Relative viscosity screening of monoclonal antibody (mAb) formulations at low concentration and low viscosity for early stage developability



Introduction

The use of therapeutic antibodies is one of the fastest growing sectors for novel biopharmaceuticals, and the industry is under increasing pressure to deliver new therapies in a timely and safe manner. Biopharmaceutical development processes are lengthy and complex with very specific analytical challenges, and particularly severely limited sample availability at the start of the pipeline. Against this constraint, the ability to assess biophysical parameters as key quality attributes as early as possible in candidate validation/early formulation development stages can reduce costs and save time, and minimize risk from prompt identification of potential problem candidates.

High viscosity issues are problematic both during bioprocessing and patient administration. The ability to screen early stage candidates and formulations for abnormal viscosity changes, using measurements at low concentrations and very low sample volumes, can help identify monoclonal antibodies (mAbs) with potential developability issues. Degradation of mAbs most commonly occurs through self-association¹, and aggregation can cause changes to the viscosity of formulations².

By setting target viscosity-concentration profiles, and using an automated relative viscosity screen with high resolution and high reproducibility at low protein concentrations, mAbs with abnormally high viscosity increases can be identified using only a small amount of sample. Lavoisier and Schlaeppi³ reported that mAbs showing abnormally high viscosity profiles with concentration could be sorted from those with low viscosities at concentration as low as 25mg/mL.

Malvern Viscosizer TD is an automated biophysical characterization tool utilizing Taylor Dispersion Analysis and providing ultra-low volume, solution-



based molecular size and stability measurement capabilities, combined with microcapillary Poiseuille flow for relative viscosity assessment.

Relative viscosity measurements are performed by injecting a sample plug of 6μ l (typically) into the capillary under constant drive pressure. The sample plug edges are detected at two fixed windows along the capillary, simply by Refractive Index change, and the time between windows is measured (Δt_s). The sample time is then referenced against the time taken for a solvent or buffer of known viscosity (e.g. water) to travel between the windows (Δt_r) to enable the viscosity of the sample to be calculated. Automatic procedures are used to clean the microcapillary and repeat measurements. Samples are held in a temperature-controlled autosampler prior to measurement.

The following work details measurements of a mAb in various formulation buffers, and with the addition of common excipients, to show the small variations in viscosity with concentration and excipient changes that the microcapillary flow technique is able to reliably measure.

Experimental

- Solutions of citrate, histidine and phosphate buffers were freshly prepared as the stock solutions
- The pH of each of these stock solutions was measured (SevenEasy pH meter, Mettler-Toledo) – see Table 1

Buffer	Measured pH
Citrate	4.035
Histidine	7.763
Phosphate	7.225

Table 1 pH measurements of buffer solutions

- Two common excipients were also used in the study:
- 1. 0.5M sucrose (Sigma-Aldrich, UK) a protein stabilizer at the concentration of
- 2. 0.05% Tween 20 (Fluka, UK) a surfactant used to help prevent surface and stress-induced aggregation ^{4,5}
- Samples were made up via the use of dialysis buttons (Hampton Research, US) to dialyze 13.5mg/mL monoclonal antibodies (mAbs) in each buffer, and to provide the samples as shown in Table 2

mAbs Concentration (mg/mL)	Excipient Added
13.5	None
5	None
1	None
1	0.5 M sucrose
1	0.05% Tween 20

Table 2 Sample concentrations for each buffer plus additional excipients

- Relative viscosity measurements were performed on a Malvern Viscosizer TD using an uncoated capillary (default measurement settings for Viscosity were used throughout); a solution of 1mg/mL caffeine in water was used as a reference
- A 0.1M NaOH wash step was used to minimize capillary interactions
- All samples were stored in the autosampler at 4°C prior to testing; measurement temperatures for each sample were 5°C, 25°C and 37°C

Results and Discussion

Figures 1-3 are radar charts for the three buffers (citrate, histidine and phosphate) showing the change in viscosity for different concentration mAb solutions, and at 1mg/mL with the addition of two excipients (as detailed in Table 2), at the three measurement temperatures.

These plots clearly show the expected change in viscosity with temperature, with the viscosity of each sample increasing as the temperature decreases.

Figures 4 is a bar chart showing the differences in mean viscosity for mAbs samples at the three different concentrations in each buffer at 25°C.

Figure 5 is a bar chart showing the differences in mean viscosity for mAbs samples at 1mg/mL concentration in each buffer and with the addition of sucrose and Tween 20 as excipients, at 25°C.

The error bars on each of these charts show the relative standard deviation of the measurements for each sample (from 5 replicates), and demonstrates the ability of the microcapillary flow technique used in Viscosizer TD to produce high resolution and highly reproducible data for relative viscosity measurements at low viscosities.

Citrate Buffer

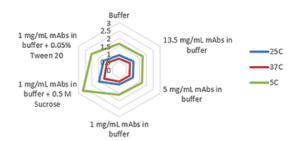


Figure 1 Radar chart of citrate buffer sample viscosity results measured at each run temperature (viscosity units are cP (or mPas))

Histidine Buffer

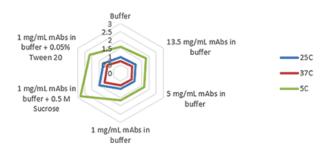


Figure 2 Radar chart of histidine buffer sample viscosity results measured at each run temperature (viscosity units are cP (or mPas))

Phosphate Buffer

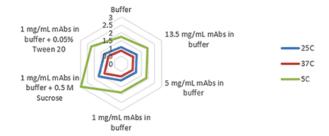


Figure 3 Radar chart of phosphate buffer sample viscosity results measured at each run temperature (viscosity units are cP (or mPas))

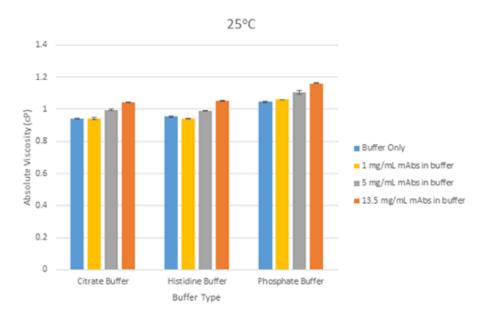


Figure 4 Bar chart showing showing the differences in mean viscosity for mAbs samples at three different concentrations, grouped by their buffer, at a measurement temperature of 25°C (error bars show the relative standard deviation of the measured viscosities)

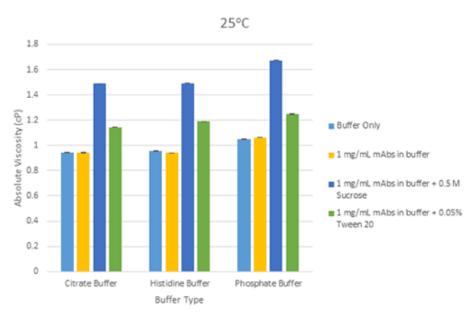


Figure 5 Bar chart showing the differences in mean viscosity for mAbs samples at 1mg/mL concentration, grouped by their buffer and with the addition of sucrose and Tween 20 as excipients, at a measurement temperature of 25°C (error bars show the relative standard deviation of the measured viscosities)

Figure 6 shows data at 25°C for citrate and phosphate buffers plotted against mAb concentration, and the expected linear increase in viscosity at low concentrations i.e. the protein molecules are isolated entities in solution at these concentrations. As mAb concentration increases to high concentrations, the protein molecules in solution will start to interfere with each other and become

entangled, causing an exponential increase in viscosity (and non-Newtonian behavior).

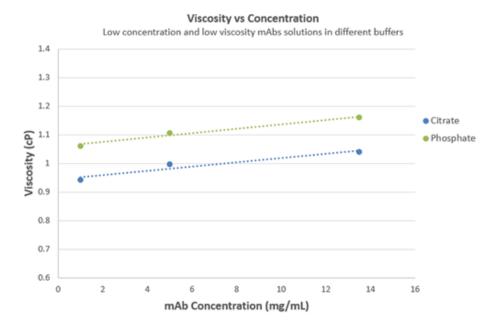


Figure 6 Plot of viscosity against concentration for mAb solution in citrate and phosphate buffers at 25°C

The increase is viscosity for 1mg/mL mAb solutions containing excipients (sucrose and Tween 20) is mainly due to the additive viscosity in the solution from the excipient. The same relative viscosity screening principle can be followed for formulations containing excipients to enable abnormally high viscosity responses to be identified, which typically arise from conformational change and/or self-association of the protein molecule in the solution conditions under evaluation.

Conclusion

The relative viscosity methodology used in the Viscosizer TD has demonstrated its ability to discern small differences between low viscosity protein formulations and give very repeatable results.

These differences include the effect concentration of sample in various buffers, the addition of excipients and the effect temperature can have on the viscosity.

These attributes are particularly well suited to early developability stages of the biopharmaceutical pipeline, given the added benefits of automation and ultra-low sample volume requirements.

By setting appropriate working limits on viscosity-concentration profiles for mAbs systems, relative viscosity screening using a high resolution technique with high reproducibility is able to highlight deviations from acceptable viscosity behavior at

low concentrations, to identify potential problem candidates as early as possible in the development pipeline.

References

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