

Supplementary Material

Identification of Cell Culture Factors Influencing Afucosylation Levels in Monoclonal Antibodies by Partial Least-Squares Regression and Variable Importance Metrics

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Table S1. List of definitions for upstream culture levers considered in the PLSR models.

| INDEPENDENT FACTOR | DEFINITION | MOLECULES |
|-----------------------------|--|--|
| ADDITIVE 1 BOLUS CONC. | The concentration of the additive 1 bolus addition. | mAb- γ |
| ADDITIVE 1 BOLUS DAY | The day of the addition of the additive 1 bolus to the bioreactor. | mAb- γ |
| ADDITIVE 1 FEED | The concentration of the additive 1 feed. | mAb- γ |
| ADDITIVE 2 CONC. | The concentration of the additive 2 supplementation. | mAb- δ |
| BASAL AMMONIA | The real concentration of ammonia in the basal media prior to inoculation. | mAb- γ |
| BASAL GLUCOSE CONC. | The real concentration of glucose in the basal media prior to inoculation. | mAb- α , mAb- β , mAb- γ , mAb- δ |
| BASAL GLUTAMATE CONC. | The real concentration of glutamate in the basal media prior to inoculation. | mAb- γ |
| BASAL GLUTAMINE CONC. | The real concentration of glutamine in the basal media prior to inoculation. | mAb- γ |
| BASAL NA ⁺ CONC. | The real concentration of sodium ions in the basal media prior to inoculation. | mAb- γ |
| BASAL OSMOLALITY | The real osmolality of the basal media prior to inoculation. | mAb- α , mAb- β , mAb- γ , mAb- δ |
| COPPER SUPP. CONC. | The concentration of the copper sulfate supplementation. | mAb- δ |
| DAILY FEED % | The percent of the feed added each day. | mAb- α , mAb- β , mAb- γ |
| FEED AMMONIA | The real concentration of ammonia in the feed media prior to addition to the bioreactor. | mAb- γ |
| FEED GLUCOSE CONC. | The real concentration of glucose in the feed media prior to addition to the bioreactor. | mAb- α , mAb- β , mAb- γ , mAb- δ |
| FEED GLUTAMATE CONC. | The real concentration of glutamate in the feed media prior to addition to the bioreactor. | mAb- γ |

| | | |
|-------------------------------------|--|--|
| FEED GLUTAMINE CONC. | The real concentration of glutamine in the feed media prior to addition to the bioreactor. | mAb- γ |
| FEED NA⁺ CONC. | The real concentration of sodium ions in the feed media prior to addition to the bioreactor. | mAb- γ |
| FEED OSMOLALITY | The real osmolality of the feed media prior to addition to the bioreactor. | mAb- α , mAb- β , mAb- γ , mAb- δ |
| FEED PH | The real pH of the feed media prior to addition to the bioreactor. | mAb- α , mAb- β , mAb- γ , mAb- δ |
| FEED START DAY | The start day to begin feeding the bioreactor after the initiation of the culture. | mAb- δ |
| GLUCOSE SETPOINT | The midpoint between the glucose action limit and glucose supplement limit. | mAb- δ |
| PH LOWER BOUND | The lower bound for the pH control range of the bioreactor. | mAb- γ |
| PH SETPOINT | The target pH at the outset of the bioreactor culture that the pH control range can be defined relative to as \pm pH units. | mAb- α , mAb- β , mAb- γ |
| PH UPPER BOUND | The upper bound for the pH control range of the bioreactor. | |
| SEEDING DENSITY | The cell density of the bioreactor culture following inoculation. | mAb- α , mAb- β , mAb- γ , mAb- δ |
| TEMPERATURE SETPOINT | The initial target temperature at the outset of the bioreactor culture. | mAb- α , mAb- β |
| TEMPERATURE SHIFT DIFFERENCE | The difference between the initial culture temperature setpoint and the new culture temperature setpoint following the application of the temperature shift. | mAb- α |

Table S2. Summary of the final PLSR model metrics for each of the molecules (Afuc refers to afucosylation, Mann refers to mannose).

| Model ID | LVs | R ² | RMSEC (% afuc) | Relative RMSEC (%) | RMSECV (% afuc) |
|-------------------------------|-----|----------------|----------------|--------------------|-----------------|
| mAb- α Total Afuc | 2 | 0.63 | 0.60 | 9.38 | 0.81 |
| mAb- α High Mann | 2 | 0.72 | 0.53 | 8.20 | 0.70 |
| mAb- α Complex Afuc | 2 | 0.79 | 0.46 | 12.56 | 0.66 |
| mAb- β Total Afuc | 3 | 0.66 | 0.58 | 8.82 | 0.95 |
| mAb- β High Mann | 4 | 0.66 | 0.58 | 8.26 | 1.56 |
| mAb- β Complex Afuc | 4 | 0.70 | 0.54 | 12.45 | 1.09 |
| mAb- γ Total Afuc | 4 | 0.82 | 0.42 | 4.07 | 0.83 |
| mAb- γ High Mann | 4 | 0.84 | 0.40 | 5.15 | 0.83 |
| mAb- γ | 6 | 0.76 | 0.48 | 12.74 | 0.96 |

| | | | | | |
|---------------|---|------|------|-------|------|
| Complex Afuc | | | | | |
| mAb- δ | 3 | 0.32 | 0.82 | 20.45 | 0.96 |
| Total Afuc | | | | | |
| mAb- δ | 3 | 0.78 | 0.46 | 17.21 | 0.55 |
| High Mann | | | | | |
| mAb- δ | 3 | 0.75 | 0.49 | 12.36 | 0.59 |
| Complex Afuc | | | | | |

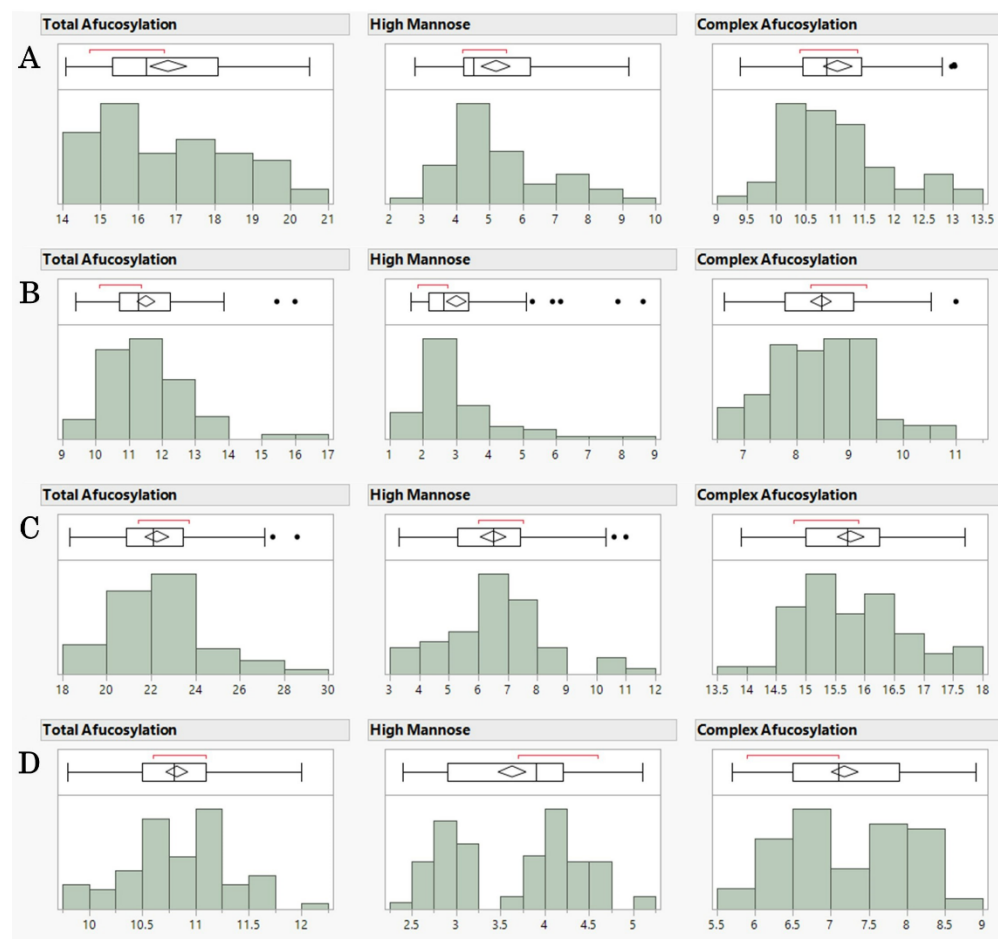


Figure S1. Distribution for the different afucosylation categories for the molecules (A) mAb- α , (B) mAb- β , (C) mAb- γ , and (D) mAb- δ . All figures were generated from JMP (Version 15.2.0, SAS Institute Inc., Cary, NC).