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Human Histocompatibility Testing by T Cell-Mediated Lympholysis: A European Standard CML Technique

Report from the European CML Study Group

on

The Third European CML Workshop

Centre D'Immunologie

INSERM-CNRS

Marseille, December 1979

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The HLA-A, B, C and DR antigenic determinants of the human major histocompatibility system have been defined by antibody mediated cytotoxicity assays using highly selected alloantisera obtained from multiparous women, immunized human volunteers, or allotransplanted patients. Using many sera which have been collated through the international histocompatibility workshops (e.g. Histocompatibility Testing 1977) it has been possible to define 60 distinct specificities genetically coded by the HLA-A, B, C and DR loci.

The concept of histocompatibility-typing with cells ("cellology") was first introduced by scientists working with the Mixed Lymphocyte Culture (MLC) technique (Bradley et al 1972, Mempel et al 1973). With this technique, the HLA-Dw determinants and the HLA-D locus were established (Histocompatibility Testing 1975).

Subsequently the use of *in vitro* generated cytotoxic T-lymphocytes for histocompatibility typing was proposed and initiated by several, separate groups (e.g. Mawas et al 1973, 1974, Kristensen et al 1974). Both products of the HLA-B, C (e.g. Grunnet et al 1975, 1976) and the D/DR loci (Albrechtsen et al 1977, Feighery & Statsny 1979, Johnsen 1978) may be recognized by cytotoxic lymphocytes, but it is not yet clear whether the actual determinants recognized by cytotoxic T lymphocytes are identical to those recognized by antibody, as had been assumed from early CML-studies (review by Eijsvoogel et al 1973). Thus, a collaborative effort to examine the relationship between the determinants defined by serology and cellology was initiated in 1977 (Kristensen 1978). The findings of investigations performed locally and the data obtained from CML-Workshop



(Goulmy et al 1976, 1977, Kristensen & Grunnet 1975, Kristensen et al 1976, Kristensen 1978, Schendel et al 1977, 1978a, Bradley et al 1978) suggested that cytotoxic T lymphocytes recognize antigen differently than do available, highly selected HLA antisera, or in fact may recognize products under different genetic control. In order to investigate these models on an international basis, two further European CML-Workshops were initiated with the aim of standardizing the CML-technique before proper immunogenetic investigations were undertaken. The results of CML-Workshop II have been published previously (Bradley et al 1980). This report, covering CML-Workshop III, presents the *European Standardized CML-technique for histocompatibility typing*. It contains detailed technical descriptions as well as procedures for data-analysis and criteria for assignment of positive and negative responses.

Recently it has become possible to propagate and clone cytotoxic T-cells (Bonnard et al 1978, Schendel et al 1978b, 1980, Charnot et al 1980, Mahissen et al 1979, 1980, Goulmy et al 1980). These techniques — much like monoclonal antibodies — may allow fine structural analysis of target cell properties both on genetic and biochemical levels, adding to our understanding of the HLA-system. Consequently, aspects of these techniques were included in Workshop III and will be employed in future CML-Workshops.

Goals of Workshop III

Based on the results of European CML-Workshop II (Bradley et al 1980) a partially standardized protocol was developed, utilizing not only similar procedures but also common sources of basic

reagents. The purpose in establishing such a protocol for Workshop III was to test (1) the same genetic combinations with common reagents and techniques to assess if more uniform results among the participating laboratories could be obtained, (2) to compare this standard technique with two variations, i.e. generation of CTL in microtiter plates with direct addition of target cells and carboxyfluorescein-diacetate labeling of target cells, (3) to compare various methods of data analysis, (4) to test various CTL expansion techniques, including, specific restimulation in 2° MLC, restimulation with pokeweed mitogen (PWM), and expansion with conditioned medium containing TCGF (T-cell growth factor).

Methods

1 — The Standardized European CML (SE CML) technique

A detailed account of the technique is given in Table 1, which also includes indications of minor discrepancies performed according to local beliefs or necessities.

The standard technique was utilized by the Aarhus, Leiden-1, Marseille and Munich laboratories.

2 — Generation of CTLs in microtiter plates with direct addition of target cells

This technique, performed by Oslo, has been published in detail elsewhere (Hirschberg et al 1977).

Briefly, MLC combinations were established in triplicates in round-bottomed microculture plates on day zero of culturing. A four fold titration of effector cells was used (1×10^5 — 1.25×10^4 cells/well), number of stimulator cells always being half of the number of effectors.

Table 1
European standardized CML technique

<i>SEPARATION OF PBL (Aarhus)</i>		Comments
Anticoagulant	Heparin Sodim Novo without preservative final concentrations 25 IU/ml	defibrinated blood (OS)
Dilution of blood	10 ml blood diluted with 10 ml TC-199 are divided between two glass tubes (100 × 15/17 mm)	Approx 70 ml diluted blood over 25 ml Ficoll (MA)
gradient solution	About 4 ml lymphoprep (Nygaard & Co, Oslo) are, through a long needle, placed below the diluted blood	
temp	All reagents, except the blood, are at 4°C before gradient centrifugation at room temperature	Room temperature for all reagents (MA)
speed	1,000 g	
time	20 min	
washing procedure	After harvest of MNC from 10 ml blood into a conical 10 ml Plastic tube (NUNC N° 1060), the cells are washed × 2 (500 g 10 min) in 10 ml TC-199 with 40 IU heparin per ml added	
<i>FREEZING PROCEDURE</i>	10 × 10 ⁶ MNC are pelleted and resuspended in 0.5 ml (4°C) cooled serum pool. Just before freezing 0.5 ml of a cooled mixture of 20% DMSO (Merck, cat n° 2950) in RPMI is added by pipette a little faster than "drop by drop". After gentle but quick resuspension, the 1 ml suspension is placed into polypropylene tubes (Cat n° 1076, Nunc, Roskilde, DK). These tubes are immediately placed in an uncovered rack in a -80°C deepfreezer. After 1 h or for convenience next day, the tubes are transferred to the final storage in the vapour phase of a nitrogen freezer (-190°C) (Ref Jørgensen & Lamm 1974).	Freezing is done in medium + 20% plasma or serum + 10% DMSO (OS-Ma) and in one step cells are resuspended in freezing medium and distributed in 1 ml ampoules (Ma)
<i>PREPARATION OF SERUM POOL (Aarhus)</i>	4 male donors, blood group AB, donated 500 ml whole blood. Immediately after coagulation at 4°C the bottles were centrifuged, and serum harvested and pooled, prior to	Plasma from heparinized blood from ~ 10 male donors may be used after inactivation at 56°C

Table 1 (contd)

PREPARATION OF SERUM POOL (Aarhus)		Comments
	separation into six aliquots of 200 ml The serum was inactivated at 56°C for 30 min and kept frozen at -20°C	
SHIPPING PROCEDURES (Aarhus)	The frozen tubes were placed in ~ 12 kg of dry ice Packing was performed immediately before shipments The cells were planned to arrive in the local center during the next 12-36 h	
THAWING TECHNIQUE (Aarhus)	The tubes are placed in a test tube rack, and thawing is performed rapidly in a 37°C waterbath, when the last ice crystal is observed the suspension is transferred to a conical plastic tube (NUNC 1030) containing 5 ml cooled TC-199 (4°C), and the cells are immediately centrifuged at 200 g for 10 min This is followed by one wash in 5 ml TC-199 with spinning at 200 g for 5 min (room temperature) After resuspension in 25% SeRPMI the cells are ready for use	Ma uses TC-199 plus heparin to prevent cell aggregation and Os uses DNase for same purpose Only one wash for Ma, Lei 1, Lei 2, Ma, Lei 1, Lei 2 and Os dilute the contents of ampules slowly to prevent osmotic shock
CULTURES FOR THE STIMULATION OF 1° CTL		
medium	RPMI 1640 (Gibco) with, L-glutamine, Penicillin 100 U/ml, Streptomycin 100 µg/ml 20% pooled serum from standardized source	
Culture vessel	50 ml culture flask/standing upright	Round bottomed culture plates — On day 0 cultures are set up in triplicates, with varying amounts of responder and stimulator cells (Os) — For few cells $\approx 2 \times 10^6$, cultures in glass tubes (Ar) or in 2 ml Cluster plates (Ma)
volume	up to 20 ml flask, when volume is less than 5 ml the cells are cultured in Costar marco-well plates with 2 ml volume	

Table 1 (Contd.)

CULTURES FOR THE STIMULATION OF 1° CTL		Comments
cell concentration	Responding cells : 1×10^6 /ml Stimulating cells : 1×10^6 /ml	
culture ratios	1 RC : 1 SC _x	
inactivation of SC	2000 rads, x-irradiation	γ -irradiation for Ar, Lei 1, Lei 2, Ma, Os.
incubation conditions	38°C, 6% CO ₂ , well-humidified incubator, minimal opening of incubator during the culture time.	37°C, 5% CO ₂ for Ar, Lei 1, Lei 2, Ma, Os.
culture duration	144 h.	
CULTURES FOR THE PRIMING OF CTL FOR USE IN 2° CML ASSAYS	The same circumstances as for the generation of primary CTLs, but the time of culture was 10 days.	No need for changing medium (Ma).
FREEZING PROTOCOL FOR PRIMED CTL	At day 10 a minimum of five 20 ml identical cultures were pooled and frozen in a number of $8-10 \times 10^6$ cells per tube (c.f. freezing procedure, vide supra).	
CULTURES FOR THE RESTIMULATION OF SPECIFIC 2° CTL		
medium	same as for 1° CTL	
culture vessel	Cluster macro-well plates with 2 ml capacity	Alternative choices: — 50 ml culture flasks standing upright (Ar — Ma). — round bottomed microplates (Os).
volume	1–2 ml/well	14 to 20 ml of medium in flasks (Ar.Ma).
cell concentration	Responding cells : 1×10^6 /ml Stimulating cells : 1×10^6 /ml	

Table 1 (Contd.)

<i>CULTURES FOR THE RESTIMULATION OF SPECIFIC 2° CTL</i>		Comments
inactivation of SC	2000 rads for 1° cultures	γ-irradiation (Ar. Lei 1, Lei 2, Ma, Os).
incubation conditions	same as for 1° cultures	
culture duration.	120 h	
<i>CULTURES FOR RESTIMULATION OF CTL WITH POKEWEED MITOGEN</i>		
medium	RPMI 1640 + Antibiotics + 10% pooled plasma	25% pooled serum (Ar).
culture vessel	50 ml culture flasks — Standing upright	Glass tubes (Ar).
volume	10 to 20 ml	5 ml (Ar).
cell concentration	1×10^6 ml	3×10^6 C/5 ml (Ar).
PWM source	GIBCO	
preparation	Lyophilised — reconstituted in 10 ml sterile distilled water — no filtration — store at 4°C.	
concentration	1% final	
feeder layer	Autologous to the responding cells — 2×10^6 cells — irradiated 4000 rads.	
incubation conditions	37°C — 5% CO ₂ — well-humidified incubator.	
culture duration	96 to 120 h	
additional comments	Feeder layer is not absolutely necessary but cell yields are much higher when adding this feeder layer.	

Table 1 (Contd)

CULTURES FOR PHA STIMULATION OF CML TARGET CLLS		Comments
medium	same as above	
culture vessel	50 ml culture flasks/standing upright	15 ml tubes, 4 ml total volume, 5×10^5 cells/ml for 0s
volume	5–10 ml	
cell concentration	1×10^6 /ml	
PHA source	Difco PHA-m 0528-57	
preparation	Stock suspended in 5 ml RPMI, dilution of this 5 ml with 45 ml medium, filtration through 0.2μ storage at -20°C	
culture concentration	1 ml of the diluted PHA to 9 ml of medium = 1% of original stock stock	
incubation	same as above	
culture duration	72 h	After 56 h, 8 ml supernate is replaced by 8 ml fresh medium, to prevent aggregation of targets at the day of the assay (Lei 2).
CML ASSAY		
Preparation of CTL		
harvesting from flasks	Flasks are gently shaken — cells are transferred to conical 50 ml glass centrifuge tubes, tubes are centrifuged at $350 \times g$ for 8 min and resuspended in fresh culture medium (2 ml combination)	
medium	Same as for 1° and 2° cultures	

Table 1 (Contd)

CML ASSAY		Comments
<i>Preparation of CTL</i>		
counting procedure	100 μ l of the cell suspension added to 100 μ l of trypan blue Live dead cell ratio determined cell concentration determined on basis of living cells only	
assay plate	V bottomed microtiter plates	Microtest plates (Terasaki type) (Le1 2)
distribution of CTL	100 μ l of the starting suspension distributed to the plates, allowing duplicate well for each target (Note 5 target cells \times 2 replicates \times 100 μ l per replicate = 1 ml of CTL suspension) After the first concentration is distributed 1 ml of culture medium is added to the remaining cell suspension giving a new volume of 2 ml The distribution is repeated as above A series of six dilutions is distributed to the plates with this sequential distribution and dilution pattern	10 μ l of starting suspension are distributed to the plates allowing triplicate wells for each target with Hamilton syringe A drop is formed at the needle tip and applied by touching the edge of the well Controls are heat treated effector cells incubated 15 min in 46°C water bath Controls are handled as above (Le1 2) Volume is 200 μ l (Le1 2)
incubation	As soon as the CTL have been distributed to the plates they are immediately placed in an incubator until the target cells can be added	The CTLs are placed at room temperature until the targets are added (Ar) After distribution of the CTLs the trays are spun at 150 \times g for 2 min and read in the automatic fluorometer (Le1 2)
<i>Preparation of target cells</i>		
harvesting	Flasks are gently shaken and cells transferred to 15 ml round bottomed plastic centrifuge tubes which are centrifuged at 350 \times g for 8 min	
resuspension	Supernatant fluid is decanted and a small volume of about 200 μ l is allowed to run back into the tube cell pellet is broken up and resuspended in this volume by scraping the tube across the top of a test tube rack	

Table 1 (Contd.)

Preparation of target cells		Comments
labeling	250 μ Ci of $\text{Na}_2\text{Cr}^{51}\text{O}_4$ in a volume of 100 μ l is added to each target sample	<ul style="list-style-type: none"> — 18 h before harvesting target cells (0s) — Targets are washed twice with Hanks The cells are labeled by resuspending in 1 ml of a fresh carboxyfluorescein-di-acetate solution in RPMI (final concentration 40 μg/ml) (Le1 2)
incubation	Tubes are placed in a beaker containing water with a temperature of 38°C and placed in a 38°C, 6% CO_2 incubator	Water bath 37°C (Ar, Le1 1, Le1 2, Ma)
incubation time	1 h and 30 min, with shaking at 30 min intervals	15 min with shaking after 7 min (Le1 2)
washing procedures	3 ml of RPMI with antibiotics and 5% pooled serum is added to each tube. This is then layered over a tube containing 5 ml of Lymphoprep ficoll solution. This gradient is centrifuged at 400 g for 20 min. The interface layer is removed and resuspended in 5 ml of the washing medium. Cells are centrifuged at 350 \times g for 8 min. This washing is performed 2 times.	<ul style="list-style-type: none"> — No lymphoprep gradient, but 2 to 3 washes with medium (without plasma or serum) (0s Ma) — 10 ml of RPMI is added and the cells are centrifuged at 350 \times g for 10 min. After decantation of the supernate the targets are resuspended in 1 ml RPMI (without serum) (Le1 2)
distribution	Cells are counted as for CTL and suspended at 1×10^5 /ml, 100 μ l of this suspension is added to each well of the microtiter plate	Viability counting is done as for CTL and cells are suspended at 8×10^5 /ml. 5 μ l of this suspension is added to wells of the microtest plate (Terasaki type). The tray is spun at 350 \times g for 5 min after which the wells are flooded with medium. After decanting the medium, effectors are added (Le1 2)
	Plates containing both CTL, and target cells are incubated for 4 h in incubation conditions identical to those described	Before incubation the plates are centrifuged 50 g, 5 min (Ar, Ma), 150 g,

Table 1 (Contd.)

CML-ASSAY-CONTINUED		Comments
harvesting of supernatants	100 μ l aliquots are removed from each well after the plates are centrifuged at 450 \times g for 5 min.	After incubation the trays are centrifuged at 450 \times g for 5 min and flooded with phosphate buffer saline. After gentle shaking to wash away released carboxy-fluorescein the saline is decanted and the trays are read in the automatic fluorometer (Lei 2).
counting	Samples are counted for 1 min in a Nuclear Chicago Counter.	Commercial filter system harvester (Os, Lei 1).
Spontaneous release	Target cells are incubated in the microtiter plaes in wells containing only culture medium containing 20% serum with no CTL.	
maximum release	1 ml of the final target cell suspension is added to a centrifuge tube containing 1 ml of a 5% Triton-X detergent solution (New England Nuclear). These tubes are centrifuged at 450 \times g for 10 min and 100 μ l aliquots ($\times 3$) are counted.	Zaponine detergent may be used. It makes no difference with Triton X (Lei 1, Ma).
additional comments		The automated fluorometer (Lei 2). Inverted epi-illuminated fluorescent microscope (Leitz) equipped with an automated stage driven by stepping motor (Leitz) and a photomultiplier (Zeiss) encased in a black plastic box with counter balanced door. Digital equipment Corporation PDP 8/E computer (4K memory), high speed paper tape reader and puncher and a teleprint 303 writer. Light source is 60 W tungsten lamp mounted outside microscope box and fed from a stabilized power supply. Filters: standard FITc filter mirror combination with 6 mm BG 38 filters in excitation light path. Objective is a 6.3 \times Neoflur.

Table 1 (Contd)

DATA ANALYSIS		Comments
raw data	cpm of duplicates recorded	Fluorometer readings before and after incubation are recorded. The value of each individual well (i.e. = percentage fluorescence) in the final reading is divided by the value of the pre-reading in the corresponding well. This ratio expressed as a percentage is used for calculation of % CML. (Le1 2)
utilization of dilution data		
linear regression analysis r values		
slope determination		
calculation of % CML	$\frac{\text{mean ER} - \text{mean SR}}{\text{mean MR} - \text{mean SR}}$	$\left(1 - \frac{\text{mean test}}{\text{mean control}} \right) \times 100 \text{ (Le1 2)}$
lytic unit size		
lytic units recovered/culture		
assignment of responses	r^2 greater than or equal to 0.90 (based on regression analysis). From this line the value of greater than or equal to 5% CML at a CTL Target Cell of 30:1 (Mu)	From dose response curves starting with an effector:target ratio 50:100:1. Positive: A minimum of two observations on the curve above 10%. Negative: All observations on the curve are less than 10%. Intermediary: Curves with only one observation above 10% (Ar). Specific ^{51}Cr release percentage $\leq 10\%$ = neg, 11–15% weakly pos, 16–40% pos $\geq 41\%$ strongly pos. (Le1 1) Specific release percentage $\leq 10\%$ = neg, 11–15% weakly pos, 16–40% pos

After a 6 day sensitization period, the supernatant in each well was aspirated, with a specially designed aspiration unit (Skatron A/S, Lierbyen, Norway) to a fixed level with 0.1 ml remaining in each well 2×10^3 ^{51}Cr -labeled target cells (see workshop) were added to the cultures, the force of the jet resuspending the cell pellet at the bottom of the well. The plates were incubated for an additional 6 h and the supernatants collected using a Skatron multiple supernatant collection system (Skatron A/S, Lierbyen, Norway). The effector to target cell ratios were assessed from the number of effector cells added to each well on day 0 and the number of targets (2×10^3) added on day 6 (E/T usually 50/1 — 6.25/1), % cytotoxicity was calculated as in the workshop, using spontaneous release (targets + supplemented medium) and maximum release (targets + 5% Triton X) harvested as above after 6 h of incubation.

3 — Carboxyfluorescein labeling of target cells

This technique was performed by Leiden. Details on target cell labeling have been published elsewhere (Bruning et al 1980).

Briefly, target cells were labeled by incubation in a 37°C waterbath for 15 min in a carboxyfluorescein-diacetate solution in RPMI (final concentration $40 \mu\text{g/ml}$). After washing with RPMI-serum, 4000 target cells were added to the wells of Terasaki type microtest plates and fixed to the bottom by centrifugation. Effectors were added and the fluorescence per well read in an automated microfluorometer. After a 4 h incubation the released carboxyfluorescein was removed by washing and fluorescence of the remaining targets determined in the fluorometer. From the difference of fluorescence between

test and control wells the degree of cytotoxicity could be calculated.

All other procedures were performed using the SE CML (Table 1).

Common Reagents used in Workshop III

1 *PBL* Unsensitized PBL from four selected blood donors (1, 2, 3, 4) were collected, frozen and shipped from Aarhus. The donors had been fully HLA-A, B, C, D/DR and red cell antigen phenotyped and were selected in such a way that CTLs raised among them, in a checkerboard fashion, would yield cytolysis both explicable and inexplicable from our present knowledge of HLA (Kristensen et al 1974, 1976) Kristensen & Grunnet 1975). These cells were also those tested during CML-Workshop II (Bradley et al 1980).

2 *Primed PBL* (1/2x, 1/3x, 1/4x, 4/1x, 4/2x, 4/3x) Cells primed *in vitro* in the Aarhus laboratory, according to the standard technique (Table 1), for 10 days were frozen and shipped to the other centers.

3 *Human serum* Pooled AB serum was prepared from four male blood donors, screened as negative for lymphocytotoxic and red cell antibodies (A gift from Aarhus).

4 *PHA M* A common batch produced by Difco Co (0528-57) was used by all centers.

5 *Radioisotope* ^{51}Cr as $\text{Na}_2^{51}\text{CrO}_4$ (500 mCi/mg Cr, 5 mCi/ml) from the same production date was provided as a gift by Amersham (Amersham Co, England).

6 *Culture medium* RPMI 1640 (Gibco Co) supplemented with L-glutamine from the same production lot was provided as a gift from Grand Island Biological Company (in France, Flobio, S A)

Format of CML Workshop III

The format of the practical part of the Workshop is given in Table 2. PBL were collected in Aarhus and shipped on dry ice as either unprimed or primed (10 days) cells. Primary cultures were established on day 0 followed by secondary cultures on day 1. Secondary cultures were restimulated either specifically (e.g. $1/2x \leftarrow 2x$) or unspecifically by PWM. Target cell cultures were established on day 3 and stimulated with PHA. Cytotoxicity was measured on day 6 using the same target cell cultures for both 1° and 2° CTLs.

For practical reasons, growth with conditioned media was initiated with the remaining CTL on day 6, hence requiring CML testing on different target cell cultures.

Results

I General Experiences

The adoption of these techniques by the participating laboratories required changes for each group, although these were variable from laboratory to laboratory, and such changes led to individual experiences and difficulties in the III European CML Workshop.

The following outlines the general assessment of the collaborating groups concerning their experiences with the standard protocol.

A Recovery of cells after thawing of shipped standard reagents

1 *Unprimed cells* No standard technique

was designated for the thawing of fresh cells. Despite variations in technique, however, uniformly high recoveries of cells with high viabilities were found by all groups (Table 3). It was felt by the various groups that the method of cryopreservation used in Aarhus was excellent and that the thawing was not detrimental to the cells.

2 *Primed cells* In contrast to the success with the unprimed cells, all groups, except Aarhus, had relatively low recoveries of cells in the shipment of primed cells. It might be overcome, in future tests of 2° CML, by performing the sensitizations in local centers rather than one laboratory providing cells. An alternative possibility to be considered is variation in the local procedures determining recovery and viability of cryopreserved cells.

B Determination of recovery and viability of cells

1 *Trypan blue or eosin dye exclusion* It was felt by most groups that utilization of such dyes which are excluded by viable cells is a useful criterion for assessment of actual cell recovery, but that it is not an accurate assessment of functional recovery since only a subpopulation of viable cells may be involved in the lytic process.

2 *Phase contrast microscopy* An alternative means of distinguishing live from dead cells is their examination by phase contrast microscopy. Again, this may be an accurate means for assessment of viability, but cannot be used to ascertain the recovered functional capacity.

3 *Lytic unit assessment* of functional activity based on linear regression analysis.

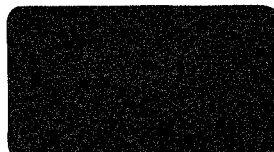


Table 2
FORMAT OF CML WORKSHOP III

		Day 0	Day 1	Day 3	Day 6	Day 17	Day 20		
PREPARATION OF CELLS FOR AARHUS SHIPMENT	{ All Labs Aa Ma Ma	1°MLC - - 2°MLC - - TC/PHA - - CML*					Submission of data to Marseille	Third European CML Workshop meeting Marseille	
		2°MLC - - - - - CML†							
		CTL-CM - - TC/PHA - - CML§					Central analysis Marseille/Aarhus		

*Unsensitized PBL were thawed and primary (1°) MLC combinations were established on day 0. Primed cells, received from Aarhus, were thawed and restimulated in secondary (2°) MLC cultures on day 1. Target cell (TC) ampules were thawed on day 3 and stimulated with PHA. Cytotoxic from 1° and 2° CML were tested on day 6 in the ⁵¹Cr-cell-mediated lympholysis (MCL) assay.

†The laboratories of Aarhus and Marseille thawed additional aliquots of primed cells which were restimulated with pokeweed mitogen (PWM) and tested in CML on day 6.

§The laboratories of Marseille and Munich initiated cytotoxic cells obtained from 1° and 2° MLC with conditioned medium (CM) containing T cell growth factor. These cells were expanded in culture for 14 days and tested in CML with target cells prepared 72 h in advance

Table 3a
Recovery of cells after thawing of shipped standard reagents (a) Unprimed cells

	Donor Number	Ar	Lei 1	Lei 2	Ma	Mu
% Viable	1					94
	2	N D	> 90%	> 90%	> 95%	93.5
	3					93
	4					93
Recovery [†]	1	7.3	7.0	6.8	9.1	6.1
	2	9.5	8.0	6.8	6.3	8.0
	3	8.2	8.2	8.0	8.3	8.0
	4	7.9	6.4	7.3	8.2	6.8

N D, Not done

[†]The concentration per frozen vial was $\approx 10 \times 10^6$ cells in 1 ml. Results are expressed as the average number $\times 10^{-6}$ of cells recovered per thawed vial

of CML activity prior to and following cryopreservation. Recovery of functional activity in cryopreserved samples can be defined by comparing the lytic unit for CTL samples prior to and following cryopreservation.

C Recovery of cells after MLC sensitization in the local laboratories

1. *1° MLC sensitization* The starting concentration of responding and stimulating cells was stipulated in the standard protocol, but culture vessels to be utilized were left to individual judgment after determination of cell recovery from the cryopreserved reagents. This led then to variation in the standard technique, with some laboratories culturing between 5 and 10×10^6 responding cells in upright standing culture flasks, with others culturing at 1×10^6 responding cells per well of a Cluster macrotiter culture plate (Costar), and still others using a combination of these two vessels. In terms of final recovery after the 6-day sensitization period, however, the results were remarkably uniform (Table 4a). Cell recovery from Oslo could not be determined because

of their variation from the standard technique.

2. *2° MLC sensitization* In spite of the recoveries of primed cells from the cryopreserved ampoules, all groups observed percentage yields after specific restimulation (Table 4b).

3. *Pokeweed mitogen restimulation of primed cytotoxic cells* As an alternative restimulation of primed cytotoxic cells with specific γ -irradiated stimulating cells, several laboratories received additional numbers of primed cells for restimulation with PWM. Two variations in this protocol were used by Aarhus and Marseille. Aarhus first used PWM at 1% final concentration with stimulation for 4 days (Mawardi, 1975). Marseille used 1% PWM with addition of autologous, γ -irradiated stimulating cells (Charmot et al. 1980). In both cases high yields of cytotoxic cells with functional activity were recovered.

4. *Expansion of CTL by growth in the presence of Conditioned Medium* (CM). Preliminary results were also obtained

Table 3b
(b) *Primed cells*

CML COMBINATION	%V	Ar Rec	%Rec	%V	Lei 1 Rec	%Rec	%V	Lei 2 Rec	%Rec	%V	Ma Rec	%Rec	%V	Mu Rec	%Rec	%V	Os Rec	%Rec
1/2x	¹	4.9 ²	54 ³	Poor	1.3	14	Poor	2.9	32	80	3.2	36	75	2.1	21	22	2.5	28
1/3x		6.4	64		0.9	9		3.1	31	15	7	7	73	2.1	22	20	2.3	23
1/4x		5.2	52		1.7	17		2.4	24	50	2.4	24	85	1.9	20	20	2.1	21
4/1x		5.3	59		3.6	40		5.1	57	80	4.4	49	88	2.6	26	26	2.6	29
4/2x		4.6	58		2.7	34		4.7	59	80	2.9	36	82	1.8	18	34	2.6	33
4/3x		5.2	61		2.7	32		3.7	44	80	3.9	46	83	2.0	20	27	1.5	18

¹ Viability not assessed with vital dye exclusion.

² Results expressed as number of cells (10^{-6}).

³ Calculated on the basis of the number of cultured small lymphocytes frozen in Aarhus ($8-10 \times 10^6$).

Table 4a
Recovery of cells after MLC sensitization in local laboratories (a) Primary MLR

TEST COMBINATION	%V	Ar Rec	%Rec	%V	Le1 1 Rec	%Rec	%V	Le1 2 Rec	%Rec	%V	Ma Rec	%Rec	%V	Mu Rec	%Rec	Os Os
1/2x	NT	18 0 ¹	180 ²	good	25 4	254	good	19 2	148	> 90	20 0	167	92	19 8	264	NO DATA ³
1/3x		20 3	203		16 9	169		13 9	126		19 8	165	95	17 1	228	
1/4x		21 1	211		13 6	136		20 3	185		37 6	313	90	20 7	276	
4/1x		15 9	159		13 6	136		13 7	125		19 0	158	77	13 8	184	
4/2x		14 8	148		12 6	126		10 1	101		15 2	127	73	13 7	144	
4/3x		15 6	156		9 3	93		11 5	105		12 4	103	74	13 3	140	
		(Flask)			(Flask)			(Flask and cluster)			(Cluster)					

¹ Results expressed as number of cells ($\times 10^6$)

² Based on the initial number of responder cells = 100%

³ Due to a different technique

NT = not tested

Table 4b
Secondary MLR

TEST COMBINATION	%V	Ar Rec	%Rec	%V	Lei 1 Rec	%Rec	%V	Lei 2 Rec	%Rec	%V	Ma Rec	%Rec	%V	Mu Rec	%Rec	Os
1/2x	NT	21.6 ¹	309 ²	Many	2.4	185		9.1	314		NT ³		64	7.6	362	NO DATA ⁴
1/3x		27.8	397	dead	2.0	222		10.3	332				75	9.5	452	
1/4x		23.1	330	cells	3.0	176		10.2	425				70	9.6	500	
4/1x		27.8	397		5.6	156	8.	8.2	161				29	2.6	100	
4/2x		26.1	373		11.6	430		10.5	223				69	9.1	500	
4/3x		33.8	483		14.5	537		9.3	251				72	5.2	260	
		(Flask)			(Cluster)			(Cluster)						(Flask)		

¹ Number of cells.

² Based on the input of responder cells = 100%.

³ Marseille used PWM expansion only.

⁴ Different technique.

Table 4c
PWM expansion of primed cells

TEST COMBINATION	NT	Ar Rec	%Rec	%V	Lei 1 Rec	%Rec	%V	Lei 2 Rec	%Rec	%V	Ma Rec	%Rec	%V	Mu [†] Rec	%Rec	%V	Os Rec	%Rec
1/2x	%V	2.5	83		NT			NT		> 90	13.0	406	80	4.3	204		NT	
1/3x		3.2	107								9.0	1286	60	2.9	138			
1/4x		3.0	100								13.0	542	32	2.6	137			
4/1x		3.8	127								19.4	441	50	1.3	50			
4/2x		3.4	113								12.6	434	58	5.3	294			
4/3x		3.4	113								23.4	600	21	1.4	70			
		Flask									Flask			Flask				
		No autologous feeder									autologous feeder			no autologous feeder				
											(-irradiated 3500R)							

Same legend as in (a) and (b).

[†]Cultures infected.

two laboratories testing the method of continued growth of CTL and their expansion by T cell growth factors present in conditioned medium. CTL obtained from both primary and secondary cultures MLC were tested after 14 days of culture in the presence of CM. CTL with very strong activity could be obtained although cytotoxic activity was not found in all combinations that were successfully expanded with TCGF (data not shown).

D. Target cells

1. *Ficoll flotation after ^{51}Cr labeling.* According to the standard protocol, labeled target cell preparations were to be layered over ficoll to separate live from dead and clumped cells. It was felt that this technique might reduce the level of spontaneous release of ^{51}Cr . The experience in this Workshop was that this procedure consumes time, and apparently did not enhance the results. The laboratories uniformly reported presence of very few clumped or dead cells in the starting populations. It was noted by some groups, however, that according to individual experience this is a valuable approach when the starting populations contain considerable numbers of dead cells. Some laboratories noted that no cells passed through the ficoll interface and suggested that this may have been due to the PHA agglutination of the target cells. It was suggested that this problem might be overcome by first washing the target cells in medium before layering over ficoll gradients. It was questioned whether DNase would be beneficial for the reduction of spontaneous release values and the elimination of clumped cells. In the experience of Aarhus this was not observed.

2. *Triton X determination of maximum release of ^{51}Cr .* According to the standard protocol, maximum release was determined by detergent lysis of target cells. Several laboratories had difficulty with this procedure, failing to obtain solutions of lysed target cells, instead they obtained colloidal suspensions. In discussion it was ascertained that this was due to the use of 100% Triton X instead of a 5% solution of Triton X in H_2O . In most laboratories, very little variation between the values of maximum release as determined by Triton X detergent lysis and total cpm of ^{51}Cr was found on a per cell basis. Thus, it appears that MR values must not be determined and calculations can be based on total cpm. These results indicate that there is not an increment of ^{51}Cr counts that is non-releasable from target cells, as has been suggested previously. This observation, however, may be dependent upon the method used for the determination of MR.

E. Harvesting of supernatants from microtiter wells.

The utilization of V-bottomed plates in the standard protocol led to some difficulties by various groups regarding means for harvesting standard aliquots of supernatant. Some local methods utilized commercial filter systems for round-bottomed plates or testing in larger volumes allowing supernatants to be decanted after centrifugation. With the V-bottomed plates there was a dilemma regarding a suitable means of supernatant collection. Some groups noted harvesting by hand to be tedious and time consuming and potentially increasing spontaneous release. Leiden 1 successfully used the commercial filter systems in the V-bottomed plates while Marseille used a locally designed multiple harvester which has 96 lcc syringes set into a plexiglass

frame which could simultaneously harvest the samples of 0.1 ml/well of an entire plate

F Alternative CML techniques

In addition to the standard protocol used by four participating groups, two additional approaches were tested with the same reagents and could be compared in Workshop III

1 Generation of CTL in microtiter plates

As presented later in the section on data analysis, this approach led to variance in the results compared to those of the standard protocol. The technique did, however, reveal that quantitative evaluation of CML activity could be made with few numbers of starting cells. It was felt that development of this approach may be beneficial in the clinical situation when tests of young children, dialysed, transplanted, aplastic, or immunodeficient patients must be performed.

2 Assessment of cytotoxic activity using

carboxyfluorescein as a viability label The second technique tested in parallel with the standard protocol was that of utilizing the viability dye, carboxyfluorescein, instead of ^{51}Cr for the determination of destruction of target cells. Basically, in this technique (Bruning et al 1980), target cells are labeled by incubation with the diacetate derivative which is taken up only by viable cells, after which the dye is liberated in the cell by intracellular hydrolysis. The target cells are fixed by centrifugation to the well bottom of Terasaki type microtest plates. Effector cells are added to the targets and total well fluorescence is determined before the incubation period using an inverted epi-illuminated fluorescence microscope with photomultiplier and a

stage driven by stepping motors. Assembly is controlled by a mini-computer. After incubation with CTLs for a period the released dye is removed and residual fluorescence is measured on an automated fluorometer. Loss of fluorescence is then used as a measure of lysis of the target cells. Comparable specificity was found using this new technique (cf. below). It was felt that great potential lies in the utilization of this approach in the future if the equipment becomes commercially available. It eliminates the requirement for radioactive material, automates the reading of results, reduces normal harvesting and counting times from 1 to approximately 30 sec, for 60 samples. Comparison of quantitation and specificity with the Workshop standard protocol is presented later.

II Data Analysis

Data were submitted from each individual laboratory in three forms: (1) raw data obtained with each CTL tested against different target cells at each of six dilutions (Oslo, submitting on fewer dilutions); (2) data calculated and expressed as percentages according to formulas of individual laboratories listed in Table 1; and (3) qualitative assignment of positive and negative responses according to judgment.

Joint analysis was performed in Marburg and Aarhus and is outlined below.

A Preliminary calculations

1. All submitted data were graphically plotted to assess the (curvi)-linear relationship between the independent variable (E/T ratio) and the dependent one (relative percentages). For most positive combinations an S-shaped curvi-linear relationship was observed. A priori it is expected

logarithmic transformation of the X-values (i.e. E/T ratios) should yield a linear relationship. Consequently, to simplify subsequent regression analysis E/T ratios were transformed to \log_{10} . In most instances this changed a curvilinear relationship into a linear one, at the same time increasing the proportion of variance of the dependent variable explained by the X-value (i.e. \log_{10} E/T), and making the distribution of the deviations from the regression line fit the normal. Assessment of data using square-root and reciprocal transformations both on X- and/or Y-axis values was unsuccessful.

2 Based on data transformed according to (1), LU_{30} and release percentages at E/T ratio = 30 were extra- or interpolated.

3 For the analysis of variance, release percentages were \log_{10} transformed.

B Main analysis

1. Analysis of Variance (Anova) was made based on (a) \log_{10} release percentages at E/T ratio 30/1, (b) LU_{30} (Lytic units i.e. the number of effectors needed to obtain 30% target cell lysis). Anova was performed to identify the variance components and their interactions both within and among laboratories.

2 Linear regression analysis of release percentages with varying \log_{10} E/T ratios were used to define the regression equations obtained by different laboratories when testing supposedly identical CTLs on identical targets.

Using least square methods, straight lines were approximated and defined by the regression equation

$$\text{Release \%} = a + b \log_{10} \text{ E/T}$$

(where b = the regression coefficient = the derivative of the function, a = the intercept

of the regression line with the Y axis = the release % at E/T = 0)

The statistical significance of b was ascertained using a t-test under the null hypothesis that b = 0.

3 Analysis of Covariance (Ancova) was used to compare the regression equations obtained within and among different laboratories using supposedly identical CTLs on identical targets.

4 2×2 comparison analyses were used to compare qualitative assignments made locally and jointly.

For basic references to the statistics used, cf Sokal & Rohlf (1969) and Bradley et al (1980).

For obvious reasons it is impossible to present all data and calculations performed in the joint analysis. The data given are thus summaries. *A total set of data may be obtained from the group.*

Table 5
Comparison of target cell data W III

	1*	2*	3*	4*	5*
Aarhus					
SR	1434	1945	1021	2690	971
MR	5813	6385	4640	8137	4466
Leiden 1					
SR	1356	1132	769	1146	1065
MR	5534	6198	4516	4657	7169
Marseille					
SR	483	600	440	1118	613
MR	6281	3944	4080	8393	6453
Munich					
SR	582	323	173	789	235
MR	3015	3580	2775	4176	2474
Oslo					
SR	526	580	572	810	564
MR	2359	2936	3662	4139	3371

SR = Spontaneous release cpm (mean of triplicates)

MR = Maximum release cpm (mean of triplicates)

Results of Joint Analysis

1 Raw data — cpm

One of the major findings from Workshop II was that there were marked differences in the ways that different laboratories handled identical target cells, as opposed to their handling of effector cells. Reasons for these differences might be the use of differential ^{51}Cr products, including different batches, labeling amounts and labeling procedures. In Workshop III these parameters were standardized and the results in terms of spontaneous release and maximum release cpm are given in Table 5.

When comparing the cpm among laboratories and targets considerable variance is observed. However, when the cpm concerning the duplicate targets (3 and 5) are compared within a given laboratory a high degree of reproducibility is found. Consequently, these findings may, at least partly, be attributed to differences between laboratories in the assessment of spontaneous and maximum releases, as well as harvesting and counting procedures (cf Table 1).

2 Analysis of variance (Anova)

Both release percentages at $E/T = 30$ and LU_{30} were subjected to Anova in which the following questions were asked (cf Table 6)

(a) How big is the variance component at the different levels?

(b) Was there a significant difference in the way in which different laboratories handled their effector cells?

(c) Was there a significant difference in the way in which different laboratories handled their target cells?

(d) Was there a significant difference in the way in which effector cells interact

with the targets? This of course was expected since it reflected the biological variance in the CML.

(e) Were any laboratories adding, relatively more to the total variance as compared to others? This was evaluated by reperforming the analysis following successive exclusion of individual laboratories.

From Table 6 (where the data are given in \log_{10} release percentages) it can be seen that the major significant variance found with the Workshop III technique is that from the CTL and target interaction, i.e. the biological variance. In addition there is variance in the way different laboratories treat target CTLs. The latter (laboratory versus target) can be attributed mainly to one laboratory (Oslo), which used a different technique (cf Table 1 and above). The variance by the laboratory and target interaction could be attributed to Leiden which used the CF-technique, and Mars. However, the "target cell problem" disclosed itself in all centers, in as much as the variance components including target component were always the largest.

Due to the overall significance of the target interaction variance no main effect variance could be tested for statistical significance. Anova-calculations performed with 10% yielded fully comparable variance components and significance values. This furnished no new information (data not shown). Comparing the overall variance components of Workshop III to the variance components obtained using local techniques during Workshop II (Bradley et al 1981) is, however, quite evident that much more variance has accumulated in the CTL-target interaction group, pointing to a more uniform and stable technique which more clearly discloses the biological interaction.

Table 6

CML-Workshop III combined analysis — three-way analysis of variance. Log. 10-transformed release percentages

Source of variation	Variance components (significance)						
	ALL	-Ar	-Lei 1	-Lei 2	-Ma	-Mu	-Os
Laboratories	3.2% (—)	1.6% (—)	2.7% (—)	3.5% (—)	3.1% (—)	2.0% (—)	1.0% (ns)
CTLs	2.4% (—)	3.1% (—)	2.5% (—)	3.6% (—)	2.7% (—)	3.8% (—)	2.8% (—)
Targets	2.8% (—)	2.7% (—)	4.2% (—)	3.6% (—)	3.3% (—)	2.2% (—)	2.9% (—)
LABS & CTLs	4.7% (●●)	6.2% (●●)	5.6% (●●)	3.1% (●●)	2.4% (●)	4.7% (●●)	1.5% (ns)
LABS & Targets	8.0% (●●)	7.1% (●●)	7.2% (●●)	6.4% (●)	6.1% (●)	9.6% (●●)	3.3% (●)
CTLs & Targets	70.0% (●●●)	69.4% (●●●)	70.4% (●●●)	68.9% (●●●)	72.3% (●●●)	69.1% (●●●)	76.6% (●●●)
LABS & CTLs & Targets	8.5% (—)	9.9% (—)	7.4% (—)	10.9% (—)	11.3% (—)	8.6% (—)	11.9% (—)

● : $P \leq 5\%$

●● : $P \leq 1\%$

●●● : $P \leq .1\%$

ns : not significant

— : cannot be tested.

3. Analysis of covariance (Ancova)

This analysis was applied in order to compare the regression equations obtained both within and among laboratories when identical CTLs were tested against identical targets in either 1° or 2° CML, i.e. whether the biology of the different tests was identical (The data from Oslo were excluded from this analysis).

The following analyses were made:

(1) within laboratories on duplicate targets (3 and 5);

(2) within laboratories in 1° and 2° CML;

(3) among laboratories with regard to specific CTL-target combinations in either 1° or 2° CML.

The results were uniformly disappointing, showing differences significant at the 1% level.

Thus, even with standardized reagents

using a standardized technique, a single laboratory is not able to fully describe the biological interaction and hence the immunology of CML.

It is easy to speculate that during freezing and thawing, MLC incubation or target cell proliferation, not all active clones survive and proliferate equally, but rather that some clones are already lost from the lymphocyte population prior to culture in a stochastic fashion varying from laboratory to laboratory. Since, however, major specificities are maintained and almost uniform positive and negative assignments can be obtained (cf. below) important (high affinity ?) clones must be selected actively during CTL sensitization whereas peripheral (low affinity ?) clones undergo stochastic selection. During target-cell culture with PHA no active selection should occur leading to more variability (the target cell

problem) These findings and speculations underline some of the difficulties which must be considered when CTLs are to be used as immunogenetic typing reagents

4 Joint analysis based on linear regression analysis and positive slope assignment

(a) *Basis for linear regression* Plotting percent release versus dilutions of CTL gives an S-shaped curve. When the dilution of CTL (E/T) is transformed to \log_{10} value a straight line should be generated. Using least square methods, straight lines can be approximated for these values of X and release percentages for the values of Y. If these have a significantly positive slope, tested as described above, the information can be used to quantify the CML response. If too few dilutions are compared the possibility to assign a significant b value is decreased (cf. above).

(b) Slope of the regression line may be used to assess the presence and subsequently the strength of a cytotoxic response. For positive CTLs it is expected that the slopes will be positive, however, some laboratories observed no (or negative) slopes, the basis of which is discussed below.

(c) Combined assignment based solely on a significantly positive slope was then used to judge the data of all laboratories. The variations between the first local assignments and these new assignments were discussed and each laboratory detailed its own exceptions and the basis for original local assignment.

5 On the basis of these discussions the problems below were encountered

(a) In some cases significance by positive slope may not be obtained if too few dilutions of CTL are tested. In the standard protocol for this experiment the use of six dilutions in duplicate was suggested.

(b) A positive slope may also fail to reach significance when cytotoxic activity reaches a plateau with most of the dilutions or, in the case of Leiden 2, when cytotoxic activity decreases with the highest E ratios. Similarly there may be a plateau at the lowest level if the first dilutions do not exceed the detection level. This could then cause false elimination of positive reactions.

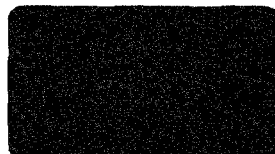
(c) Negative slopes were observed for some combinations of CTL against the autologous target cells. The basis for these observations was discussed. First, it might be due to a stabilizing effect of increasing numbers of CTL on the spontaneous release of ^{51}Cr from the target cells. This then leads to the dilemma of the most suitable control for spontaneous release (i.e. AB_x on A, unsensitized A, AA_x on A or, as was commonly used here, A in medium alone). The latter has been conventionally adopted by many laboratories to save on reagents.

(d) Alternatively, positive slopes may also be observed when the CTL is tested with its autologous target cell. Such activities should not automatically be ignored or ascribed merely to technical artefacts. They may reflect recognition of antigens of lymphocyte subpopulations or perhaps of anti-idiotypic responses.

6 Conclusion of the Joint Data Analysis Using Regression and Slope Data

(a) Positive assignments cannot be made solely on the basis of significantly positive slopes. Not all reactions are transformed to linearity by the logarithmic approach. Some difficulties may be solved by using a minimum number of E/T dilutions (greater or equal to five).

(b) Since these approaches may not work



is necessary to accept a release percentage value greater than or equal to 10% at any dilution as indicating a positive response. This perhaps can be more safely assumed if duplicate CTL cultures are also tested.

(c) For combinations tested at very low dilutions showing a slope not significantly positive extrapolations should be made to E/T ratios = 50. A release percentage greater than or equal to 10% should then be considered positive.

(d) The observation of a positive slope cannot always be taken as a means of determination since immunologically negative controls may show significantly positive slopes (cf 5 d).

Workshop Criteria for Assignments and Exclusions

As a consequence of the above mentioned

experience and discussion the following criteria were accepted for the joint assignments and exclusions

A Positive assignments were based on

1 Significantly positive slope of regression lines of release percentages on \log_{10} E/T, and/or

2 Release percentage > 10% in any dilution tested. If a CTL was tested only at E/T dilutions below 50/1 extrapolation was performed. An extrapolated release percentage > 10% at E/T = 50/1 was then considered positive.

B Exclusions were based on

1 A positive assignment (cf above) on autologous targets (e.g. 1/2x → 1)

2 A negative assignment (cf above) on the specific target (e.g. 1/2x → 2)

Table 7
CML Workshop III — combined assignments

CTLs	TARGET 1					TARGET 2					TARGET 3					TARGET 4					TARGET 5					
	A	L1	L2	Ma	M	O	A	L1	L2	Ma	M	O	A	L1	L2	Ma	M	O	A	L1	L2	Ma	M	O		
1° 1/2x	-	-	-	-	-	-	+	+	+	+	+	+	+	+	+	+	+	-	+	+	+	-	+	-	+	+
2° 1/2x	-	-	-	-	-	/	+	+	+	+	+	/	+	+	+	+	+	/	+	+	+	+	+	/	+	+
1° 1/3x	-	-	-	-	-	/	+	+	+	+	+	/	+	+	+	+	+	/	+	+	+	+	+	/	+	+
2° 1/3x	-	/	-	-	-	/	+	/	+	+	+	/	+	/	+	+	+	/	+	/	+	+	+	/	+	+
1° 1/4x	-	/	-	-	-	/	+	/	+	+	+	/	+	/	+	+	+	/	+	/	+	+	+	/	+	+
2° 1/4x	-	/	-	-	-	-	+	/	+	+	+	+	+	+	/	+	+	+	+	+	/	+	+	+	+	-
1° 4/1x	+	+	+	/	+	+	-	+	+	/	+	-	-	-	-	/	-	-	-	-	-	/	-	-	-	-
2° 4/1x	+	+	+	+	/	+	-	+	-	+	/	-	-	-	-	/	-	-	-	-	/	-	-	-	/	-
1° 4/2x	+	+	+	-	+	+	+	+	+	+	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
2° 4/2x	/	+	+	+	+	-	/	+	+	+	+	+	/	-	-	-	-	-	/	-	-	-	-	-	/	-
1° 4/3x	-	+	-	+	+	+	-	-	+	+	+	-	+	+	+	+	+	+	-	-	-	-	-	-	+	+
2° 4/3x	-	-	/	-	-	/	-	-	/	-	-	/	+	+	/	+	+	/	-	-	/	-	-	/	+	+

+ = | positive assignment |

- = | negative assignment |

/ = | excluded |

Table 8a
CML-Workshop III - 1° CML 2 × 2 analysis of
combined assignments

	++	+-	-+	--	N
Ar vs Lei 1	13	0	2	10	25
Lei 2	17	0	2	11	30
Ma	13	3	2	7	25
Mu	16	0	3	11	30
Os	5	4	1	10	20
Lei 1 vs Lei 2	14	1	1	9	25
Ma	10	3	1	6	20
Mu	15	0	1	9	25
Os	6	5	0	9	20
Lei 2 vs Ma	14	3	1	7	20
Mu	19	0	1	10	30
Os	5	6	1	8	20
Ma vs Mu	15	0	3	7	25
Os	4	3	1	7	15
Mu vs os	6	6	0	8	20
Excluded CTLs : Ar	: 1				
Lei 1	: 3				
Lei 2	: 1				
Ma	: 1				
Mu	: 1				
Os	: 3				

3. Direct indication from the laboratory in question (infection, not enough cells, technical failure, . . .).

2 × 2 Table Analysis of Combined
Assignments

Based on the assignment criteria outlined above, all local assignments were reassessed leading to exclusion of some CTLs due to positive and/or negative control values, while others (locally excluded) were accepted into the analysis. The combined assignments are shown in Table 7.

Based on these assignments, 2 × 2 tables were constructed as shown in Table 8.

The conclusions are as follows:

- 1° CML : good correlations, with Munich showing the broadest assignment of positive values (Table 8).

2. 2° CML : very few discrepancies except with Oslo showing few positive reactions (Table 8b).

3. 1° versus 2° CML (Table 8c) : excellent agreement. There may be a tendency to obtain more positives in the 1° CML than 2°, except for Marseille.

Marseille as standard uses a 2° CML so may be better at this technique.

4. CFA versus ⁵¹Cr : no major discrepancies with all other groups except Os in both 1° and 2° CML (Table 8d).

5. 2 × 2 analysis of reactions of all CTL against duplicate targets : excellent agreement (Table 8e).

Table 8b
CML-Workshop III - 2° CML 2 × 2 analysis
combined assignments

	++	+-	-+	--	N
Ar vs Lei 1	7	0	1	7	15
Lei 2	13	0	0	12	25
Ma	15	0	1	9	25
Mu	14	0	0	6	20
Os	4	1	0	5	10
Lei 1 vs Lei 2	7	1	0	7	15
Ma	10	0	0	10	20
Mu	8	0	0	7	15
Os	2	2	0	6	10
Lei 2 vs Ma	15	0	1	9	25
Mu	14	0	0	6	20
Os	5	2	0	8	15
Ma vs Mu	16	0	0	9	25
Os	5	3	0	7	15
Mu vs Os	4	2	0	4	10
Excluded CTLs : Ar	: 1				
Lei 1	: 3				
Lei 2	: 1				
Ma	: 1				
Mu	: 1				
Os	: 3				

•The result from Ma is not directly comparable as 2° CML was performed only using expansion of 1° CTLs with PWM and irradiated, autologous feeder cells. No specific restimulation was performed.

Table 8c

* CML-Workshop III - 1° vs 2° CML 2 × 2 analysis of combined assignments

	++	+-	-+	--	N
Ar vs Ar	15	0	0	10	25
Lei 1 vs Lei 1	10	1	0	9	20
Lei 2 vs Lei 2	15	1	0	9	25
Ma vs Ma	13	2	3	7	25
Mu vs Mu	16	2	0	7	25
Os vs Os	2	1	0	7	10
TOTAL	71	7	3	49	130

6. Specific 2° restimulation versus PWM : (fair agreement). Data not shown since available only from Aarhus and Marseille.

On the Specifications Detected by Workshop III CTLs

The cells circulated in this Workshop were identical to those circulated during Workshop II. An account of their total red and white cell phenotypes as defined by serology is given in the report from Workshop II (Bradley et al. 1980).

In terms of immunogenetic specificity

Table 8d

CML-Workshop III - CFA vs ⁵¹Cr targets 2 × 2 analysis of combined assignments

	++	+-	-+	--	N
1° CML					
Lei 2 vs Ar	17	2	0	11	30
Lei 1	14	1	1	9	25
Ma	14	3	1	7	25
Mu	19	0	1	10	30
Os	5	6	1	8	20
2° CML					
Lei 2 vs Ar	7	1	0	7	15
Lei 1	7	0	1	7	15
Ma	15	0	1	9	25
Mu	14	0	0	6	20
Os	5	2	0	8	15
TOTAL	117	15	6	82	220

Table 8e

CML-Workshop III - duplicate targets (3 vs 5) 2 × 2 analysis of combined assignments

	++	+-	-+	--	N
1° CML					
Ar	4	0	0	2	6
Lei 1	3	0	0	2	5
Lei 2	4	0	0	2	6
Ma	3	1	0	1	5
Mu	3				
Os	0	1	0	3	4
2° CML					
Ar	4	0	0	1	5
Lei 1	2	0	0	2	4
Lei 2	3	0	0	2	5
Mu	4	0	0	1	5
Os	0	1	0	2	3
TOTAL	36	3	0	22	60

the data of Workshop II were translated as follows:

1. Positive lysis was strongly correlated to sharing of HLA-A, B, C, DR serologically defined determinants between stimulator and target cells; however,

2. No lysis was observed in the presence of HLA-antigenic sharing between stimulator and target (five combinations).

3. Two combinations (4/1x → 2 and 4/2x → 1) gave positive CML in the absence of any known serologically defined antigen. This is in concordance with earlier findings (e.g. Kristensen et al. 1974).

In this Workshop III only the combinations from Workshop II 1/2x, 1/3x, 1/4x, 4/1x, 4/2x and 4/3x were repeated with the standard technique. The conclusions (1) and (3) above are clearly confirmed, while (2) is invalidated in as much as the one previously negative combination (1/3x → 2) retested is uniformly positive with the Workshop III standard technique (cf. Table 7). The

Table 9
CTL detecting new specificities activity in CML Workshop III

1° CML of 4/1x and 4/2x with 1* and 2*				
4/1x Laboratory	b(1*)	Slope (1*)	b(2*)	Slope(2*)
Aarhus	0.96 ^{○○}	9.15	0.68	3.30
Leiden-1	0.96 ^{○○}	17.02	0.96 ^{○○}	9.40
Leiden-2	0.93 ^{○○}	15.11	0.68	9.10
Munich	0.89 ^{○○}	10.57	0.98 ^{○○}	5.37
Oslo	0.98 ^{○○}	19.51	0.93 ^{○○}	6.78
4/2x Laboratory	b(1*)	Slope (2*)	b(2*)	Slope (1*)
Aarhus	0.90 [○]	5.68	0.97 ^{○○}	5.81
Leiden-1	0.96 ^{○○}	11.67	0.97 ^{○○}	12.10
Leiden-2	0.96 ^{○○}	16.12	0.82 [○]	8.44
Marseille	0.98 ^{○○}	10.08	0.38	1.58
Munich	0.99 ^{○○}	19.92	0.97 ^{○○}	13.30
Oslo	0.91 ^{○○}	9.73	0.92 ^{○○}	5.50

○ $P \leq 5\%$.

○○ $P \leq 1\%$.

Workshop III data on CTLs 4/1x and 4/2x possibly detecting new specificities are given in Table 9.

The conclusion, therefore, that CTLs primed *in vitro* recognize HLA-coded determinants, but that these may well be different from the serologically defined determinants is still very pertinent.

It must, however, not be forgotten that standardizing our individual techniques on the ability to produce identical assignments may counterselect for virtues of local techniques which would enable identification of marginal or weak effects such as e.g. "the target cell problems".

General Conclusions and Future Goals

1. There was excellent recovery and viability in all groups regarding the unprimed responding and stimulating cells and the CTL recovered after *in vitro* stimulation.

2. Improvements are needed at the level of 2° responses, including freezing of

primed cells, or alternatively doing this the local level.

3. All groups should aim to initiate studies of alternative means of stimulating CTL 2° reactions. This would include the use of PWM and special advice can be obtained from Marseille.

4. Hopefully, all groups will attempt to make T-cell growth factor and try to grow CTL in such media. Samples for comparison will be shipped from Munich, Marseille to all other groups and the laboratories will help with special advice.

5. Techniques for cloning of human alloreactive T-cells are now available (e.g. Malissin et al. 1980). Local techniques and experience will be circulated and compared locally.

6. New information can be obtained from analyzing various target cell populations. Aarhus has developed a uniform technique for making B cell blasts (Johnsen 1980).



and it should be initiated in all laboratories. The use of cultured T cells and other target cell types should be tested according to the interests of the individuals.

7 Competitive inhibition using cold targets can be a means to overcome some of the difficulties that most groups encounter when using a panel of ^{51}Cr target cells. Before the next Workshop, individual laboratories should experiment with this approach, e.g. using early and late PHA-blasts, EBV-cell-lines, T cell lines or platelets as cold competitors.

8 In the standard technique it still appears that there is a "target cell problem". It was suggested that laboratories might try Tris or Hepes buffered medium for ^{51}Cr incubation and target cell washing. It may be advisable to first distribute the CTL to the plates and return them to an incubator to readjust temperature and pH before adding the target cells.

9 It is by now desired to expand the participation in the CML Workshops as long as the new participants are actively involved in the area of CML research and are willing to share their expertise on the same basis as the others.

10 The immediate goal for CML-Workshop IV will be discussed within the group following local assessment of the questions raised above. A preliminary decision has been made to place the next Workshop meeting in Leiden during September 1980.

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