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Title: Evaluation of additives on reversed-phase chromatography of monoclonal antibodies using a 1000 angstrom stationary phase.

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Keywords: monoclonal antibodies; mAbs; reversed-phase LC; additives.

Corresponding Author: Professor David Victor McCalley, PhD

Corresponding Author's Institution: University of the West of England, Faculty of Health and Life Sciences

First Author: David Victor McCalley, PhD

Order of Authors: David Victor McCalley, PhD; Davy Guillarme, PhD

Abstract: A wide pore (1000 Å) diphenyl stationary phase was evaluated for the analysis of monoclonal antibodies (mAbs), comparing a conventional mobile phase of acetonitrile-water containing overall 0.1 % trifluoroacetic acid (TFA) with a similar mobile phase incorporating in addition 5 % butanol. Alternatively, TFA was replaced by ammonium formate (AF) buffer (pH 3.0) and by methane sulfonic acid. Addition of 5% butanol to the mobile phase reduces the minimum temperature at which suitable UV analysis of the mAbs can be obtained from about 70 oC with TFA alone to about 60 oC thus potentially improving column lifetime and reducing the possibility of sample degradation. AF buffers produce satisfactory UV sensitivity at 70 oC and have the advantage of reducing signal suppression in mass spectrometry (MS). Some peak tailing was noted in comparison with TFA separations. Methane sulfonic acid at the same molar concentration as TFA produced the best chromatographic peaks, maintaining reasonable UV sensitivity down to 50 oC, also giving acceptable results even at only 3 mM concentration of the additive. The good results with this additive were attributed to its stronger acidity and consequent suppression of the ionisation of column silanols. Surprisingly, peak response (as measured by the size of the peaks) was rather poorly correlated with the peak capacity of the gradient analysis. A possible explanation is self-deactivation of active column sites by a portion of the sample.

- Diphenyl bonded 1000 Å column evaluated for separation of monoclonal antibodies.
- Rituximab and Bevacizumab selected as difficult probes.
- Column was tested with trifluoroacetic acid and a variety of additives.
- Ammonium formate is mass spectrometer friendly, with acceptable results at 70 °C.
- Methanesulfonic acid gives best peak capacity, and reasonable results at 50 °C.

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Evaluation of additives on reversed-phase chromatography of monoclonal antibodies using
a 1000Å stationary phase.

David V. McCalley*

* Corresponding author.

Centre for Research in Biosciences, University of the West of England, Frenchay, Bristol
BS16 1QY, UK

Tel. 0044 1173287353

Email David.Mccalley@uwe.ac.uk

Davy Guillarme,

School of Pharmaceutical sciences, University of Geneva,

Institute of Pharmaceutical Sciences of Western Switzerland

Rue Michel Servet, 1, 12011 Geneva 4, Switzerland.

Keywords: monoclonal antibodies; mAbs; reversed phase LC; acid additives.

36 **Abstract**

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40 incorporating in addition 5 % butanol. Alternatively, TFA was replaced by ammonium
41 formate (AF) buffer (pH 3.0) and by methane sulfonic acid. Addition of 5% butanol to the
42 mobile phase reduces the minimum temperature at which suitable UV analysis of the
43 mAbs can be obtained from about 70 °C with TFA alone to about 60 °C thus potentially
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48 the best chromatographic peaks, maintaining reasonable UV sensitivity down to 50 °C,
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52 of the peaks) was rather poorly correlated with the peak capacity of the gradient analysis.
53 A possible explanation is self-deactivation of active column sites by a portion of the
54 sample.

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59 **1. Introduction.**

60 Monoclonal antibodies (mAbs) represent an increasingly large proportion of world-wide
61 pharmaceutical sales, which for some compounds are in the range of billions of dollars per
62 annum. As with all pharmaceuticals, quality control of formulations is essential. In addition,
63 many mAbs are coming off-patent, and much effort has been expended on the preparation
64 of biosimilars, which have equivalent properties to the original drug, or biobetters that may
65 have improved efficacy and safety [1]. Careful assessment of these materials e.g. for
66 comparability with the original drug, is required. Characterisation may involve a number of
67 approaches and techniques such as separation by charge (e.g. by ion-exchange
68 chromatography, capillary electrophoresis); hydrophobicity/hydrophilicity (e.g. RPLC,
69 hydrophilic interaction chromatography); size (size exclusion chromatography); and affinity
70 (affinity chromatography) [2]. Some of these techniques can be combined to give high
71 resolution 2-dimensional separations of the proteins [3, 4].

72 Much information can be derived by cleaving mAbs with specific reagents to
73 produce characteristic sub-units that can be more easily analysed. In addition, the whole
74 protein can also be digested with enzymes such as trypsin to give peptides that
75 characterise the protein. However, there is interest in the separation of intact mAbs by
76 RPLC (and other LC separation mechanisms), as the structure and properties of the entire
77 drug are retained [5, 6]. Superficially porous ("shell") particles would seem an ideal choice
78 for the analysis of these large molecules (MW~150,000 Da) as mass transfer in and out of
79 the porous fraction of these particles is facilitated by the small diffusion distance required.
80 This factor, while especially beneficial for large molecules due to their low diffusion
81 coefficients, may not be important for small molecule compounds (MW<500 Da) where
82 other considerations, possibly the better packing qualities of shell compared with totally
83 porous materials may be more significant [7]. However, it is necessary to use materials
84 with sufficient pore size to accommodate these large species-otherwise size exclusion
85 effects could dominate the separation. According to Guiochon and co-workers, the pore
86 size of the stationary phase should be at least three times the protein size e.g. at least 300
87 Å for immunoglobulins [8]. Kirkland and co-workers [9] recommended the use of shell
88 particles with 1000 Å pores for the analysis of mAbs. Such large pore size materials also
89 prevent the hindered diffusion that may be experienced with smaller pore materials,
90 leading to peak shape problems. The authors reported narrower peak widths for the Sigma
91 test mAb on a 2.7 µm 1000 Å C4 phase compared with a 1.7 µm 300 Å C4 phase, using

92 an ACN gradient with 0.1% difluoroacetic acid as additive. The improvement was
93 attributed to mass transport limitations in the smaller pore size material. For small
94 molecules analysed in the isocratic mode, the large pore C4 phase also gave good
95 performance with reduced plate height $h \sim 2.1$. Fekete and co-workers [10] recently
96 evaluated a $2.7 \mu\text{m}$ 450 \AA pore size high coverage triphenyl phase (Waters) for the
97 analysis of mAbs using mainly 0.1 % trifluoroacetic acid (TFA) as additive in an ACN –
98 water gradient. The authors stressed the importance of using elevated temperatures (80-
99 $90 \text{ }^\circ\text{C}$) to obtain good “recoveries” of intact proteins. (Recovery was measured as the
100 response at a given temperature compared with that at the highest temperature used, 90
101 $^\circ\text{C}$) . At lower temperatures, proteins may adsorb too strongly onto the stationary phase,
102 giving poor peak shapes and above all, very limited sensitivity. Conversely, elevated
103 temperatures may result in sample degradation and also negatively affect the column
104 lifetime. The authors therefore emphasised the importance of inclusion of temperature as a
105 variable in column evaluation studies. It is well-known that elevated temperature can lead
106 to improvements in separation for proteins beyond those attainable for small molecules.
107 Raising the temperature may also reduce the number of protein conformers (likely by
108 denaturing the protein into its unfolded state) hence reducing peak broadening [6, 11] .

109 It has been known for many years that TFA produces good chromatographic results
110 for the separation of peptides and proteins [12], although it is possible that other additives
111 might give chromatographic improvements over TFA. With TFA, ion pair formation with
112 basic groups on side chains enhances the hydrophobic character of the peptide or protein,
113 enhancing suitability for RP analysis [13]. Ion pairing may also reduce ionic interactions of
114 the solute with ionised silanols. However, as the ion pairs may not be broken, signal
115 suppression may result in electrospray mass spectrometry. Furthermore, surface tension
116 effects and high conductivity of the eluent may cause spray instability. In the end, TFA
117 may lead to sensitivity loss up to an order of magnitude compared with more favourable
118 methods [14]. In some cases, post-column addition of counteracting agents (e.g. propionic
119 acid) has been advocated to improve sensitivity [15]. Formic acid can give reasonable
120 peak shapes for peptides while maintaining good MS detection properties, but its
121 chromatographic performance for large proteins is doubtful, leading to significant peak
122 broadening or even irreversible adsorption compared with use of TFA [10, 16]. Another
123 problem however with TFA is that restoring the LCMS system to highest sensitivity
124 operation after use may take a considerable period of flushing (e.g. 24 hours with 50:50
125 ACN/water containing 1% acetic acid [17]. In this context, the use of ammonium formate

126 (AF) buffers adjusted to acid pH with FA may provide a compromise, giving acceptable
127 chromatographic performance while maintaining good MS sensitivity. A 5 fold average
128 increase in sensitivity was reported for therapeutic proteins and their fragments [12, 13]
129 using AF buffers at pH 3 compared with TFA.

130 The aims of the present study were:

- 131 a) To evaluate a new shell 1000 Å pore size diphenyl stationary phase, comparing its
132 performance for mAbs with a C4 phase synthesised on the same base material.
- 133 b) To compare the performance of different mobile phases on the diphenyl phase, with
134 the aim of obtaining the best chromatographic results at the lowest possible
135 temperature.
- 136 c) To determine whether it is possible to improve chromatographic performance for
137 mAbs using alternatives to 0.1 % TFA.

138

139 **2. Experimental**

140 Experiments were performed using a 1290 ultra-high performance liquid chromatograph
141 (UHPLC, Agilent, Waldbronn, Germany) comprising a binary pump, autosampler and
142 photodiode array UV detector set at 280nm (0.6 µL flow cell). The gradient delay of the
143 UHPLC system was measured as 150 µL by running a gradient from 100 % water to 100
144 % water plus 10 ppm of a UV absorbing substance (uracil) using an empty tube (10 cm x
145 0.05mm internal diameter) to generate a backpressure for normal pump performance. At a
146 flow of 0.4 mL/ min. all peaks with a retention of > 0.375 min (thus all the peaks in the
147 present study) emerged under gradient (rather than isocratic) conditions. Sample injection
148 volume was 2 µL and flow rate was 0.4 mL/min. ACN (gradient UV grade), formic acid,
149 methanesulfonic acid (MSA), trifluoroacetic acid (TFA) and ammonium formate (AF) were
150 purchased from Fisher (Loughborough, U.K.). Rituximab (Mabthera) and Bevacizumab
151 (Avastin) were obtained from the School of Pharmaceutical Sciences, University of
152 Geneva. Columns used were Halo diphenyl and Halo C4 (10 x 0.21 cm, 2.7 µm particle
153 size, 1000 Å pore size (Advanced Materials Technology, Wilmington, USA). Peak capacity
154 in gradient elution was determined using the equation:

155

$$156 P = 1 + t_G/1.699w_{0.5}$$

157

158 where t_G is the gradient time, and $w_{0.5}$ the peak width at half height [12]. This equation was
159 used rather than its equivalent using peak width at base due to the difficulty of
160 reproducible measurement of this quantity at base. The mobile phase gradient was
161 typically 30:70 ACN-water to 45:55 ACN-water in 5 min. The concentration of the various
162 additives (e.g. 0.1 % TFA) was maintained constant throughout the gradient by inclusion in
163 both mobile phase reservoirs. An equilibration time of 15 min was allowed between each
164 run. mAb samples were prepared at a concentration of 500 mg/L dissolved in 0.1 %
165 aqueous TFA.

166

167 **3. Results and Discussion.**

168 *3.1 Column and antibody selection.*

169 We first evaluated two different 1000 Å stationary phases (diphenyl and C4) from the
170 same manufacturer for the determination of the mAbs Rituximab and Bevacizumab
171 (approx. 0.5 mg/mL concentration) using a 5 minutes gradient of 30-45 % ACN containing
172 0.1 % TFA (v/v) at 0.4 mL/min. From a previous work of Bobaly et al., Rituximab and
173 Bevacizumab produced the 20th and 21st lowest values of the chromatographic behaviour
174 parameter B_C among the 23 mAbs tested where

175

$$176 B_C = P_{rel} \times R_{rel}$$

177

178 P_{rel} is the peak capacity in gradient elution and R_{rel} the peak area both at 60 °C relative to
179 results obtained at 90 °C. Results were obtained using an ACN gradient with 0.1 % TFA as
180 the additive [16]. These two mAbs therefore represent amongst the most
181 chromatographically demanding probes of performance. As reported elsewhere, Rituximab
182 can give particular problems with ionic interactions with the stationary phase due to its high
183 pI of 9.4, whereas Bevacizumab can undergo strong hydrophobic interactions as illustrated
184 by its high retention in hydrophobic interaction chromatography [18, 19]. They are both
185 important biopharmaceutical products, each having current annual sales of > \$7 billion.
186 Rituximab is currently used for the treatment of non-Hodgkin's Lymphoma, whereas
187 Bevacizumab is used for ovarian, lung, colon and rectal cancers. With a column
188 temperature of 80 °C, reasonably good results with peaks of acceptable symmetry were
189 obtained both on the C4 (Fig.1) and diphenyl (Fig. 2a) columns. Note the Y axis scale was
190 constant with a maximum value of 90 mAU in each of Figures 1 and 2. However, column
191 lifetime is enhanced at lower temperatures, and degradation of mAbs may be reduced.
192 Less aggressive conditions are also a goal for mAb separations, as the protein may exist

193 fully or partially in a “native” state, retaining or partially retaining its 3D structure. Folded
194 proteins are more compact and the interior amino acids are less prone to be charged than
195 those at the surface. Thus “native” proteins have a narrower charge state envelope in
196 electrospray MS, meaning the signal intensity is distributed over fewer charged states,
197 potentially leading to increased sensitivity [20]. However, sufficiently non-aggressive
198 conditions to allow native protein analysis remain an ambition in RP separations. With the
199 diphenyl column, 70 °C appears to be an acceptable temperature in terms of response and
200 peak shape for both mAbs using TFA. In comparison this lower temperature gave reduced
201 response for both mAbs on the C4 column. At 60 °C, the diphenyl column gave reduced
202 sensitivity (only ~40 % of the response compared with higher temperatures), but the C4
203 column failed to elute Bevacizumab and gave only a very small peak for Rituximab. At 50
204 °C, the diphenyl column gave a very small response with both mAbs, while the C4 column
205 showed no response at all. Guillarme and co-workers also reported that higher
206 temperatures (85-90 °C) were necessary to achieve similar performance for intact mAbs
207 on a C4 column compared with a triphenyl column operated at 75 °C [10]. In view of its
208 superior performance, further work concentrated on the diphenyl phase.

209

210 *3.2 Evaluation of UV response with different additives.*

211 We chose peak height as a measure of response rather than peak area, as the former is
212 less susceptible to the presence of isoforms present in the standard mAbs, which
213 may/may not be completely separated from the main peak under the various conditions
214 used. For example, Fig. 2d (0.013 M MSA) shows the presence of several minor variants
215 in the chromatogram of Rituximab and Bevacizumab, whereas these peaks are merged
216 with the major peaks in Fig. 2c (5mM AF pH 3) preventing the assessment of these
217 isoforms. Use of peak area would unfortunately skew the results in favour of poorer
218 separations, as the peak area of the major peak would be increased by inclusion of these
219 isoforms. Note that while peak height varies considerably with retention in isocratic
220 analysis, peak height is approximately independent of retention time in gradient elution
221 when the flow rate, the column, the gradient time and the gradient programme are all
222 maintained constant, as is the case for all the results in the present study shown in Figs. 2-
223 4 [21].

224 Figure 2b shows the effect of incorporating 5 % n-butanol to each channel while
225 maintaining the concentration of TFA in each at 0.1% and the ACN gradient constant. It
226 has previously been suggested that incorporating additives such as propanol or butanol
227 into a TFA containing mobile phase can improve the response at lower temperatures. It is

228 possible that these alcohols may shield silanols from detrimental ionic interaction with the
229 protein [22] . Comparison with Fig 2a shows that addition of butanol clearly has some
230 beneficial effect, with greater response to both mAbs at 60 and 50 °C compared with use
231 of 0.1% TFA alone. Fig. 3a and 3b show plots of peak height obtained for Rituximab and
232 Bevacizumab respectively at temperatures from 50-80 °C using different mobile phase
233 additives. The graphs illustrate a more quantitative summary of the results under different
234 conditions compared with mere visual inspection of the chromatograms in Fig. 2. Fig. 3
235 shows more clearly that increases in peak height were obtained with butanol addition at 60
236 °C of ~65% and ~57 % for Rituximab and Bevacizumab respectively compared with use of
237 TFA alone. However this effect is rather limited as demonstrated by the poor response of
238 both mAbs at 50 °C, even in the butanol containing mobile phase.

239 In an effort to improve MS sensitivity, attempts have been made to replace some of
240 the mobile phase TFA content with FA, as FA alone gives poor chromatographic results
241 (see above). The presence of acidic silanols can result in pronounced (probably cationic)
242 adsorption of protein. Residual silanols are increasingly ionised at higher mobile phase pH.
243 The true thermodynamic s^s pH of 0.013M FA in 60 % ACN is around 3.5, whereas that of
244 the same molar concentration of TFA is about 2.0, and undoubtedly the milder acidity of
245 the former gives rise to more deleterious effects [23]. The very low ionic strength of
246 0.013M FA at 60 % ACN (~0.3 mM) compared with that of TFA (~11mM) may also lead to
247 greater possibilities of overloading with the former additive. Unfortunately, a study where
248 half of the TFA concentration was replaced with FA resulted in significant recovery
249 decrease for six representative mAbs [16], indicating that this approach was not profitable,
250 although greater success was obtained for the analysis of mAb sub-units. Similarly poor
251 results were found with FA alone in the analysis of basic peptides [12]. However, much
252 improved results were obtained when using ammonium formate buffers both for peptides
253 [12] and also for some proteins [13]. Thus, we attempted the separation using ammonium
254 formate buffers (AF, 5 mM w^w pH 3). We chose a low buffer concentration as it limits MS
255 signal suppression. It has also been reported that chromatographic peak shapes were
256 worse for mAbs at higher buffer concentration, perhaps due to salting out and irreversible
257 adsorption [13]. While AF buffers cannot allow exploration of the beneficial low pH region
258 provided by TFA, they at least solve the problem of low ionic strength of the mobile phase,
259 as the salt is completely dissociated even in 60 % ACN. Ammonium ions should also
260 compete beneficially with sample molecules for cationic sites on the column. Fig. 2c shows
261 that it is indeed possible to obtain reasonable results with AF/FA, without using fluorinated
262 acids that are detrimental to MS sensitivity. However, clearly some of the resolution of the

263 impurities/heterogeneities in the standards is lost when using AF/FA compared with TFA
264 (Fig 2a), with these various minor variants merging into the tail of the major peak. Fig. 3a
265 and 3b confirm that for AF/FA, there is some improvement in response by using 80 rather
266 than 70 °C. However, response drops off considerably at lower temperatures. For
267 Rituximab, the response using AF/FA at 60 °C was reduced to 29 % compared with that
268 obtained using the same mobile phase at 80 °C, whereas it was only 2 % for
269 Bevacizumab. At 50 °C, neither mAb could be detected using this mobile phase. Despite
270 this poorer behaviour at lower temperatures, over 80% of the average response for the two
271 mAbs was maintained using AF/FA compared with that using 0.1 % TFA when analysis
272 was carried out at 80 °C.

273 Methane sulfonic acid (MSA) has also been demonstrated as a useful additive in a
274 study of the separation of peptides. Kadar et al. [24] reported improvements in retention
275 and efficiency for peptides in HILIC analysis when compared with TFA. The improved
276 retention in HILIC was attributed to ion-pairing of peptide with MSA. MSA has slightly
277 stronger acidity than TFA at the same molar concentration even when the pH is measured
278 in the presence of high concentrations of ACN as used in HILIC separations, and provides
279 similar ionic strength to TFA [23]. Fig. 2d indicates this acid, when used at the same molar
280 concentration as 0.1% v/v TFA (0.013M), gave superior response for both Rituximab and
281 Bevacizumab. The superior response is particularly apparent at lower temperatures (50-60
282 °C) where peak shape and response is maintained to a greater extent compared with that
283 at higher temperatures using MSA. Fig. 3a indicates that the response for Rituximab at 60
284 and 50 °C using MSA was 92 and 44% of the response at 80 °C compared with
285 considerably lower values of 67 and 8 % for the same compound using 0.1% TFA/butanol.
286 Clearly, there are even greater gains in response using MSA compared with TFA alone as
287 additive. All of these results were mirrored using Bevacizumab (Fig. 3b). Note that the
288 selectivity for variants/isoforms shown using TFA was maintained when using MSA.

289 The present study uses exclusively UV detection. Indeed, for large proteins, the
290 sensitivity of UV detection is better than for MS detection (whereas the opposite is true for
291 the detection of small molecules or peptides) due to the large envelope of MS ions
292 produced from these large molecules (see above). Furthermore, MS detection in quality
293 control of biopharmaceuticals is rarely used due to complexity and cost. Particularly for
294 good sensitivity in MS detection however, the concentration of additives should be
295 maintained at the lowest possible value consistent with preservation of the
296 chromatographic peak shape. MSA was reported to be MS compatible [24], but at higher
297 concentrations led to signal suppression, at least in the HILIC mobile phases used in that

298 study which contained a high proportion of ACN. Clearly, further investigation of the MS
299 properties of MSA is required and will be the subject of a further study. Fig. 2e
300 demonstrates that good results can even be obtained with 0.0033M MSA i.e. at one
301 quarter of the previous concentration. Note that despite the low gradient retention times of
302 the peaks in Fig. 2e (1-2 min.), all peaks eluted under gradient conditions after the gradient
303 delay (0.375 min.-see Experimental), allowing comparisons to be made with the other
304 mobile phase conditions. Use of the more dilute MSA concentration resulted in a loss of
305 peak height and peak capacity of about 25% compared with 0.013M MSA, but it appears
306 that the good results can still be obtained at 60 °C using the lower concentration of acid.
307 Some small loss of selectivity for the isoforms can be seen at this lower concentration of
308 MSA.

309

310 *3.4 Effect of additive and temperature on Peak Capacity.*

311 Figs. 4a and 4b show the peak capacity of both mAbs (calculated for a gradient of 30-60 %
312 ACN in a time of 10 min.) using each of the 4 mobile phases over the same range of
313 temperatures (50-80 °C). Particularly for Bevacizumab, peak capacities are highest when
314 using MSA and lowest when using AF/FA. However, comparison of Figs. 3 and 4 does not
315 show a close correlation between peak response and peak capacity. For example,
316 Rituximab shows a rather small decrease in peak capacity at 60 °C compared with that at
317 80 °C, using AF/FA (Fig. 4a) but a much larger decrease in response (Fig. 3a).
318 Undoubtedly, results for peak capacity with AF/FA are optimistic, because peak widths
319 were recorded at half height rather than at the base, as the latter method has problems of
320 reproducibility. This means that the peak tailing evident in Fig. 2d is not taken into account.
321 However, observation of the chromatograms (e.g Fig. 2a using 0.1 % TFA alone) indicates
322 that peak shape is largely maintained as the temperature is lowered, even as the response
323 decreases considerably. It seems more likely that some fraction of the injected protein
324 sample becomes irreversibly adsorbed to the stationary phase when held at low
325 temperature, providing a more inert surface for the chromatography of the remaining
326 sample. Indeed it was observed that running a blank sample at 70-80 °C immediately after
327 a run at 50 °C using AF/FA resulted in elution of “ghost” peaks with surprisingly good
328 shape at the appropriate retention time (results not shown).

329 Fig. 5 shows the variation of peak capacity with gradient time over the range 5-60
330 min. using 0.013M MSA as additive and a column temperature of 70 °C. The limiting peak
331 capacity for both Rituximab and Bevacizumab appears to be of the order of ~300. At
332 higher gradient times, the experimental conditions approximate more closely to those in an

333 isocratic separation, with similar band broadening processes involved, which explains the
334 levelling off shown.

335

336 **4. Conclusions**

337 A 1000 Å diphenyl phase was found to give good results for the analysis of monoclonal
338 antibodies using a gradient of 0.1 % TFA in acetonitrile. The two commercial mAbs
339 Rituximab and Bevacizumab were chosen as probes, because they are generally
340 considered to be difficult to analyse as they undergo strong ionic and hydrophobic
341 interaction respectively with RP materials. Investigations of various alternative additives
342 were performed with the objective of obtaining the best chromatographic performance at
343 the lowest possible temperature. Lower temperatures increase the lifetime of the column
344 and reduce the possibility of sample degradation. Addition of butanol to the 0.1% TFA
345 mobile phase lowers the useful analysis temperature from about 70 °C to 60 °C; possibly
346 the alcohol blocks active column sites. AF buffers, which are known to give better MS
347 detection properties than TFA, produce reasonable chromatographic results at 70 °C or
348 above, although peak shape was slightly inferior to those obtained with TFA. More
349 significant chromatographic benefits can be obtained by use of methanesulfonic acid
350 which reduces the feasible analysis temperature to around 50 °C, even when low
351 concentrations of additive are utilised (~3 mM). The stronger acidity of MSA may be
352 beneficial in reducing detrimental ionised silanol interactions. Further studies are
353 necessary to assess the MS compatibility of this additive. Surprisingly, peak capacity (a
354 measure of peak narrowness in gradient elution) correlated rather poorly with response (as
355 measured by the size of peaks) when using any of the different additives. It is possible that
356 a portion of the sample is adsorbed in active sites at low temperature, allowing some sort
357 of self-deactivation of the column surface.

358

359

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362 instrument, and thank Advanced Materials Technology (Wilmington, USA) for the gift of
363 columns used in this work.

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367 **6. Legend to Figures**

368 Fig. 1 Separation of Rituximab (first eluted) and Bevacizumab on wide pore C4 column.
369 Mobile phase gradient 30-45 % ACN containing 0.1 % TFA. Column temperature from 50-
370 80 °C. Gradient time t_G =5 min. Flow rate 0.4 mL/min. Injection volume: 2 μ L. UV detection
371 at 280 nm.

372 Fig. 2 Separation of mAbs on wide pore diphenyl column using additives in 30-45 % ACN
373 gradient, t_G =5 min. Mobile phases: a) 0.1 % TFA; b) 0.1 % TFA and 5% butanol; c) 5mM
374 AF pH3 ; d) 0.013 M MSA; e) 0.0033M MSA. Column temperature, flow rate, injection
375 volume, detection wavelength as Fig. 1

376 Fig. 3 a) Effect of temperature (Black 80 °C; Red 70 °C; Green 60 °C; Blue 50 °C) on
377 response of Rituximab using 4 different mobile phases as in Figs 2a-2d.b) Same for
378 Bevacizumab. Column temperature, flow rate, injection volume, detection wavelength as
379 Fig. 1.

380 Fig. 4 a) Effect of temperature on peak capacity using 4 different mobile phases (as Figs.
381 2a-d) for Rituximab. b) Same for Bevacizumab. Column temperature, flow rate, injection
382 volume, detection wavelength as Fig.1.

383 Fig. 5 Effect of gradient time on peak capacity for diphenyl column. Column temperature
384 70 °C. Mobile phase gradient 30-60 % ACN 0.013M MSA with gradient times 5-60 min.
385 Column temperature, flow rate, injection volume, detection wavelength as Fig.1.

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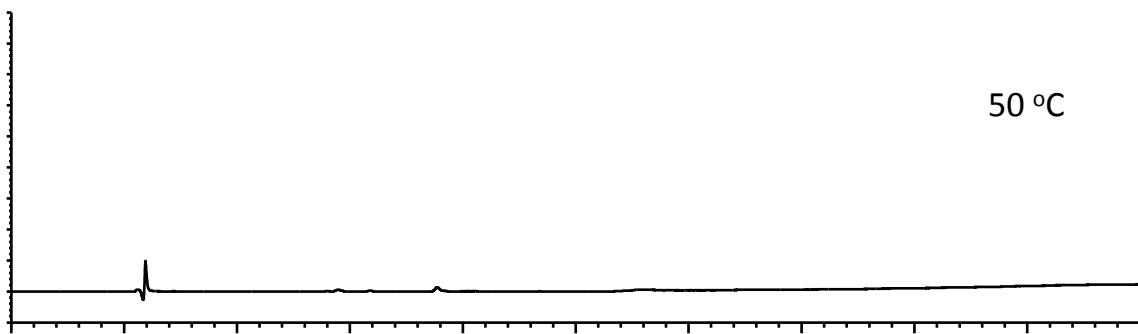
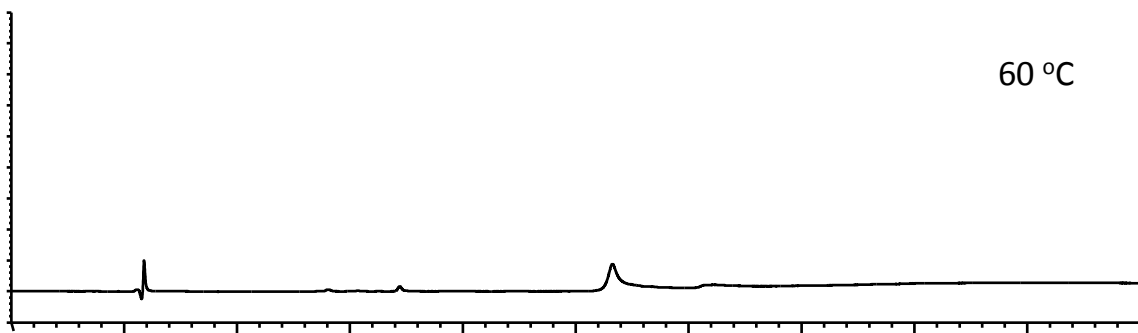
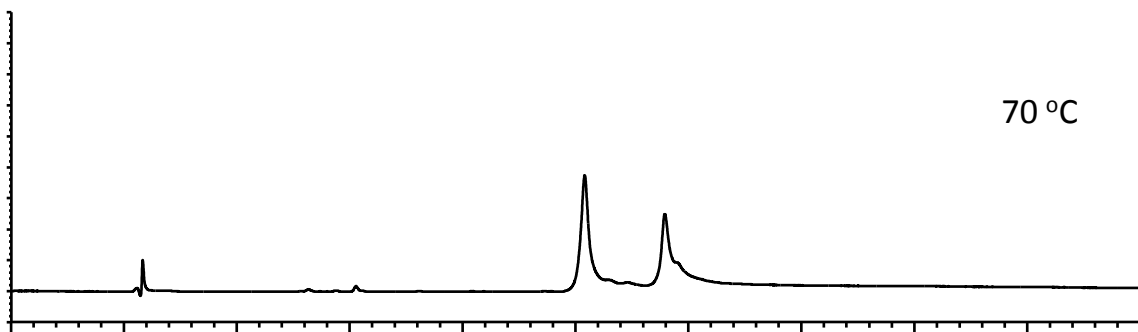
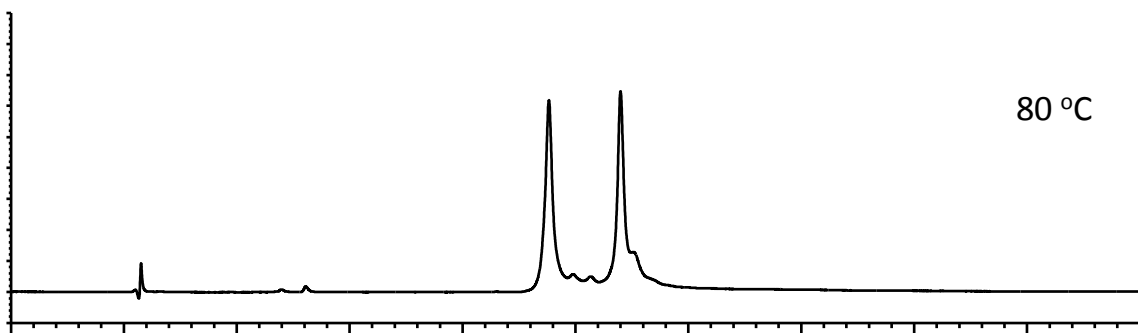
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- 463

Figure 1

Fig. 1

90 mAU

C4 0.1% TFA



5 min.

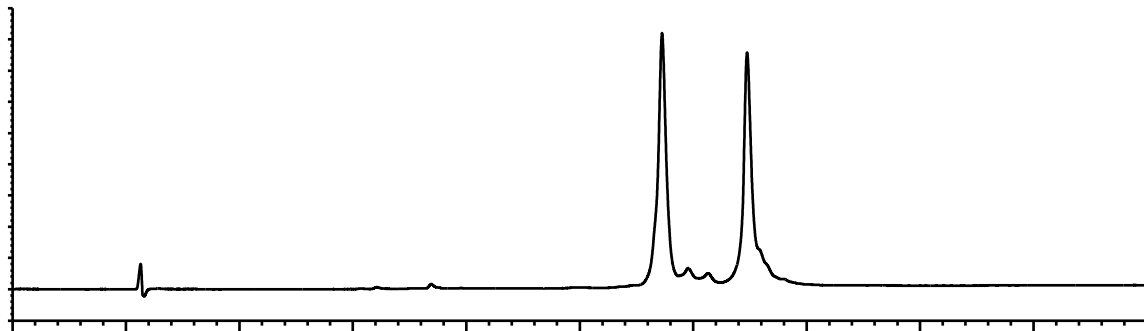
Figure 2a

Fig. 2a

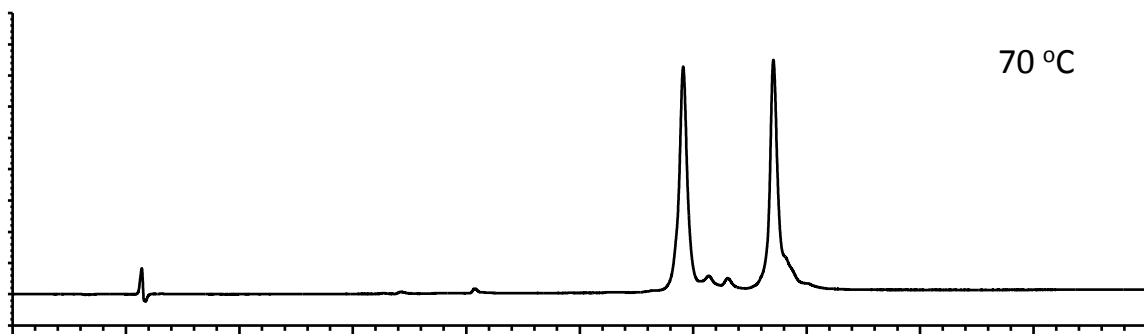
Diphen 0.1 % TFA

90 mAU

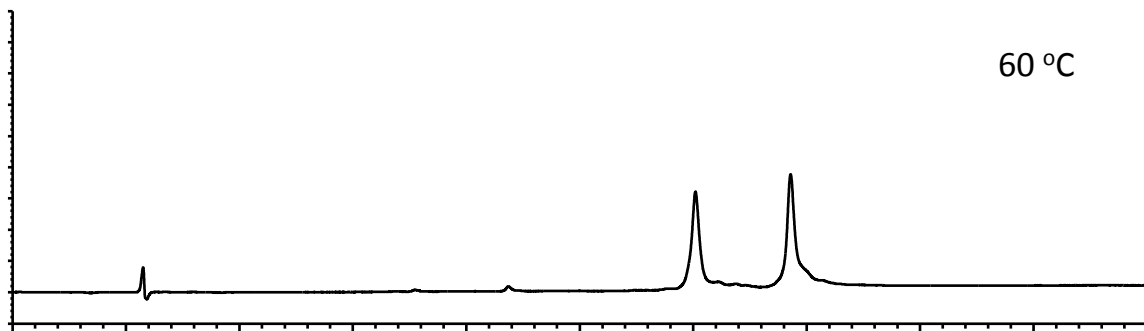
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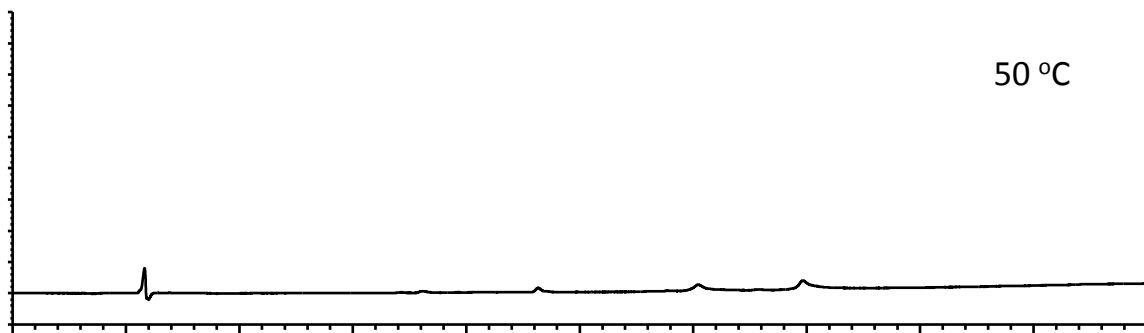
70 °C



60 °C



50 °C



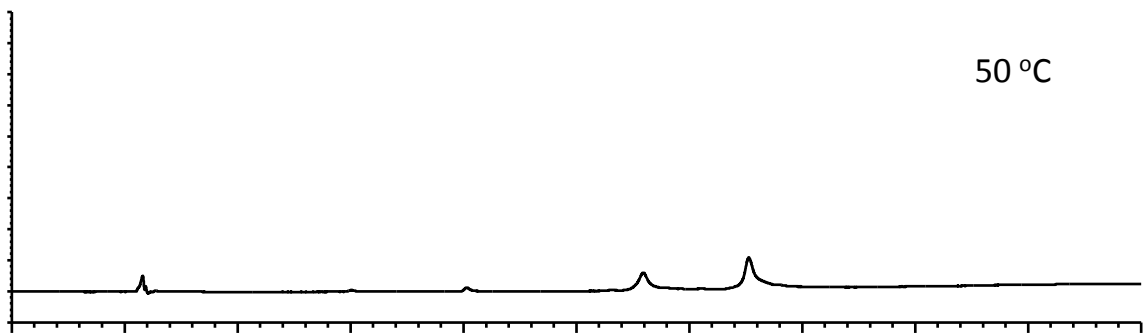
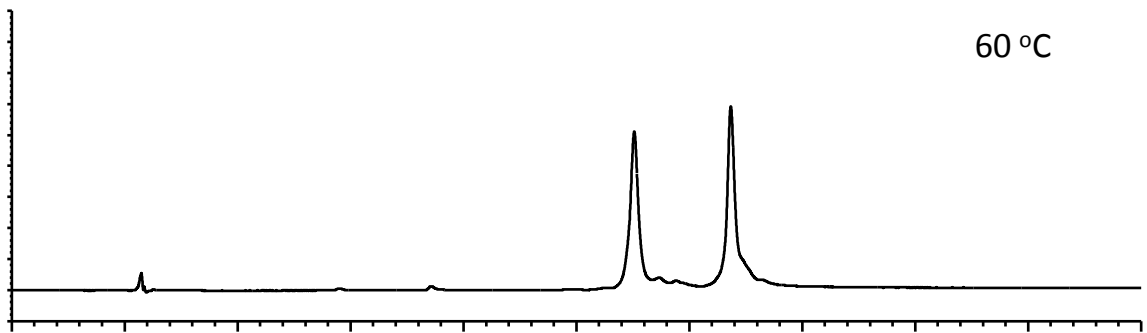
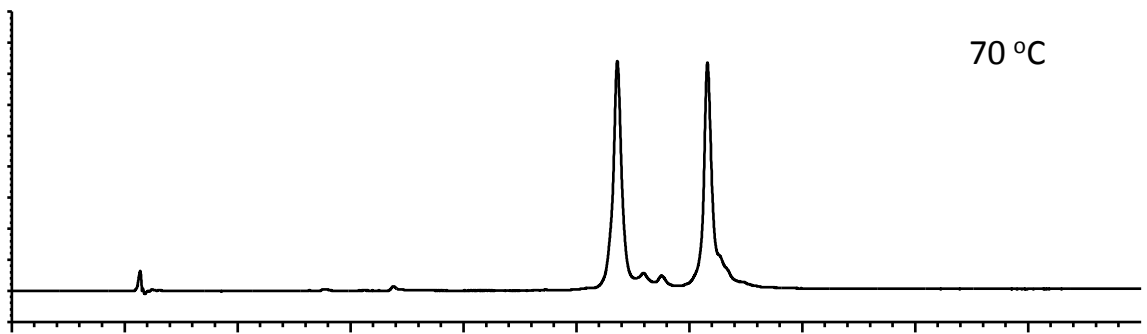
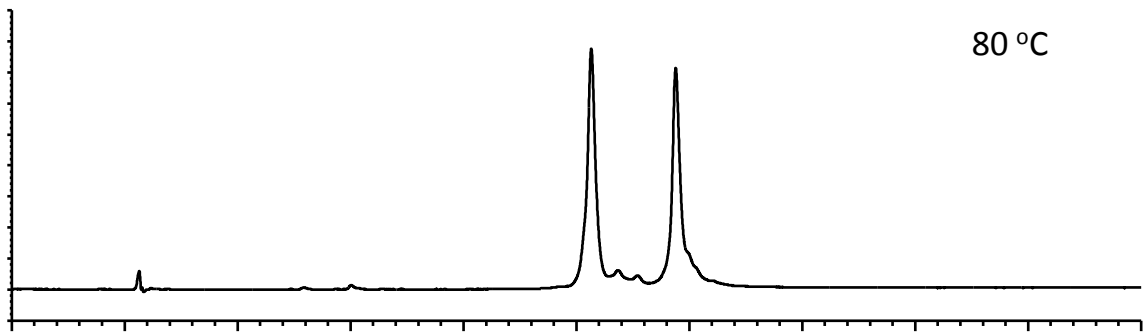
5 min.

Figure 2b

Fig. 2b

5% butanol 0.1 % TFA

90 mAU

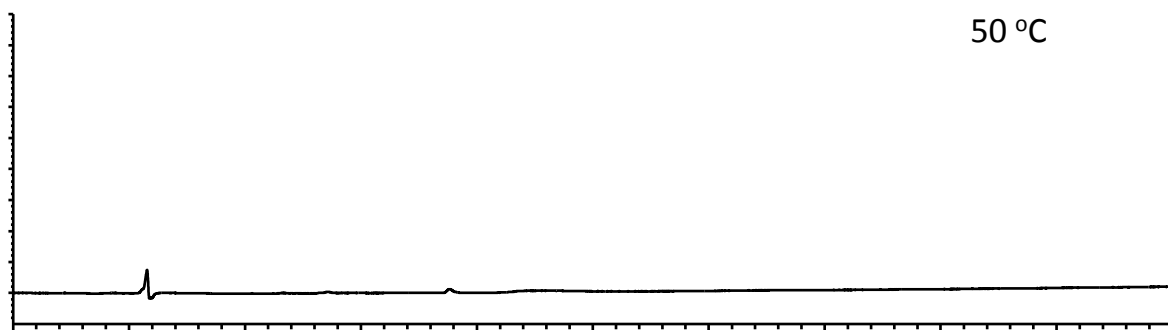
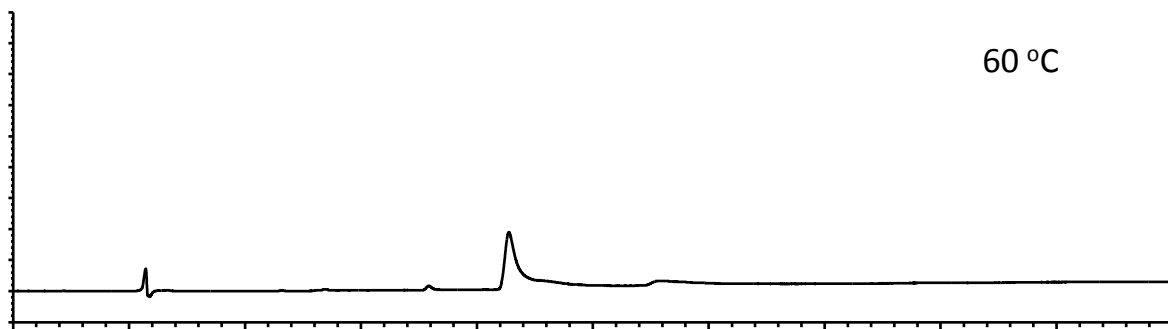
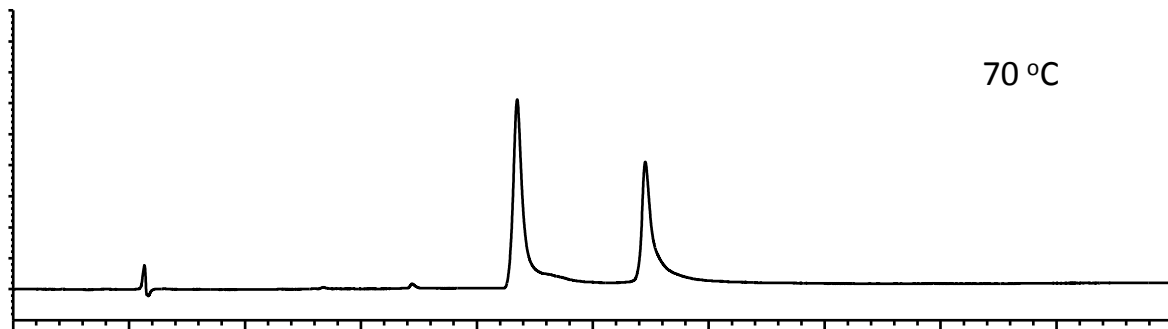
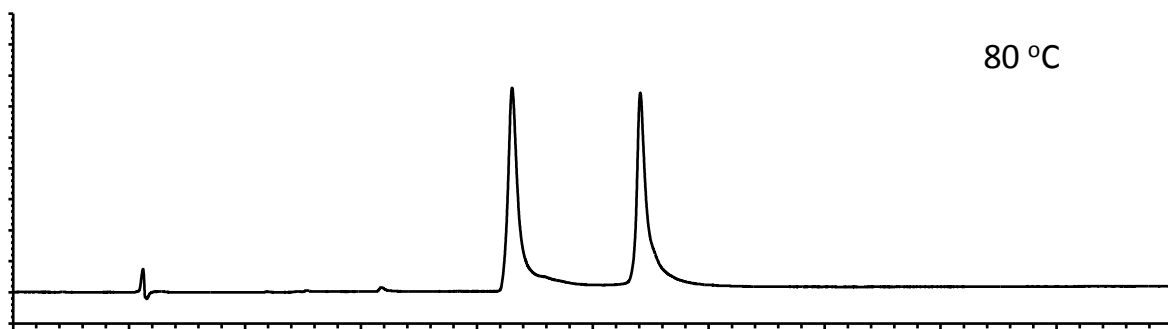


5 min.

Fig. 2c

5mM AF pH 3

90 mAU



5 min.

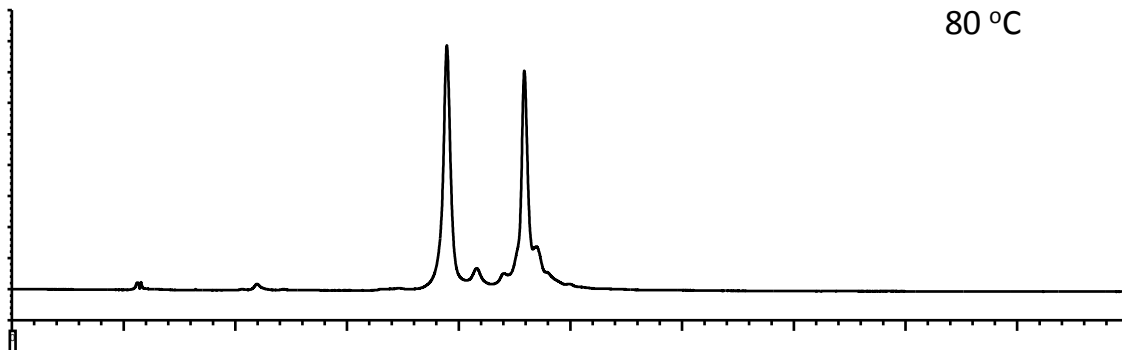
Figure 2d

Fig. 2d

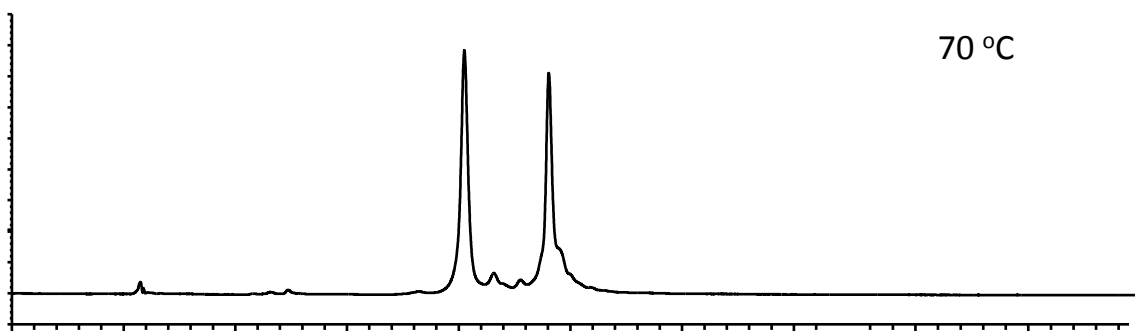
0.013M MSA

90 mAU

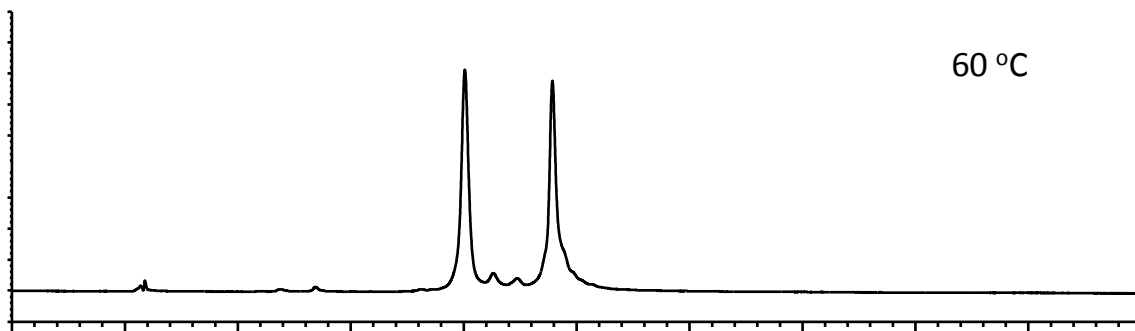
80 °C



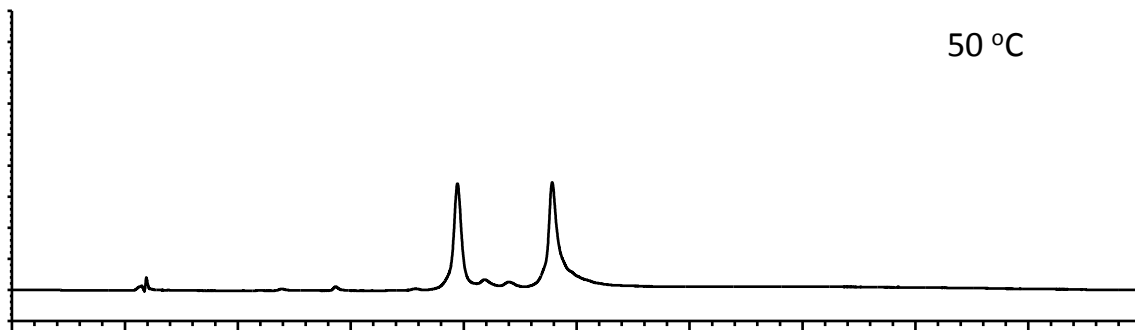
70 °C



60 °C



50 °C

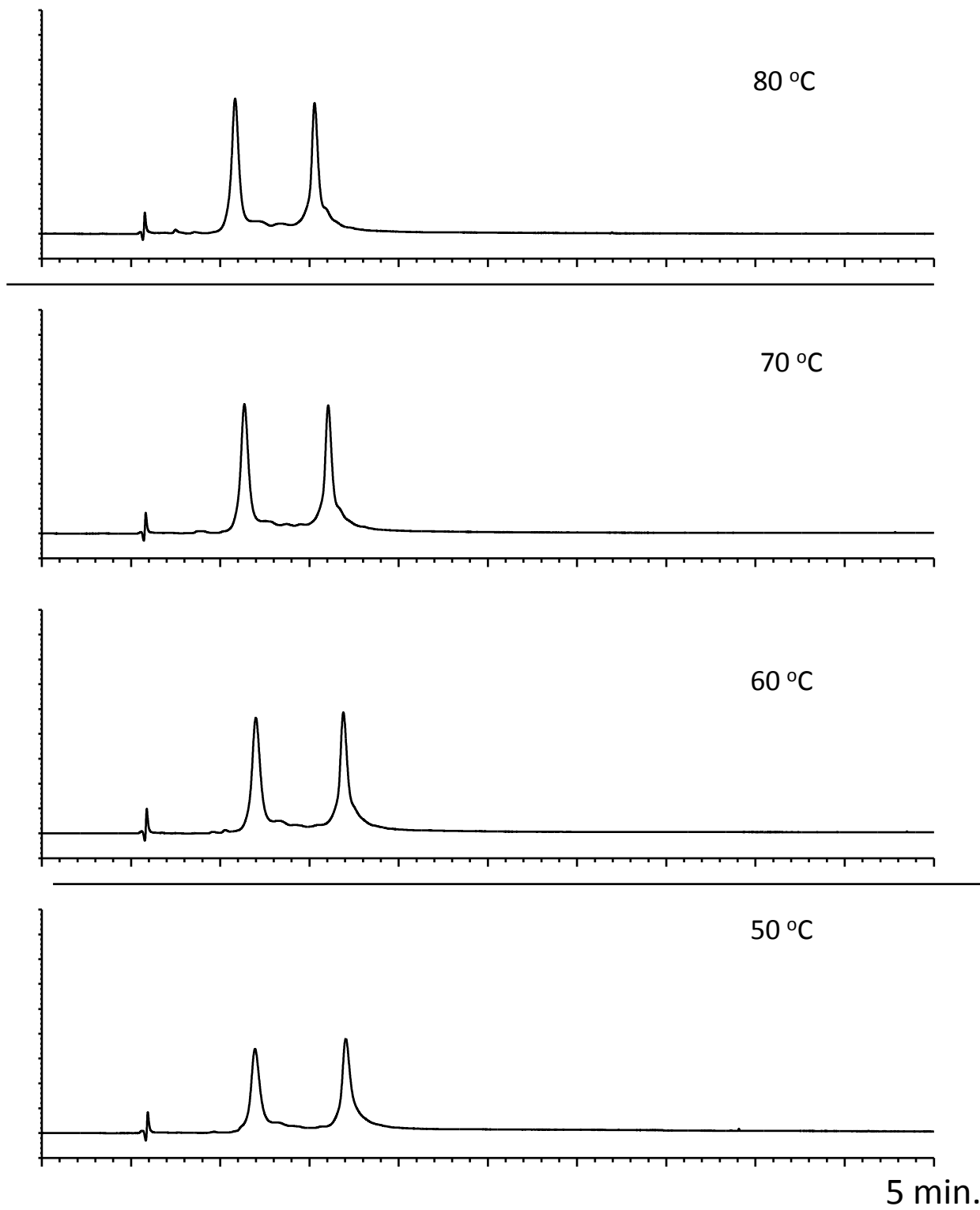


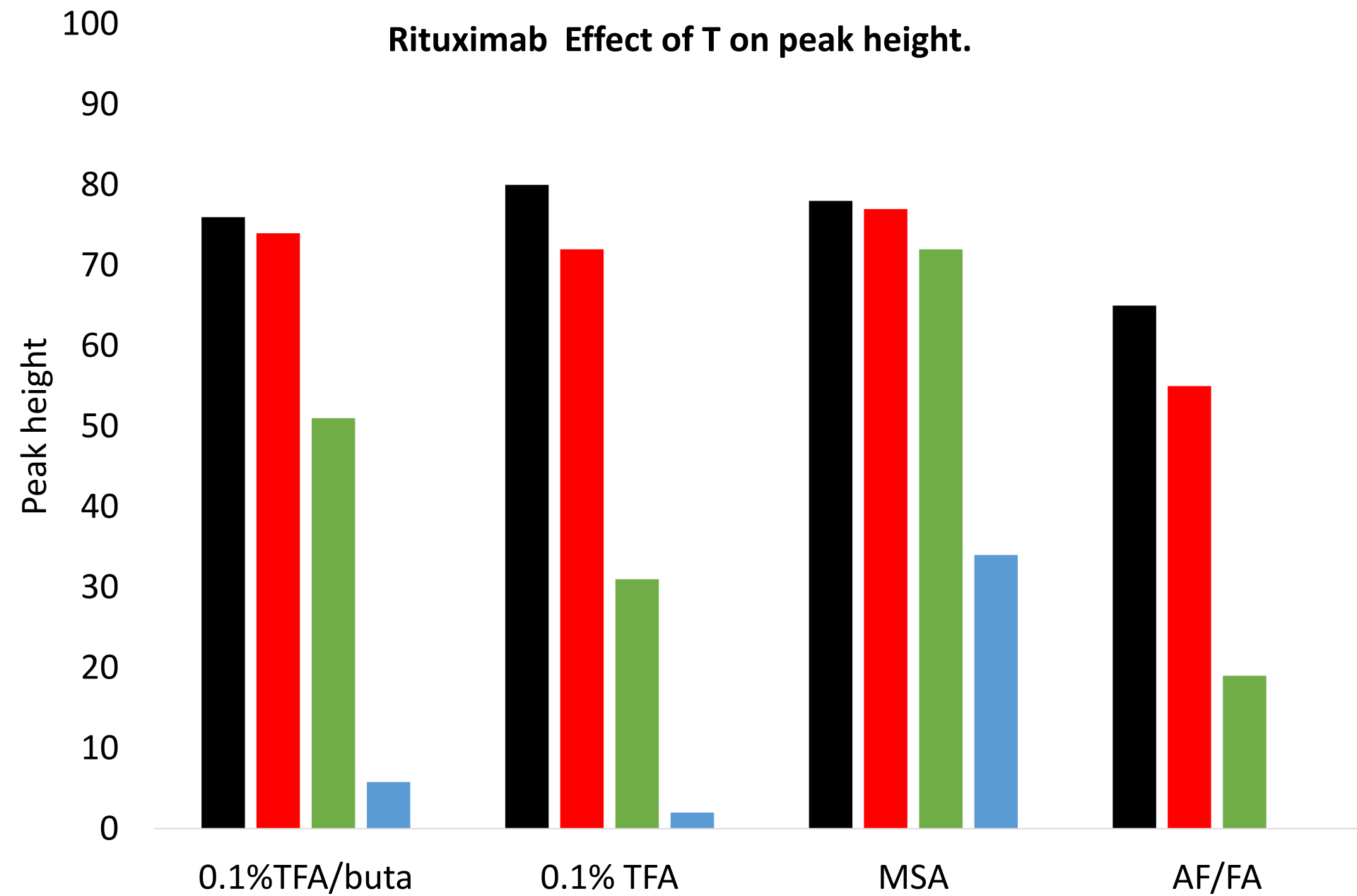
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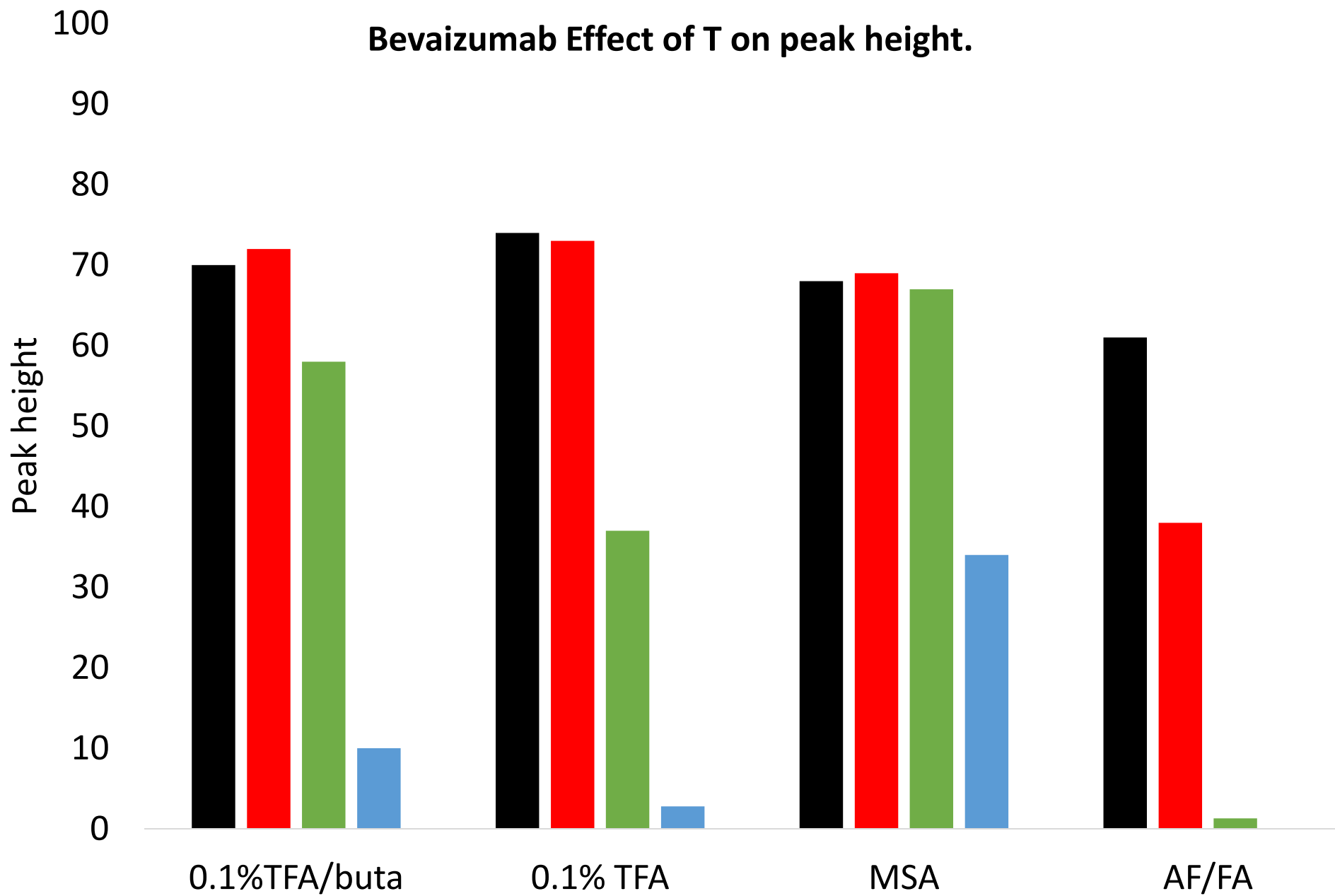
Fig. 2e

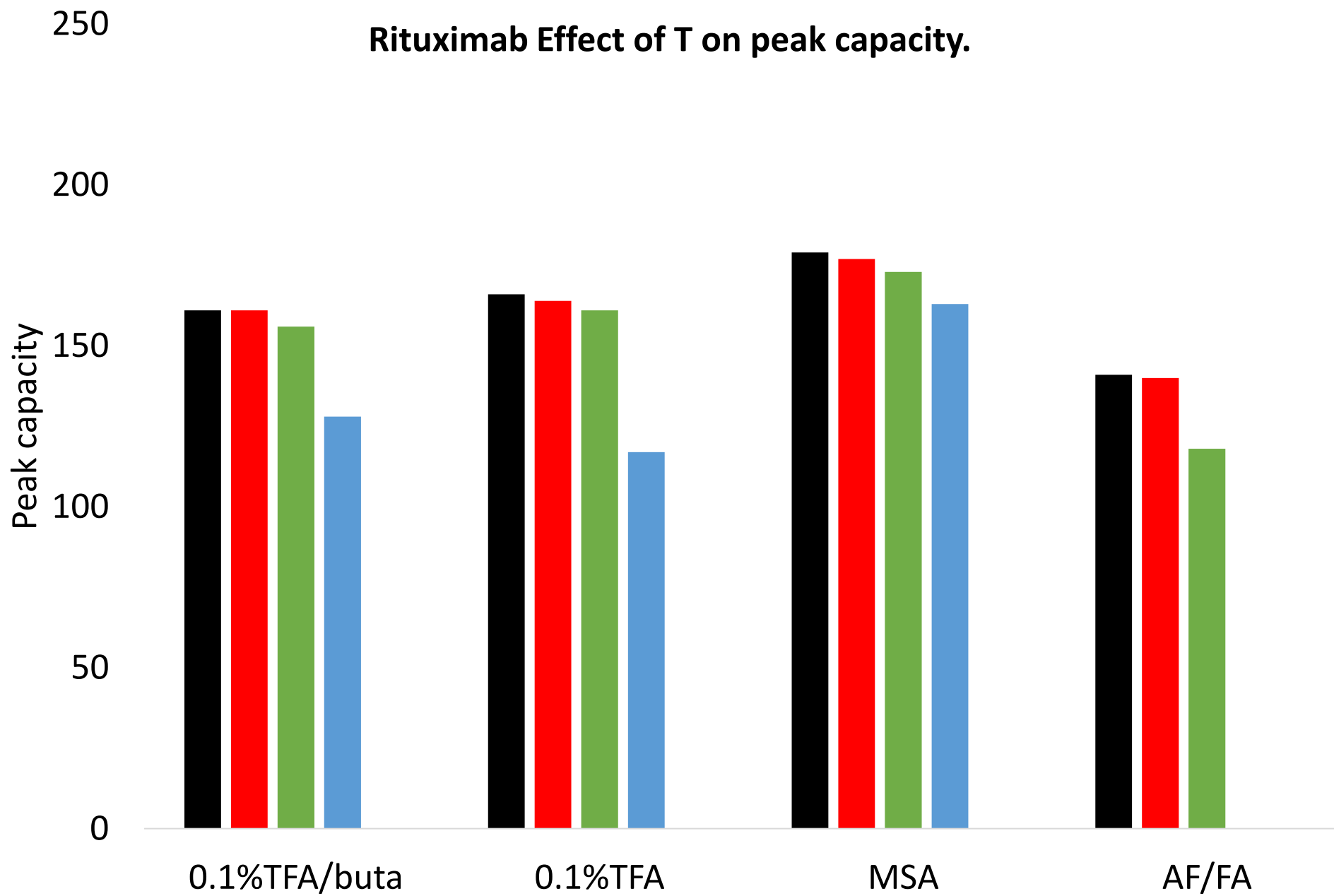
90 mAU

Diphen 0.0033M MSA









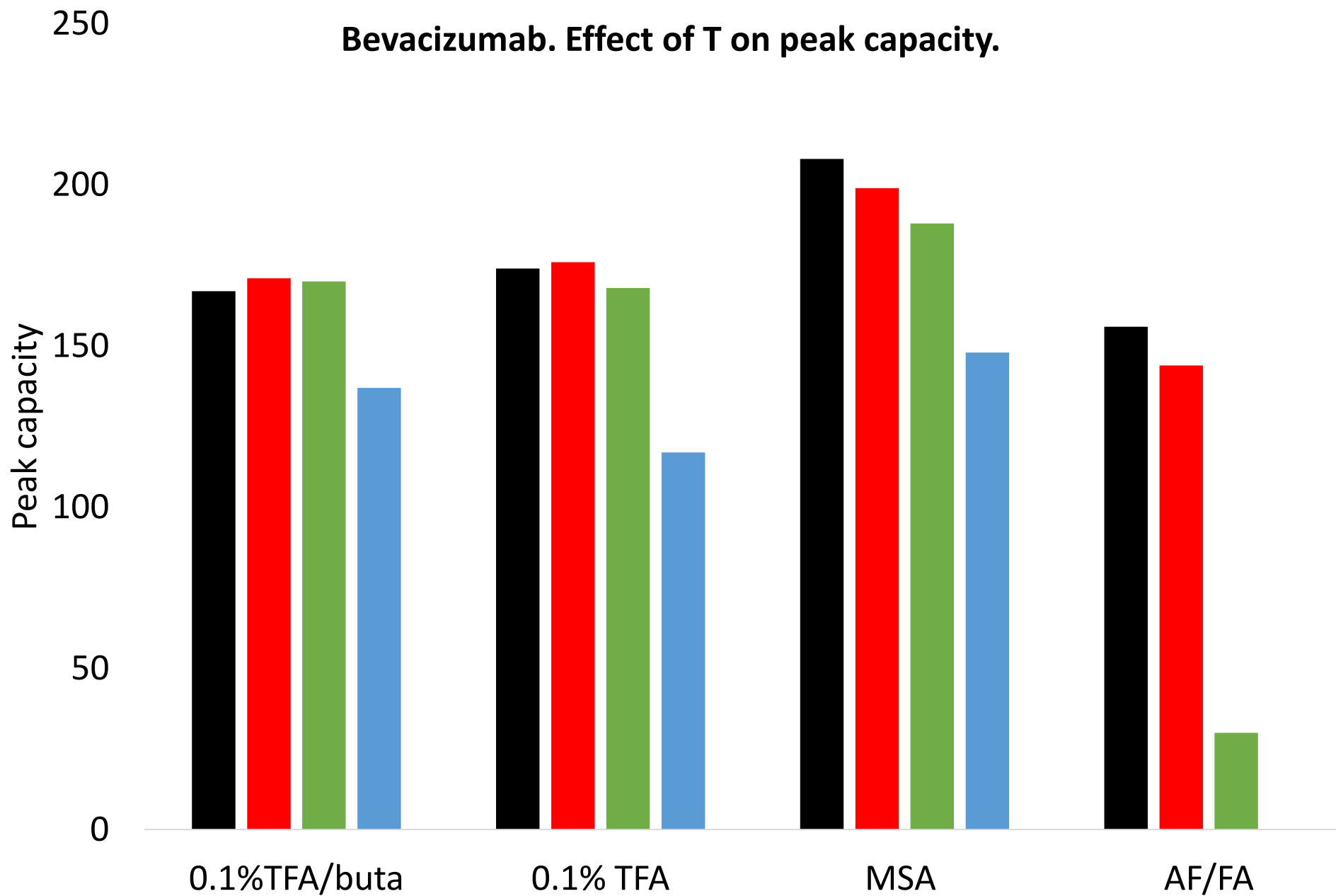
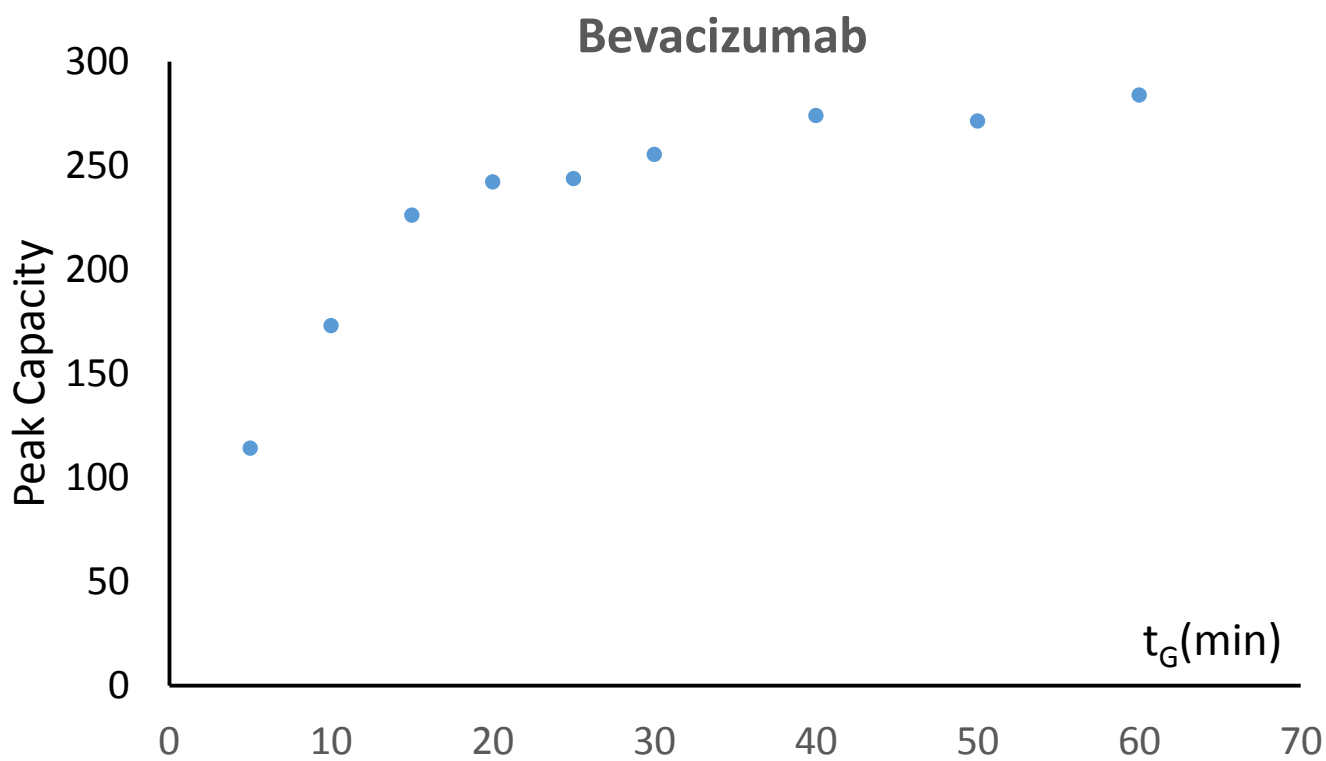
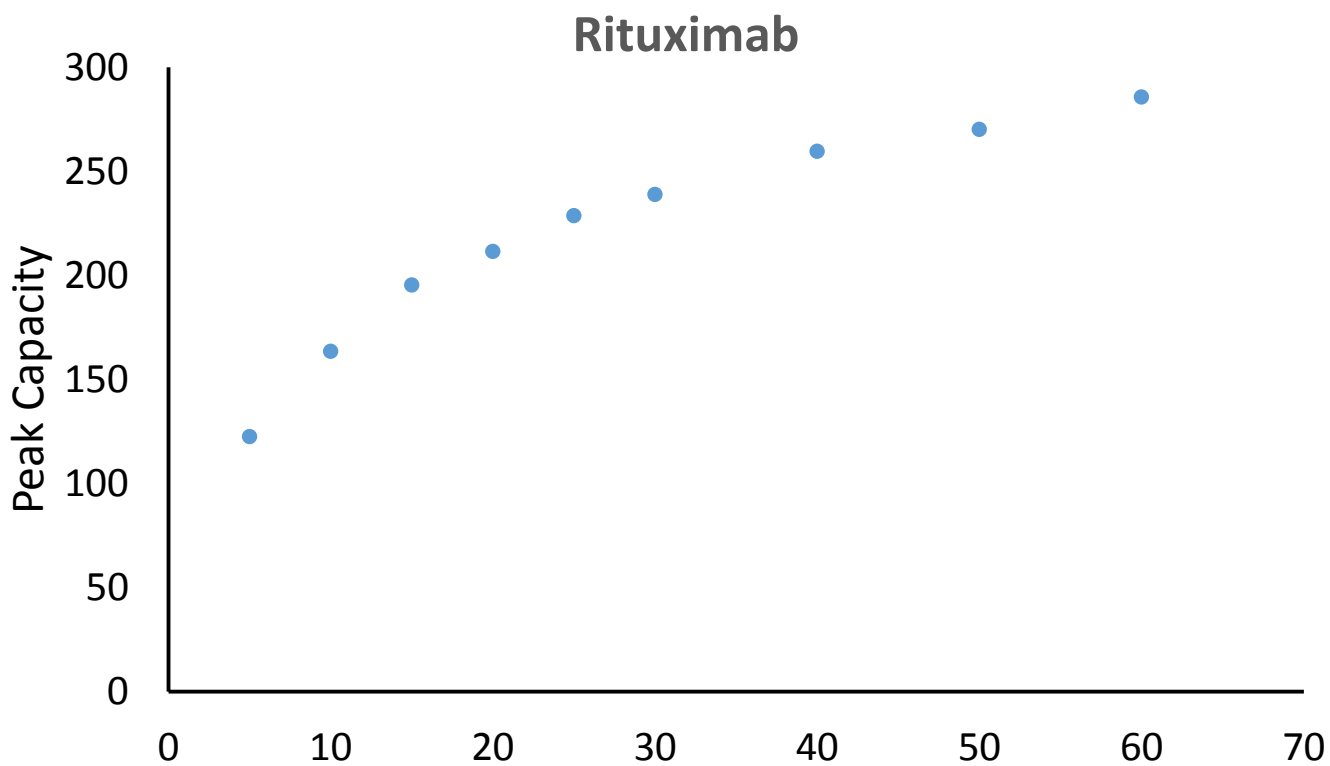


Figure 5

Fig. 5



*Conflict of Interest

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