

NIH Public Access

Author Manuscript

Cell Immunol. Author manuscript; available in PMC 2010 January 1.

Published in final edited form as:

Cell Immunol. 2009; 258(2): 131–137. doi:10.1016/j.cellimm.2009.04.002.

Increased per cell IFN-γ productivity indicates recent *in vivo* activation of T cells

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Abstract

Immunization with vaccinia virus causes long-term immunity. Efforts have been made to characterize the T cells responsible for this protection. Recently, T cell subsets were described that not only coexpress multiple cytokines, but also show increased per cell cytokine productivity. These highly productive cells are often considered to be the most protective. We used ELISPOT assays to measure per cell IFN- γ productivity of vaccinia specific T cells in childhood immunized adults immediately before and at different time points after vaccinia re-vaccination. Apart from an increase in frequency, we found a marked increase of IFN- γ productivity following vaccinia re-vaccination. However, these changes were short-lived as both parameters quickly returned to baseline values within 22 days after re-vaccination. Therefore, increased per cell IFN- γ productivity seems to be a sign of recent *in vivo* T cell activation rather than a stable marker of a distinct T cell subset responsible for long-term immune protection.

Keywords

Vaccinia; T cell memory; Interferon gamma; Cytokine productivity; ELISPOT

1. Introduction

The T cell system responds with clonal expansion to antigen stimulation, leading to an increase of antigen-specific T cells in lymphoid tissues and in the peripheral blood [1,2]. Therefore, frequency measurements of specific T cells have established themselves as a method of quantifying the magnitude of T cell immunity, particularly when assessing the success of immunizations. Initially, limiting dilution assays were used for this purpose, but these have been increasingly replaced by techniques that provide more accurate frequency information with less experimental effort. Among these newer generation techniques, ELISPOT assays and intracytoplasmic staining (ICS) have gained wide use [3]. Supernatant-based cytokine measurements, such as ELISA, cytokine bead arrays, or measurements of cytokine mRNA, can offer semi-quantitative information on the magnitude of the T cell response, but do not provide direct frequency information.

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In addition to the frequency of antigen-specific T cells, per cell cytokine productivity has recently drawn attention as a potential marker of protective immunity. The notion has evolved that those T cells that are most effective in mediating immunity in infections such as influenza, vaccinia, CMV or EBV are those T cells that co-express multiple cytokines [4-7]. Furthermore, these multiple-cytokine producing T cells have also been shown to produce a larger amount of cytokine per cell [5-7]. This observation has led to the hypothesis that in addition to memory T cells, that may or may not be able to contribute directly to host defense, there is a distinct subset of highly productive multiple-cytokine effector cells that actually convey immune protection [8-11]. These findings, however, result from studies performed with ICS, for which cells need to be treated with secretion inhibitors in order to prevent the release of cytokine. Unfortunately, this pharmacological manipulation of cells alters the very end point to be measured: the actual release of cytokine. In order to avoid this alteration in our study, we chose to assess per cell cytokine productivity using ELISPOT assays that directly measure cytokine secretion by pharmacologically untreated cells.

For this purpose, we focused on the IFN- γ productivity of vaccinia-specific T cells. Childhood immunization with vaccinia virus is thought to result in long-term protection [12-16]. Therefore, in addition to antibodies, vaccinia-specific T cells in immunized individuals can also be considered a prototype of protective T cells [17-20]. In this study, adults vaccinated in childhood were re-vaccinated and the frequencies of vaccinia-specific T cells as well as their per cell IFN- γ secretion were measured in ELISPOT assays. We observed that per cell IFN- γ productivity strongly increased fourteen (14) days after re-vaccination. However, this increase was only transient in nature, and returned to baseline levels within twenty-two (22) days post re-vaccination. Therefore, we conclude that increased per cell IFN- γ productivity of T cells is not a steady characteristic of a certain protective T cell subset, but rather indicates a temporary T cell activation state. In classic terms, these findings are compatible with memory cells secreting low amounts of cytokine, and effector memory cells secreting a larger amount of cytokine per cell. Therefore, cytokine productivity may be used as a helpful indicator of recent *in vivo* T cell reactivation. It does not appear suitable, however, for assessing long-term protection.

2. Materials and methods

2.1. Subjects and sample collection

Four healthy adult members of our lab (2 males, 2 females, ages 33-53) who had been vaccinated in childhood with a single IM dose of vaccinia virus vaccine (DryvaxTM) volunteered to be re-vaccinated with a single IM dose of DryvaxTM. Peripheral blood was drawn immediately prior to re-vaccination (day 0) as well as on days 14 and 22 after re-vaccination. For all participants, PBMC were isolated from 40 to 100 ml of heparinized blood by standard Ficoll density-gradient centrifugation (Isoprep, Robbins Scientific, Sunnyvale, CA) and cryo-preserved. PBMC samples from different time points were tested side by side in ELISPOT assays as described below.

2.2. ELISPOT assays and digital image analysis

ImmunoSpot[®] plates (Cellular Technology Limited, Cleveland, OH) were coated with IFN- γ capture antibody mAb M700A (Endogen, Woburn, MA) in PBS (3 µg/ml) and placed at 4 ° C overnight. The plates were then blocked with PBS containing 1 % BSA (Sigma, St. Louis, MO) for 1 h and washed three times with PBS. PBMC were plated in RPMI-1640 (BioWhittaker, Walkersville, MD) supplemented with 100 U/ml penicillin, 100 µg/ml streptomycin, 2 mM L-glutamine and 10 % pooled heat-inactivated human AB serum. For all experiments 2×10⁵ PBMC were plated per well in the presence of live vaccinia Lister (Applied Biosystems, Foster City, CA; used at 1 MOI) or individual vaccinia peptides (PANATecs

GmbH, Tübingen, Germany; used at 5 µg/ml). Refer to Table 1 for peptide numbers assigned, amino acid sequences, and HLA restriction [21-25]. CEF peptide pool (Cellular Technology Limited, Cleveland, OH) contained MHC-I peptides derived from Cytomegalovirus, Epstein-Barr virus, and Influenza virus and was used at a final concentration of $2 \mu g/ml$ of each peptide. Phytohemagglutinin (PHA) was obtained from Sigma and used as a positive control in all assays (10 µg/ml). Negative control wells contained PBMC with medium only. After 24 h of incubation at 37 °C and 7 % CO₂, the plates were washed with PBS and PBS/TWEEN, and IFN- γ biotinylated detection mAb M701 (Endogen, 2 µg/ml) was added. The antibody was diluted in PBS containing 1 % BSA and 0.025 % TWEEN 20 (Fisher Scientific International Inc., Hampton, NH). The plates were incubated at 4 °C overnight. The plates were then washed three times with PBS/TWEEN and subsequently streptavidin-HRP conjugate (DAKO, Carpinteria, CA) was added at 1/2000 dilution, incubated for 2 h at room temperature, and removed by washing twice with PBS and PBS/TWEEN. The spots were visualized by adding HRP substrate 3-amino-9-ethylcarbozole (Pierce, Rockford, IL). The reaction was stopped by rinsing the plate with distilled water when distinct spots were visible macroscopically. Plates were dried overnight and images of the ELISPOT wells were captured with an ImmunoSpot® Series 5 Analyzer (Cellular Technology Limited, Cleveland, OH). Image analysis of ELISPOT results was performed with the ImmunoSpot® 5.0 Analysis Software (Cellular Technology Limited, Cleveland, OH), allowing precise spot size measurements down to spot sizes of 0.001 mm². Spot size distribution curves as well as mean spot sizes were also calculated using the ImmunoSpot® 5.0 Analysis Software (Cellular Technology Limited, Cleveland, OH).

2.3. Intracytoplasmic staining and flow cytometry

Intracellular cytokine staining was performed as previously described [26]. PBMC (4×10^6 /ml) were cultured overnight with or without live vaccinia virus (1 MOI) in complete RPMI medium. The next day, duel-color flow cytometry was performed to establish the frequency of IFN- γ producing cells within the CD3 cell population. For staining of IFN- γ , FITC-labeled IFN- γ Ab (Becton Dickinson, San Jose, CA) was used; for CD3, peridinin chlorophyl protein-labeled anti-human CD3 Abs (Becton Dickinson, San Jose, CA). The isotype-matched control monoclonal antibodies were also from Becton Dickinson. The samples were analyzed on a FACScan flow cytometer (Becton Dickinson, San Jose, CA). As a protein transport inhibitor, we used Monensin or brefeldin A (PharMingen, City, State).

3. Results

3.1. Vaccinia virus-induced IFN-y secreting T cells

Four healthy adult donors were selected for this study. All four had been vaccinated in childhood with vaccinia live virus vaccine. When tested before re-vaccination (day 0), all four donors showed a weak recall response to vaccinia virus, ranging from 5 to 40 IFN- γ producing cells per 200,000 PBMC (Fig. 1). These donors also responded to CEF peptide pool. When the donors were tested on day 14 after re-vaccination, the numbers of vaccinia-induced IFN- γ spots increased significantly, ranging between 250 and 540 spots per 200,000 PBMC. This corresponds to a frequency increase between 10-fold and 50-fold compared to day 0. The numbers of CEF-induced IFN- γ spots also increased slightly in all four donors. However, this increase was less than 2.3-fold in each case. When tested on day 22, the numbers of vaccinia specific cells had already dropped to 30-58 % of their day 14 frequency. The frequency of CEF reactive cells remained unchanged or dropped to day 0 values. These data are consistent with a vaccinia-induced expansion of specific T cells, clearly detectable on day 14 post re-vaccination, and an immediate subsequent contraction phase as indicated by markedly reduced frequencies by day 22. The moderate increase of CEF-reactive cells most likely results from bystander proliferation [27].

ICS was performed in parallel to ELISPOT assays in order to determine the cell type that produces IFN- γ after antigen stimulation (Fig. 2). The vaccinia-stimulated cells were counterstained with anti-CD3. As can be seen in Fig. 2A, in un-stimulated cells, 13 per 100,000 gated events were measured within the IFN- γ +/CD3+ quadrant. In the vaccinia-stimulated population (Fig. 2B), 71 out of 100,000 gated events occurred in the IFN- γ +/CD3+ quadrant and 12 in the IFN- γ +/CD3- quadrant. These data show that the majority of IFN- γ producing cells were CD3+ T cells and that the frequency of such cells was 0.0580 % within the PBMC tested. In the ELISPOT assay performed in parallel, the medium control was 0 spots for the un-stimulated cells (Fig. 2C), compared to 115 spots per 200,000 cells after vaccinia stimulation (Fig. 2D). The resulting frequency of vaccinia-induced antigen specific T cells detected by ELISPOT was 0.0575 %; essentially identical to ICS. Therefore, using the IFN- γ ELISPOT assay, T cells can be detected at single cell resolution. The vaccinia-induced cytokine production was strictly antigen specific, since none of the 12 vaccinia-naïve donors had *in vitro* vaccinia induced cytokine spots over medium background (data not shown).

3.2. Measurement of vaccinia virus-induced per cell IFN-y productivity

The striking observation made in this study was the fact that along with the frequency increase described above, the size of IFN- γ spots induced by vaccinia also changed fundamentally between day 0 and day 14. We used digital image analysis to measure the exact spot size distribution for each donor and time point. The size distribution curve for day 0 samples of the four donors are shown in Fig. 3A with an adjoining representative well. Day 14 results are shown in Fig. 3B along with a corresponding well. Already by day 21, the spot size distribution had returned to values similar to those on day 0 (Fig. 3C). Jointly, these data clearly show that on day 14, after re-vaccination, the mean spot size was significantly increased relative to day 0, and that this increase was transient, returning to baseline value by day 22 (Fig. 3F). Therefore, increased per cell IFN- γ productivity seems to be a transient feature of recently activated antigen-specific T cells.

3.3. Vaccinia peptide-induced IFN-y recall responses

The live vaccinia virus we used in the experiments above is a highly complex antigen system. We therefore extended our studies to the use of "clean" and simple antigenic subunits: vaccinia peptides. For this purpose, we chose 50 vaccinia peptides that had been described in the literature as T cell determinants and had them commercially synthesized [21-25]. The number assigned to each peptide along with its amino acid sequence and HLA restriction is listed in Table 1. The individual peptides were tested on three of the donors on day 0 and day 14 (Fig. 4). On day 0, the peptides that did induce a recall response over medium control elicited only low frequency IFN-γ secreting cells, with the exception of peptide number 11 in donor 1 (Fig. 4A). However, on day 14 after re-vaccination, several of the peptides triggered strong recall responses in all three donors (Fig. 4B, 4D, and 4F). Although it is not the scope of this publication to match the peptide-induced responses with each donor's HLA type based on each peptide's HLA restriction, this information can be extracted from the data provided in Fig. 4 and Tables 1 and 2. Those 12 peptides that induced substantial responses on day 14 were subsequently used for spot size analysis.

The vaccinia peptide-induced responses were once again antigen specific since they did not induce significant IFN- γ responses over medium background when tested individually in 10 naïve subjects (data not shown).

3.4. Measurement of vaccinia peptide-induced per cell IFN-y productivity

We systematically compared IFN- γ spot sizes induced by vaccinia peptides on days 0, 14, and 22 (Fig. 5). The sizes of these spots were not only compared between time points but also relative to those induced by CEF peptide pool. As can be seen in Fig. 5A, vaccinia peptide-

induced spots on day 0 (decades after childhood immunization) were tiny (mean spot size 0.0036 mm²) relative to CEF peptide-induced spots (mean spot size 0.0335 mm²). On day 14 after re-vaccination, the vaccinia peptide-induced spots showed a dramatic increase in size for all three donors and all of the selected peptides (Fig. 5B). The size distribution of spots induced by the peptides showed considerable variability between individual peptides. However, each peptide curve was shifted significantly to the right compared to day 0. By day 22, the peptide-induced spots had already decreased to sizes comparable to those spots induced by CEF peptide pool (Fig. 5C). Representative well images are shown for each time point next to the corresponding spot size distribution curve.

Consistent with the observations made using live virus for *in vitro* recall, these data show that also with individual vaccinia peptides as recall antigens, IFN- γ spot size significantly increased on day 14 after re-vaccination. Again, this increase was clearly transient, as by day 22, spot sizes had returned to baseline values (Fig. 5F).

The data obtained with "clean" vaccinia peptide antigens therefore confirm the results obtained with live virus: The increased per cell IFN- γ productivity after re-vaccination seems to be merely a transient feature of recently activated antigen-specific T cells and not a stable marker of long-term protection.

4. Discussion

ELISPOT assays are widely used to measure antigen-specific T cell immunity *ex vivo*. Such studies typically focus on establishing frequencies of antigen-specific cells within the PBMC population. As shown in Fig. 2 of this publication, the frequencies measured by ELISPOT and by ICS match up rather closely, suggesting that both assays accurately measure the antigen specific recall responses to Vaccinia if the frequencies are high enough. While ICS has a detection limit around 1 in 10,000 cells, ELISPOT assays have substantially lower detection limits [28;29]. This is well-illustrated in Fig. 2, where the test sample showed a frequency of vaccinia-reactive cells at a frequency of approximately 0.06 % which was easily detectable by ELISPOT but reached the detection limit of ICS. Therefore, ELISPOT's sensitivity and ease of use gives it a unique position for frequency measurements of antigen-specific T cells.

To our knowledge, there have been no published studies that took advantage of the additional strength of ELISPOT assays, namely to monitor cytokine productivity per cell. Typically, only spot counts are used, neglecting useful spot size information. Among all techniques that permit to measure antigen-induced cytokine production, ELISPOT is the only one that detects the actual release of cytokine by each individual cell. As the cell secretes, the cytokine is immediately captured around the cell during the entire culture period. Size and density of this "footprint" are a direct reflection of the amount of cytokine secreted per cell over the measuring period. It is important to note that only that fraction of cytokine is being captured on the membrane that diffuses towards the membrane while the rest diffuses into the supernatant. In an approximation, one can therefore compare the spot left on the membrane with a cross section through a cytokine sphere. As shown by Guerkov *et al.*, the total amount of cytokine released can be calculated based upon the diameter of the spot [30]. Therefore, using ELISPOT assays in our study enabled us to precisely detect the increase of per cell IFN- γ secretion based upon the changes in spot size we measured.

In earlier studies, we showed that even for a T cell clone, per cell IFN- γ productivity follows a close to perfect log normal curve [31]. This indicates that even within a clonal population of T cells, individual cells substantially differ in the amount of cytokine they secrete. Some cells are high producers, some are low producers, and the majority lie in between. Therefore, for all the data we analyzed in this study, we used digital image analysis software (ImmunoSpot[®])

with built-in algorithms to calculate spot size distribution curves. We then compared these curves in order to assess changes in per cell productivity. As seen in Fig. 3 and 5, significant shifts of size distribution were noted due to their magnitudes expressed on a logarithmic scale. These data clearly show that the per cell cytokine productivity of vaccinia-specific T cells strongly increases by day 14 after immunization relative to day 0 values, and subsequently declines by day 22. To our knowledge, this is the first report describing the short-lived nature of this increase in T cell cytokine productivity following antigen activation *in vivo*.

The amount of cytokine produced by T cells is a function of the signal strength. When studying a T cell clone, we found that the stronger the T cell stimulation was (increasing concentrations of *in vitro* recall antigen), the more cytokine the T cells produced, indicated by a marked rightshift of the log normally distributed spot size curves [31]. Notably, in this study, identical antigen and peptide concentrations were used for all time points. In another publication we described that the type of antigen presenting cell (APC) that activates a T cell substantially influences the T cell's cytokine productivity [32]. T cells stimulated on dendritic cells (DC) not only engaged more rapidly in cytokine production, but also produced more cytokine per cell compared to T cells activated by B cells. Macrophages led to an intermediate level of T cell activation. Based on this information, the increased per cell IFN- γ productivity seen on day 14 in the present study could have also been caused by changes in the APC population: An increase of PBMC and DC caused by inflammation and leukocytosis could have led to enhanced T cell activation due to a higher density of professional APC. In that case, the increased per cell IFN- γ productivity would not be an inherent property of the T cells themselves, but rather induced by APC fluctuations. If APC changes had indeed contributed to the increased per cell productivity in our study, a right shift in spot size distribution should have also been seen in the CEF peptide-induced spots. As illustrated in Fig. 5A vs. 5B, the CEF peptide-induced curves perfectly overlap for days 0 and 14. Only the vaccinia-induced spots underwent the size increase. Therefore, we conclude that the increased per cell productivity is in fact a property of the activated T cells themselves.

Increased per cell productivity has previously been detected by ICS and linked to a multicytokine producing T cell phenotype [5-7]. A series of recent publications tried to make the point that multi-cytokine producing T cells constitute the T cell subset critical for host defense [4-11]. Some of these publications also showed that these multi-cytokine producing T cells are characterized by an increased per cell cytokine productivity. What has remained unclear, however, is whether this multi-cytokine per increased-productivity phenotype is a stable marker of a cell lineage that carries immunological memory and protection, or whether it merely reflects a transient T cell activation state. Our data do not attempt to resolve this issue. However, some conclusions can clearly be drawn. Studying the four individuals that we examined in detail - along with more than 50 additional donors immunized in childhood, but not re-vaccinated (data not shown) - we could generate substantial information on long-term memory cells. All of the spot size distributions we measured in these individuals decades after immunization were similar to those shown in Fig. 3A, where none of the spots exceeded a size threshold of 0.08 mm² (corresponding to 10^{-1.1} on the logarithmic scale). Therefore, high productivity cells (causing spots larger than this) that were commonly seen on day 14 and occasionally still on day 22, occur (if at all) in undetectably low frequencies in the long-lived memory T cell pool known to provide long-term protection. These high productivity cells were only detectable transiently after re-vaccination and then disappeared. We therefore conclude that the high producing cell type is not a protective memory T cell lineage of its own but rather represents a transiently increased activation state of long-lived memory T cells. Even if high productivity cells constituted a distinct T cell subset or blast, they would be too short-lived to convey the long-term protection seen after vaccinia immunization. The simplest explanation of our data is that long-term resting memory cells show low per cell productivity, but when stimulated with antigen, transform into effector memory cells with high cytokine productivity.

By no means do our data attempt an extensive analysis of cytokine productivity or memory T cell biology in general. The data emerged from a limited re-vaccination project and, in view of these striking results, are communicated with two intents: First, to introduce cytokine productivity measurements as a promising new application for T cell diagnostics and to show how easily it can be added to the standard frequency measurements; second, to draw attention to the fact that increased per cell productivity may signify recently *in vivo* activated T cells rather than a distinct subset of long-term memory T cells. Therefore, rather than using productivity measurements in order to distinguish between long-term memory and recently activated T cells, and thus be able to detect actively ongoing immune processes, be it in infection, allergy or autoimmune disease.

Acknowledgments

The authors wish to thank Ms. Andrea Csipak for her most valuable assistance with image analysis and statistical calculations.

This work was supported by NIAID contract N01-AI-400948.

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Fig. 1.

Frequencies of vaccinia specific T cells before and after vaccinia re-vaccination. The numbers of IFN- γ spots per 200,000 PBMC, measured by ELISPOT immediately prior to re-vaccination (day 0) as well as 14 and 22 days after re-vaccination, are shown for each of the four donors. Panels A – D correspond to donors 1 – 4. IFN- γ responses to CEF peptide pool served as control.

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Fig. 2.

Comparison of IFN- γ ELISPOT and ICS. Frequencies of vaccinia specific T cells were measured for one donor on day 0 using IFN- γ ELISPOT and ICS in parallel. Measurements obtained with ICS are shown without (A) and with (B) *in vitro* vaccinia stimulation. The number of events collected in quadrant 2 (Q2), representing CD3+/IFN- γ + cells, are given along with the resulting frequency of vaccinia specific T cells in percent. ELISPOT was performed with the same PBMC samples. Well images are shown without (C) and with (D) *in vitro* challenge with live vaccinia virus. Spot numbers are given for each well along with the resulting mean spot number and the frequency of vaccinia specific T cells in percent.



Fig. 3.

IFN- γ spot size distribution using live vaccinia virus as recall antigen. Panel A shows the individual IFN- γ spot size distribution curves for donors 1 – 4 (dotted lines), obtained with live virus *in vitro* recall of day 0 PBMC. The solid line indicates the mean spot size distribution of all four donors. For comparability, all curves are adjusted to cover a total of 500 spots. The same information is given for days 14 (B) and 22 (C). Representative well images are depicted for each time point. Panel D directly compares the mean spot size distribution curves from panels A – C, corresponding to the three time points (dotted line, day 0; solid line, day 14; dashed line, day 22). Panel E illustrates the day 14 spot size shift by showing the upper range

of the three curves from panel D on a linear scale. Panel F shows the mean spot sizes (mean of all four donors) for the three time points.



Fig. 4.

IFN- γ recall response to individual vaccinia peptides. The 50 individual vaccinia peptides listed in Table 1 were used as recall antigens to detect vaccinia peptide specific T cell responses in donors 1 – 3 before (day 0) and after re-vaccination (day 14). Donor 1: panels A and B. Donor 2: panels C and D. Donor 3: panels E and F.





Fig. 5.

IFN- γ spot size distribution using individual vaccinia peptides as recall antigens. Peptide antigens that elicited a marked frequency increase in Fig. 4 were chosen for detailed spot size analysis (n = 12). Panel A shows the individual IFN- γ spot size distribution curves elicited by each of the chosen peptides (dotted lines) when used as recall antigen on day 0 PBMC. The solid line indicates the mean spot size distribution calculated from all 12 peptides (combined data from donors 1 – 3). The dashed line shows the spot size distribution obtained with CEF peptide pool as control. For comparability, all curves are adjusted to cover a total of 500 spots. The same information is given for days 14 (B) and 22 (C). Representative well images are depicted for each time point. Panel D directly compares the mean spot size distribution curves

from panels A - C, corresponding to the three time points (dotted line, day 0; solid line, day 14; dashed line, day 22). Panel E illustrates the day 14 spot size shift by showing the upper range of the three curves from panel D on a linear scale. Panel F shows the mean spot sizes (mean of all 12 peptides) for the three time points.

Table 1

Vaccinia peptides ^a

Pentide ID	Sequence	HLA restriction	
1	VSVNNVCHMY	A *2601	
2	SOSDTVEDY	A*0101	
3	OSDTVFDYY	A*0101	
4	VTDTNKFDNY	A*0101	
5	CMI TEEL HY	A*2601	
5	ETIDEKI KY	A*2001 A*2601	
7	GTHVLLPFY	Δ*0101	
, 8	DMCDIVLLY	A*0101 *2601	
9	FGDSKEPVPY	A*0101, *2601	
10	FI SMI NI TKY	Δ*0101, *2601	
10	OSITRSLIV	A*2601	
12	AMI NGIVV	A*0201	
12	VI PEDIKKI	A *0201	
14	NI WNGIVPM	A*0201	
14		A*0201 *0202 *0202 *0202	
15	VVNAIL VOL	A*0201, *0202, *0203, *0206	
10	I VNAIL I QI	A*0201	
17	SLSATIK V	A*0201	
10		A*0201	
19		A*0201	
20	FLISVINKV	A*0201	
21		A*0201	
22	FLNISWFYI	A*0201	
23	GLNDYLHSV	A*0201	
24	SMHFYGWSL	A*0201	
25	YLYTEYFLFI	A*0201	
26	MMLVPLITV	A*0201	
27	YTYGIPLSL	A*0201	
28	AVYGNIKHK	A*0301	
29	KVLHVIDNK	A*0301	
30	ATSLDVINY	A*1101	
31	KLKIISNDYK	A*0301	
32	KVMFVIRFK	A*0301, *1101	
33	IVFNLPVSK	A*0301	
34	NQVKFYFNK	A*0301	
35	KTKNFTIDFK	A*0301	
36	YLLVKWYRK	A*3303	
37	VTSSGAIYK	A*1101	
38	GTIAGGVCYY	A*1101	
39	AVFKDSFLRK	A*1101	

Peptide ID	Sequence	HLA restriction	
40	VWINNSWKF	A*2402	
41	RYRFAFLYLI	A*2402	
42	RYYDGNIYE	A*2407	
43	KPKPAVRFAI	B*0702	
44	APNPNRFVI	B*0702	
45	RPMSLRSTII	B*0702	
46	MPAYIRNTL	B*0702	
47	GESKSYCEL	B*4001	
48	DELVDPINY	B*3701	
49	TEYDDHINL	B*4001	
50	HPRHYATVM	B*0702	

^aThe table lists the assigned ID numbers, amino acid sequences (single letter code), and HLA restrictions for all 50 vaccinia peptides used as recall antigens in Fig. 4 and 5.

Table 2

HLA genotypes^a

HLA Class I	Donor 1	Donor 2	Donor 3
A1	A3-0301	A2-0201	A2-0201
A2	A11-1101	A24-2402	A24-2402
B1	B35-3511	B44-4403	B15-1535
B2	B55-5513	B51-5101	B40-4001
C1	Cw9-0303	Cw1-0102	Cw7-0702
C2	Cw4-0401	Cw4-0401	Cw15-1502

aThe table lists the HLA genotypes of donors 1 - 3.