Synthesis of Sulfo-Sialic Acid Analogues: Potent Neuraminidase Inhibitors in Regards to Anomeric Functionality

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The design, synthesis and application of N-acetylneuraminic acid-derived compounds bearing anomeric sulfo functional groups are described. These novel compounds, which we refer to as sulfo-sialic acid analogues, include 2-decarboxy-2-deoxy-2-sulfo-N-acetylneuraminic acid and its 4-deoxy-3,4-dehydrogenated pseudoglycal. While 2-decarboxy-2-deoxy-2-sulfo-N-acetylneuraminic acid contains no further modifications of the 2-deoxy-pyranose ring, it is still a more potent inhibitor of avian-origin H5N1 neuraminidase (NA) and drug-resistant His275Tyr NA as compared to the oxocarbenium ion transition state analogue 2,3-dehydro-2-deoxy-N-acetylneuraminic acid. The sulfo-sialic acid analogues described in this report are also more potent inhibitors of influenza NA (up to 40-fold) and bacterial NA (up to 8.5-fold) relative to the corresponding anomeric phosphonic acids. These results confirm that this novel anomeric sulfo modification offers great potential to improve the potency of next-generation NA inhibitors including covalent inhibitors.

Neuraminidase (NA, EC 3.2.1.18), also referred to as sialidase, is a retaining glycosidase that hydrolyzes α-ketosidic linkages to sialic acids. Common substrates recognized by NA contain terminal N-acetylneuraminic acid (Neu5Ac, NANA) or N-glycolylneuraminic acid (Neu5Gc, NGNA) with an α2-3 or α2-6 ketosidic bond to galactose. NA is found in many types of pathogens including Trypanosoma cruzi, Clostridium perfringens, Streptococcus pneumoniae, Vibrio cholerae, mumps virus, parainfluenza virus and influenza virus. By destroying sialic acid-containing glycosides, NA facilitates the movement of pathogens through biological environments rich in sialic acid receptors, including mammalian tissues and mucus. Therefore inhibition of NA can hinder the spread of sialic acid binding pathogens, and NA inhibitors offer great potential as anti-microbial agents. In the case of sialic acid binding viruses, NA also facilitates the release of progeny virions from infected cells. Accordingly, NA inhibitors have been especially successful as anti-influenza pharmaceuticals1, 2.

The first generation of NA inhibitors was based upon Neu5Ac and its dehydrated product 2,3-dehydro-2-deoxy-N-acetylneuraminic acid (Neu5Ac2en or DANA). Currently, all clinically used NA inhibitors contain an anomeric carboxy group, which forms strong electrostatic interactions with three structurally conserved arginine residues in the NA active site (Fig. 1)3. Despite a lack of sequence identity among NAs from unrelated species, many active site structural features, including the triarginyl cluster, are highly conserved across all known NAs. For example, in the case of influenza A NA, a glycoside hydrolase family 34 member, this triarginyl cluster consists of Arg118, Arg292 and Arg371 (Fig. 1)3. In the case of Clostridium perfringens NanI NA, a glycoside hydrolase family 33 member, the arginine triad consists of Arg266, Arg555 and Arg615.

Replacement of the inhibitor carboxy group with a phosphono group has previously been reported to improve NA inhibitory activity and therefore phosphonic acid analogues of sialic acid are currently under development4–10. This increase in potency has been attributed to stronger electrostatic interactions of the anomeric phosphono group with the conserved active site NA triarginyl cluster (Fig. 1)5, 6.

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Based upon the higher acidity and electronegativity of sulfo relative to carboxy and phosphono functional groups, we predicted that sialic acid anomer sulfonyl acid analogues, which we refer to as sulfo-sialic acid analogues, should form the strongest electrostatic interactions with the conserved NA active site triarginyl cluster (Fig. 1). This strategy has been considered in an in silico analysis, which also indicated that the sulfo group should produce the strongest binding inhibitors. Yet, the actual synthesis of sulfo-sialic acid analogues is quite challenging and has remained unreported. In fact, with exception to 2-amino and 2-acetamido anomer sulfonates, no 2-deoxy sugar with an anomic sulfo group has been reported to our knowledge.

We first became interested in the synthesis of sulfo-sialic acid analogues during our research on covalent NA inhibitors, which have utilized an electronegative 3-flouro group to destabilize formation of oxocarbenium ion transition states (Fig. 1). In addition to enhancing electrostatic interactions with the triarginyl cluster, a strong electron withdrawing sulfo group might also destabilize oxocarbenium ion formation. In this report we reveal the first synthesis of sulfo-sialic acid analogues and their application as potent NA inhibitors.

Results and Discussion
Sulfo-sialic acid analogues were synthesized via oxidation of a mixture of acetylthio intermediates according to Fig. 2. Neu5Ac (1)-derived peracetylated octoses 2a and 2b, first reported by Potter and von Itzstein, were selected as the substrate for sulfur addition. The decarboxylated Neu5Ac derivatives 2a and 2b were synthesized based on the method of Shie et al. with modifications.

Nucleophilic substitution of the 2 anomic position with HSAC was mediated by either BF3·Et2O or TMSOTf, using CH3CN, CHCl3, or 1,2-dichloroethane (1,2-DCE) as solvent (Table 1). Although our original intention was to produce only 3, during NMR analysis of the acetylthio reaction products, a surprisingly large peak was always observed at 5.80 ppm, with coupling to another singlet at 6.27 ppm. These NMR peaks were later attributed to H-1, H-2 and H-3 of the 2,3-unsaturated pseudoglycal 4b, similar to compounds resulting from the Ferrier rearrangement of glycals in which the axial orientation of the anomeric acetylthio group is also preferred. The axial orientation of the acetylthio group of 4b is indicated by the lack of NOE correlation between H-1 and H-5. The structure of 4b was further supported by high-resolution mass spectrometry (HRMS), and characterization of the oxidation products 6b and 8b. The dominant change in the axial acetylthio orientation of this pseudoglycal could be explained by the anomic effect. Less polar reaction conditions were used to produce a higher ratio of 4b to 3 (Table 1). More polar reaction conditions could be used to weaken the anomic effect and increase the proportion of equatorial acetylthio compound 3a (Table 1).

Oxidation of the acetylthio functionality of 3 and 4 to the sulfo functionality was achieved using excess Oxone (potassium peroxymonosulfate, KHSO5·0.5KHSO4·0.5K2SO4) in AcOH buffered with KOAc, based upon the method of Reddie and modified with NH4OAc in place of KOAc for the oxidation of 3 to 5, which appeared to be quantitative. Yields for the oxidation of 4 to 6 were low due to concurrent oxidation of the pseudoglycal ring double bond by the excess Oxone. During HRMS analysis of polar Oxone oxidation products, negative ions m/z 452.0826 (6 + O), m/z 454.0985 (6 + O + 2H) and m/z 470.0949 (6 + O + 4H) were detected, indicating that 6 was further oxidized.

Sulfonates 5a and 5b could also be converted to the corresponding methyl esters by reaction with trimethylsilyldiazomethane, a toxic reagent that should be handled with caution. Active sulfo-sialic acid analogues, 7a, 7b and 8b were obtained after deprotection of the precursors in the presence of NaOH.

The structures of sulfo-sialic acid analogues 5-8 were determined using HRMS, 1H, 13C, 1H-1H COSY, NOESY and 13C-13C data. H-1, H-2 and H-5 are all shifted downfield in the axial anomer sulfonate 5b relative to the equatorial isomers 5a and 7a, which is also observed in 2, 3, and the phosphonic acid analogues of 5 and 7. The H-1 to H-2 coupling constants of 5a, 5b, 7a and 7b match those of the corresponding phosphonic acids and phosphonates.
carboxylic acids as well. A clear H-1 to H-5 NOE correlation is observed in the equatorial sulfo compound 7a. In contrast, the lack of any NOE correlation between H-1 and H-5 in 8b is consistent with the axial anomeric functional group orientations of 4b, 6b and 8b. A clear H-1 to H-2 NOE correlation in 8b further supports the equatorial orientation of H-1. Synthesis of pseudo-glycals with anomeric sulfones has also been reported with axial anomeric stereoselectivity. Isomerization of the anomeric position was never observed for any of the sulfo-sialic acid analogues.

Due to the high electronegativity of the sulfo group, the diagnostic H-1, H-2, H-5 and C-1 chemical shifts of 5 and 7 are shifted downfield relative to the corresponding carboxylic acids and phosphonic acids (Table 2). This observance of a strong electron-withdrawing inductive effect of the anomeric sulfo group supports our hypothesis that sulfo-sialic acid analogues will be useful for the development of next-generation covalent NA inhibitors.

In order to assess the relative effectiveness of sulfo-sialic acid analogues, the anomeric phosphonic acids corresponding to 7a (equPO₃H, ePANA) and 7b (axPO₃H, aPANA) were synthesized based upon the methods of Shie et al., with modifications. Inhibition of influenza A/Anhui/1/2005 (H5N1) NA (N1), A/RI/5+/1957 (H2N2) NA, Clostridium perfringens Nan) NA (CpNA) and Streptococcus 6646K NA (StrepNA), was quantified using a fluorogenic assay based upon the method by Potier et al. with modifications. Inhibition of influenza and bacterial NA by sulfo-sialic acid analogues confirms that the sulfo group enables more potent NA inhibition relative to the analogous phosphonic acids (Table 3, Fig. 3). The NA inhibitory activity of axPO₃H was difficult to quantify due to its low potency. For inhibition of influenza N2, the equatorial anomeric sulfonate 7a was 40-fold more...
against highly-prevalent drug-resistant NA and highly-pathogenic avian-origin NA. The Km Arg292Lys substitution is well documented, we showed that it results in an 88-fold increase and a decrease in Km.[1.62 (1.08–2.27) −]− −−−

Table 2. Diagnostic $^{13}$C and $^1$H chemical shifts (ppm) of sulfo-sialic acid analogues compared to the corresponding phosphonic acids$^4$, and carboxylic acids$^{20}$. Sialic acid numbering has been changed to assign the anomeric position as 1.

|          | C-1 Anomeric | C-2 | H-1 Anomeric | H-2ax | H-2eq | H-5 | Solvent |
|----------|--------------|-----|--------------|-------|-------|-----|---------|
| 7a       | 87.6         | 36.4| 4.28         | 1.79  | 2.45  | 3.63| CD$_2$OD |
| 7b       | 86.7         | 34.3| 4.70         | 1.92  | 2.62  | 4.25| CD$_2$OD |
| eqPO$_3$H| 73.6         | 36.2| 3.75         | 1.70  | 2.24  | 3.49| CD$_2$OD |
| axPO$_3$H| 71.9         | 34.4| 4.25         | 1.94  | 2.36  | 3.93| CD$_2$OD |
| eqCO$_2$H*| 76.8         | 37.8| 3.87         | 1.56  | 2.36  | 3.58| D$_2$O   |
| axCO$_2$H*| 74.8         | 35.2| 4.44         | 1.87  | 2.56  | 3.62| D$_2$O   |

Table 3. Km (inhibitory constant) and IC$_{50}$ (50% inhibition constant) values for the inhibition of N1, N2, CpNA, and StrepNA. All values are given in μM units and 95% confidence intervals are listed inside parenthesis.

|          | 7a                  | 7b                   | 8b                   | eqPO$_3$H |
|----------|---------------------|----------------------|----------------------|-----------|
| N1-K$_i$ | 1.62 (1.08–2.27)    | −                     | −                     | −         |
| N1-His275Tyr K$_i$ | 2.67 (1.63–2.62) | −                     | −                     | −         |
| N2-K$_i$ | 2.47 (1.95–3.14)    | 33.5 (26.0–43.1)     | 88.1 (72.8–107)      | 103 (82.6–128) |
| N2IC$_{50}$ | 8.86 (6.99–11.3) | 120 (93.0–155)       | 316 (260–384)        | 368 (296–458) |
| CpNA IC$_{50}$ | 37.5 (31.2–45.0)  | −                     | −                     | 320 (221–460) |
| StrepNA IC$_{50}$ | 3.74 (2.56–5.43) | −                     | −                     | −         |

The sialic acid anomeric sulfo modification is advantageous in terms of drug resistance because of the essential role of the triarginy1 binding site for natural NA substrate binding. Substitution of Arg118, Arg292 and Arg371 is known to result in impaired viral fitness due to decrease in NA activity and substrate affinity$^{24}$. Although the Arg292Lys substitution is well documented, we showed that it results in an 88-fold Km increase and a decrease in the enzymatic activity of H7N9 NA$^{24}$. One of the most common forms of NA inhibitor resistance is caused by a His275Tyr substitution, which confers a high level of resistance to oseltamivir and moderate resistance to peramivir$^{24,26}$. Consistent with the results for A/RI/5+1957(H2N2) NA, 7a inhibits H5N1 N1 and its drug-resistant His275Tyr form at the same level (Table 3, Fig. 4), demonstrating the effectiveness of sulfo-sialic acid analogues against highly-prevalent drug-resistant NA and highly-pathogenic avian-origin NA.

NMR Analysis. Compounds were analyzed using Varian 400 MHz and 600 MHz systems (now Agilent) at the Okayama University Collaboration Center, and also using a JEOL ECZ400 spectrometer at the Kobe University Department of Chemical Science and Engineering. Measurements were taken with 2-4 dissolved in CDCl$_3$, and 5-8 dissolved in CD$_2$OD. NMR chemical shifts ($\delta$) are provided in parts per million (ppm).$^1$H-NMR coupling constants (J) are listed in Hz. Residual peaks of chloroform (7.26 ppm) and methanol (3.31 ppm) were used as$^1$H-NMR references. The residual peaks of chloroform (77.0 ppm) and methanol (49.0 ppm) were also used as$^{13}$C-NMR references. For 2D experiments, the standard pulse sequences programmed into the Varian NMR systems were used. 2D-NOESY experiments were performed at room temperature with a spectral width of 9615.4 Hz (sw), an acquisition time of 0.15 seconds (at), 2844 data points (np), a filter bandwidth of 4000 Hz (fb), a relaxation delay of 2.000 seconds (d1), 4 transients (nt), 128 or 200 experiments (ni), a pulse width of 9.700 or 10.900 microseconds (pw) and a mixing time of 0.500 seconds (mixN).
High Resolution Mass Spectrometry (HRMS). Compound identities were confirmed with HRMS, which was performed using an Agilent 6520 Accurate-Mass Q-TOF at the Okayama University Collaboration Center.

Neuraminidase Inhibition. Inhibition of neuraminidase (NA) activity was quantified using a fluorogenic assay based upon the method by Potier et al. with modifications. 4-methylumbelliferyl-α-d-Neu5Ac
Modeling Sulfo-Sialic Acid Analogue Binding. Sulfo-sialic acid analogue binding to NA active sites was modeled with Molecular Operating Environment (MOE) version 2016.0802 from Chemical Computing Group. The structure of 7a was built in the A/RI/5+1957(H2N2) N1 (PDB ID: 3CKZ) active sites using the MOE Builder function. This was followed by running Structure Preparation, Protonate 3D, and Energy Minimization with the Amber10:EHT forcefield. Binding of 7a to wildtype N1 was performed in the same manner as that of the N1-His275Tyr structure (PDB ID: 3CKZ) after substitution of Try275 with histidine using the MOE Protein Builder function.

Synthesis of 4-Acetamido-3,6,7,8-tetra-O-acetyl-1,2,4-trideoxy-d-glycero-d-galacto-octopyranose (3) and 4-Acetamido-6,7,8-tri-O-acetyl-1,2,4-trideoxy-d-glycero-d-galacto-octopyranose (4). The known compound 4-acetamido-1,3,6,7,8-penta-O-acetyl-2,4-dideoxy-d-glycero-d-galacto-octopyranose (2) was synthesized based on the methods of Shi et al. with a 49% yield. 2 (0.4-1 g, 0.69-2 mmol) was dissolved in 1,2-dichloroethane (1,2-DCE) or CH₃CN dried with excess molecular sieves. After cooling, HSAc and Lewis acid (BF₃·Et2O or TMSOTf) were carefully added to the solution to the reaction mixture, which was then filtered through a fritted funnel. The crude mixture was evaporated under reduced pressure. Acetylthio reaction products were obtained from 4-MU-Neu5Ac. Fluorescence of liberated 4-methylumbelliferyl (Ex. 355-360 nm, Em. 460 nm) was detected at room temperature using a Thermo Scientific Varioskan™ Flash or a Molecular Devices SpectraMax Paradigm microplate reader. IC₅₀ values and confidence intervals were calculated by analyzing inhibition data using the log(inhibitor) vs. response (three parameters) function of Prism 7 software. N1 Kᵦ values were calculated using the Michaelis-Menten function of Prism 7, and N2 Kᵦ values were calculated using the Cheng-Prusoff equation. A/Anhui/1/2005 (H5N1) NA was purchased from Sino Biological. Influenza A/RI/5+/1957(H2N2) NA was produced according to our previously reported methods.

Synthesis of 4-Acetamido-3,6,7,8-tetra-O-acetyl-1-sulfo-1,2,4-trideoxy-d-glycero-d-galacto-octopyranose (5). (4-MU-Neu5Ac) substrate was utilized at a final concentration of 120 µM for inhibition of N2 and Streptococcus 6646K NA (Amsbio). For inhibition of Clostridium perfringens NanJ NA (Sigma) the 4-MU-Neu5Ac concentration was 60 µM when using 7a and 120 µM when using eqPO₃H. For IC₅₀ determination, 5 inhibitor concentrations were used with 1–4 replicates for each condition. For N1 Kᵦ determination, 5 inhibitor concentrations and 3 substrate concentrations were used. NA and inhibitor were pre-incubated for 20 minutes at room temperature before starting each assay with addition of 4-MU-Neu5Ac. Fluorescence of liberated 4-methylumbelliferyl (Ex. 355–360 nm, Em. 460 nm) was detected at room temperature using a Thermo Scientific Varioskan™ Flash or a Molecular Devices SpectraMax Paradigm microplate reader. IC₅₀ values and confidence intervals were calculated by analyzing inhibition data using the log(inhibitor) vs. response (three parameters) function of Prism 7 software. N1 Kᵦ values were calculated using the Michaelis-Menten function of Prism 7, and N2 Kᵦ values were calculated using the Cheng-Prusoff equation. A/Anhui/1/2005 (H5N1) NA were purchased from Sino Biological. Influenza A/RI/5+/1957(H2N2) NA was produced according to our previously reported methods.
Synthesis of 4-Acetamido-6,7,8-tri-O-acetyl-1-sulfo-2,3-didehydro-d-glycero-d-galacto-octopyranose (6). 4b (76 mg, 176 μmol) was dissolved in AcOH (3 mL). KOAc (Wako, 190 mg, 1.94 mmol, 9.85 eq) and OXone (378 mg, 1.23 mmol, 6.24 eq) were mixed in to form a slurry. After 4.66 hours MeOH was added to the reaction mixture, which was then filtered through a fritted funnel. The crude mixture was evaporated under reduced pressure, followed by purification of 6b (10.5 μmol, 6% yield) using normal phase column chromatography.

Preparation of 4-Acetamido-1-sulfo-2,4-trideoxy-d-galacto-d-galacto-octopyranose (7) and 4-Acetamido-1-sulfo-2,3-didehydro-1,2,3,4-tetradeoxy-d-glycero-d-galacto-octopyranose (8). To prepare active sulfo-sialic acid analogues, 5a (10 mg), 5b (10 mg), and 6b (4 mg) were treated with a solution of 0.1 M NaOH in MeOH. After deprotection reached completion, the pH was neutralized with AcOH and solvent was evaporated under reduced pressure resulting in quantitative final product yields. 7a, 7b and 8b were purified to homogeneity using an Inertsil® ODS-3 semi-preparative HPLC (GL Sciences) with a mobile phase of 10–15% MeOH supplemented with NH₄OAc.

Data Availability. The datasets generated and analyzed during the current study are available from the corresponding authors upon reasonable request.

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Author Contributions
C.J.V. and H.K. planned the research, designed the experiments and wrote the manuscript. C.J.V. synthesized all novel compounds and carried out the NA inhibition assays. C.M. synthesized the phosphonate analogues used in this study. T.H. and Y.K. assisted with inhibition assays. M.A. helped analyze sulfo-sialic acid analogue binding in silico. Y.W. and G.F.G. provided the recombinant influenza N2 used in this study. H.O. and M.I. assisted with the synthetic strategies.

Additional Information
Competing Interests: The authors declare that they have no competing interests.

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