

Clinical Study

Voluntary driven exoskeleton as a new tool for rehabilitation in chronic spinal cord injury: a pilot study

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Abstract

BACKGROUND CONTEXT: Treadmill training after traumatic spinal cord injury (SCI) has become an established therapy to improve walking capabilities. The hybrid assistive limb (HAL) exoskeleton has been developed to support motor function and is tailored to the patients' voluntary drive.

PURPOSE: To determine whether locomotor training with the exoskeleton HAL is safe and can increase functional mobility in chronic paraplegic patients after SCI.

DESIGN: A single case experimental A-B (pre-post) design study by repeated assessments of the same patients. The subjects performed 90 days (five times per week) of HAL exoskeleton body weight supported treadmill training with variable gait speed and body weight support.

PATIENT SAMPLE: Eight patients with chronic SCI classified by the American Spinal Injury Association (ASIA) Impairment Scale (AIS) consisting of ASIA A (zones of partial preservation [ZPP] L3–S1), n=4; ASIA B (with motor ZPP L3–S1), n=1; and ASIA C/D, n=3, who received full rehabilitation in the acute and subacute phases of SCI.

OUTCOME MEASURES: Functional measures included treadmill-associated walking distance, speed, and time, with additional analysis of functional improvements using the 10-m walk test (10MWT), timed-up and go test (TUG test), 6-minute walk test (6MWT), and the walking index for SCI II (WISCI II) score. Secondary physiologic measures including the AIS with the lower extremity motor score (LEMS), the spinal spasticity (Ashworth scale), and the lower extremity circumferences.

FDA device/drug status: Not applicable.

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MS-K: Nothing to disclose. **OH:** Nothing to disclose. **RCM:** Nothing to disclose. **MT:** Nothing to disclose. **PS:** Nothing to disclose. **YS:** Royalties: University of Tsukuba (E); Stock Ownership (E, Paid directly to institution); Private Investments: (E, Paid directly to institution); Consulting: (E, Paid directly to institution); Speaking / Teaching Arrangements: (E, Paid directly to institution); Trips/Travel: (E, Paid directly to institution); Board of Directors: CYBERDYNE, Inc. (E); Grants: Cabinet Office (I). **TAS:** Nothing to disclose.

The disclosure key can be found on the Table of Contents and at www.TheSpineJournalOnline.com.

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Authors' contributions: MA and OC carried out the experiments and data analysis as well as drafting of the manuscript. RCM, PS and MT helped with the experimental set up. MS-K and OH contributed to the data analysis. TAS participated in study design and coordination of the study.

All authors read and approved the final manuscript.

YS is a founder, shareholder, and the CEO of Cyberdyne, Inc., which produces the HAL.

YS and Cyberdyne were neither involved in study funding, design, data collection, and analysis, nor in writing or submitting this article, therefore concluding in no specific influence on the trial. We certify that no party having a direct interest in the results of the research supporting this article has or will confer a benefit on us or on any organization with which we are associated.

YS and Cyberdyne as the manufacturer of the device provided exclusively technical and advisory support.

YS as the CEO of Cyberdyne has been involved exclusively in terms of an advisory capacity, regarding technical support and the limitations of the exoskeleton. Therefore, the inclusion and exclusion criteria have been modified (eg, body weight and contractures).

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METHODS: Subjects performed standardized functional testing before and after the 90 days of intervention.

RESULTS: Highly significant improvements of HAL-associated walking time, distance, and speed were noticed. Furthermore, significant improvements have been especially shown in the functional abilities without the exoskeleton for over-ground walking obtained in the 6MWT, TUG test, and the 10MWT, including an increase in the WISCI II score of three patients. Muscle strength (LEMS) increased in all patients accompanied by a gain of the lower limb circumferences. A conversion in the AIS was ascertained in one patient (ASIA B to ASIA C). One patient reported a decrease of spinal spasticity.

CONCLUSIONS: Hybrid assistive limb exoskeleton training results in improved over-ground walking and leads to the assumption of a beneficial effect on ambulatory mobility. However, evaluation in larger clinical trials is required. © 2014 Elsevier Inc. All rights reserved.

Keywords:

Exoskeleton; Treadmill training; Rehabilitation; Paraplegia; Hybrid assistive limb; Spinal cord injury

Introduction

About 1,200 people suffer a traumatic spinal cord injury (SCI) each year in Germany. Recent statistics indicate that more than 50% of these injured patients have a motor incomplete lesion [1]. In patients with initial motor incomplete SCI, at least 75% regain some kind of ambulatory function. Better functional outcome is associated with age, level of lesion, and the classification in the American Spinal Injury Association (ASIA) Impairment Scale [2]. In the first 2 months after initial SCI, approximately half of the recovery occurs. Within the following 4 months, a decreasing rate of recovery has been observed. One year after injury, neurologic recovery is assumed to be nearly complete [3]. Although conventional rehabilitation programs enhance the performance of functional tasks, the loss of strength and coordination substantially limit one's capacity for over-ground ambulation training [4]. In the past two decades, body weight supported treadmill training (BWSTT) has been proposed as a useful adjunct to enhance locomotor function after motor incomplete SCI [5]. In patients with incomplete or complete SCI, a bilateral leg muscle activation combined with coordinated stepping movements can be induced in partially unloaded patients, standing on a moving treadmill. Body weight supported treadmill training enables early initiation of gait training and integration of weight-bearing activities, stepping and balance, by the use of a task-specific approach and a systematic gait pattern [6]. To facilitate the delivery of BWSTT in SCI patients, the locomotor training evolved over the last 12 years and a motorized robotic driven gait orthosis (DGO) has been developed [7]. The advantages over conventional BWSTT methods are considered to be less effort for attending physiotherapists [8], longer duration, more physiologic and reproducible gait patterns, and the possibility to measure a patients' performance. Several studies pointed out that DGO training improves over-ground walking [9–13]. However, there was no reported difference in the outcome of DGO training compared with conventional training. A significant switch in the ASIA classification has not been found [10,14].

Over the last 5 years, exoskeletal systems became available for SCI patients. These systems offer different possibilities. Three exoskeletons (Ekso [EksoBionics, Richmond, CA, USA], Rex [Rex Bionics, Auckland, New Zealand] and Re-Walk [ARGO Medical Technologies, Israel]) allow SCI patients to stand up, walk with a defined pattern, and even climbing stairs mainly on a basis of passive range of motion (ROM). The exoskeleton hybrid assistive limb (HAL; Cyberdyne, Inc., Japan) offers the possibility of getting connected with the SCI patient through electromyography electrodes on the skin at the extensor/flexor muscle region of the lower extremities. This allows voluntary machine supported ROM of incomplete SCI patients by using minimal bioelectrical signals, recorded and amplified from hip and knee flexors and extensors [15–17]. More recently, these various exoskeletal systems allow the patients mobilization outside the treadmill. A former study by Kawamoto et al. [18] concerning locomotion improvement using HAL in chronic stroke patients, emphasized the feasibility for rehabilitation of these particular patients.

The aim of this pilot study was to evaluate the possibilities of exoskeletal locomotor training (HAL; Cyberdyne, Inc.) under voluntary control and identify beneficial effects on functional mobility of the patients. The hypothesis was that exoskeleton treadmill training is feasible and safe in application and capable of improving ambulatory mobility in chronic SCI patients.

Materials and methods

Patients

We enrolled eight patients (two women, six men). The mean \pm standard deviation age at the time of enrollment was 48 ± 9.43 years. All patients were in the chronic stage of traumatic SCI according to the time since injury of 1 to 19 years (97.2 ± 88.4 months). Inclusion criteria were traumatic SCI with chronic incomplete (ASIA B/C/D) or complete paraplegia (ASIA A) after lesions of the conus medullaris/cauda equine with zones of partial preservation. Independent of ASIA classification, the enrolled patients

must present motor functions of hip and knee extensor and flexor muscle groups to be able to trigger the exoskeleton. Exclusion criteria were as follows: nontraumatic SCI, pressure sores, severe limitation of ROM regarding hip and knee joints, cognitive impairment, body weight more than 100 Kg, nonconsolidated fractures, and mild or severe heart insufficiency. Two patients suffered from an incomplete thoracic SCI (ASIA C/D) from 3 to 13 years. Two patients suffered from an incomplete lumbar SCI (ASIA B/C) from 12 to 13 months and four patients had a complete SCI with zones of partial preservation in L3–S1 after lesions of the conus medullaris. The classification according to the ASIA was carried out before the treadmill training was initiated. The study was approved by Ethical Board Committee of Bergmannsheil Hospital and the University of Bochum and followed strictly the declaration of Helsinki.

All patients provided written informed consent. The study design was a single case experimental A-B (pre-post) design by repeated assessments of the same patients (Table 1).

Intervention

During this study, the patients underwent a BWSTT five times per week using the HAL exoskeleton (Cyberdyne, Inc., Japan). The study was performed between June 2013 and September 2013 in the BG University Hospital Bergmannsheil, Bochum.

Neither adverse nor severe adverse events occurred during the intervention.

The exoskeleton

The HAL robot suit (Cyberdyne, Inc., Japan) is an exoskeleton with a frame and robotic actuators that attach to the patients' legs. The joint movement is supported by electric motors. Voluntary initiated minimal bioelectrical signals recovered from extensor and flexor muscles of hip and knee are detected via electromyography electrodes (Fig. 1).

Through a cable connection between the exoskeleton and patient, this system allows voluntary robotic supported ROM (cybernic voluntary control mode). Also a passive, nonvoluntary ROM (cybernic autonomous control mode) is possible (Fig. 2).

The treadmill

The treadmill system (Woodway USA, Inc., Waukesha, WI, USA) includes a body weight support system with a harness. The speed can be adjusted from 0 Km/h to approximately 4.5 Km/h. During treatments, the velocity of the treadmill was set individually between comfortable and maximum speed tolerated by the patients. Approximately 50% of each patient's body weight needed to be supported by the harness system, individually reduced during the following sessions as tolerated without substantial knee buckling or toe drag.

EVIDENCE & METHODS

Context

The authors present a series of patients treated with an assistive exoskeleton developed to facilitate treadmill exercise in patients with spinal cord injury (SCI).

Contribution

In a series of eight patients with SCI graded ASIA A to C/D, improvements in walking time, distance and speed were noted after treatment with assistive exoskeleton.

Implications

This study is a case series of eight patients with heterogeneous clinical characteristics, including the severity of their spinal cord injury. The findings are limited to clinical contexts specific to these patients and clearly cannot be translated to the care of other individuals. This is simply a report that may show proof of concept. It should be noted that one of the authors reports a substantive conflict of interest (founder and shareholder of the company that produces the exoskeleton device).

—The Editors

The training

The patients underwent a 90-day period of HAL exoskeleton (Cyberdyne, Inc.) training (five per week), including a mean number of sessions of 51.75 ± 5.6 . The training was performed on a treadmill with individually adjustable body weight support and speed, recording walking speed, time, and distance. It included a 10-m walk test (10MWT) before and after each session and regular physiotherapy that lasted approximately 90 minutes. The training was supervised by a physiotherapist and a medical doctor.

Measurements

Walking capabilities and neurologic status

All patients were assessed on admission by medical doctors involved in this trial. The outcomes were assessed

Table 1
Subject demographics and clinical characteristics

| Case | Sex | Time | | Etiology | Level | ASIA/ZPP | WISCI | |
|------|-----|---------|-----------|-----------|-------|----------|-------|----------|
| | | Age (y) | trauma, y | | | | II | Ashworth |
| 1 | M | 40 | 13 | # T7/T8 | T8 | C | 13 | 4 |
| 2 | M | 63 | 1 | # T12 | L1 | B/L3 | 6 | 0 |
| 3 | M | 36 | 1.16 | # T11/T12 | T12 | A/L3 | 6 | 0 |
| 4 | F | 55 | 1.08 | # L1 | L1 | C | 13 | 0 |
| 5 | M | 42 | 16 | # L1 | L1 | A/L3 | 9 | 0 |
| 6 | M | 52 | 10 | # L3 | L2 | A/L3 | 6 | 0 |
| 7 | F | 40 | 19 | # L1 | T11 | A/S1 | 9 | 0 |
| 8 | M | 53 | 3 | # T12 | T12 | D | 18 | 0 |

M, male; F, female; #, fracture; ASIA, American Spinal Injury Association; ZPP, zones of partial preservation; WISCI, walking index for spinal cord injury; T, thoracic; L, lumbar; S, sacral.

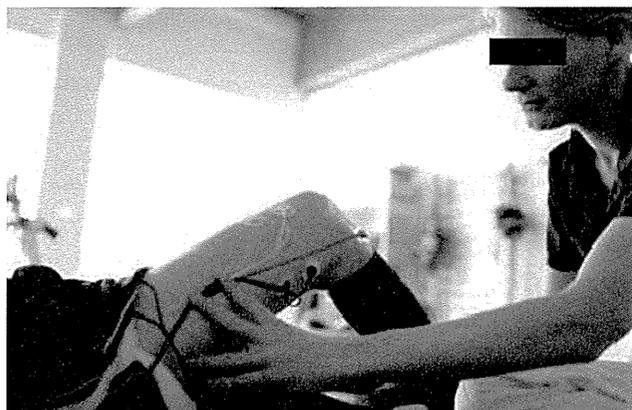


Fig. 1. Positioning of the electromyography electrodes on the knee extensor and flexor muscles.

by physiotherapists neither involved in the study design nor analysis after 45 days and on discharge from the training period. An assessment through the ASIA classification



Fig. 2. Patient performing treadmill locomotion training with body weight support and hybrid assistive limb exoskeleton.

was already done on admission and on discharge from the SCI department, Bergmannsheil, Bochum, within the initial therapy after acute SCI. The 10MWT, done before and after each session, detected the needed time, the number of steps, and the required assistance to walk a 10 m distance [19,20]. The timed-up and go test (TUG test) describes the time and assistance required for standing up from the wheelchair, walk 3 m, turn around, walk back, and sit down. It was performed every 2 weeks. The 6-minute walk test (6MWT) was done at the beginning, at half time, and at the end if possible, depending on the patient. It evaluates the distance and assistance while walking for 6 minutes [21]. The main outcome was the functional motor assessment by the walking index for SCI II (WISCI II) [22,23]. The WISCI II score is a 20-item scale, measuring the walking capabilities of a patient based on the requirements of assistance because of walking aids, personal assistance, or braces. Grade 0 means that the patient has neither standing nor walking abilities. Grade 20 means that no assistance is needed to walk a distance of 10 m. The neurologic status was assessed using the ASIA Impairment Scale modified from the Frankel classification and classifies motor and sensory impairments that result from a SCI [3]. The lower extremity motor score (LEMS) acquired in this study was obtained by the addition of the impairment scores (0–5) of the lower extremity key muscles of both sides. Muscle volume was assessed by manual measurements, 20/10 cm above and 15 cm below inner knee gap.

Statistical analysis

Descriptive analysis of the demographic and injury characteristics was done using frequency distribution for categorical data and mean for continuous variables. Differences between pre- and posttraining sessions were assessed by a paired *t* test (for continuous variables). Treatment effects on functional performance as the WISCI II are all ordinal scales. Medians were used as descriptive statistics for these outcomes, and nonparametric tests were used to assess the relative effect of the treatments.

Results were considered statistically significant when the *p* value was $\leq .05$.

Results

Treadmill associated results

All patients improved in treadmill training by using HAL (Cyberdyne, Inc., Japan). The mean walking speed increased from 0.91 ± 0.41 m/s (0.5–1.8 m/s) in the first session up to 1.59 ± 0.5 m/s (0.8–2.1 m/s) in the last session after 3 months. The progress in speed after 6 weeks of training was lower than in the first weeks. The range was located between 0 km/h and 0.8 km/h. The mean walking time at the beginning was 12.37 ± 4.55 minutes. The average walking time at the

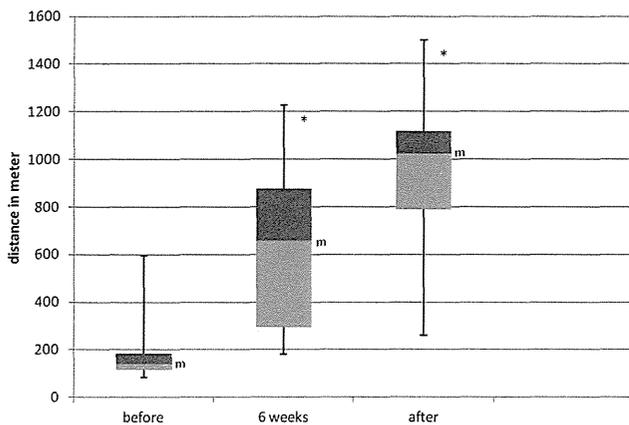


Fig. 3. Changes in treadmill-associated walking distance in pre-, mid-, and postevaluations. m, median. *pre-post difference, $p < .05$.

end was 31.97 ± 9.45 minutes. The mean ambulated distance at the first session was 195.9 ± 166.7 m and increased to 954.13 ± 380.4 m on discharge (Fig. 3).

Functional outcome

Although the mean improvement concerning the WISCI II score was not statistically significant, three patients showed functional improvement in gait abilities. Two subjects needed braces, a walker, and support by a physiotherapist at the beginning and were able to walk after the training series only with a walker and braces (WISCI II score increased from 6 to 9). One patient increased from 9 to 12 and, therefore, was able to walk with two crutches and a brace compared with a walker and a brace before the training. At baseline, the mean WISCI II score was 10 ± 4.3 . At the end of the 90 days trial, the mean WISCI II was 11.13 ± 6.68 . Improvements in speed and endurance in over-ground gait assessments in all participants have been achieved. The 10MWT showed a significant increase in mean gait speed at the end of the training period compared with baseline (0.28 ± 0.28 m/s vs. 0.50 ± 0.34 m/s) (Fig. 4).

The improvement corresponded to a 44% faster walking than in initial evaluation. It also includes the reduction of

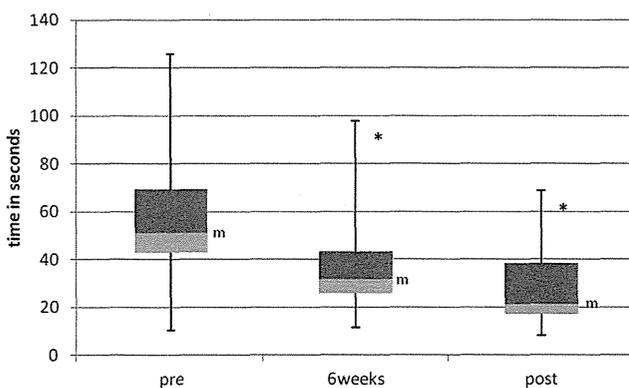


Fig. 4. Changes in 10-m walk test in pre-, mid-, and postintervention evaluations. m, median. *pre-post difference, $p < .05$.

Table 2
Comparison of pre- and postinterventions

| Outcome measurements | Before training | After training | n |
|----------------------|---------------------|-----------------------|---|
| 10MWT speed (m/s) | 0.28 ± 0.28 | $0.5 \pm 0.34^*$ | 8 |
| Number of steps | 29.88 ± 7.85 | $19.38 \pm 3.16^*$ | 8 |
| 6MWT distance (m) | 70.1 ± 130 | $163.3 \pm 160.6^*$ | 8 |
| TUG test (s) | 55.34 ± 32.20 | $38.18 \pm 25.98^*$ | 8 |
| Distance (m) | 195.88 ± 166.71 | $954.13 \pm 380.35^*$ | 8 |
| WISCI-II | 10 ± 4.34 | 11.12 ± 3.68 | 8 |

10MWT, 10-m walk test; 6MWT, 6-minute walk test; TUG, timed-up and go; WISCI, walking index for spinal cord injury.

Note: Values are means \pm standard deviation.

* Pre-post difference, $p < .05$.

support needed detected by the WISCI II score. The mean number of steps decreased from 29.8 ± 7.85 to 19.4 ± 3.16 . We observed significant increase in gait speed from pre- to midtraining and from mid- to posttraining assessments. Similar results were detected for the TUG test. The mean time needed for the TUG test decreased from 55.34 ± 32.2 seconds to 38.18 ± 25.98 seconds. The 6MWT was done with a constant walking time of 6 minutes without any break. Only three patients were able to perform the 6MWT before the training with a mean walking distance of 187 ± 162.2 m. The subjects in this subgroup improved their performance and increased the walking distance to 287.3 ± 229.4 m. After completing the training, all eight patients could be evaluated, therefore the overall mean distance increased from 70.1 ± 130 m to 163.3 ± 160.6 m (Table 2).

The LEMS increased in all patients. The mean LEMS before the training increased significantly from 21.75 ± 8.3 to 24.38 ± 7.6 after the intervention (Fig. 5).

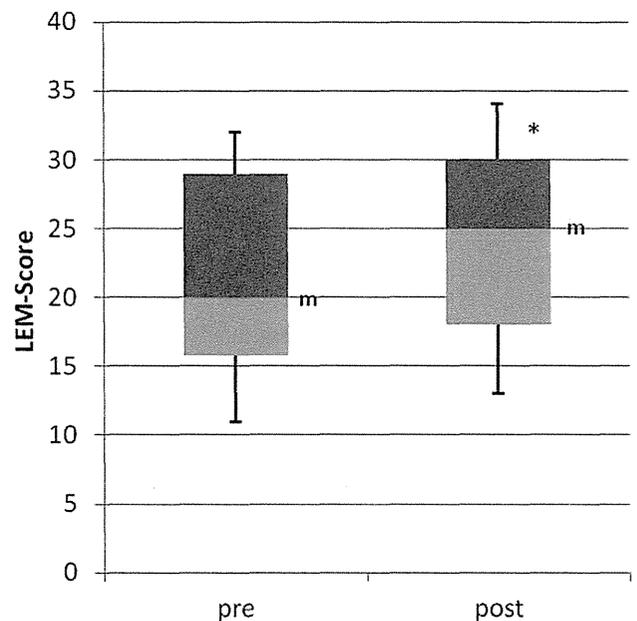


Fig. 5. LEMS in pre- and postevaluation. LEM, lower extremity motor score; m, median. *pre-post difference, $p < .05$.

One patient switched in the ASIA scale from ASIA B to C, he was at the beginning of the training 12 months posttrauma.

Others

To describe muscle volume, measurements of the circumferences 10/20 cm cranial of the inner knee joint gap and 15 cm distal of it have been done before and after the 90 days of training. Seven participants showed a gain of muscle circumference from 5 mm up to 50 mm. In one participant with edema in his lower legs, we observed a loss of circumference up to 25 mm. One patient suffering from a thoracic SCI presented a significant spinal spasticity. For spastic motor behaviors, we used the modified Ashworth scale to evaluate the involuntary resistance to passive stretch of the quadriceps muscle group. At pretraining evaluation, he showed an extensor spasm with high resistance to passive stretch according to Ashworth 4. After the training sessions, the resistance was reduced according to Ashworth 2. This level lasted for about 6 to 8 hours with a new maximum level at the next morning. All other patients showed no spastic motor behaviors.

Discussion

The objective of the study was to determine whether locomotor training with the exoskeleton HAL is feasible and safe in application, improves functional mobility, and increases motor functions in chronic paraplegic patients after SCI. The results obtained revealed a highly significant improvement for over-ground walking abilities evaluated by the 10MWT, the 6MWT, and the TUG test and the partial reduction of physical assistance and walking aids in the WISCI II score. Muscle strength, measured with the LEMS increased in all patients.

The results acquired in this clinical trial imply that HAL-supported locomotion training can improve walking abilities in terms of speed, gait, and distance. Furthermore, it improves motor functions.

Thus far there is insufficient evidence and only a few articles addressing the main hypothesis of this study that locomotor training improves walking function for patients with SCI [24].

The present study is according to the knowledge of the authors the first to investigate the impact of HAL-supported locomotor training in chronic SCI patients, where referring to the current state of knowledge no further functional improvements are to be expected.

In the subject population consisting of eight patients including patients suffering from SCI from 1 to 19 years (8.03 ± 7.4 years), all patients improved significantly regarding treadmill-associated walking distance and speed and functional improvement was detected in the over-ground walking tests.

Although no significant influence was seen on the requirements of assistance in the 10MWT, three patients attained improvement in walking abilities according to the

WISCI II, under condition of a comfortable and stable gait. A further reduction of assistance was not forced because of more pathologic gait or higher risk of falling [24].

Although the evidence is still insufficient, the effectiveness of automated locomotor training using the DGO in patients with chronic SCI is being investigated and considered promising in several systematic reviews including a Cochrane review [25,26]. The results mentioned previously add to the wealth of that data presuming that HAL-assisted locomotion training is useful in terms of functional mobility and a safe adjunct to the treatment of patient with chronic SCI.

Our study had several limitations: the relatively small number of patients ($n=8$) and the mixture of complete and incomplete SCIs.

However, all the patients were treated in the same facility by the same multidisciplinary team, according to a standardized protocol.

In summary, our study provides the first data demonstrating the clinical potential of HAL-locomotor training based on voluntary drive in patients suffering from chronic SCI.

It was proven to be a safe device for locomotion therapy as neither adverse nor severe adverse events occurred.

However, continued research in the form of large randomized trials to compare the efficacy of HAL-assisted training with well established, conventional therapies is necessary.

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Neuroimmunological aspects of human T cell leukemia virus type 1-associated myelopathy/tropical spastic paraparesis

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Abstract Human T cell leukemia virus type 1 (HTLV-1) is a human retrovirus etiologically associated with adult T cell leukemia/lymphoma and HTLV-1-associated myelopathy/tropical spastic paraparesis (HAM/TSP). Only approximately 0.25–4 % of infected individuals develop HAM/TSP; the majority of infected individuals remain lifelong asymptomatic carriers. Recent data suggest that immunological aspects of host–virus interactions might play an important role in the development and pathogenesis of HAM/TSP. This review outlines and discusses the current understanding, ongoing developments, and future perspectives of HAM/TSP research.

Keywords HTLV-1 · HAM/TSP · Host immune response

Introduction

Human T cell leukemia virus type 1 (HTLV-1) is a replication-competent human retrovirus associated with two distinct types of disease: a malignancy of mature CD4⁺ T cells called adult T cell leukemia/lymphoma (ATL) (Hinuma et al. 1981; Poiesz et al. 1980; Yoshida et al. 1984) and a chronic inflammatory central nervous system disease HTLV-1-associated myelopathy/tropical spastic paraparesis (HAM/TSP) (Gessain et al. 1985; Osame et al. 1986). Like human immunodeficiency virus (HIV), HTLV-1 is never eliminated from the host despite vigorous cellular and humoral immune responses. However, in contrast to HIV infection, few with HTLV-1 develop disease; only approximately 2–3 % of infected persons develop ATL (Tajima 1990), another 0.25–4 % develop HAM/TSP (Hisada et al. 2004; Kramer et al. 1995; Nakagawa et al. 1995; Osame et al. 1990), and the majority of infected individuals remain lifelong

asymptomatic carriers (ACs). Therefore, evaluation of the individual risk of developing disease in ACs would certainly be of considerable importance, especially in HTLV-1 endemic areas.

The viral, host, and environmental risk factors as well as the host immune response against HTLV-1 infection appear to regulate the development of HTLV-1-associated diseases (Bangham and Osame 2005). In particular, a strong immune response, especially the cytotoxic T lymphocyte (CTL) response, to HTLV-1 is seen in patients with HAM/TSP and suggested to be strongly associated with the pathogenesis of HTLV-1-associated diseases (Matsuura et al. 2010; Saito et al. 2012). For more than two decades, the investigation of HTLV-1-mediated immunopathogenesis has focused on Tax, an HTLV-1-encoded viral oncoprotein, because Tax activates many cellular genes by binding to groups of transcription factors and co-activators and is necessary and sufficient for cellular transformation. However, recent reports have identified that another regulatory protein, HTLV-1 basic leucine zipper factor (HBZ), also has a critical role in the development of ATL and HAM/TSP (Matsuoka and Jeang 2011). This review summarizes past and recent studies of HAM/TSP, attempting to answer the following fundamental questions: Why do some HTLV-1-infected people develop disease whereas the vast majority remain healthy? How does HTLV-1 persist in the individual host despite a strong host immune response? How is the inflammatory lesion in HAM/TSP initiated and maintained?

History and epidemiology of HTLV-1

HTLV-1 belongs to the *Deltaretrovirus* genus of the *Orthoretrovirinae* subfamily and infects 10–20 million people worldwide (de The and Bomford 1993; Proietti et al. 2005; Uchiyama 1997). HTLV-1 can be transmitted through sexual contact (Roucoux et al. 2005), intravenous drug use (Proietti et al. 2005), and breastfeeding from mother to child (Hino

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et al. 1985; Kinoshita et al. 1987). At present, the infection is endemic in southwest Japan, the Caribbean, Sub-Saharan Africa, and South America, with smaller foci in Southeast Asia, South Africa, and northeast Iran (Verdonck et al. 2007). HTLV-1 was initially isolated in 1980 from two T cell lymphoblastoid cell lines and the blood of a patient originally thought to have a cutaneous T cell lymphoma (Poesz et al. 1980). It was the first retrovirus ever associated with cancer in a human. Three years before the isolation of HTLV-1, Takatsuki et al. reported ATL, a rare form of leukemia endemic to southwest Japan, as a distinct clinical entity (Uchiyama et al. 1977). In 1981, Hinuma et al. clearly demonstrated that ATL was caused by a new human retrovirus, originally termed ATL (Hinuma et al. 1981; Miyoshi et al. 1981). Since then, ATL and HTLV have been shown to be identical, and a single name, HTLV-1, has been adopted. In the mid-1980s, epidemiological data linked HTLV-1 infection to a chronic progressive neurological disease, which was termed tropical spastic paraparesis in the Caribbean (Gessain et al. 1985) and HTLV-1-associated myelopathy in Japan (Osame et al. 1986). HTLV-1-positive TSP and HAM were subsequently found to be clinically and pathologically identical, and the disease was given a single designation as HAM/TSP (Hollberg and Hafler 1993). To date, more than 3,000 cases of HAM/TSP have been reported in HTLV-1 endemic areas. Sporadic cases have also been described in non-endemic areas such as the USA and Europe, mainly in immigrants from an HTLV-1 endemic area (Araujo and Silva 2006). HTLV-1 can cause other chronic inflammatory diseases such as uveitis (Mochizuki et al. 1992), arthropathy (Nishioka et al. 1989), pulmonary lymphocytic alveolitis (Maruyama et al. 1988; Sugimoto et al. 1989; Sugimoto et al. 1987), polymyositis (Higuchi et al. 1992; Morgan et al. 1989), Sjögren syndrome (Terada et al. 1994), and infective dermatitis (LaGrenade et al. 1990), although there is no clear evidence for an etiological role of HTLV-1 in these diseases.

Clinical and pathological features of HAM/TSP

HAM/TSP is a chronic progressive myelopathy characterized by spastic paraparesis, sphincter dysfunction, and mild sensory disturbance in the lower extremities (Nakagawa et al. 1996). The period from initial HTLV-1 infection to the onset of HAM/TSP is assumed to range from months to decades, a shorter time than that for the onset of ATL (Nakagawa et al. 1995; Olindo et al. 2006). HAM/TSP occurs both in vertically infected individuals and in those who become infected later in life, that is, through sexual contact almost exclusively from male to female, intravenous drug use, contaminated blood transfusions, etc. The mean age at onset is 43.8 years, and like autoimmune diseases, the frequency of cases of HAM/TSP is greater in women than in men (the male-to-female

ratio of occurrence is 1:2.3) (Nakagawa et al. 1995). In addition to HTLV-1 antibody (Ab) positivity both in serum and cerebrospinal fluid (CSF), the presence of atypical lymphocytes (the so-called “flower cells”) in peripheral blood and CSF, a moderate pleocytosis, and raised protein content in CSF is observed in patients with HAM/TSP (Araujo and Silva 2006). Oligoclonal immunoglobulin bands in the CSF; raised concentrations of inflammatory markers such as neopterin, tumor necrosis factor (TNF)- α , interleukin (IL)-6, and interferon (IFN)- γ ; and increased intrathecal Ab synthesis specific for HTLV-1 antigens have also been described (Jacobson 2002). Clinical progression of HAM/TSP is associated with an increase in the proviral load (PVL) in individual patients, and a high ratio of PVL in CSF cells/peripheral blood mononuclear cells (PBMCs) is also significantly associated with clinically progressive disease (Takenouchi et al. 2003). Thus, a pro-inflammatory environment associated with increased numbers of HTLV-1-infected cells is a characteristic immunological profile of HAM/TSP.

Pathological analysis of HAM/TSP autopsy materials showed the loss of myelin and axons in the lateral, anterior, and posterior columns of the spinal cord. These lesions are associated with perivascular and parenchymal lymphocytic infiltration with the presence of foamy macrophages, reactive astrocytosis, and fibrillary gliosis, predominantly at the thoracic level (Iwasaki 1990; Izumo et al. 2000; Yoshioka et al. 1993), suggesting that the immune response against HTLV-1 causes the inflammatory spinal cord damage seen in patients with HAM/TSP (Bangham 2000). In patients with active chronic lesions in the spinal cord, perivascular inflammatory infiltration with similar composition of cell subsets was also seen in the brain (Aye et al. 2000). The peripheral nerve pathology of patients with HAM/TSP with sensory disturbance showed varying degrees of demyelination, remyelination, axonal degeneration, regeneration, and perineural fibrosis (Bhigjee et al. 1993; Kiwaki et al. 2003).

Treatment of HAM/TSP

To date, no generally agreed standard treatment regimen has been established for HAM/TSP, and no treatment has proven to be consistently effective on a long-term basis. Therefore, clinical practice for treatment of patients with HAM/TSP is based on case series and open, nonrandomized, uncontrolled studies. Mild to moderate beneficial effects have been reported for a number of agents in open-label studies including corticosteroids (Nakagawa et al. 1996), danazol (Harrington et al. 1991), pentoxifylline (Shirabe et al. 1997), immunosuppressants such as ciclosporin A (Martin et al. 2012), high-dose intravenous gamma globulin (Kuroda et al. 1991), plasmapheresis (Matsuo et al. 1988), antibiotics (erythromycin and fosfomycin), and vitamin C (Nakagawa et al. 1996). It should be noteworthy that oral prednisolone was effective in 81.7 %

of 131 patients in a large-scale case series study (Nakagawa et al. 1996). However, the complications of corticosteroids limit their use, particularly in post-menopausal women, who are at higher risk for developing HAM/TSP. Multicenter double-blind randomized placebo-controlled trials for the IFN- α treatment indicate that IFN- α is an effective therapy with an acceptable side effects profile (Izumo et al. 1996), although the benefit of long-term IFN- α therapy has not been well studied. In regard to oral antiviral drugs zidovudine plus lamivudine, no evidence of significant benefit yet exists from randomized placebo-controlled trials (Taylor et al. 2006). Recently, oral administration of histone deacetylase inhibitor valproic acid (VPA) has been conducted as a single-center, open-label trial (Olindo et al. 2011). Although administration of VPA induced a transient increase of HTLV-1 expression to expose virus-positive cells to the host immune response, clinical measures and PVL were stable overall. It has also been reported that the antibiotic minocycline significantly inhibited spontaneous lymphocyte proliferation and degranulation/IFN- γ expression in CD8⁺ T cells of patients with HAM/TSP, suggesting its potential for treatment (Enose-Akahata et al. 2012). Overall, more clinical trials with adequate power are needed in the future.

Risk factors for developing HAM/TSP

It is well-known that HAM/TSP occurs in only a minority of HTLV-1-infected individuals. A previous population association study in HTLV-1 endemic southwest Japan revealed that one of the major risk factors is the HTLV-1 PVL, because the PVL is significantly higher in patients with HAM/TSP than in ACs (Nagai et al. 1998). A higher PVL in patients with HAM/TSP than in ACs was also observed in other endemic areas such as the Caribbean (Manns et al. 1999), South America (Adaui et al. 2006), and the Middle East (Sabouri et al. 2005). In southwest Japan, it was suggested that genetic factors such as the human leukocyte antigen (HLA) genotype are related to the high PVL in patients with HAM/TSP and genetic relatives. Namely, possession of the HLA class I genes HLA-A*02 and Cw*08 was associated with a statistically significant reduction in both HTLV-1 PVL and the risk of HAM/TSP, whereas possession of HLA class I HLA-B*5401 and class II HLA-DRB1*0101 predispose to HAM/TSP in the same population (Jeffery et al. 2000; Jeffery et al. 1999). Because the function of class I HLA proteins is to present antigenic peptides to CTL, these results imply that individuals with HLA-A*02 or HLA-Cw*08 mount a particularly efficient CTL response against HTLV-1, which may be an important determinant of PVL and the risk of HAM/TSP. In accordance with this observation, it has been reported that CTL spontaneously kill autologous HTLV-1-infected cells *ex vivo* (Hanon et al. 2000), granzymes and perforin are more highly expressed in individuals with a low

PVL (Vine et al. 2004), and the lytic efficiency of the CD8⁺ T cell response (i.e., the fraction of autologous HTLV-1-expressing cells eliminated per CD8⁺ T cell per day) was inversely correlated with both PVL and the rate of spontaneous proviral expression (Kattan et al. 2009). Furthermore, the major histocompatibility complex (MHC) class I tetramer analysis of lymphocytes isolated from the CSF of patients with HAM/TSP showed even higher frequencies of HTLV-1 Tax11-19-specific, HLA-A*02-restricted CD8⁺ lymphocytes compared with PBMCs (Nagai et al. 2001b). These findings indicate that an increased proliferation or migration of HTLV-1-infected and/or HTLV-1-specific lymphocytes to the central nervous system might be closely associated with the pathogenesis of HAM/TSP (Hayashi et al. 2008a), and the CTLs against HTLV-1 reduce both PVL and the risk of HAM/TSP. Recently, using a combination of computational and experimental approaches, MacNamara et al. reported that a CTL response against HBZ restricted by protective HLA alleles such as HLA-A*02 or Cw*08, but not a response to the immunodominant protein Tax, also determines the outcome of HTLV-1 infection (Macnamara et al. 2010).

Meanwhile, analysis of non-HLA host genetic factors by candidate gene approaches revealed that non-HLA gene polymorphisms also affect the risk of developing HAM/TSP. Namely, the TNF- α promoter -863 A allele (Vine et al. 2002) and the longer CA repeat alleles of matrix metalloproteinase 9 promoter (Kodama et al. 2004) predisposed to HAM/TSP, whereas IL-10 -592 A (Sabouri et al. 2004), stromal-derived factor 1 +801A (Vine et al. 2002), and IL-15 +191 C alleles (Vine et al. 2002) conferred protection against HAM/TSP. The polymorphisms in the matrix metalloproteinase 9 and IL-10 promoters were each associated with differences in the HTLV-1 Tax-mediated transcriptional activity of the respective gene (Kodama et al. 2004; Sabouri et al. 2004). However, the contributions of these non-HLA genes to the pathogenesis of HAM/TSP are largely unknown and these data have not yet been reproduced in different populations. Further candidate gene studies together with genome-wide association studies in different ethnic populations in a larger sample size may provide evidence for the association of non-HLA genes with the pathogenesis of HAM/TSP.

It has been reported that the lifetime risk of developing HAM/TSP differs among ethnic groups, ranging between 0.25 and 4 %. The annual incidence of HAM/TSP is higher among Jamaican subjects than among Japanese subjects (20 versus 3 cases/100,000 population), with a two to three times greater risk for women in both populations (Hisada et al. 2004; Kramer et al. 1995; Nakagawa et al. 1995; Osame et al. 1990). Although most studies of HTLV-1 genotype have reported no association between variants of HTLV-1 and the risk of HAM/TSP, Furukawa et al. reported the association between HTLV-1 Tax gene variation and the risk of HAM/TSP (Furukawa et al. 2000). Tax subgroup A, which belongs to cosmopolitan

subtype A, was more frequently observed in patients with HAM/TSP, and this association was independent of the protective effect of HLA-A*02. Interestingly, HLA-A*02 appeared to give protection against only one of the two prevalent sequence variants of HTLV-1, Tax subgroup B, which belongs to cosmopolitan subtype B, but not against Tax subgroup A in the Japanese population (Furukawa et al. 2000). Jamaican subjects, who had a higher annual incidence of HAM/TSP, also have cosmopolitan subtype A, whereas approximately 80 % of Japanese subjects, who had a lower annual incidence of HAM/TSP, have cosmopolitan subtype B. Interestingly, HLA-A*02 did not appear to provide protection against HAM/TSP development with cosmopolitan subtype A in a population in Iran (Sabouri et al. 2005).

To test whether the genomic integration site determines the abundance and the pathogenic potential of an HTLV-1-positive T cell clone, Gillet et al. recently reported the results of high-throughput mapping and quantification of HTLV-1 proviral integration in the host genome (Gillet et al. 2011). They mapped >91,000 unique insertion sites (UISs) of the provirus in primary PBMCs from 61 HTLV-1-infected individuals and showed that a typical HTLV-1-infected host carries between 500 and 5,000 UISs in 10 µg of PBMC genomic DNA. They calculated an oligoclonality index to quantify the clonality of HTLV-1-infected cells in vivo and found that the oligoclonality index did not distinguish between ACs and patients with HAM/TSP and that there was no correlation between the oligoclonality index and HTLV-1 PVL in either ACs or patients with HAM/TSP. These results indicate that the higher PVL observed in patients with HAM/TSP was attributable to a larger number of UISs but not, as previously thought, to a difference in clonality. They also obtained evidence that the abundance of established HTLV-1 clones is determined by genomic features of the host DNA flanking the provirus. Namely, HTLV-1 clonal expansion in vivo is favored by a proviral integration site near a region of host chromatin undergoing active transcription or same-sense transcriptional orientation of the provirus. In contrast, negative selection of infected clones, probably by CTLs during chronic infection, favors establishment of proviruses integrated in transcriptionally silenced DNA, and this selection is more efficient in ACs than in HAM/TSP, indicating the selection of HTLV-1-infected T cell clones with low pathogenic potential. More recent reports indicate that circulating HTLV-1-positive cells each contain a single integrated proviral copy (Cook et al. 2012), and cells expressing HTLV-1 Tax protein (i.e., viral protein expression) were significantly more frequent in clones of low abundance in vivo, whereas certain transcription start sites immediately upstream of the viral integration site were associated with virus latency (i.e., no viral protein expression). In particular, Tax-expressing, more “pathogenic” clones were efficiently controlled by the immune response, especially CTLs, whereas non-Tax-expressing “invisible” infected clones were associated with mitotic clonal expansion in vivo (Melamed et al. 2013).

The innate immune response in HAM/TSP

Type I IFN is a key innate immune cytokine produced by cells in response to viral infection. The type I IFN response protects cells against invading viruses by inducing the expression of IFN-stimulated genes, which execute the antiviral effects of IFN (Samuel 2001). The IFN-stimulated genes then generate soluble factors including cytokines that activate adaptive immunity or directly inhibit the virus itself (Liu et al. 2011). In PBMCs of HTLV-1-infected individuals, the level of HTLV-1 messenger RNA is very low and viral protein is not detectable, but these molecules are rapidly expressed after a short time in culture in vitro (Hanon et al. 2000). However, the mechanisms of this phenomenon are largely unknown. Recently, it has been reported that HTLV-1 expression in HTLV-1-infected T cells is suppressed by stromal cells (i.e., epithelial cells and fibroblasts) in culture through type I IFNs (Kinpara et al. 2009). Namely, HTLV-1 Gag protein expression was suppressed when contacted with stromal cells and restored when separated from the stromal cells. Although neutralizing antibodies against human IFN- α/β receptor only partly abrogated this phenomenon, the results indicate that the innate immune system suppresses HTLV-1 expression in vitro and in vivo, at least through type I IFN. More recently, it has been reported that IFN-stimulated genes were overexpressed in circulating leukocytes and the expression correlated with the clinical severity of HAM/TSP (Tattermusch et al. 2012).

Previous reports indicated that patients with HAM/TSP had both a lower frequency and a lower activity of natural killer (NK) cells (especially the CD3⁺ CD16⁺ subset) than ACs, although the results were not normalized with respect to the PVL (Yu et al. 1991). Because an important mechanism of induction of NK cell-mediated killing is recognition by the NK cell of a complex of the non-polymorphic MHC molecule HLA-E bound to a peptide derived from the signal sequence of some other MHC class I molecules, the synthetic tetramers of HLA-E with the HLA-G signal sequence peptide were used to identify NK cells in patients with HAM/TSP (Saito et al. 2003). The results clearly showed a lower frequency of HLA-E tetramer-binding cells in patients with HAM/TSP than in ACs; as in the earlier studies (Yu et al. 1991), this reduction in frequency was particularly notable in the CD3⁺ cells, whereas there was no significant difference in the frequency of HLA-E tetramer-binding CD3⁻ cells between patients with HAM/TSP and ACs (Saito et al. 2003). Recent reports also suggest that the frequency of invariant natural killer T (NKT) cells in the peripheral blood of patients with HAM/TSP is significantly decreased when compared with that in healthy subjects and/or ACs (Azakami et al. 2009; Ndhlovu et al. 2009). These findings indicated that the activity of the NK or NKT cell response was associated with the presence or absence of HAM/TSP. Interestingly, a previous uncontrolled preliminary trial of viable *Lactobacillus casei* strain Shirota-containing

fermented milk in patients with HAM/TSP resulted in a significant increase in NK cell activity with improvements in clinical symptoms (Matsuzaki et al. 2005). Thus, circulating NK and NKT cells might also play an important role in the disease progression and pathogenesis of HAM/TSP.

The acquired immune response in HAM/TSP

It has been reported that patients with HAM/TSP generally have higher anti-HTLV-1 Ab titers than ACs with a similar PVL (Ishihara et al. 1994; Kira et al. 1992; Nagasato et al. 1991), suggesting the existence of an augmented humoral immune response to HTLV-1. Interestingly, although Ab responses to the immunodominant epitopes of the HTLV-1 Envelope (Env) proteins were similar in all three clinical groups of HTLV-1 infection (HAM/TSP, ATL, and ACs), reactivity to four Tax immunodominant epitopes was higher in patients with HAM/TSP (71–93 %) than in patients with ATL (4–31 %) or ACs (27–37 %) (Lal et al. 1994). A recent report indicates that the Ab response against HBZ was associated with reduced CD4⁺ T cell activation in patients with HAM/TSP, and HBZ-specific Ab inhibited spontaneous *in vitro* lymphocyte proliferation in the PBMCs of patients with HAM/TSP (Enose-Akahata et al. 2013). Among these anti-HTLV-1 antibodies, anti-Env Ab is particularly important because some anti-Env Abs have neutralizing activity against HTLV-1. Antisera raised against recombinant HTLV-1 Env polypeptides (Kiyokawa et al. 1984; Nakamura et al. 1987), vaccinia virus containing the HTLV-1 env gene (Hakoda et al. 1995; Shida et al. 1987), immunization with neutralizing epitope peptides (Tanaka et al. 1994), and passive transfer of human immunoglobulin G that has neutralizing activity (Murata et al. 1996; Tanaka et al. 1993) were all shown to neutralize HTLV-1 infectivity. In HTLV-1 infection, the roles of HTLV-1 neutralizing Ab *in vivo* are still largely unknown. It will be interesting to examine whether HTLV-1 neutralizing Ab titers correlate with disease status and PVL in infected individuals. Because the mutation rate of HTLV-1 provirus is significantly lower than that of HIV-1, passive immunization with human monoclonal Ab may be a beneficial and effective method to prevent HTLV-1 infection.

Antiviral CD4⁺ T cell responses are of central importance in driving B cell and CD8⁺ T cell responses *in vivo*. The most common HTLV-1 antigen recognized by CD4⁺ T cells is the Env protein (Goon et al. 2004b; Kitze et al. 1998), in contrast to the immunodominance of Tax in the CD8⁺ T cell response (Goon et al. 2004a; Jacobson et al. 1990; Kannagi et al. 1991). At a similar PVL, patients with HAM/TSP had a significantly increased frequency of virus-specific CD4⁺ T cells compared with ACs (Goon et al. 2004b; Nose et al. 2007). The antiviral T-helper (Th) 1 phenotype is also dominant among HTLV-1-specific CD4⁺ T cells in both ACs and patients with HAM/TSP (Goon et al. 2002), and there is a higher frequency of

IFN- γ , TNF- α , and IL-2 production by CD4⁺ T cells in patients with HAM/TSP compared with ACs of a similar PVL (Goon et al. 2002; Goon et al. 2003). A role for CD4⁺ T cells in initiating and causing HAM/TSP is also consistent with the immunogenetic observations that the possession of HLA-DRB1*0101, which restricts the immunodominant epitope of HTLV-1 Env gp21, was associated with susceptibility to HAM/TSP in independent HTLV-1-infected populations in southern Japan (Jeffery et al. 1999, 2000) and northeast Iran (Sabouri et al. 2005). Accordingly, a synthetic tetramer of DRB1*0101 and the immunodominant HTLV-1 Env380-394 peptide was used to analyze Env-specific CD4⁺ T cells directly *ex vivo* (Nose et al. 2007). The results showed that the frequency of tetramer⁺ CD4⁺ T cells was significantly higher in patients with HAM/TSP than in ACs with a similar PVL. Furthermore, direct *ex vivo* analysis of tetramer⁺ CD4⁺ T cells from two unrelated DRB1*0101-positive patients with HAM/TSP indicated that certain T cell receptor V β s were utilized and antigen-specific amino acid motifs were identified in complementarity determining region 3 from both patients. These results suggest that the observed increase in virus-specific CD4⁺ T cells in patients with HAM/TSP, which may contribute to CD4⁺ T cell-mediated antiviral immune responses and to an increased risk of HAM/TSP, was not simply due to the rapidly growing HTLV-1-infected CD4⁺ T cells but was the result of *in vivo* selection by specific MHC-peptide complexes, as observed in freshly isolated HLA-A*0201/Tax11-19 tetramer⁺ CD8⁺ T cells (Saito et al. 2001) and muscle-infiltrating cells from patients with HAM/TSP and HTLV-1-infected patients with polymyositis (Saito et al. 2002).

Previous reports indicated that HTLV-1-specific CD8⁺ CTLs are typically abundant, chronically activated, and mainly targeted to the viral transactivator protein Tax (Bangham 2000). Further, as already mentioned, the median PVL in PBMCs of patients with HAM/TSP was more than 10 times higher than that in ACs, and a high PVL was also associated with an increased risk of progression to disease (Nagai et al. 1998). Furthermore, HLA-A*02 and HLA-Cw*08 genes were independently and significantly associated with a lower PVL and a lower risk of HAM/TSP (Jeffery et al. 2000; Jeffery et al. 1999), and CD8⁺ T cells efficiently kill autologous Tax-expressing lymphocytes in fresh PBMCs in HTLV-1-infected individuals (Hanon et al. 2000). These data have raised the hypothesis that the class I-restricted CD8⁺ CTL response plays a critical part in limiting HTLV-1 replication *in vivo* and that genetically determined differences in the efficiency of the CTL response to HTLV-1 account for the risk of developing HAM/TSP. The analysis of gene expression profiles using microarrays in circulating CD4⁺ and CD8⁺ lymphocytes indicated that granzymes and perforin are more highly expressed in individuals with a low PVL (Vine et al. 2004), suggesting that a strong CTL response is associated

with a low PVL and a low risk of HAM/TSP. In accordance with this observation, the lytic capacity of HTLV-1-specific CTLs in patients with HAM/TSP and ACs quantified by a CD107a mobilization assay showed significantly lower CD107a staining in HTLV-1-specific CTLs in patients with HAM/TSP than in ACs (Sabouri et al. 2008); this suggests that patients with HAM/TSP have a high frequency of HTLV-1-specific CD8⁺ T cells with poor lytic capacity, whereas ACs have a lower frequency of cells with high lytic capacity. Moreover, it has been reported that the high CTL avidity, which is closely associated with the lytic efficiency of CTLs, correlates with low PVL and proviral gene expression (Kattan et al. 2009), indicating that the efficient control of HTLV-1 *in vivo* depends on the quality of CTLs, which determines the position of virus–host equilibrium and also the outcome of persistent HTLV-1 infection. More recently, MacNamara et al. (Macnamara et al. 2010) showed that HLA class I alleles, which strongly bind oligopeptides from the HBZ protein, enable the host to have a more effective immune response against HTLV-1; therefore, such individuals have a lower PVL and are more likely to be asymptomatic. Another recent report showed the presence of HBZ-specific CD4⁺ and CD8⁺ cells *in vivo* in patients with HAM/TSP and in ACs and a significant association between the HBZ-specific CD8⁺ cell response and asymptomatic HTLV-1 infection (Hilburn et al. 2011). These findings provide strong evidence to support the hypothesis of the crucial role of CTLs and confirm the importance of HBZ for persistent infection. However, because the frequency of HTLV-1-specific CD8⁺ T cells was significantly elevated in patients with HAM/TSP compared with ACs (Greten et al. 1998; Nagai et al. 2001a), and these cells have the potential to produce proinflammatory cytokines (Kubota et al. 1998), there is debate on the role of HTLV-1-specific CD8⁺ T cells, namely, whether these cells contribute to the inflammatory and demyelinating processes of HAM/TSP or whether the dominant effect of such cells *in vivo* is protective against disease, although a protective role and a pathogenic role of CTLs are not mutually exclusive. Indeed, there are other examples of viral infections in which the virus-specific CTLs exert both beneficial (antiviral) and detrimental (inflammatory) effects, such as lymphocytic choriomeningitis virus infection in the mouse (Klenerman and Zinkernagel 1997). It is difficult to separate cause and effect in analyzing the association between T cell attributes and the efficiency of viral control in a persistent infection at equilibrium.

Regulatory T cells (Tregs) are important mediators of peripheral immune tolerance and play an important role in chronic viral infections. HTLV-1 preferentially and persistently infects CD4⁺ CD25⁺ lymphocytes *in vivo* (Yamano et al. 2005), which contain the majority of the Foxp3⁺ Tregs (Sakaguchi et al. 2006). In patients with HAM/TSP, the percentage of Foxp3⁺ Tregs in CD4⁺ CD25⁺ cells is lower than that in ACs and uninfected healthy controls (Oh et al. 2006; Yamano et al. 2005), whereas the percentage of Foxp3⁺ cells in the CD4⁺

population tends to be higher in patients with HAM/TSP than in ACs (Best et al. 2009; Hayashi et al. 2008b; Toulza et al. 2008). Because CD25 is induced by HTLV-1 Tax oncoprotein (Inoue et al. 1986), the proportion of Foxp3⁺ cells falls in the CD4⁺ CD25⁺ population, which contains both Tregs and activated non-Tregs, in HTLV-1-infected individuals, especially patients with HAM/TSP. Therefore, it is inappropriate to use CD25 as a marker of Tregs in HTLV-1 infection, and the best current working definition of Treg phenotype is CD4⁺ Foxp3⁺. The high frequency of CD4⁺ Foxp3⁺ T cells in HTLV-1-infected individuals is maintained by CCL22 produced by HTLV-1-infected PBMCs (Toulza et al. 2010). The frequency of HTLV-1-negative Foxp3⁺ CD4⁺ cells positively correlated with the HTLV-1 PVL (Hayashi et al. 2008a; Toulza et al. 2008), and the CTL activity negatively correlated with the frequency of HTLV-1-negative Foxp3⁺ CD4⁺ cells (Toulza et al. 2008). These results suggest that an increase in HTLV-1-negative Foxp3⁺ CD4⁺ Tregs is one of the chief determinants of the efficiency of T cell-mediated immune control of HTLV-1. If such Tregs reduce CTL activity, which in turn increases the HTLV-1 PVL, this activity increases the risk of developing HAM/TSP.

Dendritic cells and the other reservoirs of HTLV-1

Dendritic cells (DCs) are antigen-presenting cells that play a critical role in the regulation of the adaptive immune response. In HTLV-1 infection, it has been shown that the DCs from patients with HAM/TSP were infected with HTLV-1 (Macatonia et al. 1992), and the development of HAM/TSP is associated with rapid maturation of DCs (Ali et al. 1993). *In vitro* culture of lymphocytes from HTLV-1-infected individuals results in “spontaneous lymphocyte proliferation” (SLP), which is the *in vitro* proliferation of PBMCs without any exogenous stimuli such as antigen or mitogen. In patients with HAM/TSP, the levels of SLP reflect the severity of the disease (Ijichi et al. 1989; Itoyama et al. 1988). Interestingly, depletion of DCs from the PBMCs of patients with HAM/TSP abolished SLP, whereas supplementing DCs restores proliferation (Macatonia et al. 1992); supplementing B cells or macrophages had no effect. A DC-dependent mechanism of SLP was further supported by data showing that antibodies to MHC class II, CD86, and CD58 can block SLP (Makino et al. 1999). Recently, Jones et al. demonstrated that human-derived myeloid and plasmacytoid DCs are susceptible to infection with cell-free HTLV-1 and that HTLV-1-infected DCs can rapidly transfer virus to autologous primary CD4⁺ T cells (Jones et al. 2008). Furthermore, it was recently demonstrated that transmission of HTLV-1 from DCs to T cells was mediated primarily by DC-SIGN (Jain et al. 2009), and the DCs are the major cell type responsible for the generation and maintenance of Tax-specific CD8⁺ T cells both *in vitro* and *in vivo* (Manuel et al. 2009).

These findings suggest that the interaction of DCs with HTLV-1 is also crucial for the pathogenesis of HAM/TSP. Moreover, using transgenic mouse models that permit conditional transient depletion of CD11c+ DCs, and a chimeric HTLV-1 that carries the envelope gene from Moloney murine leukemia virus, Rahman et al. demonstrated the critical role of DCs in their ability to mount both innate and adaptive immune responses during early cell-free HTLV-1 infection (Rahman et al. 2011, 2010). Because HTLV-1 can impair the differentiation of monocytes into DCs (Nascimento et al. 2011), the interaction of DCs with HTLV-1 plays a central part in the persistence and pathogenesis of HTLV-1.

Conclusions

During the three decades since the discovery of HTLV-1, advances in research have successfully helped us to understand the clinical features of HTLV-1-associated diseases and the virological properties of HTLV-1, although the precise mechanism of disease pathophysiology is still incompletely understood and treatment is still unsatisfactory. Accumulating evidence suggests that the virus–host immunological interactions play a pivotal role in the pathogenesis of HAM/TSP. A genetically determined, less efficient CTL response against HTLV-1 may cause higher PVL and antigen expression in infected individuals, which lead to the activation and expansion of antigen-specific T cell responses, subsequent induction of large amounts of proinflammatory cytokines and chemokines, and progression of the development of HAM/TSP. Future studies should be conducted to identify the precise mechanism of disease development to allow effective treatment and prevention of disease. This will require the development of a humanized small animal model that could be exploited as a tool for screening and evaluation of HTLV-1-associated diseases.

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Elimination of Human T Cell Leukemia Virus Type-1-Infected Cells by Neutralizing and Antibody-Dependent Cellular Cytotoxicity-Inducing Antibodies Against Human T Cell Leukemia Virus Type-1 Envelope gp46

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Abstract

Human T cell leukemia virus type-1 (HTLV-1) is prevalent worldwide with foci of high prevalence. However, to date no effective vaccine or drug against HTLV-1 infection has been developed. In efforts to define the role of antibodies in the control of HTLV-1 infection, we capitalized on the use of our previously defined anti-gp46 neutralizing monoclonal antibody (mAb) (clone LAT-27) and high titers of human anti-HTLV-1 IgG purified from HAM/TSP patients (HAM-IgG). LAT-27 and HAM-IgG completely blocked syncytium formation and T cell immortalization mediated by HTLV-1 *in vitro*. The addition of these antibodies to cultures of CD8⁺ T cell-depleted peripheral blood mononuclear cells (PBMCs) from HAM/TSP patients at the initiation of culture not only decreased the numbers of Tax-expressing cells and the production of HTLV-1 p24 but also inhibited the spontaneous immortalization of T cells. Coculture of *in vitro*-HTLV-1-immortalized T cell lines with autologous PBMCs in the presence of LAT-27 or HAM-IgG, but not an F(ab')₂ fragment of LAT-27 or non-neutralizing anti-gp46 mAbs, resulted in depletion of HTLV-1-infected cells. A 24-h ⁵¹Cr release assay showed the presence of significant antibody-dependent cellular cytotoxicity (ADCC) activity in LAT-27 and HAM-IgG, but not F(ab')₂ of LAT-27, resulting in the depletion of HTLV-1-infected T cells by autologous PBMCs. The depletion of natural killer (NK) cells from the effector PBMCs reduced this ADCC activity. Altogether, the present data demonstrate that the neutralizing and ADCC-inducing activities of anti-HTLV-1 antibodies are capable of reducing infection and eliminating HTLV-1-infected cells in the presence of autologous PBMCs.

Introduction

HUMAN T CELL LEUKEMIA VIRUS type-1 (HTLV-1) is the first human retrovirus that was etiologically associated with adult T cell leukemia (ATL) and HTLV-1-associated myelopathy/tropical spastic paraparesis (HAM/TSP).¹⁻⁴ HTLV-1 is prevalent worldwide with foci of high prevalence in southwest Japan, the Caribbean islands, South America, and a part of Central Africa. The total number of HTLV-1 carriers is currently estimated to be 10–20 million.⁵ The majority of HTLV-1 carriers remain asymptomatic throughout their lives, and approximately 5% of HTLV-1-infected individuals will develop either ATL or HAM/TSP after prolonged latency periods.

HTLV-1 is transmitted through contact with bodily fluids containing infected cells most often from mother to child through breast milk or via blood transfusion. It has been previously established that HTLV-1 efficiently spreads from cell to cell via the formation of virological synapses.⁶ More recently, however, the formation of extracellular HTLV-1 viral particles similar to the formation of bacterial films has also been shown to be effective in viral transmission.⁷ HTLV-1-antigen-expressing cells are difficult to detect at least in fresh peripheral blood mononuclear cells (PBMCs) from HTLV-1-infected individuals.⁸ However, when these PBMCs are isolated from the blood and cultured *in vitro*, some T cells begin to produce HTLV-1 antigen^{9,10} followed by spontaneous immortalization of the cells in media containing interleukin-2 (IL-2).¹¹

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Although it has been suggested that HTLV-1 can stay dormant in infected cells and become resistant to immune effector mechanisms by ratcheting down its antigen production,¹² the continued presence of strong CD8⁺ cytotoxic T lymphocyte (CTL) responses¹³ and readily detectable levels of antibodies specific for HTLV-1 antigens in HTLV-1-infected people¹⁴ indicates that persistent production of HTLV-1 must occur *in vivo* to maintain such effector mechanisms. Escape from immune effector mechanisms by spontaneous mutation of key residues is unlikely, due to the high degree of genomic stability that is characteristic of the HTLV-1 genome.¹⁵ It has been suggested that HTLV-1-infected cells expressing HTLV-1 antigens occur at a low enough frequency that they are constantly being eliminated by HTLV-1-specific CTL *in vivo*¹⁰ without leading to immune exhaustion. Besides CTL and virus neutralizing antibodies, there has been renewed interest in the potential role of antibody-dependent cellular cytotoxicity (ADCC) as an effector mechanism against a number of viral infections. This view has been highlighted by the recent demonstration of the potential role of ADCC in the only known partially successful human RV144 trial of a vaccine against human immunodeficiency virus type-1 (HIV-1).¹⁶ The ADCC activity against HTLV-1 was first reported by Miyakoshi *et al.* in 1984¹⁷ followed by a number of other reports.¹⁸⁻²¹

So far, several lines of evidence show that the HTLV-1 envelope gp46 antigen serves as a major target of ADCC.²²⁻²⁴ Antibodies against gp46 antigen are commonly detected in the sera of HTLV-1-infected individuals.²⁵⁻²⁷ However, the precise role of ADCC effector mechanism(s) in controlling HTLV-1 infection has been lacking. A possible involvement of anti-HTLV-1 antibodies in the suppression of spontaneous HTLV-1 antigen expression by HTLV-1-infected cells was first reported by Tochikura *et al.*²⁸ These investigators showed that serum IgG from HTLV-1-infected donors interfered with HTLV-1 antigen expression by *in vitro*-cultured PBMCs from both ATL patients and healthy HTLV-1 carriers. However, the precise mechanism by which this was mediated remained unclear.

In efforts to define the role of antibodies with neutralizing and ADCC-inducing activities in the control of HTLV-1 infection, we capitalized on the use of our previously defined rat anti-gp46 neutralizing monoclonal antibody (mAb) (LAT-27)²⁹ and pooled human anti-HTLV-1 IgG purified from HAM/TSP patients (HAM-IgG). Studies were conducted to evaluate the potential of these antibodies to block HTLV-1 infection and eliminate HTLV-1-infected cells from autologous T cell cultures that had previously been infected with HTLV-1 *in vivo* or *in vitro*. Results of these studies show that monoclonal LAT-27 and the polyclonal HAM-IgG are not only capable of mediating neutralization and ADCC, but are also highly effective in the elimination of HTLV-1-infected cells in the presence of fresh autologous PBMCs while preventing *de novo* infection with HTLV-1.

Materials and Methods

Reagents

The medium used throughout was RPMI 1640 medium (Sigma-Aldrich, Inc., St. Louis, MO) supplemented with

10% fetal calf serum (FCS), 100 U/ml of penicillin, and 100 µg/ml of streptomycin (hereinafter called RPMI medium). Anti-human CD3 (clone OKT-3) and anti-CD28 (clone 28.2) mAbs were purchased from the American Type Culture Collection (Rockville, MD) and Biologend (San Diego, CA), respectively.

The rat and mouse mAbs utilized in the studies reported herein were produced and characterized by our laboratory previously.²⁹⁻³⁴ These antibodies were rat IgG2b anti-gp46 (clones LAT-27 and LAT-25), rat IgG2a anti-gp46 (clone LAT-12), rat IgG2b anti-HCV (clone Mo-8), rat IgG2a anti-HTLV-1 p24 (clone WAG-24), mouse IgG1 anti-HTLV-1 gp46 (clone MET-3), mouse IgG3 anti-HTLV-1 Tax (clone Lt-4), mouse IgG1 anti-p24 (clone NOR-1), and mouse IgG1 anti-HIV-1 p24 (clone 2C2). These in-house mAbs were purified from the ascites fluids of groups of CB.17-SCID mice carrying the appropriate hybridoma cell line. The ascites fluid was subjected to ammonium sulfate precipitation followed by gel filtration using Superdex G-200 (GE Healthcare, Tokyo, Japan). Aliquots of these mAbs were labeled with either fluorescein isothiocyanate (FITC), Alexafluor 488, Alexafluor 647, HRP (Dojindo, Kumamoto, Japan), or Cy-5 (GE Healthcare) according to the manufacturer's instructions. The FITC- or phycoerythrin (PE)-labeled mouse mAbs against human CD3, CD4, CD8, CD14, CD16, CD19, or CD56 and unlabeled mouse anti-CD16 and anti-CD32 mAbs were purchased from Abcam.

For cell depletion, magnetic beads labeled with anti-CD4, CD8, CD14, CD16, CD19, and antimouse IgG (Dyna) and those labeled with anti-CD56 mAb (LifeTec) were used according to the manufacturer's recommendations. Mitomycin-C (MMC) was commercially purchased from Kyowa Kirin (Tokyo, Japan) and used at 50 µg/ml in RPMI medium. A purified F(ab')₂ fragment of LAT-27 IgG generated by enzymatic digestion of LAT-27 IgG was purchased from IBL Inc. (Gunma, Japan). Human IgG was purified from pooled plasma from three normal donors (normal IgG) and three HAM patients (HAM-IgG) using protein-G affinity purification kits (GE Healthcare).

The protocols for the use of human PBMCs and animals were approved by the Human IRB and the Institutional Animal Care and Use Committee (IACUC) on clinical and animal research of the University of the Ryukyus prior to initiation of the present study.

Cell cultures

PBMCs were isolated from heparinized blood by standard density gradient centrifugation using Lympholyte (Cedarlane, Burlington, Canada). Some PBMCs were cryopreserved using a cell freezing media (Cell reservoir, Nakarai Tesque Inc., Kyoto, Japan). The method to activate PBMCs with anti-CD3 and CD28 mAbs has been described previously.³⁴ The HTLV-1-producing T cell lines utilized included MT-2, HUT102, IL-2-dependent CD4⁺CD8⁺ILT-M1 cells derived from an HAM/TSP patient, CD4⁺CD8⁻ILT-H2 cells, ATL-3 cells derived from ATL patients, and a number of other T cell lines derived from normal PBMCs following *in vitro* immortalization by cocultivation with MMC-treated ILT-M1 cells. These cell lines were maintained in culture using RPMI medium containing 20 U/ml IL-2.