

T-Select MHC Tetramer

HLA-A*02:01 WT1₁₂₆₋₁₃₄ Tetramer -RMFPNAPYL (50 tests)

For Research Use Only. Not for use in diagnostic procedures.

These T-Select MHC Tetramers use patented technology (Japanese patent No. P4422903).

Background

T lymphocytes play a central role in immune system. Total T cell and T cell subset counts are measured by detection of various cell surface molecules. Enumeration of CD8⁺ antigen-specific T cells requires cognate recognition of the T cell receptor (TCR) by a class I MHC/peptide complex. This can be done using T-Select MHC class I Tetramers which are composed of four MHC class I molecules each bound to the specific peptide and conjugated with a fluorescent protein. Thus, T-Select MHC Tetramer assays allow quantitation of the total T cell population specific for a given peptide complexed with a particular MHC molecule. Furthermore, since binding does not depend on functional pathways, this population includes specific CD8⁺ T cells regardless of functional status. Measurements may be performed in whole blood or isolated lymphocyte/mononuclear cell preparations. In some cases where frequency is low, it may be necessary to perform an *in vitro* cell expansion. Specific cell staining is accomplished by incubating the sample with the T-Select MHC Tetramer reagent, then washing away excess Tetramer. The number of Tetramer positive lymphocytes is then determined by flow cytometry.

This Tetramer reagent comprises human class I HLA-A*02:01 and WT1-derived epitope peptide, WT1₁₂₆₋₁₃₄, and it can detect HLA-A*02:01-restricted WT1₁₂₆₋₁₃₄-specific CD8⁺ T cells.

Wilms' tumor gene 1 (WT1) is a zinc finger transcription factor with limited expression in normal adult tissues, but is overexpressed in the majority of leukemias and various types of solid tumors. In 2009, WT1 was ranked first in a list of 75 representative cancer antigens in a National Cancer Institute prioritization project. Many clinical trials of cancer immunotherapy targeting the WT1 have been carried out around the world.

The RMFPNAPYL peptide from WT1 is presented by both HLA-A*02:01 and HLA-A*02:06 in humans, as well as by H-2D^b in C57BL/6 mice, and has been shown to be the target of WT1-specific responses in both species.

A Tetramer, which is constructed with the same allele (HLA-A*02:01) of interest and an irrelevant peptide, may be used as a negative control Tetramer.

HLA Restriction: HLA-A*02:01

Origin and Sequence of CTL Epitope
WT1 (126-134 aa, RMFPNAPYL)

Reagents

500 µL liquid - 10 µL/test

The Tetramer is dissolved in an aqueous buffer containing 0.5 mM EDTA, 0.2% BSA, 10 mM Tris-HCl (pH 8.0), 150 mM NaCl, and 0.09% NaN₃.

Conjugates

TS-M016-1

Streptavidin-Phycoerythrin (SA-PE)

Excites at 486-580 nm

Emits at 586-590 nm

TS-M016-2

Streptavidin-Allophycocyanin (SA-APC)

Excites at 633-635 nm

Emits at 660-680 nm

Storage Conditions

Store at 2 to 8°C. Do not freeze. Minimize exposure to light. The expiration date is indicated on the vial label.

Evidence of Deterioration

Any change in the physical appearance of this reagent may indicate deterioration and the reagent should not be used. The normal appearance is a clear, colorless to pink (SA-PE), or light blue (SA-APC).

Usage

This reagent is for use with standard flow cytometry methodologies.

References for Products

- 1) Call KM, et al. *Cell* **60**: 509-520 (1990)
- 2) Gessler M, et al. *Nature* **343**: 774-778 (1990)
- 3) Oka Y, et al. *Immunogenetics* **51**: 99-107 (2000)
- 4) Oka Y, et al. *J Immunol* **164**: 1873-1880 (2000)
- 5) Bellantuono I, et al. *Blood* **100**: 3835-3837 (2002)
- 6) Oka Y, et al. *Int J Hematol* **78**: 56-61 (2003)

- 7) Doubrovina ES, et al. *Clin Cancer Res* **10**: 7207-7219 (2004)
- 8) Oka Y, et al. *Curr Med Chem* **13**: 2345-2352 (2006)
- 9) Thomas S, et al. *J Immunol* **179**: 5803-5810 (2007)
- 10) Kobayashi M, et al. *J Ovarian Res* **7**: 48 (2014)

High Specificity

The T cell surface CD8 enhances T cell antigen recognition by binding to HLA class I molecules. Therefore, MBL produced T-Select MHC class I human Tetramers with one point mutation at the HLA α3 domain known to alter the interaction with CD8. These mutated Tetramers showed a greatly diminished nonspecific binding but retained specific binding. Alterations of CD8 binding by mutation of the MHC greatly improved the specificity of MHC-peptide multimers, thus providing efficient tools to sort specific human T cells for immunotherapy.

(French application Number; FR9911133)

References for T-Select MHC Tetramer

- Altman JD, et al. *Science* **274**: 94-96 (1996)
McMichael AJ, et al. *J Exp Med* **187**: 1367-1371 (1998)
Bodinier M, et al. *Nat Med* **6**: 707-710 (2000)

Statement of Warnings

1. This reagent contains 0.09% sodium azide. Sodium azide under acid conditions yields hydrazoic acid, an extremely toxic compound. Azide compounds should be flushed with running water while being discarded. These precautions are recommended to avoid deposits in metal piping in which explosive conditions can develop. If skin or eye contact occurs, wash excessively with water.
2. Specimens, samples and material coming in contact with them should be handled as if capable of transmitting infection and disposed of with proper precautions.
3. Never pipette by mouth and avoid contact of samples with skin and mucous membranes.
4. Minimize exposure of reagent to light during storage or incubation.
5. Avoid microbial contamination of reagent or erroneous results may occur.
6. Use Good Laboratory Practices (GLP) when handling this reagent.

Materials Required But Not Supplied

- 12 x 75 mm polypropylene test tubes
- Transfer pipettes
- Pipettors and disposable pipette tips
- Vortex mixer
- Centrifuge capable of 150 x g or 400 x g
- Aspirator
- PBS
- Red blood cell lysis reagent
- Anti-CD8-FITC (T8), Beckman Coulter, Inc., PN 6603861
- Anti-CD8-PC5 (T8), Beckman Coulter, Inc., PN 6607011
- 7-AAD Viability Dye, Beckman Coulter, Inc., PN

A07704

- Clear Back (human FcR blocking reagent), MBL, PN MTG-001

Procedure for Whole Blood

1. Collect blood by venipuncture into a blood collection tube containing an appropriate anti-coagulant.
2. Add 10 µL of T-Select MHC Tetramer to each 12 x 75 mm test tube.
3. Add 200 µL of whole blood into each test tube.
4. Vortex gently.
5. Incubate for 30-60 minutes at 2-8°C or room temperature (15-25°C) protected from light.
6. Add any additional antibodies (e.g. anti-CD8) and vortex gently.
7. Incubate for 30 minutes at 2-8°C protected from light.
8. Lyse red blood cells using commercially available reagents.
9. Prepare samples according to description of the package insert.
10. Analyze prepared samples by flow cytometry. If necessary, store the samples at 2-8°C protected from light for a maximum of 24 hours prior to analysis.

Procedure for Peripheral Blood Mononuclear Cells

1. Prepare peripheral blood mononuclear cells (PBMC) according to established procedures. Cells should be re-suspended at a concentration of 2 x 10⁷ cells/mL. 50 µL of sample is required for each T-Select MHC Tetramer determination.
2. Add 10 µL of Clear Back (human FcR blocking reagent, MBL, PN MTG-001) to each 12 x 75 mm test tube.
3. Add 50 µL PBMC into each test tube (e.g. 1 x 10⁶ cells per tube).
4. Incubate for 5 minutes at room temperature.
5. Add 10 µL of T-Select MHC Tetramer and vortex gently.
6. Incubate for 30-60 minutes at 2-8°C or room temperature (15-25°C) protected from light.
7. Add any additional antibodies (e.g. anti-CD8) and vortex gently.
8. Incubate for 30 minutes at 2-8°C protected from light.
9. Add 3 mL of PBS or FCM buffer (2% FCS/0.09% NaN₃/PBS).
10. Centrifuge tubes at 400 x g for 5 minutes.
11. Aspirate or decant the supernatant.
12. Resuspend the pellet in 500 µL of PBS with 0.5% formaldehyde.
13. Analyze prepared samples by flow cytometry. If necessary, store the samples at 2-8°C protected from light for a maximum of 24 hours prior to analysis.

Limitations

1. For optimal results with whole blood, retain specimens in blood collection tubes at room temperature, while rocking, prior to staining and analyzing. Refrigerated specimens may give aberrant results.
2. Recommended cell viability for venous blood specimens is > 90%.
3. Prolonged exposure of cells to lytic reagents may cause white blood cell destruction and loss of cells in the population of interest.
4. All red blood cells may not lyse under the following conditions: nucleated red blood cells, abnormal protein concentration or hemoglobinopathies. This may cause falsely decreased results due to unlysed red blood cells being counted as leukocytes.

Technical Hints

- A. If PBMC culture is needed, we recommend the use of heparin as an anti-coagulant.
- B. Clear Back reagent (human FcR blocking reagent) may effectively block non-specific binding caused by macrophages or endocytosis, resulting in clear staining when cells are stained with MHC Tetramer and antibodies. Please refer to the data sheet (MBL, PN MTG-001) for details.
- C. A Tetramer that is constructed with the same allele of interest and an irrelevant peptide may be used as a negative control.
- D. We recommend the use of anti-CD8 antibody, clone SFCI21Thy2D3 (T8, Beckman Coulter, Inc.), because some anti-CD8 antibodies inhibit Tetramer-specific binding to TCR.
- E. The use of CD45 antibody and gating of the lymphocyte population are recommended in order to reduce contamination of unlysed or nucleated red blood cells in the gate.
- F. Apoptotic, necrotic, and/or damaged cells are sources of interference in the analysis of viable cells by flow cytometry. Cell viability should be determined by 7-aminoactinomycin D (7-AAD) staining; intact viable cells remain unstained (negative).
- G. Cells do not require fixation prior to analysis if the stained cells are analyzed by flow cytometry within several hours.

Related Products

WT1 Tetramers and Peptide

- | | |
|-----------|---|
| TS-M140-1 | HLA-A*02:01 WT1 ₃₇₋₄₅ Tetramer-VLDFAPPAGA-PE |
| TS-M140-2 | HLA-A*02:01 WT1 ₃₇₋₄₅ Tetramer-VLDFAPPAGA-APC |
| TS-M140-P | HLA-A*02:01 WT1 ₃₇₋₄₅ peptide |
| TS-M016-1 | HLA-A*02:01 WT1 ₁₂₆₋₁₃₄ Tetramer-RMFPNAPYL-PE |
| TS-M016-2 | HLA-A*02:01 WT1 ₁₂₆₋₁₃₄ Tetramer-RMFPNAPYL-APC |
| TS-M014-1 | HLA-A*24:02 modified WT1 Tetramer-CYTWNQMLN-PE |
| TS-M014-2 | HLA-A*24:02 modified WT1 Tetramer-CYTWNQMLN-APC |
| TS-M504-1 | H-2D ^b WT1 ₁₂₆₋₁₃₄ Tetramer-RMFPNAPYL-PE |
| TS-M504-2 | H-2D ^b WT1 ₁₂₆₋₁₃₄ Tetramer-RMFPNAPYL-APC |

Human Tetramers

Cancer

- TS-M141-1 HLA-A*24:02 ACC-1 Tetramer-DYLQYVLQI-PE
 TS-M137-1 HLA-A*01:01 AIM-2 Tetramer-RSDSGQQARY-PE
 TS-M112-1 HLA-A*24:02 CA9₂₁₉₋₂₂₇ Tetramer-EYRALQLHL-PE
 TS-M103-1 HLA-A*02:01 CEA Tetramer-YLSGANLNL-PE
 TS-M080-1 HLA-A*02:01 CEA (N6D) Tetramer-YLSGADLN-PE
 TS-M101-1 HLA-A*02:01 CD33 Tetramer-AIISGDSPV-PE
 TS-M102-1 HLA-A*02:01 CD33 A65Y Tetramer-YIISGDSPV-PE
 TS-M084-1 HLA-A*02:01 EphA2 Tetramer-TLADFDPRV-PE
 TS-0014-1C HLA-A*02:01 gp100 (wild) Tetramer-ITDQVPFSV-PE
 TS-0013-1C HLA-A*02:01 gp100 (mutant) Tetramer-IMDQVPFSV-PE
 TS-0035-1C HLA-A*02:01 gp100₁₅₄₋₁₆₂ Tetramer-KTWGQYWQV-PE
 TS-M082-1 HLA-A*02:01 gp100 Tetramer-YLEPGPVTA-PE
 TS-M089-1 HLA-A*24:02 gp100-intron 4 Tetramer-VYFFLPDH-PE
 TS-0016-1 HLA-A*02:01 Her-2/heu Tetramer-RLLQETELV-PE
 TS-0015-1C HLA-A*02:01 Her-2/heu E75 Tetramer-KIFGSLAFL-PE
 TS-M083-1 HLA-A*02:01 HM1.24 Tetramer-KLQDASAEV-PE
 TS-M010-1 HLA-A*24:02 hTERT Tetramer-VYGFVRACL-PE
 TS-M115-1 HLA-A*02:01 hTERT Tetramer-ILAKFLHWL-PE
 TS-M086-1 HLA-A*02:01 IDO Tetramer-ALLEIASCL-PE
 TS-M070-1 HLA-A*02:01 MAGE-A1 Tetramer-KVLEYVIK-PE
 TS-M071-1 HLA-B*07:02 MAGE-A1 Tetramer-RVRRFFPSL-PE
 TS-M072-1 HLA-A*02:01 MAGE-A2 Tetramer-YLQLVFGIEV-PE
 TS-M073-1 HLA-A*24:02 MAGE-A2 Tetramer-EYQLQVFGI-PE
 TS-M075-1 HLA-A*02:01 MAGE-A3₁₁₂₋₁₂₀ Tetramer-KVAELVHFL-PE
 TS-M076-1 HLA-A*02:01 MAGE-A3₂₇₁₋₂₇₉ Tetramer-FLWGPRLV-PE
 TS-M077-1 HLA-A*24:02 MAGE-A3 Tetramer-IMPKAGLLI-PE
 TS-M078-1 HLA-A*02:01 MAGE-A10 Tetramer-GLYDGMEHL-PE
 TS-M138-1 HLA-A*02:01 MAGE-C1 Tetramer-ILFGISLREV-PE
 TS-0009-1C HLA-A*02:01 Mart-1 Tetramer-ELAGIGILTV-PE
 TS-M091-1 HLA-A*24:02 MCPyV large Tag Tetramer-EWWRSGGFSF-PE
 TS-M088-1 HLA-A*02:01 MUC1 Tetramer-LLLTVLTV-PE
 TS-M011-1 HLA-A*02:01 NY-ESO-1 Tetramer-SLLMMVITQC-PE
 TS-M105-1 HLA-A*02:01 NY-ESO-1 C9V Tetramer-SLLMMVITQV-PE
 TS-M109-1 HLA-B*07:02 P2X5 Tetramer-TPNQRQNVC-PE
 TS-M081-1 HLA-A*02:01 p53 Tetramer-LLGRNSFEV-PE
 TS-M107-1 HLA-A*02:01 PAP₂₉₉₋₃₀₇ Tetramer-ALDVYNGLL-PE
 TS-M136-1 HLA-A*24:02 PBF A24.2 Tetramer-AYRPVSRNI-PE
 TS-M117-1 HLA-A*02:01 PRAME₁₀₀₋₁₀₈ Tetramer-VLDGLDVLL-PE
 TS-M119-1 HLA-A*02:01 PRAME₁₄₂₋₁₅₁ Tetramer-SLYSFPEPEA-PE
 TS-M116-1 HLA-A*02:01 PRAME₃₀₀₋₃₀₉ Tetramer-ALYVDSLFFL-PE
 TS-M118-1 HLA-A*02:01 PRAME₄₂₅₋₄₃₃ Tetramer-SLLQHLIGL-PE
 TS-M120-1 HLA-A*02:01 PSA₁₄₁₋₁₅₀ Tetramer-FLTPKKLQCV-PE
 TS-0017-1 HLA-A*02:01 PR-1 Tetramer-VLQELNVTV-PE
 TS-M087-1 HLA-A*02:01 PSA Tetramer-KLQCVDLHV-PE
 TS-M104-1 HLA-A*02:01 RHAMM Tetramer-ILSLELMKL-PE
 TS-M095-1 HLA-A*02:01 PP2A Tetramer-SLLPAIVEL-PE
 TS-M079-1 HLA-A*02:01 SSX-2 Tetramer-KASEKIFYV-PE
 TS-M025-1 HLA-A*24:02 survivin-2B Tetramer-AYACNTSTL-PE
 TS-M085-1 HLA-A*02:01 Survivin (T2M) Tetramer-LMLGEFLKL-PE
 TS-0019-1C HLA-A*02:01 Tyrosinase Tetramer-YMDGTMMSQV-PE
 TS-M090-1 HLA-A*24:02 Tyrosinase Tetramer-AFLPWHLF-PE

Adenovirus

- TS-M058-1 HLA-A*02:01 Adv11 Hexon₉₁₃₋₉₂₁ Tetramer-YLLFEVFDV-PE
 TS-M059-1 HLA-A*02:01 Adv11 Hexon₉₁₄₋₉₂₂ Tetramer-LLFVFDFDV-PE
 TS-M061-1 HLA-A*02:01 AdV Hexon₉₁₇₋₉₂₅ Tetramer-YLFVFDFV-PE
 TS-M062-1 HLA-A*24:02 Adv11 Hexon₃₇₋₄₅ Tetramer-TYFNLGNKF-PE
 TS-M064-1 HLA-A*24:02 Adv11 Hexon₆₉₆₋₇₀₄ Tetramer-VYSGSIPYL-PE
 TS-M063-1 HLA-A*24:02 Adv5 Hexon₃₇₋₄₅ Tetramer-TYFSLNNKF-PE
 TS-M067-1 HLA-B*35:01 AdV Hexon₁₂₀₋₁₂₉ Tetramer-MPNRPNHYIAF-PE
 TS-M068-1 HLA-B*35:01 AdV Hexon₇₀₅₋₇₁₃ Tetramer-IPYLDGTFY-PE
 TS-M065-1 HLA-B*07:02 AdV Hexon₁₁₄₋₁₂₄ Tetramer-KPYSGTAYNSL-PE
 TS-M066-1 HLA-B*07:02 AdV Hexon₁₁₄₋₁₂₄ Tetramer-KPYSGTAYNAL-PE

CMV

- TS-M057-1 HLA-A*02:01 CMV IE1₃₁₆₋₃₂₄ Tetramer-VLEETSVML-PE
 TS-M100-1 HLA-A*03:01 CMV IE1₁₈₄₋₁₉₂ Tetramer-KLGGAQAK-PE
 TS-0026-1C HLA-B*08:01 CMV IE1 Tetramer-ELRRKMMYMY-PE
 TS-0024-1C HLA-A*01:01 CMV pp50 Tetramer-VTEHDTLLY-PE

TS-0010-1C	HLA-A*02:01 CMV pp65 Tetramer-NLVPVMVATV-PE
TS-0020-1C	HLA-A*24:02 CMV pp65 Tetramer-QYDPVAALF-PE
TS-M012-1	HLA-A*11:01 CMV pp65 Tetramer-ATVGQQLNLK-PE
TS-M099-1	HLA-B*07:02 CMV pp65 Tetramer-RPHERNGFTVL-PE
TS-0025-1C	HLA-B*07:02 CMV pp65 Tetramer-TPRVTGGGAM-PE
TS-0027-1C	HLA-B*35:01 CMV pp65 Tetramer-IPSINVHHY-PE
TS-M013-1	HLA-B*15:01 CMV pp65 Tetramer-KMQVIGDQY-PE

EBV

TS-0011-1C	HLA-A*02:01 EBV BMLF1 Tetramer-GLCTLVAML-PE
TS-M003-1	HLA-A*24:02 EBV BMLF1 Tetramer-DYNFVKQLF-PE
TS-M002-1	HLA-A*24:02 EBV BRLF1 Tetramer-TYPVLEEMF-PE
TS-M124-1	HLA-A*03:01 EBV BRLF1 Tetramer-RVRAYTYSK-PE
TS-M036-1	HLA-B*08:01 EBV BZLF1 ₁₉₀₋₁₉₇ Tetramer-RAKFKQLL-PE
TS-M037-1	HLA-B*35:01 EBV BZLF1 ₅₄₆₄ Tetramer-EPLPQGQLTAY-PE
TS-M033-1	HLA-A*03:01 EBV EBNA3A ₃₃₉₋₃₆₁ Tetramer-RLRAEAQVK-PE
TS-M004-1	HLA-A*24:02 EBV EBNA3A Tetramer-RYSIFFDYM-PE
TS-M123-1	HLA-B*08:01 EBV EBNA3A Tetramer-FLRGRAYGL-PE
TS-M028-1	HLA-A*11:01 EBV EBNA3B ₃₉₉₋₄₀₈ Tetramer-AV/FDRKSDAK-PE
TS-M029-1	HLA-A*11:01 EBV EBNA3B ₄₁₆₋₄₂₄ Tetramer-IVTDFSVIK-PE
TS-M005-1	HLA-A*24:02 EBV EBNA3B Tetramer-TYSAGIVQI-PE
TS-M006-1	HLA-A*02:01 EBV LMP1 Tetramer-YLQQNWVTL-PE
TS-M030-1	HLA-A*02:01 EBV LMP2 ₂₄₃₋₂₅₁ Tetramer-TVCGGIMFL-PE
TS-M031-1	HLA-A*02:01 EBV LMP2 ₃₂₉₋₃₃₇ Tetramer-LLWTLVLLVLL-PE
TS-M069-1	HLA-A*02:01 EBV LMP2 ₃₅₆₋₃₆₄ Tetramer-FLYALALLL-PE
TS-M032-1	HLA-A*02:01 EBV LMP2 ₄₂₆₋₄₃₄ Tetramer-CLGGLLTMV-PE
TS-M034-1	HLA-A*24:02 EBV LMP2 ₁₃₁₋₁₃₉ Tetramer-PYLFWLAAI-PE
TS-M001-1	HLA-A*24:02 EBV LMP2 Tetramer-IYVLVMLVL-PE
TS-M035-1	HLA-A*24:02 EBV LMP2 ₄₁₉₋₄₂₇ Tetramer-TYGPVFMSL-PE
TS-M038-1	HLA-B*35:01 EBV LMP2 ₁₋₉ Tetramer-MGSLEMPVM-PE
TS-M111-1	HLA-A*11:01 EBV LMP2 Tetramer-SSCSSCPLSK-PE
TS-M135-1	HLA-A*11:01 EBV LMP2 S9T Tetramer-SSCSSCPLTK-PE
TS-M009-1	HLA-A*24:02 EBV Mix Tetramer-PE

HBV

TS-0018-1C	HLA-A*02:01 HBV core Tetramer-FLPSDFFPSV-PE
TS-0022-1C	HLA-A*24:02 HBV core Tetramer-EYLVSFGVW-PE
TS-M051-1	HLA-A*02:01 HBV env ₃₃₅₋₃₄₃ Tetramer-WLSLLVPFV-PE
TS-M052-1	HLA-A*02:01 HBV env ₃₄₈₋₃₅₇ Tetramer-GLSPTVWLSV-PE
TS-M053-1	HLA-A*02:01 HBV pol Tetramer-FLLSLGIHL-PE
TS-0023-1C	HLA-A*24:02 HBV pol Tetramer-KYTSFPWLL-PE

HCV

TS-M044-1	HLA-A*24:02 HCV E2 ₇₁₇₋₇₂₅ Tetramer-EYVLLLFL-PE
TS-M039-1	HLA-A*02:01 HCV NS3 ₁₀₇₃₋₁₀₈₁ Tetramer-CINGVCWTV-PE
TS-M040-1	HLA-A*02:01 HCV NS3 ₁₄₀₆₋₁₄₁₅ Tetramer-KLVALGINAV-PE
TS-M041-1	HLA-A*02:01 HCV NS4B ₁₉₂₂₋₂₀₀₀ Tetramer-VLSDFKTWL-PE
TS-M042-1	HLA-A*02:01 HCV NS5B ₂₅₉₄₋₂₆₀₂ Tetramer-ALYDVVTKL-PE
TS-M043-1	HLA-A*02:01 HCV NS5B ₂₅₉₄₋₂₆₀₂ Tetramer-ALYDVVSKL-PE

HIV

TS-M007-1	HLA-A*24:02 HIV env Tetramer-RYLRDQQQL-PE
TS-M027-1	HLA-A*02:01 HIV gag ₇₇₋₈₅ Tetramer-SLYNTVATL-PE
TS-M139-1	HLA-A*02:01 HIV gag ₉₋₂₇ Tetramer-TLNAAWVKKV-PE
TS-M110-1	HLA-A*24:02 HIV nef ₁₃₄₋₁₄₁ Tetramer-RYPLTFGW-PE
TS-M054-1	HLA-B*07:02 HIV nef Tetramer-TPGPGVRYPL-PE
TS-M106-1	HLA-B*35:01 HIV nef ₇₄₋₈₁ Tetramer-VPLRPMTY-PE
TS-0008-1C	HLA-A*02:01 HIV pol Tetramer-ILKEPVHGV-PE
TS-M055-1	HLA-B*35:01 HIV RT Tetramer-NPDIVIYQY-PE

HPV

TS-0031-1	HLA-A*02:01 HPV16 E7 Tetramer-YMLDLQPET-PE
TS-M047-1	HLA-A*02:01 HPV16 E6 Tetramer-KLPQLCTEL-PE
TS-M049-1	HLA-A*24:02 HPV16 E6 Tetramer-VYDFAFRDL-PE
TS-M048-1	HLA-A*02:01 HPV16 E7 Tetramer-YMLDLQPETT-PE

HTLV

TS-M022-1	HLA-A*24:02 HTLV-1 Env ₁₁₋₁₉ Tetramer-FFQFCPLIF-PE
TS-M017-1	HLA-A*02:01 HTLV-1 Tax ₁₁₋₁₉ Tetramer-LLFGYPVYV-PE
TS-M019-1	HLA-A*02:01 HTLV-1 Tax ₁₇₈₋₁₈₆ Tetramer-QLGAFLTNV-PE
TS-M024-1	HLA-A*11:01 HTLV-1 Tax ₂₇₂₋₂₈₀ Tetramer-QSSSFIFHK-PE
TS-M023-1	HLA-A*11:01 HTLV-1 Tax ₃₈₉₋₃₉₆ Tetramer-KVLTTPPIH-PE
TS-M020-1	HLA-A*24:02 HTLV-1 Tax ₁₂₂₀ Tetramer-LFGYPVYVF-PE
TS-M021-1	HLA-A*24:02 HTLV-1 Tax ₁₈₇₋₁₉₅ Tetramer-PYKRIEELL-PE
TS-M018-1	HLA-A*24:02 HTLV-1 Tax ₃₀₁₋₃₀₉ Tetramer-SFHSLHLLF-PE

Influenza

TS-M045-1	HLA-A*01:01 Influenza NP Tetramer-CTELKLSDY-PE
TS-M046-1	HLA-B*35:01 Influenza NP Tetramer-LPFEKSTVM-PE
TS-0012-1C	HLA-A*02:01 Influenza M1 Tetramer-GILGFVFTL-PE

Measles virus

TS-M092-1	HLA-A*02:01 measles virus HA Tetramer-KLWCRHFCV-PE
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RSV

TS-M056-1	HLA-A*01:01 RSV M ₂₂₉₋₂₃₇ Tetramer-YLEKESIYY-PE
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VZV

TS-M122-1	HLA-A*02:01 VZV IE62 Tetramer-ALWALPHAA-PE
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Mycobacterium tuberculosis

TS-M026-1	HLA-A*02:01 Mtb MPT51 Tetramer-TLAGKGISVW-PE
TS-M128-1	HLA-A*02:01 Mtb Ag85A ₄₈₋₅₆ Tetramer-GLPVEYLVQV-PE
TS-M129-1	HLA-A*02:01 Mtb Ag85A ₂₄₂₋₂₅₀ Tetramer-KLIANTRV-PE
TS-M131-1	HLA-A*02:01 Mtb Ag85B Tetramer-KLVANNTRL-PE
TS-M134-1	HLA-B*35:01 Mtb Ag85C Tetramer-WPTLIGLAM-PE
TS-M125-1	HLA-A*02:01 Mtb ESAT-6 Tetramer-AMASTEGNV-PE
TS-M127-1	HLA-A*02:01 Mtb Rv1614 Tetramer-FLYELIWNV-PE
TS-M130-1	HLA-A*02:01 Mtb Hsp65 Tetramer-KLQERLAKL-PE
TS-M132-1	HLA-A*02:01 Mtb 16 kDa Tetramer-GILTWSVAV-PE
TS-M133-1	HLA-A*02:01 Mtb 19 kDa Tetramer-VLTDGNPVEV-PE

Control

TS-M007-1	HLA-A*24:02 Negative Tetramer-RYLRDQQQL-PE
TS-M007-2	HLA-A*24:02 Negative Tetramer-RYLRDQQQL-APC
TS-M007-3	HLA-A*24:02 Negative Tetramer-RYLRDQQQL-FITC
TS-0029-1C	HLA-A*02:01 Negative Tetramer-PE
TS-0029-2C	HLA-A*02:01 Negative Tetramer-APC

Others

TS-M097-1	HLA-A*02:01 BTG1 Tetramer-TLWVDPYEV-PE
TS-M093-1	HLA-A*02:01 HA-1 Tetramer-VLHDDLLEA-PE
TS-M108-1	HLA-A*02:01 HA-2 Tetramer-YIGEVLVSV-PE
TS-M098-1	HLA-A*02:01 HA-8 Tetramer-RTLDKVLEV-PE
TS-M094-1	HLA-A*02:01 H-Y Tetramer-FIDSYICQV-PE
TS-M096-1	HLA-A*02:01 RNA helicase Tetramer-YLLPAIVHI-PE

Please check our web site (<http://ruo.mbl.co.jp>) for up-to-date information on products and custom MHC Tetramers.

T-Select MHC Tetramers use patented technology (US patent No. 5,635,363, French application No. FR9911133, and Japanese patent No. P3506384) of Beckman Coulter Inc..

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