対側耳後部、腹部に針電極を刺入し、検査 側外耳道入口にスピーカーを設置し刺激音 を与え、検査側耳後部皮下と腹部皮下間で ABR を測定した。

ABR の測定には PowerLab system (ADInstruments CastleHill/Australia)を用いた。刺激発生装置は RP2.1and PA5 (Tucker-Davis Technologies FL/USA)、スピーカーはES1 spc(BioResearchCenter Nagoya/Japan)を使用した。刺激音は12000Hz の tone burst(0.2ms rise/fall time (cosine gate) and 1ms flat segment)とし、10dBSPL ごとに刺激音圧を変化させ、63dBSPL の刺激音で波形が明確に確認できないものを聴力障害ありと判定した。できるかぎり電磁波などの影響の少ない環境とするために、金網で囲まれた聴力検査室においてABRの測定をおこなった。

3.組織学的観察

感染モデルマウスは聴覚障害が確認できた時点で両側の蝸牛軸が評価できる面を含んだ側頭骨病理組織標本を作製し、HE 染色および免疫組織学的染色を行い、観察をおこなった。

(倫理面への配慮)

実験動物の取り扱いに関しては福島県立 医科大学動物実験ガイドラインに沿って、 愛護的に行った。

C. 研究結果

ウイルス投与量が 1.7 ×102pfu の脳内

接種モデルでは脳内接種後三週で聴力評価 を行ったところ21匹中6匹で両側聴覚障害 が出現し、5匹では片側聴覚障害を示した。 さらに 5 匹の片側聴覚障害をきたした感染 マウスの 4 匹はさらに一週間後 (脳内接種 4 週後) に両側聴覚障害をきたしていた。 脳内接種後三週で聴覚障害を認めていなか った10匹のうち4週間後に一側聴覚障害を きたしたものは5匹であった。脳内接種4 週間後に聴覚障害をきたさなかった 4 匹の うち脳内接種後 6 週までに聴覚障害をきた さなかったものは2匹おり、脳内接種を行 い 6 週まで飼育できたマウスは 19 匹中 17 匹(89.5%)において両側もしくは片側の聴 覚障害をきたした。コントロールとして同 じウイルス量を腹腔内投与したが、聴覚障 害をきたしたものは20匹中1匹であった。 本方法により確立されたCMV感染聴覚障 害マウスの組織学的な検討を行った。HE染 色による観察では形態学的な変化は軽度の 炎症細胞の浸潤を鼓室階、前庭階に認める 程度であった。明からなCMV感染巨細胞な どは確認できなかった。MCMV IE3抗体に よる染色ではラセン神経節、外リンパ腔、 髄膜に感染細胞の局在が確認された。

D. 考察

先天性ヒトサイトメガロウイルス感染症において、ウイルスに感染しただけでは聴覚障害をきたすとは限らず、どのような条件がそろうと聴覚障害が生じるのか解明されていない。聴覚障害を引き起こすメカニズムを解明する上で高率に聴覚障害をきた

すサイトメガロウイルス感染動物モデルの 作製が重要である。本邦において、モルモ ットを使用した垂直感染モルモットモデル 1)2)やマウスにおける感染モデル3)が聴覚障 害解明メカニズムのために作製され、側頭 骨内でのサイトメガロウイルスの局在等に ついて組織学的な検討がなされてきたが、 その聴覚障害がどの程度の比率で生じるの かは論じられていなかった。我々の作製し たウイルス感染モデルは高率に聴覚障害を きたすことが示された。今後本幹線モデル を用い、より詳細な感染部位の同定、聴覚 障害をきたすメカニズムの解析など、まだ 明らかにされていないサイトメガロウイル ス感染に伴う聴覚障害に関して有用な情報 を提供することが期待される。

E. 結論

生後24時間以内にマウスサイトメガロウイルスの脳内接種をおこなうことで、高率に聴覚障害モデルひきおこす動物モデルを作製した。接種6週後には80%以上のマウスにおいて聴性脳幹反応にて聴覚障害が確認された。光学顕微鏡による観察ではごく軽度の形態学的な変化が確認された。免疫組織学的な手法により、ラセン神経節に感染細胞が確認された。

F. 健康危険情報

なし

G. 研究発表

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2. 学会発表

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H. 知的財産権の出願・登録状況

なし

IV. 研究成果の刊行に関する一覧表

研究成果の刊行に関する一覧表

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IV. 研究成果の刊行物・別刷

2 Hearing Loss in Children with Congenital

3 Cytomegalovirus Infection

- 4 Satoshi Iwasaki and Shin-ich Usami
- 5 Additional information is available at the end of the chapter

7 1. Introduction

- 8 Sensorineural hearing loss (SNHL) is a common birth defect. The genetic origins of SNHL can
- 9 be identified in half of the prelingual cases; in the others, SNHL is caused by environmental
- $10\,$ $\,$ or unidentified genetic factors. The most common environmental cause of SNHL is congenital
- 11 cytomegalovirus (CMV) infection. CMV is also the most common cause of intrauterine and
- 12 congenital viral infection, affecting 0.5% to 2.5% of all live neonates [1]. While 90% of CMV-
- infected children are asymptomatic at birth, 10% of those exhibit clinically apparent sequelae
- at birth, including SNHL, mental retardation, motor disability, and microcephaly [1-4]. Recent
- studies have revealed that children with asymptomatic congenital CMV infection are at risk
- of late-onset SNHL and/or deterioration of SNHL during early childhood. These developments
- may not appear until months or even years following birth. The frequency of SNHL associated
- may not appear until months of even years following birth. The frequency of 514112 associated
- with asymptomatic congenital CMV infection reportedly ranges from 13% to 24% [5-9].

 Although asymptomatic CMV infection is associated with a lower incidence of SNHL than
- Although asymptomatic CMV infection is associated with a lower incidence of SNHL than symptomatic CMV infection, SNHL caused by congenital CMV often remains undiagnosed
- 21 because maternal screening for CMV infection is not routinely conducted and the detection of
- 22 SNHL during newborn hearing screening (NHS) tests is difficult [7, 10].
- 23 Hearing loss is detected in approximately 50% of children with symptomatic congenital
- 24 CMV infection. In 66% of these patients, hearing loss will deteriorate [3, 11]. Children
- 25 with symptomatic congenital CMV infection are easily identified at birth. In children with
- 26 symptomatic infection, intrauterine growth retardation and petechiae have been associat-
- 27 ed with the development of hearing loss [12]. SNHL is diagnosed in 7%-25% of children
- 28 with asymptomatic congenital CMV infection. Rates of delayed-onset SNHL, progressive
- 29 SNHL, and improvement of SNHL are reported to be 11%-18%, 23%-62%, and 23% -
- 30 47%, respectively [5-9].



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2 Manifestations of Cytomegalovirus Infection

- 1 Thus, the incidence of asymptomatic CMV infection and resulting SNHL may be higher,
- 2 making it the leading cause of SNHL in children. Treatment of children with congenital CMV
- 3 infection can prevent late-onset SNHL and/or deterioration of SNHL during early childhood.
- 4 Cochlear implantation is also effective for the development of speech perception and auditory
- 5 skills for deaf children with congenital CMV infection. Therefore, early identification of
- 6 congenital CMV infection is very important.

7 2. Epidemiology of hearing-impaired children with congenital CMV

8 infection

- 9 Of the 12,599 pregnant women included in a prospective study [13] conducted where from
- 10 June 1996 to December 2003, maternal ages were as follows: <20 years, 1.6%; 20–24 years, 14.7%;
- 11 25-29 years, 41.4%; 30-34 years, 28.6%; 35-39 years, 7.9%; and >40 years, 0.8%. The annual
- 12 seropositivity rate decreased over the 8-year study period, particularly during the last 4 years.
- 13 The seropositivity rate of CMV immunoglobulin G (IgG) antibody was 75.3% in the sample as
- 14 a whole. The seronegativity rate was 23.6%, and the percentage of cases borderline positive
- for IgG antibody was 1%. The seronegativity rate of CMV IgM antibody was 94.8% in the
- sample as a whole. The seropositivity rate was 2.2%, and 3% of cases were borderline positive
- 17 for CMV IgM antibody. During the study period, in the cases positive for IgM antibody (n =
- 18 146), borderline positive for IgM antibody (n = 73), and borderline positive for IgG antibody
- 19 (n = 14) and in cases with seroconversion of IgG antibody (n = 3), neonatal urine was analyzed
- 20 for CMV DNA. Seroconversion of CMV IgG antibody occurred in 0.32% of the 929 cases
- 21 negative for IgG antibody. Congenital CMV infection was identified in 18 infants by polymer-
- 22 ase chain reaction (PCR) analysis of urine. Follow-up was conducted in these cases.
- 23 The symptoms at birth and sequelae observed during the first 6 months of life in the 18 children
- with congenital CMV infection are shown in Table 1. Among these infants, 2 children (11.1%)
- 25 were symptomatic and the remaining 16 (88.9%) were asymptomatic. In this study, newborn
- 26 infants were considered symptomatic if central nervous system involvement such as micro-
- 27 cephaly or ventricular dilatation was detected. SNHL was detected in 1 child (50%) with
- 28 symptomatic infection and in 4 children (25%) with asymptomatic infection. Profound
- 29 unilateral SNHL had developed in the child with symptomatic infection. In the 4 children with
- 30 asymptomatic infection, the severity of SNHL varied from mild unilateral loss to profound
- 31 bilateral loss. Of the 4 children, unilateral SNHL was identified in 3 (75%). Mild unilateral
- 32 SNHL occurred in 2 children (66.7%), and profound unilateral loss occurred in 1 child (33.3%).
- Profound bilateral SNHL occurred in 1 child with asymptomatic infection. The unilateral hearing loss in Case 1 was detected by a neonatal automatic auditory brainstem response (ABR)
- hearing loss in Case 1 was detected by a neonatal automatic auditory brainstem response (ABR)
 screener. SNHL in the other 3 children was detected by conventional ABR. Table 2 shows a
- summary of the findings from longitudinal audiological evaluations in the 5 children with
- asymptomatic congenital CMV infection. On subsequent audiological testing, delayed-onset
- 38 SNHL was detected in 2 children who had passed the newborn hearing screening (NHS) test
- 39 (1 bilateral and 1 unilateral). Two cases (40%) had progressive hearing loss and 2 (40%) had

- improvement of hearing loss from the initial abnormal ABR (profound unilateral loss and
- profound bilateral loss, respectively).

	Symptoms	Audiologic examinations				
Case 1	Not found	Automatic ABR: unilateral REFER				
		ABR: unilateral moderate hearing loss				
Case 2	Not found	ABR: unilateral moderate hearing loss				
Case 3	Not found	ABR: unilateral profound hearing loss				
Case 4	Not found	ABR: bilateral severe hearing loss				
Case 5	Not found	Automatic ABR: bilateral PASS				
Case 6-16	Not found	ABR: normal				
Case 17	Microcephaly	ABR: unilateral profound hearing loss				
	Ventricular dilatation					
Case 18	Microcephaly	ABR: normal				
	Ventricular dilatation					
	Heart anomaly					

3 ABR: auditory brainstem response. This table is cited from reference [11].

Table 1. Initial symptoms and audiologic results during the first 6 months of life in 18 children with congenital CMV infection.

	Initial hearing loss	Results	of follow-up aud	Outcome		
		Age	Hearing loss	Characteristic		
Case 1	Unilateral moderate (Unilateral REFER)	36 mo	Bilateral profound	Delayed-onset Progressive	Cochlear implantation (39 mo)	
Case 2	Unilateral moderate	53 mo	Unilateral moderate	Fluctuating	Normal speech development	
Case 3	Unilateral profound	53 mo	Unilateral mild	Fluctuating Improvement	Normal speech development	
Case 4	Bilateral severe	17 mo	Normal	Fluctuating Improvement	Normal speech development	
Case 5	Normal (Bilateral PASS)	26 mo	Bilateral profound	Delayed-onset Progressive	Cochlear implantation (29 mo)	

6 SNHL: sensorineural hearing loss. This table is cited from reference [11].

Table 2. Results of longitudinal audiologic examinations in 5 children with SNHL caused by asymptomatic CMV

4 Manifestations of Cytomegalovirus Infection

- 1 In this prospective study, the rates of delayed-onset SNHL, progressive SNHL, and improve-
- 2 ment of SNHL were 12%, 40%, and 40%, respectively. Although a low rate of fetal CMV
- 3 infection was observed, the results of the present study regarding the rate of SNHL are in
- 4 accordance with the findings of those previous studies. The prevalence of congenital CMV
- 5 infection is affected by the socioeconomic and geographic differences, but it seems to be no
- differences on characteristics of hearing loss induced by congenital CMV infection.
- 7 Because they develop later, both delayed-onset and progressive hearing loss frequently remain
- 8 undiagnosed during universal newborn hearing screening (NHS) test [7, 10]. The 1994 Joint
- 9 Committee on Infant Hearing [14] pointed out that additional hearing evaluations after
- 10 universal NHS are required to detect delayed-onset hearing loss. Combined neonatal screening
- 11 for CMV infection and repeated auditory evaluation should be considered, particularly for
- 12 children with asymptomatic congenital CMV infection. Counseling of pregnant women on
- 13 prevention of CMV infection is also important.

14 2.1. Retrospective study of congenital CMV infection

- 15 Hearing loss in children with congenital CMV infection often presents at birth; however, in
- 16 many instances, it may develop after months or even years. One report stated that children
- 17 with normal hearing at 6 months of age develop hearing loss at a rate of approximately 1% per
- year; the cumulative risk of late-onset hearing loss is substantial (6.9%) in a population of in-
- 19 fants with asymptomatic congenital CMV infection [15]. Speech is often delayed in children
- 20 with bilateral hearing loss. For cases of severe bilateral SNHL, Ogawa et al. [16] reported that
- 21 congenital CMV infection could be diagnosed through the detection of CMV DNA in the dried
- 22 umbilical cord. In addition, genetic defects (particularly those related to GIB2) were identified
- 23 in 15% and 30% of the children, respectively. However, the etiology of pediatric SNHL, in-
- 24 cluding mild to moderate and unilateral SNHL, remains uncertain. In a study of congenital
- 25 CMV infection retrospectively diagnosed by the detection of CMV DNA extracted from dried
- 26 umbilical cord specimens, the prevalence of CMV in children with unilateral or bilateral
- 27 SNHL was investigated. In many of these cases, SNHL developed several months or even
- 28 years after birth.
- 29 In total, 134 patients (70 males and 64 females) with bilateral (n = 46; 34.3%) or unilateral (n =
- 30 88; 65.7%) SNHL were evaluated. These cases were referred to the Department of Otolaryng-
- 31 ology, Shinshu University School of Medicine from May 2008 to September 2009 (Table 3) [17].
- 32 The age of these children ranged from 1 month to 138 months (mean age: 37.7 ± 36.2 months).
- 33 In children with bilateral SNHL, both genetic testing for deafness and CMV DNA analysis
- 34 were performed. For children with unilateral SNHL, CMV DNA analysis and genetic testing
- 35 for gene mutations of GJB2, Mitochondrial1555 were performed. Objective audiometric
- 36 evaluation was performed for each patient using ABR and auditory steady-state evoked
- 37 response systems (MASTER 580-NAVPRO; NIHON KOHDEN Co., Ltd, Tokyo, Japan).
- 38 Behavioral audiological tests and/or pure-tone audiometry were also performed. Hearing
- 39 levels were classified into 2 categories on the basis of the severity of hearing loss in the worse
- 40 ear as severe (>70 dB) to profound (>90 dB) and mild (20-40 dB) to moderate (41-70 dB). Follow-
- 41 up hearing assessments were performed at intervals of 6–12 months. Progressive hearing loss

was defined as a decrease in hearing of ≥10 dB at 1 or more frequencies. Fluctuating hearing loss was defined as a decrease in hearing of >10 dB followed by an improvement of >10 dB at 1 or more frequencies. To analyze congenital CMV infection, CMV DNA quantitative PCR (qPCR) analysis was performed. Prior to qPCR analysis, total DNA, including genomic DNA and CMV DNA, was extracted from preserved dried umbilical cords. The results of this study revealed that in 9.0% (12/134) of children, SNHL could be attributed to congenital CMV infection. CMV DNA from preserved umbilical cords was detected in 8.7% (4/46) of children with bilateral SNHL and 9.1% (8/88) of those with unilateral SNHL. Congenital CMV infection caused bilateral severe-to-profound SNHL, bilateral mild-to-moderate SNHL, unilateral severe-to-profound SNHL, and unilateral mild-to-moderate SNHL in 14.3% (4/28), 0% (0/18), 9.6% (7/73), and 6.7% (1/15) of hearing-impaired children, respectively. This study also revealed that both congenital and late-onset SNHL could be caused by congenital CMV infection.

Hearing loss	Gender	Hearing level	Severe-pr	rofound HL	Mild-moderate HL		
	(n)	(dB)	n	Diagnostic age	n	Diagnostic age	
Total	M: 70, F: 64		101	34.4±34.7 mo	33	48.8±38.7 mo	
(N=134)			(75.4%)		(24.6%)		
Bilateral HL	M: 31, F: 15	71.8 dB [R]	28	16.6±19.9 mo	18	11.1±39.1 mo	
(N=46)		71.7 dB [L]	(20.9%)		(13.4%)		
Unilateral	M: 39, F: 49	89.5 dB (W)	72	41.2±36.6 mo	15	40.3±36.8 mo	
(N=88)		13.6 dB (B)	(54.5%)		(11.2%)		

HL: hearing loss. Diagnostic age: age diagnosed as hearing loss.

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15 M: male, F: female. R: right, L: left. B: better ear, W: worse ear. This table is cited from reference [16].

16 Table 3. Summary of characteristics of children with bilateral or unilateral hearing loss.

17 Table 4 shows the clinical characteristics of 12 children in whom CMV DNA was identified. 18 Of these 12 children, bilateral SNHL was detected in 4 and unilateral SNHL in 8. All 4 children 19 with bilateral SNHL had late-onset profound SNHL. Hearing fluctuation and PASS at the NHS 20 test were confirmed in 3 children (75%). Of the 8 children with unilateral SNHL, detectable 21 defects were confirmed in 2 children. Hearing fluctuation was detected in only 1 child (12.5%). 22 No inner ear anomaly was found in any of the 8 children with unilateral SNHL.

Retrospective diagnosis of congenital CMV infection is important to improve our understanding of the etiology of pediatric SNHL. In previous reports (Table 5), the frequency of congenital CMV infection in children with bilateral SNHL has varied from 3% to 36% because of variations in parameters (number of subjects, severity of SNHL) and methods [CMV IgM testing, DNA urinalysis, DNA from dried blood spots (DBS) in Guthrie cards] [19-24]. In 2 Japanese studies based on the retrospective diagnostic method of analysis of preserved dried umbilical cords, congenital CMV infection was detected in 10%-12% of children with bilateral SNHL [25, 26];

6 Manifestations of Cytomegalovirus Infection

however, these studies included few subjects (10–26 cases). In children with unilateral SNHL, CMV DNA from preserved umbilical cords was detected in 9.1% (8/88). The frequency of congenital CMV infection was similar in children with unilateral and bilateral SNHL. It has been speculated that approximately 10% of SNHL in children is caused by congenital CMV infection. Few reports have examined the frequency of congenital CMV infection using retrospective diagnostic methods in children with unilateral SNHL. However, using the CMV DNA detection method, 25% (1/4) [16] and 19% (8/42) [19] of children with unilateral SNHL were diagnosed with congenital CMV infection.

Case no.	Sex	Diagnostic age	Bilateral/ Unilateral	Severity	Average HL (R/L: dB)	Onset	NHS
1	F	60 mo	Bilateral	Profound	87.5/108.8	Late	Pass
2	F	52 mo	Bilateral	Profound	87.5/110.0	Late	Pass
3	М	50 mo	Bilateral	Profound	100.0/100.0	Late	Pass
4	М	62 mo	Bilateral	Profound	110.0/46.3	Likely late	-
5	М	6 mo	Unilateral	Profound	32.5/103.8	Congenital	Refer (L)
6	М	65 mo	Unilateral	Profound	107.5/17.5	Unknown	_
7	М	50 mo	Unilateral	Profound	6.3/100.0	Unknown	-
8	F	98 mo	Unilateral	Profound	110.0/15.0	Unknown	_
9	F	55 mo	Unilateral	Profound	15.0/92.5	Late	Pass
10	F	2 mo	Unilateral	Profound	90.0/18.3	Congenital	Refer (R)
11	М	80 mo	Unilateral	Severe	13.3/70.0	Unknown	-
12	F	44 mo	Unilateral	Moderate	15.0/58.3	Late	Pass

⁹ F: female, M: male. Mo: month. HL: hearing loss. R: right, L: left. NHS: newborn hearing screening. Diagnostic age: age diagnosed as hearing loss. This table is cited from reference [16].

12

2.2. Genetic hearing loss and congenital CMV infection

13 Genetic testing for deafness has become valuable for precise diagnosis of hearing loss. The 14 most frequently implicated gene in nonsyndromic hearing loss is GJB2, the most prevalent 15 gene responsible for congenital hearing loss worldwide. GJB2, SLC26A4, CDH23, and mito-16 chondrial 12s ribosomal RNA (rRNA) are the other major genes that cause hearing loss in 17 Japan. One study stated that genetic mutations were responsible for deafness in 40%-45% of 18 children with congenital hearing loss [27]. In our study [17], 10 gene mutations associated with 19 deafness (GJB2, n = 7; SLC26A4, n = 3) were identified in 21.7% (10/46) of children with bilateral 20 SNHL. In children with bilateral severe-to-profound SNHL, gene mutations causing deafness

¹¹ Table 4. Clinical data of CMV DNA-positive children

Reference	Year	Subjects	CMV positive rate			Diagnostic methods	Country
			Total	Bilateral	Unilateral		
Barbi et al. [19]	2003	> 40 dBHL	9/79	1/37 (2.7%)	8/42 (19%)	DBS, qPCR	Italy
Ogawa et al. [16]	2007	> 20dB, nonsyndromic	(11.4%)	9/63	1/4 (25%)	US, PCR	Japan
Samileh et al. [21]	2008	SNHL	10/67	(14.3%)	NR/20	Cerologic test	Iran
Stehel et al. [22]	2008	> 40 dBHL	(10.5%)	NR/75	NR	DNA from	USA
Walter et al. [43]	2008	NHS refer	33/95	16/256	NR	urine	UK
Mizuno et al. [44]	2008	unexplained SNHL	(34.7%)	(6%)	0	DSS, qPCR	Japan
Jakubikova et al.	2009	only bilateral	16/256	NR	0/16 (0%)	UC, qPCR	Slovak Re.
[20]	2009	> 60 dBHL, NHS refer	(6%)	3/45 (6.7%)	NR	Cerologic test	Belgium
Boudewyns et al.	2009	NHS refer, > 20 dB	8/35	4/55 (7.3%)	NR	DBS, qPCR	USA
[45]	2009	NHS refer	(22.9%)	NR	0 (0%)	DBS, qPCR	Japan
Choi et al. [18]	2010	> 70 dB, deaf school	3/45 (6.7%)	13/479	3/17	UC, qPCR	USA
Tagawa et al. [26]	2010	children	4/71 (5.6%)	(2.7%)	(17.6%)	DBS, qPCR	Japan
Kimani et al. [46]		NHS refer	4/55 (7.3%)	3/26	0	US, qPCR	
Adachi et al. [47]		NHS refer, >35dB,	13/479	(11.5%)			
		bilateral	(2.7%)	8/92 (8.8%)			
			3/26	13/77			
			(11.5%)	(17%)			
			11/109				
			(10.1%)				
			13/77				
			(17%)				

NR: not reported. NHS: newborn hearing screening. DBS: dried blood spot. UC: umbilical cord. qPCR: quantitative PCR. HL: hearing level. SNHL: sensorineural hearing loss. Re.: republic. This table is cited from reference [16].

40 **Table 5.** List of previous reports on children with congenital CMV nfection.

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and CMV DNA positivity were detected in 32.1% (9/28) and 14.3% (4/28) of patients, respectively [17]. The diagnostic rate has been concluded to be 46.4% (13/28). If analysis of CMV DNA from preserved dried umbilical cords could be combined with genetic testing for deafness, approximately 50% of cases of bilateral severe-to-profound hearing loss in children could be detected.

Congenital CMV infection is also often diagnosed by detecting CMV DNA in urine within the first 2 weeks of life and serological testing for CMV-specific IgM antibody from mother and child [28]. In recent years, the detection of CMV DNA by retrospective methods has been more valuable not only in diagnosing congenital CMV infection during later stages of life but also in identifying children at highest risk of late-onset and progressive SNHL. Some reports have stated that DBS stored on Guthrie cards has been used for the retrospective diagnosis of congenital CMV infections [18, 29]. Similarly, preserved umbilical cords have been recently

- Manifestations of Cytomegalovirus Infection
- used in Japan [25, 26, 30]. The sensitivity varies widely depending on the DNA extraction 1
- 2 method in the DBS case. Some investigators have reported sensitivities of 71%-100% and
- 3 specificities of 99%-100% [19, 29]. In this study, the qPCR method and preserved umbilical
- cords were used because they were useful for more accurate detection of CMV DNA.

3. Diagnosis of congenital CMV infection 5

6 3.1. Detection methods

- 7 The gold standard for diagnosis of congenital CMV infection is isolation of the virus from urine
- 8 or saliva in the first 2 weeks of life. However, asymptomatic congenital CMV infection in
- 9 children who develop SNHL after the first 2 weeks following birth cannot be diagnosed on the
- 10 basis of viral isolation from urine or saliva. Detection of CMV DNA in infant blood or the
- 11 umbilical cord using PCR assays is a more feasible method for identifying children with late-
- 12 onset SNHL. The method involves analysis of blood stored as DBS on Guthrie cards. In
- 13 Japanese culture, the dried umbilical cord is generally stored at home as a memento of the
- 14 birth. These specimens are suitable for retrospective diagnosis of congenital CMV infection.
- 15 The sensitivity varied widely depending on the DNA extraction method from DBS on Guthrie
- 16 cards. Some investigators reported sensitivities of 71-100% and specificities of 99-100% [19,
- 17 29]. The qPCR method and dried umbilical cord could be useful for more precise detection of
- 18 CMV DNA.

19 3.2. Serological method

- 20 Diagnosis of symptomatic CMV infection is easier in children who display cognitive or
- 21 neuromuscular abnormalities than in asymptomatic children with CMV infection. Without
- 22 neonatal viral screening, the prevalence of SNHL caused by asymptomatic CMV infection
- 23 remains undetermined. To diagnose primary CMV infection, a serological method has been
- 24 used [31]. Pregnant women who test positive for CMV IgG seroconversion or CMV IgM
- antibody may transmit the virus to the fetus. Production of IgM antibody persists for 6-9 25
- 26 months [28]; therefore, a CMV IgM-positive result alone does not accurately predict the risk
- 27 of fetal infection.

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28 3.3. Detection of CMV DNA from umbilical cord

- 29 For the detection of congenital CMV infection, CMV DNA qPCR analysis was performed.
- 30 Prior to qPCR analysis, total DNA, including genomic DNA and CMV DNA, was extract-
- 31 ed from preserved dried umbilical cords. The procedure is as follows. Each 5-mm tissue
- section was incubated in a lysis buffer containing proteinase K and incubated overnight 33 at 56°C. Total DNA was extracted using the DNeasy® Blood & Tissue Kit (Qiagen
- 34 GmbH, Hilden, Germany), according to the manufacturer's instructions. The total amount
- 35 of DNA was measured using the Qubit® Fluorometer with Quant-iT™ dsDNA BR Assay
- Kit (Life technologies-Invitrogen, Carlsbad, CA, USA). Total DNA (10 pg) was analyzed

- 1 using the Step One Real-Time PCR System (Applied Biosystems, Foster City, CA, USA)
- 2 and TagMan® Universal Master Mix II (Applied Biosystems). The qPCR primers and
- 3 TaqMan® probe used for CMV DNA qPCR analysis were as follows: US14-1F: 5'-
- 4 ACGTCCACGTTAGGATGAGG-3', US14-1R: 5'-GTATGTGGCGCTTCTCTCGT-3', and
- 5 US14-1 TagMan probe: 5'-FAM- AACCTGTGCACCACAGCGCC -TAMRA-3'. To quantify
- 6 the input DNA amount in each sample, qPCR of each genomic region was also per-
- 7 formed using the following primers and TaqMan® probe: GJB2-2F: 5'-ACGTCCACGT-
- 8 TAGGATGAGG-3', GJB2-2: 5'-GTATGTGGCGCTTCTCTCGT-3', and GJB2-2 TaqMan
- 9 probe: 5'-FAM- AACCTGTGCACCACAGCGCC -TAMRA-3'. The initial preheating steps
- 10 were performed for 2 min at 50°C and 10 min at 95°C. Following this, qPCR was per-
- 11 formed for 43 cycles of 15 s at 95°C and 60 s at 60°C. After qPCR analysis, relative CMV
- 12 concentrations in each sample were evaluated as ΔCt (delta cycle threshold), which was
- 13 calculated by determining the threshold cycle of CMV qPCR minus that of GJB2 qPCR.
- 14 The invader assay described by Abe [32] was used for genetic testing for deafness.

4. Treatment for hearing loss induced by

is important for successful speech perception [34].

16 congenital CMV infection

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17 4.1. Cochlear implantation in children deafened by symptomatic CMV infection

18 Cochlear implantation for the correction of congenital deafness is an effective way to ensure 19 the development of speech recognition. Cochlear implantation in children deafened by 20 symptomatic CMV infection has been reported [33, 34]. The prognosis of children with 21 symptomatic CMV infection is worse than that of those with asymptomatic CMV infection 22 with regard to cognitive and neurological development. It has been suggested that cochlear 23 implantation should be contraindicated for infants with symptomatic CMV infection and 24 deafness because they are less likely to develop spoken language [35]. In contrast, other reports 25 [33, 34] have suggested that cochlear implantation may improve quality of life, even if progress 26 is slower or lesser than that expected in congenitally deaf children not infected with CMV. 27 Pyman et al. [35] suggested that the prognosis in terms of linguistic outcome after cochlear 28 implantation is poorer for CMV-infected deaf children than for other congenitally deaf 29 children because of coexisting central disorders. Wide variation in speech perception and 30 intelligibility after cochlear implantation has also been reported in children deafened by 31 symptomatic CMV infection [33]. In that report, poor development in these areas was observed 32 in 50% of children with symptomatic CMV infection, whereas development similar to that in 33 congenitally deaf children not infected with CMV was evident in 31% of children and devel-34 opment better than that in noninfected congenitally deaf children was evident in 19% of 35 children. In addition, a recent study has shown that deafness caused by symptomatic congen-36 ital CMV infection associated with motor and cognitive delays is not a contraindication for 37 cochlear implantation. Early diagnosis of hearing loss and subsequent cochlear implantation

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1 4.2. Cochlear implantation in children deafened by asymptomatic CMV infection

The effectiveness of cochlear implantation in children deafened as a result of symptomatic congenital CMV infection has been evaluated by various groups, but there are only limited outcome data for deaf children with asymptomatic CMV infection. Children with asymptomatic congenital CMV infection have a better prognosis than symptomatic children, but it is difficult to evaluate the SNHL because children with asymptomatic congenital CMV infection are at risk of development of delayed onset SNHL and progressive SNHL. As a result, they are also at risk of late-onset learning difficulties and/or progressive learning difficulties.

9 A prospective study was conducted on deaf children with asymptomatic CMV infection to 10 assess the development of speech perception and auditory skills. This study examined 2 deaf 11 infants before and after cochlear implantation using the Infant/Toddler Meaningful Auditory 12 Integration Scale (IT-MAIS) [36]. Vocalization behavior in Case 1 was observed 6 months after 13 implementation and showed slow improvement but finally overtook after 36 months. After 3 months of cochlear implant use, the 2 children responded to speech and environmental sounds 15 in everyday situations and interpreted sounds in a meaningful way. They continued to 16 improve at 36 months postoperatively. IT-MAIS scores in these 2 children were similar to the 17 mean scores in the 5 congenitally deaf children without CMV infection. No difference was 18 observed in the effect of early cochlear implantation for deafness induced by CMV infection 19 between the groups of children. Another group reported that significant improvement in 20 auditory and language skills could be achieved in cochlear implanted children with asymp-21 tomatic CMV infection, but they did not achieve the same levels of outcome as congenitally 22 deaf children without CMV infection [37]. They found a wide variation in the outcome of 23 cochlear implantation in these children and speculated that the variation is related to the 24 degree of cognitive impairment. There are only a few studies available on outcomes of cochlear 25 implanted children with asymptomatic CMV infection. Therefore, more studies will be needed 26 to evaluate the effectiveness of cochlear implantation in these children.

4.3. Treatment for hearing-impaired children with congenital CMV infection

28 To prevent late-onset and/or deterioration of SNHL, treatment with intravenous ganciclovir 29 (GCV) and/or oral valganciclovir (VGCV) has been recommended in children with sympto-30 matic congenital CMV disease involving the central nervous system [38-41]. In previous 31 reports, treatment with intravenous GCV was initiated within the first 10-14 days of life for 32 2-6 weeks, and GCV doses ranged from 5 to 12 mg/kg twice daily. One report revealed that 33 in 5 of 9 children with congenital CMV infection and SNHL, treatment with intravenous GCV 34 induced improvement of SNHL in 2 children and prevented deterioration of SNHL in 5 35 children [38]. Another report revealed that in 4 of 6 children with congenital CMV infection and SNHL, treatment with intravenous GCV induced improvement of SNHL in 2 children and 37 no deterioration of SNHL in 4 children during the 21-month observation period [39]. Im-38 provement of SNHL or maintenance of normal hearing was reported in 84% of children treated 39 with intravenous GCV and 59% of untreated children. Deterioration of SNHL was reported in 40 21% of treated children and 68% of untreated children [40]. According to these reports, good 41 results have been observed in the group of children treated with GCV. Treatment with

- intravenous GCV and oral VGCV can prevent the development of SNHL during an 18-month
- 2 administration period [41]. Treatment with intravenous GCV has been investigated in hearing-
- 3 impaired children with asymptomatic congenital CMV infection. No SNHL was found for 4 –
- 4 11 years in 12 children with asymptomatic congenital CMV infection treated with intravenous
- 5 GCV, but SNHL developed in 2 of 11 untreated children [42]. Unfortunately there is no
- evidence for the efficacy of longer treatment with oral VGCV.

5. Conclusion 7

- 8 Congenital CMV infection is a major cause of bilateral and unilateral SNHL in children. In
- 9 total, 9.0% of SNHL cases of unknown causes (bilateral SNHL: 8.7%, unilateral SNHL: 9.1%)
- 10 are attributed to congenital CMV infection. Screening tests such as the detection of CMV DNA
- 11 from preserved dried umbilical cords and genetic testing are important for the detection of
- 12 SNHL in children. Using this combined methodology, detection of the cause of SNHL is
- 13 possible in approximately 50% of children with hearing loss.
- 14 Cochlear implantation is effective to ensure the development of speech perception and
- 15 auditory skills in deaf children with asymptomatic congenital CMV infection. No significant
- 16 difference in growth of meaningful auditory integration was observed between the overall
- 17 pediatric cochlear implant population not infected with CMV and that with asymptomatic
- 18 CMV infection. Implementation of CMV screening models is important to prevent late-onset
- 19 SNHL and deterioration of hearing loss.

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