

Identification of Lck-derived peptides applicable to anti-cancer vaccine for patients with human leukocyte antigen-A3 supertype alleles

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The identification of peptide vaccine candidates to date has been focused on human leukocyte antigen (HLA)-A2 and -A24 alleles. In this study, we attempted to identify cytotoxic T lymphocyte (CTL)-directed Lck-derived peptides applicable to HLA-A11⁺, -A31⁺, or -A33⁺ cancer patients, because these HLA-A alleles share binding motifs, designated HLA-A3 supertype alleles, and because the Lck is preferentially expressed in metastatic cancer. Twenty-one Lck-derived peptides were prepared based on the binding motif to the HLA-A3 supertype alleles. They were first screened for their recognisability by immunoglobulin G (IgG) in the plasma of prostate cancer patients, and the selected candidates were subsequently tested for their potential to induce peptide-specific CTLs from peripheral blood mononuclear cells of HLA-A3 supertype⁺ cancer patients. As a result, four Lck peptides were frequently recognised by IgGs, and three of them – Lck_{90–99}, Lck_{449–458}, and Lck_{450–458} – efficiently induced peptide-specific and cancer-reactive CTLs. Their cytotoxicity towards cancer cells was mainly ascribed to HLA class I-restricted and peptide-specific CD8⁺ T cells. These results indicate that these three Lck peptides are applicable to HLA-A3 supertype⁺ cancer patients, especially those with metastasis. This information could facilitate the development of peptide-based anti-cancer vaccine for patients with alleles other than HLA-A2 and -A24.

British Journal of Cancer (2007) 97, 1648–1654. doi:10.1038/sj.bjc.6604071 www.bjcancer.com

Published online 27 November 2007

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Keywords: Lck; cytotoxic T lymphocyte; peptide; HLA-A3 supertype

Many tumour antigens and their peptides that are recognised by cytotoxic T lymphocytes (CTLs) have been identified from a variety of cancer types (Renkvist *et al*, 2001), and subsequent clinical applications, including a form of peptide-based immunotherapy, have been performed (Yamada and Itoh, 2006). Most of these peptides are limited to human leukocyte antigen (HLA)-A2 or -A24 alleles, primarily because of the higher worldwide frequency of these alleles (Imanishi *et al*, 1992). Several HLA alleles are known to have structural similarities according to peptide-binding motif analyses, and four superotypes have been proposed: HLA-A2, -A3, -B7, and -B44 supertype alleles (Sette and Sidney, 1999). Among them, the A3 supertype alleles include the allelic products of at least five common HLA-A alleles, including A3, A11, A31, A33, and A68. Thirty-eight percent of Caucasians,

53% of Chinese, 46% of Japanese, and 43% of North American African Americans and Hispanics were positive for the HLA-A3 supertype alleles (Sette and Sidney, 1999). Therefore, the identification of peptide vaccine candidates for HLA-A3 supertype⁺ cancer patients could facilitate the development of peptide-based immunotherapy for many ethnic populations.

The Lck protein (p56^{lck}), the *src* family tyrosine kinase, is known to be essential for both T-cell development and function (Veillette *et al*, 1989). It is notable that the Lck protein is aberrantly expressed in several malignancies, including colon carcinoma, small cell lung carcinoma, and prostate carcinoma with a trend of preferential expression in metastatic lesions (Robinson *et al*, 1996; McCracken *et al*, 1997; Krystal *et al*, 1998; Lutz *et al*, 1998). Although biological roles of the Lck protein in cancer cells have not been fully confirmed, several lines of evidence suggest that this protein contributes to the process of neoplastic transformation (Marth *et al*, 1988; Robinson *et al*, 1996; McCracken *et al*, 1997; Krystal *et al*, 1998; Lutz *et al*, 1998). In support of this idea, several studies have reported that the Lck protein may contribute to the anchorage-independent growth of TGF- β -initiated tumour cells through the transcription of p56^{lck} with a type I promoter

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Received 14 June 2007; revised 9 October 2007; accepted 9 October 2007; published online 27 November 2007

(Amundadottir and Leder, 1998). In addition, we previously reported that the Lck-derived peptides can be recognised by cancer-reactive CTLs of cancer patients, and that Lck peptide-specific CTLs can be induced from patients with distant metastases, but not from those without distant metastases (Harashima *et al*, 2001). In this study, we further attempted to identify novel Lck-derived peptide candidates that would be applicable to cancer patients with HLA-A3 supertype alleles to expand the possibilities of a peptide-based vaccine for metastatic cancer patients with alleles other than HLA-A2 and -A24.

MATERIALS AND METHODS

Patients

Peripheral blood mononuclear cells (PBMCs) were obtained from prostate cancer patients who had provided written informed consent. These patients included HLA-A11⁺, -A31⁺, and -A33⁺ patients, but PBMCs from HLA-A3⁺ or -A68.1⁺ patients were not included because of their extremely low frequency (1.6 and 0.5%) in the Japanese population (Aizawa, 1986). None of the subjects were infected with HIV. Twenty millilitres of peripheral blood was obtained, and PBMCs were prepared by Ficoll-Conray density gradient centrifugation. All of the samples were cryopreserved until they were used for the experiments. The expression of HLA-A11, -A31, and -A33 molecules on PBMCs was determined by flow cytometry using the following antibodies: anti-HLA-A11 monoclonal antibody (mAb) (cat. no. 0284HA; One Lambda Inc., Canoga, CA, USA), anti-HLA-A31 mAb (cat. no. 0273HA; One Lambda), and anti-HLA-A33 mAb (cat. no. 0612HA; One Lambda).

Peptides

The Lck-derived peptides that are provided in Table 1 were prepared on the basis of the binding motifs to the HLA-A11, -A31, and -A33 molecules (Parker *et al*, 1994). All peptides were of > 90% purity and were purchased from the Biologica Co. (Nagoya, Japan). Epstein-Barr virus (EBV)-derived, tyrosinase-related protein 2 (TRP-2)-derived, and HIV-derived peptides were used as controls binding to HLA-A3 supertype alleles. All peptides were dissolved with dimethyl sulphoxide at a dose of 10 mg ml⁻¹.

Measurement of IgGs reactive to Lck peptides

The levels of immunoglobulin Gs (IgGs) reactive to Lck peptides were measured by the Luminex™ method, as reported previously (Komatsu *et al*, 2004). In brief, 100 µl of diluted plasma was incubated with 5 µl of colour-coded beads (Luminex Corp., Austin, TX, USA) coated with each of the Lck peptides on 96-well filter plates (MABVN1250; Millipore Corp., Bedford, MA, USA) for 2 h at room temperature on a plate shaker. The plates were then washed with Tween-PBS and incubated with 100 µl of biotin-conjugated goat anti-human IgG (BA-3080; Vector Laboratories, Burlingame, CA, USA) for 1 h at room temperature on a plate shaker. After the plates were washed, 100 µl of streptavidin-PE was added to the wells, and the samples were incubated for 30 min at room temperature on a plate shaker. The bound beads were washed four times, and 100 µl of Tween-PBS was added to each well. Fifty microlitres of sample was examined using the Luminex system.

Induction of peptide-specific CTLs from PBMCs

The assay for the detection of peptide-specific CTLs was performed according to a previously reported method with several modifications (Hida *et al*, 2002). In brief, PBMCs (1 × 10⁵ cells per well) were incubated with 10 µl ml⁻¹ of each peptide in quadruplicate in a U-bottom-type 96-well microculture plate (Nunc, Roskilde, Denmark) in 200 µl of culture medium. The culture medium

Table 1 Summary of Lck-derived peptide candidates binding to the HLA-A3 supertype alleles

Peptides	Amino-acid sequence	Bind to ^b	Binding score ^a				
			A3	A11	A31	A33	A68
27–35	IVRLDGKGR		0.2	0.2	1	15	200
36–45	LLIRNGSEVR		6	0.12	4	9	10
37–45	LIRNGSEVR		0.4	0.08	2	15	5
90–99	ILEQSGEWWK		60	0.8	1	0.3	4.5
131–139	NLSRKDAER		4	0.08	2	9	7.5
146–154	NTHGSFLIR		1.8	0.4	4	3	50
198–207	TFPGLHELVR		0.012	0.08	1.2	3	0.75
290–299	NLMKQLQHQR		6	0.16	6	9	10
291–299	LMKQLQHQR		4	0.08	2	15	5
293–302	KQLQHQRLLVR		1.08	1.08	24	0.9	10
294–302	QLQHQRLLVR		8	0.16	4	9	5
354–363	FIEERNYIHR		1.2	0.16	4	15	7.5
379–387	KIADFGLAR		7.2	0.48	12	4.5	10
388–397	LIEDNEYTAR		0.4	0.08	2	15	5
429–438	LLTEIVTHGR		9	0.08	2	9	5
430–438	LTEIVTHGR		0.3	0.2	1	3	50
449–458	VIQNLERGYR		0.12	0.08	2	15	5
450–458	IQNLERGYR		0.036	0.12	2	3	5
452–461	NLERGYRMYR		24	0.16	4	9	5
471–480	QLMRLCWKER		3	0.08	3	9	10
472–480	LMRLCWKER		2	0.04	1	15	
EBV	IVTDFSVIK	A11	10.0	4.0	0.6	0.5	240
Flu	NVKNLYEKVK	A11	3.0	1.0	0.1	0.5	180
TRP-2	LLGPGRPYR	A31/ A33	6.0	0.1	2.0	9.0	15
HIV	RLRDLIIIVTR	A31	—	—	—	—	—

EBV = Epstein-Barr virus; HLA = human leukocyte antigen; TRP-2 = tyrosinase-related protein 2. ^aThe peptide-binding score was calculated based on the predicted half time of dissociation from HLA class I molecules as obtained from a website (Bioinformatics and Molecular Analysis Section, Computational Bioscience and Engineering Laboratory, Division of Computer Research and Technology, NIH). The binding score of the HIV peptide was not calculated because the peptide consisted of 11-mer amino acids. ^bPreviously reported HLA class I alleles in which the peptides have immunogenicity are shown.

consisted of 45% RPMI 1640, 45% AIM-V medium (Gibco BRL, Gaithersburg, MD, USA), 10% FCS, 100 U ml⁻¹ of interleukin-2 (IL-2), and 0.1 mM MEM nonessential amino-acid solution (Gibco BRL). Every 3 or 4 days, half the culture medium was removed and replaced by new medium containing the corresponding peptide (20 µg ml⁻¹) and 100 U ml⁻¹ IL-2. On the fourteenth day of the culture, the cultured cells were separated into four wells. Two wells were used for the culture with the corresponding peptide-pulsed C1R-A11, -A31, or -A33 cells, and the other two were used for the culture with HIV peptide-pulsed C1R-A11, -A31, or -A33 cells. After an 18-h incubation, the supernatant was collected, and the level of interferon (IFN)-γ was determined by enzyme-linked immunosorbent assay. The successful induction of peptide-specific CTLs was judged to be positive when a value of *P* < 0.05 was reached by a two-tailed Student's *t*-test.

Cell lines

SQ-1 is an HLA-A11⁺ lung carcinoma cell line. LC-1 is an HLA-A31⁺ and HLA-A33⁺ lung carcinoma cell line. COLO 201 and LNCaP are HLA-A3 supertype negative colon carcinoma and prostate carcinoma cell lines, respectively. All tumour cell lines were maintained in RPMI 1640 (Invitrogen) with 10% FCS. The expression of the Lck protein on these cell lines was examined by flow cytometry using anti-Lck mAb (mouse IgG2b) (sc-433; Santa Cruz Biotechnology Inc., Santa Cruz, CA, USA), followed by FITC-conjugated goat anti-mouse IgG no. 55493, Cappel ICN, Aurora,

OH, USA). Normal mouse IgG (sc-3879; Santa Cruz Biotechnology Inc.) was used as a control for anti-Lck mAb.

Cytotoxicity assay

Peptide-stimulated PBMCs were tested for their cytotoxicity against COLO 201 (HLA-A2⁺), SQ-1 (HLA-A11⁺), and LC-1 (HLA-A31⁺/A33⁺) by a standard 6-h ⁵¹Cr-release assay. Phytohaemagglutinin (PHA)-activated T cells were used as a negative control. Two thousand ⁵¹Cr-labelled cells per well were cultured with effector cells in 96-round well plates at the indicated effector/target ratio. The specific ⁵¹Cr release was calculated according to the following formula: (test c.p.m. – spontaneous c.p.m.). Spontaneous release was determined by the supernatant of the sample incubated with no effector cells, and the total release was then determined by the supernatant of the sample incubated with 1% Triton X (Wako Pure Chemical Industries, Osaka, Japan). In some experiments, CD8⁺ T cells were positively isolated using a CD8 Positive Isolation Kit (Dynal, Oslo, Norway). An measure of 10 μg ml⁻¹ of either anti-HLA class I (W6/32: mouse IgG2a) or anti-HLA-DR (L243: mouse IgG2a) mAb was added into wells at the initiation of the culture.

Cold inhibition assay

The specificity of peptide-stimulated CTLs against cancer cells was confirmed by a cold inhibition assay. In brief, ⁵¹Cr-labelled target cells (2 × 10³ cells per well) were cultured with the effector cells (2 × 10⁴ cells per well) in 96-round well plates with 2 × 10⁴ cold target cells. CIR-A11, -A31, and -A33, which were pre-pulsed with either the HIV peptide or a corresponding peptide, were used as cold target cells.

Statistics

The statistical significance of the data was determined using a two-tailed Student's *t*-test. A *P*-value of less than 0.05 was considered statistically significant.

RESULTS

Measurement of IgGs reactive to the Lck peptides in the plasma of cancer patients

First, we prepared 21 peptides derived from the Lck protein based on the binding motifs to the HLA-A3 supertype alleles (Table 1). These include 9- and 10-mer peptides. Although the A3 supertype alleles include HLA-A3 and -A68, we preferentially considered the binding capacity to HLA-A11, -A31, and -A33 molecules because the HLA-A3 and -A68 alleles are very rare in the Japanese population (Aizawa, 1986). We first screened these peptide candidates based on their recognisability by the IgGs of prostate cancer patients, since we previously observed that IgGs reactive to CTL-directed peptides are detectable in the plasma of patients with different types of cancer (Nakatsura *et al*, 2002; Ohkouchi *et al*, 2002). In addition, the number of available PBMCs from prostate cancer patients was limited, and 21 peptides was too large a number of candidates to individually test their potential to generate peptide-specific CTLs from the PBMCs of cancer patients. The results were that IgGs reactive to the Lck_{90–99}, Lck_{449–458}, Lck_{450–458}, and Lck_{452–461} peptides were detected in the plasma of 9, 11, 7, and 5 out of the 20 prostate cancer patients, respectively (Table 2). Immunoglobulin Gs reactive to the other 17 Lck peptides were less frequently observed in the plasma of cancer patients (data not shown).

Table 2 IgG reactive to Lck peptides in the plasma of prostate cancer patients

Patient	Lck _{90–99}	Lck _{449–458}	Lck _{450–458}	Lck _{452–461}	No peptide
	<i>Immunofluorescence intensity</i>				
A	49	48	36	32	47
B	99	113	98	87	64
C	66	74	56	43	26
D	33	132	63	40	46
E	92	123	96	82	47
F	15	16.5	12	10	7
G	1044	1361	1138	1096	447
H	54	69	48	55	62
I	100	86	75	134	49
J	44	52	37	34	36
K	64	65	46	40	24
L	36	43	33	29	20
M	319	274	228	213	80
N	3415	3489	3206	3217	1672
O	23	25	19	15	13
P	45	49	35	33	21
Q	52	55	44	40	37
R	42	53	37	41	49
S	81	79	59	53	53
T	146	178	115	121	253
Positive/total	9/20	11/20	7/20	5/20	

IgG = immunoglobulin G. We measured the levels of peptide-specific IgG in the plasma of 20 patients. The fluorescence intensity of the plasma (1 : 100 dilution) was measured by the Luminex method. Positive results (> no peptide × 1.5) are shown in bold.

Induction of peptide-specific CTLs from the PBMCs of prostate cancer patients with HLA-A3 supertype alleles

We next determined whether or not these four Lck peptides, which were more frequently recognised by IgGs in cancer patients than were the other 17 peptides, could induce peptide-specific CTLs from the PBMCs of HLA-A11⁺, -A31⁺, or -A33⁺ prostate cancer patients. The PBMCs were stimulated *in vitro* with each of the Lck-derived peptides or with a control peptide. The peptide-stimulated PBMCs were then assessed for their IFN-γ production in response to corresponding peptide-pulsed CIR-A11, -A31, or -A33 cells. The results of 17 patients (seven with HLA-A11, five with HLA-A31, and five with HLA-A33) are shown in Table 3. Successful induction of peptide-specific CTLs was judged to be positive when the *P*-value was < 0.05. As a result, the Lck_{90–99}, Lck_{449–458}, Lck_{450–458}, and Lck_{452–461} peptides induced corresponding peptide-specific CTLs from the PBMCs of five, two, four, and three out of seven HLA-A11⁺ cancer patients. These peptides induced peptide-specific CTLs from the PBMCs of three, three, four, and one out of five HLA-A31⁺ cancer patients, and of two, three, three, and one out of five HLA-A33⁺ cancer patients, respectively. In total, the Lck_{90–99}, Lck_{449–458}, Lck_{450–458}, and Lck_{452–461} peptides induced corresponding peptide-specific CTLs from the PBMCs of 10, 8, 11, and 5 out of 17 HLA-A3 supertype⁺ cancer patients, respectively. These rates of peptide-specific CTL induction were comparable with those of the positive control EBV and TRP-2 peptides. These results suggest that the Lck_{90–99}, Lck_{449–458}, and Lck_{450–458} peptides have the potential to efficiently generate peptide-specific CTLs in the PBMCs of prostate cancer patients with HLA-A3 supertype alleles.

Induction of cancer-reactive CTLs from the PBMCs of prostate cancer patients with HLA-A3 supertype alleles

Before the cytotoxicity assay, we selected target cancer cells that express the HLA-A3 supertype alleles and the Lck protein. Flow cytometric analysis revealed that HLA-A3 supertype negative colon

Table 3 Induction of peptide-specific CTLs from the PBMCs of HLA-A11, -A31, and -A33 cancer patients

Patient	Peptide					
	Lck ₉₀₋₉₉	Lck ₄₄₉₋₄₅₈	Lck ₄₅₀₋₄₅₈	Lck ₄₅₂₋₄₆₁	EBV TRP-2 HIV	
<i>HLA-A11</i>						
1	10	—	1	—	443	8
2	40	15	—	54	3	36
3	150	57	37	—	—	2
4	57	9	30	—	2	28
5	—	8	—	—	1	2
6	60	116	44	34	1000	—
7	87	2	232	133	583	140
	5/7	2/7	4/7	3/7	3/7	3/7
<i>HLA-A31</i>						
8	15	6	—	26	11	4
9	22	17	72	15	43	17
10	34	98	23	30	32	40
11	127	34	341	11	40	89
12	299	96	118	31	12	—
	3/5	3/5	4/5	1/5	3/5	2/5
<i>HLA-A33</i>						
13	151	55	28	13	48	19
14	14	—	24	—	207	2
15	61	60	105	15	39	157
16	9	100	16	26	—	19
17	3	6	—	—	67	20
	2/5	3/5	3/5	1/5	3/5	1/5
Positive/total	10/17	8/17	11/17	5/17	9/17	6/17

CTL = cytotoxic T lymphocyte; HLA = human leukocyte antigen; IFN- γ = interferon- γ ; PBMC = peripheral blood mononuclear cell. The PBMCs from HLA-A11, -A31, and -A33 cancer patients were stimulated *in vitro* with the indicated Lck peptides. On day 14, the cultured PBMCs were tested for their reactivity to C1R-A11, -A31, or -A33 cells, which were pre-pulsed with a corresponding peptide or the HIV peptide. The values represent the results of positive wells among four wells, and the background IFN- γ production in response to the HIV peptide was subtracted. Significant values ($P < 0.05$ by two-tailed Student's *t*-test) are shown in bold.

carcinoma COLO 201, HLA-A11⁺ lung carcinoma SQ-1, and HLA-A31⁺/A33⁺ lung carcinoma LC-1 were positive for the Lck protein (Figure 1). Although we had established several LNCAp transfectants that express each of HLA-A11, -A31, and -A33 molecules (Minami *et al*, 2007), the LNCAp cell line was negative for the Lck protein. Therefore, we used SQ-1 as an Lck-expressing HLA-A11⁺ target, and LC-1 as an Lck-expressing HLA-A31⁺/A33⁺ target. COLO 201 was used as an Lck-expressing HLA-A3 supertype negative target.

We then determined whether or not the CTLs induced by *in vitro* stimulation with each of the Lck₉₀₋₉₉, Lck₄₄₉₋₄₅₈, and Lck₄₅₀₋₄₅₈ peptides could show cytotoxicity against cancer cells (Figure 2). The PBMCs from HLA-A11⁺ patients (patients 2, 6, and 3), which were stimulated *in vitro* with each of the Lck₉₀₋₉₉, Lck₄₄₉₋₄₅₈, and Lck₄₅₀₋₄₅₈ peptides, exhibited higher levels of cytotoxicity against HLA-A11⁺ SQ-1 cells than against HLA-A11⁻ COLO 201 cells and HLA-A11⁺ PHA-stimulated T-cell blasts. Similarly, these peptides possessed the ability to induce LC-1 (HLA-A31⁺/A33⁺)-reactive CTLs from the PBMCs of HLA-A31⁺ and -A33⁺ patients (patients 10, 11, and 13). Each of the peptide-specific CTLs showed higher levels of cytotoxicity against LC-1 cells than against COLO 201 cells or T-cell blasts. Taken together, these results indicate that the PBMCs that were stimulated *in vitro* with each of the Lck₉₀₋₉₉, Lck₄₄₉₋₄₅₈, and Lck₄₅₀₋₄₅₈ peptides can exhibit cytotoxicity against cancer cells in an HLA-A11-, -A31-, or -A33-restricted manner.

Peptide-specific and CD8⁺ T-cell-dependent cytotoxicity against cancer cells

We further attempted to identify the cells responsible for the cytotoxicity of Lck peptide-stimulated PBMCs. Purified CD8⁺ T cells were used in the following experiments. As shown in Figure 3, the cytotoxicity of purified CD8⁺ T cells from the peptide-stimulated PBMCs against SQ-1 and LC-1 was significantly decreased by the addition of anti-HLA class I mAb, but not by the addition of anti-HLA class II (HLA-DR). In addition, the cytotoxicity of these cells against SQ-1 and LC-1 was significantly inhibited by the addition of corresponding peptide-pulsed unlabelled C1R-A11, C1R-A31, and C1R-A33 cells, but not by the addition of HIV peptide-pulsed unlabelled C1R-A11, C1R-A31, or C1R-A33 cells (Figure 4). These results suggested that the cytotoxicity of peptide-stimulated PBMCs against Lck-expressing cancer cells was mainly dependent on HLA class I-restricted and Lck peptide-specific CD8⁺ T cells.

DISCUSSION

Although the Lck protein is known to be essential for both T-cell development and function (Veillette *et al*, 1989), this protein is aberrantly expressed in several malignancies, including colon carcinoma, small cell lung carcinoma, and prostate carcinoma with a trend of preferential expression in metastatic lesions (Robinson *et al*, 1996; McCracken *et al*, 1997; Krystal *et al*, 1998; Lutz *et al*, 1998). We previously reported that HLA-A24-restricted tumour-reactive CTLs from patients with distant metastases can recognise Lck-derived peptides as a tumour antigen (Harashima *et al*, 2001). We also identified Lck-derived peptides that were applicable to metastatic cancer patients positive for HLA-A2 molecules (Imai *et al*, 2001). Thereafter, we carried out peptide-based immunotherapy against various types of cancer in which Lck-derived peptides were vaccinated into HLA-A24⁺ or -A2⁺ cancer patients (Sato *et al*, 2003; Mine *et al*, 2004). The primary purpose of the present study was to identify new Lck-derived peptides that would be applicable to patients with HLA-A3 supertype alleles to expand the possibilities for peptide-based anti-cancer vaccine for many ethnic populations, as described in the Introduction.

We first screened 21 Lck peptide candidates based on their ability to be recognised by the IgGs of cancer patients before testing their ability to induce HLA class I-restricted CTLs, because this approach was successful in our previous studies (Matsueda *et al*, 2005; Minami *et al*, 2007). As a result, we selected four Lck peptides that were frequently recognised by IgG. These candidates were then tested for their ability to induce peptide-specific CTLs from PBMCs of HLA-A3 supertype⁺ cancer patients, and three of them were found to effectively induce peptide-specific and cancer-reactive CTLs. Although we did not include peptides that were recognised by IgG less frequently in the CTL induction assay in this study, we previously included them and compared to those that were recognised by IgG more frequently. As a result, we observed a correlation between their recognisability by IgG and the induction rate of CTLs (Harada *et al*, 2004; Matsueda *et al*, 2005). In our clinical trials, peptide vaccination frequently resulted in the induction of IgGs reactive to the administered CTL-directed peptides, and the induction of IgGs reactive to vaccinated peptides was positively correlated with clinical responses and with the survival of vaccinated patients (Sato *et al*, 2003; Mine *et al*, 2004). Peptides that can be recognised by both CTLs and IgG might be more useful in peptide-based immunotherapy than those that are recognised only by CTLs.

Although we selected three Lck peptides from 21 candidates, the most important point was to determine whether or not the selected candidates could have the potential to induce cancer-reactive CTLs in cancer patients. In this study, we demonstrated that three Lck

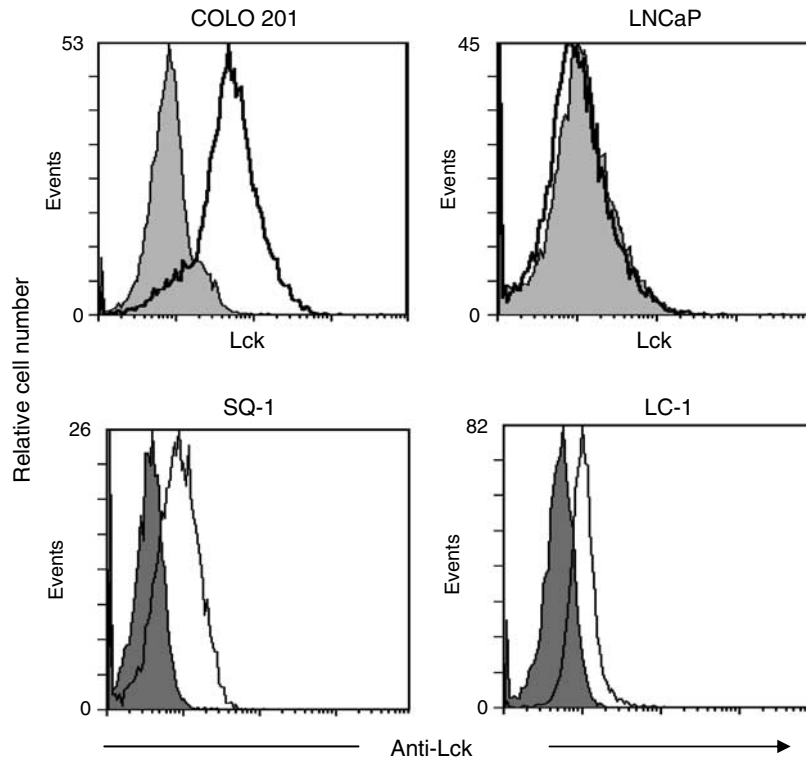


Figure 1 The expression of the Lck protein in four tumour cell lines. These tumour cell lines were analysed by flow cytometry for their expression of the Lck protein. These cells were stained first with anti-Lck mAb, followed by staining with FITC-conjugated anti-mouse IgG mAb. The grey background represents staining first with normal mouse IgG, followed by staining with FITC-conjugated anti-mouse IgG mAb.

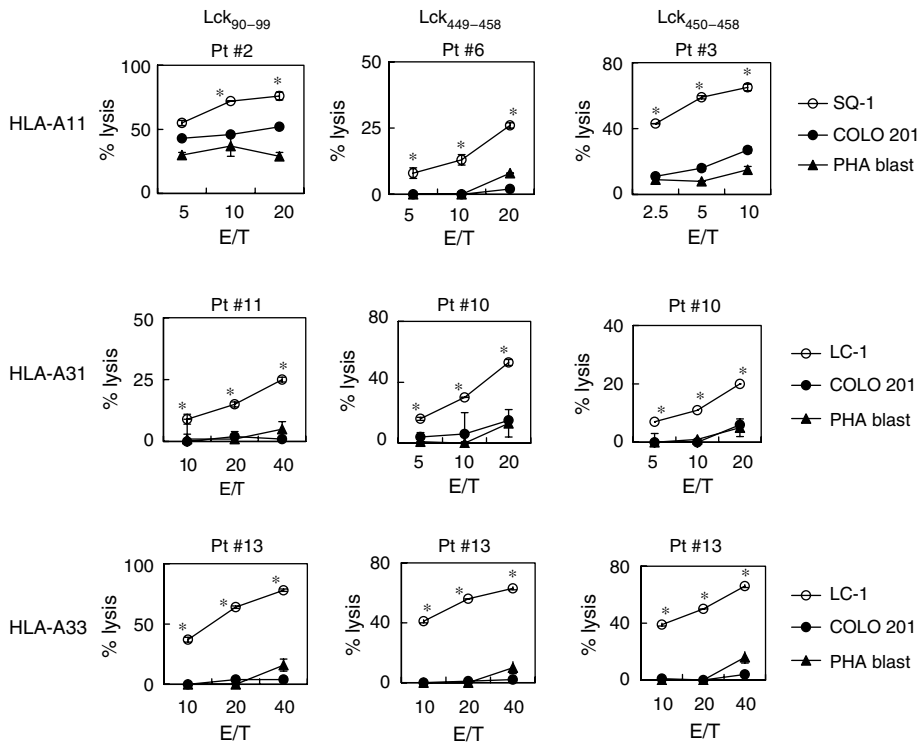


Figure 2 Cytotoxicity of peptide-stimulated PBMCs from HLA-A3 supertype⁺ prostate cancer patients. Peptide-stimulated PBMCs from HLA-A3 supertype⁺ prostate cancer patients were tested for their cytotoxicity towards three different targets by a 6-h ⁵¹Cr-release assay. Phytohaemagglutinin (PHA)-stimulated T-cell blasts from HLA-A3 supertype⁺ healthy donors were used as a control. *Statistically significant at *P* < 0.05.

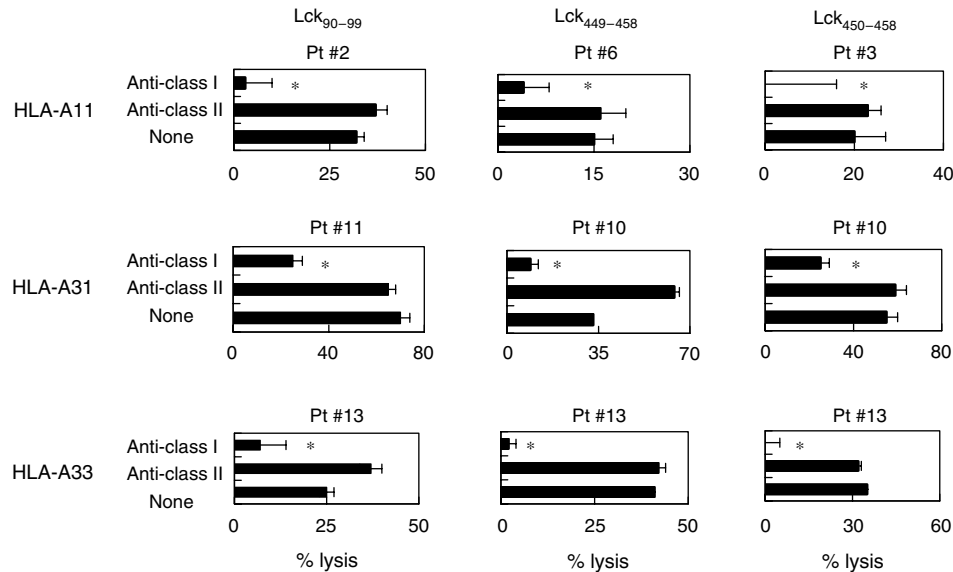


Figure 3 Human leukocyte antigen class I-restricted cytotoxicity of peptide-stimulated PBMCs against cancer cells. Purified CD8⁺ T cells from peptide-stimulated PBMCs of HLA-A3 supertype⁺ patients were tested for their cytotoxicity against HLA-A11⁺ SQ-1 cells or HLA-A31⁺/A33⁺ LC-1 cells in the presence of the indicated mAbs by a 6-h ⁵¹Cr-release assay. *Statistically significant at $P < 0.05$.

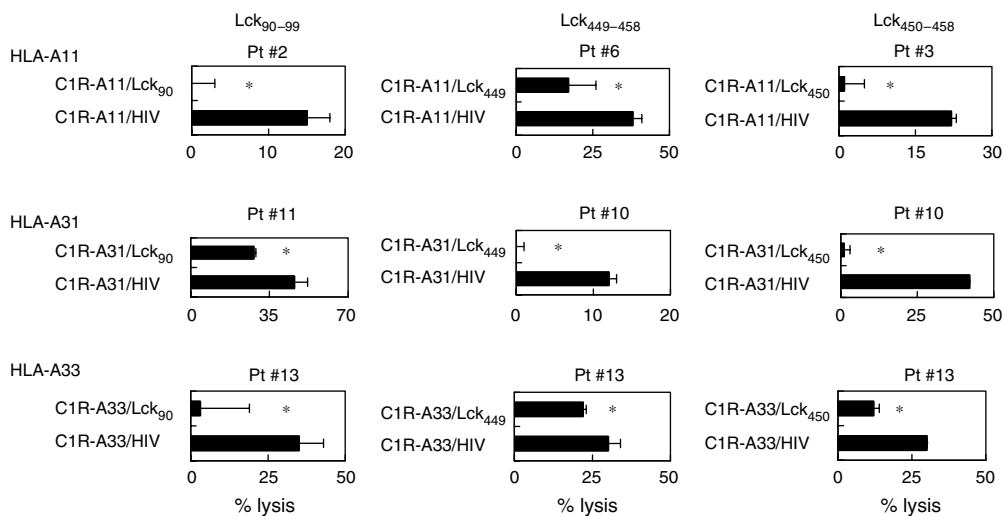


Figure 4 Peptide-specific cytotoxicity against cancer cells. Purified CD8⁺ T cells from peptide-stimulated PBMCs of HLA-A3 supertype⁺ patients were tested for their cytotoxicity against HLA-A11⁺ SQ-1 cells or HLA-A31⁺/A33⁺ LC-1 cells in the presence of unlabelled C1R-A11, -A31, or -A33 cells, which were pre-loaded with either the corresponding peptide or the HIV peptide, by a 6-h ⁵¹Cr-release assay. *Statistically significant at $P < 0.05$.

peptides, Lck₉₀₋₉₉, Lck₄₄₉₋₄₅₈, and Lck₄₅₀₋₄₅₈, could induce cancer-reactive CTLs from the PBMCs of HLA-A3 supertype⁺ cancer patients. In addition, a cold inhibition assay revealed that the cytotoxicity towards HLA-A3 supertype⁺ cancer cells was mainly dependent on Lck peptide-specific CD8⁺ T cells. These lines of evidence indicate that these Lck peptides would be useful as a peptide-based anti-cancer vaccine for HLA-A3 supertype⁺ cancer patients.

The optimal COOH-terminal amino acid of A11-binding peptides is lysine, and that of A31- or A33-binding peptides is arginine (Rammensee *et al*, 1995; Sidney *et al*, 1996). Among the three Lck peptides identified in this study to be candidates for HLA-A3 supertype alleles, the Lck₉₀₋₉₉ peptide carries lysine at the COOH terminus, and both the Lck₄₄₉₋₄₅₈ and Lck₄₅₀₋₄₅₈ peptides carry arginine at the COOH terminus. However, our results

revealed that the Lck₉₀₋₉₉ peptide induced peptide-specific CTLs from the PBMCs of HLA-A31⁺ or -A33⁺ cancer patients and, on the other hand, both the Lck₄₄₉₋₄₅₈ and Lck₄₅₀₋₄₅₈ peptides induced peptide-specific CTLs from the PBMCs of HLA-A11⁺ cancer patients. We reported a similar observation in our previous study (Matsueda *et al*, 2005). These observations might suggest that peptides carrying lysine or arginine at the COOH terminus fit the binding motif for all HLA-A11, -A31, and -A33 molecules.

In conclusion, we identified three new peptide candidates that were applicable to HLA-A3 supertype⁺ cancer patients. In combination with known Lck peptides for HLA-A2⁺ or -A24⁺ cancer patients (Harashima *et al*, 2001; Imai *et al*, 2001), those identified in the present study enable us to develop a peptide-based anti-cancer vaccine for cancer patients with metastases in diverse ethnic populations.

ACKNOWLEDGEMENTS

This study was supported in part by Grants-in-Aid from the Ministry of Education, Science, Sports and Culture of Japan (no. 12213134 to KI, and no. 18591449 to MH); from the Research

Center of Innovative Cancer Therapy of 21st Century COE Program for Medical Science to KI; and from the Ministry of Health, Labor and Welfare, Japan (no. H14-trans-002, 11–16, and H12-cancer-004 to KI).

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