

# Localisation and distribution of *O*-acetylated *N*-acetylneuraminic acids, the endogenous substrates of the hemagglutinin-esterases of murine coronaviruses, in mouse tissue

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**Abstract** Infections by mouse hepatitis viruses result in disease of the liver, the gastrointestinal tract, respiratory tract, and the central nervous system. Coronaviruses related to mouse hepatitis virus express a hemagglutinin-esterase surface glycoprotein, which specifically hydrolyses either 5-*N*-acetyl-4-*O*-acetyl neuraminic acid (Neu4,5Ac<sub>2</sub>) or 5-*N*-acetyl-9-*O*-acetyl neuraminic acid (Neu5,9Ac<sub>2</sub>). Moreover, these sialic acids represent potential cellular receptor determinants for murine coronaviruses. Until now, the distribution of these sialic acids in mouse brain was not thoroughly investigated. Particularly Neu4,5Ac<sub>2</sub> was not yet found in mouse brain. Using a sensitive method of gas chromatography coupled to mass spectrometry in the electron impact mode of ionization this manuscript demonstrates the occurrence of 13 different sialic acids varying in their alkyl and acyl substituents in mouse tissues including 5-*N*-acetyl-4-*O*-acetyl-9-*O*-lactyl-neuraminic acid (Neu4,5Ac<sub>2</sub>9Lt), 5-*N*-acetyl-9-

*O*-lactyl-neuraminic acid (Neu5Ac9Lt), 5-*N*-acetyl-8-*O*-methyl-neuraminic acid (Neu5Ac8Me) and the 1,7-lactone (Neu5Ac1,7L) of neuraminic acid. Neu4,5Ac<sub>2</sub>, relatively abundant in the gut, was present as a minor compound in all tissues, including liver, olfactory lobe, telencephalon, metencephalon and hippocampus. Neu5,9Ac<sub>2</sub> was also found in these tissues, except in the liver. It is suggested that these sialic acids represent the endogenous substrate and receptor determinants for murine coronaviruses.

**Keywords** Gas chromatography · Mass-spectrometry · Mouse hepatitis virus · Mouse tissues · 5-*N*-acetyl-4-*O*-acetylneuraminic acid · Sialic acid

## Abbreviations

amu	atomic mass unit
EI	electron impact ionization
GC	gas chromatography
MS	mass-spectrometry
HFB	heptafluorobutyrate
HFBA	heptafluorobutyric acid anhydride
Kdn	3-deoxy-D-glycero-D-galactonulosonic acid;
	the nomenclature of the other sialic acids is after Schauer and Kamerling [36]
TIC	total ion count
HE	hemagglutinin esterase
MHV	mouse hepatitis virus

## Introduction

Viral infections are initiated by binding of the virus to cellular receptors. In many instances the virus-receptor recognition is highly specific, conferring species specificity and/or

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tissue specificity. Coronaviruses, which belong to the order of Nidovirales, infect different animal species. They are enveloped viruses with a single-stranded plus-sense RNA genome of approximately 30 kilobases. Their genome is characterized by a 5'-terminal cap structure and a 3' poly A tract. The coronaviridae are divided into the toroviruses and the coronaviruses. The coronavirus genus is divided into three serogroups [1]. Most recently novel human coronaviruses were identified during the search on the etiological agent causing Severe Acute Respiratory Syndrome (SARS) [2–5]. SARS-CoV is an early split off of group 2 coronaviruses [6], HCoV-NL63 belongs to group 1 [7], and CoV-HKU I is a group 2 coronavirus [8]. Mouse hepatitis viruses (MHV) belong to group 2, together with rat coronaviruses, puffinosis coronavirus, human coronavirus OC43 (HCoV OC43), bovine coronavirus (BCoV), and porcine hemagglutinating encephalomyelitis virus (HEV). A major characteristic of group 2 coronaviruses is the presence of a hemagglutinin-esterase (HE) surface glycoprotein in addition to the viral spike protein. It was previously demonstrated [9] that the HE protein of the MHV strain *S* is a sialate-4-*O*-acetyltransferase, which is able to de-*O*-acetylate Neu4,5Ac<sub>2</sub>. In addition to this strain, puffinosis coronavirus and sialodacryoadenitis virus exhibit a similar substrate specificity [10–12]. The HE proteins of group 2 coronaviruses interact either with Neu4,5Ac<sub>2</sub> or sialic acids with *O*-acetylation at position 9 (Neu5,9Ac<sub>2</sub>). While human coronaviruses related to the reference strain HCoV OC43 and BCoV specifically interact with Neu5,9Ac<sub>2</sub> via the HE [13,14] and the spike glycoprotein [15], the murine coronavirus strains *S* and JHM recognize Neu4,5Ac<sub>2</sub> [9] and a cellular glycoprotein termed MHV-receptor or CEACAM 1a [16–19]. For Influenza *C* viruses and BCoV it was shown that they utilize Neu5,9Ac<sub>2</sub> as receptor determinant, which may be part of glycoproteins or the sialic-acid-containing gangliosides [14,20]. Although it is known that MHV-like coronaviruses bind and hydrolyse either Neu4,5Ac<sub>2</sub> or Neu5,9Ac<sub>2</sub> [9,21], the significance for infections by murine coronaviruses remained unclear.

Neu4,5Ac<sub>2</sub> is known to be expressed to high levels in guinea pigs and horses [22] as well as in a few other animals [23]. No detailed analysis of the occurrence of this sialic acid in mice is available. Recently, Neu4,5Ac<sub>2</sub> was detected in the colon and trachea of mice, although sialic acids *O*-acetylated at the glycerol side-chain prevailed [24]. MHV strains exhibit different virulence and tissue tropisms. MHV strains JHM and *S* represent neurotropic viruses, which cause either chronic demyelinating disease or rapidly fatal infection of the CNS, associated with a virus dissemination to the liver, bowel and other organs [25,26]. MHV strain A59 produces a mild to moderate meningoencephalitis and chronic demyelination, as well as moderate to severe hepatitis [27,28]. Enterotropic MHV strains preferentially replicate in the intestinal mucosa [29]. Since different strains of MHV preferentially infect ei-

ther the gastrointestinal tract, the liver, the respiratory tract or the CNS, we were aiming to analyse the distribution of *O*-acetylated sialic acids in all mouse tissues susceptible for infection by MHV.

A method was developed for the identification and quantification of the sialic acid diversity [30]. It involved, after liberation of sialic acids using mild acid hydrolysis, the formation of methyl esters with diazomethane, followed by the blockage of all free hydroxyl-(amino-) groups with heptafluorobutyric acid anhydride. Subsequent GC/MS analysis of these volatile derivatives allowed, in the electron impact mode of ionization, to identify 38 different sialic acids in samples from different origins. This method was applied to the study of the sialic acids glycosidically bound to glycoproteins and glycolipids from human erythrocyte membranes [31]. It allowed the unambiguous identification and quantification of a large variety of sialic acids including Neu4,5Ac<sub>2</sub>, which was never detected before in human tissues. The same methodology was used to demonstrate an increasing gradient of *O*-acetylation in the descending human gut [32]. Because of its sensitivity it became possible to analyze the occurrence and distribution of Neu4,5Ac<sub>2</sub> in less than 1 mg of initial total homogenate from different mouse tissues.

This study clearly demonstrates the presence of this sialic acid in mice and discusses the implications of tissue- and development-specific variations of the different sialic acids.

## Materials and methods

### Chemicals and media

Diazogen<sup>TM</sup> was from Aldrich (Milwaukee, USA). Heptafluorobutyric acid anhydride (HFBA; puriss) was from Fluka (Buchs, Switzerland). Heavy walled screw cap tubes (10 × 100 mm) and Teflon-lined caps (GL14) were from Schott (Mainz, Germany). Diazomethane was prepared in a Wheaton<sup>TM</sup> apparatus according to the procedure proposed by the manufacturer and the diazomethane solution was kept as aliquots in tubes identical to the reaction vials at room temperature in a ventilated hood.

### Animals

For all our studies we used NMRI (Naval Medical Research Institute, USA) mice.

### Preparation of primary astrocytes cell culture

Astrocytes were isolated and derived from newborn (NMRI) mouse brain as described by Tontsch and Bauer [33]. Brain halves were dissected under sterile conditions. Meninges were carefully removed, the tissue minced in cold DiSG solution containing 10 mM HEPES, 17 mM glucose and

22 mM sucrose (pH 6.8) and dissociated in 2 ml 0.3% Trypsin/DiSG for 5–10 min. Then 7 ml warm astrocyte medium was added, and the tissue fragments were dissociated by trituration through a Pasteur pipette. The cells were resolved in 8 ml astrocytes medium and equally plated in four 25 cm<sup>2</sup> flasks (Greiner, Austria). Astrocyte medium was Dulbecco's Modified Eagle's Medium (DMEM) purchased from Gibco with glucose (6 g/l) containing 10% FCS and supplemented with penicillin (50 IU/ml) and streptomycin (50 µg/ml). The astrocytes were incubated at 37°C in a humidified atmosphere of 5% CO<sub>2</sub> in air. After 6 h, medium was partly aspirated to remove floating cells and neurons and changed to fresh astrocytes medium. Cells were cultured for 12 days to confluency with regular changes of the culture medium every third day.

Astrocytes were harvested with a sterile cell scraper, washed 5 times in warm 1 × PBS and centrifuged at 300 g for 10 min. Supernatant was aspirated and the weight of the cell pellet was determined.

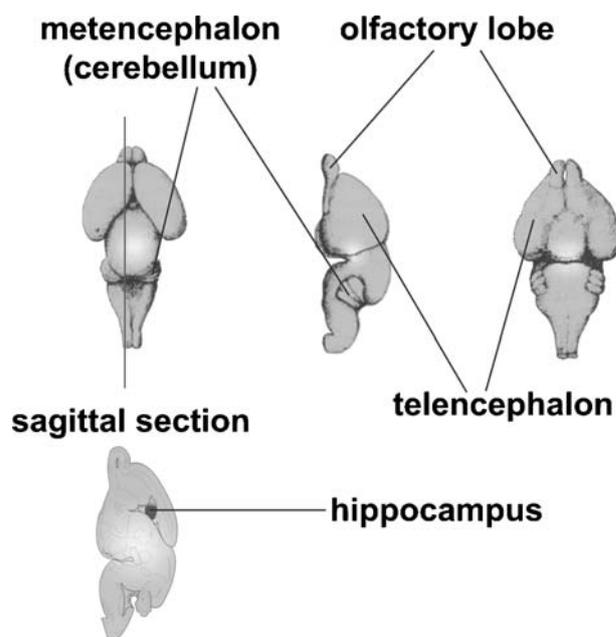
#### Dissection and delipidation of mouse tissues

Pre- and perinatal mice were sacrificed by decapitation, adult mice by cervical dislocation. After dissection, the brains were washed in cold 1 × PBS and the meninges were removed. Brains were from embryos at stage 13 (E13), 15 (E15), 17 (E17), from newborn (P0), postnatal day 7 (P7) and adult. The wet weight of each sample was determined. The hippocampus, olfactory lobe, metencephalon and the telencephalon (Figure 1) as well as liver and gut (duodenum, jejunum, ileum, cecum, colon and rectum) were prepared from an adult, male NMRI mouse.

Mouse tissues were minced in cold methanol, homogenized and samples containing 10 mg wet weight of the initial tissue were taken for delipidation. Cold MeOH was added to the homogenized tissue samples to a final volume of 1 ml and transferred into glass screw top vials. After stirring the samples for 15 min, 500 µl chloroform was added to each vial. To separate the glycoproteins and glycolipids, the samples were stirred for another 30 min and centrifuged at 3600 g for 10 min at room temperature. The supernatants, containing the glycolipids, were transferred into new glass vials. Pellets were resolved in 1 ml chloroform/methanol (2:1 v/v), stirred for 60 min and centrifuged at 3600 g for 10 min at room temperature. The combined supernatants (glycolipids) and pellet (glycoproteins) were dried separately under a light stream of nitrogen. The tissue samples were then ready for GC/MS analysis.

#### GC/MS analysis of sialic acid from mouse tissue

The GC/MS analysis of sialic acid was performed essentially as previously described [30] with a few modifications



**Fig. 1** Brain sections. Sections were taken from an adult NMRI mouse. For the preparation of the hippocampus a sagittal section was taken as indicated.

as follows. The dried samples were submitted to mild acid hydrolysis (2 M acetic acid for 105 min at 80°C) with intermittent agitation during the first 15 min of the reaction. The lipid samples were then lyophilized to dryness. The protein/glycoprotein samples were centrifuged for 60 min at 4°C (60 min at 3000 rpm) and the supernatant transferred to reaction vials, followed by lyophilisation.

All samples were methyl-esterified with diazomethane. Briefly, samples were supplemented with 200 µl of anhydrous redistilled methanol, then 200 µl of the diazomethane reagent was added in a ventilated hood. The samples were left in this reagent until analysis. Before GC/MS analysis, the samples were dried under a stream of nitrogen in a ventilated hood, and then supplemented with 400 µl of acetonitrile and 50 µl of heptafluorobutyric acid anhydride (Fluka). After heating for 5 min at 150°C, samples were evaporated in a stream of nitrogen until dryness, and then taken up in the desired volume of acetonitrile, which had been kept dry on calcinated calcium chloride. The samples (1 µl) were directly injected on the Ross injector of the GC/MS apparatus and analyzed in the impact mode of ionization.

The GC separation was performed on a Carlo Erba GC 8000 gas chromatograph equipped with a 25 m × 0.32 mm CP-Sil5 CB Low bleed/MS capillary column, 0.25 µm film phase (Chrompack France, Les Ulis, France). The temperature of the Ross injector was 260°C and the samples were analyzed using the following temperature program: 90°C for 3 min and then 5°C/min until 260°C. The column was coupled to a Finnigan Automass II mass spectrometer (mass detection limit 1000). The analyses were

performed in the electron impact mode (ionization energy 70 eV; source temperature 150°C). In order to preserve the filament of the ionization source, the GC/MS records were performed 5 min after the injection of the sample. For quantifications, the chromatograms were analyzed using the Xcalibur software (Thermoquest-Finnigan) integrating the signals at  $m/z = 169$ , a reporter base ion specific for heptafluorobutyrate (HFB) derivatives. This procedure allows overcoming the problem of contaminations by derivatives unrelated to sialic acids present in the samples derived from total homogenates. Each constituent was identified through its retention time and fragmentation profile. The major contamination interferences observed with the analysis of sialic acids were due to disaccharides of hexoses which were particularly abundant in the material derived from the liver lipid fraction.

## Results

### Identification and quantification of sialic acids

The GC/MS analysis of sialic acids as HFB derivatives allows separation of the different sialic acids, which are identified by their retention times and fragmentation patterns in the electron impact mode of ionization starting from very small amounts of total homogenates. Indeed, quantities as low as 1 pg of each sialic acid can be unambiguously identified with the MS detector through specific fragmentation mass spectra [30].

In the different mouse samples, 13 different sialic acids were identified and shown to be present at various levels (Tables 1 and 2). Six different mono-*O*-acylated sialic acids were detected, including 5-*N*-acetyl-4-*O*-acetyl neuraminic acid (Neu4,5Ac<sub>2</sub>), 5-*N*-acetyl-7-*O*-acetyl neuraminic acid (Neu5,7Ac<sub>2</sub>), 5-*N*-acetyl-9-*O*-acetyl neuraminic acid (Neu5,9Ac<sub>2</sub>), 5-*N*-acetyl-9-*O*-lactyl neuraminic acid (Neu5Ac9Lt), 5-*N*-glycolyl-9-*O*-lactyl neuraminic acid (Neu5Gc9Lt) and 5-*N*-acetyl-8-*O*-sulfo neuraminic acid (Neu5Ac8S). Besides 5-*N*-acetyl neuraminic acid (Neu5Ac), generally present as the major sialic acid, traces of 2-keto-3-deoxyxnononic acid (Kdn) were detected as a minor compound in the glycoproteins of the gut and in the glycolipids of the liver. A single di-*O*-acylated sialic acid was detected as 5-*N*-acetyl-4-*O*-acetyl-9-*O*-lactyl neuraminic acid (Neu4,5Ac<sub>2</sub>9Lt). Two compounds of the 5-*N*-glycolyl neuraminic acid (Neu5Gc) family, Neu5Gc and its lactylated derivative Neu5Gc9Lt were especially abundant in the glycoproteins and glycolipids of the liver. 5-*N*-acetyl-8-*O*-methyl neuraminic acid (Neu5Ac8Me) and neuraminic acid (Neu) were also present, the latter especially in the glycolipid fraction from various tissues. The intra-molecular lactone of Neu5Ac (Neu5Ac1,7L) was particularly abundant in

the glycolipids of the mesencephalon (more than 90%) and of the brain at P7.

### Sialic acids in different mice tissues

Neu4,5Ac<sub>2</sub> was abundant in glycoproteins (16%) and glycolipids (27.4%) of the gut (Tables 1 and 2). This finding correlates with data obtained by LC/MS analysis, when Morimoto *et al.* observed that 15.8% of the total peak area of sialic acids in the mouse colon represent this neuraminic acid derivative [24]. Smaller, but significant levels of Neu4,5Ac<sub>2</sub> were present in glycoproteins of the olfactory lobe, metencephalon and telencephalon. Remarkably, in the glycolipid fraction the relative amount of this sialic acid was lower in the neuronal tissues. In the gut, Neu4,5Ac<sub>2</sub> represented 27.4% of the total sialic acids in the glycolipid fraction, compared to 16% in the glycoprotein fraction. In almost all tissues examined, Neu5Ac and its derivatives were the major sialic acids with the exception of the glycolipid fraction of the adult metencephalon and the P7 brain in which Neu5Ac1,7L was the major compound (Table 2). Neu5Gc and its lactylated derivative Neu5Gc9Lt were especially abundant in the glycoproteins and glycolipids of liver (about 70% of total sialic acids). In contrast, Neu5Gc and Neu5Gc9Lt were poorly expressed in adult brain (Tables 1 and 2). Significant amounts of Neu5Ac8Me (7.3 and 12.8%) were observed with a particular abundance in gut glycoproteins and glycolipids (Tables 1 and 2). Neu was particularly abundant in the olfactory lobe and in the gut glycolipids (Table 2). Indirect evidence for the presence of this compound in a GD<sub>3</sub> ganglioside isolated from human melanoma was previously forwarded [34].

### Distribution of sialic acids during mouse brain development

As shown in Table 1, the different glycoprotein-bound sialic acids showed variable patterns during total brain development. Glycolipid-bound Neu4,5Ac<sub>2</sub> was only detectable in newborn mouse brain (P0) and seven day old mice (P7), whereas the glycoprotein-bound form was present during all stages of the brain development. The other mono-*O*-acetylated sialic acids (Neu5,7Ac<sub>2</sub> and Neu5,9Ac<sub>2</sub>) also showed significant quantities of their glycolipid bound forms at P0 (Tables 1 and 2).

Neu5Ac was found in very high amounts (60–98%) in almost all stages of the developing brain, except the glycolipids of P7 brain. Here, we found high amounts (57.7%) of the 1,7 lactone of Neu5Ac and significant amounts of Neu (19.3%). Glycolipid-bound Neu5Gc was only verifiable in the embryonic stages E15 and E17 of the mouse brain, whereas glycoprotein-bound Neu5Gc decreased till day of birth and then increased again. Neu5Ac8Me and Neu5Ac9Lt showed a maximal expression at birth, while Neu4,5Ac<sub>2</sub>9Lt showed

**Table 1** Distribution of glycoprotein-bound sialic acids (%) in different mouse tissues

	Brain E13	Brain E15	Brain E17	Brain P0	Brain P7	Brain adult	Hippo	Olf	Met	Tel	Astro	Gut	Liver
Kdn	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.12	0.00
Neu5Ac	87.87	90.96	89.54	88.75	87.66	87.68	92.47	84.21	85.33	87.61	87.81	68.02	24.62
Neu5Ac8Me	2.17	0.95	1.29	2.97	0.95	0.42	0.81	1.80	4.31	1.69	1.04	7.29	0.00
Neu5Ac9Lt	2.34	2.02	2.35	4.47	2.70	2.99	2.35	6.18	2.92	3.58	3.86	0.00	0.00
Neu5,9Ac <sub>2</sub>	0.89	1.45	1.57	0.58	1.15	1.29	0.60	1.11	2.47	1.65	1.17	2.55	0.00
Neu5Ac1,7L	0.00	0.00	0.00	0.00	0.35	0.00	0.00	0.00	0.00	0.00	0.00	0.00	3.54
Neu4,5Ac <sub>2</sub>	0.76	0.63	0.58	0.18	0.35	0.67	0.00	0.46	0.13	0.46	0.22	16.01	1.46
Neu	3.67	1.87	2.51	1.68	3.60	2.87	2.17	3.19	1.65	2.07	3.35	0.42	0.00
Neu5,7Ac <sub>2</sub>	0.00	0.00	0.00	0.15	0.09	0.46	0.39	0.00	0.38	0.00	0.00	0.00	0.00
Neu4,5Ac <sub>2</sub> 9Lt	1.11	0.91	1.82	1.21	2.39	2.37	0.96	1.60	1.06	1.08	1.10	0.00	0.00
Neu5Gc	1.19	1.13	0.00	0.02	0.13	0.47	0.11	0.43	1.21	0.67	0.76	5.60	41.98
Neu5Gc9Lt	0.00	0.00	0.17	0.00	0.00	0.50	0.00	0.50	0.55	0.86	0.34	0.00	28.40
Neu5Ac8S	0.00	0.08	0.16	0.00	0.63	0.29	0.14	0.52	0.00	0.32	0.35	0.00	0.00
% total sialic acid ⊗	57.67	10.70	11.96	17.85	48.97	33.60	41.41	37.53	69.25	51.14	51.48	65.52	33.40

Abbreviations: Hippo = hippocampus; Olf = olfactory lobe; Met = metencephalon; Tel = telencephalon; Astro = astrocytes; P0 = postnatal day 7; P7 = postnatal day 7; P0 = newborn; E17 = embryo day 17; E15 = embryo day 15; E13 = embryo day 13; ⊗ glycoprotein-bound + glycolipid-bound = 100%.

**Table 2** Distribution of glycolipid-bound sialic acids (%) in different mouse tissues

	Brain E13	Brain E15	Brain E17	Brain P0	Brain P7	Brain adult	Hippo	Olf	Met	Tel	Astro	Gut	Liver
Kdn	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.72
Neu5Ac	60.59	98.20	91.02	84.94	10.22	89.87	87.35	73.79	0.00	87.43	97.82	11.82	16.03
Neu5Ac8Me	2.36	0.00	1.78	0.00	4.37	1.97	3.82	4.14	9.14	1.86	0.68	12.82	0.00
Neu5Ac9Lt	3.89	0.00	1.98	1.07	0.00	0.00	0.68	1.24	0.00	4.09	0.00	0.00	0.00
Neu5,9Ac <sub>2</sub>	2.09	0.47	0.23	1.13	0.00	0.72	1.64	3.27	0.00	1.88	0.00	12.19	0.00
Neu5Ac1,7L	1.18	0.00	0.00	0.00	57.73	0.20	1.81	0.00	90.51	0.00	0.00	0.00	0.00
Neu4,5Ac <sub>2</sub>	0.00	0.00	0.00	0.36	1.01	0.06	0.46	0.00	0.00	0.00	1.50	27.42	0.00
Neu	27.79	0.90	4.37	10.65	19.38	5.55	0.88	16.31	0.22	1.90	0.00	35.75	0.00
Neu5,7Ac <sub>2</sub>	0.00	0.00	0.00	0.73	0.35	0.54	0.00	0.00	0.00	0.00	0.00	0.00	0.00
Neu4,5Ac <sub>2</sub> 9Lt	0.00	0.00	0.43	0.91	2.92	0.82	0.78	0.00	0.00	1.10	0.00	0.00	0.00
Neu5Gc	0.00	0.35	0.17	0.00	0.00	0.00	0.00	0.00	0.00	0.56	0.00	0.00	59.70
Neu5Gc9Lt	0.00	0.00	0.00	0.00	0.00	0.00	0.04	0.00	0.00	0.80	0.00	0.00	23.55
Neu5Ac8S	2.09	0.09	0.02	0.24	4.01	0.28	2.54	1.26	0.13	0.37	0.00	0.00	0.00
% total sialic acid ⊗	42.33	89.30	88.04	82.15	51.03	66.40	58.59	62.47	30.75	48.86	48.52	34.48	66.60

Abbreviations: Hippo = hippocampus; Olf = olfactory lobe; Met = metencephalon; Tel = telencephalon; Astro = astrocytes; P7 = postnatal day 7; P0 = newborn; E17 = embryo day 17; E15 = embryo day 15; E13 = embryo day 13; ⊗ glycoprotein-bound + glycolipid-bound = 100%.

a lower relative abundance at the same period. A similar pattern was also observed for Neu5Gc9Lt and Neu5Ac8S.

## Discussion

Due to methods used in the past, the analysis of *O*-acetylated sialic acids in glycoconjugates was difficult. The first structural analysis of various sialic acids, especially *O*-acetylated ones by mass-spectrometry, was reported by Kamerling *et al.* [35], using pertrimethylsilyl derivatives of the sialic acid methyl esters. This method significantly contributed to the discovery of a series of previously unknown naturally occurring sialic acids [36]. The GC/MS method used in this report differed essentially from other methodologies using HPLC separation of fluorescent compounds [22,37,38] for which the retention time is the major criterion, the coupling with mass spectrometry being difficult. These techniques require an initial purification of the liberated sialic acids, a time-consuming step performed starting from much higher quantities of initial material (at least 3 orders of magnitude). Furthermore, due to the hydrolysis and isolation procedure, minor quantities of sialic acids may be lost. Even if pooled mice brain were used for the delipidation step (needed for having data on glycoprotein and glycolipid-bound sialic acids), all data presented here were obtained on samples corresponding to 1 mg of initial protein or less.

The identity of the different compounds was ascertained by their fragmentation mass spectra and retention times. As previously described, the identification of the different sialic acids was performed using a chromatogram reconstitution for the ion at  $m/z = 169$  specific of HFB derivatives (Figure 2). This procedure allowed the elimination of the contaminants like glucose necessarily present in total homogenate samples. The manual integration of the areas of the different peaks allowed a quasi-perfect reproducibility for the major compounds (generally Neu5Ac; less than 0.001% of the reported values) and a good reproducibility for minor compounds (in all cases less than 1% of the reported values). Furthermore, the re-injection of a sample after the first analysis gave the same results within less than 1% error. The second step of identification was performed using chromatogram reconstitution with ions specific of the different classes of sialic acids ( $m/z = 861$  for mono-*O*-acetylated sialic acids;  $m/z = 707$  for di-*O*-acetylated sialic acids;  $m/z = 227$  for Neu5Gc derivatives;  $m/z = 112$  for 9-*O*-lactylated sialic acids;  $m/z = 122$  for 8-*O*-sulfated sialic acids;  $m/z = 805$  for 8-*O*-methyl sialic acids and  $m/z = 505$  for Neu). Finally, the exact position of the *O*-acetyl group of mono-*O*-acetyl sialic acids was determined by their precise fragmentation patterns [30].

Based on these criteria, the population of the sialic acids of mice tissues presents a high diversity. Although we did not

analyze the mouse gut mucins in particular (homogenates contained the mucus as well as subjacent tissues) it should be emphasized that the sialic acid composition of the mouse gut glycoproteins is significantly different from the sialic acid composition of human gut mucins [32]. A characteristic of human gut is its high level of oligo-*O*-acetylation. In the mouse gut, we could not detect significant amounts of such sialic acid. Careful examinations of the GC/MS chromatogram indicated that these compounds were never present at a level 20 times lower than minor sialic acids observed in mouse gut *i.e.* Neu. Therefore these compounds, although detectable, were not considered.

In conclusion, the diversity of sialic acids is concerned with all investigated mouse tissues and was observed both in glycoproteins and in glycolipids.

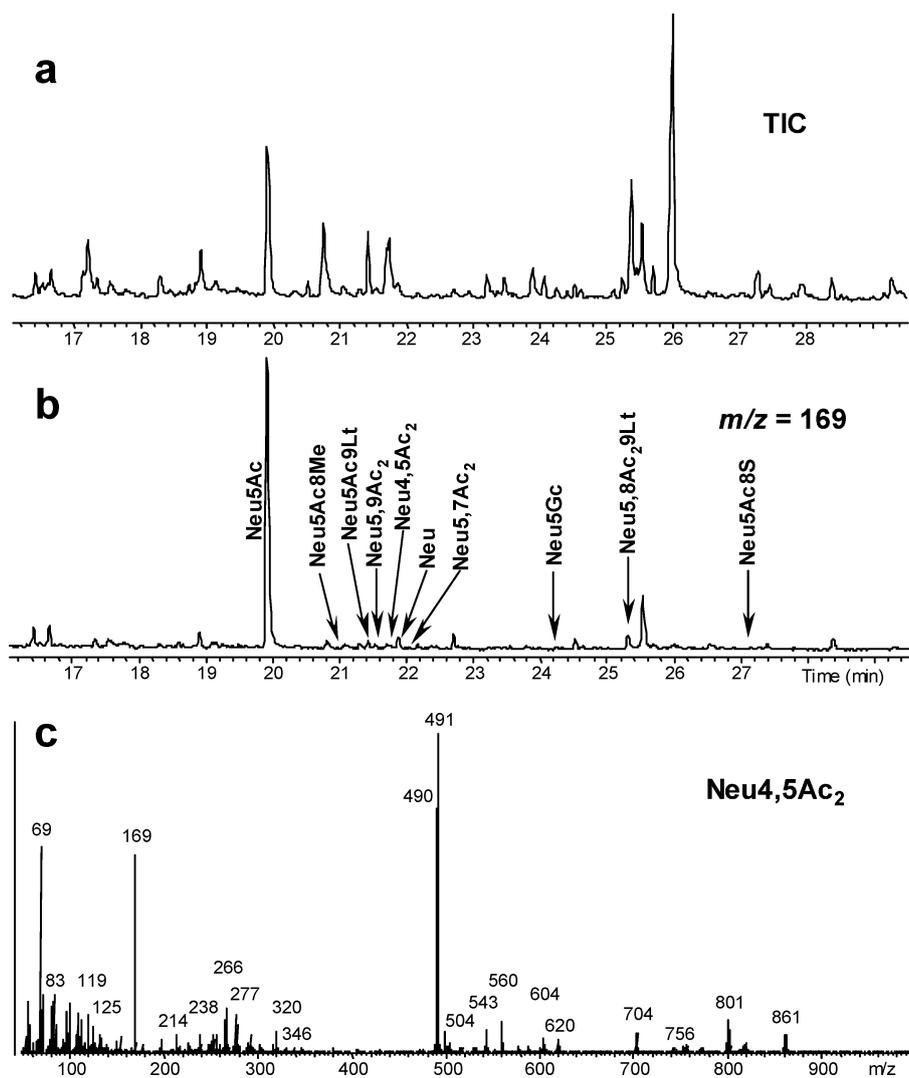
Neu4,5Ac<sub>2</sub> has been detected for the first time in mouse brain and it was observed in all target tissues of MHV. We observed a complete absence of glycoprotein-bound Neu4,5Ac<sub>2</sub> in the hippocampus.

The occurrence of *O*-acetylated sialic acids in glycolipids has been documented in many publications [39,40]. Earlier identification of *O*-acylation in gangliosides was possible by using two-dimensional thin-layer chromatography with an intermediate step of exposure to ammonia vapors [41]. Nevertheless, this method does not allow the identification of the precise nature and position of *O*-acyl groups. The use of antibodies or lectins specific for 9-*O*-acetylated sialic acids in gangliosides allowed detection of these compounds [42,43]. In our analyses of mammalian tissues with the new GC/MS technique, 9-*O*-lactylated compounds are generally more abundant than 9-*O*-acetylated sialic acids. This is the case for human [31,32], murine (here) and rat brain synaptosomal plasma membranes (Zanetta, unpublished results). The presence of Neu4,5Ac<sub>2</sub> is also a common feature in these three species, although the levels being lower compared to horse tissues [36,44]. However, as is best illustrated with the high level in the murine gut and the low level in the other mouse tissues, its expression apparently is tissue-specific. Therefore, the enzyme acetyl-CoA: Sialate-4-*O*-acetyltransferase (E.C. 2.3.1.44) involved in the synthesis of this compound should be present in these species and expressed in different activities *e.g.* in the various mouse tissues. This enzyme has recently been solubilized from guinea-pig liver Golgi membranes and partially characterized [45].

Studies on the sialic acids of the human erythrocyte membranes [31] emphasized the fact that individual variations are present for apparently healthy individuals. This was also observed for sialic acids from gut mucins [32], although an increasing gradient of *O*-acetylation in the descending gut was always observed. Evidently, the reasons for these individual variations remain speculative.

From the quantitative point of view, it is emphasized that the glycoprotein-bound level of sialic acids is low at the

**Fig. 2** GC/MS analysis of the sialic acids found in P7 mouse brain glycoprotein fraction. (a) Chromatogram obtained for the total ion count (TIC). (b) Chromatogram reconstitution for the ion at  $m/z = 169$  specific for HFB derivatives. (c) EI fragmentation mass spectrum of Neu4,5Ac<sub>2</sub>. Note in (a) that only Neu5Ac can be safely detected on the TIC chromatogram. In (b), the different minor sialic acids can be extracted from the majority of the contaminants and can be safely analyzed. Neu4,5Ac<sub>2</sub> is easily identified through a series of reporter fragment ions at  $m/z = 861$ , 704 and 491



period between E15 and P0 in the mouse brain. In the case of Neu4,5Ac<sub>2</sub>, the changes from 0.6% at E17 to 0.2% at P0 and to 0.7% in the adult is significant, especially considering the absolute amount of this compound in the samples. Analysis of sialic acids in cultures of pure astrocytes, which are susceptible to infection by MHV, also showed the presence of Neu4,5Ac<sub>2</sub> in these cells (Tables 1 and 2). The low amounts of Neu4,5Ac<sub>2</sub> in brain tissue raised the question of its cell localization. It may be specific for certain cell types but no evidence for this is provided in this study.

Studies of 9-*O*-acetylation showed a specific association of Neu5,9Ac<sub>2</sub> with synaptophysin [42]. This hypothesis may be verified using the HE protein as a lectin for detecting endogenous ligands and identification of these ligands using proteomic techniques.

The poor expression of Neu5Gc in the neuronal probes studied, in contrast to e.g. mouse liver, corresponds with similar observations made in other mammals [46]. It is discussed that this phenomenon is related to the function of the brain. A

rare sialic acid is 8-*O*-sulfated Neu5Ac, which was found in small to minor quantities in the neuronal probes investigated. Strikingly, it was absent both in glycoproteins and glycolipids from gut and liver. It was also described to occur, by using the same analytical method, in human erythrocytes [31]. By LC-ESI-MS, Morimoto *et al.* [24] identified Neu5Ac8S also in various tissues of mice and rats in relatively small quantities.

Another rare sialic acid, known to occur in relatively large amounts in some echinoderm species (reviewed in [47,36]) is 8-*O*-methylated Neu5Ac. Since the sialic acids were derivatised by diazomethane in the analytical method applied here, a mere chemical origin cannot be excluded. However, this is not very likely since the methyl group is exclusively found at *O*-8 and the amounts in the various mouse tissues differ, with both glycoproteins and glycolipids of the gut sharing a maximum of almost 13% Neu5Ac8Me. With regard to the finding of Neu (de-*N*-acetylated Neu5Ac) in most of the mouse tissues analyzed, this again, is possibly not due to an artefact. Neu has been discovered in various mammalian,

including human cells like leucocytes, melanoma and neuroblastoma cells [48,39]. Data on the biosynthesis and regulation of Neu, which is believed to be involved in transmembrane signaling [49], are scarce. Manzi *et al.* [50] reported a rapid turnover rate of the *N*-acetyl groups in the sialic acid moiety of melanoma gangliosides as studied by radioactive labelling experiments. The occurrence of Neu5Ac1,7L in sometimes large quantities in the brain and liver is particularly puzzling. This compound was never detected before the initial study [30], the major reason being that this compound was missed in all analyses involving ion-exchange purification of sialic acids before analysis. Furthermore, it does not react with fluorescent reagents specific for  $\alpha$ -keto-carboxylic acids and enables a sensitive HPLC analysis of sialic acids [37]. This substance is the only sialic acid being identified in the GC/MS analysis when the diazomethane methyl-esterification was omitted. Furthermore, this lactone having a pseudo-dioxane ring, is stable to acidic and alkaline treatments compared with other esters. It can be recovered partially after acid-catalysed methanolysis. The abundance of this compound in the glycolipid fraction of P7 contrasted with its quasi absence in all other glycoprotein and glycolipid fractions, except in liver glycoproteins. In the P7 glycolipids but not in the other tissues studied this abundance is associated with the abundance of Neu, as well as in other tissues with Neu5Ac1,7L occurrence (Tables 1 and 2). However, in the corresponding glycoprotein fraction of the same sample, the abundance of the lactone is not associated with the presence of Neu. It is emphasized that Neu5Ac1,7L was found to be the ligand of human IL-4 [51]. Although it is not known whether the murine IL-4 has the same lectin activity as the human one, it may be speculated that gangliosides possessing this type of sialic acid at this specific stage of development may be involved in signalling processes related to cell proliferation or apoptosis.

In summary, this manuscript provides the first evidence that many different sialic acids are present in mouse tissues. In fact, except the presence of Neu5Gc, it resembled that of human tissues in which the same mono-*O*-acylated compounds are present [31,32]. For instance Neu5,9Ac<sub>2</sub>, the predominant mono-*O*-acetylated sialic acid in human gut is also present in the mouse gut, even though in much lower amounts. There are also noticeable differences. Neu5,7Ac<sub>2</sub> could only be detected in very low amounts in the mouse brain but not in the gut. Remarkably, Neu4,5Ac<sub>2</sub>, is present at a high level in the gut, at lower but significant levels in glycoproteins of the liver and in glycoproteins and glycolipids of the brain, two tissues known to be targets of the mouse hepatitis virus strain S. Di- and oligo-*O*-acetylated sialic acids could not be detected in any kind of mouse tissue tested.

In an investigation on the transneural spread of the neurotropic MHV strain JHM within the mouse brain, Barnett *et al.* found that MHV JHM spreads in a characteristic way

throughout the rodent CNS without infection of the hippocampal complex [52]. Several lines of evidence suggest that the presence of CEACAM1 is required for infection by murine coronaviruses. CEACAM1 is a member of the immunoglobulin family, which is involved in intercellular adhesion and angiogenesis. The isoform CEACAM1a is also involved in binding and penetration of murine coronaviruses. The presence of this molecule is apparently essential for infection *in vivo*: Ceacam1a<sup>-/-</sup> mice are resistant to infection by MHV strain A59 [53]. CEACAM1a is a highly glycosylated surface protein [18], which is present in apical membranes of many epithelial cells [54]. In the case of MHV strain A59, which does not express an HE protein, binding is mediated by the viral spike protein S. The binding site is located at the aminoterminal region of the S protein, which recognizes the first loop of CEACAM1. During biosynthesis and maturation, the S protein is proteolytically hydrolyzed in the Golgi into the subunits S1 and S2, which remain non-covalently linked in mature virions. Binding of the S1 subunit to CEACAM1 results in a conformational change of S1 by generation of alternative disulfide bonds. In consequence, the S1 fragment dissociates from S2, thereby activating the fusion activity of the viral S2 fragment [55]. The binding site on the CEACAM1a molecule is flanked by two N-linked glycans. It was suggested that these glycans may help to delineate the region for the viral spike docking [18]. We have shown in the past that MHV strains S and JHM, which possess an HE protein, bind to Neu4,5Ac<sub>2</sub>, presumably via the spike protein [12]. Most recently, we found that the MHV strain DVIM expresses an HE protein which is specific for Neu5,9Ac<sub>2</sub> [21]. It is currently unknown whether N-glycans of murine CEACAM1a contain *O*-acetylated sialic acids. If so, the viruses might first bind to the glycans near the docking site on CEACAM1a, which then would direct the viral S1 to the docking site on CAECAM1a. In this case the first interaction between *O*-acetylated sialic acids and the viral S1 may not be essential *in vitro*. CEACAM1a, expressed in insect cells, was shown to bind to S1 and to trigger the activation of the S2 fusion domain. Since proteins expressed in Sf9 insect cells do not contain sialic acids, binding of MHV to recombinant CEACAM1a is most likely independent of the presence of Neu4,5Ac<sub>2</sub> or Neu5,9Ac<sub>2</sub>.

For HIV, the receptor is the surface molecule CD4, which is sufficient for binding. Once HIV has bound to CD4, the chemokine receptors CCR5 and CXCR4 are the major coreceptors, which are required for penetration (for review: [56–58]). Compared with the situation observed for HIV, we may speculate that infection by murine coronaviruses involves at least two steps. In general terms the receptor is essential for initial binding, and additional binding to the coreceptor then allows fusion and penetration. A similar situation may exist for MHV: Depending on the virus strain, the primary receptors are either 4- or 9-*O*-acetylated sialic acids.

Binding to these sugars, which we have shown to represent ligands for MHV, is the first step of infection. In order to gain entry into target cells, the viruses have then to bind to the specific coreceptor CEACAM1a. In cell culture systems with high expression rates of CEACAM1a the presence of *O*-acetylated sialic acids for infection by MHV may be not an absolute requirement. In the *in vivo* situation both the expression of *O*-acetylated sialic acids at the cell surface as well as the expression of HE proteins and the affinity of the viral S protein to *O*-acetylated sialic acids most likely determine the organ- and tissue tropism of different MHV strains. In the future it will be interesting to test this hypothesis.

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