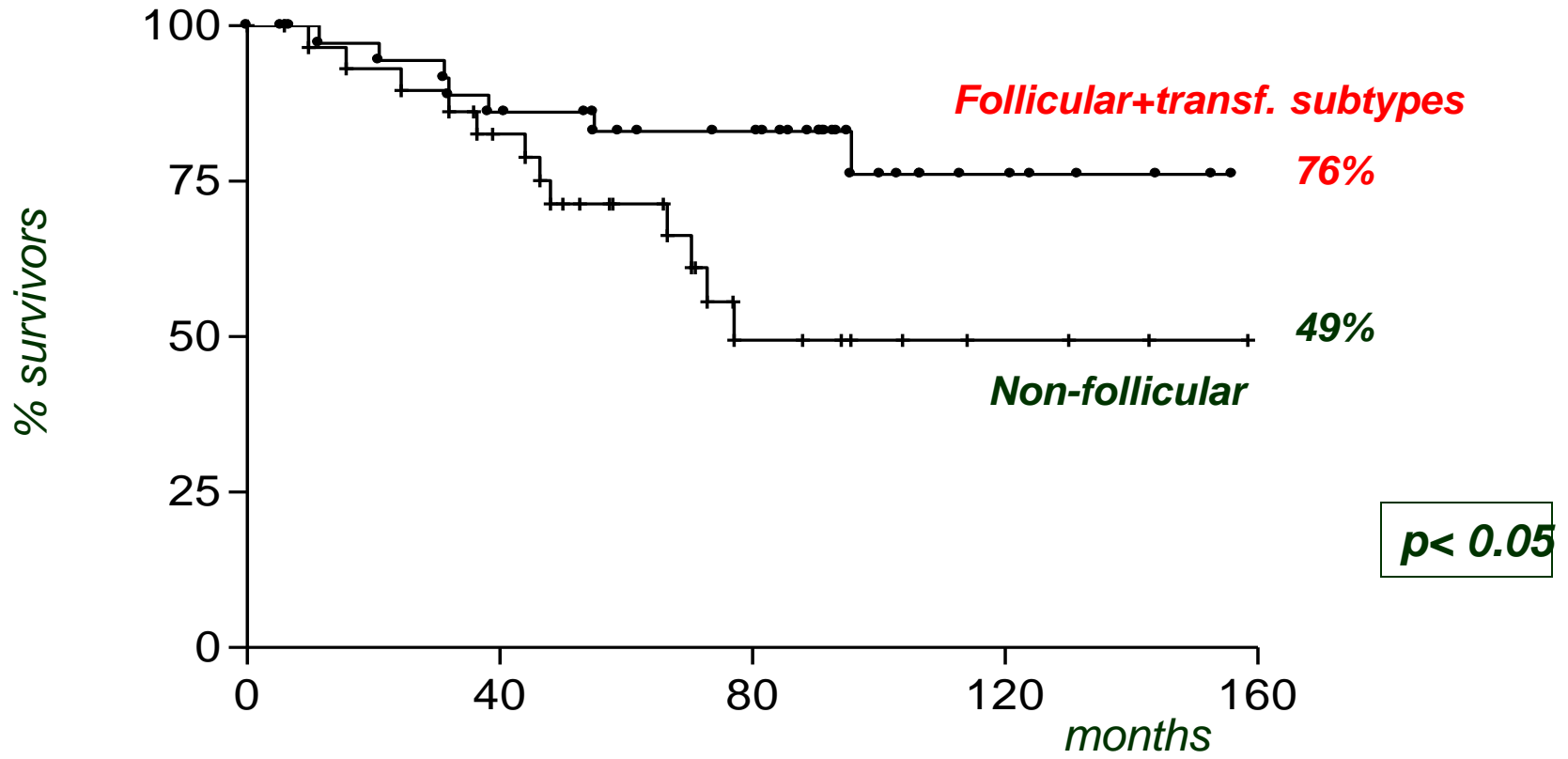
Corradini et al. J Clin Oncol, Apr 15, 2004

Table 1. Main clinical features of the patients entered in the study

<i>Parameter</i>	<i>n</i>	<i>(%)</i>
• Age, median (range), years	47	(32-61)
• Male/female ratio	43/27	
• Diag./rel.	61/9	
• Histology		
lymphocitic	14	20
Follicular	29	41
Transformed	11	16
Mantle cell	16	23
• Stage IV	61	87
• Tumor-related symptoms	45	65
• Bone marrow involvement	57	81
• aalPI score ≥ 2	30	43

Figure 2 A

OVERALL SURVIVAL



Long term molecular monitoring

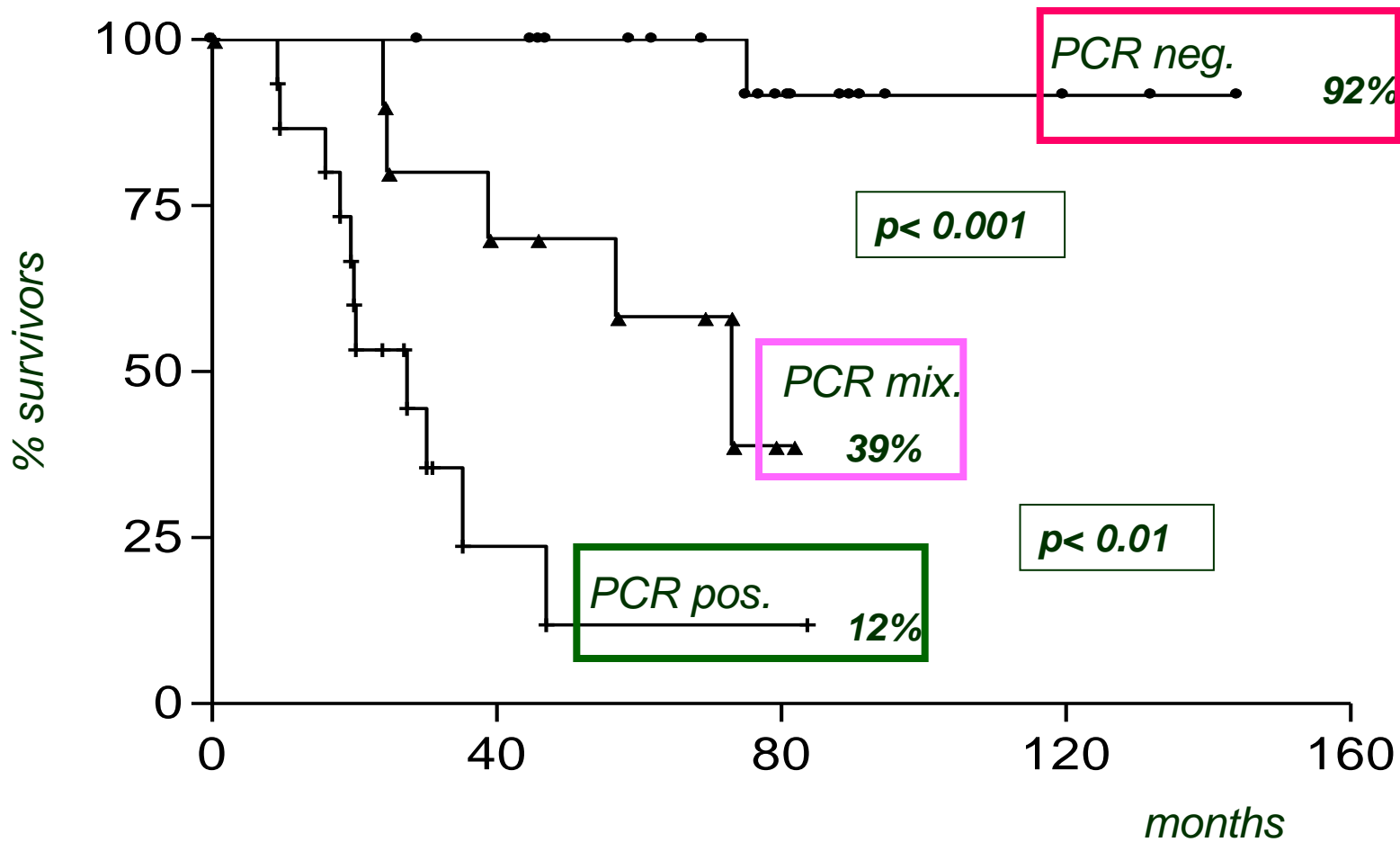
(Median follow-up : 75 months)

• **Patients with a molecular marker: 60 (86%)**

PCR negativity

	<i>post-harvest</i>	<i>post-ABMT</i>	<i>last follow-up</i>
	<i>n= (%)</i>	<i>n= (%)</i>	<i>n= (%)</i>
Follicular	19/35 (54)	21/30 (70)	18/22 (82)
Non follicular	3/25 (12)	5/20 (25)	1/20 (5)

Disease free survival according to PCR status during molecular follow-up



Multivariate analysis

Characteristics	Overall survival as end point			Event-free survival as end point		
	Risk ratio	95% CI	p value	Risk ratio	95% CI	p value
CR achievement: -NO	6.29	1.9-21.2	<u>0.003</u>	9.8	3.7–25.6	<u><0.0001</u>
MR achievement: -NO	46.9	5–465	0.001	24.4	3.1 – 191	0.0023

CONCLUSIONS - 1

- FCL e non-FCL patients display a significantly different outcome in terms of:
 - PCR-negative harvests
 - Achievement of post-autograft Molecular Remission
- Post autograft **MR** persistence is highly predictive of prolonged overall and disease-free survival



Thus, **MR** achievement seems an essential “target” in the autograft setting

are high-dose therapies with autograft reproducible at a multicenter setting ?

1997: GITMO study to evaluate reproducibility in a multicenter setting of i-HDS as front-line therapy for advanced-stage FCL < 60 y.o.

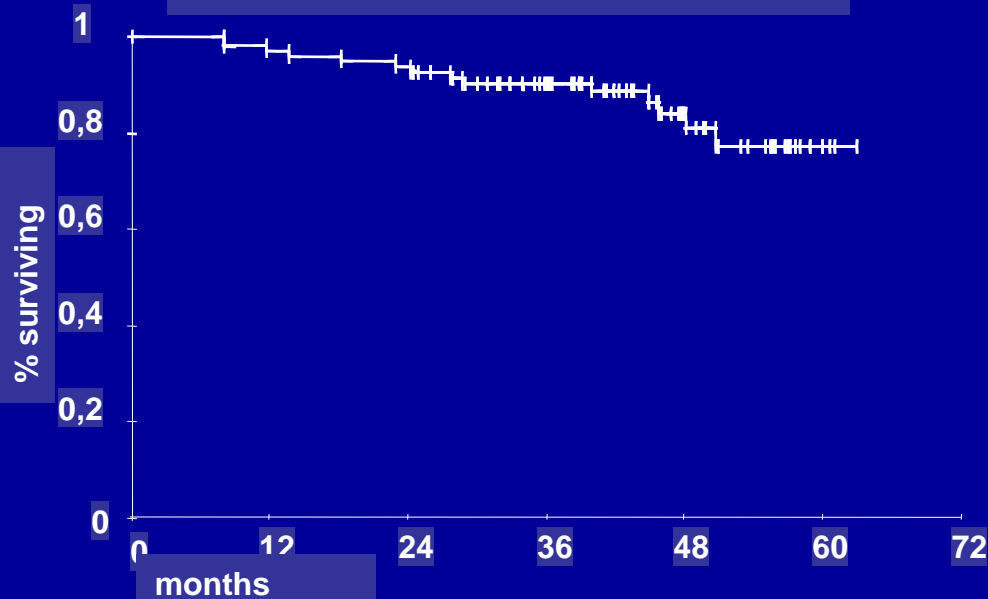
**THE MULTICENTER GITMO TRIAL HAS
BEEN LAUNCHED IN DECEMBER 1996**

**20 ITALIAN CENTERS HAVE BEEN
INVOLVED IN THIS STUDY**

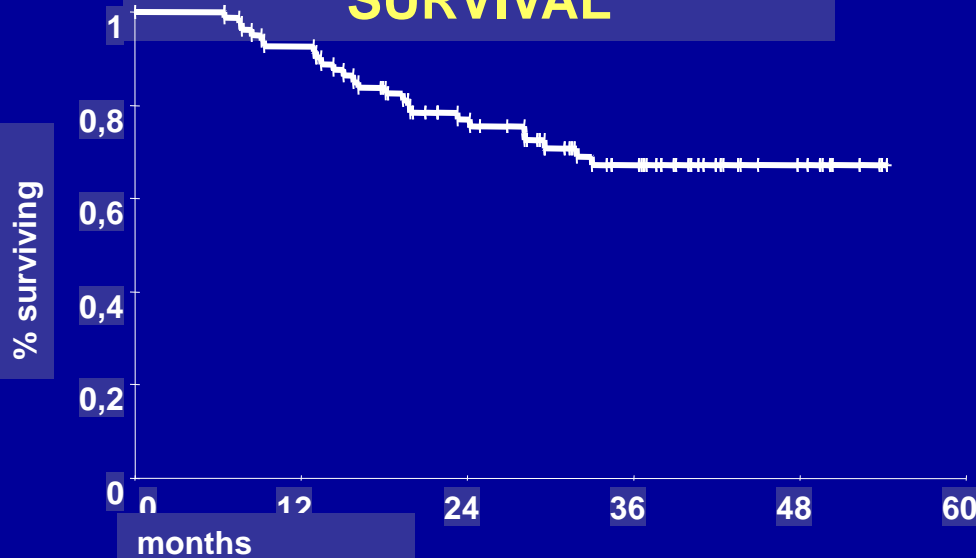
**92 PATIENTS WERE CONSIDERED
EVALUABLE ON AN
INTENTION TO TREAT BASIS**

*three patients were excluded due to inappropriate inclusion
(1MCL, 1CLL, 1CNS involvement)*

OVERALL SURVIVAL



DISEASE-FREE SURVIVAL



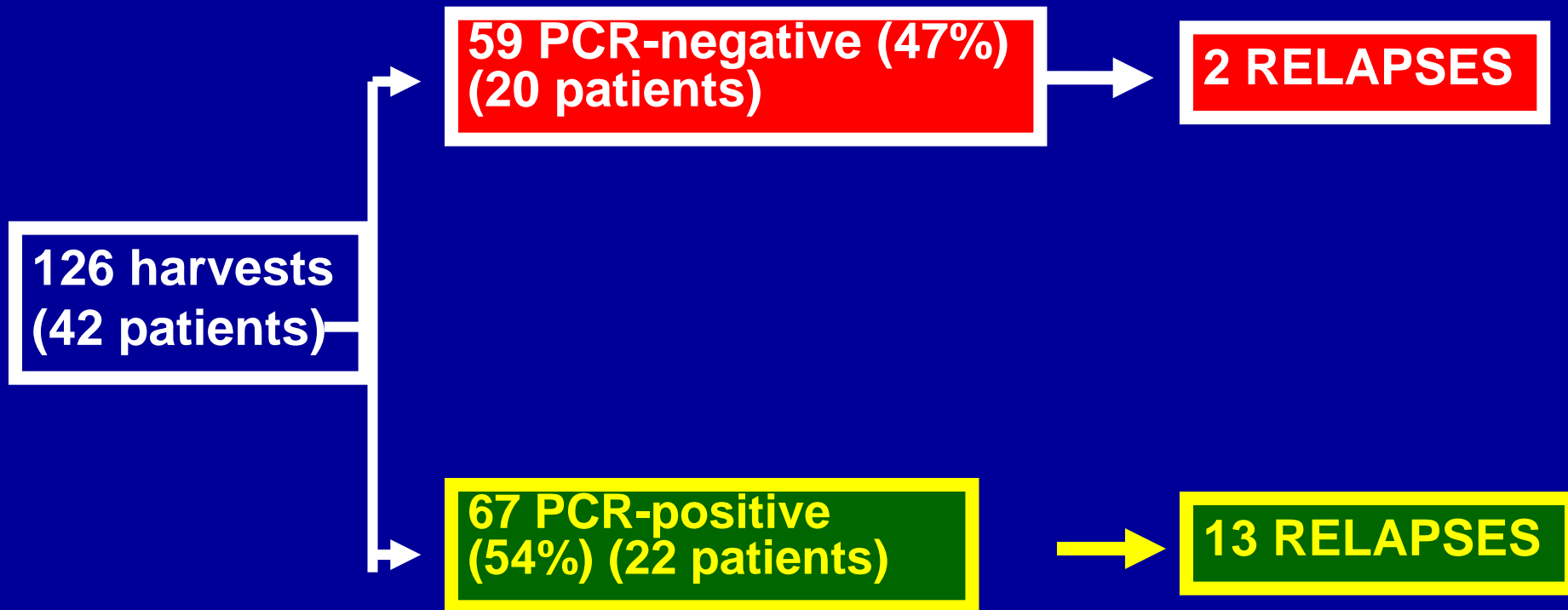
High rate of clinical and molecular remissions in follicular lymphoma patients receiving high-dose sequential chemotherapy and autograft at diagnosis: a multicenter prospective study by the Gruppo Italiano

LADETTO et al, Blood 2002, 100: 1559-65

GITMO multicenter study

MOLECULAR RESPONSE assessed by PCR

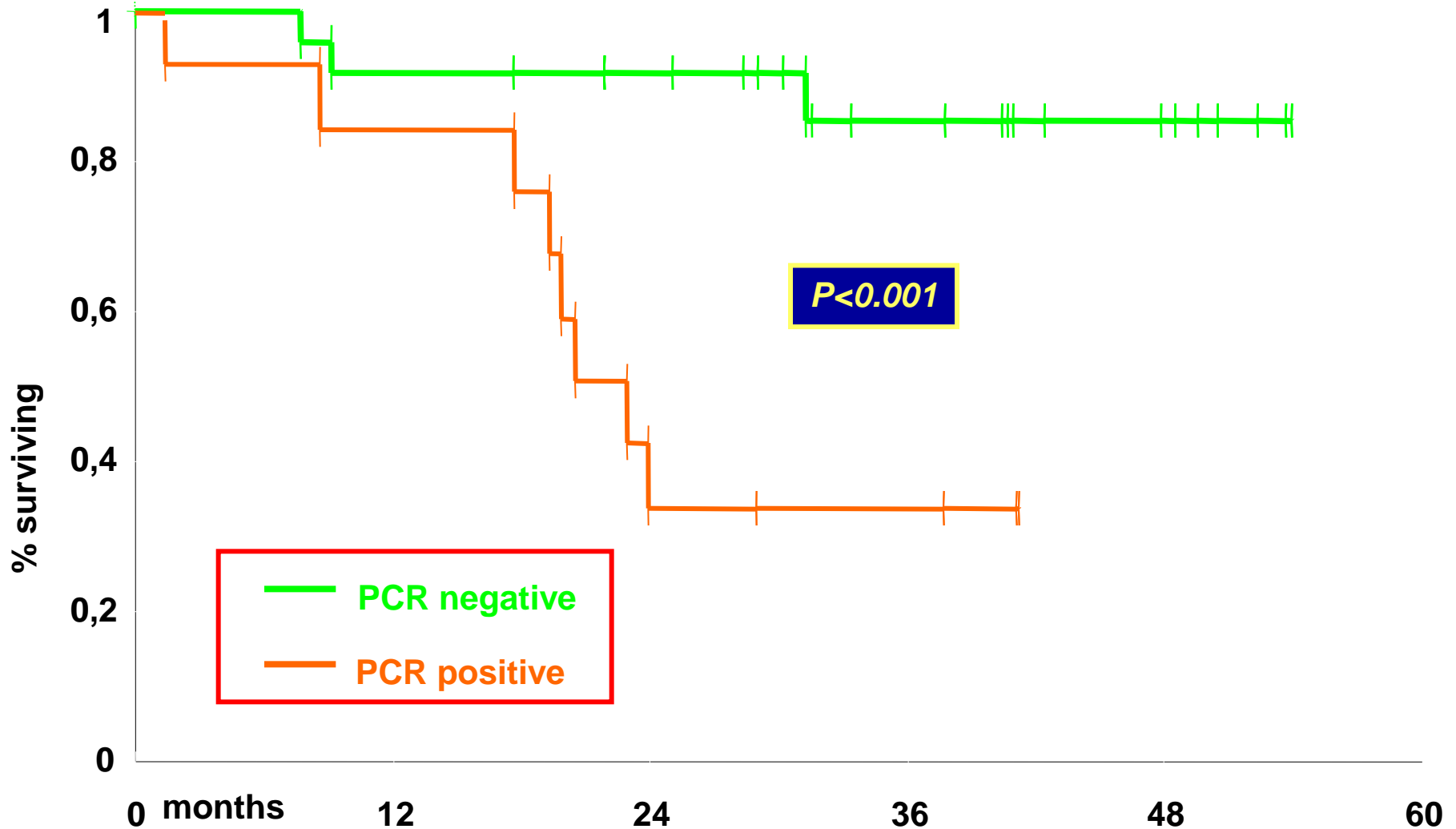
Analysis of PBPC harvests



P=0.06

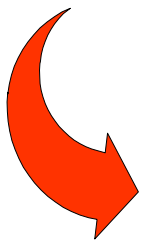
GITMO multicenter study

DISEASE FREE SURVIVAL according to post-transplant PCR status



CONCLUSION - 2

Both single-center and multicenter studies have demonstrated that *MR* achievement is an essential “target” in indolent lymphoma patients undergoing an intensified program with autograft



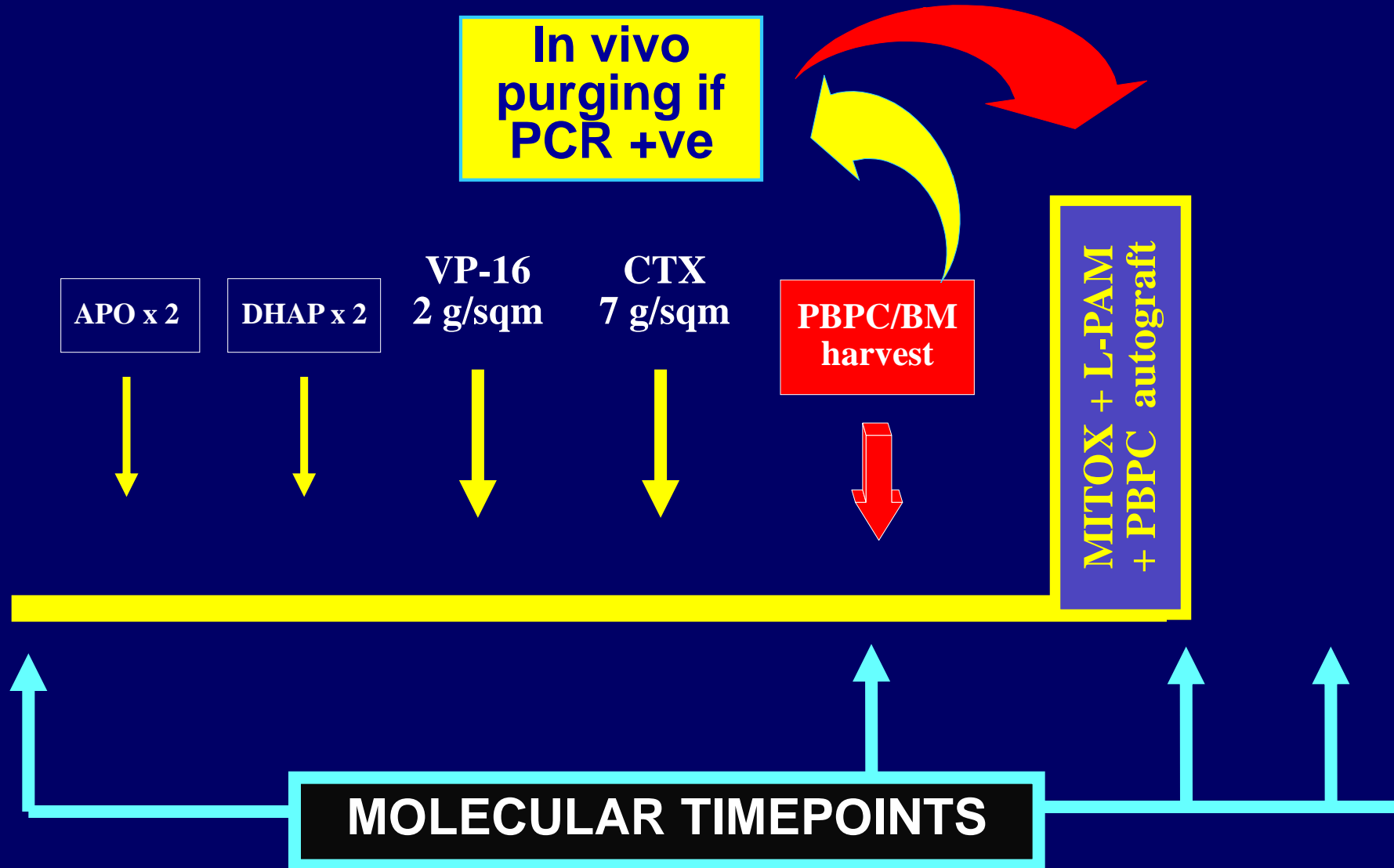
May we increase the number of patients achieving MR

IN VITRO MANIPULATION ("ex-vivo purging") OF CIRCULATING HEMOPOIETIC PROGENITORS

in follicular and mantle-cell lymphoma patients treated with i-HDS, and displaying molecularly detectable minimal residual disease in the harvests

Tarella C. et al.,
Leukemia 1999, 13: 1456

I-HDS SCHEME FOR BM+, INDOLENT LYMPHOMAS



immunomagnetic negative ex-vivo purging procedure

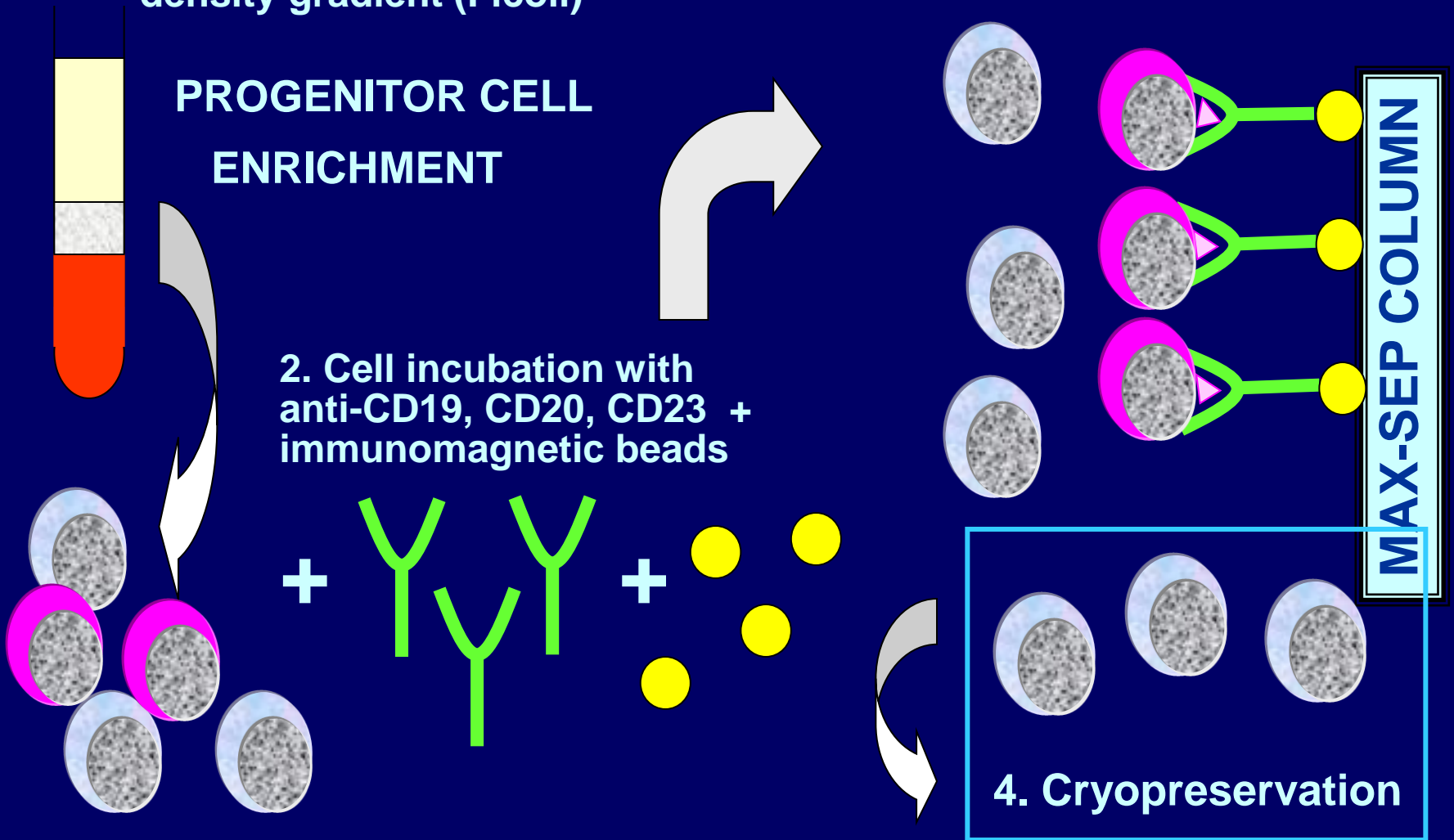
1. Cell separation on density gradient (Ficoll)

PROGENITOR CELL ENRICHMENT

2. Cell incubation with anti-CD19, CD20, CD23 + immunomagnetic beads

3. Lymphoma Cell separation through a passage on a Max-Sep column

4. Cryopreservation



PCR assessment of the “ex-vivo purging” efficiency

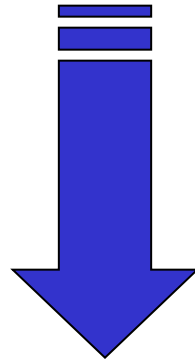
	<u>Evaluable pts.</u>
Follicular Lymphoma	8
• post-purging PCR neg	3
• post-autograft PCR neg	3
• <i>total achieving PCR neg</i>	6
<u>Mantle Cell Lymphoma</u>	3
• <i>post-purging PCR Neg</i>	<i>none</i>
• <i>post-autograft PCR Neg</i>	<i>none</i>

Andersen NS, et al. (Dana Farber group, Boston)

Blood 90: 4212-4221, 1997:

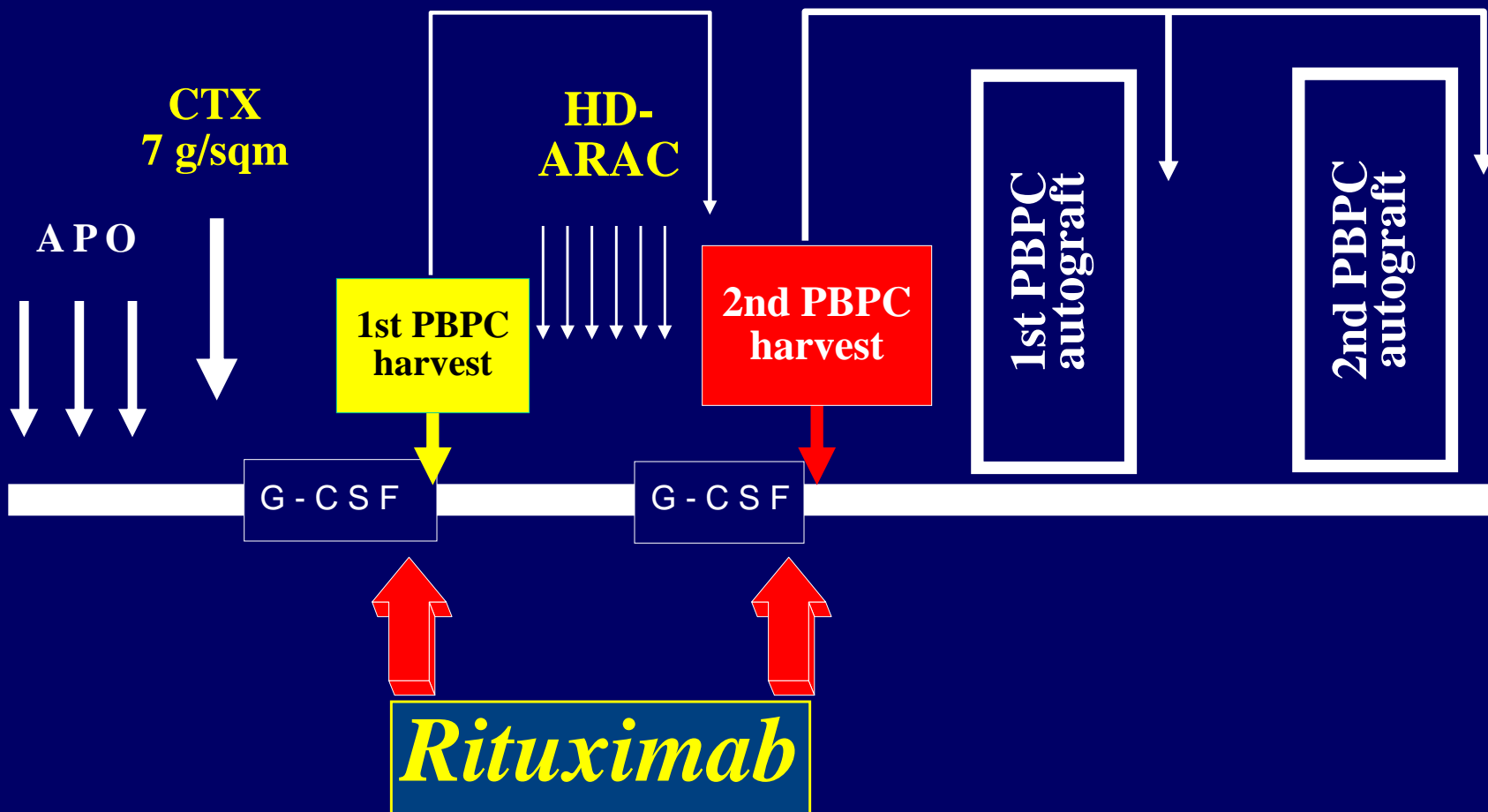
**Failure of immunologic purging in
Mantle-cell lymphoma assessed by
polymerase chain reaction detection
of minimal residual disease**

Searching for new approaches able to increase the chances of harvesting PCR- PBPC

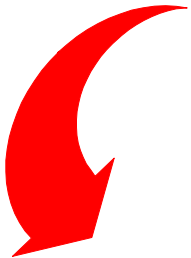


*the potential use of Rituximab in
combination with autograft with
the aim of exploiting an
in vivo purging effect
prior to PBPC harvesting*

MODIFIED HDS WITH RITUXIMAB (R-HDS) GIVEN PRIOR TO PBPC COLLECTIONS



**A RITUXIMAB-SUPPLEMENTED HDS REGIMEN
HAS BEEN PROVED TO BE FEASIBLE AND
EFFECTIVE IN MCL AND IN RELAPSED FCL**



**SUCCESSFUL IN VIVO PURGING OF CD34-CONTAINING
PERIPHERAL BLOOD HARVESTS IN MANTLE CELL AND
INDOLENT LYMPHOMA: EVIDENCE FOR A ROLE OF
BOTH CHEMOTHERAPY AND RITUXIMAB INFUSION**

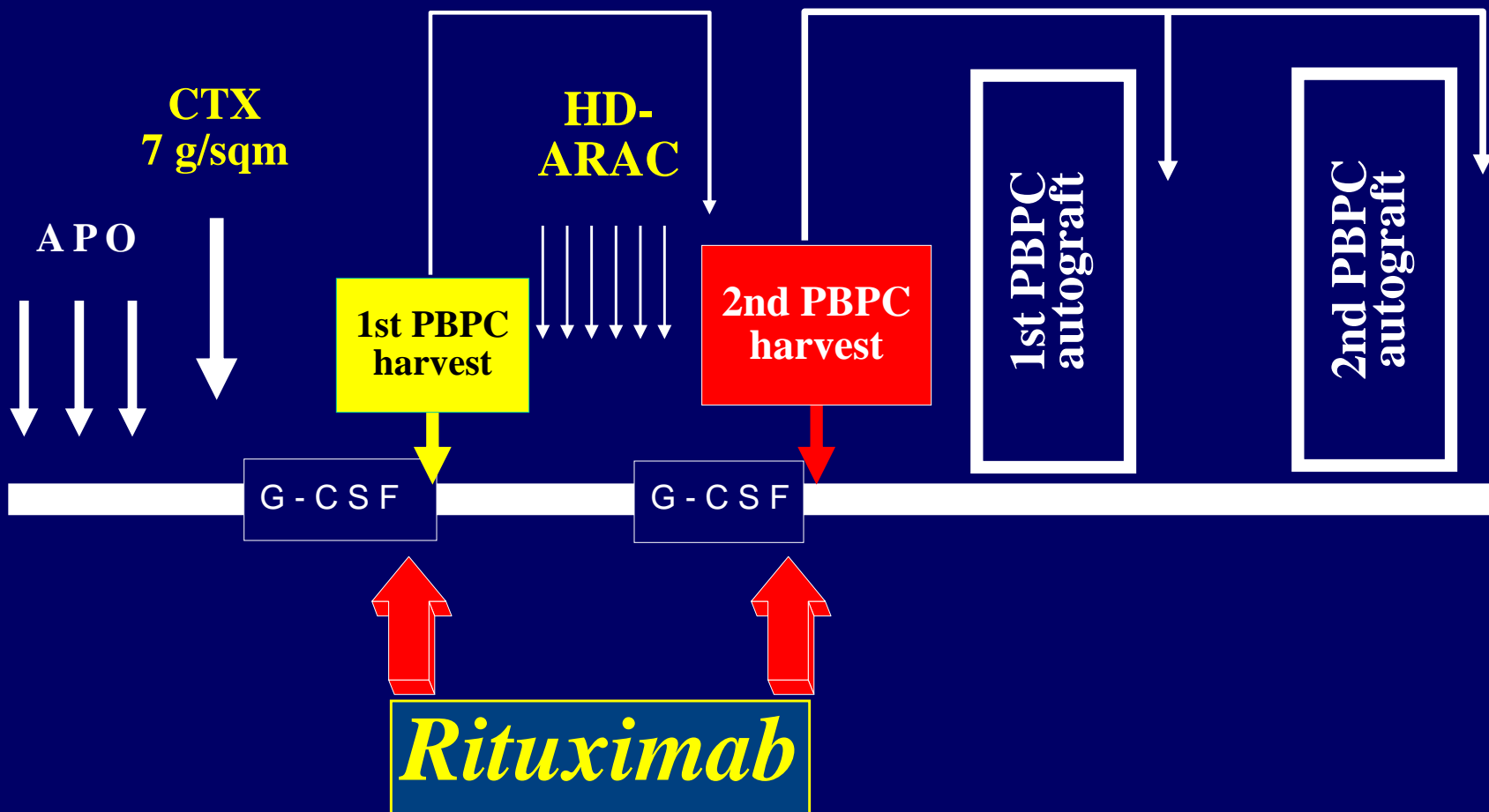
*Magni M, Di Nicola M, Devizzi L, Matteucci P, Lombardi F, Gandola L,
Ravagnani F, Giardini R, Dastoli G, Tarella C, Pileri A, Bonadonna G,
Gianni. Blood 2000, 96:864*

R-HDS vs. HDS pilot study

Main Patient Characteristics

	R-HDS <i>n=15</i>	HDS <i>n=10</i>
Age, median (range)	43 (34-58)	46 (36-53)
Female/Male (No.)	4/11	5/5
Histology (No.)		
- follicular	7	7
- mantle	7	3
- marginal	1	-
BM involvement	11	7
PCR+ only	4	3
PB lymphoma cells	6	3

MODIFIED HDS WITH RITUXIMAB (R-HDS) GIVEN PRIOR TO PBPC COLLECTIONS



Quantity and quality of harvested PBPC in R-HDS treated and group patients

	<i>R-HDS</i> <i>(n=15)</i>	<i>Controls</i> <i>(n=10)</i>
CD34+x10⁶ /kg p-CY	10,5	14,5
PCR neg p-CY	57%	20%
CD34+x10⁶ /kg p-AraC	23	22,5
PCR neg p-AraC	93%	44%

***Preliminary experience at the Turin Center
With the Rituximab-HDS approach
As 1st line therapy for MCL
(april 1999 – november 2003)***



- 11 patients have been treated, all of them achieved CR, with MR achievement in 6/6***
- at a median follow-up of 36 mos., 11/11 patients are alive***
- so far, 1 single recurrence has been recorded: this patients is now in his 2nd remission after allogeneic transplant)***

***efficacy of the Rituximab-supplemented
approach in high-risk FCL and
in relapsed/progressing B-DLCL***



**CONCURRENT ADMINISTRATION OF HIGH-DOSE
CHEMOTHERAPY AND RITUXIMAB IS A FEASIBLE AND
EFFECTIVE CHEMO/IMMUNOTHERAPY FOR PATIENTS
WITH HIGH-RISK NON-HODGKIN'S LYMPHOMA.**

*Marco Ladetto, Francesco Zallio, Sonia Vallet, Irene Ricca, Alessandra
Cuttica, Daniele Caracciolo, Paolo Corradini, Monica Astolfi, Selina
Sametti, Federica Volpato, Paola Bondesan, Umberto Vitolo, Mario
Boccadoro, Alessandro Pileri, Alessandro M. Gianni, Corrado Tarella.*

LEUKEMIA, 2001, 15: 1941

R-HDS in high-risk FCL results

	<i>refr./rel</i> <i>n=</i>	<i>at diag.</i> <i>n=</i>
<i>Clinical CR</i>	5/7	10/11
<i>Molecular CR</i>	5/6	10/10
<i>TRM</i>	1	0
<i>CCR</i>	5/7	10/11

4-yr. Overall Survival: 84% (median follow-up: 2 yrs.)

**CURRENTLY ONGOING RANDOMIZED
GITMO TRIAL
*IN POOR-RISK FCL AT DISEASE ONSET***

**RITUXIMAB-supplemented HDS
(R-HDS)**

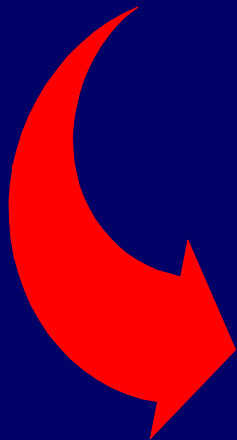
versus

**RITUXIMAB-supplemented CHOP
(CHOP then R)**

**involving 50 Centers
associated to GITMO**

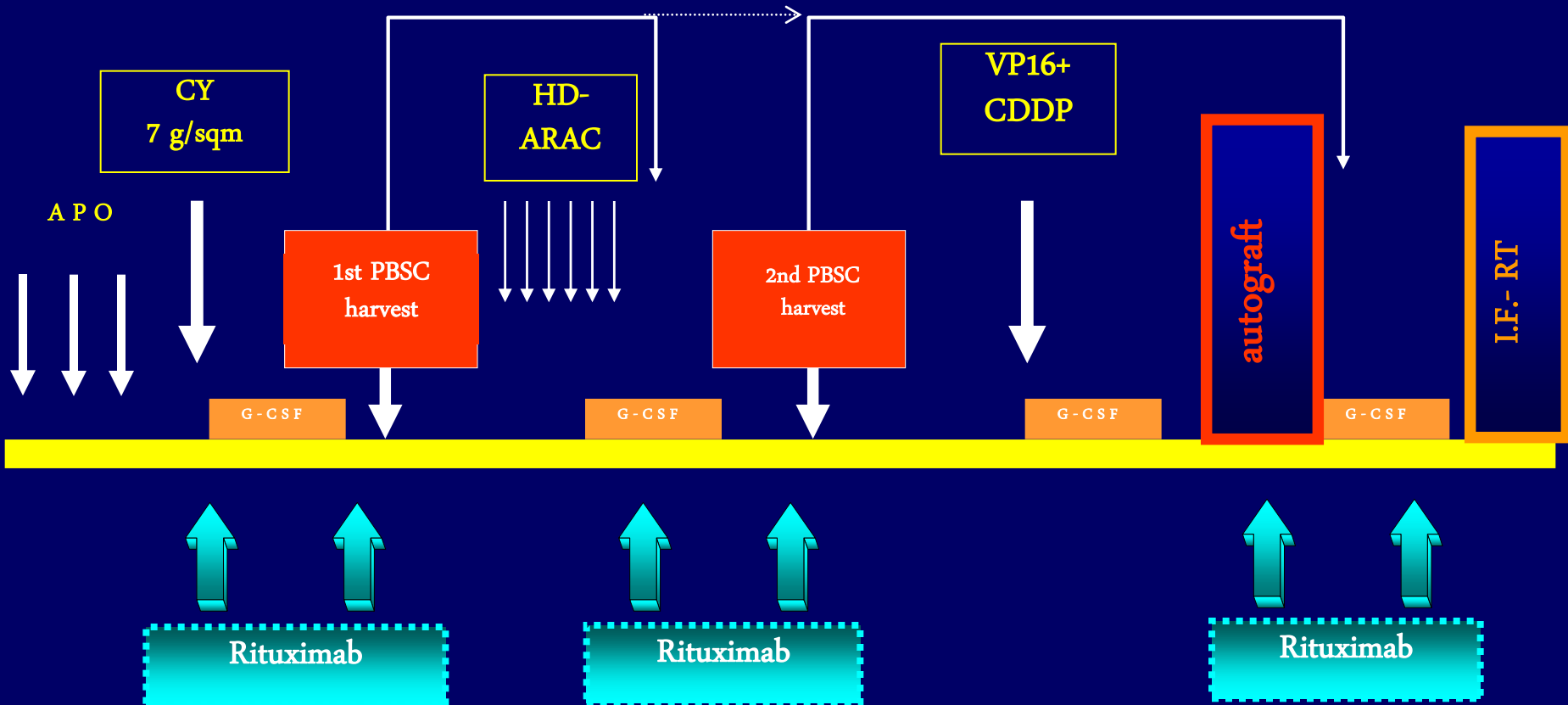
R – HDS in aalPI 2-3 DLCL

*a multicenter experience of
the GRUPPO ITALIANO
TERAPIE INNOVATIVE NEI
LINFOMI (GITIL)*



Preliminary results

RITUXIMAB-SUPPLEMENTED HDS SCHEME FOR HIGH-RISK B-CELL DLCL



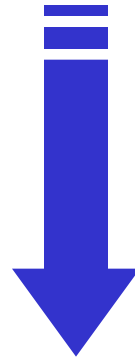
R-HDS multicenter study protocol

Main Patient characteristics

<i>Parameter</i>	<i>patients (n =95)</i>
• Age, median (range), yrs.	48 (18-66)
• aalPI 2 vs 3	58 vs 33
• Extranodal sites	57
- BM invasion	29
• Histological transformation	7

18 out of 29 (62 %) B-DLCL patients
with 2-3 aalPI score and BM involvement
are presently alive in CCR following
R-HDS, at + 10 up to + 35 mos.
since treatment conclusion

Several recent studies have confirmed the efficacy of Rituximab for “in vivo purging” purposes, allowing the collection of PCR – in most patients



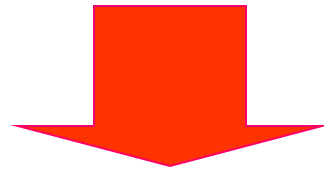
The use of Rituximab is advisable whenever patients with B-cell lymphoma, with BM involvement, are enrolled in intensive programs with PBPC autograft

Is there any indication for the early treatment of “molecular recurrence” ?



Again, at present the treatment of molecular recurrence should be considered only for high-risk patients, if managed with programs aimed to disease eradication or at least maximal tumor cytoreduction

Molecular Recurrence of the Bcl-2/IgH Rearrangement in FL Following ABMT



- IS AN UNCOMMON BUT WELL DEFINED EVENT
- IS ASSOCIATED WITH A HIGH RISK OF RELAPSE



EARLY TREATMENT OF MOLECULAR RELAPSE?

Table 1. Sequence Analysis of *BCR/ABL* Rearrangements From Normal Individuals

Sample	<i>BCR/ABL</i> fusion	Allele	Reference
27	gag-myc	12/16	Shigenaga 1993-94
301	gag-myc	12/16	Shigenaga 1993-94
302	gag-myc	12/16	Shigenaga 1993-94
427	gag-myc	12/16	Shigenaga 1993-94
501	gag-myc	12/16	Shigenaga 1993-94
502	gag-myc	12/16	Shigenaga 1993-94
503	gag-myc	12/16	Shigenaga 1993-94
504	gag-myc	12/16	Shigenaga 1993-94
505	gag-myc	12/16	Shigenaga 1993-94
506	gag-myc	12/16	Shigenaga 1993-94
507	gag-myc	12/16	Shigenaga 1993-94
508	gag-myc	12/16	Shigenaga 1993-94
509	gag-myc	12/16	Shigenaga 1993-94
510	gag-myc	12/16	Shigenaga 1993-94
511	gag-myc	12/16	Shigenaga 1993-94
512	gag-myc	12/16	Shigenaga 1993-94
513	gag-myc	12/16	Shigenaga 1993-94
514	gag-myc	12/16	Shigenaga 1993-94
515	gag-myc	12/16	Shigenaga 1993-94
516	gag-myc	12/16	Shigenaga 1993-94
517	gag-myc	12/16	Shigenaga 1993-94
518	gag-myc	12/16	Shigenaga 1993-94
519	gag-myc	12/16	Shigenaga 1993-94
520	gag-myc	12/16	Shigenaga 1993-94

NOTE: Multiple sequences of the *BCR/ABL* rearrangement located in patients with chronic Philadelphia chromosome-positive leukemia. Patients of Shigenaga were identified in relation to *BCR/ABL* rearrangement in 1993 and 1994. Sequence accession no. U00001 and U00002.

frequency of more than one in 10⁶ cells, the mean age was 45.8 years.

Comparison of the *BCR/ABL* Rearrangements in NP and FL

Sequence analysis of the *BCR/ABL* rearrangement, performed on PCR products isolated from seven normal samples having the highest frequency of the rearrangement, confirmed that these products were unique and specific for the *BCR/ABL* rearrangement (Table 1).

DISCUSSION

The *BCR/ABL* rearrangement can be used as a marker to monitor the course of the disease in patients with FL. However, the finding that the rearrangement is detected in normal individuals poses a potential complication when used to quantify MRD. In this study, the incidence and frequency of the *BCR/ABL* rearrangement was determined in 481 DNA samples from normal individuals using real-time PCR.

The rearrangement was detected in 2.5% (11) of the 481 samples analyzed. In at least 7% of normal samples (two of 84 lymphoblastoid lines, 12 of 187 blood DNA samples), the *BCR/ABL* rearrangement was present at frequencies as low as that seen in NP cells, which is higher than that observed in many patients. The lower incidence of the rearrangement compared with that observed in other studies^{1,2} probably reflects the smaller amount of DNA analyzed for each normal individual. Furthermore, the real-time PCR assay used in this study requires a minimum of 10 target copies for the reproducible detection of the rearrangement, increasing the likelihood of false-negative results. The occurrence of the *BCR/ABL* rearrangement in 25% of normal peripheral blood samples is likely therefore to represent a lower limit for the incidence of the rearrangement in normal individuals.

Previous studies have shown an increasing incidence of the *BCR/ABL* rearrangement with age that suggested a link between the incidence of FL and the occurrence of the *BCR/ABL* rearrangement.^{1,2} The normal individuals described here as being no apparent age-associated increase in

the incidence and frequency of circulating *BCR/ABL* bearing cells. The high detection rate of the rearrangement in the peripheral blood of normal individuals and the low incidence of FL^{1,2} would also seem to rule out the possibility that these cells represent a subclinical form of the disease.^{1,2}

The high incidence of the *BCR/ABL* rearrangement in normal individuals will make it difficult to quantify the frequency of the rearrangement close to patients with FL who may similarly carry a background of *BCR/ABL* cells unrelated to their disease. The difficulty involved in discriminating between the rearrangement close to other *BCR/ABL* bearing cells, usually determined by sequence comparison with the known clonal rearrangement, can also be overcome using disease-specific primers designed to bind target and junction region *Bcr/abl* alleles.¹² This can be necessary to reach the quantities of the rearranged clone in patients, particularly those having low levels of detectable *BCR/ABL* cells. An especially direct quantitation of the rearrangement is only obtained with a minimum viable copy number of five to 10 *BCR/ABL* cells.¹² The current real-time PCR protocols used to measure peaks in the ratio between the number of *BCR/ABL* in FL. However, efforts to further optimize the reproducibility of the real-time PCR assay will be worthwhile as the technique offers a potential real-time method of quantifying the *BCR/ABL* rearrangement when compared with conventional semiquantitative nested PCR approaches.

In conclusion, this study quantifies the presence of *BCR/ABL* rearrangement in a population of normal individuals at levels comparable to or greater than that found in patients with FL. It is likely that the presence of a background of *BCR/ABL* cells in patients could confound the detection and quantitation of MRD, particularly at low levels of tumor burden.

ACKNOWLEDGMENT

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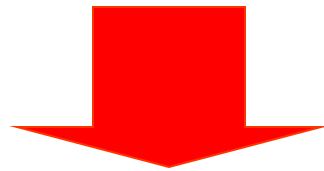
THE HIGH INCIDENCE OF NNBR IN THE PB OF HEALTHY SUBJECTS



MAJOR CONFOUNDING FACTOR IN THE SETTING OF PCR-ANALYSIS

...the current PCR protocols seem to be inappropriate as the sole technique for monitoring MRD in FL. (Summers et al JCO, 2001 15: 420-424)

CAN AT LEAST A PROPORTION OF
THE SO-CALLED MOLECULAR
RELAPSES BE ASSOCIATED TO
NLABR AND NOT TO THE
REAPPARENCE OF THE ORIGINAL
TUMOR CLONE ?



Recent study to verify the nature
of molecular relapses occurring in
our patient population

Ladetto M et al, Exp Hematol 2003, 31(9):784-8

PATIENT SAMPLE

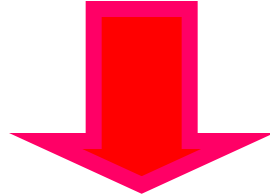
119 FCL patients enrolled in prospective clinical trials including high-dose sequential chemotherapy + autologous transplantation with or without Rituximab supplementation (16 vs 103)

DEFINITIONS

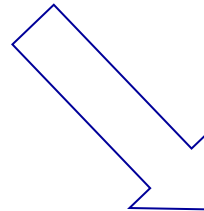
MOLECULAR REMISSION (MR): any CR patient achieving PCR-negativity on two separate BM samples obtained after an interval of at least three months

MOLECULAR RELAPSE: any MR patient reverting to PCR positivity on two separate BM samples taken after an interval of one month.

119 patients with Bcl-2/IgH+ FL



8 MOLECULAR RECURRENCES



6 PATIENTS HAD
THE SAME
REARRANGEMENT
DETECTED
AT DIAGNOSIS

2 PATIENTS HAD A
NOVEL UNRELATED
REARRANGEMENT
(different VH usage
and/or N-insertions)

Summary of the Results

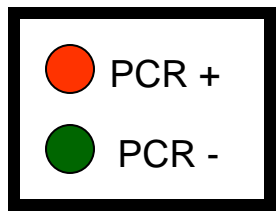
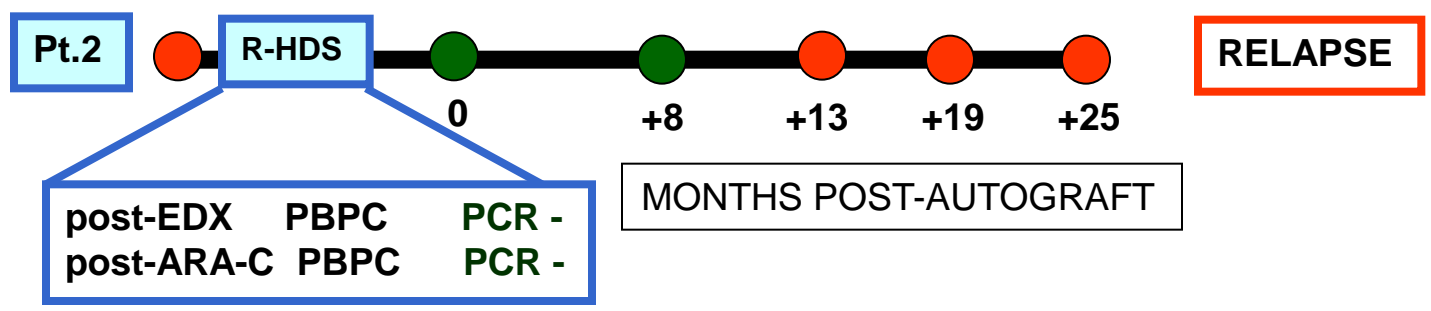
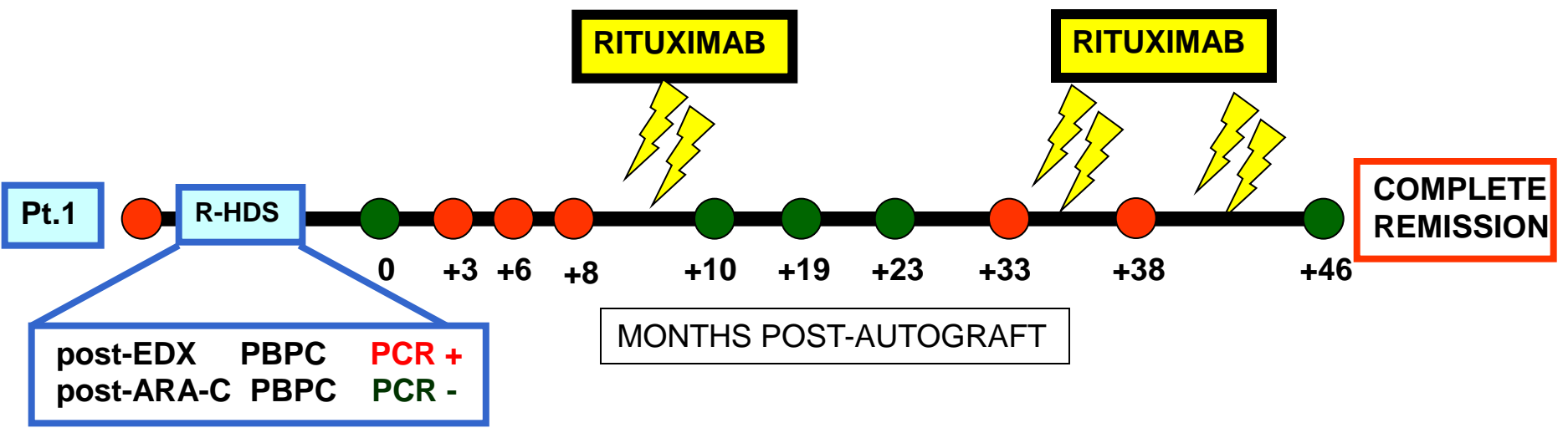
- MOLECULAR RECURRENCE DUE TO UNRELATED REARRANGEMENTS OCCURRED ALWAYS AFTER THREE OR MORE YEARS FROM TRANSPLANT
- THESE NOVEL UNRELATED REARRANGEMENTS PERSISTED FOR SEVERAL MONTHS (20 in one subject)
- NONE OF THE SUBJECTS WITH UNRELATED REARRANGEMENTS SHOWED EVIDENCE OF ACTIVE LYMPHOPROLIFERATIVE DISEASE

R-HDS protocol for Mantle Cell Lymphoma (MCL)

***the risk of a “misdiagnosed” molecular relapse due to
a non-neoplastic bcl-2 rearrangement does not apply
to mantle-cell lymphoma***

Relationship between post-graft PCR status and clinical outcome

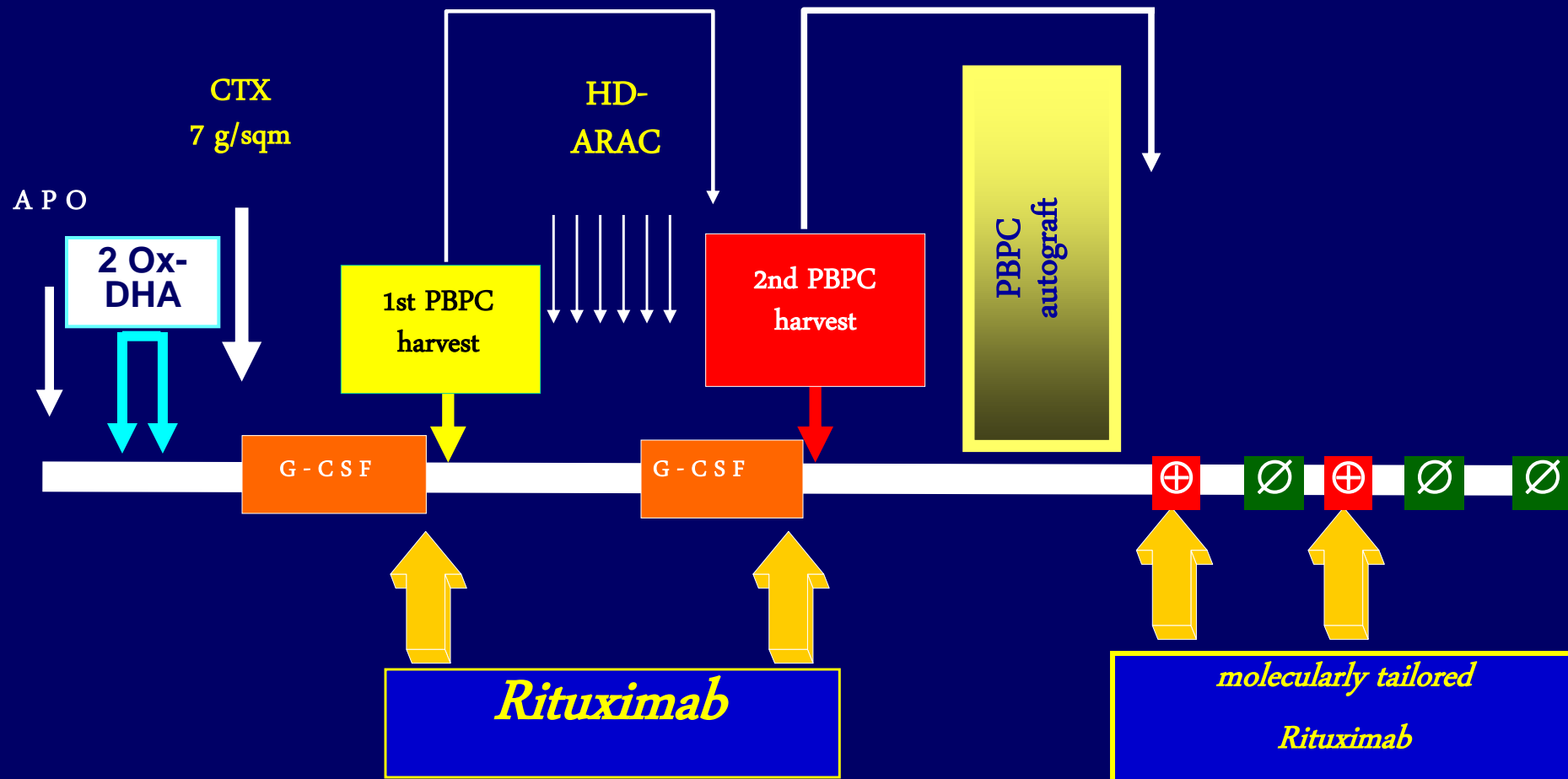
the lesson learned from two cases treated at diagnosis with R-HDS



R-HDS protocol for ***Mantle Cell Lymphoma (MCL)***

***the possible post-autograft Rituximab administration
“tailored” on the molecular monitoring of the minimal
residual disease***

NOVEL MULTICENTER STUDY PROGRAM FOR LMANTLE-CELL LYMPHOMA



MR as an essential “target” in the management of non-Hodgkin’s Lymphoma

conclusions

- **although MR achievement seems associated with a better outcome, at present there are no sufficient data to support the need of MR achievement in indolent lymphoma patients treated with conventional chemo-immunotherapy**
- **On the other hand, MR achievement seems an essential goal for patients enrolled in intensive programs with autograft**
- **In B-cell lymphoma, MR of PBPC harvests can be easily achieved by adding Rituximab as an “in vivo purging” agent**
- **Treatment of “molecular relapse” is not a minor issue in patients potentially curable with intensive treatments**

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Div Ematologia

Università di Torino

Prof. Alessandro Pileri

**ALL CENTERS
PARTICIPATING TO HDS-
BASED STUDY PROTOCOLS
FOR HIGH-RISK NHL**

Clinical significance of MRD in MM

Molecular monitoring of MRD has been used for the following purposes:

- **to evaluate the effectiveness of purging procedures**
- **to evaluate whether PCR negativity predicted a reduced risk of relapse**

Clinical significance of molecular analysis in MM

- **Molecular disease in MM is currently assessed by PCR-based methods**
- **In nearly 70% of MM patients it is possible to identify a molecular marker (IgH specific rearrangement)**
- **Molecular disease significance has been explored in different settings**

Prognostic significance of MRD

Auto-SCT can induce in a small proportion of patients a long term survival (>5 yrs)

It has been hypothesized that molecular status
can predict long term survival after auto-SCT

MRD after auto-SCT

Corradini et al.

- **12/12 patients** were PCR positive after auto-SCT. (Blood 1995)
 - **14/15 patients** were PCR positive after auto-SCT. (JCO 1999)
-

Martinelli et al.

- **25/30 patients** were PCR positive after auto-SCT and PCR negativity was not predictive of prolonged survival . (JCO 2000)

Prognostic significance after autologous stem cell transplantation (auto-SCT)

- **Early studies demonstrated that molecular remission after auto-SCT is uncommon.** (Corradini Blood 1995, Martinelli JCO 2000)
- **Molecular follow up after auto-SCT is not usually performed since the rare molecular remissions are not predictive of long term disease free survival.** (Corradini JCO 1999, Martinelli JCO 2000)

In vitro purging procedures

Autologous stem cell transplantation (auto-SCT) has been identified as the gold standard therapy for patients aged less than 65 years.

However, virtually all patients relapse after auto-SCT.

PCR analysis of purified stem cells (2)

Stewart et al.

(JCO 2001)

- **190 patients** were randomized to receive an autograft of CD34-selected or unselected PBSC
- a median **3 log** depletion (limiting dilution assay) of plasma cells was achieved
- 50% were **PCR-negative**
- selected and unselected patients had similar outcomes.

PCR analysis of purified stem cells (3)

Voena et al.

(BJH 2002)

- **6/12 patients** of CD34 selected PBSC were PCR positive in small scale selection
- Taqman analysis of large scale CD34 selected PBSC showed a 2 log reduction of tumor cell contamination.

Evaluation of purging procedures

- **Purging manipulations of PBSC (CD34 positive selection +/- negative selection) have been shown to be ineffective, since residual tumor cells persist after purging.** (Lemoli BJH 1999, Lemoli Blood 2000)
- **Despite a 3 log reduction of tumor contamination, patients transplanted with purged PBSC have similar outcome respect to non-purged patients.** (Stewart, JCO 2001)

Conclusions

- Molecular monitoring of MRD by means of PCR-based methods is feasible in 70-80% of cases
- PCR analysis of purged stem cells is, at present, poorly useful
- PCR follow up of auto-transplanted patients seems unnecessary

Allo-SCT can induce long term survival.

Identification of patients at impending risk of relapse can facilitate early immunotherapeutical interventions (↓toxicity, ↑efficacy)

It has been hypothesized that molecular status can precede disease changes after allo-SCT

Prognostic significance after allogeneic stem cell transplantation (allo-SCT)

- **Early studies have shown that molecular negativity after allo-SCT is achieved in 50% of patients in clinical remission.** (Corradini Blood 1995, Martinelli JCO 2000)
- **A recent analysis of 48 patients in clinical remission has shown that molecular negativity correlates with a low risk of relapse** (Corradini Blood, in press)

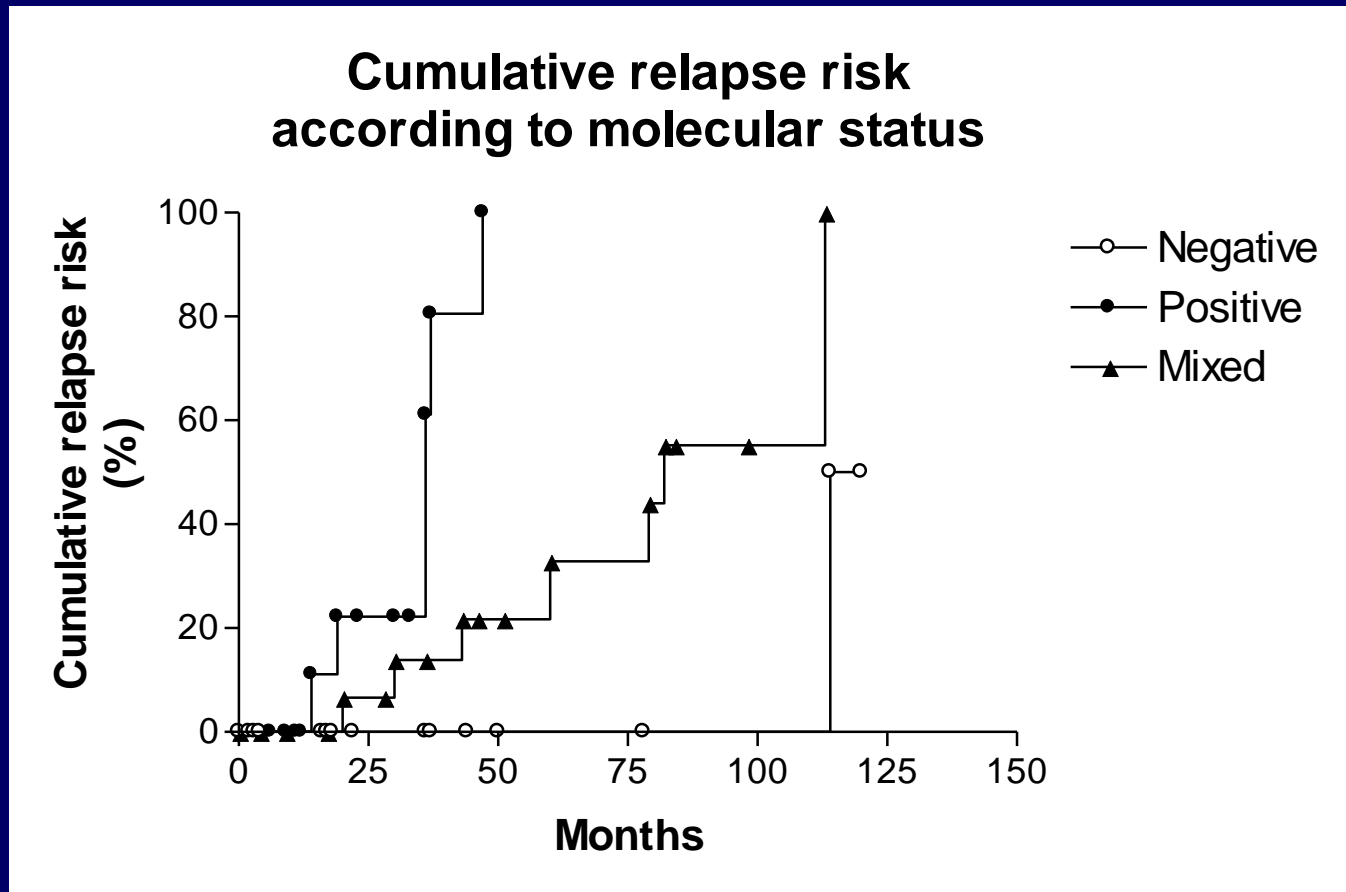
**MOLECULAR REMISSION AFTER MYELOABLATIVE
ALLOGENEIC STEM CELL TRANSPLANTATION PREDICTS
A BETTER RELAPSE-FREE SURVIVAL IN MULTIPLE
MYELOMA: an EBMT study**

70 patients in CR after allo-SCT:

48/70 (68%) had a molecular marker

- 16 pts (33%) were persistently PCR-negative**
- 13 pts (27%) were persistently PCR-positive**
- 19 pts (40%) had a mixed pattern**

Molecular remission after myeloablative alloH SCT predicts a better relapse-free survival in MM



The cumulative risk of relapse at 5 years for NEG, MIX and POS patients was 0%, 33% and 100%, respectively

**Continuous PCR negativity is predictive of
a reduced risk of relapse respect to
continuous PCR positivity (P=0.0001)
and mixed pattern (P=0.001).**

- **PCR analysis of allo-transplanted patients has a prognostic significance and could have clinical implications**
- **Quantitative molecular monitoring of MRD after allo-SCT seems promising**
- **New drugs can be highly effective against MM and molecular monitoring can play a role in assessing the disease response.**

Autograft and Molecular Monitoring in FCL

Conclusions

- the autograft experience has shown that disease eradication may be pursued in indolent lymphoma, specifically in FCL
- the notion of disease eradication is mainly supported by molecular monitoring results
- due to its high predictive value, a post-transplant PCR positive status needs careful clinical evaluation and possibly an early treatment
- however, for FL patients reverting to PCR-positivity following a prolonged period of molecular remission confirmatory direct-sequencing analysis is recommended

Figure 2 B

DISEASE FREE SURVIVAL

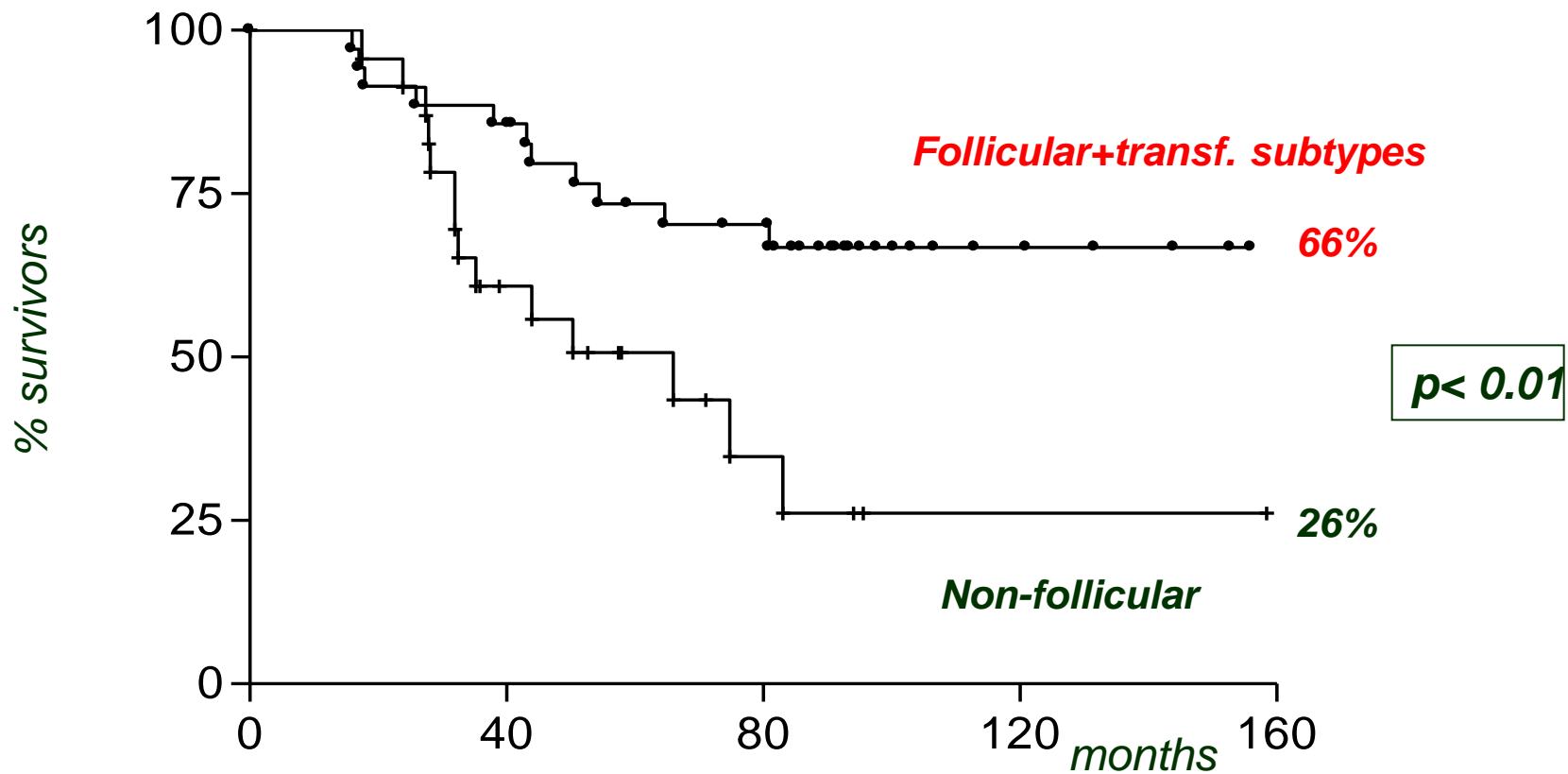
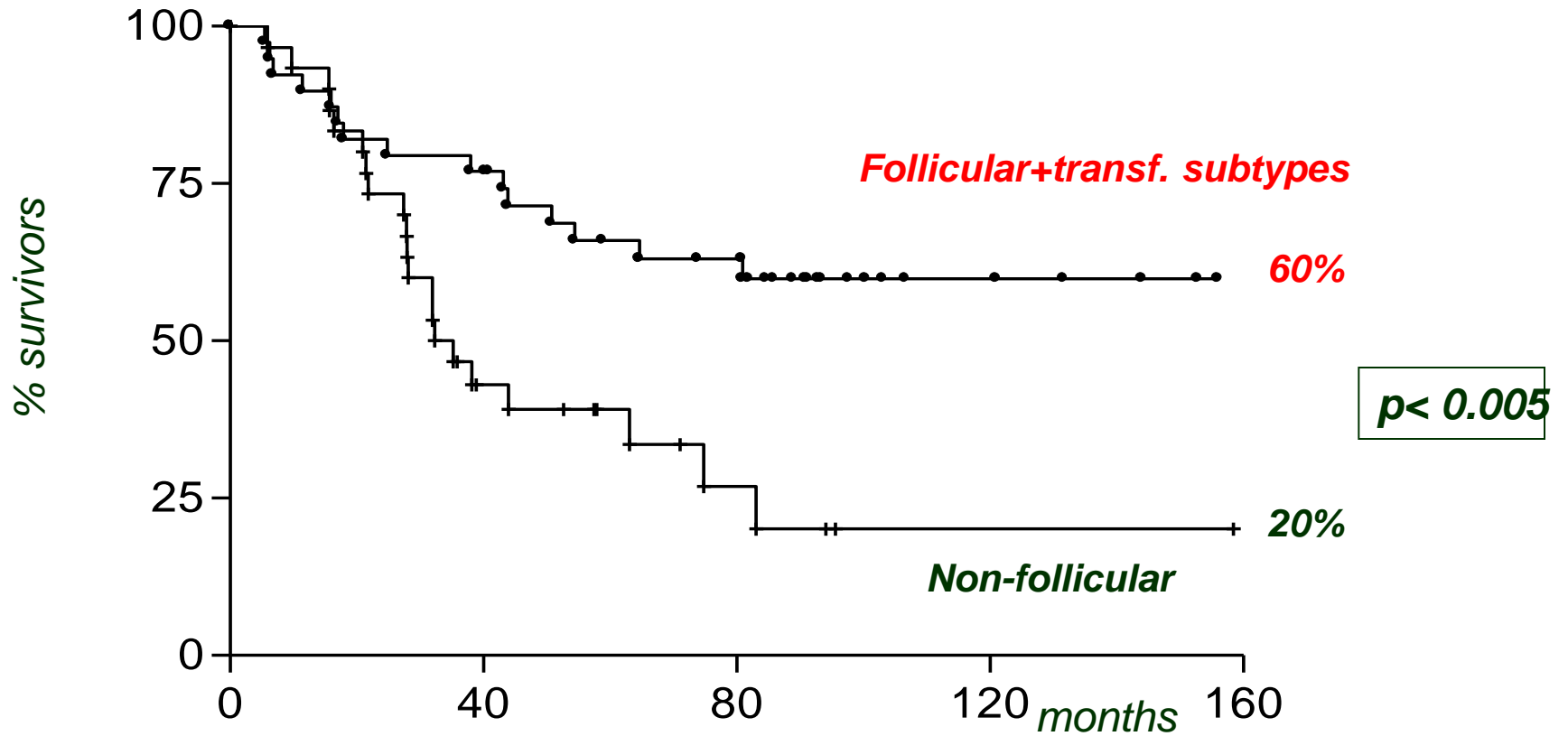
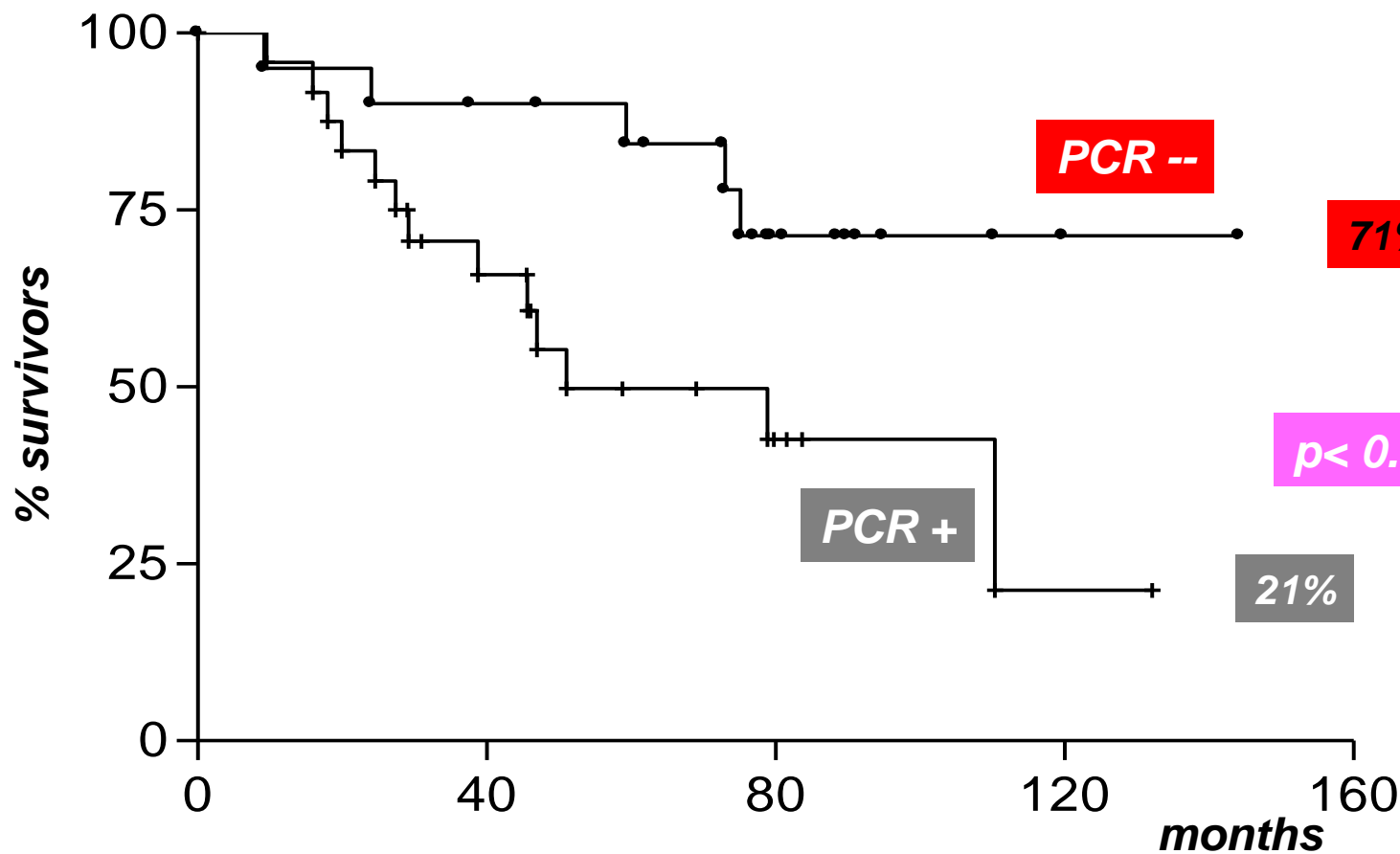


Figure 2 C

PROGRESSION FREE SURVIVAL



Disease free survival according to collection of one or more PCR-negative harvests



Long-Term Follow-Up of Autologous Bone Marrow Transplantation in Patients With Relapsed Follicular Lymphoma

By Freedman A et al.: Blood, 94: 3325-3333, 1999



FFR after ABMT for 113 informative patients who were either PCR- or PCR+ after ex vivo purging