

SUPPLEMENTARY TABLES

Supplementary Table 15. Description of directly measured subclass-specific Fc IgG glycan traits measured by LC-MS with mass list.

| Glycan trait ¹ | IgG1 glycopeptide m/z ³ [M+2H] ²⁺ [M+3H] ³⁺ | IgG2&3 glycopeptide m/z ⁴ [M+2H] ²⁺ [M+3H] ³⁺ | IgG4glycopeptide m/z ⁵ [M+2H] ²⁺ [M+3H] ³⁺ | Glycan trait description | Glycan trait calculation ⁶ | | | | |
|---------------------------|--|---|---|--------------------------|---------------------------------------|----------|----------|--|---|
| G0F | | 1317,527 | 878,687 | 1301,532 | 868,024 | 1309,529 | 873,356 | Fraction of FA2 glycan in total subclass Fc glycans | <i>G0F/total subclass Fc glycans*100</i> |
| G1F | | 1398,553 | 932,705 | 1382,558 | 922,042 | 1390,556 | 927,373 | Fraction of FA2G1 glycan in total subclass Fc glycans | <i>G1F/total subclass Fc glycans*100</i> |
| G2F | | 1479,58 | 986,722 | 1463,585 | 976,059 | 1471,582 | 981,391 | Fraction of FA2G2 glycan in total subclass Fc glycans | <i>G2F/total subclass Fc glycans*100</i> |
| G0FN | | 1419,067 | 946,38 | 1403,072 | 935,717 | 1411,069 | 941,049 | Fraction of FA2B glycan in total subclass Fc glycans | <i>G0FN/total subclass Fc glycans*100</i> |
| G1FN | | 1500,093 | 1000,398 | 1484,098 | 989,735 | 1492,096 | 995,066 | Fraction of FA2BG1 glycan in total subclass Fc glycans | <i>G1FN/total subclass Fc glycans*100</i> |
| G2FN | | 1581,119 | 1054,416 | 1565,125 | 1043,752 | 1573,122 | 1049,084 | Fraction of FA2BG2 glycan in total subclass Fc glycans | <i>G2FN/total subclass Fc glycans*100</i> |
| G1FS1 | | 1544,101 | 1029,737 | 1528,106 | 1019,073 | 1536,104 | 1024,405 | Fraction of FA2G1S1 glycan in total subclass Fc glycans | <i>G1FS1/total subclass Fc glycans*100</i> |
| G2FS1 | | 1625,127 | 1083,754 | 1609,133 | 1073,091 | 1617,13 | 1078,423 | Fraction of FA2G2S1 glycan in total subclass Fc glycans | <i>G2FS1/total subclass Fc glycans*100</i> |
| G1FNS1 | | 1645,641 | 1097,430 | 1629,646 | 1086,767 | 1637,643 | 1092,098 | Fraction of FA2BG1S1 glycan in total subclass Fc glycans | <i>G1FNS1/total subclass Fc glycans*100</i> |
| G2FNS1 | | 1726,667 | 1151,447 | 1710,672 | 1140,784 | 1718,67 | 1146,116 | Fraction of FA2BG2S1 glycan in total subclass Fc glycans | <i>G2FNS1/total subclass Fc glycans*100</i> |
| G0 | | 1244,498 | 830,001 | 1228,503 | 819,338 | 1236,501 | 824,67 | Fraction of A2 glycan in total subclass Fc glycans | <i>G0/total subclass Fc glycans*100</i> |
| G1 | | 1325,524 | 884,019 | 1309,529 | 873,356 | 1317,527 | 878,687 | Fraction of A2G1 glycan in total subclass Fc glycans | <i>G1/total subclass Fc glycans*100</i> |
| G2 | | 1406,551 | 938,036 | 1390,556 | 927,373 | 1398,553 | 932,705 | Fraction of A2G2 glycan in total subclass Fc glycans | <i>G2/total subclass Fc glycans*100</i> |
| G0N | | 1346,038 | 897,694 | 1330,043 | 887,031 | 1338,04 | 892,363 | Fraction of A2B glycan in total subclass Fc glycans | <i>G0N/total subclass Fc glycans*100</i> |
| G1N | | 1427,064 | 951,712 | 1411,069 | 941,049 | 1419,067 | 946,38 | Fraction of A2BG1 glycan in total subclass Fc glycans | <i>G1N/total subclass Fc glycans*100</i> |
| G2N | | 1508,090 | 1005,730 | 1492,096 | 995,066 | 1500,093 | 1000,398 | Fraction of A2BG2 glycan in total subclass Fc glycans | <i>G2N/total subclass Fc glycans*100</i> |
| G1S1 | | 1471,072 | 981,051 | 1455,077 | 970,387 | 1463,075 | 975,719 | Fraction of A2G1S1 glycan in total subclass Fc glycans | <i>G1S1/total subclass Fc glycans*100</i> |
| G2S1 | | 1552,098 | 1035,068 | 1536,104 | 1024,405 | 1544,101 | 1029,737 | Fraction of A2G2S1 glycan in total subclass Fc glycans | <i>G2S1/total subclass Fc glycans*100</i> |
| G1NS1 | | 1572,612 | 1048,744 | 1556,617 | 1038,081 | 1564,614 | 1043,412 | Fraction of A2BG1S1 glycan in total subclass Fc glycans | <i>G1NS1/total subclass Fc glycans*100</i> |
| G2NS1 | | 1653,638 | 1102,761 | 1637,643 | 1092,098 | 1645,641 | 1097,43 | Fraction of A2BG2S1 glycan in total subclass Fc glycans | <i>G2NS1/total subclass Fc glycans*100</i> |

¹Glycan composition: N (N-acetylglucosamine), F (fucose), G (galactose) and S (N-acetylneuraminic acid) followed by a number representing the number and type of monosaccharides attached to A2 glycan.

²Glycan structures are drawn in GlycoWorkbench version 2. blue square = N-acetylglucosamine, red triangle = fucose, green circle = mannose, yellow circle = galactose, purple diamond = N-acetylneuraminic acid.

³IgG1 tryptic peptide sequence carrying glycan: E293EQYNSTYR301

⁴IgG4 tryptic peptide sequence carrying glycan: E293EQFNSTFR301

⁵IgG2&3 tryptic peptide sequence carrying glycan: E293EQFNSTYR301

⁶total subclass Fc glycans = sum of all 20 glycopeptides in one IgG subclass

Supplementary Table 16. Description of derived subclass-specific Fc IgG glycan traits measured by LC-MS with mass list.

| Derived glycan trait | Derived trait description | Derived trait calculation |
|----------------------|--|--|
| Core fucosylation | Fraction of structures containing core fucose in subclass specific Fc glycans | G0F+G1F+G2F+G0FN+G1FN+G2FN+G1F S1+G2FS1+G1FNS1+G2FNS1 |
| Bisecting GlcNAc | Fraction of structures containing bisecting GlcNAc in subclass specific Fc glycans | G0FN+G1FN+G2FN+G1FNS1+G2FNS1+ G0N+G1N+G2N+G1NS1+G2NS1 |
| Agalactosylation | Fraction of agalactosylated structures in subclass specific Fc glycans | G0+G0F+G0N+G0FN |
| Monogalactosylation | Fraction of structures containing one galactose in subclass specific Fc glycans | G1F+G1FN+G1FS1+G1FNS1+G1+G1N+G 1S1+G1NS1 |
| Digalactosylation | Fraction of structures containing two galactoses in subclass specific Fc glycans | G2F+G2FN+G2FS1+G2FNS1+G2+G2N+G 2S1+G2NS1 |
| Sialylation | Fraction of structures containing sialic acid in subclass specific Fc glycans | G1FS1+G2FS1+G1FNS1+G2FNS1+G1S1+ G2S1+G1NS1+G2NS1 |

Supplementary Table 17. Correlations between derived glycan traits measured by HILIC-UPLC.

| | Agalactosylation | Monogalactosylation | Digalactosylation | Bisecting GlcNAc | Sialylation | Core fucosylation |
|---------------------|------------------|---------------------|-------------------|------------------|-------------|-------------------|
| Agalactosylation | 1,00 | -0,23 | -0,89 | 0,38 | -0,70 | 0,17 |
| Monogalactosylation | -0,23 | 1,00 | 0,21 | -0,07 | -0,41 | 0,56 |
| Digalactosylation | -0,89 | 0,21 | 1,00 | -0,39 | 0,43 | 0,04 |
| Bisecting GlcNAc | 0,38 | -0,07 | -0,39 | 1,00 | -0,23 | -0,09 |
| Sialylation | -0,70 | -0,41 | 0,43 | -0,23 | 1,00 | -0,63 |
| Core fucosylation | 0,17 | 0,56 | 0,04 | -0,09 | -0,63 | 1,00 |

Supplementary Table 18. Correlations between derived subclass-specific Fc IgG glycan traits measured by LC-MS.

| | | IgG4_Bisecting |
|--------------------------|--|----------------|
| | IgG4_Sialylation | |
| | IgG4_Digalactosylation | |
| | IgG4_Monogalactosylation | |
| | IgG4_Agalactosylation | |
| | IgG2_Bisecting | |
| | IgG2_Sialylation | |
| | IgG2_Digalactosylation | |
| | IgG2_Monogalactosylation | |
| | IgG2_Agalactosylation | |
| | IgG1_Bisecting | |
| | IgG2_Agalactosylation | |
| | IgG2_Monogalactosylation | |
| | IgG2_Digalactosylation | |
| | IgG2_Sialylation | |
| | IgG2_Bisecting | |
| | IgG4_Agalactosylation | |
| | IgG4_Monogalactosylation | |
| | IgG4_Digalactosylation | |
| | IgG4_Sialylation | |
| | IgG4_Bisecting | |
| IgG1_Agalactosylation | 1,00 -0,64 -0,90 -0,62 0,26 0,81 -0,63 -0,76 -0,57 0,21 0,67 -0,39 -0,62 -0,47 | 0,22 |
| IgG1_Monogalactosylation | -0,64 1,00 0,47 -0,09 0,00 -0,42 0,65 0,29 0,08 -0,02 -0,33 0,50 0,22 0,02 | -0,06 |
| IgG1_Digalactosylation | -0,90 0,47 1,00 0,46 -0,30 -0,81 0,56 0,87 0,53 -0,26 -0,62 0,31 0,67 0,40 | -0,23 |
| IgG1_Sialylation | -0,62 -0,09 0,46 1,00 -0,26 -0,51 0,15 0,47 0,62 -0,17 -0,50 0,04 0,44 0,60 | -0,19 |
| IgG1_Bisecting | 0,26 0,00 -0,30 -0,26 1,00 0,28 -0,15 -0,31 -0,23 0,84 0,28 -0,12 -0,31 -0,19 | 0,60 |
| IgG2_Agalactosylation | 0,81 -0,42 -0,81 -0,51 0,28 1,00 -0,72 -0,93 -0,77 0,28 0,61 -0,30 -0,60 -0,46 | 0,24 |
| IgG2_Monogalactosylation | -0,63 0,65 0,56 0,15 -0,15 -0,72 1,00 0,56 0,20 -0,17 -0,44 0,41 0,39 0,18 | -0,12 |
| IgG2_Digalactosylation | -0,76 0,29 0,87 0,47 -0,31 -0,93 0,56 1,00 0,68 -0,28 -0,58 0,23 0,64 0,43 | -0,23 |
| IgG2_Sialylation | -0,57 0,08 0,53 0,62 -0,23 -0,77 0,20 0,68 1,00 -0,22 -0,46 0,10 0,42 0,50 | -0,23 |
| IgG2_Bisecting | 0,21 -0,02 -0,26 -0,17 0,84 0,28 -0,17 -0,28 -0,22 1,00 0,23 -0,11 -0,25 -0,15 | 0,58 |
| IgG4_Agalactosylation | 0,67 -0,33 -0,62 -0,50 0,28 0,61 -0,44 -0,58 -0,46 0,23 1,00 -0,56 -0,91 -0,75 | 0,25 |
| IgG4_Monogalactosylation | -0,39 0,50 0,31 0,04 -0,12 -0,30 0,41 0,23 0,10 -0,11 -0,56 1,00 0,34 -0,06 | -0,39 |
| IgG4_Digalactosylation | -0,62 0,22 0,67 0,44 -0,31 -0,60 0,39 0,64 0,42 -0,25 -0,91 0,34 1,00 0,69 | -0,19 |
| IgG4_Sialylation | -0,47 0,02 0,40 0,60 -0,19 -0,46 0,18 0,43 0,50 -0,15 -0,75 -0,06 0,69 1,00 | 0,00 |
| IgG4_Bisecting | 0,22 -0,06 -0,23 -0,19 0,60 0,24 -0,12 -0,23 -0,23 0,58 0,25 -0,39 -0,19 0,00 | 1,00 |