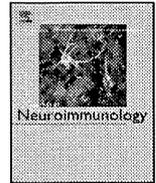




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Antibodies to LM1 and LM1-containing ganglioside complexes in Guillain–Barré syndrome and chronic inflammatory demyelinating polyneuropathy

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ABSTRACT

LM1 is localized in human peripheral nerve myelin. Antibodies to ganglioside complexes (GSCs) have been reported in Guillain–Barré syndrome (GBS). We investigated IgG antibodies to LM1 and two GSCs (GM1 and LM1, or GD1b and LM1) in the sera of each 40 patients with chronic inflammatory demyelinating polyneuropathy (CIDP) and GBS, using ELISA. We detected anti-LM1 antibody in five with GBS and seven with CIDP; anti-GM1/LM1 antibody in three with GBS and one with CIDP; and anti-GD1b/LM1 antibody in two with CIDP. Antibodies to LM1 and LM1-containing GSCs may be among the targets for autoimmunity in GBS and CIDP.

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1. Introduction

Guillain–Barré syndrome (GBS) is an acute immune-mediated polyneuropathy that is often preceded by an infection. The acute-phase sera of approximately 60% of patients with GBS contain anti-ganglioside antibodies. Each ganglioside is uniquely localized. Therefore, each anti-ganglioside antibody may influence clinical features by specifically binding to the regions where the target ganglioside is localized. Although there are reports of antibodies to several antigens such as PO, PMP22, Gal-C, and sulfatide, a positive reaction is rarely detected in patients with chronic inflammatory demyelinating polyneuropathy (CIDP) (Meléndez-Vásquez et al., 1997; Yan et al., 2001; Allen et al., 2005).

Antibodies to ganglioside complexes (GSCs) have recently been reported in patients with GBS and Fisher syndrome (FS) (Kaida et al., 2004; Kaida et al., 2006; Kaida et al., 2007; Kaida et al., 2008; Kanzaki et al., 2008; Ogawa et al., 2009). These antibodies may recognize novel epitopes that are formed by two different gangliosides. Gangliosides are components of the lipid raft on the plasma membrane. Anti-GSC antibodies may cause neuropathy by binding to the GSCs formed by the clustered gangliosides on the lipid rafts. Among the anti-GSC antibodies, the IgG antibodies to the GD1a/GD1b complex and the GD1b/GT1b complex are associated with GBS in

which a patient requires artificial ventilation (Kaida et al., 2007). IgG antibodies to the GM1/GalNAc-GD1a complex are associated with acute motor conduction block neuropathy (AMCBN), which is characterized by frequent conduction blocks (CBs) at the intermediate nerve segments and no sensory involvement (Kaida et al., 2008). IgM antibodies to GSCs have also been reported in patients with CIDP and IgM monoclonal gammopathy (Nobile-Orazio et al., 2010).

LM1 is a ganglioside localized in peripheral nerve myelin (Ogawa-Goto et al., 1992). It is a candidate target molecule in the demyelinating peripheral neuropathies. However, antibodies against LM1-containing GSCs have not been examined in patients with GBS or CIDP. Therefore, we investigated the activity of IgG antibodies to LM1 and LM1-containing GSCs (e.g., GM1/LM1 and GD1b/LM1) in patients with GBS and CIDP.

2. Materials and methods

2.1. The serum samples

Serum samples were obtained from 40 patients with GBS who were diagnosed in accordance with the established criteria (Asbury and Cornblath, 1990); 40 patients with CIDP who were diagnosed in accordance with the criteria proposed by European Federation of Neurological Societies and Peripheral Nerve Society (EFNS/PNS) (Joint Task Force of the EFNS and the PNS., 2005); 15 normal control subjects; and 40 patients with other neurological diseases (ONDs), which included 10 patients with motor neuron disease, four patients with Parkinson disease, three patients with spinocerebellar

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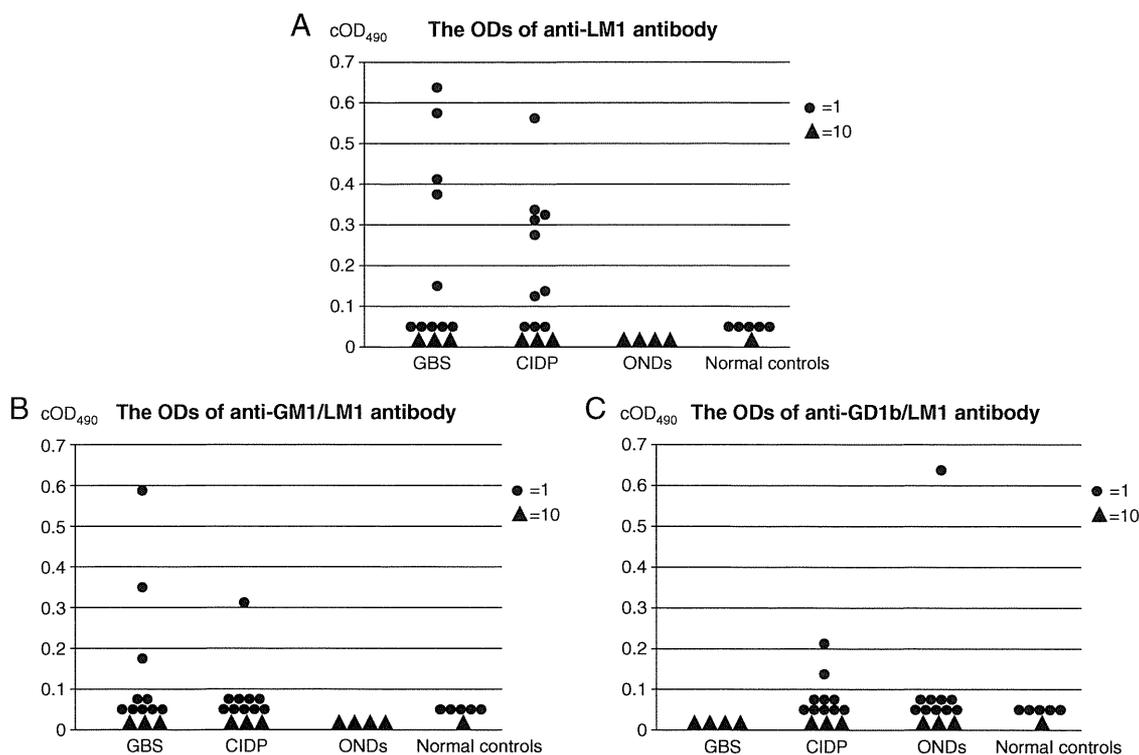


Fig. 1. The ODs of antibodies to LM1, GM1/LM1 and GD1b/LM1. Abbreviations: OD = optical density; CIDP = chronic inflammatory demyelinating polyneuropathy; GBS = Guillain-Barré syndrome; ONDs = other neurological diseases.

degeneration, two patients with multiple system atrophy, one patient with corticobasal degeneration, six patients with multiple sclerosis, four patients with neuromyelitis optica (NMO), eight patients with

myasthenia gravis, and two patients with Lambert–Eaton syndrome. The sera from GBS patients were obtained in the acute phase (within 4 weeks after the onset of the disease) and the sera from CIDP patients were obtained in relapse. The sera from GBS and CIDP patients were collected before immunotherapy.

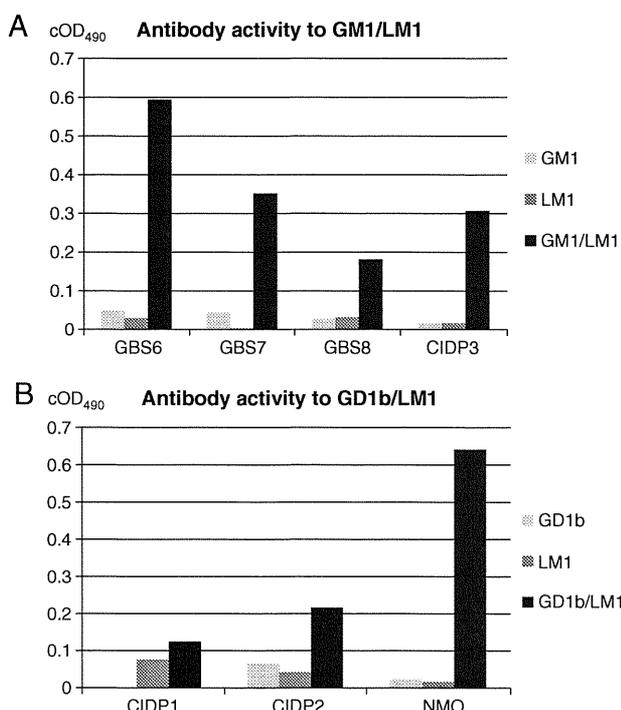


Fig. 2. The ODs of IgG antibodies to LM1-containing GSCs in antibody-positive patients. Abbreviations: OD = optical density; CIDP = chronic inflammatory demyelinating polyneuropathy; GBS = Guillain-Barré syndrome; NMO = neuromyelitis optica.

2.2. Enzyme-linked immunosorbent assay for detecting antibodies against gangliosides and LM1-containing ganglioside complexes

IgG antibodies to GM1, GD1b, LM1, the GM1/LM1 complex, and the GD1b/LM1 complex in the patients' sera were tested by enzyme-linked immunosorbent assay (ELISA), as previously reported (Kaida et al., 2004). A sample was considered positive for the presence of GM1 and GD1b antibodies when its optical density (OD) was greater than 0.1, as previously described (Kaida et al., 2004). A sample was considered positive for the presence of IgG antibodies to LM1 when its OD was greater than 0.12. (The OD was the value of the mean + 3 standard deviation [SD] of the ODs of 15 normal control subjects.) The antibodies to other gangliosides (e.g., GM2, GM3, GD1a, GD3, GT1b, GQ1b, and GalNAc-GD1a) were also examined in the same way, as previously described (Kaida et al., 2004). A sample was considered positive for the presence of antibodies to GM1/LM1 complex if one of the three criteria was fulfilled; 1) it was negative for antibodies to GM1 alone and LM1 alone and the ODs were greater than 0.12 in the wells that had been coated with both GM1 and LM1, 2) it was positive for antibodies to GM1 alone or LM1 alone and the ODs of the wells coated with both GM1 and LM1 were 0.2 higher than the ODs of the wells coated with GM1 alone and LM1 alone, 3) it was positive for antibodies to GM1 alone and LM1 alone and the ODs of the wells coated with both GM1 and LM1 were greater than the sum of the ODs of the wells coated with GM1 alone and those with LM1 alone. The same criteria were applied for the presence of antibodies to GD1b/LM1 complex.

2.3. The clinical and electrophysiological features of Guillain-Barré syndrome patients with anti-LM1 or LM1-containing GSCs IgG antibodies

Clinical and electrophysiological features were assessed in GBS patients with IgG antibodies to LM1 or LM1-containing GSCs. The patients were classified as having acute inflammatory demyelinating polyneuropathy (AIDP), acute motor axonal neuropathy (AMAN), equivocal, or inexcitable based on the criteria of Ho et al (Ho et al., 1995).

This study was approved by the internal review board of Kinki University School of Medicine.

3. Results

3.1. Antibodies to LM1 or LM1-containing GSCs

The presence of anti-LM1 IgG antibody was positive in five patients with GBS and seven patients with CIDP. In patients with ONDs and in normal control subjects, anti-LM1 IgG antibody was not detected (Fig. 1). Anti-GM1/LM1 IgG antibody was detected in three patients with GBS and in one patient with CIDP. Anti-GD1b/LM1 IgG antibody was detected in two patients with CIDP and in one OND patient who had NMO (Fig. 2). IgG antibody to LM1 alone was not present in the patient who had antibodies to LM1-containing GSCs. No patient had antibodies to both GM1/LM1 and GD1b/LM1. The prevalence of LM1, GM1/LM1, and GD1b/LM1 antibodies in patients and in normal control subjects was shown in Table 1.

3.2. Clinical and electrophysiological features of GBS patients with anti-LM1 or anti-GM1/LM1 IgG antibodies

All the eight sera from anti-LM1 or -GM1/LM1 antibody-positive GBS patients were obtained within 4 weeks and five of those were within 2 weeks after the onset of disease. The GBS patients with either anti-LM1 (five patients) or anti-GM1/LM1 (three patients) antibodies were all men. Of the five patients positive for anti-LM1 antibody, two had an antecedent respiratory infection and one had an antecedent gastrointestinal infection, whereas two of the three patients positive for anti-GM1/LM1 antibody had antecedent gastrointestinal infections. Four of the five patients positive for anti-LM1 antibodies and one of the three patients positive for anti-GM1/LM1 antibodies were classified as having AIDP. The remaining three were classified as equivocal. IgG antibodies to GD1b were detected in three patients. Among them, two patients were positive for anti-GM1/LM1 antibody and one patient was positive for anti-LM1 antibody. IgG antibody to GalNAc-GD1a was detected in one patient who had anti-LM1 antibody. No antibodies to other gangliosides were detected in four patients (Table 2).

4. Discussion

LM1 is the predominant ganglioside in human peripheral nerve myelin. GM1 is also present in human myelin and is mainly contained in motor nerve myelin (Ogawa-Goto et al., 1992). There is some indication that GM1 is present in the nodes of Ranvier (Kusunoki et al.,

Table 1
The prevalence of LM1, GM1/LM1, and GD1b/LM1 antibodies in patients and in normal control subjects.

Patient	LM1	GM1/LM1	GD1b/LM1	Total
GBS	5/40 (12.5%)	3/40 (7.5%)	0/40 (0%)	8/40 (20%)
CIDP	7/40 (17.5%)	1/40 (2.5%)	2/40 (5%)	10/40 (25%)
ONDs	0/40 (0%)	0/40 (0%)	1/40 (2.5%)	1/40 (2.5%)
Normal controls	0/15 (0%)	0/15 (0%)	0/15 (0%)	0/15 (0%)

Table 2

The clinical and electrophysiological features of anti-LM1 IgG antibody-positive or GM1/LM1 IgG antibody-positive patients with GBS.

Patient no.	Age/sex	Antecedent infection*	Involved cranial nerve(s)	Classification of NCS†	Presence of antibody to LM1 or GM1/LM1	Presence of antibodies to other gangliosides
GBS1	59/M	R	(-)	AIDP	LM1	(-)
GBS2	63/M	R	(-)	AIDP	LM1	(-)
GBS3	59/M	(-)	(-)	AIDP	LM1	GalNAc-GD1a
GBS4	47/M	GI	(-)	AIDP	LM1	GD1b
GBS5	43/M	(-)	(-)	Equivocal	LM1	(-)
GBS6	28/M	(-)	(-)	AIDP	GM1/LM1	GD1b
GBS7	31/M	GI	12	Equivocal	GM1/LM1	(-)
GBS8	36/M	GI	9, 10	Equivocal	GM1/LM1	GD1b

* GI, gastrointestinal infection; R, respiratory tract infection.

† NCS, nerve conduction study; AIDP, acute inflammatory demyelinating polyneuropathy; AMAN, acute motor axonal neuropathy;

1993). GD1b has been detected in human peripheral nerve myelin, predominantly in the sensory nerves (Ogawa-Goto et al., 1992). Immunolocalization of GD1b also has been reported in the paranodal myelin of human peripheral nerves and in the large neurons in the dorsal root ganglia (Kusunoki et al., 1993). In this study, we detected IgG antibody to LM1 in the sera of five of 40 patients (12.5%) with GBS and seven of 40 patients (17.5%) with CIDP, and we detected IgG antibodies to LM1-containing GSCs in the sera of three patients (7.5%) with GBS and three patients (7.5%) with CIDP. None of these antibodies were present in the normal controls or the patients with ONDs (except for one patient [2.5%] affected with NMO).

Four of five patients who were diagnosed with GBS and who had anti-LM1 IgG antibody were classified as having AIDP. Of the three GBS patients who were positive for anti-GM1/LM1 antibody, one was classified as having AIDP and the remaining two were equivocal. Considering that LM1 is distributed predominantly in the myelin of human peripheral nerve (Ogawa-Goto et al., 1992), anti-LM1 IgG antibody may be involved in the demyelinating process in AIDP and CIDP (Fredman et al., 1991; Ilyas et al., 1992; Yako et al., 1999; Harukawa et al., 2002; Susuki et al., 2002). Antibodies to LM1-containing GSCs may also cause demyelination by binding to the epitopes formed by two contiguous gangliosides in the plasma membrane.

IgG antibody to GD1b/LM1 was detected in one patient with NMO. In NMO, it is known that anti-AQP4 antibody and other antibodies associated with collagen diseases are frequently present. It is considered that humoral immunity is activated in NMO. LM1 is present in human microvascular endothelial cells that form the blood-brain barrier (BBB) (Kanda et al., 2004). It is therefore possible that the destruction of the BBB in patients with NMO may secondarily raise the serum level of the antibodies to LM1-containing GSCs. In contrast, all the sera from anti-LM1 or -GM1/LM1 antibody-positive patients with GBS were obtained in the acute phase, indicating that the antibodies were associated with the pathogenesis of GBS.

Target molecules have rarely been identified in patients with CIDP. The anti-ganglioside antibodies are usually associated with the AMAN form of GBS. Only a few gangliosides are known to be targeted by serum antibodies in patients with AIDP. The present investigation shows that LM1 and LM1-containing GSCs are among the target antigens in the sera of patients with CIDP and AIDP. Future studies involving a larger number of patients are necessary to elucidate how these antibodies cause demyelination in the peripheral nervous system.

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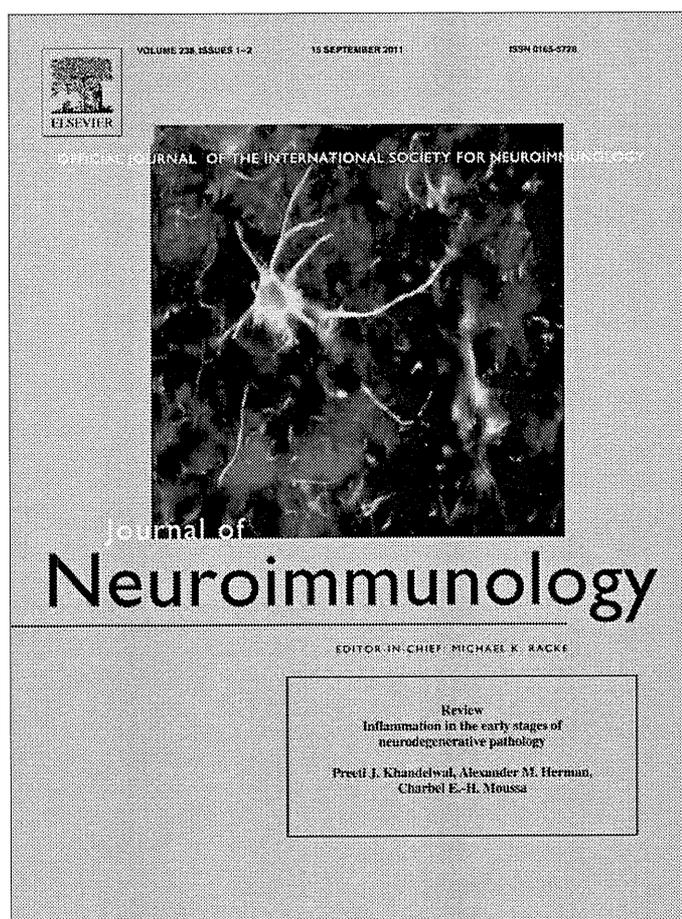
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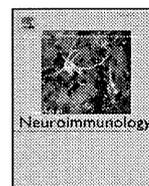
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Short communication

Four cases of anti-ganglioside antibody-positive neuralgic amyotrophy with good response to intravenous immunoglobulin infusion therapy

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ABSTRACT

Neuralgic amyotrophy (NA), which is an idiopathic disorder in the peripheral nerves, is characterized by an acute onset of unilateral pain in the proximal limbs followed by muscular weakness and wasting. Some cases of NA are thought to be related to immune pathogenic disorders such as Guillain-Barré syndrome (GBS). We report the case of four patients with NA who were positive for anti-N-acetylgalactosaminyl GD1a (anti-GalNAc-GD1a) antibodies, had a preceding infection, and showed a good response to intravenous immunoglobulin infusion therapy. Anti-ganglioside antibodies, especially the anti-GalNAc-GD1a antibody, may be a useful marker for predicting response to immune therapy.

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1. Introduction

Neuralgic amyotrophy (NA), or Parsonage–Turner syndrome, is an idiopathic disorder in the peripheral nerves that is characterized by an acute onset of shoulder and arm pain, which is followed by muscular weakness and wasting (Parsonage and Turner, 1948). Similar symptoms in the lower extremities have been reported and are considered to be a variant of NA (Takata et al., 2007). Although some cases have developed subsequent to surgery or trauma, the frequent association of this disorder with preceding infections and immunizations indicates an autoimmune pathogenesis (Suarez et al., 1996).

Frequent findings of increased levels of anti-ganglioside antibodies in sera obtained from patients in the acute phase of Guillain-Barré syndrome (GBS) and other inflammatory neuropathies have been reported, and clinical features of these diseases were associated with the reactivity of anti-ganglioside antibodies (Hartung et al., 1995; Kaida et al., 2000). Herein, we report the case of four patients with NA who presented with increased levels of anti-ganglioside antibodies in their sera and who had a good response to intravenous immunoglobulin infusion therapy (IVIg).

Abbreviations: NA, neuralgic amyotrophy; GalNAc-GD1a, N-acetylgalactosaminyl GD1a; GBS, Guillain-Barré syndrome; IVIg, immunoglobulin infusion therapy; CMV, cytomegalovirus; MRC, Medical Research Council.

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2. Case Report

2.1. Case 1

A 34-year-old woman presented with fever and vomiting which was followed by right shoulder pain that radiated to the forearm after 2 weeks and weakness in the right arm after 4 weeks. Her medical history was normal. Physical examination showed slightly decreased strength in the proximal and distal muscles of the right arm (Medical Research Council [MRC] score, 4). There was a slight diminution of touch sensation in the right hand. Tendon reflexes were normal. Laboratory tests were normal for complete blood counts, liver and renal function tests, thyroid hormone tests, and immunological screening. An enzyme-linked immunosorbent assay (ELISA) was performed for antibodies to the ganglioside as described previously (Kusunoki et al., 1994). Ganglioside antigens used in the ELISA were 200 ng each of GM1, GM2, GD1a, GD1b, GD3, GT1a, GT1b, GQ1b, and N-acetylgalactosaminyl GD1a (GalNAc-GD1a). The ELISA was also performed for antibodies to ganglioside complex containing 100 ng each of GD1a and GD1b.

ELISA revealed an IgM antibody to GalNAc-GD1a and an IgG antibody to cytomegalovirus (CMV) in her serum. IgM antibody against GM2 was negative. An examination of cerebrospinal fluid (CSF) showed slightly increased proteins (0.42 g/L). An electrophysiological investigation showed low F-wave persistence in the right median nerve (25%). Needle electromyography was unremarkable. Brain and cervical magnetic resonance images (MRIs) were normal. The patient underwent a course of 5-day IVIg (400 mg/kg) because her symptoms

were suspected to be caused by an immune-mediated mechanism. Neuropathic symptoms improved during the regimen. On day 2 of the IVIg, muscle strength returned to the preillness state. The right arm pain subsided gradually, and 2 months later, she showed complete recovery (Fig. 1A).

2.2. Case 2

A previously healthy 26-year-old woman developed pain in both legs 2 weeks after an upper respiratory infection. Upon examination, muscle strength testing revealed mild weakness of both distal dominant legs (MRC, 4). Sensory examination showed hyperalgesia in her legs. Tendon reflexes were normal. The routine blood test results were normal. Serum antibody testing showed high titers of the anti-GalNAc-GD1a IgM antibody and the anti-CMV IgG antibody. IgM antibody against GM2 was negative. CSF examination was normal. An electrophysiological study showed decreased motor conduction velocity and F-wave persistence in the right ulnar nerve (50%). Brain and cervical MRIs were normal. Treatment with IVIg was effective for alleviating her pain, and muscle strength improved. On the 14th day after onset, she showed complete recovery (Fig. 1B).

2.3. Case 3

A previously healthy 54-year-old man presented with fever and developed left shoulder and forearm pain the next day. On day 2, he developed weakness in left arm elevation, and his pain continued to progress. Examination showed muscular weakness and atrophy in only the left deltoid muscle (MRC, 3). A sensory examination showed a moderate loss of vibration sensation and a mild loss of touch and pinprick sensation in the left upper extremity. Tendon reflexes were normal. Blood tests showed liver function disorder (AST, 137 IU/L;

ALT, 234 IU/L) and high titers of the anti-GalNAc-GD1a IgM antibody, anti-GM2 antibody, and anti-CMV IgG antibody. A CSF examination showed slightly increased proteins (0.48 g/L). An electrophysiological study showed low F-wave persistence in the left median nerve (45%). Brain, cervical, and thoracic MRIs were normal. The patient recovered completely within 30 days without any treatment.

2.4. Case 4

A 37-year-old man had a history of three operations to his right arm when he was 16, 25, and 28 years old. He had a 2 years and 5 months history of right neck pain that extended to the elbow. From the seventh month after the onset of the right arm pain, he had developed weakness and muscular atrophy in his right chest and arm. Although prednisolone was administered by his local doctor, his symptoms were aggravated gradually. After his pain and weakness spread to his right leg, he visited our hospital. An examination showed muscular weakness in the right neck extensor and the right upper and lower extremities. A sensory examination showed a mild loss of vibration, touch, and pinprick sensations in the right upper and lower extremities. Deep tendon reflexes were decreased in both upper and lower extremities. Laboratory studies were normal except for the positivity of the IgG anti-GalNAc-GD1a antibody and the anti-CMV IgG antibody. IgM antibody against GM2 was negative. CSF examination on admission showed slightly increased proteins (0.5 g/L). An electrophysiological investigation, including needle electromyography, showed no remarkable findings. Brain and cervical MRIs were normal. Because his clinical course was different from that of other 3 cases, it may be difficult to have a diagnosis of NA. However, the existence of anti-GalNAc-GD1a antibodies and neurological symptoms inexplicable by his history of operations allowed us to consider as immune-pathogenesis. The patient was treated with IVIg, which

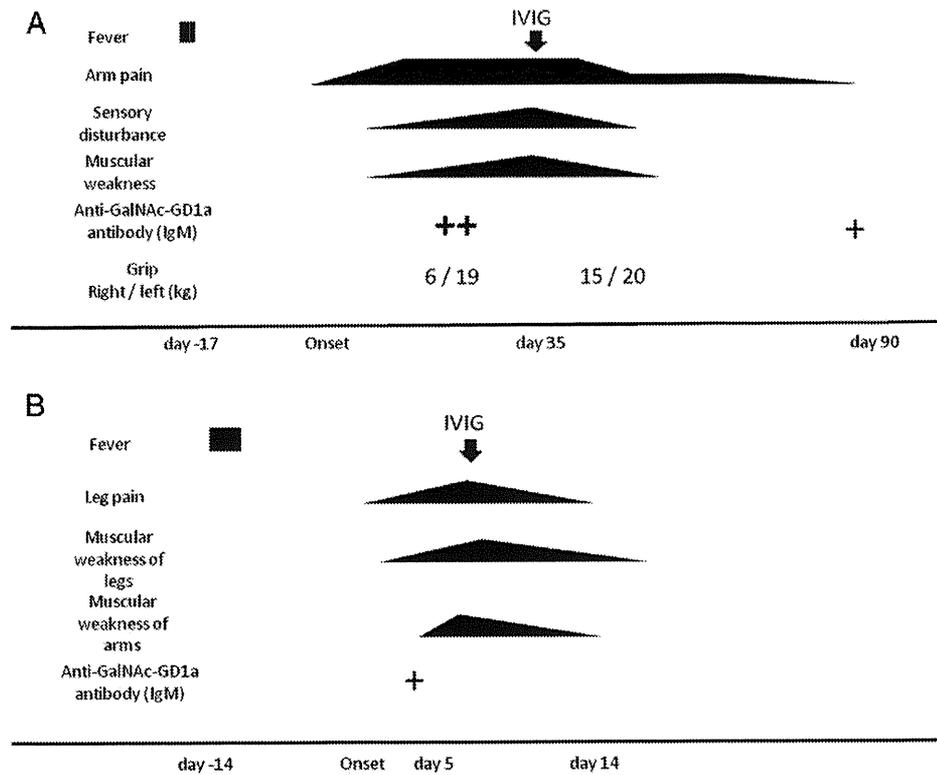


Fig. 1. The clinical course of Case 1 and Case 2. A) Case 1: The acute onset of arm pain and muscular weakness after a preceding infection is illustrated. The patient's neurological symptoms improved gradually after immunoglobulin infusion therapy (IVIg), and the titers of the serum anti-N-acetylgalactosaminyl GD1a (GalNAc-GD1a) antibody decreased. Her clinical course was considered to be monophasic. B) Case 2: The patient developed acute leg pain and muscular weakness of the extremities after an antecedent infection. Her symptoms improved immediately after IVIg.

Table 1
Summary of clinical features, serological findings, and treatment of neuralgic amyotrophy.

Patient	Age(y)/sex	Preceding infection	Clinical features	Nerve conduction study	Anti-ganglioside antibodies	Anti-CMV antibodies	Treatment	Outcome	Course
1	34/F	Fever and vomiting	Painful right arm neuropathy	Low F-wave persistence	IgM GalNac-GD1a	IgG	IVIg	Improvement	Acute monophasic
2	26/F	Upper airway inflammation	Painful bilateral leg neuropathy	Low F-wave persistence	IgM GalNac-GD1a	IgG	IVIg	Improvement	Acute monophasic
3	54/M	Fever	Painful left arm neuropathy	Low F-wave persistence	IgM GalNac-GD1a and GM2	IgG	Nothing	Spontaneous remission	Acute monophasic
4	37/M	Unknown	Painful right arm neuropathy	Normal	IgG GalNac-GD1a	IgG	IVIg	No change	Chronic

GalNac-GD1a, N-acetylgalactosaminyl GD1a; IVIg, immunoglobulin infusion therapy.

was complicated by aseptic meningitis. Thus, IVIg was performed for only 3 days. The aseptic meningitis was resolved immediately after treatment was discontinued. Pain and muscular weakness were not improved, but no further aggravation was observed.

3. Discussion

The etiology of NA remains to be clarified. In some cases, however, preceding events, the monophasic course, and the good response to immunomodulative therapy implicate an autoimmune mechanism. Pathologic findings with brachial plexus neuropathy in which conspicuous mononuclear inflammatory infiltrates are observed surrounding epineurial and endoneurial vessels within the brachial plexus support an immune-mediated mechanism (Suarez et al., 1996). However, NA in the presence of herpes zoster virus infection has been reported (Ohtake et al., 1991; Ismail et al., 2009), suggesting viral reactivation may also be a cause of NA. In addition, NA sometimes occurs on a hereditary basis and with genetic abnormalities (Tsao et al., 2004; Kuhlenbäumer et al., 2005). Thus, NA should be considered as a syndrome that is caused by a variety of etiologies.

Here, we reported the case of four patients with NA who were positive for anti-GalNac-GD1a antibody and who had a preceding infection. Three of the four patients had an antecedent infection. All patients had elevated titers of antibodies against GalNac-GD1a. Two of the four patients responded to IVIg. A complication forced discontinuation of IVIg in one patient, but his symptoms had stopped progressing before discontinuation. The clinical features, serological findings, and treatment of NA are summarized in Table 1.

GalNac-GD1a, which is a minor component of the gangliosides in the nervous system, was reported as a target antigen for the acute-phase serum antibody in GBS (Kusunoki et al., 1994). Further investigation showed that high titers of IgG antibodies to GalNac-GD1a were closely associated with the pure motor variant of GBS (Kaida et al., 2000). In contrast, IgM anti-GalNac-GD1a antibodies were grouped into two types. Those in the patients with antecedent CMV infection were frequently cross-reactive with GM2 and characterized by frequent facial and sensory deficits, whereas those in the patients with preceding gastrointestinal infections were specifically reactive with GalNac-GD1a and characterized by motor GBS (Kaida et al., 2001). In addition, anti-GalNac-GD1a antibodies were sometimes associated with multifocal motor neuropathy and chronic motor axonal neuropathy (Sugie et al., 1998; Kaji et al., 2000). The present investigation showed that some patients with NA exhibited anti-GalNac-GD1a antibodies with or without cross-reactivity to GM2. We could not detect evidence of recent CMV infection in serological markers or antibodies cross-reactive to GalNac-GD1a and GM2, however these findings might indicate that autoimmune mechanisms similar to GBS were involved in the pathogenesis of some cases of NA.

The pathogenic role of anti-GalNac-GD1a antibodies is still unclear. The location of GalNac-GD1a has been shown in the nodes of Ranvier, the paranodal regions, and in the small-diameter dorsal

root fibers (Kaida et al., 2003). Anti-GalNac-GD1a antibodies may function by binding to these sites and causing atrophy, weakness, and severe pain. Thus we hypothesized that anti-GalNac-GD1a antibodies could cause part of NA. However, the binding of anti-ganglioside antibodies is known to be affected by the local environment that surrounds the ganglioside antigens and the avidity of the antibodies (Kaida and Kusunoki, 2010).

Future investigations are necessary to elucidate whether the anti-GalNac-GD1a antibody is actually involved in the pathogenesis. In any case, anti-GalNac-GD1a antibodies could be useful markers in NA to predict response to immune therapy, especially IVIg.

Acknowledgments

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Human dental pulp-derived stem cells promote locomotor recovery after complete transection of the rat spinal cord by multiple neuro-regenerative mechanisms

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Spinal cord injury (SCI) often leads to persistent functional deficits due to loss of neurons and glia and to limited axonal regeneration after injury. Here we report that transplantation of human dental pulp stem cells into the completely transected adult rat spinal cord resulted in marked recovery of hind limb locomotor functions. Transplantation of human bone marrow stromal cells or skin-derived fibroblasts led to substantially less recovery of locomotor function. The human dental pulp stem cells exhibited three major neuroregenerative activities. First, they inhibited the SCI-induced apoptosis of neurons, astrocytes, and oligodendrocytes, which improved the preservation of neuronal filaments and myelin sheaths. Second, they promoted the regeneration of transected axons by directly inhibiting multiple axon growth inhibitors, including chondroitin sulfate proteoglycan and myelin-associated glycoprotein, via paracrine mechanisms. Last, they replaced lost cells by differentiating into mature oligodendrocytes under the extreme conditions of SCI. Our data demonstrate that tooth-derived stem cells may provide therapeutic benefits for treating SCI through both cell-autonomous and paracrine neuroregenerative activities.

Introduction

The development of effective treatments for spinal cord injury (SCI) has been stifled by this injury's complicated pathophysiology (1). During the acute phase, the focal mechanical insult disrupts tissue homeostasis. This triggers secondary injury processes in which multiple destructive cascades cause the necrotic and apoptotic death of neurons, astrocytes, and oligodendrocytes, which spreads beyond the initial injury site and leads to irreversible axonal damage and demyelination (2, 3). Subsequently, reactive astrocytes and oligodendrocytes near the site of injured spinal cord (SC) respectively produce chondroitin sulfate proteoglycans (CSPGs) and myelin proteins (including myelin-associated glycoprotein [MAG], Nogo, oligodendrocyte myelin glycoprotein [OMgp], netrin, semaphorin, and ephrin). These extracellular molecules function as axon growth inhibitors (AGIs), acting through the intracellular Rho GTPase signaling cascade (4). These multiple pathogenic signals synergistically accelerate the progressive deterioration after SCI. Therefore, therapeutic strategies for functional recovery from SCI must exert multifaceted reparative effects against a variety of pathogenesises (2).

Stem cell-based transplantation therapy holds great promise for establishing such a multifaceted therapeutic strategy. In the last decade, a variety of cell types, including human neural stem cells (5), embryonic stem cell derivatives (6–8), and adult bone marrow

stromal cells (BMSCs) (9, 10), have been transplanted into the injured SC of rats or mice, and their neuroregenerative activities evaluated. These preclinical studies showed that engrafted stem cells promote substantial functional recovery after SCI through both cell-autonomous/cell-replacement and paracrine/trophic effects (11). However, the previously tested stem cells show poor survival (6–8, 12) and/or differentiation under the severe conditions of SCI (9, 13, 14), and the transplantation of individual stem cells has led to only modest therapeutic benefits. Furthermore, although the trophic factors derived from these stem cells promote *in vitro* neurite extension and survival, their roles in the functional recovery of SCI are still largely unknown.

Human adult dental pulp stem cells (DPSCs) and stem cells from human exfoliated deciduous teeth (SHEDs) are self-renewing stem cells residing within the perivascular niche of the dental pulp (15). They are thought to originate from the cranial neural crest and express early markers for both mesenchymal and neuroectodermal stem cells (16, 17). Since naturally exfoliated deciduous and impacted adult wisdom teeth are not usually needed, DPSCs and SHEDs can be obtained without adverse health effects. Similar to BMSCs, these cells are able to differentiate into osteoblasts, chondrocytes, adipocytes, endothelial cells, and functionally active neurons *in vitro*, under defined conditions (16–19). Trophic factors expressed by them promote neuronal survival, proliferation, differentiation, and migration (20–23). Thus, these previous reports support the use of tooth-derived stem cells as a unique cellular resource for neuroregeneration therapies. However, their ability to promote functional recovery in neurological disorders remains largely unknown.

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Table 1
Flow cytometry of stem cells from humans

	SHEDs (n = 3)		DPSCs (n = 3)		BMSCs (n = 3)	
	Positive (%)	SD	Positive (%)	SD	Positive (%)	SD
MSC markers						
CD90	98.25	0.91	98.96	0.95	≥90	
CD73	91.45	8.44	96.60	2.14	≥90	
CD105	98.20	2.44	98.23	0.54	≥90	
Negative markers						
CD45	0.33	0.28	0.11	0.09	≤10	
CD34	0.36	0.32	0.07	0.03	≤10	
CD11b	0.02	0.02	0.03	0.02	≤10	
HLA-DR	0.45	0.39	0.12	0.10	≤10	
Neural markers						
DCX	95.42	0.66	84.45	0.45	91.37	8.20
Nestin	92.71	10.46	95.40	1.52	35.76	8.06
GFAP	92.93	8.30	97.50	3.54	4.49	3.11
βIII-Tubulin	99.69	0.21	85.43	0.77	99.24	0.73
NeuN	31.93	7.25	26.61	4.28	2.97	1.74
A2B5	94.84	3.72	96.34	0.33	35.47	15.07
CNPase	99.21	0.11	98.19	0.46	21.35	7.81
APC	0.20	0.01	0.36	0.02	2.75	2.05
MBP	0.68	0.04	0.32	0.02	3.02	2.00

Here we examined the neuroregenerative activities of DPSCs and SHEDs by transplanting them into a completely transected rat SCI model during the acute phase, in which axonal regeneration rather than axonal sprouting can be evaluated accurately. Our data show that these tooth-derived stem cells promoted functional recovery after SCI by multifaceted neuro-regenerative activities, via both cell-autonomous/cell replacement and paracrine/trophic mechanisms.

Results

Characterization of isolated human SHEDs and DPSCs for use in transplantation studies. Flow cytometry analysis showed that the SHEDs and DPSCs expressed a set of mesenchymal stem cell (MSC) markers (i.e., CD90, CD73, and CD105), but not endothelial/hematopoietic markers (i.e., CD34, CD45, CD11b/c, and HLA-DR) (Table 1). Like human BMSCs, both the SHEDs and DPSCs exhibited adipogenic, chondrogenic, and osteogenic differentiation as described previously (refs. 16, 17, and data not shown). The majority of SHEDs and DPSCs coexpressed several neural lineage markers: nestin (neural stem cells), doublecortin (DCX; neuronal progenitor cells), βIII-tubulin (early neuronal cells), NeuN (mature neurons), GFAP (neural stem cells and astrocytes), S-100 (Schwann cells), and A2B5 and CNPase (oligodendrocyte progenitor cells), but not adenomatous polyposis coli (APC) or myelin basic protein (MBP) (mature oligodendrocytes) (Figure 1A and Table 1). This expression profile was confirmed by immunohistochemical analyses (Figure 1B).

Next, we examined the expression of representative neurotrophic factors by real-time PCR. Both the SHEDs and DPSCs expressed glial cell-derived neurotrophic factor (*GDNF*), brain-derived neurotrophic factor (*BDNF*), and ciliary neurotrophic factor (*CNTF*) at more than 3 to 5 times the levels expressed by skin-derived fibroblasts or BMSCs (Figure 1C).

We further characterized the transcriptomes of SHEDs and BMSCs by cDNA microarray analysis. This gene expression analysis revealed a 2.0-fold difference in the expression of 3,318 of 41,078 genes between SHEDs and BMSCs. Of these, 1,718 genes were expressed at higher levels in the SHEDs and 1,593 genes were expressed at lower levels (data not shown). The top 30 genes showing higher expression in the SHEDs were in the following ontology categories: extracellular and cell surface region, cell proliferation, and tissue/embryonic development (Table 2).

SHEDs and DPSCs promoted locomotor recovery after SCI. To compare the neuroregenerative activities of human SHEDs and DPSCs with those of human BMSCs and human skin fibroblasts, we transplanted the cells into the completely transected SCs, as described in Methods, and evaluated locomotion recovery using the Basso, Beattie, Bresnahan locomotor rating scale (BBB scale) (24). Remarkably, the animals that received SHEDs or DPSCs exhibited a significantly higher BBB score during the entire observation period, compared

with BMSC-transplanted, fibroblast-transplanted, or PBS-injected control rats (Figure 2A). Importantly, their superior recoveries were evident soon after the operation, during the acute phase of SCI. After the recovery period (5 weeks after the operation), the rats that had received SHEDs were able to move 3 joints of hind limb coordinately and walk without weight support ($P < 0.01$; Supplemental Videos 1 and 2), while the BMSC- or fibroblast-transplanted rats exhibited only subtle movements of 1–2 joints. These results demonstrate that the transplantation of SHEDs or DPSCs during the acute phase of SCI significantly improved the recovery of hind limb locomotor function. Since the level of recovery was similar in the SHED- and DPSC-transplanted rats, we focused on the phenotypical examination of SHED-transplanted rats to elucidate how tooth-derived stem cells promoted the regeneration of the completely transected rat SC.

SHEDs regenerated the transected corticospinal tract and raphespinal serotonergic axons. To examine whether engrafted SHEDs affect the preservation of neurofilaments, we performed immunohistochemical analyses with an anti-neurofilament M (NF-M) mAb, 8 weeks after transection. Compared with the PBS-treated control SCs, the SHED-transplanted SCs exhibited greater preservation of NF-positive axons from 3 mm rostral to 3 mm caudal to the transected lesion site (Figure 2, B and C; asterisk indicates epicenter). The percentages of NF-positive axons in the epicenter of the SHED-transplanted and control SCs were $35.8\% \pm 13.0\%$ and $8.7\% \pm 3.4\%$, respectively, relative to sham-treated SCs (Figure 2D).

Regeneration of both the corticospinal tract (CST) and the descending serotonergic raphespinal axons is important for the recovery of hind limb locomotor function in rat SCI. We therefore examined whether these axons had extended beyond the epicenter in the SHED-transplanted SCs. The CST axons were traced with the anterograde tracer biotinylated dextran amine (BDA), which was injected into the sensorimotor cortex. The serotoner-

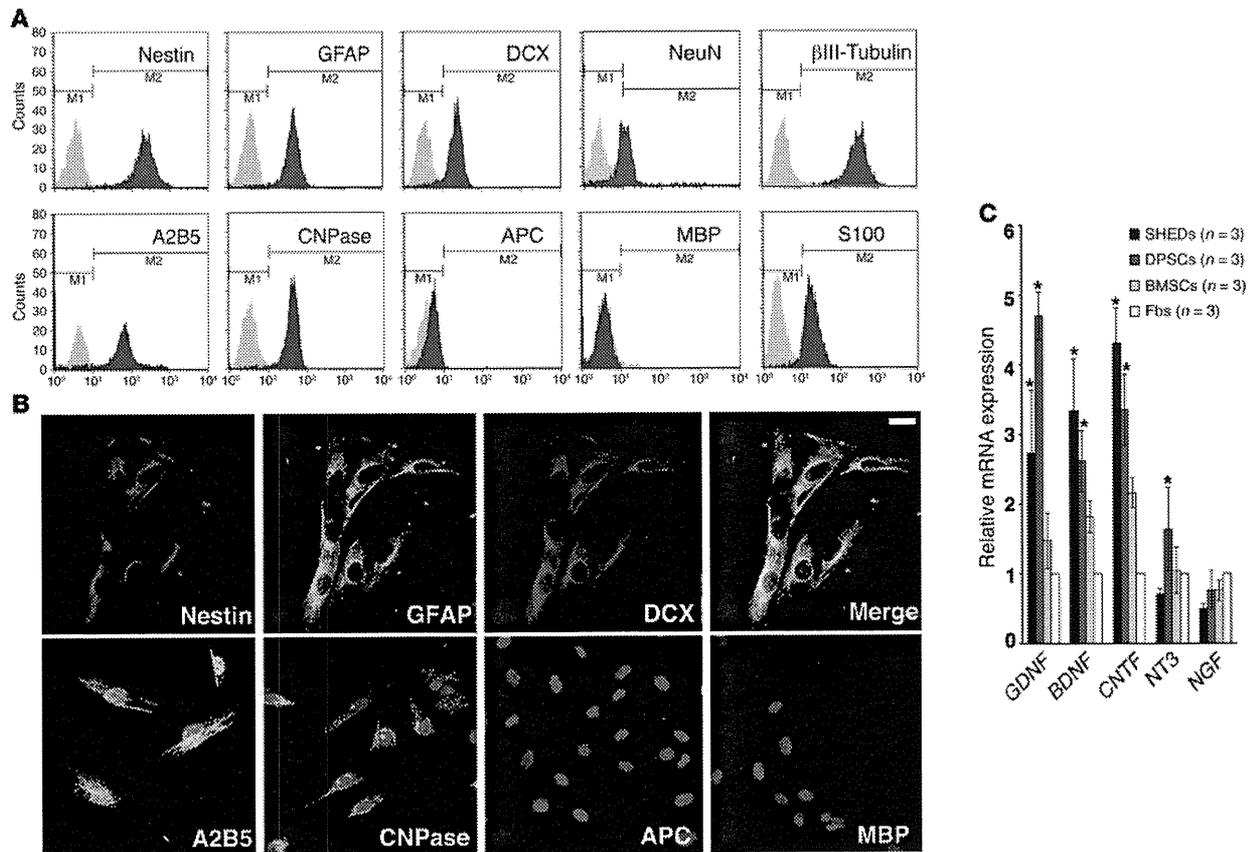


Figure 1

Characterization of the SHEDs and DPSCs used for transplantation. (A) Flow cytometry analysis of the neural cell lineage markers expressed in SHEDs. Note that most of the SHEDs and DPSCs coexpressed neural stem and multiple progenitor markers, but not mature oligodendrocytes (APC and MBP). (B) Confocal images showing SHEDs coexpressed nestin, GFAP, and DCX. SHEDs also expressed markers for oligodendrocyte progenitor cells (A2B5 and CNPase), but not for mature oligodendrocytes (APC and MBP). Scale bar: 10 μ m. (C) Real-time RT-PCR analysis of the expression of neurotrophic factors. Results are expressed as fold increase compared with the level expressed in skin fibroblasts. Data represent the average measurements for each cell type from 3 independent donors. This set of experiments was repeated twice and yielded similar results. Data represent the mean \pm SEM. * $P < 0.01$ compared with BMSCs and fibroblasts (Fbs).

gic raphespinal axons were immunohistochemically detected by a mAb that specifically reacts with serotonin (5-hydroxytryptamine [5-HT]), which is synthesized within the brain stem. We found that both BDA- and 5-HT-positive fibers extended as far as 3 mm caudal to the epicenter in the SHED-transplanted but not the control group (Figures 3 and 4). Furthermore, some BDA- and 5-HT-positive boutons could be seen apposed to neurons in the caudal stump (Figure 3D and Figure 4C), suggesting that the regenerated axons had established new neural connections. Notably, although the number of descending axons extending beyond the epicenter was small, we observed many of them penetrating the scar tissue of the rostral stump (Figure 3A and Figure 4A). The percentages of 5-HT-positive axons of the SHED-transplanted SCs at 1 and 3 mm rostral to the epicenter were $58.9\% \pm 3.9\%$ and $78.3\% \pm 7.4\%$ relative to sham-treated SC, respectively (Figure 4D). These results demonstrate that the engrafted SHEDs promoted the recovery of hind limb locomotion via the preservation and regeneration of transected axons, even in the microenvironment of the damaged CNS.

SHEDs inhibited the Rho GTPase activity induced by SC transection. The apparent axon regeneration in the SHED-transplanted SCs suggested that the SHEDs might modulate multiple AGI signals generated from oligodendrocytes and reactive astrocytes forming the glial scar. We therefore measured the activity level of Rho GTPase, which is an intracellular target of multiple AGIs, by pull-down assay. The injured SCs were isolated 7 days after transection and subjected to immunoprecipitation with GST-tagged Rho-binding domain (RBD). The level of active Rho (GTP-bound Rho [GTP-Rho]) in the transected control SCs increased; however, the engrafted SHEDs remarkably inhibited the activation of Rho (Figure 4E). These results strongly suggest that SHEDs promoted axon regeneration through the inhibition of multiple AGI signals.

Serum-free conditioned medium from both SHEDs and DPSCs antagonizes CSPG- or MAG-mediated neurite growth inhibition. Next, to analyze the roles of trophic mechanisms in the SHED-mediated axon regeneration, we examined whether the conditioned medium (CM) from SHEDs (SHED-CM) or DPSCs (DPSC-CM) could promote the neurite extension of cerebral granular neurons (CGNs)



Table 2
Functional gene classification in SHEDs versus BMSCs

Term	Changed gene up	Total gene	P
Extracellular region	343	2,865	2.52×10^{-14}
Skeletal system development	104	661	1.46×10^{-9}
Extracellular matrix	101	678	9.20×10^{-9}
Extracellular space	147	1,134	2.00×10^{-8}
Extracellular matrix organization	43	195	4.86×10^{-8}
Multicellular organismal development	643	6,683	9.36×10^{-8}
Collagen fibril organization	20	57	4.97×10^{-7}
Anatomical structure morphogenesis	346	3,339	9.52×10^{-7}
Mitotic cell cycle	146	1,184	1.11×10^{-6}
Proteinaceous extracellular matrix	82	578	1.36×10^{-6}
Organ morphogenesis	144	1,182	2.43×10^{-6}
Vasculature development	98	732	3.76×10^{-6}
Embryonic morphogenesis	96	728	7.04×10^{-6}
Cell proliferation	245	2,288	7.17×10^{-6}
Cell cycle	230	2,135	9.74×10^{-6}
Blood vessel development	93	707	1.31×10^{-5}
Response to wounding	191	1,738	2.02×10^{-5}
Receptor protein serine/threonine kinase signaling	56	369	2.12×10^{-5}
M phase of mitotic cell cycle	77	567	2.40×10^{-5}
Cell surface	86	671	3.26×10^{-5}
Organ development	362	3,675	3.68×10^{-5}
Collagen binding	21	90	3.90×10^{-5}
Glycosaminoglycan binding	42	262	4.65×10^{-5}
Mitotic spindle organization	12	33	7.15×10^{-5}
Cell adhesion	183	1,693	7.76×10^{-5}
Skeletal system morphogenesis	42	260	8.16×10^{-5}
Tissue development	185	1,720	8.76×10^{-5}
Cell surface receptor linked signaling pathway	368	3,785	8.98×10^{-5}
Mitosis	73	554	9.98×10^{-5}
Regulation of cell cycle	127	1,103	0.000109

on dishes coated with an AGI. CGNs isolated from newborn rats extended neurites on poly-L-lysine (PLL), but not on CSPG or MAG. Remarkably, both SHED-CM and DPSC-CM restored neurite extension activity of CGNs, while CM from fibroblasts (fibroblast-CM) or BMSCs (BMSC-CM) exhibited only subtle extension (Figure 5). Quantitative analysis showed that neurite extension through the inhibition of multiple AGIs was a unique characteristic of the tooth-derived stem cell (Figure 5, L and M). These results demonstrate that both SHEDs and DPSCs promote the regeneration of transected axons through direct inhibition of the multiple AGI signals by paracrine mechanisms.

SHEDs inhibited myelin degeneration. Next, we examined whether transplanted SHEDs preserved myelination in the transected SC by immunohistochemical staining with the fluorescent dye FluoroMyelin. In transverse sections of sham-operated SCs, white matter was clearly labeled by FluoroMyelin, and gray matter was not (data not shown). The control SCs exhibited little or no staining at the epicenter or 3 mm caudal to it (Figure 6, C and D). In contrast, we found significant FluoroMyelin-positive spots in the epicenter of the SHED-transplanted SCs, indicating that the SHEDs caused the regeneration of myelin structures in the transected region (Figure 6A). Notably, the myelin-positive areas of the SHED-transplanted SCs at 3 and 4 mm caudal to the epicenter constituted $55.3\% \pm 4.5\%$ and $78.0\% \pm 4.1\%$, respectively, of the

same areas in the sham-operated SCs, demonstrating that the SHEDs exerted remarkable myelin preservation activity (Figure 6E).

SHEDs survived and specifically differentiated into oligodendrocytes in the injured SC. In the FluoroMyelin-stained sections, we observed a myelin-expressing cell cluster in the gray matter of SHED-transplanted SCs (Figure 6B). We anticipated that these myelin-expressing cells would be mature oligodendrocytes derived from the transplanted SHEDs. To assess this possibility, we performed immunohistochemical analyses using anti-human nuclear antigen (HuNu) and two mature oligodendrocyte markers, APC and MBP (25, 26). Eight weeks after grafting, $32.3\% \pm 3.1\%$ of the transplanted SHEDs still survived in the injured SCs (data not shown). Of these cells, $86.2\% \pm 6.2\%$ and $90.2\% \pm 4.6\%$ expressed APC and MBP, respectively (Figure 7). In addition, 10% of the HuNu-positive cells were negative for MBP and APC, but their fate is currently unknown (data not shown). Before the transplantation, SHEDs expressed many early neural cell lineage markers (Figure 1 and Table 1). However, the surviving transplanted SHEDs did not express NF-M or GFAP (Figure 7), indicating that they specifically differentiated along the oligodendrocyte lineage in the injured SCs.

SHEDs inhibited neuronal and glial apoptosis after SCI. SCI-induced cell death is a major contributor to secondary injury, in which irreversible tissue damage spreads across the SC. Twenty-four hours after injury, at 1 mm caudal to the epicenter, most of the cells expressing NeuN, GFAP, or MBP were costained with TUNEL, showing that massive multicellular apoptosis occurred immediately after SCI (Figure 8). The engrafted SHEDs significantly decreased the TUNEL staining in all 3 of these lineages (Figure 8, C, D, G, H, K, and L): The total number of TUNEL-positive cells in the SHED-transplanted SCs was approximately 20% of that in the control SCs (Figure 8M). The percentages of TUNEL-positive cells in the control and SHED-transplanted SCs were $87.7\% \pm 3.1\%$ and $3.1\% \pm 3.2\%$, respectively (Figure 8N). These results demonstrate that the transplanted SHEDs minimized the expansion of secondary injury through strong neuroprotection of all the neural cell lineages.

Discussion

We report here the remarkable neuroregenerative activity of tooth-derived stem cells, SHEDs and DPSCs, for functional recovery after SCI. Previous studies have dealt with the differentiation characteristics of tooth-derived stem cells (16–19) and their trophic effects on the proliferation, migration, and survival of particular subsets of neurons (20–23). However, few studies have considered the therapeutic benefits of these stem cells for a particular neurological disorder. Our study revealed that engrafted SHEDs exhibited three major therapeutic benefits for recovery after SCI, including (a) inhibition of SCI-induced apoptosis of neurons, astrocytes, and oligodendrocytes, which promoted the preservation of neural fibers and myelin sheaths; (b) regeneration of the transected axon through the direct inhibition of multiple AGI signals, such as CSPGs and MAG,

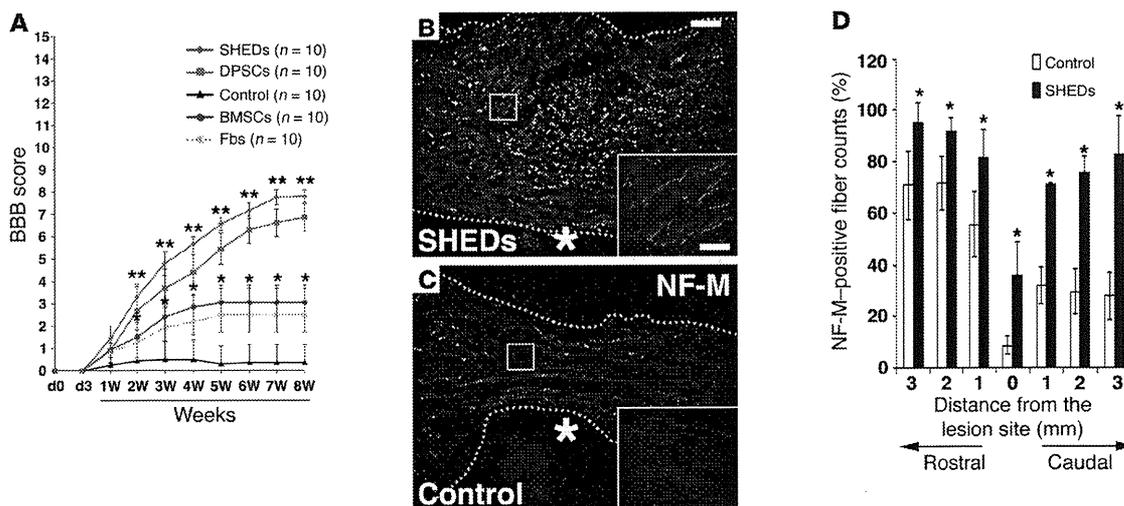


Figure 2 Engrafted SHEDs promote functional recovery of the completely transected SC. (A) Time course of functional recovery of hind limbs after complete transection of the SC. A total of 1×10^6 SHEDs, DPSCs, BMSCs, or fibroblasts were transplanted into the SCI immediately after transection. Data represent the mean \pm SEM. $**P < 0.001$, $*P < 0.01$ compared with SCI models injected with PBS. (B–D) Representative images (B and C) and quantification (D) of NF-M-positive nerve fibers in sagittal sections of a completely transected SC, at 8 weeks after SCI. Dashed lines outline the SC. Insets are magnified images of boxed areas in B and C. (D) Nerve fiber quantification, representing the average of 3 experiments performed under the same conditions. The x axis indicates specific locations along the rostrocaudal axis of the SC (3 mm rostral and caudal to the epicenter), and y axis indicates the percentage of NF-M-positive fibers compared with that of the sham-operated SCs at the ninth thoracic spinal vertebrate (Th9) level. Data represent the mean \pm SEM. $*P < 0.05$ compared with SCI models injected with PBS. Scale bars: 100 μ m and inset 20 μ m (B) and 50 μ m (C). Asterisks in B and C indicate the epicenter of the lesion.

by paracrine mechanisms; and (c) replacement of lost or damaged oligodendrocytes after SCI through specific differentiation into mature oligodendrocytes under the extreme conditions of SCI. To our knowledge, the latter two neuroregenerative activities (b and c) are unique to tooth-derived stem cells and are not exhibited by any other previously described stem cells. Thus, our data demonstrate that tooth-derived stem cells may provide significant therapeutic benefits for treating the acute phase of SCI through both cell-autonomous and paracrine/trophic regenerative activities.

Adult MSCs have been isolated from various tissues, including bone marrow, adipose tissue, skin, umbilical cord, and placenta (27–30). The therapeutic benefits of these stem cells have drawn intense attention in the field of translational medicine. Nevertheless, their biological equivalency/heterogeneity and identity are largely unknown (31). Tooth-derived stem cells exhibited BMSC-like multipotency and cell surface marker expression; however, they expressed a distinct set of multiple early neural lineage markers (Table 1 and Figure 1). A cDNA microarray gene expression analysis showed that the SHEDs expressed many genes in the categories of extracellular and cell surface region, cell proliferation, and tissue/embryonic development, at levels at least 2-fold higher than BMSCs (Table 2). These data indicate that tooth-derived stem cells belong to a highly proliferative ectomesenchymal stem cell-like population that actively communicates with neighboring cells. These characteristics raise the question of what the role of these stem cells is in tooth development and maintenance. Although we do not have a clear answer at present, future analyses using model animals such as dogs and pigs may clarify their precise origin and normal functions, as well as identifying the physiological system that maintains the “stemness” of these cells in vivo.

Both axon regeneration and the reformation of appropriate neuronal connections are prerequisites for functional recovery from SCI. However, multiple AGIs block the inherent regenerative activities of injured axons (2–4). It is well known that multiple AGIs constitute a remarkably intricate molecular network in the extracellular space of the injured CNS, in which they activate a common intracellular signaling mediator, Rho GTPase, and its effector, Rho-associated kinase (ROCK) (32–36). The activation of the Rho-ROCK cascade induces growth cone collapse and axonal repulsion (37). The inactivation of Rho by C3 transferase or of ROCK by Y-27632 downregulates AGI signaling and promotes functional recovery after SCI (38–40). Thus, Rho/ROCK signaling is an important target for SCI treatments; however, no reports have yet described an effect of stem cell transplantation on regulating the multiple AGIs/Rho/ROCK signaling cascades. We found that engrafted SHEDs promoted the regeneration of two major types of descending axons (CST and 5-HT) beyond the lesion epicenter and concomitantly inhibited the SCI-induced Rho activation (Figure 4). Furthermore, both SHED-CM and DPSC-CM promoted the neurite extension of CGNs cultured on two different AGIs, CSPG and MAG (Figure 5). Taken together, these results strongly suggest that tooth-derived stem cells promote the regeneration of transected axons through the direct inhibition of multiple AGI signals by paracrine mechanisms. Notably, in contrast to the CMs from tooth-derived stem cells, BMSC-CM showed only a subtle anti-AGI activity in the neurite extension assay, in good agreement with the level of functional recovery observed in BMSC-transplanted rats. Thus, the anti-AGI activity of tooth-derived stem cells is one of their major therapeutic benefits for the treatment of SCI.

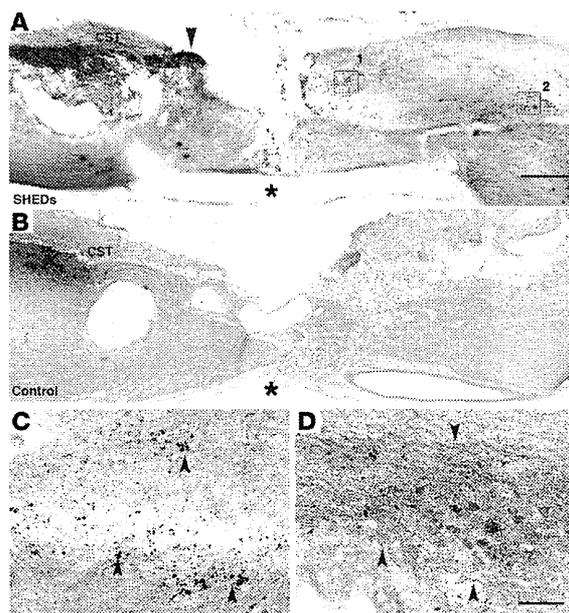


Figure 3

SHEDs regenerate CST fibers. Representative images of BDA-labeled CST axons. BDA-positive axons extended beyond the epicenter in the SHED-transplanted (A), but not the control SC (B). C and D are high-magnification views of boxed areas 1 and 2 in A, respectively. BDA-positive boutons were detected on the neurons of the caudal stump. Scale bars: 500 (A) and 100 μ m (D). Arrowhead in A indicates abundant penetration of CST axons into the scar tissue of the rostral stump. Arrowheads in C and D indicate regenerated CST axons extended beyond the epicenter. Asterisks in A and B indicate the epicenter of the lesion.

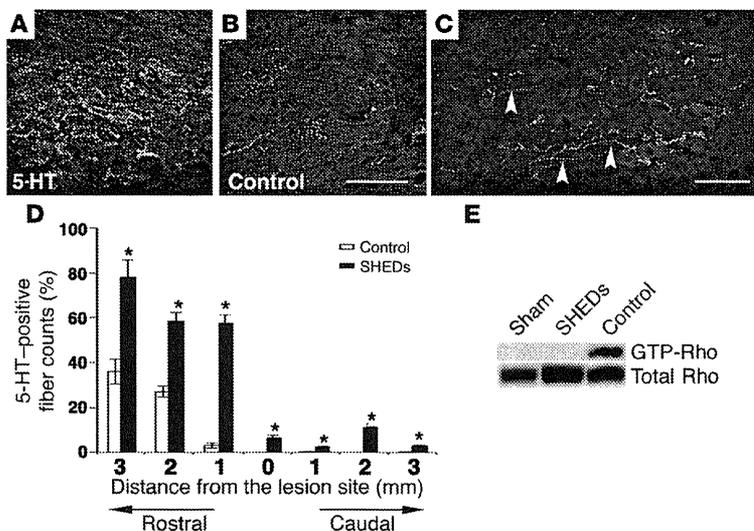
derived stem cells regulate multiple AGI activities to promote the regeneration of injured axons.

It has been shown that pharmacological blockade of neuron and/or oligodendrocyte apoptosis by erythropoietin (41, 42), inhibitors of purine receptor P2X7 (OxATP and PPADS) (43), a neutralizing Ab against CD95 (FAS) antigen (44), or minocycline (45, 46) promotes functional recovery after SCI. We found that engrafted SHEDs suppressed the apoptosis of both neurons and oligodendrocytes (Figure 8), which resulted in the remarkable preservation of neurofilaments and myelin sheaths in the region surrounding the epicenter (Figures 2 and 6). Notably, in addition to these two cell lineages, SHEDs strongly inhibited the apoptosis of astrocytes recruited to the lesion. In the classical view, reactive, CSPG-generating astrocytes are considered to be an obstacle to axon regeneration; however, recent genetic studies in mice have shown that the conditional ablation of astrocytes after SCI resulted in larger lesions, failure of blood brain barrier repair, expansion of the inflammatory response and tissue disruption, severe demyelination, and profound cell death of neurons and oligodendrocytes (47–51). Thus, the accumulated evidence demonstrates that, in addition to their anti-regenerative activity, astrocytes play an important role in the neuroprotection during the acute phase of SCI. We found that SHEDs suppressed the apoptosis of astrocytes and minimized secondary injury but inhibited AGI activity of CSPG derived from activated astrocytes. Thus, these results demonstrate that SHEDs promote the neuroprotective role but inhibit the anti-neuroregenerative activity of astrocytes to promote functional recovery after SCI.

The mechanisms that underlie the inhibition of multiple AGIs by SHED-CM and DPSC-CM are currently unknown. Although both SHEDs and DPSCs expressed an array of neurotrophic factors (Figure 1), our preliminary analysis showed that these trophic factors alone failed to promote the neurite extension of CGNs cultured on CSPG-coated dishes (K. Sakai and A. Yamamoto, unpublished observations). These results suggest that unknown factors, rather than neurotrophic factors, expressed by SHEDs and/or DPSCs may play major roles in the inhibition of multiple AGI signaling pathways. Since the strong anti-AGI activity was unique to the tooth-derived stem cells, but not to BMSCs, extracellular-related genes being preferentially expressed in SHEDs relative to BMSCs (Table 2) is a possible candidate anti-AGI factor. Future functional analysis of these genes will be required to reveal the molecular mechanisms by which tooth-

Figure 4

SHEDs regenerate 5-HT fibers and inhibit SCI-induced activation of Rho GTPase. (A–D) Representative images (A–C) and quantification (D) of serotonergic raphe axons stained with 5-HT mAb in the sagittal sections of the transected SC. A large number of 5-HT axons penetrated the scar tissue of the rostral stump in the SHED-transplanted SC (A), while only a few did in the control transected SC (B). (C) 5-HT-positive boutons were in contact with neurons of the caudal stump. Arrowhead indicates 5-HT-positive fiber extended beyond the epicenter. Quantification of regenerated 5-HT axons (D) was carried out as described in Figure 2D, except the y axis indicates the percentage of 5-HT axons compared with that in the sham-operated SC. Data represent the mean \pm SEM. * $P < 0.01$ compared with SCI models injected with PBS. Scale bars: 50 μ m (B and C). (E) SCI-induced Rho GTPase activation 7 days after SCI was suppressed by engrafted SHEDs. The level of active Rho in lysate from the samples indicated at the top (sham-operated, control, and SHED-transplanted) was examined by RBD pull-down assay.



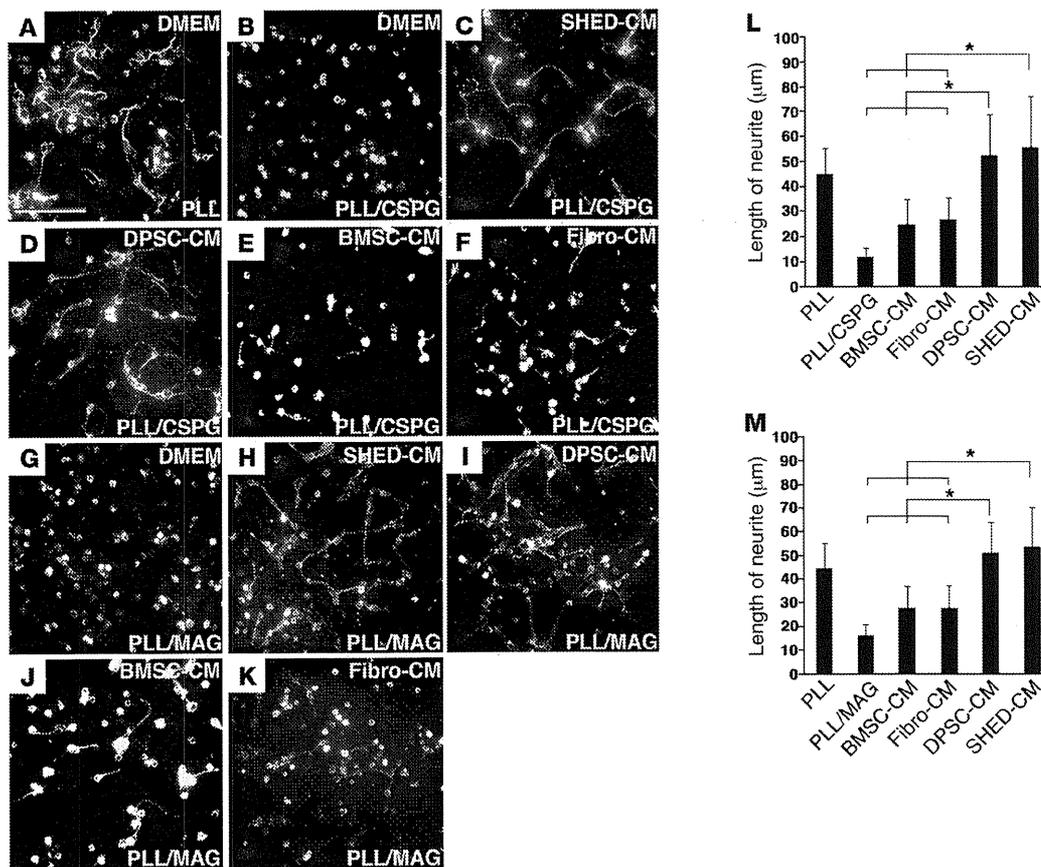


Figure 5

SHED-CM and DPSC-CM promote the neurite extension of CGNs on CSPG or MAG. CGNs were plated on PLL (A), PLL/CSPG (B–F), or PLL/MAG (G–K), with the CM indicated at the top of each panel. CGNs plated with SHED-CM or DPSC-CM extended their neurites on the CSPG- (C and D) and MAG-coated (H and I) dishes, while BMSC-CM and fibroblast-CM (Fibro-CM) elicited only marginal extension on CSPG (E and F) and MAG (J and K). Quantification of the neurite length of CGNs plated on CSPG (L) and MAG (M). The y axis indicates the neurite length. Data represent the average measurements for each cell type from 3 independent donors. This set of experiments was performed 3 times and yielded similar results. Error bars represent SD. * $P < 0.05$. Scale bar in A is 100 µm.

Our data revealed two major advantages of using SHEDs for cell replacement in SCI treatment. First, we observed good survival of the engrafted SHEDs: more than 30% of the engrafted SHEDs survived as a cell mass in the injured SC. A previous study reported that, although the experimental details differed from ours, the survival rate of human ES cell-derived oligodendrocytes or motor neurons, transplanted just after complete SC transection, is less than 1% (8). We speculate that the SHED-mediated minimization of secondary injury and/or the formation of cohesive cell clusters of engrafted SHEDs may be attributable to their excellent cell survival rate. Second, we observed that the engrafted SHEDs specifically differentiated toward mature oligodendrocytes, expressing APC and MBP. It has been shown that DPSCs and SHEDs differentiate in vitro toward functionally active neurons that express voltage-gated Na^+ channels and in vivo toward neuron-like cells 48 hours after their transplantation into the mesencephalon of avian embryos (18). Taken together with our findings, these results support the idea that tooth-derived stem cells exhibit neural stem cell-like characteristics and that unknown environmental cues are important for their fate determination. Since cell-based remy-

elination strategies can restore saltatory conduction and promote functional recovery after SCI (52), the SHED's strong cell survival and oligodendrocyte-specific differentiation potential, particularly under the extreme conditions of SCI, would be great advantages in using these cells to treat SCI. It is hoped that in the future, clarification of the regulatory cues for the specific differentiation of SHEDs will help us to establish efficient therapeutic protocols for SCI patients based on precise cell fate control.

The aim of this study was to address the neuroregenerative activity of tooth-derived stem cells in a particular CNS injury model, SCI. We used the rat complete transection model, because it provides good reproducibility and permits a more accurate assessment of the effects of treatment than do other SCI models. Although contusion and crush models would provide experimental conditions that are closer to the SCIs seen clinically in humans, the amount of injury in these models is not consistent from animal to animal. Furthermore, these incomplete transection models permit spontaneous recovery after SCI, and the residual SC tissues may provide routes for the compensatory sprouting of uninjured SC axons (53). Thus, the transection model was cho-

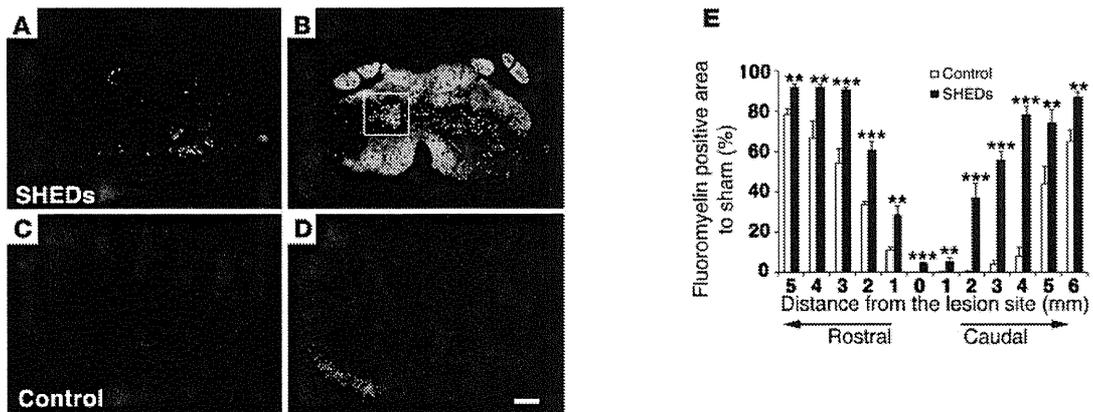


Figure 6

SHEDs preserve myelin sheath and differentiate into mature oligodendrocytes in the transected SC. Representative images (A–D) and quantification (E) of the myelinated area 8 weeks after SCI. Transverse sections of the epicenter (A and C) and 3 mm caudal to it (B and D) were stained with FluoroMyelin. The myelinated area in both regions was significantly preserved in the SHED-transplanted SC (A and B), but abolished in the control SC (C and D). Scale bar: 100 μ m (D). (E) Quantification of the myelinated area showing the average of 3 experiments performed in parallel. The x axis indicates specific locations along the rostrocaudal axis of the SC. The y axis indicates the percentage of the myelin-positive area compared with that of the sham-operated SC at the Th9 level. Error bars represent SD. ** $P < 0.01$, *** $P < 0.001$ compared with SCI models injected with PBS.

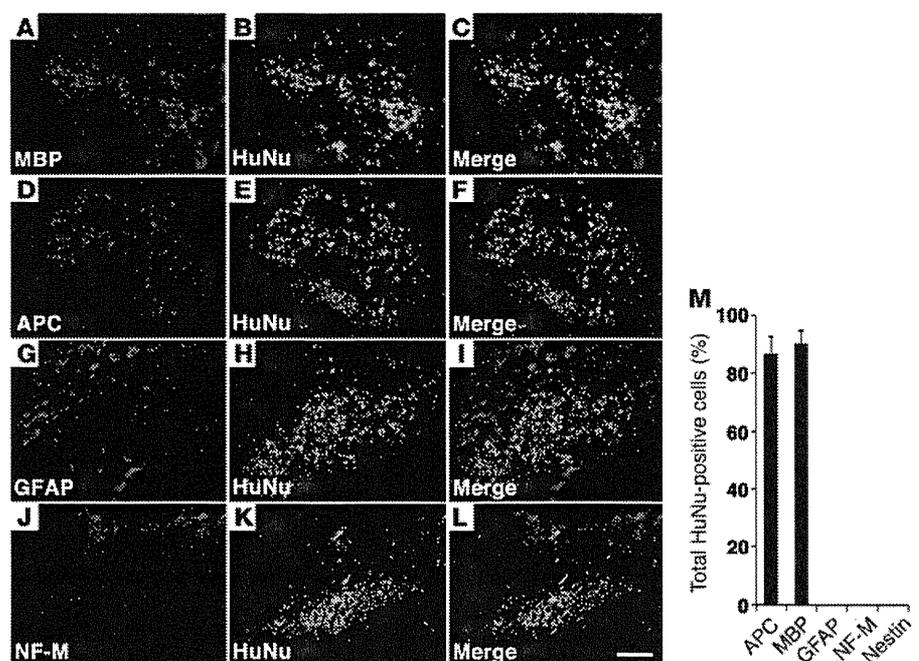
sen as most appropriate for the precise assessment of the axonal regeneration activity of tooth-derived stem cells.

In this study, we transplanted cells into the injured SC immediately after surgical transection, which is impractical for most human SCI cases. We chose this system to examine the therapeutic benefits of the transplanted cells in countering the multiple pathogenic signals that function synergistically during the early phase of SCI. Our future studies will analyze the neuroregenerative activities of tooth-derived stem cells in SCI under more clinically relevant experimental conditions.

In conclusion, we demonstrated multifaceted neuroregenerative activities of tooth-derived stem cells that fulfill many requirements for functional recovery after SCI. Not only did engrafted SHEDs have remarkable neuroregenerative activities, they also showed no malignant transformation 8 weeks after implantation (data not shown). Furthermore, SHEDs and DPSCs can be obtained from exfoliated deciduous and impacted adult wisdom teeth without adverse health effects. Thus, there are few ethical concerns regarding their clinical use. We propose that tooth-derived stem cells may be an excellent and practical cellular resource for the treatment of SCI.

Figure 7

SHEDs differentiate into mature oligodendrocytes in the transected SC. A myelin-positive cell cluster ectopically identified in the medulla of a SHED-transplanted SC (boxed area in Figure 6B) was characterized by immunohistochemical staining with an anti-human nuclei mAb (HuNu) together with Abs against neural cell lineage markers: anti-MBP (A–C), anti-APC (D–F), anti-GFAP (G–I), or anti-NF-M (J–L). The data indicate that SHEDs specifically differentiate into mature oligodendrocytes. The percentage of the lineage marker-positive to total HuNu-positive cell number (M) represents the average of 3 experiments performed in parallel. Error bars represent SD. Scale bar: 100 μ m (L).



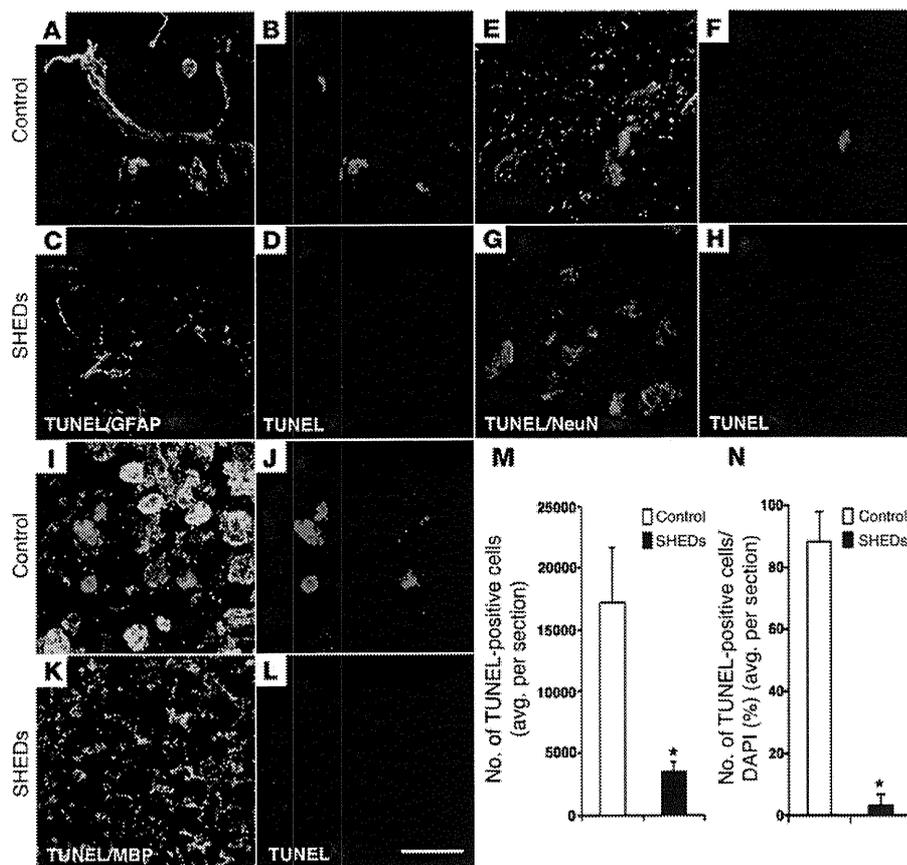


Figure 8

SHEDs suppress the apoptosis of neural cell lineages and secondary injury after SCI. Representative images (A–L) and quantifications (M and N) of apoptotic cell death 24 hours after SCI. Transverse sections 1 mm caudal to the epicenter of PBS-injected (A, B, E, F, I, and J) and SHED-transplanted SCs (C, D, G, H, K, and L) were stained with TUNEL and then subjected to immunohistochemical analysis with an anti-GFAP mAb (A–D), anti-NeuN mAb (E–H), or anti-MBP mAb (I–L). The engrafted SHEDs decreased the apoptotic cell death of all 3 neural cell lineages. (M) Quantification of the total TUNEL-positive cell number within 3 mm rostral and caudal to the epicenter shows the average of 3 experiments performed in parallel. (N) The percentage of TUNEL-positive relative to total DAPI-positive cell number in the same area as in M. Error bars represent SD. **P* < 0.01 compared with SCI models injected with PBS. Scale bar: 20 μm (L).

Methods

Isolation of SHEDs and DPSCs, and cell culture. Human SHEDs and DPSCs were isolated as described previously (16, 17). Briefly, exfoliated deciduous teeth (from individuals 6–12 years old) and adult third molars (18–30 years old) extracted for clinical purposes were collected. After separation of the crown and root, the dental pulp was isolated and then digested in a solution of 3 mg/ml collagenase type I and 4 mg/ml dispase for 1 hour at 37°C. Single-cell suspensions (1 × 10⁴ to 2 × 10⁴ cells/ml) were plated on culture dishes in DMEM supplemented with 10% fetal calf serum, then incubated at 37°C in 5% CO₂. Mesenchymal stem cells of three human bone marrow lines (hBMSCs, from individuals 20–22 years old) at passage 5 and three human skin-fibroblast lines (hFbs, 36–40-years old) at passage 5 were obtained from Lonza and the Health Science Research Resources Bank Japan, respectively.

Real-time PCR and microarray analysis. Total RNA was quantified by a spectrophotometer, and RNA integrity was checked on 1% agarose gels. RT reactions were carried out with Superscript III reverse transcriptase (Invitrogen) using 1 μg of total RNA in a 50 μl total reaction volume. Real-time PCR was performed using the THUNDERBIRD SYBR qPCR Mix (Toyobo) driven by the StepOnePlus Real-Time PCR System (Applied Biosystems). Primers were designed using DNADynamo (BlueTractorSoftware Ltd) and primer 3, as follows: *BDNF* forward (5'-GGGAAAAGGGAACAG-GAAAA-3'), *BDNF* reverse (5'-AACAGACAGGATGGGCAGAA-3'), *GDNF* forward (5'-CGAACTCTTGCCCTGACCT-3'), *GDNF* reverse (5'-ACAGC-CACGACATCCCATAAC-3'), *CNTF* forward (5'-CCTTCTCTTCTTCTT-GCTTTCTCTT-3'), *CNTF* reverse (5'-TGTCCTGCTCCACTCTCT-3'), *NT-3* forward (5'-TCAAACGGCAACTCTCCT-3'), *NT-3* reverse (5'-CTC-

GACAAGGCACACACACA-3'), *NGF* forward (5'-TTCCTTGACACT-GCCCTTC-3'), *NGF* reverse (5'-GATGATGACCGCTTGCTCCT-3'). Microarray experiments were carried out using a CodeLink Human Whole Genome Bioarray (Applied Microarrays Inc.) at Filgen Inc. The arrays were scanned using a GenePix4000B Array Scanner (Molecular Devices). The data were analyzed by using MicroArray Data Analysis Tool version 3.2 (Filgen Inc.) and deposited in the GEO database (accession GSE32403).

Flow cytometry analysis. For flow cytometry, 1 × 10⁶ cells were incubated with FITC-conjugated primary mAbs against CD34, CD45, and CD11b and PE-conjugated against HLA-DR, CD105, CD73, and CD90 (BD Biosciences) at 4°C for 30 minutes and then washed twice with PBS containing 0.1% bovine serum albumin. The expression of intracellular markers was examined by indirect immunostaining. Cells were fixed with 4% (w/v) PFA for 5 minutes and permeabilized with 0.1% (v/v) Triton X-100 in PBS for 5 minutes. After blocking with 10% (v/v) goat serum for 30 minutes, the cells were incubated with primary Abs: anti-GFAP (mouse IgG, 1:500, Millipore), anti-βIII-tubulin (mouse IgG, 1:1,000, R&D Systems), anti-NeuN (mouse IgG, 1:100, Millipore), anti-CNPase (mouse IgG, 1:500, Millipore), anti-nestin (rabbit IgG, 1:500, Millipore), anti-DCX (guinea pig IgG, 1:500, Millipore), anti-APC (rabbit IgG, 1:300, Abcam), anti-MBP (rabbit IgG, 1:500, Abcam), anti-A2B5 mAb (mouse IgG, 1:500, Millipore). The secondary Abs were anti-mouse IgG, anti-rabbit IgG, and anti-guinea pig IgG-conjugated with Alexa Fluor 448 (Invitrogen), used at 1:1,000. Cell fluorescence was evaluated by flow cytometry using a FACSCalibur (BD Biosciences).

Animal model and surgical procedure. Adult female Sprague-Dawley rats were anesthetized with a mixture of xylazine (100–150 mg/kg) and ketamine (60–90 mg/kg). After laminectomy at the 9th–11th thoracic vertebral lev-



els, the dura was opened, and the SC was completely transected using a surgical blade (Feather surgical blade stainless steel no. 11). The severed ends of the SCs typically retracted about 1–2 mm. The rostral and caudal stumps were lifted to ensure complete transection. Then, 1×10^6 cells were drawn into a glass pipette with a tip diameter of 50–70 μm mounted onto a 10- μl Hamilton syringe attached to a micromanipulator. First, the cells were deposited into two injection sites at the rostral and the caudal stumps, 2 mm from the lesion and 0.5 mm lateral to the midline, at a depth of 1.5 mm. A 2.5- μl sample containing 2.5×10^5 cells in PBS was grafted into each site (injection rate, 0.8 $\mu\text{l}/\text{min}$). Next, 1×10^5 cells in fibrin glue were implanted into the 1- to 2-mm gap to fill the lesion site in the severed SC. After surgery, the rats were placed in temperature- and humidity-controlled incubation chambers until they awoke. They were then transferred to cages, and bladder evacuation was applied daily. Antibiotics (sodium ampicillin, 10 mg/kg body weight) were injected into the rats daily for a week. The rats were maintained under postoperative care for 8 weeks. All rats were given cyclosporine (Novartis) at 10 mg/kg/d on the day before surgery transplantation, then every day after surgery.

Immunohistochemical analysis. Cells were plated on PLL-coated 8-chamber slides and then incubated with the primary Abs listed above. For histological examination of the treated SCs, the animals were anesthetized and transcardially perfused with 4% PFA in 0.1 M PBS, 8 weeks after transplantation. The SCs were embedded in OCT compound (Sakura Finetek) and sectioned in the sagittal or transverse plane at 20 μm on a cryostat (Leica). The sections were incubated with primary Abs against human nuclei (mouse IgG, 1:100), NF-M (rabbit IgG, 1:300, Millipore), and 5-HT (rabbit IgG, 1:500, Sigma-Aldrich) in addition to the Abs listed above. Secondary Abs were anti-mouse IgG–Alexa Fluor 488, anti-rabbit IgG–Alexa 546, and anti-guinea pig IgG–Alexa 647. Myelin was stained by FluoroMyelin green dye (Invitrogen), according to the manufacturer's instructions. After counterstaining with DAPI (Sigma-Aldrich), cell images were captured with a confocal laser scanning microscope (A1Rsi, Nikon), while tissue images were taken with a universal fluorescence microscope (BZ9000, Keyence).

The differentiation activity of the engrafted SHEDs and the cells staining positive for MBP, APC, NF-M, or GFAP among the anti-human nuclei-positive transplanted cells were quantified. Cells were counted in at least 15 confocal images from 3 individuals in parallel experiments, with error bars representing SD.

Anterograde neuronal tracing study. For tracing of the CSTs, 0.5 μl of 5% biotinylated dextran amine (BDA; MW 10,000, Molecular Probes, Invitrogen; 5% in PBS) was injected into 4 sites in the hind limb area of the sensorimotor cortex at a 1.2-mm depth, following the rat brain atlas (54). Two weeks after the injections, sagittal cryosections (20 μm thick) of the SCs were prepared and processed by diaminobenzidine (DAB) staining with the ABC reaction protocol (VECTASTAIN Elite ABC, Vector Laboratories).

BBB open field locomotor score. Hind limb neurobehavioral testing was performed using the BBB locomotor rating scale (24). The 22-point (from 0 to 21) BBB scale was used to assess hind limb locomotor recovery, including joint movements, stepping ability, coordination, and trunk stability. A score of 21 indicates unimpaired locomotion as observed in uninjured rats. Two examiners who were blinded to the animal's treatments performed the tests. The duration of each session was 4 minutes per rat. The scores were analyzed by repeated-measures ANOVA with Tukey's multiple comparison tests at each time point.

CM. At 70%–80% confluence, the cell culture medium was changed to serum-free DMEM. After 48 hours incubation at 37°C in 5% CO₂, the CM was collected and centrifuged for 4–5 minutes at 4°C, 22,140 g. After the brief re-centrifugation, the supernatant was collected and used as CM.

Neurite outgrowth assays. Forty-eight-well tissue culture plates (Falcon, BD) were coated with 20 $\mu\text{g}/\text{ml}$ PLL (Sigma-Aldrich) and then with 300 ng/ml

extracellular CSPG mixture (Millipore) or 400 ng/ml MAG/Fc Chimera (MAG; Sigma-Aldrich) for 4 hours at 37°C. Rat CGNs were seeded onto PLL-, PLL/CSPG-, or PLL/MAG-coated 48-well tissue culture plates at 2.0×10^4 cells/well and cultured at 37°C in 5% CO₂ with SHED-CM, DPSC-CM, BMSC-CM, or fibroblast-CM. After 24 hours incubation, cells were fixed in 4% paraformaldehyde/PBS and stained with anti-neuron-specific β III-tubulin (R&D Systems) to visualize neurites. Cell processes were defined as neurites when they were longer than the diameter of the cell body. Neurite length was evaluated by manually tracing neurite per cell using ImageJ software (version 1.29, <http://rsbweb.nih.gov/ij/>) and referenced to a known length. Each experiment was conducted in triplicate, and images were taken with 20 or more cells per field. For each experiment, at least 100 cells were randomly counted and measured.

Analysis of apoptosis. Apoptotic cell death was analyzed by TUNEL assay (In Situ Cell Death Detection kit, Roche). TUNEL-positive cells were counted on sections from sham-treated, PBS-injected, and SHED-transplanted animals. A researcher blinded to the experimental protocol determined the number of TUNEL-positive cells in the entire serial parasagittal section. The average number of TUNEL-positive cells per section was calculated from the values obtained by counting serial sagittal sections through the lesion site of each animal, with 3 animals examined per group.

Statistics. An unpaired 2-tailed Student's *t* test was used for single comparisons. For analysis of the real-time PCR results and open-field scores, we used repeated-measures ANOVA with Tukey's post hoc test (SPSS 19.0). A *P* value less than 0.05 was considered significant.

Study approval. The animal studies were carried out in accordance with the NIH Guidelines for the Care and Use of Laboratory Animals and approved by the Animal Research Committee of Nagoya University. Extracted teeth were collected at the Nagoya University School of Medicine, under approved guidelines set by Nagoya University (H-73, 2003). Ethical approval was obtained from the ethics committee of Nagoya University (permission number 8-2). All participants provided written informed consent.

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Premature Ligand-Receptor Interaction during Biosynthesis Limits the Production of Growth Factor Midkine and Its Receptor LDL Receptor-related Protein 1*

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Protein production within the secretory pathway is accomplished by complex but organized processes. Here, we demonstrate that the growth factor midkine interacts with LDL receptor-related protein 1 (LRP1) at high affinity (K_d value, 2.7 nM) not only at the cell surface but also within the secretory pathway during biosynthesis. The latter premature ligand-receptor interaction resulted in aggregate formation and consequently suppressed midkine secretion and LRP1 maturation. We utilized an endoplasmic reticulum (ER) retrieval signal and an LRP1 fragment, which strongly bound to midkine and the LRP1-specialized chaperone receptor-associated protein (RAP), to construct an ER trapper. The ER trapper efficiently trapped midkine and RAP and mimicked the premature ligand-receptor interaction, *i.e.* suppressed maturation of the ligand and receptor. The ER trapper also diminished the inhibitory function of LRP1 on platelet-derived growth factor-mediated cell migration. Complementary to these results, an increased expression of RAP was closely associated with midkine expression in human colorectal carcinomas (33 of 39 cases examined). Our results suggest that the premature ligand-receptor interaction plays a role in protein production within the secretory pathway.

Protein production within the secretory pathway is tightly regulated by complex but organized processes, which include folding in the endoplasmic reticulum (ER),² glycosylation, and vesicular transport to the cell surface through the ER and Golgi apparatus. Although these processes play important roles in quality control and maturation of proteins, it is not fully understood how production of proteins in the secretory pathway is regulated. This regulation mechanism is also important to understand some diseases, as disorganized production of

growth factors can be causative events in some pathological conditions, such as cancer (1). A fundamental question yet to be carefully addressed is whether a ligand and its receptor come into contact during their biosyntheses. To address this, we focused on two molecules in the present study, the growth factor midkine (MK) and its receptor low density lipoprotein (LDL) receptor-related protein 1 (LRP1).

MK is a heparin-binding growth factor, which was originally discovered as the product of a retinoic acid-responsive gene (2, 3). MK and pleiotrophin/heparin-binding growth-associated molecule form a family distinct from other heparin-binding growth factor families (4–6). MK plays important biological roles mainly in four areas, *i.e.* cancer, inflammation, neuro- and cardioprotection, and hypertension. To date, the biological functions in which MK has been implicated are all attributable to the secreted form of MK, and thus exogenous MK reverses phenotypes observed in MK-deficient mice. For example, MK-deficient mice are more resistant to vascular restenosis and hypertension associated with chronic kidney disease, whereas exogenous MK promotes these pathogenesis (7, 8). It is noteworthy that not only MK expression in carcinoma tissues but also the blood MK level is increased in cancer patients (9, 10). Both the MK expression level and MK blood level are closely related to clinical outcome (11–13). Furthermore, MK expression knockdown leads to tumor growth suppression (14, 15). These data suggest that the MK secreted by cancer cells plays a critical role in cancer progression.

LRP1 is a large endocytosis receptor belonging to the LDL receptor family (16). Because LRP1 recognizes many different ligands, the physiological roles of LRP1 govern a diverse array of biological functions, including metabolism of lipoproteins and proteinases, cellular entry of viruses and toxins, activation of lysosomal enzymes, and neurotransmission. Disruption of the LRP1 gene is embryonic-lethal, supporting the fundamental role of LRP1 in development (17). The receptor-associated protein (RAP) is a specialized ER chaperone for members of the LDL receptor family (18–20). Maturation of LRP1 is severely impaired in RAP-deficient mice, supporting an important role of RAP in the biosynthesis of LRP1 (20).

LRP1 is a receptor for MK and is required for MK-mediated cell survival (21, 22). Because MK and LRP1 are produced

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² The abbreviations used are: ER, endoplasmic reticulum; APP, amyloid precursor protein; LRP1, LDL receptor-related protein 1; MK, midkine; mLRP, mini-LRP; RAP, receptor-associated protein.