

Supporting Information

**Enhanced Carbohydrate Structural Selectivity in Ion Mobility-Mass Spectrometry
Analyses by Boronic Acid Derivatization**

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Abstract: Boronic acid derivatization of carbohydrates and glycans is demonstrated as an ion mobility shift strategy to improve confidence in the identification and characterization of carbohydrate assignments using ion mobility-mass spectrometry.

Table S1. Common names, systematic names, average molecular weights, and derivatized mass of the carbohydrates reported in this work.

| Common name or abbreviation | Systematic name | Molecular weight (M_r , Da) | [M+ Na] ⁺ (Da) | [M+FBA+Na] ⁺ (Da) | [M+PBA+Na] ⁺ (Da) |
|-----------------------------------|--|--------------------------------------|------------------------------|---------------------------------|---------------------------------|
| Maltose | 4-O- α -D-Glucose-D-glucose | 342.3 | 365.3 | 559.1 | 641.5 |
| Lactose | β -D-galactose-(1 \rightarrow 4)- α -D-glucose | 342.3 | 365.3 | 559.1 | 641.5 |
| LN | β -D-Galactose-(1 \rightarrow 4)-D-N-acetylglucosamine | 383.4 | 383.4 | 600.2 | 682.5 |
| Gala3-type1 | Galactose- α -(1 \rightarrow 3)-galactose- β -(1 \rightarrow 3)-N-acetylglucosamine- β -azide | 614.6 | 637.6 | 831.4 | 913.7 |
| P1 | Galactose- α -(1 \rightarrow 4)-galactose- β -(1 \rightarrow 4)-N-acetylglucosamine- β -azide | 614.6 | 637.6 | 831.4 | 913.7 |
| Lec-Lec | Galactose- β -(1 \rightarrow 3)-N-acetylglucosamine- β -(1 \rightarrow 3)-galactose- β -(1 \rightarrow 3)-N-acetylglucosamine- β -azide | 817.8 | 840.8 | 1034.6 | 1116.9 |
| LNT | Galactose- β -(1 \rightarrow 3)-N-acetylglucosamine- β -(1 \rightarrow 3)-galactose- β -(1 \rightarrow 4)-N-acetylglucosamine- β -azide | 817.8 | 840.8 | 1034.6 | 1116.9 |
| LNFP1 | α -L-fucose-(1 \rightarrow 2)- β -D-galactose-(1 \rightarrow 3)- β -D-N-acetylglucosamine-(1 \rightarrow 3)- β -D-galactose-(1 \rightarrow 4)-D-glucose | 853.8 | 876.8 | 1070.6 | 1152.9 |
| LNFP2 | β -D-Galactose-(1 \rightarrow 3)-(α -L-fucose-[1 \rightarrow 4])- β -D-N-acetylglucosamine-(1 \rightarrow 3)- β -D-galactose-(1 \rightarrow 4)-D-glucose | 853.8 | 876.8 | 1070.6 | 1152.9 |

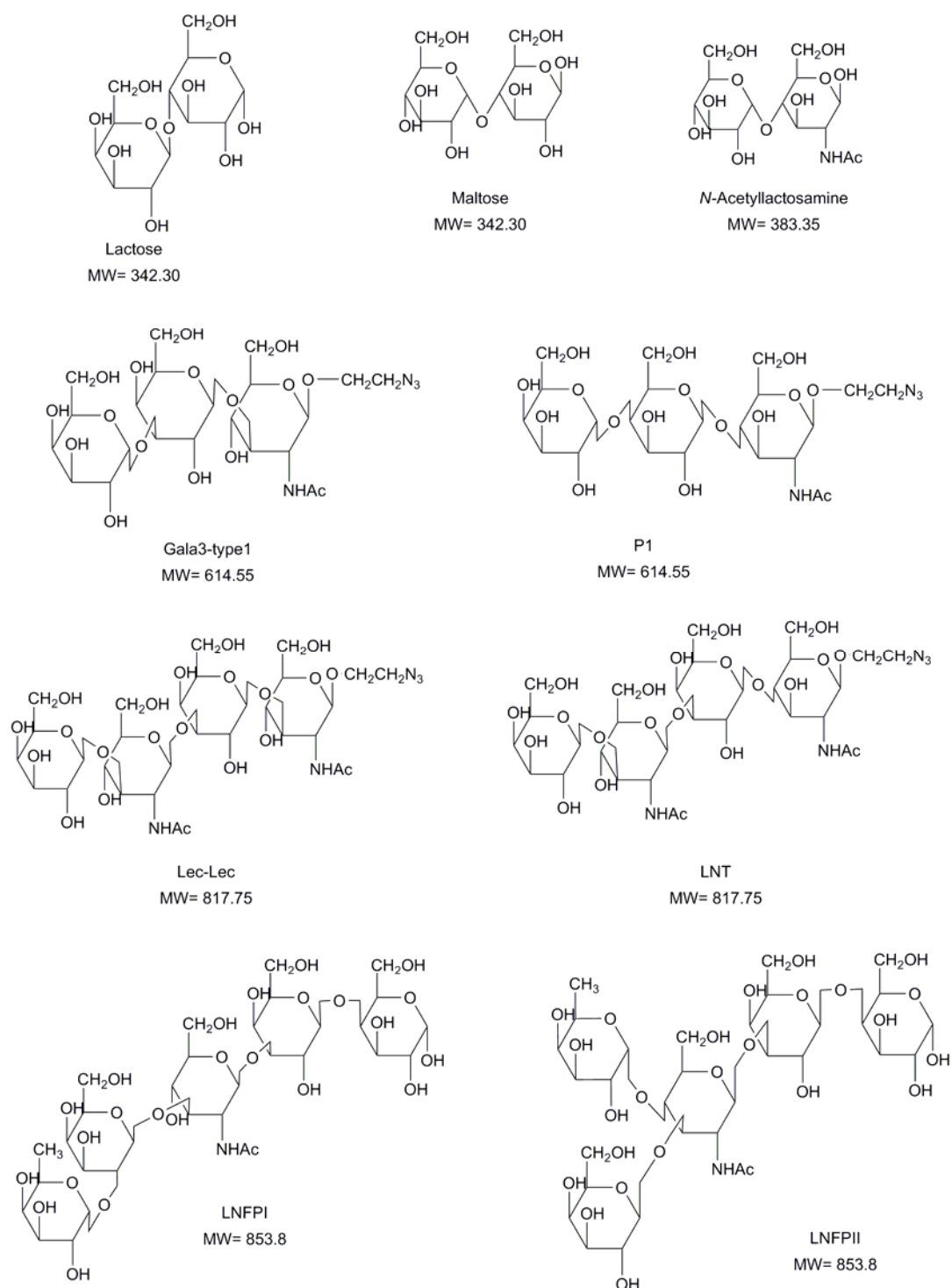


Figure S1 Structures of the underivatized carbohydrates studied.

Table S2 Collision cross sections determined for protonated and/or sodium coordinated carbohydrates prior to and following boronic acid derivatization.

| Carbohydrate | Collision cross section (\AA^2) ^a | | |
|--------------|---|-------------------------------------|-------------------------------------|
| | $[\text{M}+\text{Na}]^+$ | $[\text{M}+\text{FBA}+\text{Na}]^+$ | $[\text{M}+\text{PBA}+\text{Na}]^+$ |
| Maltose | 103.1 ± 4.1 (21) | 140.3 ± 7.7 (5) | 156.3 ± 5.7 (16) |
| Lactose | 106.8 ± 5.1 (35) | 151.5 ± 8.4 (16) | 163.7 ± 3.4 (30) |
| LN | 117.4 ± 1.8 (23) | 156.7 ± 3.4 (15) | 195.0 ± 1.5 (30) |
| Gala3-type1 | 160.2 ± 2.1 (26) | 197.9 ± 4.4 (31) | 229.8 ± 7.3 (18) |
| P1 | 166.9 ± 1.2 (70) | 200.5 ± 2.8 (39) | 255.4 ± 2.3 (40) |
| Lec-Lec | 183.2 ± 1.6 (51) | 220.0 ± 2.6 (15) | 252.2 ± 2.1 (21) |
| LNT | 195.9 ± 1.4 (42) | 217.9 ± 2.8 (19) | 254.8 ± 8.3 (19) |
| LNFP1 | 204.3 ± 1.4 (147) | 222.5 ± 3.4 (9) | 230.2 ± 6.1 (26) |
| LNFP2 | 201.4 ± 1.0 (166) | 226.1 ± 2.3 (11) | 233.7 ± 3.1 (21) |

a. Error represents $\pm 1\sigma$ for n measurements indicated in parenthesis. The protonated form of PBA derivatized carbohydrates is not included as the relative abundance for $[\text{M}+\text{PBA}+\text{H}]^+$ species was <2% of the base peak ($[\text{M}+\text{PBA}+\text{Na}]^+$) in the spectra.