

Table S1. The table illustrates the formulas utilized to calculate derived glycan traits from integrated peaks generated by CGE-LIF and Empower 3 software.

| Glycan trait | Description | Calculation |
|--------------|-----------------------------|---|
| G0 | agalactosylation | P14+P15+P18 |
| G1 | monogalactosylation | P16+P17+P19+P20+P21+P22+P23+P24 |
| G2 | digalactosylation | P25+P26+P27 |
| G | galactosylation (G1+G2) | P16+P17+P19+P20+P21+P22+P23+P24+P25+P26+P27 |
| S0 | asialylation | P14+P15+P16+P17+P18+P19+P20+P21+P22+P23+P24+P25+P26+P27 |
| S1 | monosialylation | P5+P6+P7+P8+P9+P10+P11+P12+P13 |
| S2 | disialylation | P1+P2+P3+P4 |
| S | sialylation (S1+S2) | P1+P2+P3+P4+P5+P6+P7+P8+P9+P10+P11+P12+P13 |
| CF | fucosylation | P3+P4+P7+P8+P12+P13+P15+P18+P21+P22+P23+P24+P26+P27 |
| CF0 | afucosylation | P1+P2+P5+P6+P9+P10+P11+P14+P16+P17+P19+P20+P25 |
| B | bisecting GlcNAc | P2+P4+P11+P13+P14+P18+P19+P20+P23+P24+P25+P27 |
| B0 | absence of bisecting GlcNAc | P1+P3+P5+P6+P7+P8+P9+P10+P12+P15+P16+P17+P21+P22+P26 |

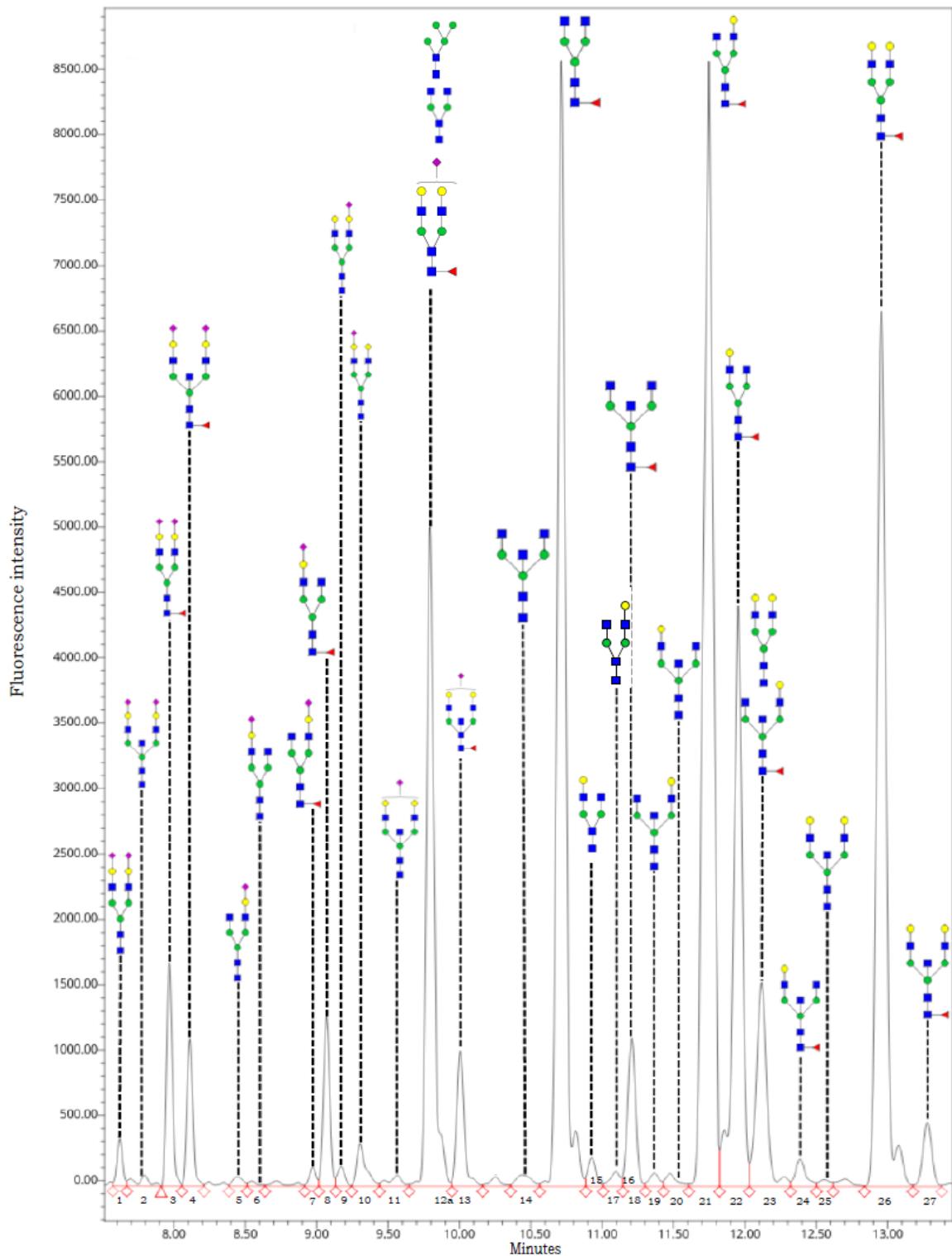
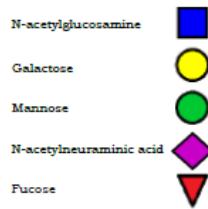


Figure S1. The image depicts the modified integrated glycan profile created using Empower 3 software. Each of the 27 peaks is clearly labelled in sequence, and the glycans associated with each peak are accurately identified.