



Optimized protocol for the detection of multifunctional epitope-specific CD4+ T cells combining MHC-II tetramer and intracellular cytokine staining technologies

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1	Optimized protocol for the detection of multifunctional epitope-specific CD4 ⁺ T cells combining
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4	Gabiria Pastore, Monica Carraro, Elena Pettini, Emanuele Nolfi, Donata Medaglini and Annalisa
5	Ciabattini [*]
6	
7	Laboratory of Molecular Microbiology and Biotechnology (LA.M.M.B.), Department of Medical
8	Biotechnologies, University of Siena, Siena, Italy.
9	
10	*Correspondence:
11	Università di Siena, Dipartimento di Biotecnologie Mediche
12	Policlinico Le Scotte, V lotto piano 1,
13	Viale Bracci, 4
14	Siena 53100, Italy
15	e-mail: annalisa.ciabattini@unisi.it
16	
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21 Abstract

22

Analysis of multifunctional CD4⁺ T cells is fundamental for characterizing the immune responses to 23 vaccination or infections. Peptide-MHC tetrameric complexes represent a powerful technology to 24 detect antigen-specific T cells by the specific binding to their T cell receptor, and their combination 25 with functional assays is fundamental for characterizing the antigen-specific immune response. Here 26 27 we optimized a protocol for the detection of multiple intracellular cytokines within epitope-specific CD4⁺ T cells identified by the MCH class II tetramer technology. The optimal procedure for assessing 28 the functional activity of tetramer-binding CD4⁺ T is based on the simultaneous intracellular staining 29 with both MHC tetramers and cytokine-specific antibodies upon in vitro restimulation of cells with 30 the vaccine antigen. The protocol was selected among procedures that differently combined the steps 31 of cellular restimulation and tetramer staining with intracellular cytokine labelling. 32 This method can be applied to better understand the complex functional profile of CD4⁺ T cell 33

34 responses upon vaccination or infection.

- 35 **1. Introduction**
- 36

The study of the CD4⁺ T cell activation and effector function is fundamental in the characterization of immune responses to vaccination (1).

CD4⁺ T cells play a central role in mediating vaccine immune responses by shaping both the 39 humoral and cellular immunity (2). Activated CD4 T cells are critically involved in providing 40 41 cognate help to B cells for production of protective antibodies, and modulate the functions of macrophages and CD8⁺ cytotoxic T cells through cytokines secretion. The characterization of the 42 cytokine production of antigen-specific T cells is therefore of critical importance to profile 43 vaccine immune response. The direct and specific method for identifying antigen-specific CD4⁺ 44 45 T cells is based on the major histocompatibility complex (MHC) tetramer staining technique (3). This procedure allows the identification of specific T cells due to the selective and multivalent 46 47 binding of tetramer MHC-peptide complexes to the T cell receptors (TCR) (3,4) and has been used for characterizing the primary and recall antigen-specific CD4⁺ T cell responses in many 48 49 pre-clinical and human studies (1,5–9).

The effector function of antigen-reactivated T cells is commonly measured by flow cytometry-50 based intracellular cytokine staining (ICS) that allows the simultaneous phenotypic 51 52 characterization and cytokine detection within single cells (10,11). The characterization of intracellular cytokines allows to identify activated CD4⁺ T cells capable of producing more than 53 54 one cytokine, and the analysis of these multifunctional/polyfunctional cells is important for characterizing the immune response elicited by the vaccination or natural infections (12). 55 Polyfunctional CD4⁺ T cells secreting IFN- γ , TNF- α and IL-2 have been proposed as a major 56 component of immune response that correlates with mouse protection against challenge with 57 *Leishmania major* (13). In tuberculosis (TB), it has not been clarified if the frequency and quality 58 of polyfunctional CD4⁺ T cell responses elicited in mice by different types of vaccines correlate 59 with protective immunity (14–17), while human studies have shown that a consistent response of 60 61 CD4⁺ T cells co-expressing IFN- γ , TNF- α and IL-2 was associated with acute TB infection (18). Tetramer labeling and intracellular cytokine staining are generally not recommended to be 62 performed concurrently since the *in vitro* antigen restimulation can induce TCR internalization, 63 thus losing the possibility of detecting epitope-specific $CD4^+$ T cells using tetramers (19). 64

In order to identify a protocol for the detection of intracellular cytokine production within the activated epitope-specific CD4⁺ T cells, we assessed different strategies that combined cellular restimulation (with the vaccine antigen or tetramers) and tetramer staining (extracellular or intracellular) with intracellular cytokine labelling. The different procedures were tested in

- splenocytes from mice immunized with the chimeric tuberculosis vaccine antigen H56 (20) mixed 69 70 with the adjuvant CAF01 (21), a model vaccine formulation deeply characterized in preclinical studies for its capacity of inducing both humoral and cellular responses (9,22–24). H56 is a fusion 71 protein of *M. tuberculosis* antigens Ag85B, ESAT-6, and Rv2660, and the H56-specific CD4⁺T 72 cell response can be monitored by employing Ag85B₂₈₀₋₂₉₄-complexed MHC class II tetramers 73 (8). The different procedures were tested also in another experimental setting, in which mice were 74 immunized with the model chicken ovalbumin antigen, and the CD4⁺ T cell response was assessed 75 76 employing tetramers specific for the epitope₃₂₅₋₃₃₅ (25). 77 The comparative analysis of the different protocols has permitted to optimize the procedure for
- identifying the multifunctional profile of tetramer-specific CD4⁺ T cells performing intracellular
 staining with both tetramers and cytokine-specific antibodies, upon antigen restimulation. This
 method represents a helpful tool for identifying epitope-specific CD4⁺ T cells and analyzing their
- 81 82

specific effector function.

83 **2. Materials and methods**

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85 **2.1 Mice**

Female C57BL/6 mice, purchased from Charles River (Lecco, Italy) were housed under specific
pathogen-free conditions in the animal facility of the Laboratory of Molecular Microbiology and
Biotechnology (LA.M.M.B.), Department of Medical Biotechnologies at University of Siena, and
treated according to national guidelines (Decreto Legislativo 26/2014). The protocol was approved
by the Italian Ministry of Health (authorization n° 1004/2015-PR, 22 September 2015).

91

92 **2.2 Immunizations**

93 Groups of 10-12 mice were immunized by the subcutaneous route at the base of the tail with the 94 chimeric tuberculosis vaccine antigen H56 (2 μ g/mouse) combined with the adjuvant CAF01 (250 μ g 95 dimethyldioctadecylammonium and 50µg trehalose dibehenate/mouse, both kindly provided by Statens Serum Institut, Denmark) and boosted with a lower dose of H56 alone (0.5 µg/mouse) 4 96 97 weeks later. Another experiment was performed immunizing mice with albumin from hen egg white (OVA, 25 µg/mouse, Sigma-Aldrich) combined with the adjuvant CAF01, and boosted with OVA 98 99 alone. The formulations containing antigens and CAF01 were injected in a volume of 150µl/mouse 100 of Tris 10 mM, while the formulations containing H56 and OVA alone in a volume of 100µl/mouse 101 of 1X Dulbecco's Phosphate Buffered Saline (1X PBS). Mice were sacrificed 5 days after boosting.

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103 **2.3 Sample collection and cell preparation**

Spleens collected from mice were mashed onto 70µm nylon screens (Sefar Italia, Italy) and washed in complete RPMI (cRPMI) medium [RPMI (Lonza, Belgium), 100 U/ml penicillin/streptomycin, and 10% fetal bovine serum (Gibco, USA)] for 10 min at 300 g at 4°C. Splenocytes were treated with red blood cell lysis buffer (1X, eBioscience, USA) for 4 min. Following centrifugation at 300 x g at 4°C for 10 min, cells were washed with 1X PBS and counted with cell counter (Bio-Rad, USA).

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110 **2.4 Protocols and reagents**

Six different protocols for detecting intracellular cytokines within activated epitope-specific CD4⁺ T splenocytes that differently combined cellular restimulation, tetramer staining and cytokine labelling, were assessed (Figure 1). Protocols were assessed in two different experimental settings, in which mice were immunized with H56 or OVA antigens, and the CD4 T cell response was analysed employing two different tetramers, specific for the H56 or OVA epitopes, respectively.

Protocol 1: Splenocytes (2x10⁶/well) were cultured in a round-bottom 96-well plate with H56 protein 116 (2 µg/ml) or OVA (50 µg/ml), anti-CD28 and anti-CD49d costimuli (both 2 µg/ml, eBioscience) at 117 37° C, 5%CO₂ for 6h, with Brefeldin A (BFA, 5 µg/ml, Sigma-Aldrich) and monensin solution (1×, 118 eBioscience) added during the last 5 h of incubation. Cells were washed with cRPMI for 7 min at 300 119 g at 4°C, labelled with Fixable Viability Stain 780 staining (FVS780, BD Biosciences, 1:1000, 100 120 µl/well) for 20 min at RT in the dark, and washed twice in PBS. Cells were fixed and permeabilized 121 for 20 min at 4°C with BD Cytofix/Cytoperm (Becton Dickinson). Samples were blocked for 30 min 122 at 4°C in Fc-blocking solution (5 µg/ml of CD16/CD32 mAb, eBioscience, USA) and stained for 1 h 123 124 at room temperature (RT) with PE-conjugated I-A (b) M. tuberculosis Ag85B precursor 280-294 (FQDAYNAAGGHNAVF) tetramer (diluted 1:80, hereafter Tet-Ag85B) or with PE-conjugated I-A 125 (b) chicken ova 325-335 (QAVHAAHAEIN) tetramer (diluted 1:50, hereafter Tet-OVA; both 126 tetramers were kindly provided by NIH MHC Tetramer Core Facility, Emory University, Atlanta, 127 GA, USA) diluted in Perm/wash buffer. In the last 20 minutes of tetramer incubation, the following 128 mix of fluorescent antibodies was added: APC-conjugated anti-CD3 (clone 145-2C11), BB700-129 conjugated anti-CD4 (clone RM-5), APC-R700-conjugated, anti-CD44 (clone IM-7), BV786-130 131 conjugated anti-IFN-γ (clone XMG1.2), BV650-conjugated anti-TNF-α (clone MP6-XT22), BV421conjugated anti-IL-17A (clone TC11-18H10), PE-CF594-conjugated anti-IL-2 (clone JES6-5H4), 132 133 (all antibodies were purchased from BD Biosciences). All antibodies and tetramer were titrated for

134 optimal dilution.

Protocol 2: Splenocytes were cultured with the respective antigens and costimuli as in protocol 1, then were washed and stained with the respective tetramers for 1h at RT. Cells were labelled with FVS780, fixed and permeabilized with BD Cytofix/Cytoperm and stained with the mix of fluorescent antibodies for 20 minutes at RT.

Protocol 3: Splenocytes were stained with the respective tetramers for 1h at RT, washed and stimulated with the respective antigens and anti-CD28 and anti-CD49d costimuli at 37°C for 6h, with Brefeldin A and monensin solution added during the last 5 h of incubation. Cells were labelled with FVS780, fixed and permeabilized with BD Cytofix/Cytoperm and stained with the mix of fluorescent antibodies for 20 minutes at RT.

- *Protocol 4*: Splenocytes were cultured with the respective tetramers and costimuli for 1h at RT,
 washed and added with BFA and monensin solution at 37°C for 5 h. Cells were labelled with FVS780,
- 146 fixed and permeabilized with BD Cytofix/Cytoperm and stained with the mix of fluorescent

147 antibodies for 20 minutes at RT.

Protocol 5: Splenocytes were cultured with the respective tetramers and costimuli for 1h at RT and
for 5h at 37°C in the presence of BFA and monensin. Cells were labelled with FVS780, fixed and

- permeabilized with BD Cytofix/Cytoperm and stained with the mix of fluorescent antibodies for 20minutes at RT.
- *Protocol 6*: Splenocytes were cultured with the respective tetramers and costimuli for 6h at 37°C,
 with BFA and monensin during the last 4h of incubation. Cells were labelled with FVS780, fixed and
 permeabilized with BD Cytofix/Cytoperm and stained with the mix of fluorescent antibodies for 20
 minutes at RT.
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159 **2.5 Flow cytometry**

About 7x10⁵ stained cells from each protocol were acquired on BDTM LSRFortessa X20 flow cytometer (BD Biosciences) and stored. Data analysis was performed using FlowJo v10 (TreeStar, USA), and the evaluation of different cytokines co-expression was performed using the FlowJo Boolean gate platform. Fluorescence minus one (FMO) controls were performed for all fluorescence and used for gating setting.

165 166

167 **2.6 Statistical analysis**

168 Kruskal-Wallis test, followed by Dunn's post test for multiple comparisons, was used to assess the

- statistical difference between protocols. A P value ≤ 0.05 was considered significant. Analysis were
- 170 performed using GraphPad Prism v7 (GraphPad Software, USA).

171 **3. Results**

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In order to optimize the protocol for the detection of intracellular cytokines within activated epitope-173 specific CD4⁺ T cells, we tested different procedures in splenocytes from mice parenterally 174 immunized with two different antigens, the chimeric TB vaccine antigen H56 or OVA, combined 175 with the liposome adjuvant CAF01, 5 days after the booster immunization. The induction of Ag-176 specific CD4⁺ T cells producing cytokines was assessed combining antigen restimulation and 177 tetramer staining, followed by intracellular cytokine detection (Figure 1). In protocols 1-3 splenocytes 178 179 were restimulated with the respective antigens, added before (protocols 1 and 2) or after (protocol 3) tetramer staining. In protocols 4-6 the restimulation step was performed directly with epitope-180 181 complexed MHC II tetramers, that were therefore used not only as staining tool, but also as functional stimulus. The comparison of results obtained following the different strategies, and tested with two 182 183 different antigens, has permitted to optimize the procedure for identifying the cytokine profile of tetramer-specific CD4⁺ T cells. 184

185

186 **3.1 Identification of tetramer-specific CD4**⁺ **T cells producing cytokines**

187 H56-specific CD4⁺ T cells were identified using the Ag85B₂₈₀₋₂₉₄-complexed MHC class II tetramers specific for the immunodominant epitope of Ag85B (26), which is part of the chimeric H56 protein, 188 while OVA-specific CD4⁺ T cells using the chicken OVA₃₂₅₋₃₃₅ -complexed MHC class II 189 tetramersTetramer positive cells (Tet-Ag85B⁺ or Tet-OVA⁺) were identified as live single CD3⁺ 190 CD4⁺ CD44⁺ cells, and TNF-α, IFN-γ, IL-17 and IL-2 cytokines were detected within gated Tet-191 Ag85B⁺ cells. All gates were defined on the bases of the respective FMO controls. Staining specificity 192 was determined using a control tetramer complexed with an unrelated antigen that showed a level of 193 staining below 0.02% (data not shown). The identification of Tet-Ag85B⁺ cells in the different six 194 protocols (Figure 2 A), and their intracellular cytokine production (Figure 2 B) are shown. 195

ICS protocol typically includes an antigen stimulation step, that is crucial for activation of effector 196 function of CD4⁺ T cells. Nevertheless, this step induces the internalization of TCR molecules, thus 197 198 negatively impacting on the tetramer staining procedure. To overcome this limitation, we assessed a strategy based on the antigen stimulation phase followed by permeabilization and fixation of cells 199 200 and subsequent tetramer staining (Figure 1, protocol #1). Using this procedure, that allows to identify both extra and intracellular TCR molecules, we detected 0.53% of Tet-Ag85B⁺ cells (Figure 3A, 201 202 orange box). This frequency was significantly higher compared to protocol 2, in which splenocytes 203 were firstly stimulated with H56 antigen and then labelled with the specific tetramer (Figure 1, protocol #2) allowing an identification of only a 0.2 % of tetramer-positive T cells (Figure 3A, lightgreen).

The impact of the tetramer staining performed before antigen restimulation was also evaluated (Figure 1, #3). This procedure allowed to detect a frequency of 0.37% of Tet-Ag85B⁺ T cells that was higher compared to protocol 2 while was lower respect to protocol 1(Figure 3A dark green box). The higher number of Tet-Ag85B⁺ T cells detected in protocol 1 could be due to the effect of prior antigen restimulation that is known to induce the formation of large clusters of TCR molecules thus increasing tetramer binding avidity (27).

In protocols 4, 5 and 6, there was no antigen stimulation and the Ag85B₂₈₀₋₂₉₄-complexed MHC class 212 II tetramers were used not only for identifying but also for stimulating antigen-specific CD4⁺ T cells 213 (Figure 1, protocols #4, 5 and 6). As expected, the frequencies of Tet-Ag85B⁺ T cells detected in 214 protocol 4 were comparable to those of protocol 3 (Figure 3A, light blue and dark green box). In 215 216 protocols 5 and 6, in which a tetramer incubation phase of 6 hours was performed, frequencies of 1.07 and 1% of Tet-Ag85B⁺ T cells were observed, significantly higher respect to H56-stimulated 217 samples (Figure 3A, $P \le 0.05$ and $P \le 0.001$ compared to protocols 3 and 2, respectively). A similar 218 behavior was observed in mice immunized with the OVA antigen, in which the OVA-specific CD4⁺ 219 T cells were identified using the OVA₃₂₅₋₃₃₅ peptide-complexed tetramer (Figure 3B). This shows that 220 tetramers with different peptide specificity, respond in a very similar way to the *in vitro* staining 221 procedures assessed in the different protocols. 222

The effector function of both Tet-Ag85B⁺ and Tet-OVA⁺ T cells identified with the different 223 strategies, was analyzed by measuring the intracellular production of 4 different cytokines using 224 multiparametric flow cytometry. As shown in figure 3 C and D, the highest percentage of cells 225 producing intracellular cytokines was detected for both tetramers in protocols 1, 2 and for Tet-226 Ag85B⁺ also in protocol 3, compared to protocols 4, 5 and 6. These data show that in absence of 227 antigen stimulation, despite the high frequencies of tetramer-binding CD4⁺ T cells (protocols 5 and 228 6, Figure 3 A and B), a significant lower cytokines production is induced (Figure 3 C and D) 229 230 highlighting the importance of the *in vitro* re-stimulation step with the vaccine antigen to effectively 231 stimulate the effector function of antigen-specific CD4⁺ T cells.

232

3.2 Evaluation of tetramer-specific CD4⁺ **T cells mutlifunctional profile.**

In order to have a picture of the multifunctional profiles of T cells elicited by immunization and detected by the different experimental procedures, a Boolean analysis of data was performed within Tet-Ag85B⁺ cells (Figure 4A). A significant amount of cells positive for all the four cytokines or for TNF- α , IFN- γ , and IL-2 were observed in protocols 1 and 3 compared to protocols 4 and 5 (P \leq 0.05).

Cells producing only IFN- γ were instead significantly higher in protocols 5 and 6 compared to 238 protocol 2 (P \leq 0.01). The analysis of the frequency of cells producing two or more cytokines 239 (multifunctional), a single one (single) or no cytokines in each protocol shows that the frequency of 240 multifunctional Tet-Ag85B⁺ T cells was higher in protocols 1, 2 and 3 (79, 68, 69 % respectively) 241 that included the antigen restimulation. Lower frequencies of multifunctional cells were observed in 242 protocols 4, 5 and 6 (12, 7, 18 % respectively) in which most of cells did not produce any cytokine 243 (75, 74, 68 %) (Figure 4B). The intracellular staining with the tetramer performed in protocol 1, 244 allowed to detect the highest percentage of multifunctionality, while no differences were observed 245 among protocols 2 and 3. Therefore the optimal strategy of staining that allows to identify 246 multifunctional T-helper cells among tetramer-specific CD4⁺ T cells was the protocol #1. 247

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In conclusion, our comparative analysis, confirmed for two different antigens and their respective tetramers, has shown that the optimal strategy for identifying the multifunctional cytokine profile of tetramer-specific CD4⁺ T cells is the procedure #1, in which the antigen restimulation phase is followed by the intracellular tetramer staining. Indeed, this protocol allows to detect a significant amount of tetramer-specific T cells, their multifunctional activity, and it allows to reduce the staining time, by adding cytokine-specific antibodies in the last 20 minutes of tetramer incubation. A detailed description of protocol #1 is reported in Figure 5.

256 **Discussion**

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In this study we optimized a flow cytometric protocol for identifying at the single-cell level 258 multifunctional epitope-specific CD4⁺ T cells, elicited by immunization. Demonstrating pros and 259 260 cons of different protocols, we showed that the optimal procedure for the simultaneous detection of epitope-specific CD4⁺ T cells and their effector function is based on the antigenic stimulation of cells 261 combined with a single step of cytokine and tetramer staining in permeabilized cells (Figure 5). Our 262 analysis was based on the comparison of different experimental procedures, tested with two different 263 264 epitope-specific tetramers, in which the steps of antigen restimulation, tetramer and cytokine staining were differently combined. The systematic analysis of different procedures performed in the same 265 266 samples has offered the possibility of selecting the optimal protocol among different strategies. The results have been confirmed with tetramers specific for two different antigens, thus strengthening the 267 268 possible application of the selected procedure to the characterization of the complex functional profile of CD4⁺ T cell responses upon vaccination or infection. 269

270 Most of the studies of intracellular cytokine production within tetramer positive cells, have been conducted in CD8⁺T cells (28–30), while few works have been performed in CD4⁺T cells with MHC 271 272 class II tetramers (31–33) and none has compared different protocols in a systematic way. Even though a direct comparison with the present study is difficult due to different experimental settings, i.e the 273 use of T cell clones or human CD4⁺ T cells, prolonged incubation with antigen for cell activation, 274 magnetic bead enrichment of tetramer positive cells before ICS staining, we can generally observe 275 276 that the tetramer staining was performed extracellularly, often before the antigen stimulation step. 277 Our analysis clearly demonstrates that antigenic stimulation is necessary for an efficient reactivation 278 of the cellular effector function, and the same stimulation effect can not be obtained with the direct 279 incubation of cells with epitope-complexed MHC tetramers, also when prolonged for 6 hours 280 (protocols 5-6). Nevertheless, many studies have demonstrated that ligation of TCR by processed antigen induces TCR internalization and a subsequent down-modulation of its cell surface expression 281 282 (19,34). Indeed, in protocol 2, in which antigen stimulation was performed before the extracellular 283 tetramer staining the frequency of tetramer positive cells was significantly lower compared to protocol 1. 284

Here, using tetramers specific for two different antigens, we have shown the efficiency of tetramer staining performed in permeabilized cells (protocol 1) that allows to detect both surface expressed and internalized TCR molecules resulting in the identification of the highest percentages of tetramerbinding cells. This procedure stained epitope-specific CD4⁺ T better than protocol 3 in which labeling with tetramer was performed before antigen stimulation. This can be due to the lower avidity of tetramer binding to TCR molecules in the absence of cell activation by antigen stimulation. Indeed,
cellular activation is known to induce the TCR reorganization with the generation of large clusters
of TCR molecules (27) that increase the strength of tetramer binding.

Analysis of multifunctional CD4⁺ T cells is of critical importance for in depth characterization of 293 immune responses to vaccination both in pre-clinical and clinical studies. It is therefore essential to 294 295 have a protocol that optimally combines the identification of antigen-specific T cells with the analysis of their cytokine profile. Here, we show the possibility to combine, upon antigen stimulation, tetramer 296 297 and intracellular cytokine staining in permeabilized cells allowing the identification of an higher 298 number of polyfunctional tetramer positive CD4 T cells. The amount of cells producing all the four cytokines, or coexpressing two or three cytokines (especially TNF- α , IFN- γ , and IL-2) was indeed 299 300 higher compared to the other protocols tested. Significant lower levels of multifunctional cells were observed when tetramers were used both as stimulus and as staining (protocols 4, 5 and 6). Indeed, 301 302 even though a higher percentage of tetramer-binding T cells was identified by protocols 5 and 6, about 70% were negative for cytokine production, respect to 14% observed in protocol 1, demonstrating 303 304 that the binding of epitope-complexed MHC class II molecules to TCR in the presence of CD49d and CD28 costimuli is not sufficient for effectively reactivating multifunctional antigen-specific CD4⁺ T 305 cells. 306

The functional characterization of $CD4^+$ T cells described by the protocols analyzed here, is particularly suitable for pre-clinical studies, in which sufficient quantity of $CD4^+$ T cells can be easily identified in draining lymphoid organs such as lymph nodes or spleens, while in humans it is generally more complicated because the frequency of antigen-specific $CD4^+$ T cells in blood are low and often undetectable (6). Moreover, the use of MHC tetramers requires prior knowledge of the peptide epitope and host MHC haplotype, a limitation that can be easily circumvented in inbred animals.

In conclusion, in the present work we have selected an optimized protocol for identifying epitopespecific CD4⁺ T cells and their effector function, combining antigenic stimulation of cells with the intracellular staining of TCR molecules and cytokines. Antigenic restimulation, performed at the beginning of the procedure, allows the activation of cells and elicits multiple cytokine production, but at the same time it promotes the down regulation of surface TCR expression that is resolved by the intracellular tetramer staining. This procedure allows also to reduce the total protocol time, since tetramer, surface marker and cytokine staining are combined in a single staining step.

This protocol allows to better understand the complex functional profile of T cell responses upon vaccination or natural infection, and it can be instrumental for the dissecting the immune response to vaccination.

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334 Author Contributions

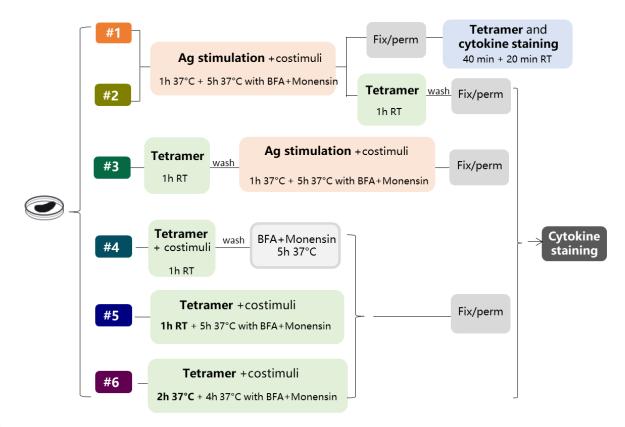
- GP, DM, AC conceived and designed the experiments; GP, MC, EP: performed the experiments; GP,
- 336 MC, AC, EP, EN: analyzed the data; GP, AC: wrote the paper; DM, AC: critically revised the
- 337 manuscript. All authors read and approved the final manuscript.

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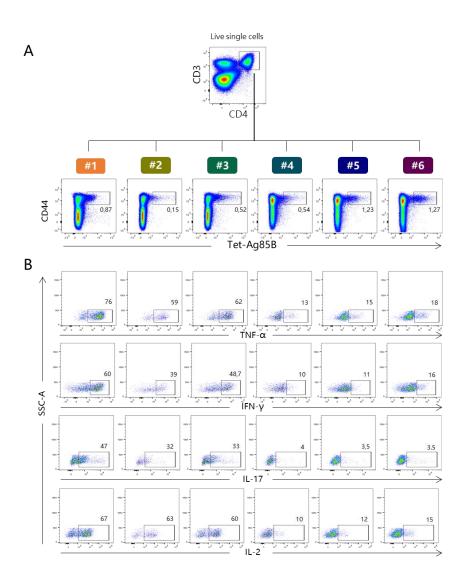
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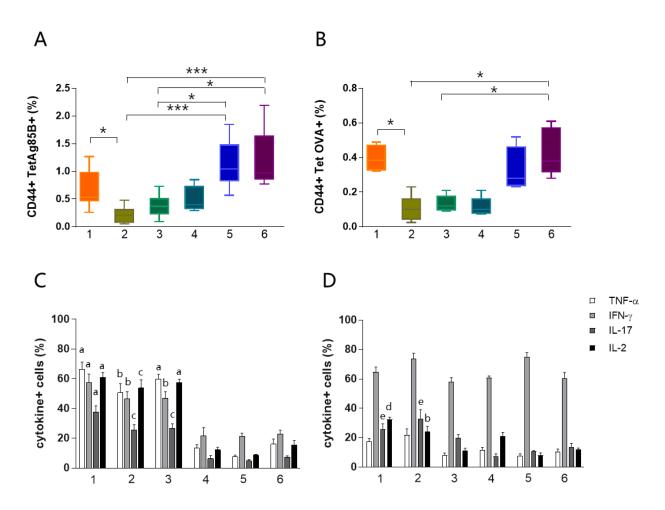


437 Figure 1. Study design. Six different protocols, combining antigen stimulation and tetramer staining with intracellular cytokine labelling, were used for detecting antigen-specific CD4+ T cell producing cytokines in 438 splenocytes of mice immunized with two different antigens, H56 or OVA, and CAF01 adjuvant, 5 days after 439 booster immunization. In protocols 1-3 splenocytes were restimulated with the respective antigen (Ag, pink 440 441 box), added before (protocols 1 and 2) or after (protocol 3) tetramer staining (green box), while in protocols 442 4-6 the restimulation step was performed directly with Ag85B or OVA epitope-complexed MHC II tetramers. Anti-CD28 and anti-CD49d (co-stimuli) were added with Ag (protocol 1-3) or with tetramers (protocols 4-6). 443 444 After Ag or tetramer incubation, cells were treated with Brefeldin A and Monensin for 4-5 h at 37°C, fixated 445 and permeabilized (gray box) and finally stained with anti-cytokines antibodies (dark gray box), except for 446 protocol 1, in which tetramer staining was performed in fixed and permeabilized cells together with cytokine staining. In protocol 6, tetramer staining was performed for 2 h at 37°C. 447

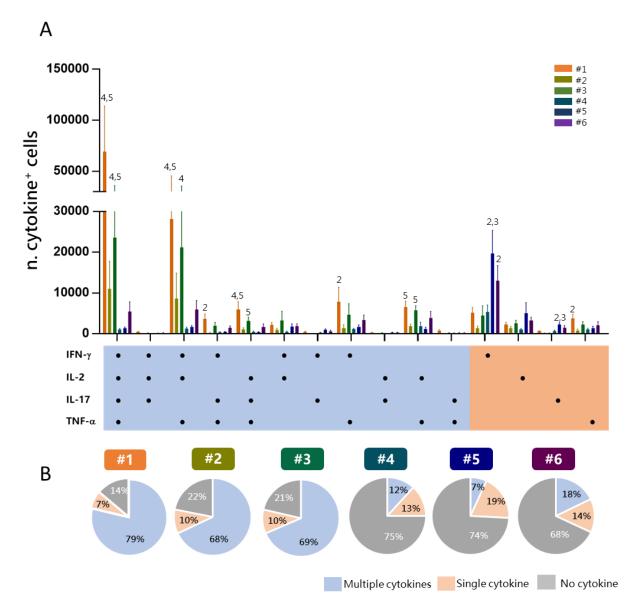


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Figure 2. Flow cytometric analysis of Tet-Ag85B+ T cells producing cytokines. A. Ag85B-tetramer binding T cells were identified among live single CD3+ CD4+, as CD44high Tet-Ag85B+ cells in the six different protocols, and the frequencies of positive cells are reported within the dot plots. B. Intracellular production of TNFα, IFNγ, IL-17 and IL-2 cytokines assessed within the Tet-Ag85B+ T cells in the six different protocols. Frequencies of positive cells are reported within the dot plots. Gates were defined on the respective FMO controls.



456 Figure 3. Identification of tetramer-specific CD4+ T cells and their cytokine production. Tetramer-457 specific CD4+ T cells and their cytokine production were assessed in splenocytes treated with the different 458 protocols reported in Figure 1. A-B. Box plots of the frequencies of Tet-Ag85B+ (A) and Tet-OVA+ (B) T 459 cells respect to CD4+ T cells, detected employing protocols 1-6, as reported in x axis. Values are reported as 460 mean \pm SEM of 10-12 mice, obtained in three independent experiments. Kruskal-Wallis test, followed by Dunn's post test for multiple comparisons, was used to assess the statistical difference between protocols (*P 461 ≤ 0.05 , **P ≤ 0.01 , ***P ≤ 0.001). C-D. Frequencies of TNF- α , IFN γ , IL-17 and IL-2 positive cells among 462 Tet-Ag85B+ (C) and Tet-OVA+ (D) cells, employing protocols 1-6, as reported in x axis. Values are reported 463 464 as mean \pm SEM of 10-12 mice, obtained in three independent experiments. The significant difference between each cytokine among the different protocols, according to the Kruskal-Wallis test followed by Dunn's post test 465 for multiple comparisons (P \leq 0.05), is reported with letters above the error bars; "a", significant difference 466 467 versus protocols 4, 5 and 6; "b" versus protocol 5; "c" versus protocols 4 and 5; "d" versus protocols 3, 5 and 468 6; "e" versus protocol 4.



470 Figure 4. Multifunctional response of Tet-Ag85B+ T cells. Multifunctional profiles of Tet-Ag85B+ T cells detected by the different experimental procedures. A. Histograms represent the number of Tet-Ag85B+ T cells 471 producing different combinations of cytokines shown on the x axis, detected employing the different protocols. 472 473 Responses are grouped and color coded according to the functionality (orange for single cytokine, light blu for 474 two or more cytokines). Values are reported as mean ± SEM of 10-12 mice, obtained in three independent experiments, and the numbers above the error bars indicate which protocols are significantly different 475 according to the Kruskal-Wallis test, followed by Dunn's post test for multiple comparisons ($P \le 0.05$). B. Pie 476 charts of the 6 protocols, in which each slice of the pie represents the fraction of Tet-Ag85B+T cells producing 477 478 two or more cytokines (multiple cytokines, light blue), a single one (orange) or none (grey). Frequencies are 479 reported within each slice.

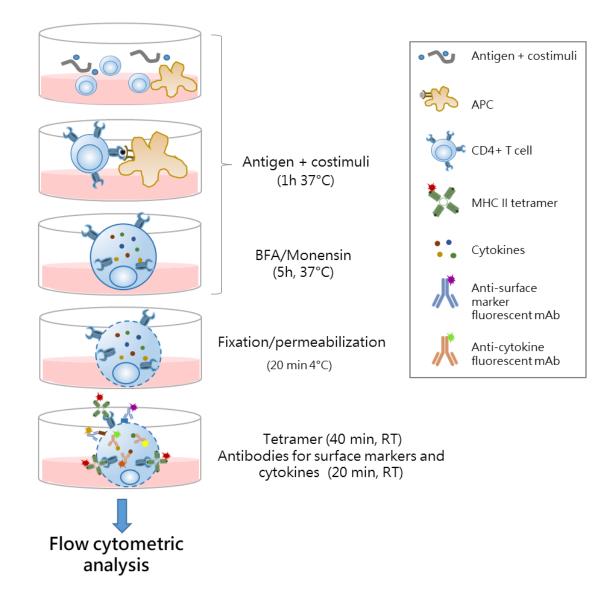


Figure 5. Optimal procedure for identifying multifunctional tetramer-specific CD4+ T cells. Schematic 481 482 overview of the protocol optimized for the detection of multifunctional epitope-specific CD4+ T cells. 483 Splenocytes are cultured in 96-well plates with antigen and costimuli for 1h at 37°C in order to allow antigen 484 presentation by APC to cognate epitope-specific CD4+ T cells. Antigen stimulation elicits reactivation of 485 effector function of CD4+ cells, and TCR internalization. Brefeldin A (BFA) and monensin solution are added for the last 5 h of incubation to block cytokines secretion. Cells are fixed and permabilized for 20 min at 4°C, 486 and then simultaneously stained with MHC II tetramers (1 h, at RT) and surface markers/cytokine-specific 487 antibodies (last 20 minutes). This allows to detect both surface expressed and internalized TCR molecules, 488 489 with intracellular cytokines. The single-step staining procedure allow to reduce the complex time of the 490 protocol. Stained samples are then analysed by flow cytometry.