

# **Rel. Impurity of trifloxystrobin**

## **CIPAC Peer Validation**

CIPAC peer validation of an analytical method  
for the determination of CGA 344605  
in trifloxystrobin technical, SC, EC and WG formulations  
by High Performance Liquid Chromatography

Report to CIPAC  
by  
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## 1 Introduction

### 1.1 Scope

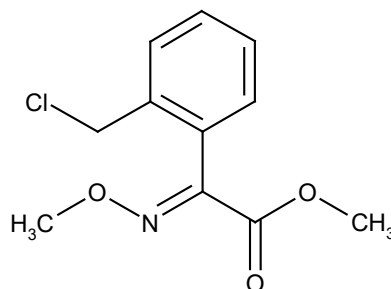
The results of the peer validation of an analytical method for the determination of CGA 344605 content in trifloxystrobin in technical grade active ingredient (TGAI), trifloxystrobin suspension concentrate (SC), trifloxystrobin emulsifiable concentrate (EC) and trifloxystrobin water dispersible granule (WG) formulations are reported.

The peer validation was performed according to CIPAC guideline for analytical methods for the determination of relevant impurities referred to in FAO and/or WHO specification for pesticide technical grade active ingredients and formulations.

### 1.2 Analyte to be determined

Analyte Name	CGA 344605
Synonyms	AE 1344136
Chemical name (IUPAC)	methyl (E)-(2-chloromethylphenyl)-methoxy-iminoacetate

Structural Formula



Empirical Formula	C <sub>11</sub> H <sub>12</sub> Cl <sub>1</sub> N <sub>1</sub> O <sub>3</sub>
Molecular Weight	241.7 g/mol
CAS No	189813-45-4

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### 1.3 Samples

Four test samples, three blank formulations and one reference standard were sent to the participants:

1. Prothioconazole technical (TC)  
1.1 trifloxystrobin TC  
Batch ID: EUTE002446  
Content: 96.0 % (w/w)
  
2. trifloxystrobin suspension concentrate (SC)  
2.1 trifloxystrobin SC 500 (500 g/L)  
Batch ID: EM4L026661  
Declared content: 500 g/L, 43.9 % (w/w)  
2.2 trifloxystrobin SC 0 (0 g/L) (blank formulation)  
Batch ID: 2020-008522  
Declared content: 0 g/L, 0 % (w/w)
  
3. trifloxystrobin emulsifiable concentrate (EC)  
3.1 fluoxastrobin + prothioconazole + trifloxystrobin EC 300 (75+150+75 g/L)  
Batch ID: EM4L028674  
Declared content: 75 g/L, 6.47 % (w/w)  
3.2 fluoxastrobin+prothioconazole+trifloxystrobin EC 0 (0+0+0 g/L) (blank formulation)  
Batch ID: 2020-009165  
Declared content: 0 g/L, 0 % (w/w)
  
4. trifloxystrobin water dispersible granule (WG)  
4.1 trifloxystrobin WG 50 (50 g/kg)  
Batch ID: EM20001194  
Declared content: 500 g/kg, 50 % (w/w)  
4.2 trifloxystrobin WG 0 (0 g/kg) (blank formulation)  
Batch ID: 2020-008527  
Declared content: 0 g/kg, 0 % (w/w)

CGA 344605 reference standard (purity 98.4% w/w)

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## 1.4 Participants

Dr. Trevor Bowen	Bayer AG, Crop Science Division Product Chemistry Analytics 5 65926 Frankfurt am Main Germany
Dr. Rolf Foerster	BASF SE Carl-Bosch-Strasse 38 67056 Ludwigshafen Germany
Dr. Michael Haustein	Currenta GmbH & Co. OHG CUR-ANT-PDA 41538 Dormagen Germany
Dr. Christian Mink	Syngenta Crop Protection AG Breitenloh 5 CH-4333 Münchwilen Switzerland
Developing Lab Schulz, Friedhelm	Bayer AG, Crop Science Division Formulation Analytics 1 Alfred-Nobel-Strasse 50 40789 Monheim am Rhein Germany

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## 2 Analytical Method

### 2.1 Outline of the Method

Reversed Phase High Performance Liquid Chromatography (RP-HPLC) hyphenated to an UV detector was used for quantitation of CGA 344605.

The CGA 344605 is separated from formulation components and active substances using a reversed phase column and gradient elution. The quantitative evaluation is carried out by comparing the peak areas with those of reference standards, using an external standard method.

### 2.2 Method Development Prior to Peer Validation

The analytical method was developed by Bayer AG, Crop Science Division and accordingly the following procedure

1) Confirmation of Analyte identification

The UV spectra of standard and sample solutions were measured under the operating conditions of the method.

2) Specificity

Interferences at the retention time of the analyte in the technical material and in the formulations were checked by comparing the chromatograms of the reference standard CGA 344605 and the test solutions of technical material, formulations and blank formulations.

3) Calibration

Calibration solutions were prepared using CGA 344605 standard. The solutions were analyzed and the peak areas of CGA 344605 were plotted against the concentration of the calibration solutions to establish a calibration line.

4) Repeatability

Six independently prepared sample solutions for each technical material and formulation were analyzed according to the method.

The content of CGA 344605 found in the technical material and in the formulations SC and EC and WG was not enough to evaluate the repeatability. Therefore, the test solutions were spiked with the analyte CGA 344605 at a concentration of 0.1 mg/50 mL to determine the repeatability. Mean and relative standard deviations (RSD) were calculated.

5) Recovery

The technical material or the blank formulations were fortified with the analyte CGA 344605 at two levels, fortification level I 0.02 mg/50 mL and fortification level II 0.22 mg/50 mL. These solutions were analyzed, and the recoveries were calculated.

6) Limit of Quantification (LOQ)

The concentration of the lowest tested fortification level was defined as LOQ, as acceptable recovery and precision under repeatability conditions were successfully demonstrated.

### **2.3 Peer Validation**

The peer validation was conducted in four laboratories. The participants are shown in 1.4. We requested the collaborators to conduct the peer validation according to the prescribed protocol, describe the operating conditions in detail and attach the calibration curve and the chromatograms for each sample.

The investigations included specificity, calibration, repeatability, recovery and LOQ. The details of each procedure are the same as those described in 2.2.

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### 3 Remarks of the participants

Laboratory	Remarks
Developing Lab	n.a.
Lab 1	n.a.
Lab 2	n.a.
Lab 3	n.a.
Lab 4	n.a.



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## 4 Results and Discussion

### 4.1 Peer Validation

#### 1) Confirmation of Analyte identification

The DAD spectra obtained by the developing laboratory during the method development prior to peer validation are shown in Figure 1 to Figure 5.

#### 2) Specificity

The specificity of the analytical method for CGA 344605 is assessed sufficient as no significant interfering compounds were detected in the chromatograms at the retention time of the analyte.

#### 3) Calibration

The calibration lines are shown in Figure 48 to Figure 57. The calibration equation and the correlation factor are reported in addition. The calibration function of the analytical method is assessed as linear over a concentration range of 8% of the max. accepted level (MAL) in the formulation (0.016 mg/50 mL) to 120% MAL (0.24 mg/50 mL), as the correlation factor was satisfactory  $> 0.99$  (in accordance with SANCO/3030/99).

#### 4) Repeatability

The precision of the method was satisfactory with RSD values of 0.1 – 2. % as shown in Table 1 to Table 4.

In all cases, the Horrat value (Horwitz ratio, Hr) was  $\leq 1$ .

#### 5) Recovery

The recoveries were satisfactory as shown in Table 5 to Table 12

The technical material or the blank formulations were fortified with the analyte CGA 344605 at two levels, fortification level I (LOQ) 0.02 mg/50 mL and fortification level II (close to the max. accepted level (MAL) in the formulation, see Appendix 1) 0.22 mg/50 mL. These concentrations are equal to:

	LOQ Level I	Level II
trifloxystrobin TC	0.040 % (w/w) <sup>1</sup>	0.44 % (w/w) <sup>1</sup>
trifloxystrobin SC 500 (500 g/L)	0.017 % (w/w) <sup>2</sup>	0.19 % (w/w) <sup>2</sup>
Fluoxastrobin+Prothioconazole+trifloxystrobin EC 300 (75+150+75 g/L)	0.0026 % (w/w) <sup>3</sup>	0.028 % (w/w) <sup>3</sup>
trifloxystrobin WG 50 (500 g/kg)	0.020% (w/w) <sup>4</sup>	0.22% (w/w) <sup>4</sup>

#### 6) Limit of Quantification (LOQ)

A concentration of 0.02 mg/50 mL CGA 344605 was the lowest tested fortification level for which acceptable recovery and precision under repeatability conditions were successfully demonstrated. This concentration is equal to:

trifloxystrobin TC	0.040 % (w/w)
trifloxystrobin SC 500 (500 g/L)	0.017 % (w/w)
fluoxastrobin+prothioconazole+trifloxystrobin EC 300 (75+150+75 g/L)	0.0026 % (w/w)
trifloxystrobin WG 50 (500 g/kg)	0.020% (w/w)

## 5 Conclusion

For all samples, the analytical method was peer-validated in terms of specificity, calibration, repeatability, recovery and limit of quantification. The RSDs of repeatability for technical material and formulations were found acceptable according to the Horrat criteria  $\leq 1$ .

In conclusion, the proposed method was successfully peer-validated and was considered appropriate for the determination of CGA 344605 in technical material trifloxystrobin and SC, EC and WG formulations.

<sup>1</sup> Referred to a nominal sample concentration of 50 mg/50 mL

<sup>2</sup> Referred to a nominal sample concentration of 115 mg/50 mL

<sup>3</sup> Referred to a nominal sample concentration of 770 mg/50 mL

<sup>4</sup> Referred to a nominal sample concentration of 100 mg/50 mL

## 6 Determination of relevant impurity CGA 344605

### 6.1 Tables of results

Table 1: Results trifloxystrobin TC (Repeatability)

	CGA 344605 [mg/50 mL] <sup>1</sup>				
	Developing Lab	Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	0.1107	0.1093	0.1026	0.1020	0.1102
Weighing no. 2	0.1085	0.1087	0.1031	0.1019	0.1098
Weighing no. 3	0.1010	0.1097	0.1026	0.1021	0.1101
Weighing no. 4	0.1104	0.1081	0.1026	0.1022	0.1101
Weighing no. 5	0.1078	0.1090	0.1040 <sup>3</sup>	0.1027	0.1104
Weighing no. 6	0.1131	0.1089	0.1031	0.1021	0.1104
<b>Mean value [mg/50 mL]</b>	<b>0.110</b>	<b>0.109</b>	<b>0.103</b>	<b>0.102</b>	<b>0.110</b>
<b>Mean value [% w/w]<sup>2</sup></b>	<b>0.220</b>	<b>0.218</b>	<b>0.206</b>	<b>0.204</b>	<b>0.220</b>
SD	0.00186	0.000543	0.000548	0.00028	0.000225
RSD [%]	1.69	0.50	0.54	0.27	0.20
Horwitz-Value RSD (r) <sub>max</sub>	3.37	3.37	3.40	3.40	3.37
<b>Horrat value H<sub>r</sub></b>	<b>0.50</b>	<b>0.15</b>	<b>0.16</b>	<b>0.08</b>	<b>0.06</b>
Outliers	no	no	yes <sup>3</sup>	no	no

<sup>1</sup> The amount of CGA 344605 determined in the test item was < LOQ. Therefore, the precision was determined using test item solutions spiked with reference item CGA 344605. For the calculation of the RSD, the results in mg/50 mL were used to eliminate the error of the variability of the test item weight in the test solutions (error propagation)

<sup>2</sup> Calculated from mean value in [mg/50 mL] for a nominal sample concentration of 50 mg/50 mL

<sup>3</sup> The results of laboratory 2 show an upper outlier at 95% confidence according to Dixon Test. However, this value was considered valid and therefore it was not disregarded to calculate the mean value and the RSD.

**Table 2: Results trifloxystrobin SC 500 (500 g/L) (Repeatability)**

	CGA 344605 [mg/50 mL] <sup>1</sup>				
	Developing Lab	Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	0.1060	0.1062	0.1003	0.0995	0.1068
Weighing no. 2	0.1071	0.1066	0.0996	0.0996	0.1070
Weighing no. 3	0.1053	0.106	0.0981	0.0993	0.1076
Weighing no. 4	0.1044	0.1065	0.1029	0.0994	0.1069
Weighing no. 5	0.1055	0.1059	0.0995	0.0995	0.1073
Weighing no. 6	0.1054	0.1059	0.0989	0.0994	0.1055
<b>Mean value [mg/50 mL]</b>	<b>0.106</b>	<b>0.106</b>	<b>0.0999</b>	<b>0.0995</b>	<b>0.107</b>
<b>Mean value [% w/w]<sup>2</sup></b>	<b>0.0918</b>	<b>0.0918</b>	<b>0.0869</b>	<b>0.0865</b>	<b>0.0929</b>
SD	0.000906	0.000306	0.00165	0.000105	0.000723
RSD [%]	0.86	0.29	1.65	0.11	0.68
Horwitz-Value RSD (r) <sub>max</sub>	3.84	3.84	3.87	3.87	3.83
<b>Horrat value H<sub>r</sub></b>	<b>0.22</b>	<b>0.08</b>	<b>0.43</b>	<b>0.03</b>	<b>0.18</b>
Outliers	no	no	no	no	no

<sup>1</sup> The amount of CGA 344605 determined in the test item was < LOQ. Therefore, the precision was determined using test item solutions spiked with reference item CGA 344605. For the calculation of the RSD, the results in mg/50 mL were used to eliminate the error of the variability of the test item weight in the test solutions (error propagation)

<sup>2</sup> Calculated from mean value in [mg/50 mL] for a nominal sample concentration of 115 mg/50 mL

**Table 3: Results fluoxastrobin+prothioconazole+trifloxystrobin EC 300 (75+150+75 g/L) (Repeatability)**

	CGA 344605 [mg/50 mL] <sup>1</sup>				
	Developing Lab	Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	0.1124 <sup>3</sup>	0.1004	0.0934	0.0932	0.1098 <sup>3</sup>
Weighing no. 2	0.1001	0.1018	0.0908	0.0934	0.1027
Weighing no. 3	0.1036	0.0991	0.0903	0.0941	0.1034
Weighing no. 4	0.1025	0.0999	0.0908	0.0936	0.1026
Weighing no. 5	0.1026	0.1043	0.0932	0.0936	0.1029
Weighing no. 6	0.1034	0.0984	0.0907	0.0935	0.1026
<b>Mean value [mg/50 mL]</b>	<b>0.102</b>	<b>0.101</b>	<b>0.0915</b>	<b>0.0936</b>	<b>0.103</b>
<b>Mean value [% w/w]<sup>2</sup></b>	<b>0.0133</b>	<b>0.0131</b>	<b>0.0119</b>	<b>0.0122</b>	<b>0.0134</b>
SD	0.00137	0.00213	0.00138	0.000301	0.000336
RSD [%]	1.34	2.12	1.51	0.32	0.33
Horwitz-Value RSD (r) <sub>max</sub>	5.13	5.15	5.22	5.21	5.13
<b>Horrat value H<sub>r</sub></b>	<b>0.26</b>	<b>0.41</b>	<b>0.29</b>	<b>0.06</b>	<b>0.06</b>
Outliers	yes <sup>3</sup>	no	no	no	yes <sup>3</sup>

<sup>1</sup> The amount of CGA 344605 determined in the test item was < LOQ. Therefore, the precision was determined using test item solutions spiked with reference item CGA 344605. For the calculation of the RSD, the results in mg/50 mL were used to eliminate the error of the variability of the test item weight in the test solutions (error propagation)

<sup>2</sup> Calculated from mean value in [mg/50 mL] for a nominal sample concentration of 770 mg/50 mL

<sup>3</sup> The results of the developing laboratory and Lab 4 show an upper outlier at 95% confidence according to Dixon Test. This value was disregarded for the calculation of the mean value and the standard deviation

**Table 4: Results trifloxystrobin WG 50 (500 g/kg) (Repeatability)**

	CGA 344605 [mg/50 mL] <sup>1</sup>				
	Developing Lab	Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	0.1115	0.1077	0.1040	0.1031	0.1105
Weighing no. 2	0.1095	0.1059	0.1039	0.1033	0.1111
Weighing no. 3	0.1099	0.1057	0.1036	0.1034	0.1102
Weighing no. 4	0.1096	0.1049	0.1022	0.1037	0.1112
Weighing no. 5	0.1095	0.1051	0.1022	0.1053	0.1090
Weighing no. 6	0.1088	0.1041	0.1020	0.1040	0.1107
<b>Mean value [mg/50 mL]</b>	<b>0.110</b>	<b>0.106</b>	<b>0.103</b>	<b>0.104</b>	<b>0.110</b>
<b>Mean value [% w/w]<sup>2</sup></b>	<b>0.110</b>	<b>0.106</b>	<b>0.103</b>	<b>0.104</b>	<b>0.110</b>
SD	0.000919	0.00122	0.000943	0.0008	0.000802
RSD [%]	0.84	1.16	0.92	0.77	0.73
Horwitz-Value RSD (r) <sub>max</sub>	3.74	3.76	3.77	3.77	3.73
<b>Horrat value H<sub>r</sub></b>	<b>0.22</b>	<b>0.31</b>	<b>0.24</b>	<b>0.20</b>	<b>0.20</b>
Outliers	no	no	no	no	no

<sup>1</sup> The amount of CGA 344605 determined in the test item was < LOQ. Therefore, the precision was determined using test item solutions spiked with reference item CGA 344605. For the calculation of the RSD, the results in mg/50 mL were used to eliminate the error of the variability of the test item weight in the test solutions (error propagation)

<sup>2</sup> Calculated from mean value in [mg/50 mL] for a nominal sample concentration of 100 mg/50 mL

**Table 5: Results trifloxystrobin TC (Recovery Level I - LOQ)**

	CGA 344605				
	Developing Lab	Recovery [%]			
		Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	101.8	103.3	95.64	98.70	99.95
Weighing no. 2	99.17	105.5	96.13	98.70	98.08 <sup>1</sup>
Weighing no. 3	98.86	106.1	96.13	98.27	99.48
Weighing no. 4	98.95	105.6	96.13	100.0	99.48
Weighing no. 5	102.2	103.0	95.64	99.57	99.95
Weighing no. 6	101.3	105.3	99.60 <sup>1</sup>	98.27	99.95
<b>Mean value</b>	<b>100.4</b>	<b>104.8</b>	<b>96.6</b>	<b>98.9</b>	<b>99.5</b>
SD	1.55	1.31	1.52	0.71	0.72
RSD [%]	1.54	0.96	1.57	0.72	0.73
Outliers	no	no	yes <sup>1</sup>	no	yes <sup>1</sup>
<sup>1</sup> The results of laboratory 2 show an upper outlier and the results of laboratory 4 show a lower outlier at 95% confidence according to Dixon Test. However, these values were considered valid and therefore they were not disregarded to calculate the mean value and the RSD.					

Mean recovery rates are within the requested range of 75-125 % according to SANCO/3030/99 for a nominal analyte content  $\geq 0.01$  -  $< 0.1\%$  (w/w)

Remark: The initial concentration of the analyte found in the unfortified sample, which was significant at the concentration of recovery level I, was subtracted from the measured concentration of the fortified sample to calculate the recovery [%].

**Table 6: Results trifloxystrobin TC (Recovery Level II - MAL)**

	CGA 344605				
	Developing Lab	Recovery [%]			
		Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	101.9	103.3	100.5	100.8	100.5
Weighing no. 2	101.4	103.0	101.2	100.8	100.5
Weighing no. 3	100.8	103.2	100.5	101	100.3
Weighing no. 4	101.0	103.2	100.1	100.7	100.3
Weighing no. 5	102.2	103.7	100.8	101.1	100.4
Weighing no. 6	100.9	103.2	100.6	100.9	100.4
<b>Mean value</b>	<b>101.4</b>	<b>103.3</b>	<b>100.6</b>	<b>100.9</b>	<b>100.4</b>
SD	0.58	0.23	0.37	0.15	0.09
RSD [%]	0.57	0.23	0.36	0.15	0.09
Outliers	no	no	no	no	no

Mean recovery rates are within the requested range of 80-120 % according to SANCO/3030/99 for a nominal analyte content  $\geq 0.1$  to  $< 1\%$  (w/w)



**Table 7: Results trifloxystrobin SC 500 (500 g/L) (Recovery Level I - LOQ)**

	CGA 344605				
	Developing Lab	Recovery [%]			
		Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	98.34	96.73	94.15	99.13	97.14
Weighing no. 2	104.2	99.21	94.15	100.0	96.20
Weighing no. 3	98.56	98.42	94.65	100.0	97.61
Weighing no. 4	102.1	98.66	93.16	101.3	97.14
Weighing no. 5	99.34	97.48	95.14	100.9	103.2 <sup>1</sup>
Weighing no. 6	101.7	98.42	94.15	100.4	95.26
<b>Mean value</b>	<b>100.7</b>	<b>98.2</b>	<b>94.2</b>	<b>100.3</b>	<b>97.8</b>
SD	2.33	0.894	0.659	0.764	2.80
RSD [%]	2.31	0.91	0.70	0.76	2.86
Outliers	no	no	no	no	yes <sup>1</sup>

<sup>1</sup> The results of laboratory 4 show an upper outlier at 95% confidence according to Dixon Test. However, this value was considered valid and therefore it was not disregarded to calculate the mean value and the RSD.

Mean recovery rates are within the requested range of 75-125 % according to SANCO/3030/99 for a nominal analyte content  $\geq 0.01$  -  $< 0.1\%$  (w/w)

**Table 8: Results trifloxystrobin SC 500 (500 g/L) (Recovery Level II - MAL)**

	CGA 344605				
	Developing Lab	Recovery [%]			
		Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	100.7	99.24	100.3	98.79	96.31 <sup>1</sup>
Weighing no. 2	99.50	99.34	98.97	98.53	97.72
Weighing no. 3	99.82	99.33	99.78	98.57	97.63
Weighing no. 4	100.7	99.11	99.33	98.62	97.72
Weighing no. 5	101.1	98.97	99.92	98.57	97.29
Weighing no. 6	100.4	99.25	99.42	98.66	97.50
<b>Mean value</b>	<b>100.4</b>	<b>99.2</b>	<b>99.6</b>	<b>98.6</b>	<b>97.4</b>
SD	0.60	0.14	0.47	0.09	0.54
RSD [%]	0.60	0.14	0.48	0.09	0.55
Outliers	no	no	no	no	yes <sup>1</sup>

<sup>1</sup> The results of laboratory 4 show a lower outlier at 95% confidence according to Dixon Test. However, this value was considered valid and therefore it was not disregarded to calculate the mean value and the RSD.

Mean recovery rates are within the requested range of 80-120 % according to SANCO/3030/99 for a nominal analyte content  $\geq 0.1$  to  $< 1\%$  (w/w)

**Table 9: Results fluoxastrobin+prothioconazole+trifloxystrobin EC 300 (75+150+75 g/L) (Recovery Level I - LOQ)**

	CGA 344605				
	Developing Lab	Recovery [%]			
		Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	178.3 <sup>1</sup>	108.5	87.22	93.07	100.4
Weighing no. 2	94.41	105.3	85.73	93.51	93.85
Weighing no. 3	99.65	115.3	87.22	93.94	92.91
Weighing no. 4	99.00	106.4	83.75	92.64	95.73
Weighing no. 5	98.56	107.0	85.73	93.51	94.32
Weighing no. 6	95.68	110.6	84.24	93.51	95.26
<b>Mean value</b>	<b>97.5</b>	<b>108.9</b>	<b>85.7</b>	<b>93.4</b>	<b>95.4</b>
SD	2.28	3.66	1.45	0.45	2.64
RSD [%]	2.34	3.58	1.70	0.48	2.77
Outliers	yes <sup>1</sup>	no	no	no	no

<sup>1</sup> The results of the developing laboratory show an upper outlier at 95% confidence according to Dixon Test. This value was disregarded for the calculation of the mean value and the standard deviation

Mean recovery rates are within the requested range of 70-130 % according to SANCO/3030/99 for a nominal analyte content < 0.01% (w/w)

**Table 10: Results fluoxastrobin+prothioconazole+trifloxystrobin EC 300 (75+150+75 g/L) (Recovery Level II)**

	CGA 344605				
	Developing Lab	Recovery [%]			
		Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	98.91	101.5	98.16	97.32	95.59 <sup>1</sup>
Weighing no. 2	99.03	101.2	98.12	97.49	96.95
Weighing no. 3	100.4	101.3	97.98	97.40	96.69
Weighing no. 4	98.82	100.8	98.34	97.23	96.91
Weighing no. 5	99.03	100.9	99.51 <sup>1</sup>	97.10	96.82
Weighing no. 6	100.0	100.7	97.76	97.32	96.91
<b>Mean value</b>	<b>99.4</b>	<b>101.1</b>	<b>98.3</b>	<b>97.3</b>	<b>96.7</b>
SD	0.66	0.31	0.62	0.14	0.53
RSD [%]	0.67	0.31	0.63	0.14	0.54
Outliers	no	no	yes <sup>1</sup>	no	yes <sup>1</sup>
<sup>1</sup> The results of laboratory 2 show an upper outlier and the results of laboratory 4 show an lower outlier at 95% confidence according to Dixon Test. However, these values were considered valid and therefore they were not disregarded to calculate the mean value and the RSD.					

Mean recovery rates are within the requested range of 75-125 % according to SANCO/3030/99 for a nominal analyte content  $\geq 0.01$  to  $< 0.1\%$  (w/w)

**Table 11: Results trifloxystrobin WG 50 (500 g/kg) (Recovery Level I - LOQ)**

	CGA 344605				
	Developing Lab	Recovery [%]			
		Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	98.43	85.54	92.67 <sup>1</sup>	94.37	106.5
Weighing no. 2	108.2	79.75	94.65	95.67	115.4
Weighing no. 3	100.3	73.27	94.65	103.5	114.0
Weighing no. 4	103.4	74.11	94.65	94.81	114.0
Weighing no. 5	106.9	78.81	95.64	102.6	109.8
Weighing no. 6	105.3	75.59	95.64	96.54	107.0
<b>Mean value</b>	<b>100.3</b>	<b>77.9</b>	<b>94.7</b>	<b>97.9</b>	<b>111.1</b>
SD	3.81	4.56	1.08	4.06	3.87
RSD [%]	3.66	5.85	1.15	4.14	3.49
Outliers	no	no	yes <sup>1</sup>	no	no

<sup>1</sup> The results of laboratory 2 show a lower outlier at 95% confidence according to Dixon Test. However, this value was considered valid and therefore it was not disregarded to calculate the mean value and the RSD.

Mean recovery rates are within the requested range of 75-125 % according to SANCO/3030/99 for a nominal analyte content  $\geq 0.01$  to  $< 0.1\%$  (w/w)

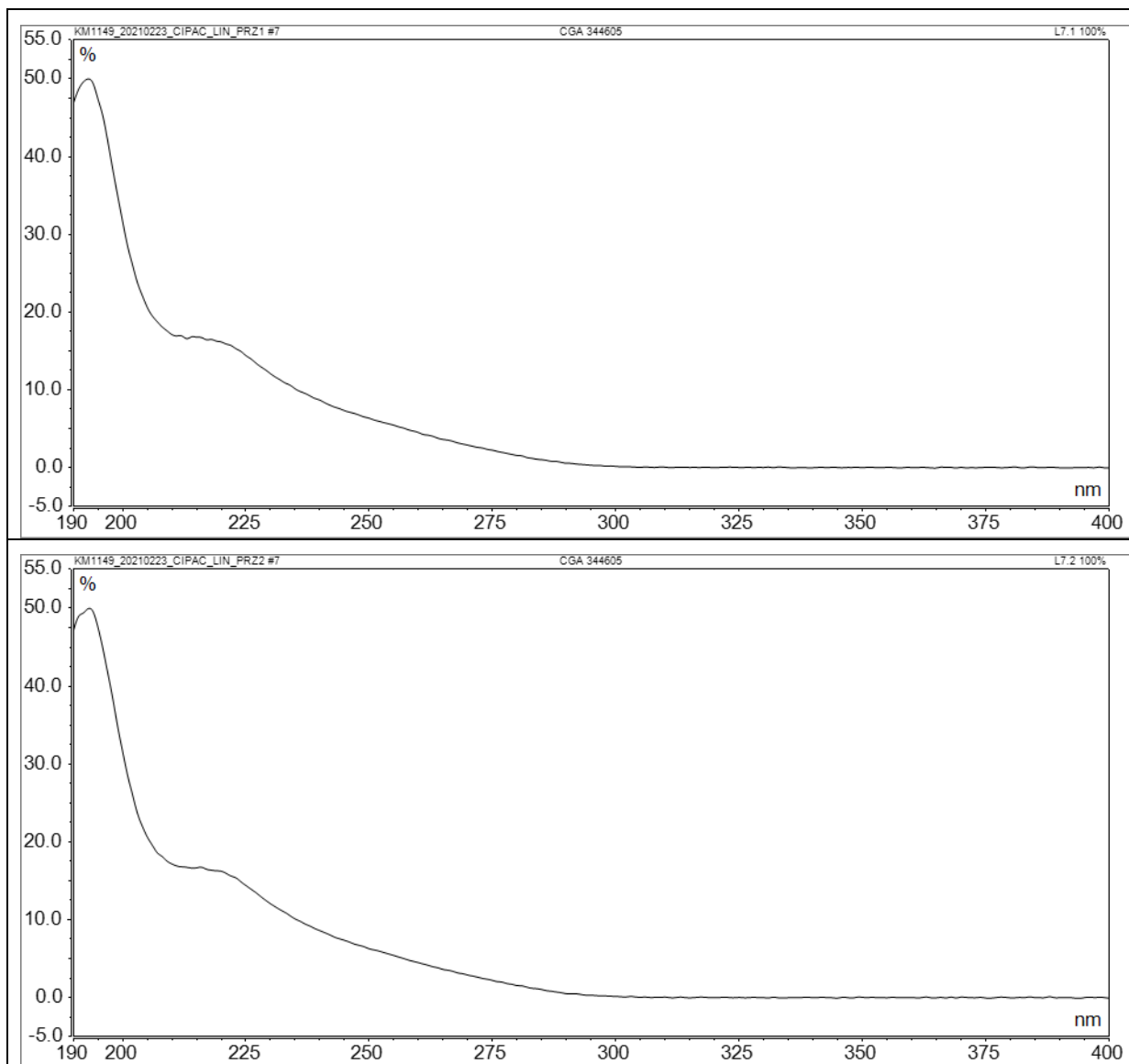
**Table 12: Results trifloxystrobin WG 50 (500 g/kg) (Recovery Level II)**

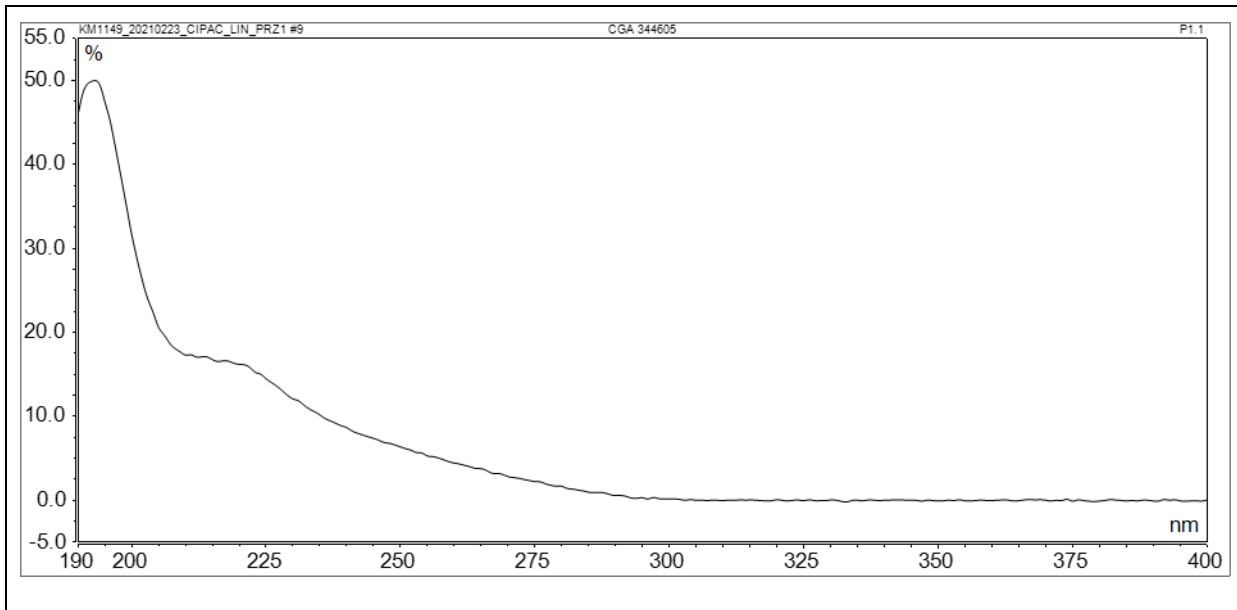
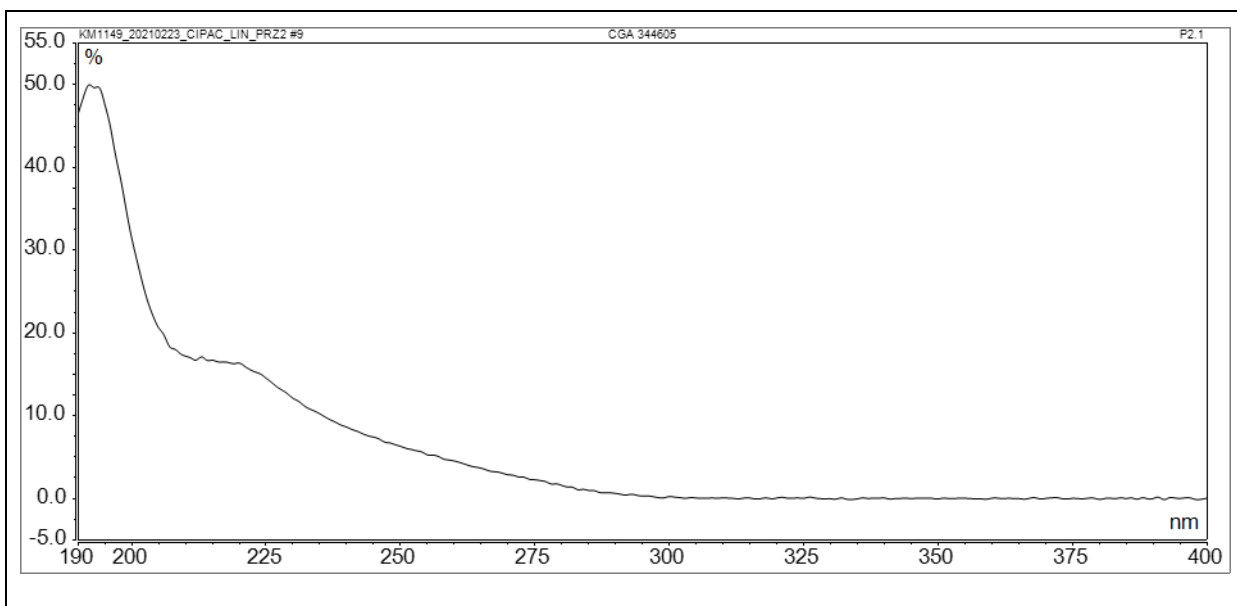
	CGA 344605				
	Developing Lab	Recovery [%]			
		Lab 1	Lab 2	Lab 3	Lab 4
Weighing no. 1	99.59	93.99	97.35	102.3 <sup>1</sup>	99.38
Weighing no. 2	100.7	95.33	98.39	98.05	99.42
Weighing no. 3	99.63	96.40	97.31	97.23	99.68
Weighing no. 4	101.0	95.66	97.4	97.84	99.42
Weighing no. 5	100.4	94.60	99.06	98.49	99.38
Weighing no. 6	99.49	95.22	101.7	98.62	99.72
<b>Mean value</b>	<b>100.1</b>	<b>95.2</b>	<b>98.5</b>	<b>98.8</b>	<b>99.5</b>
SD	0.65	0.84	1.70	1.81	0.16
RSD [%]	0.65	0.88	1.73	1.83	0.16
Outliers	no	no	no	yes <sup>1</sup>	no
<sup>1</sup> The results of laboratory 3 show an upper outlier at 95% confidence according to Dixon Test. However, this value was considered valid and therefore it was not disregarded to calculate the mean value and the RSD.					

Mean recovery rates are within the requested range of 80-120 % according to SANCO/3030/99 for a nominal analyte content  $\geq 0.1$  to  $< 1\%$  (w/w)

## 6.2 UV-Spectra

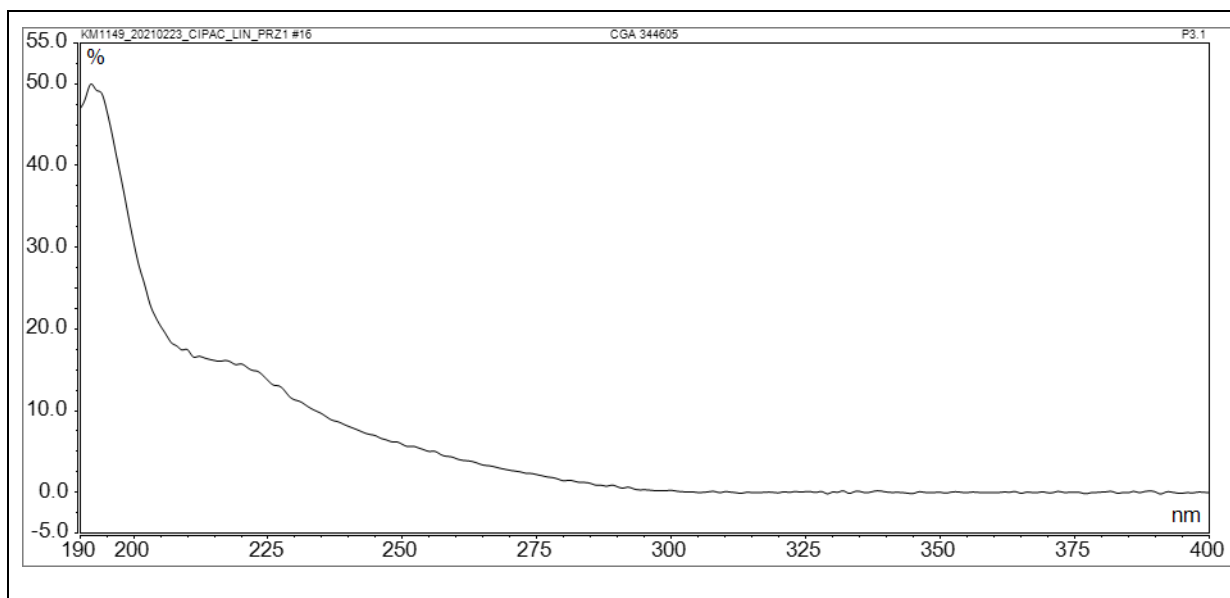
**Figure 1: UV spectra of reference standard CGA 344605**  
(Top: in acetonitrile; Bottom: in Acetonitrile/Water)



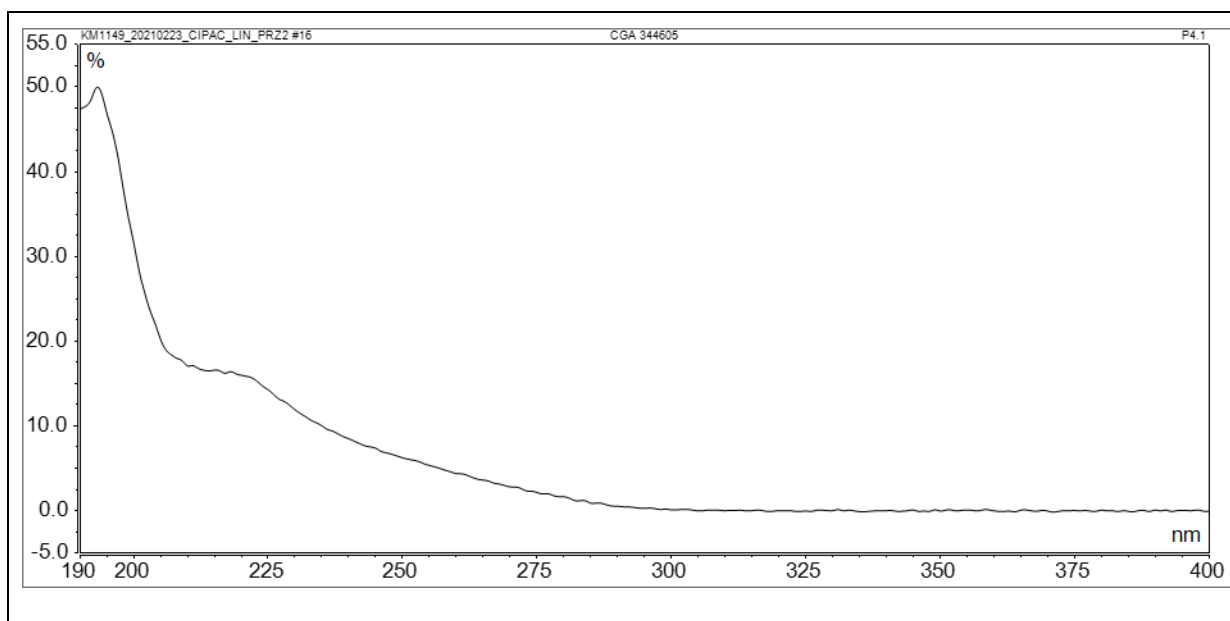
**Figure 2: UV spectra of CGA 344605 in spiked test item trifloxystrobin TC****Figure 3: UV spectra of CGA 344605 in spiked test item trifloxystrobin SC 500 (500 g/L)**



**Figure 4: UV spectra of CGA 344605 in spiked test item fluoxastrobin + prothioconazole + trifloxystrobin EC 300 (75+150+75 g/L)**

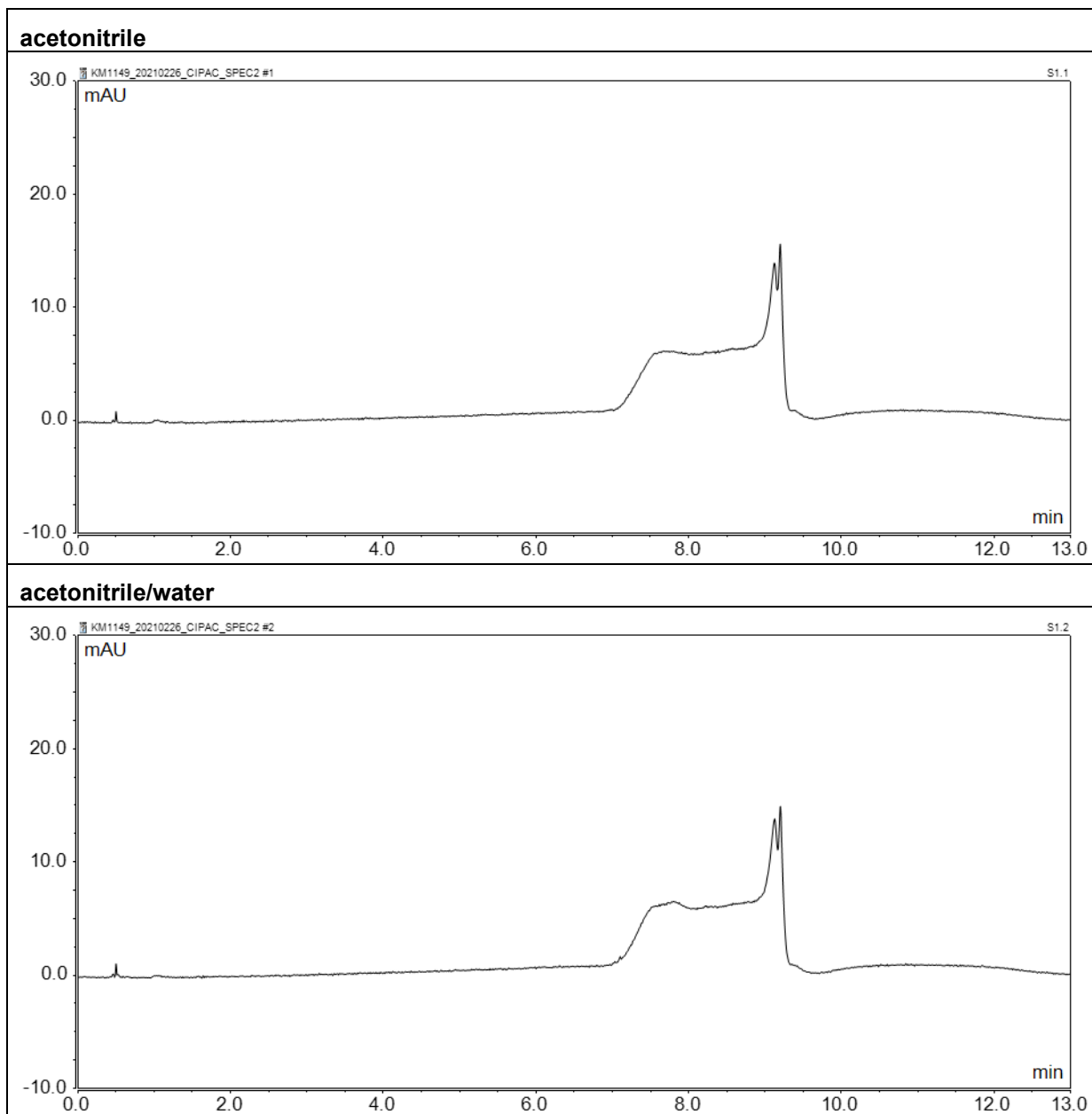


**Figure 5: UV spectra of CGA 344605 in spiked test item trifloxystrobin WG 50 (50 g/kg)**



### 6.3 Chromatograms

Figure 6: Chromatograms of sample solvent  
Developing Lab



**Figure 7: Chromatograms of reference standard CGA 344605  
Developing Lab**

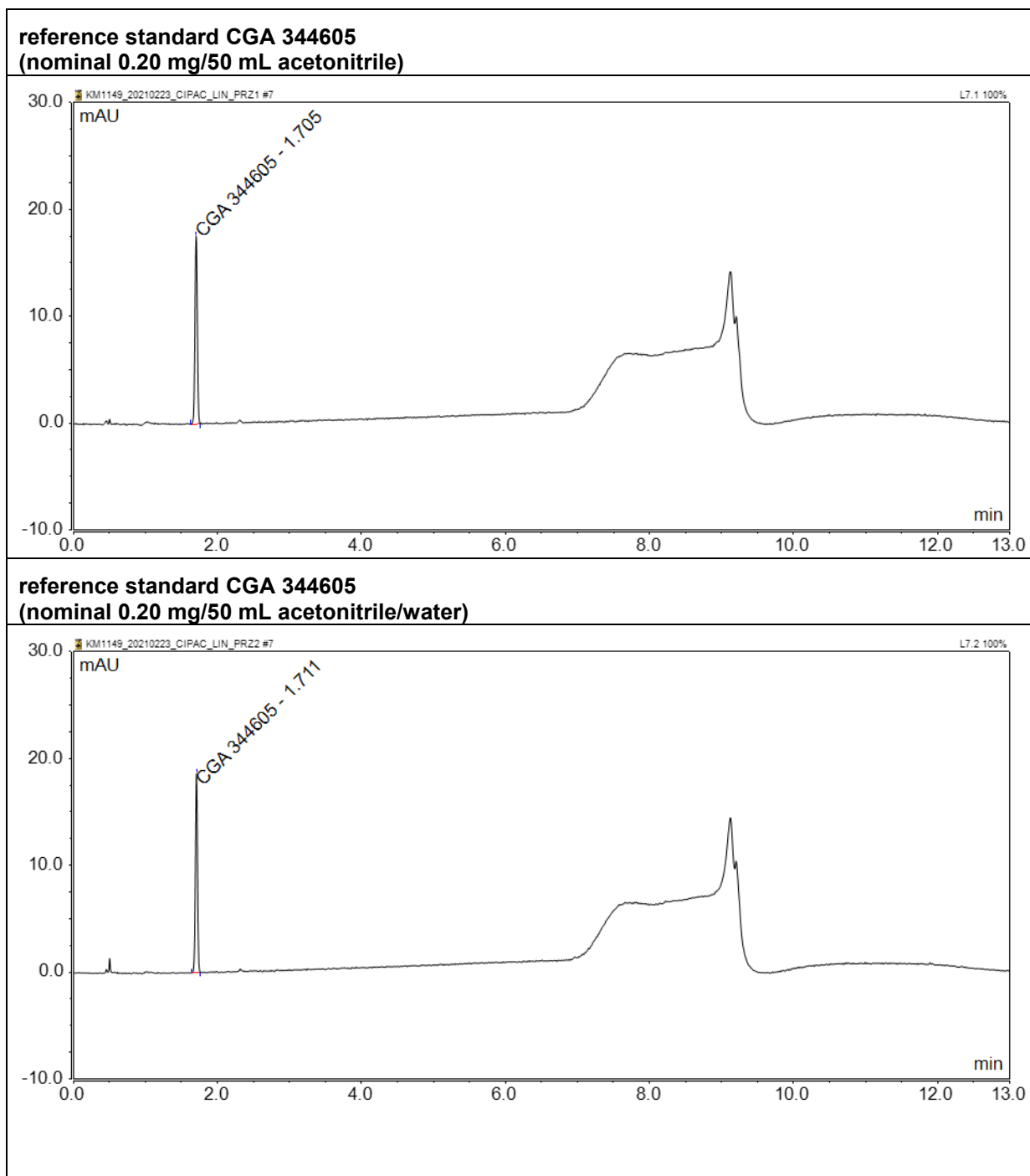
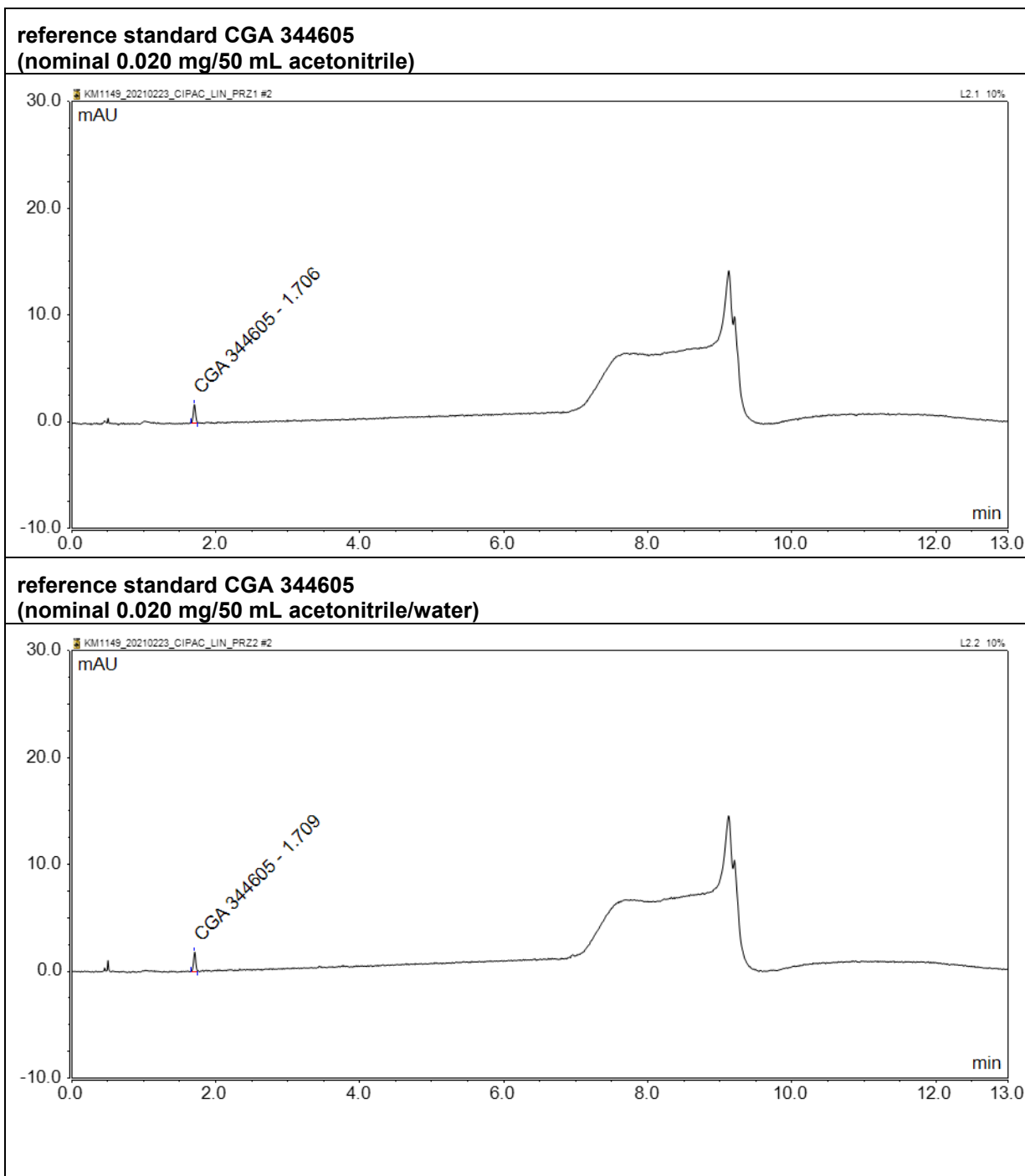
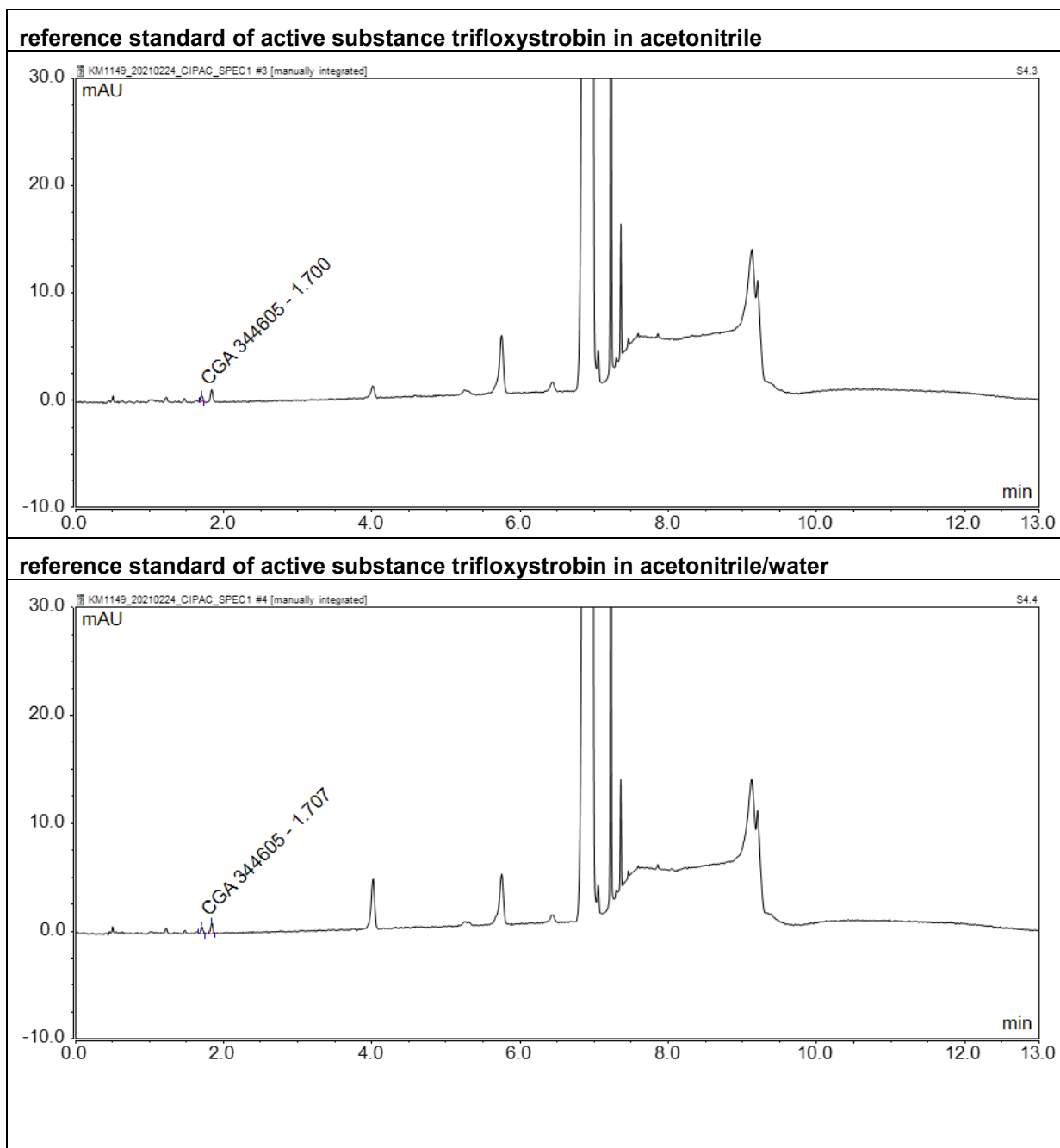


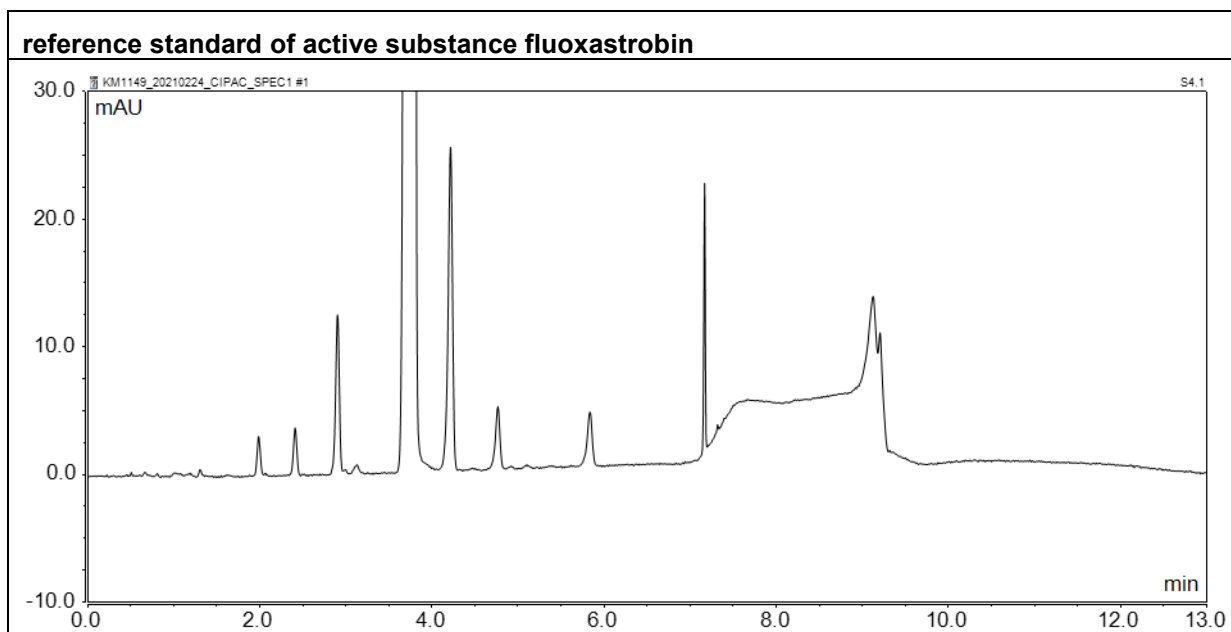
Figure 7. Continuation



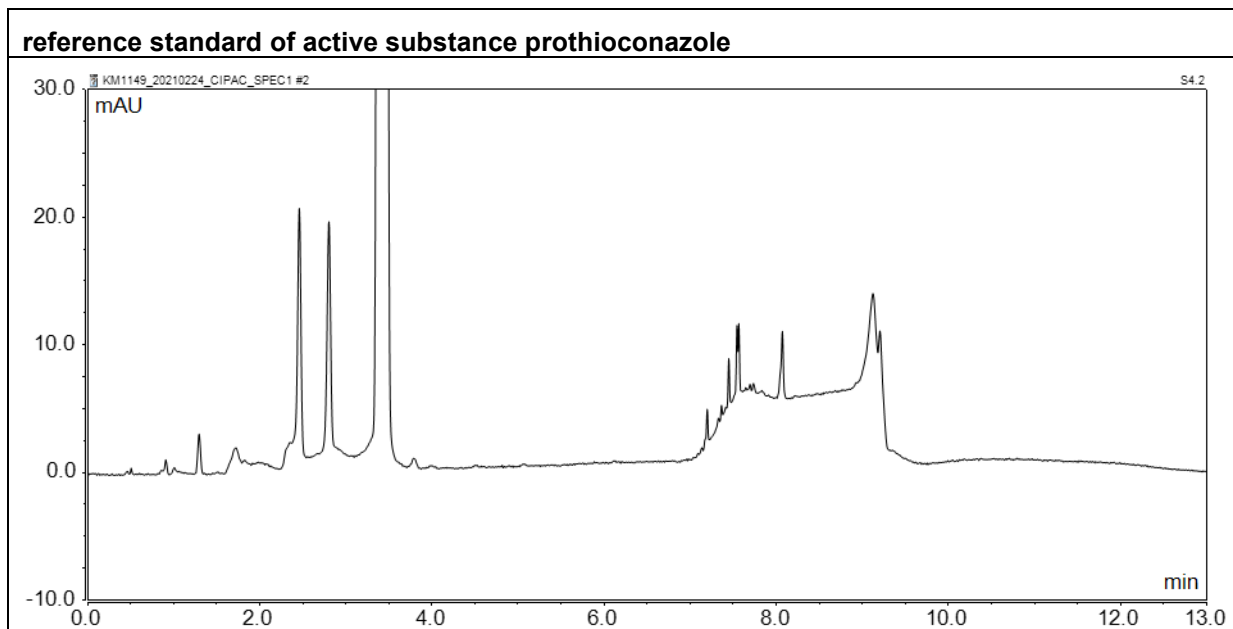
**Figure 8: Chromatograms of reference standard of active substance trifloxystrobin. Developing Lab**



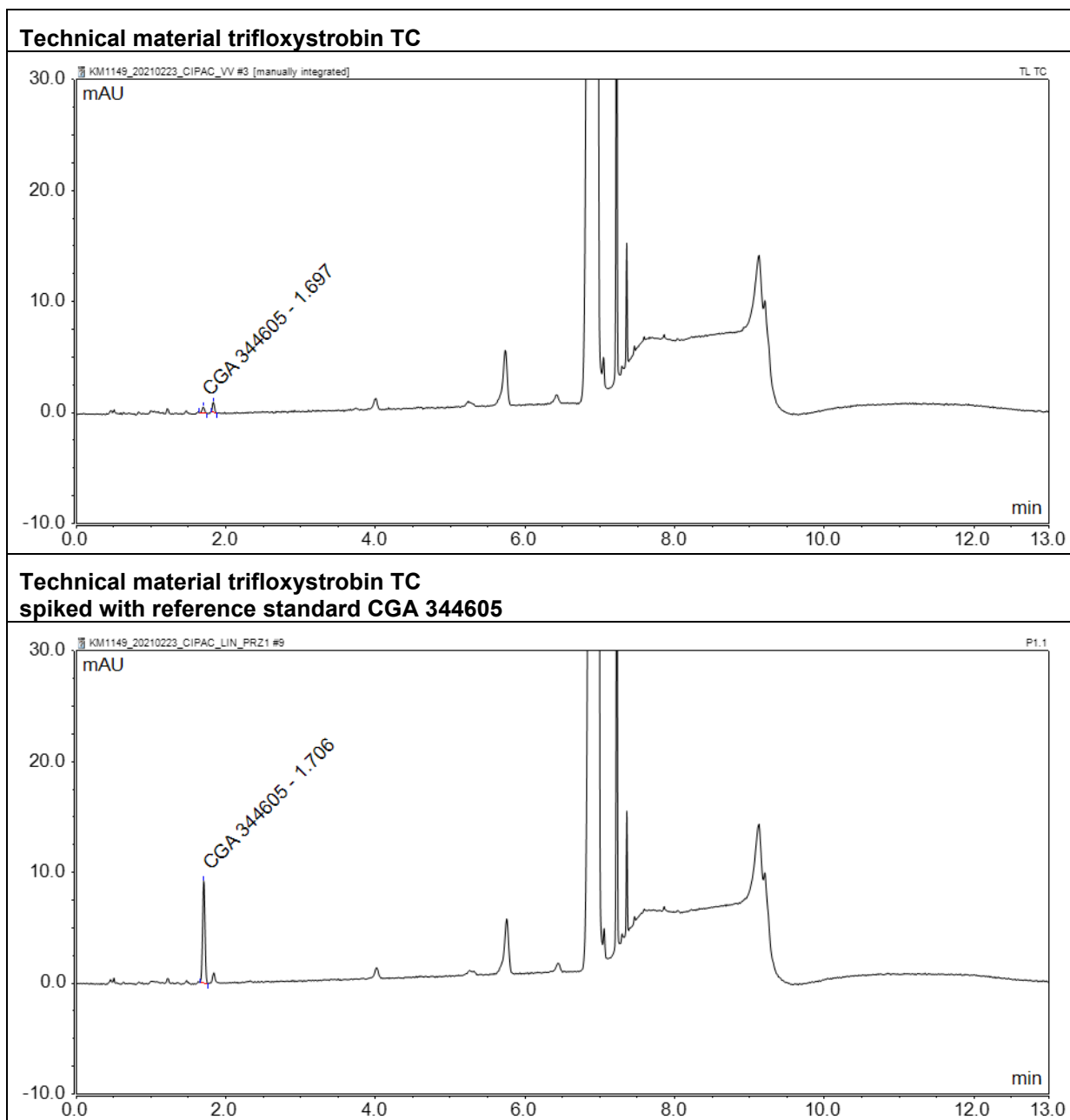
**Figure 9: Chromatograms of reference standard of active substance fluoxastrobin. Developing Lab**



**Figure 10: Chromatograms of reference standard of active substance prothioconazole. Developing Lab**



**Figure 11: Chromatograms of trifloxystrobin TC  
Developing Lab**



**Figure 12: Chromatograms of trifloxystrobin SC 500 (500 g/L)  
Developing Lab.**

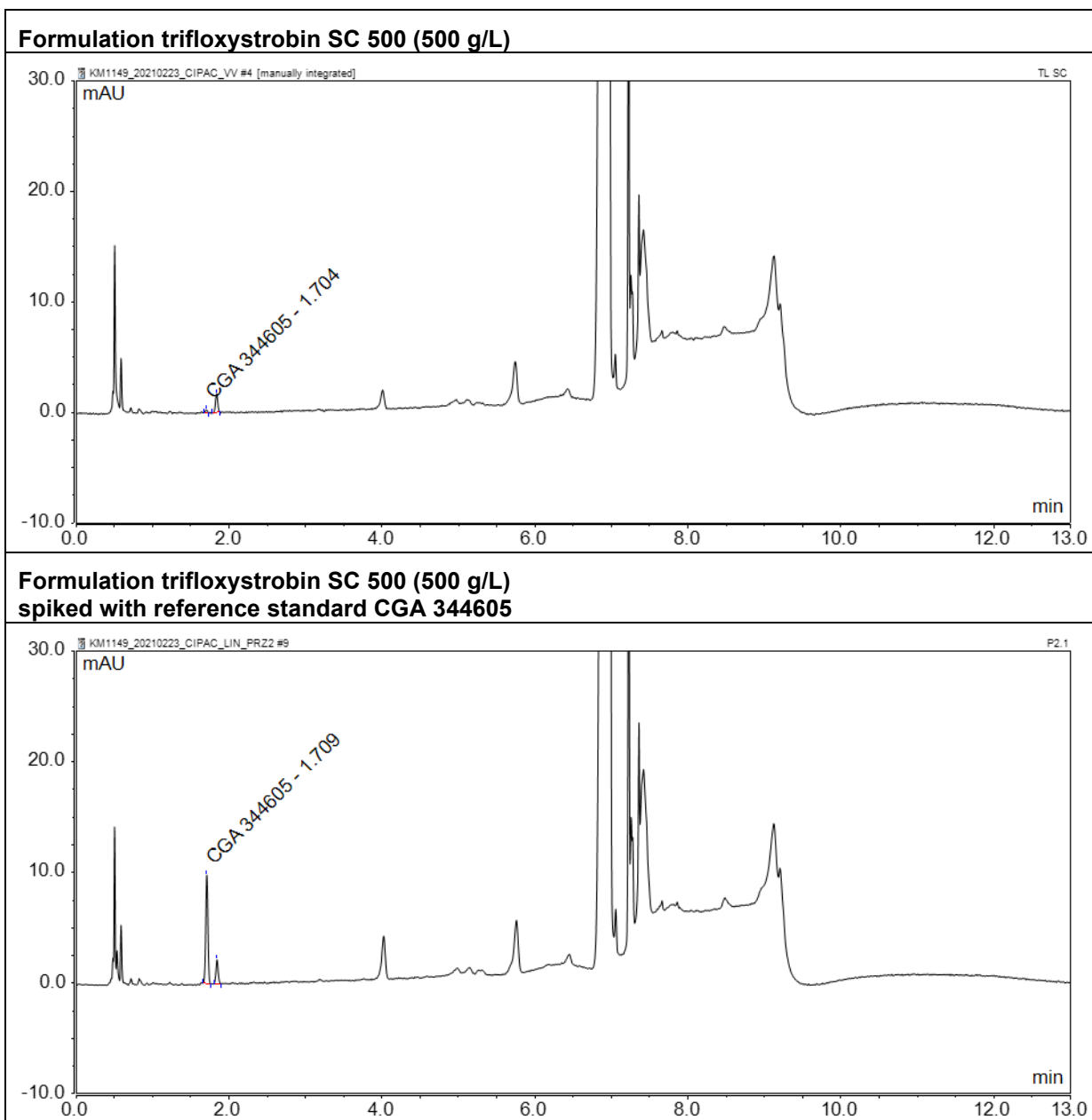




Figure 12. Continuation

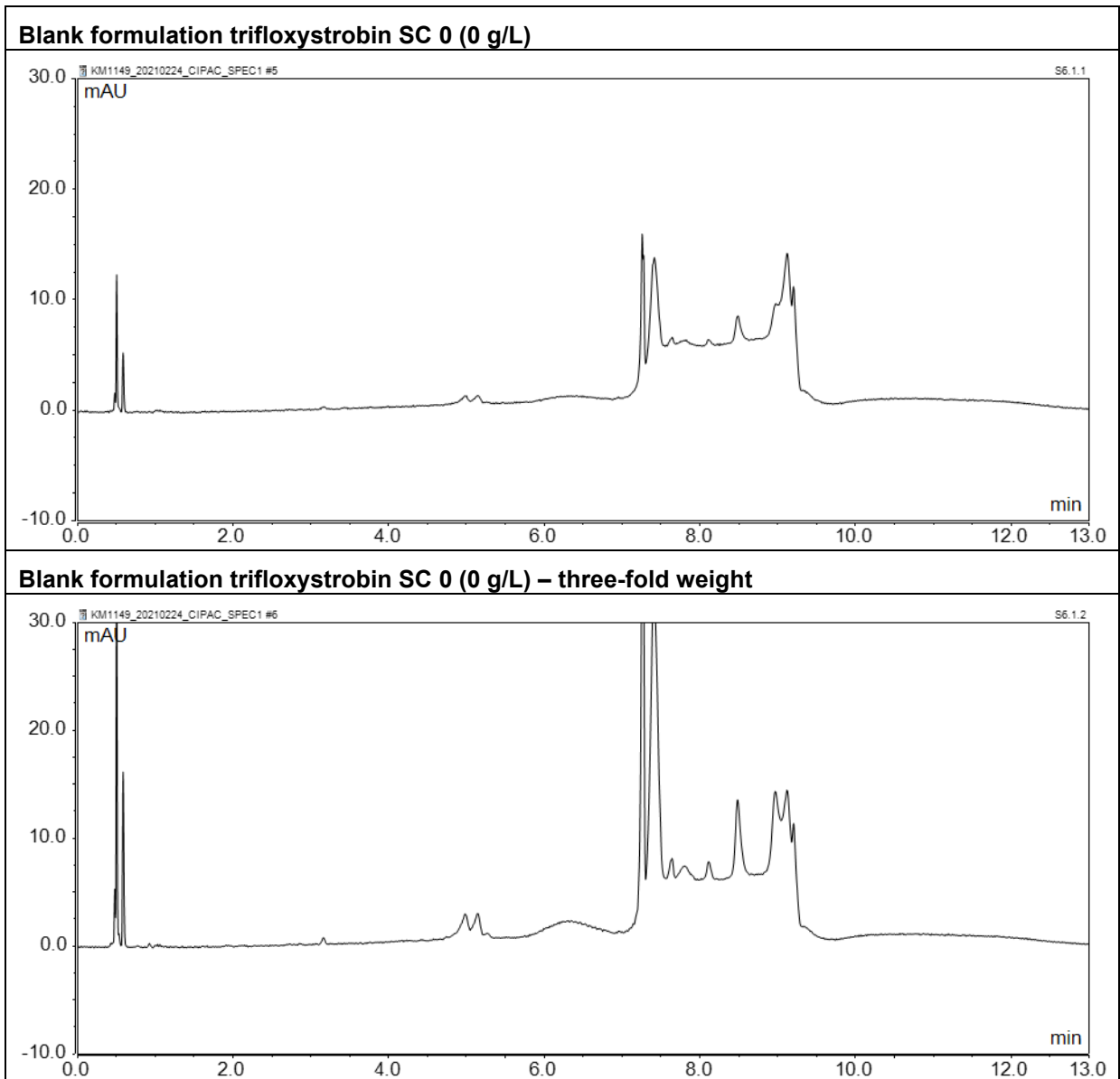
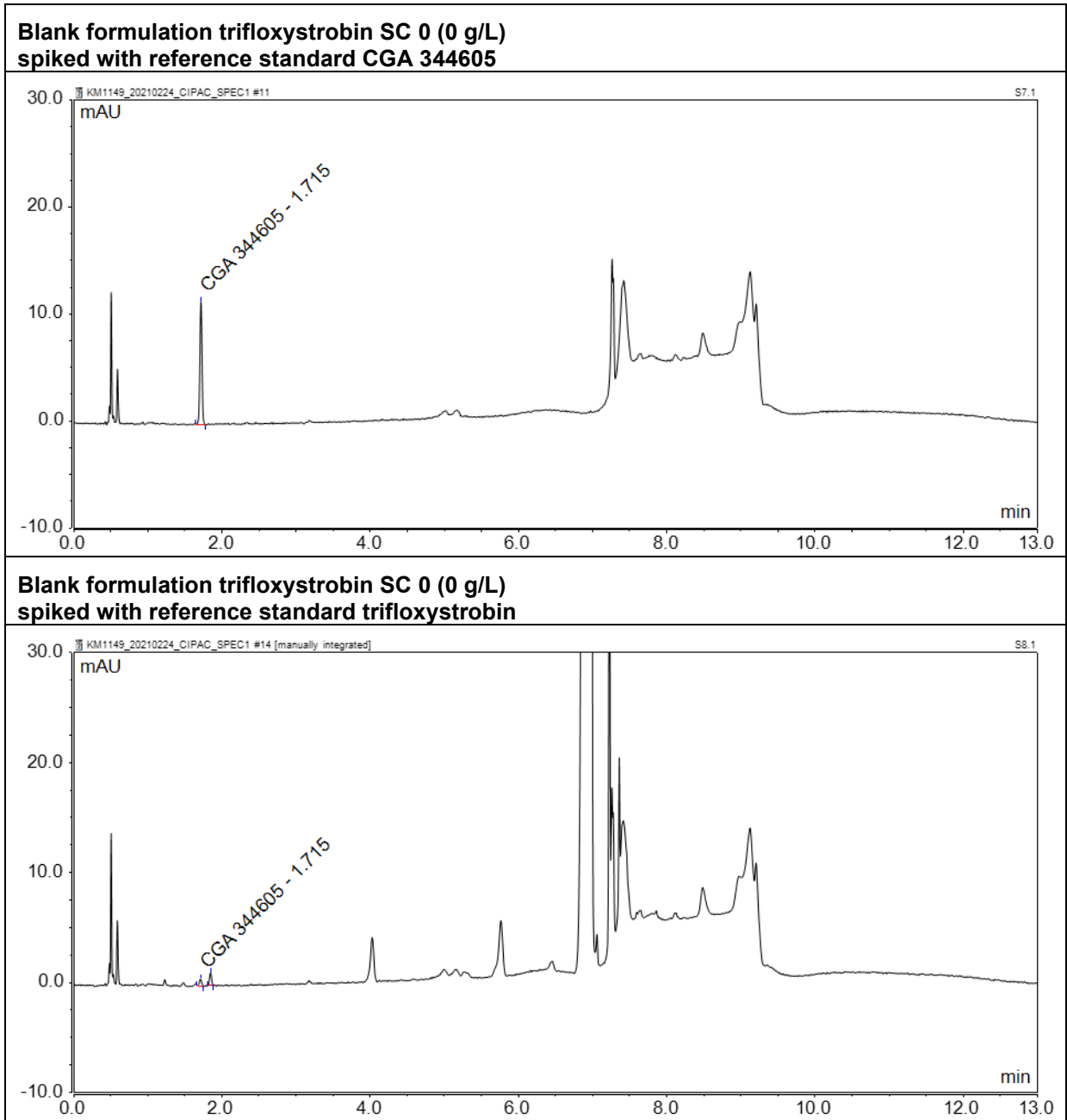


Figure 12. Continuation



**Figure 13: Chromatograms of fluoxastrobin + prothioconazole + trifloxystrobin EC 300 (75+150+75 g/L) Developing Lab**

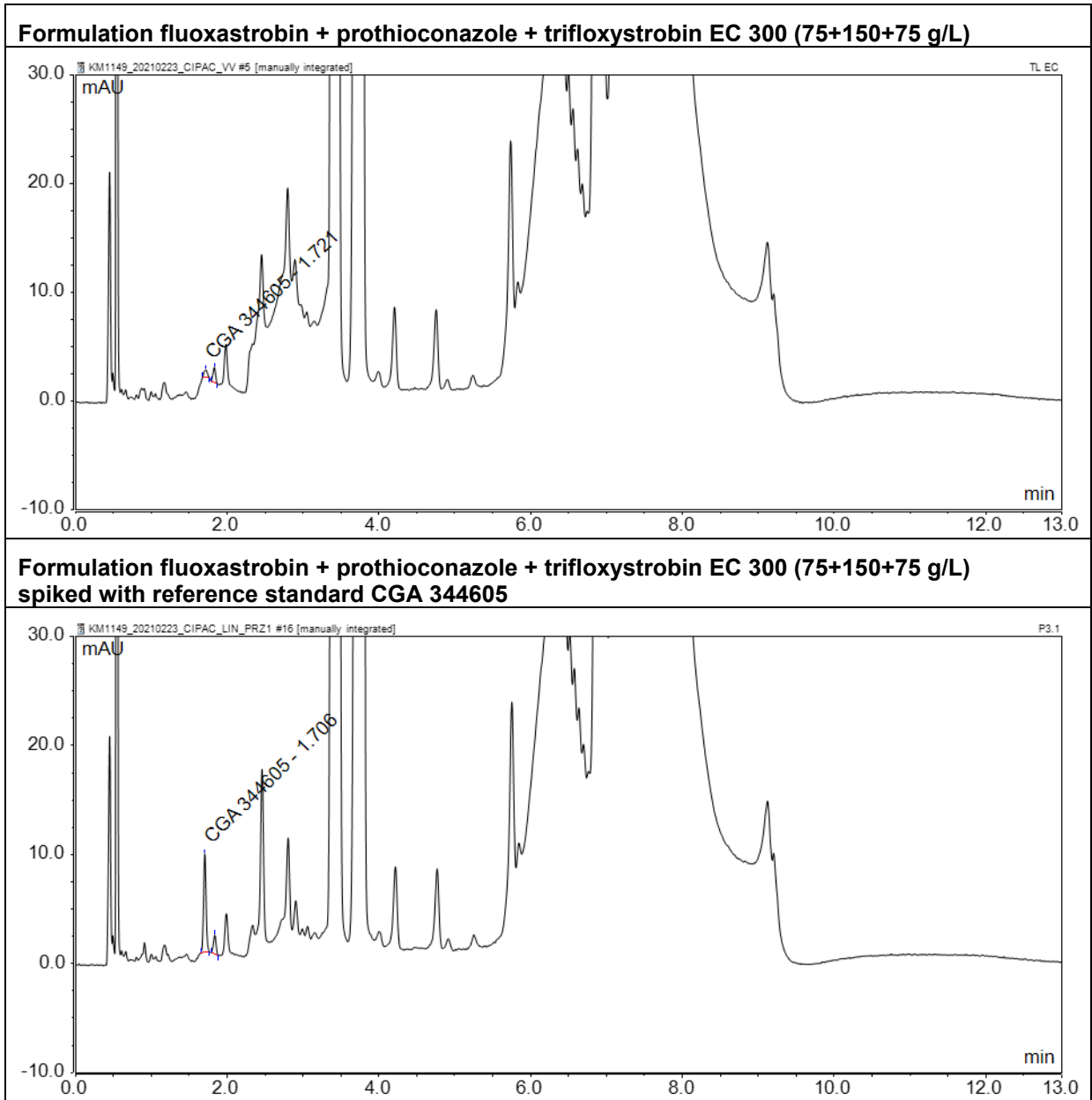


Figure 13. Continuation

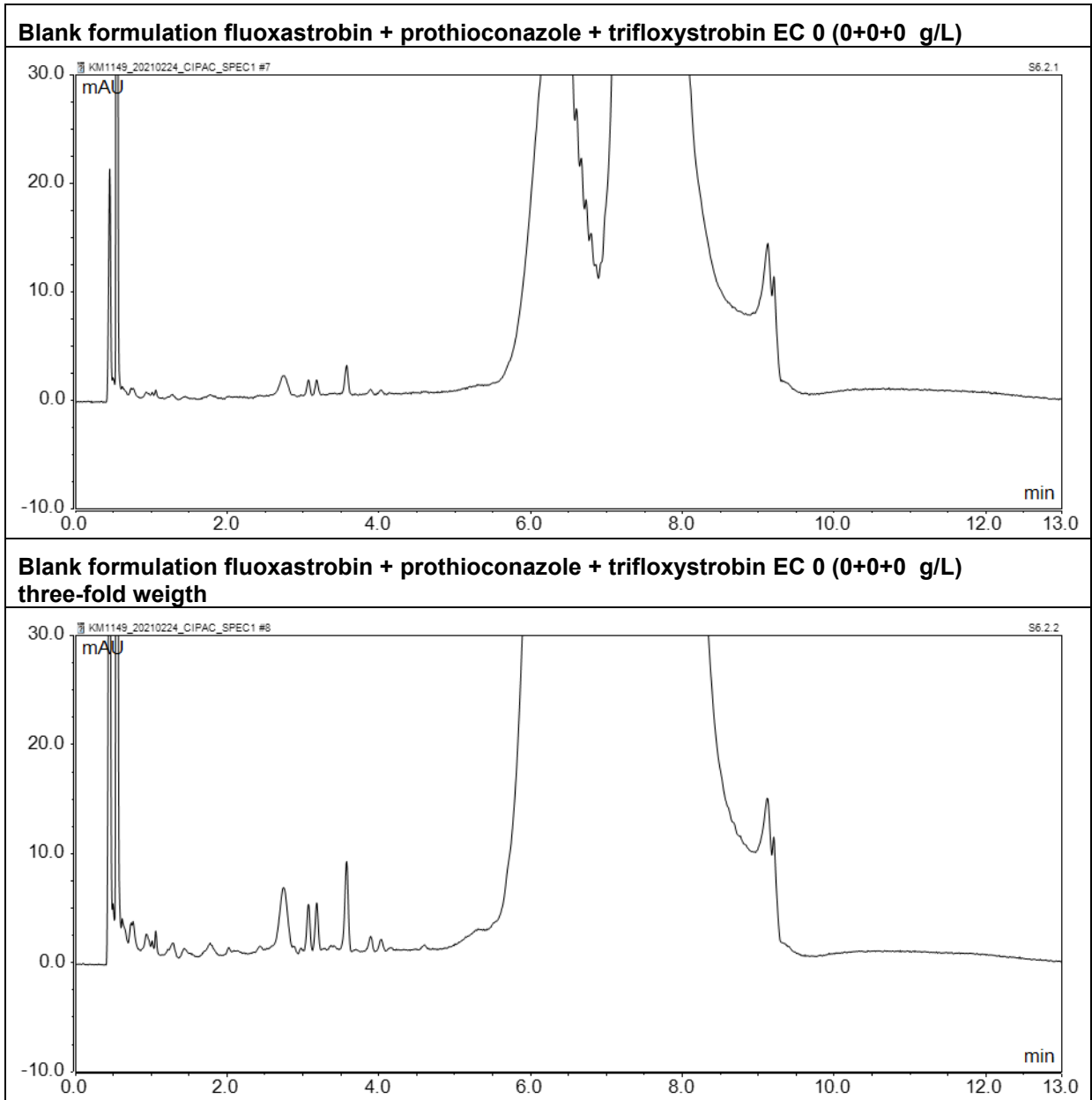
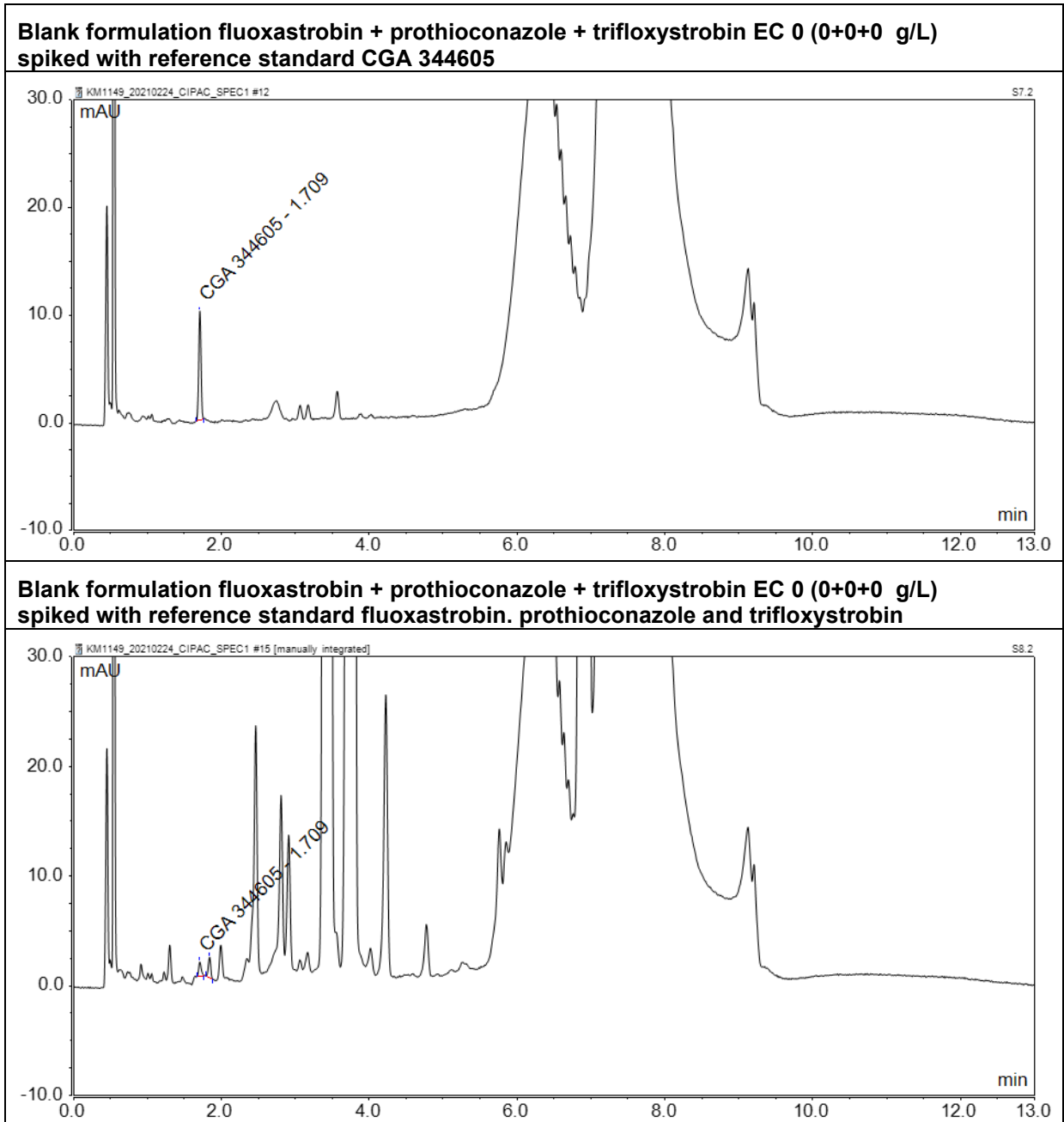


Figure 13. Continuation



**Figure 14: Chromatogram of trifloxystrobin WG 50 (50 g/kg)  
Developing Lab**

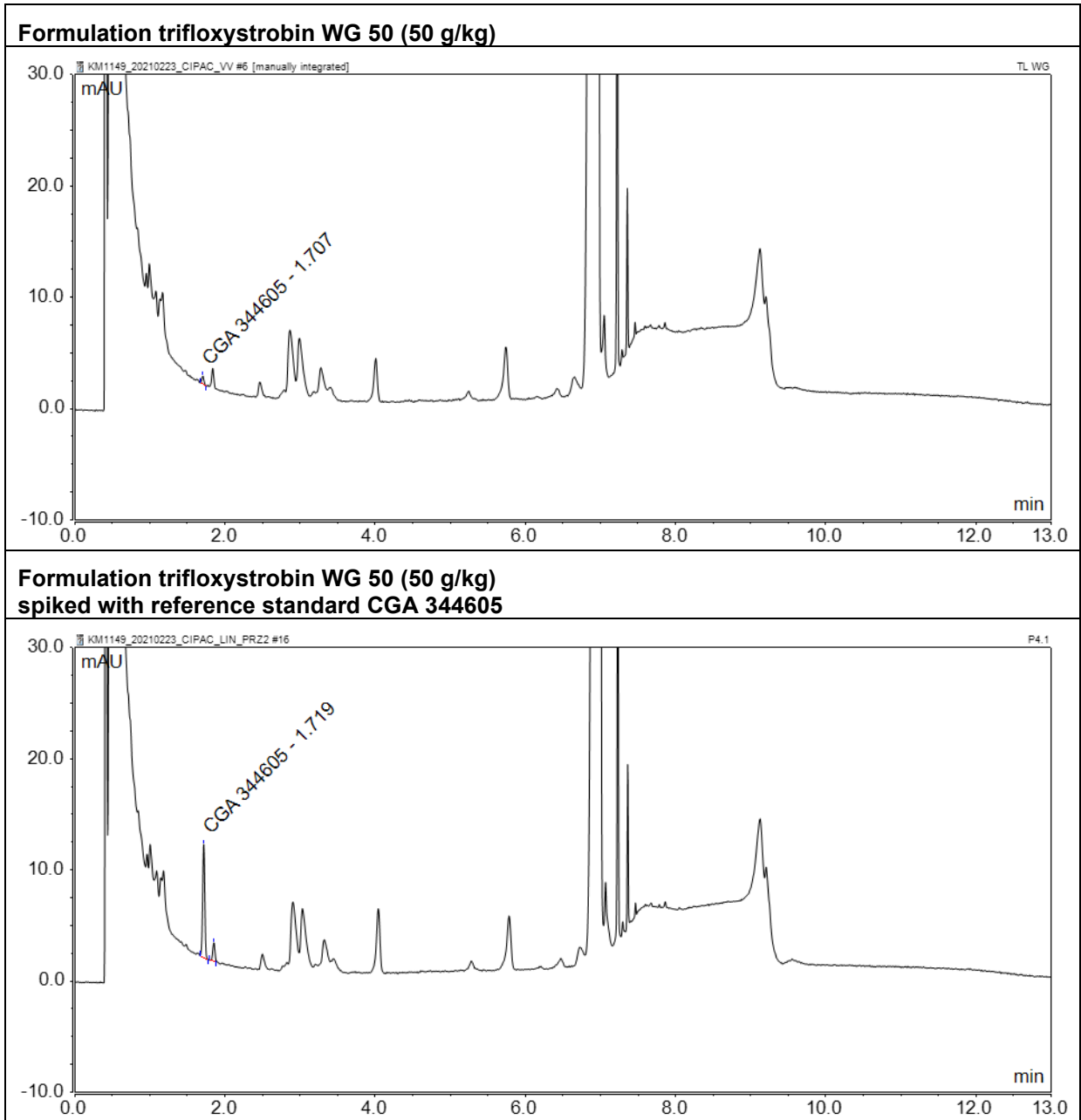


Figure 14. Continuation

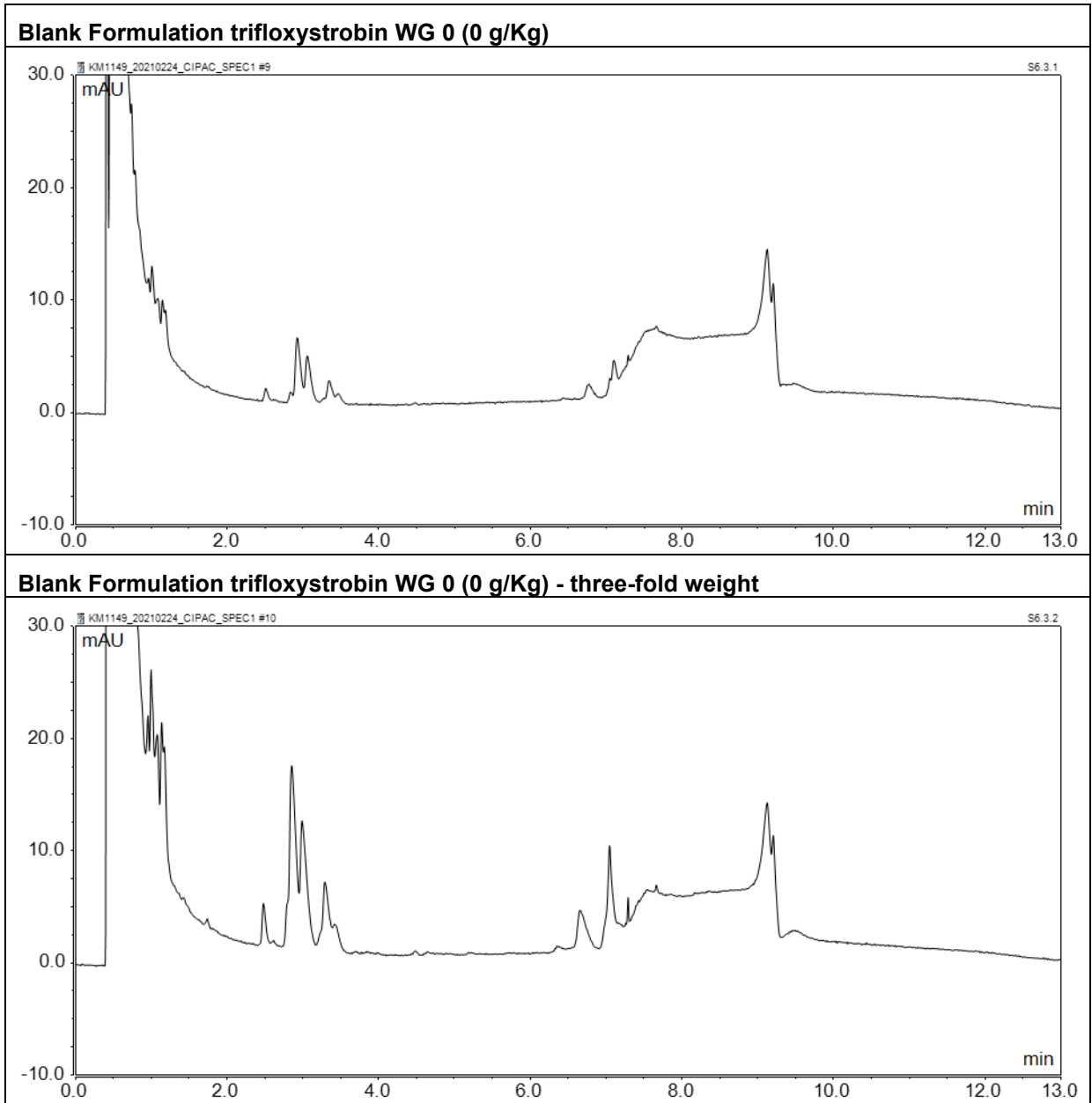
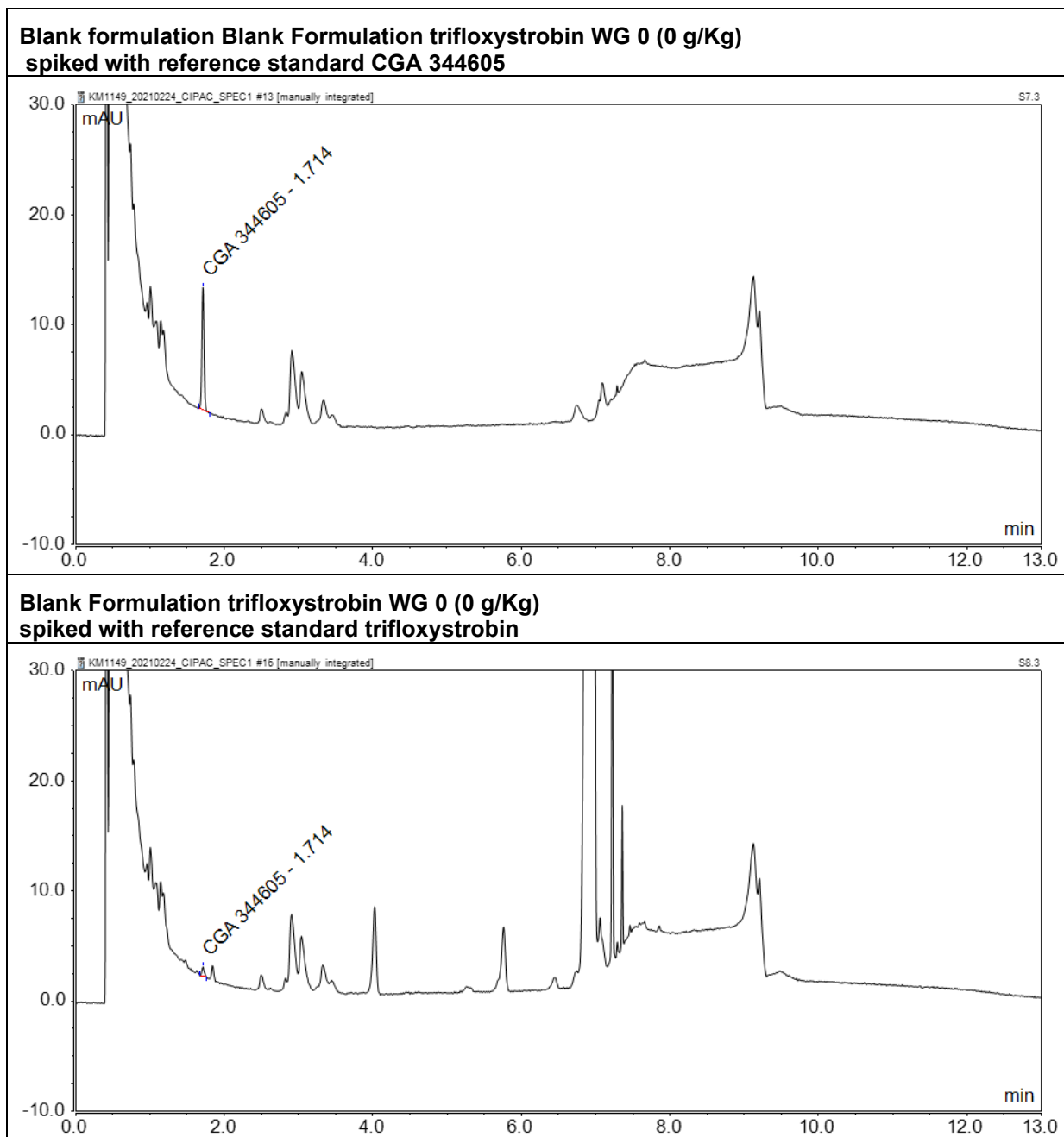
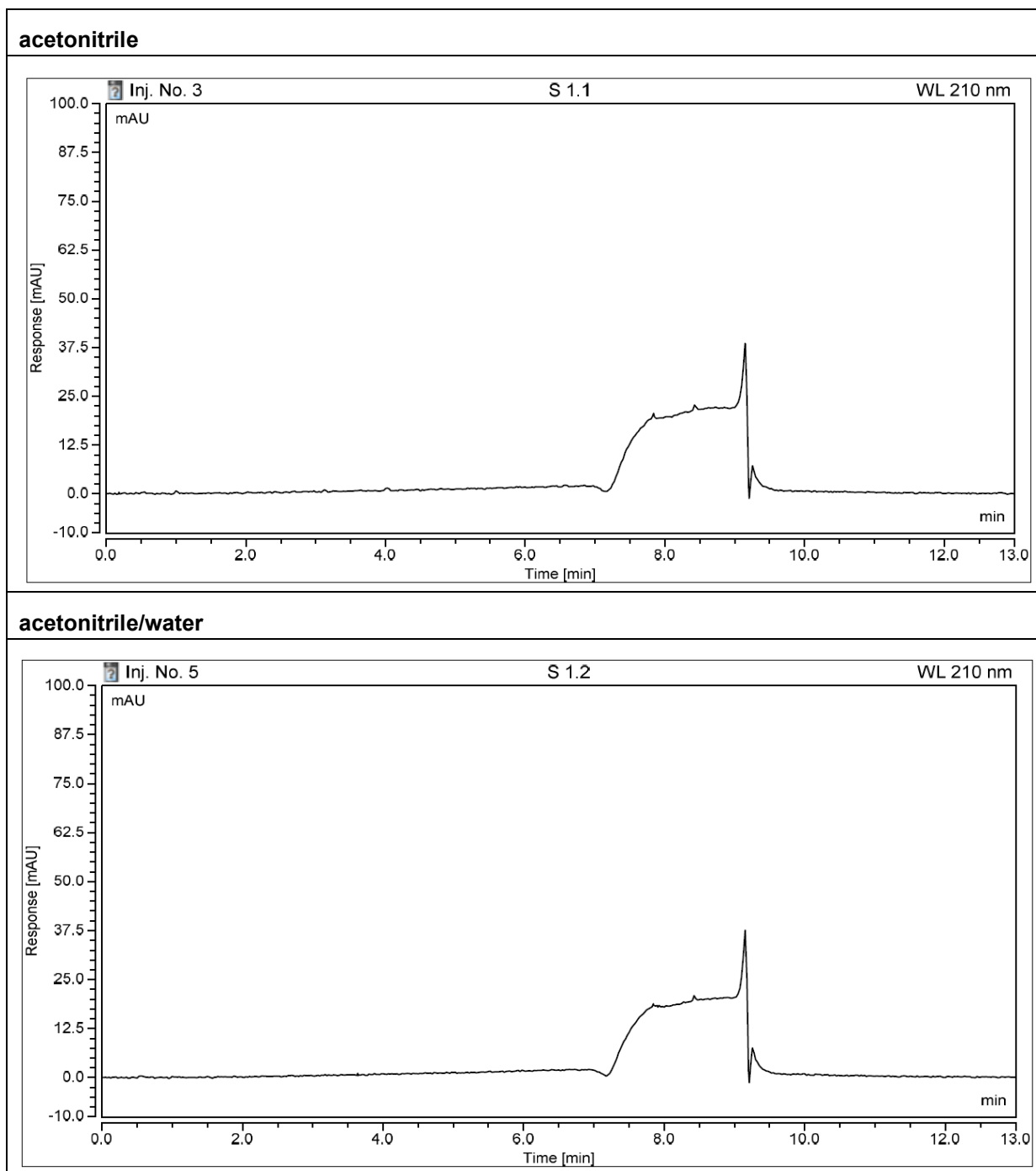


Figure 14. Continuation





**Figure 15: Chromatograms of sample solvent  
Lab 1**



**Figure 16: Chromatograms of reference standard CGA 344605  
Lab 1**

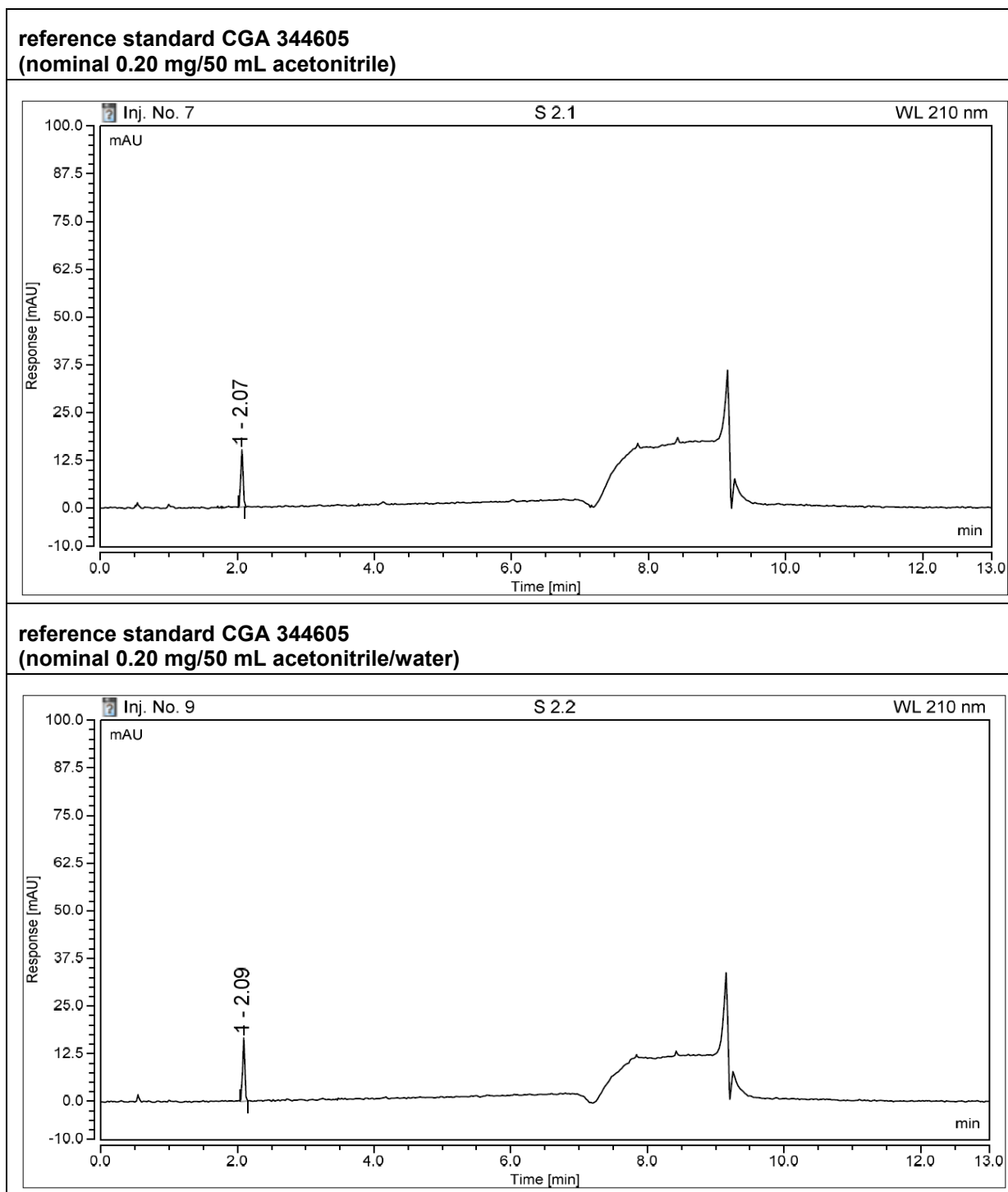
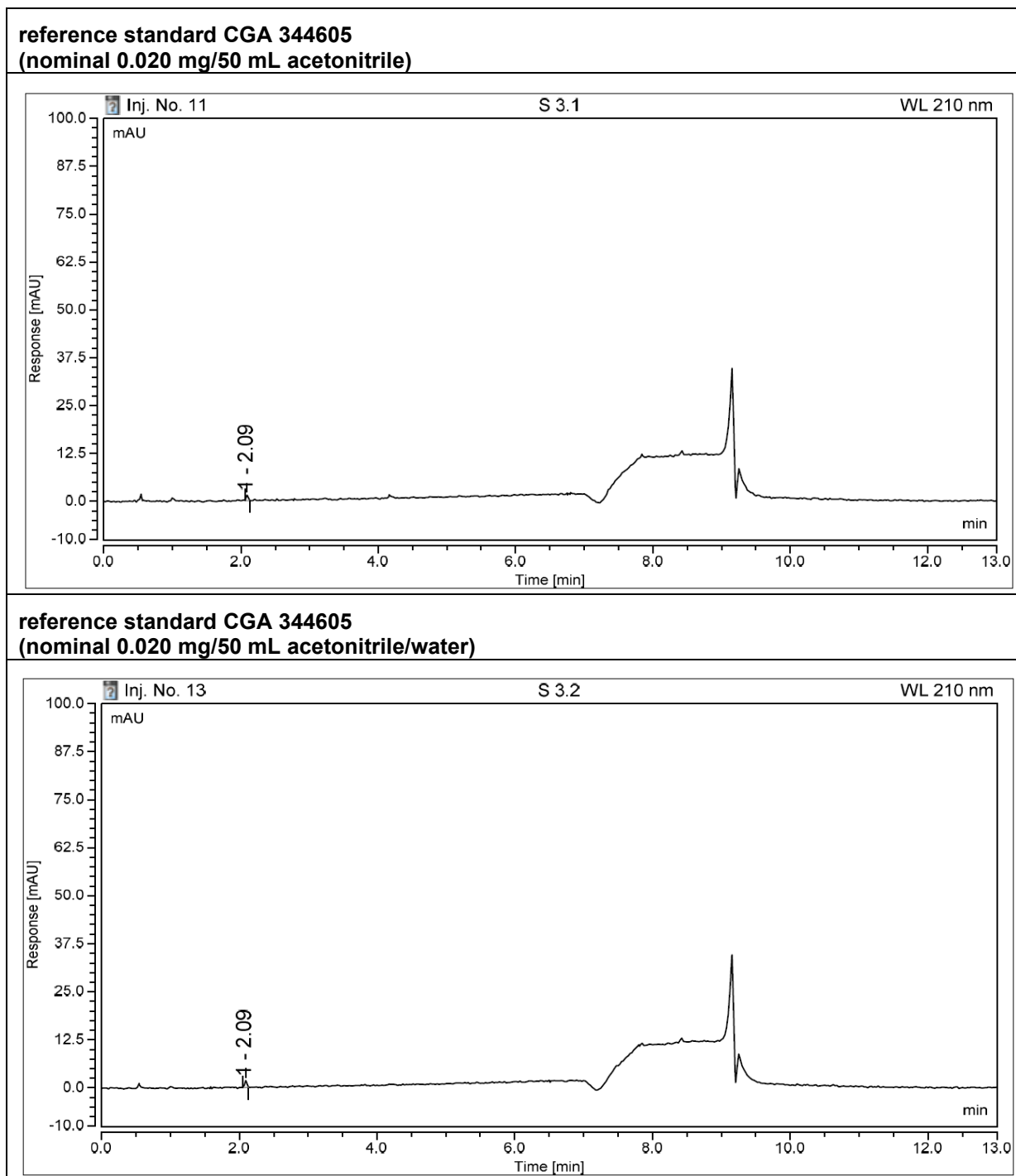
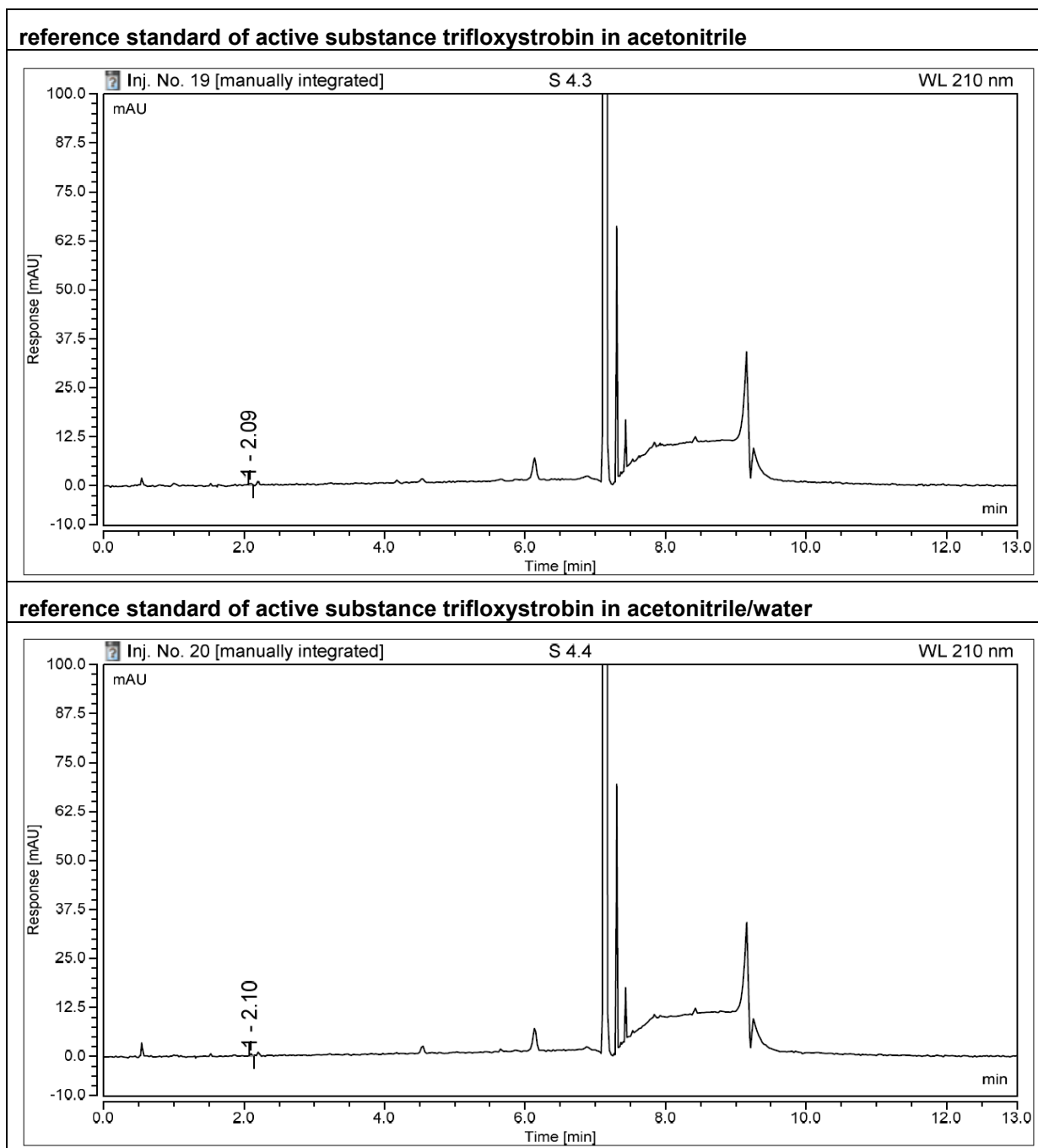


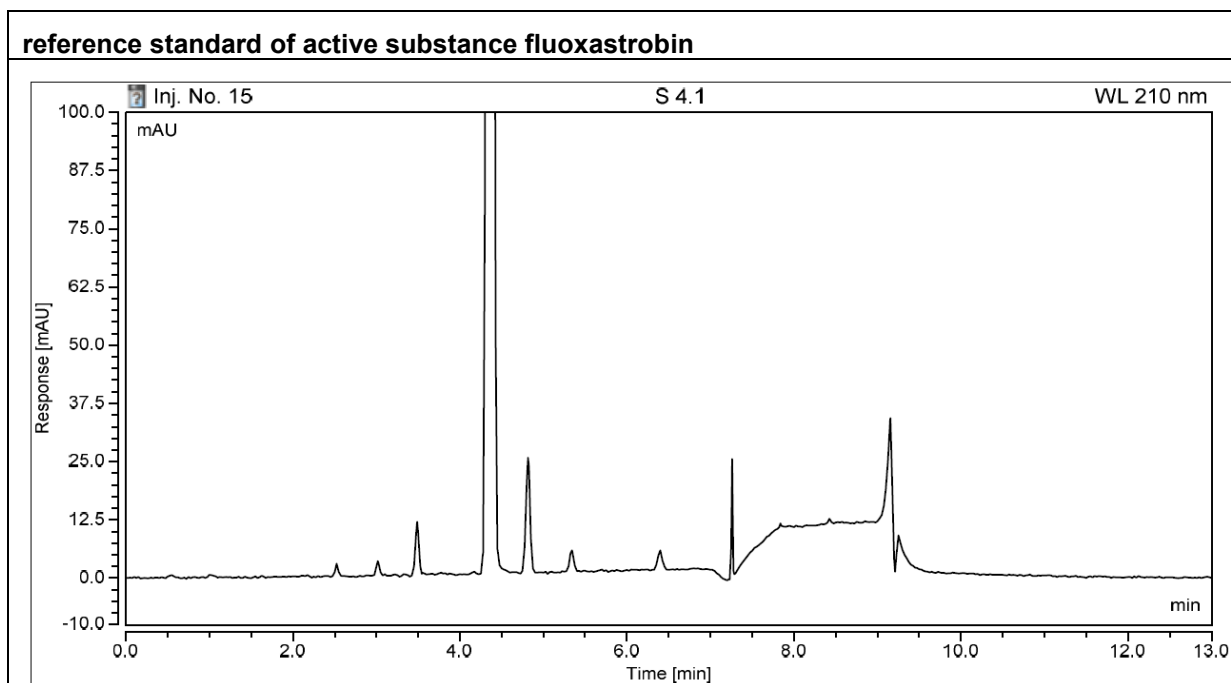
Figure 16. Continuation



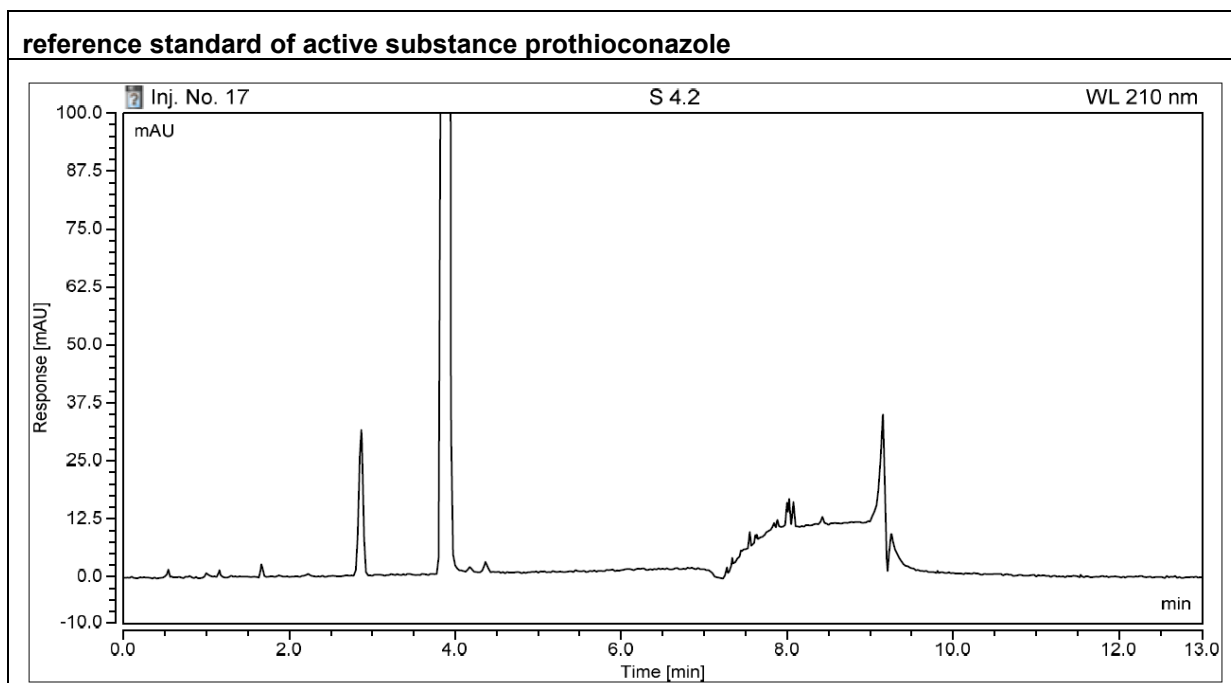
**Figure 17: Chromatograms of active substance trifloxystrobin  
Lab 1**



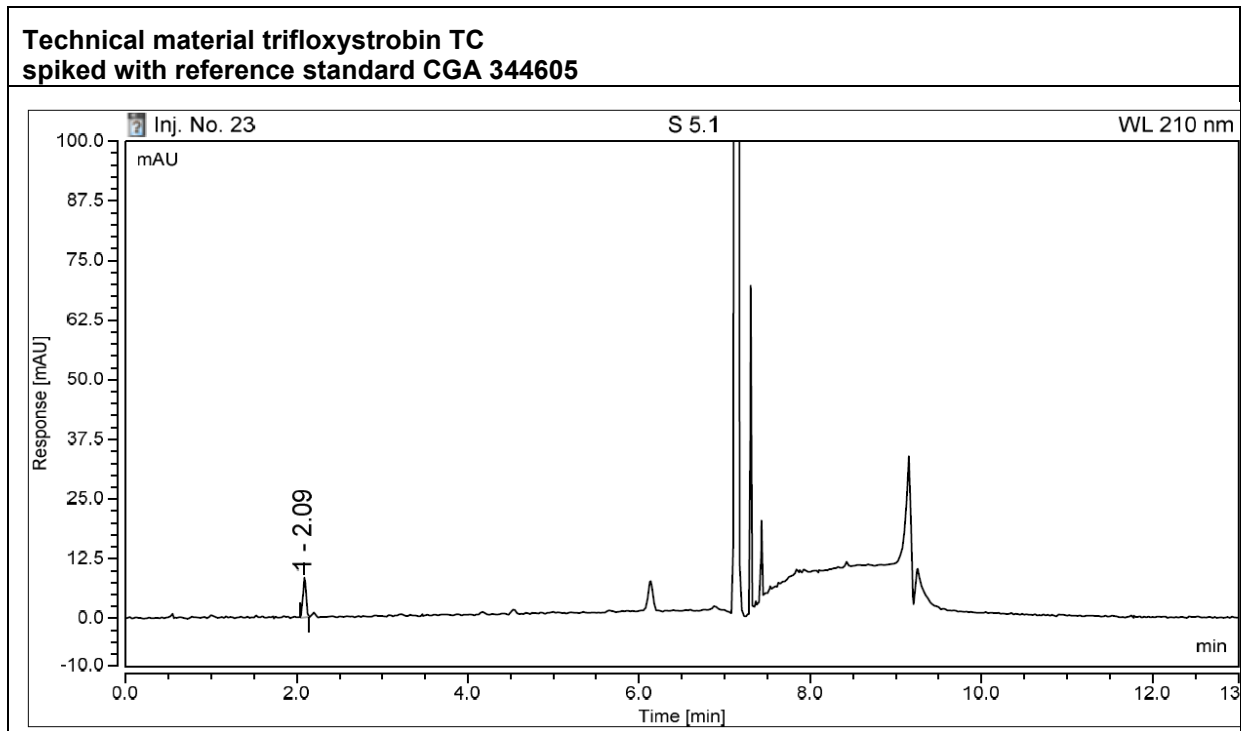
**Figure 18: Chromatograms of active substance fluoxastrobin.  
Lab 1**



**Figure 19: Chromatograms of active substance prothioconazole  
Lab 1**



**Figure 20: Chromatograms of trifloxystrobin TC  
Lab 1**



**Figure 21: Chromatograms of trifloxystrobin SC 500 (500 g/L)  
Lab 1**

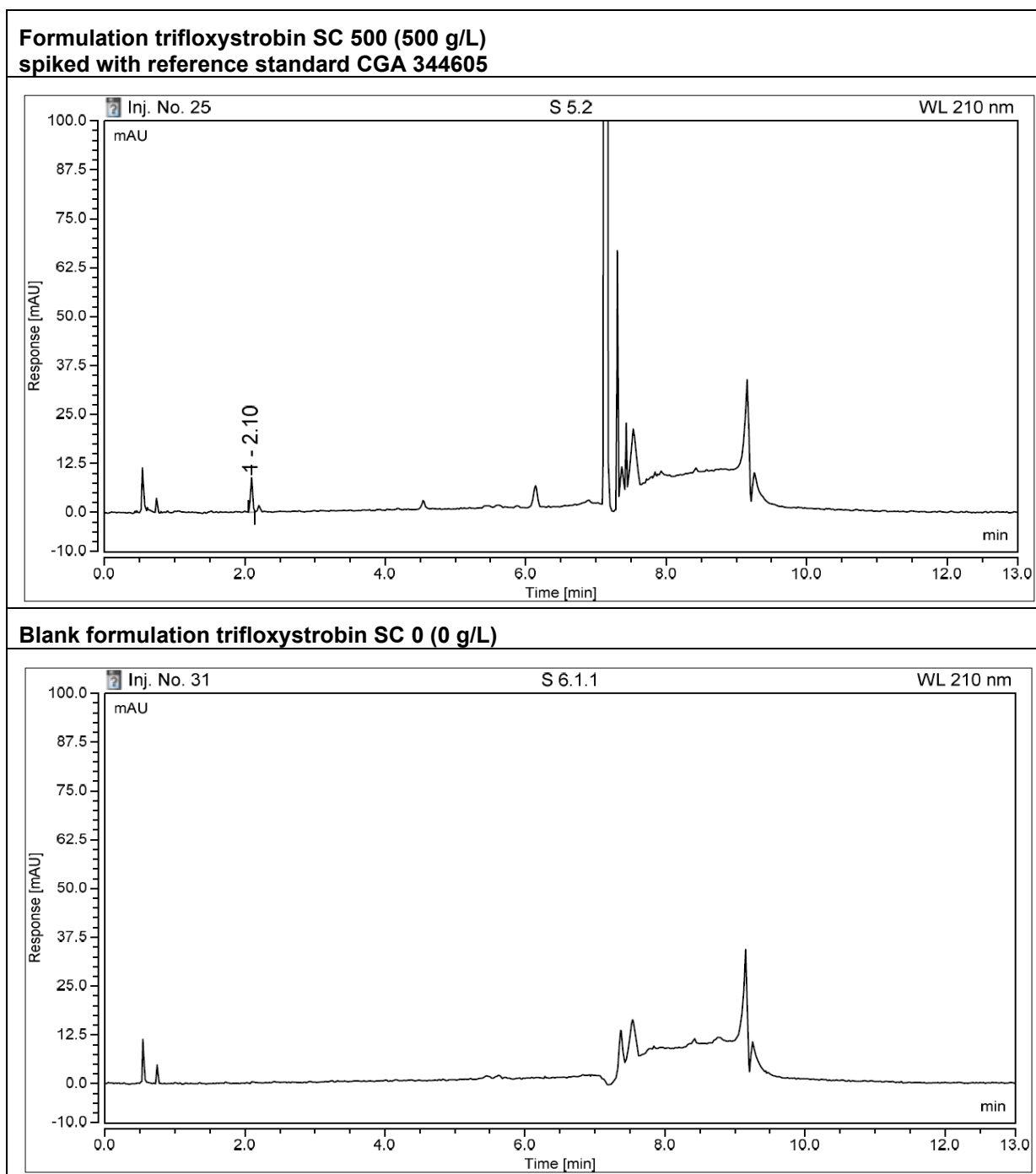
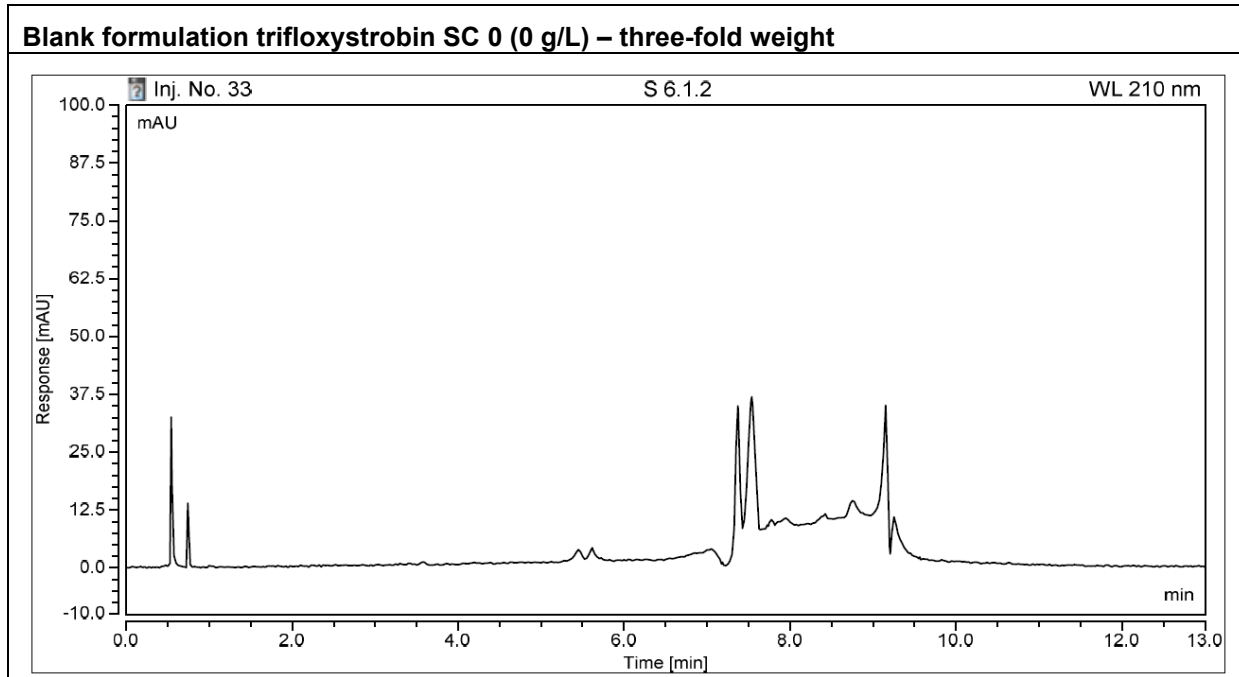


Figure 21. Continuation





**Figure 22: Chromatograms of fluoxastrobin + prothioconazole + trifloxystrobin EC 300 (75+150+75 g/L) Lab 1**

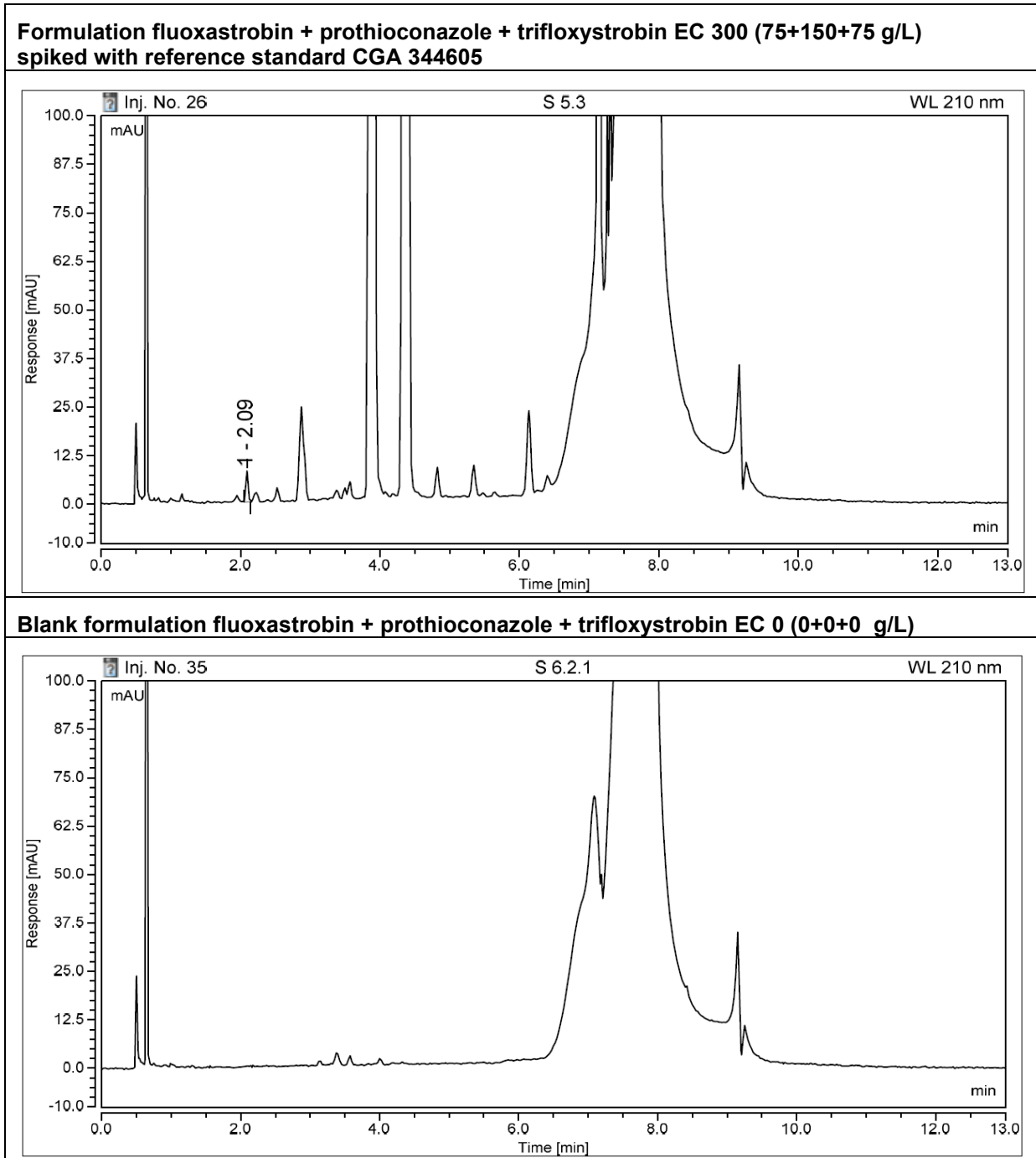
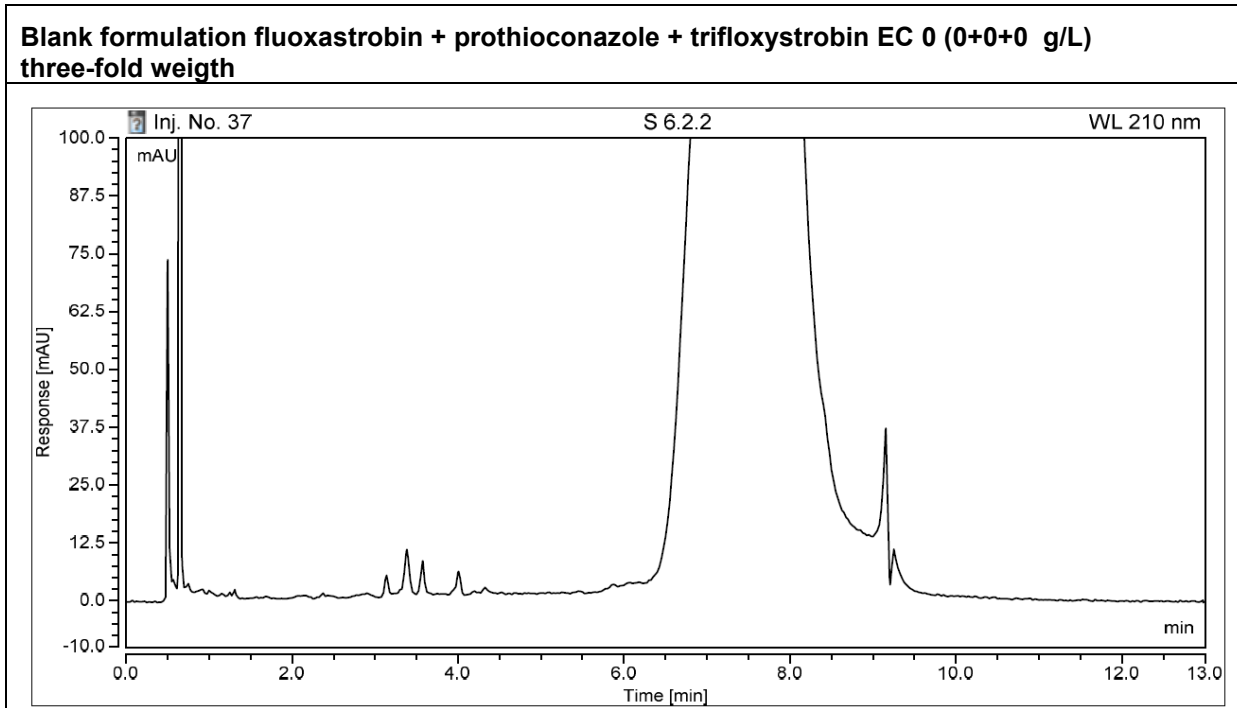


Figure 22. Continuation



**Figure 23: Chromatogram of trifloxystrobin WG 50 (50 g/kg)  
Lab 1**

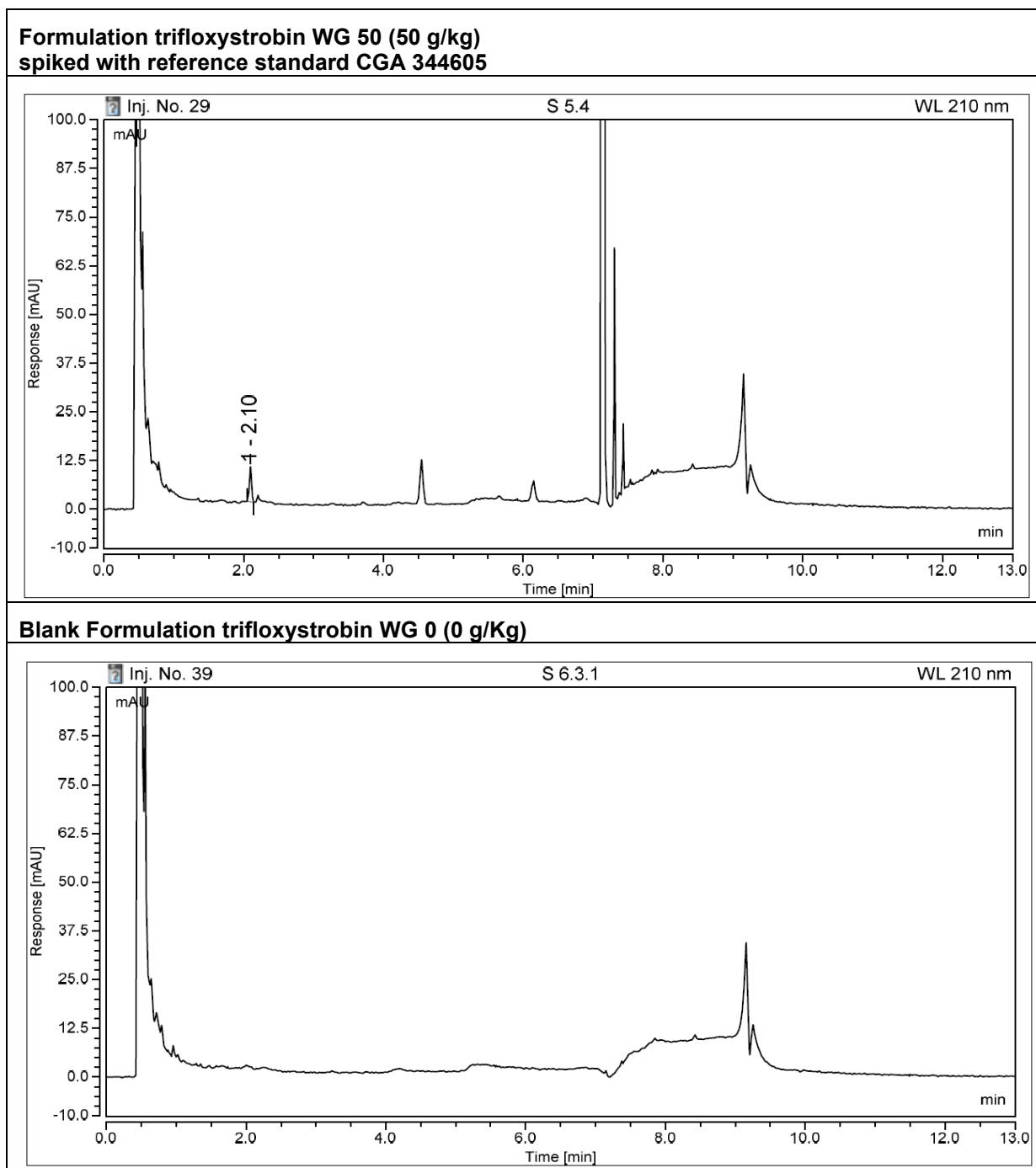
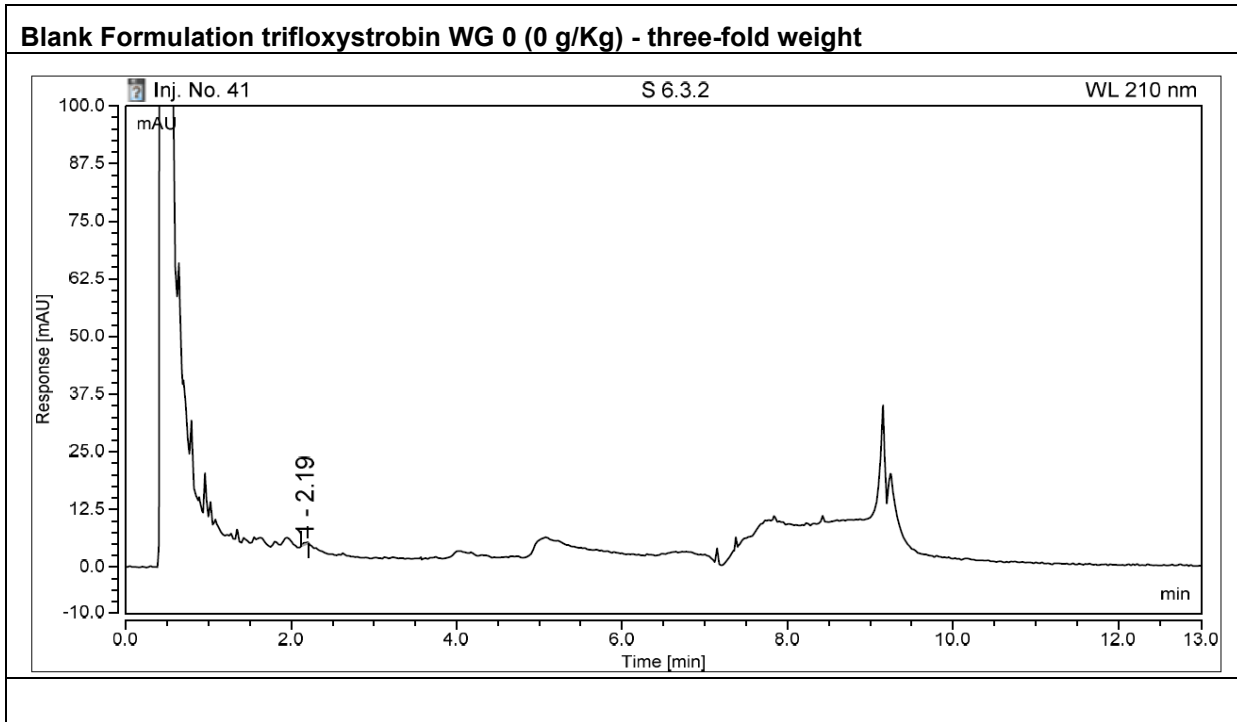
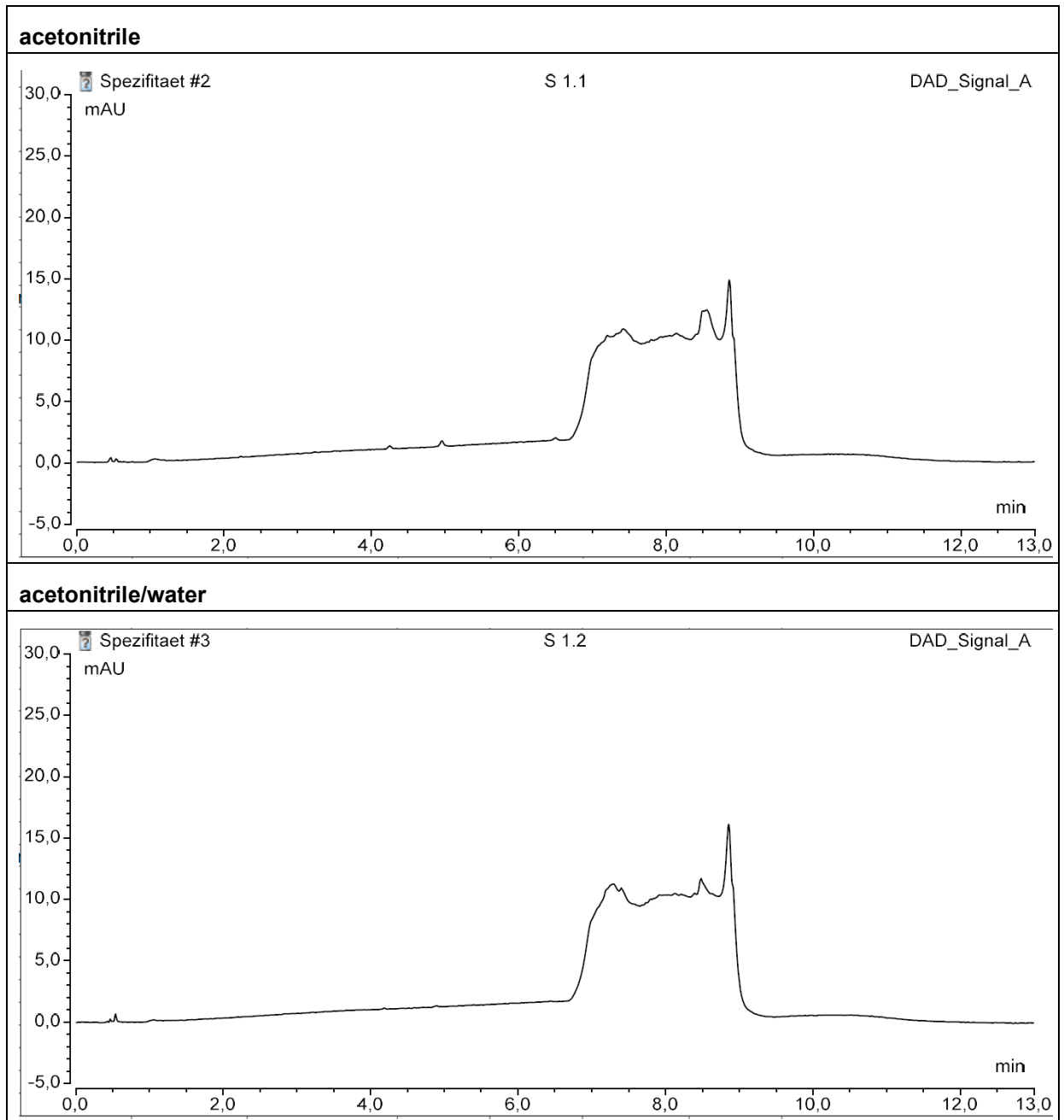


Figure 23. Continuation



**Figure 24: Chromatograms of sample solvent  
Lab 2**



**Figure 25: Chromatograms of reference standard CGA 344605  
Lab 2**

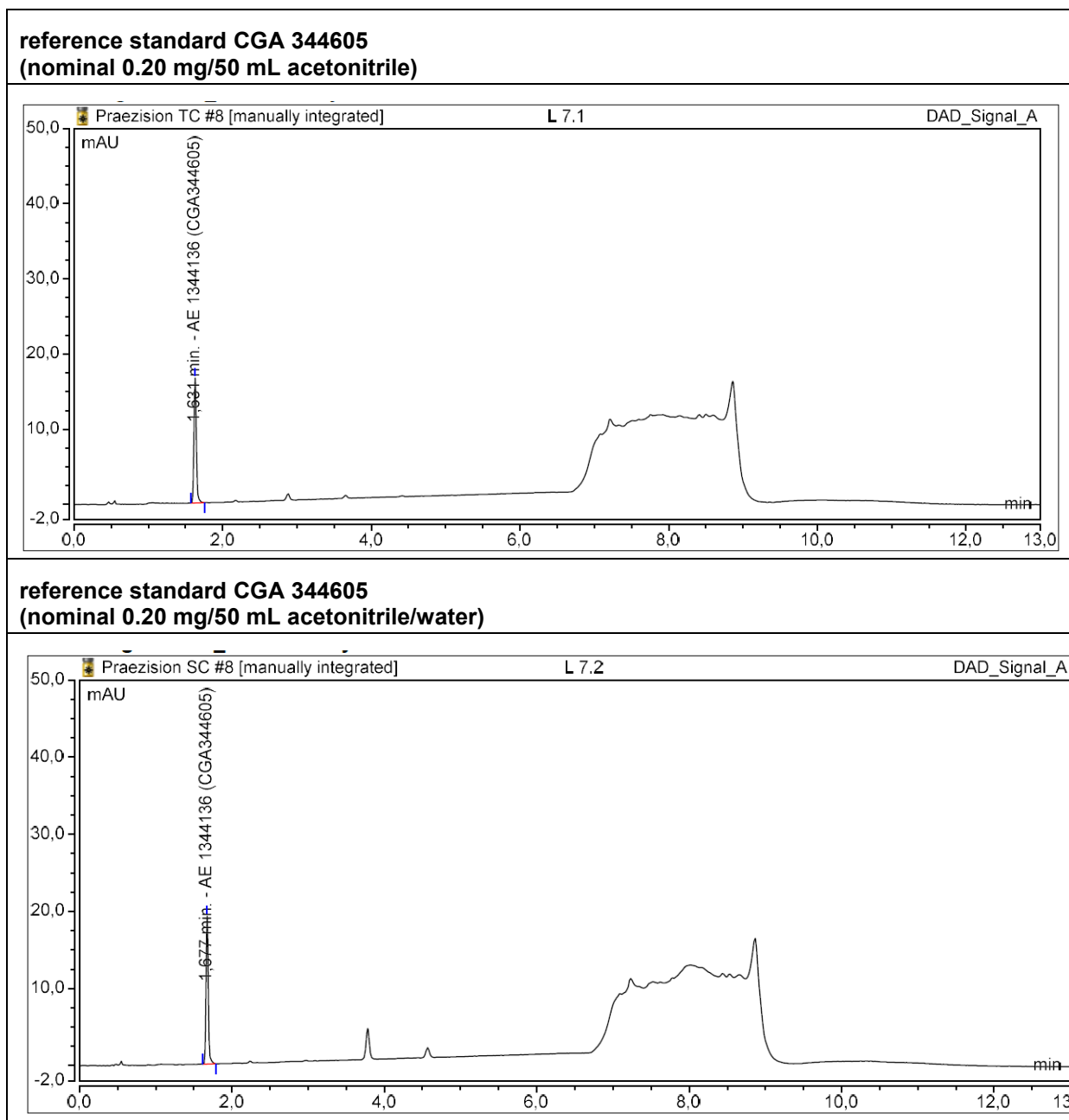
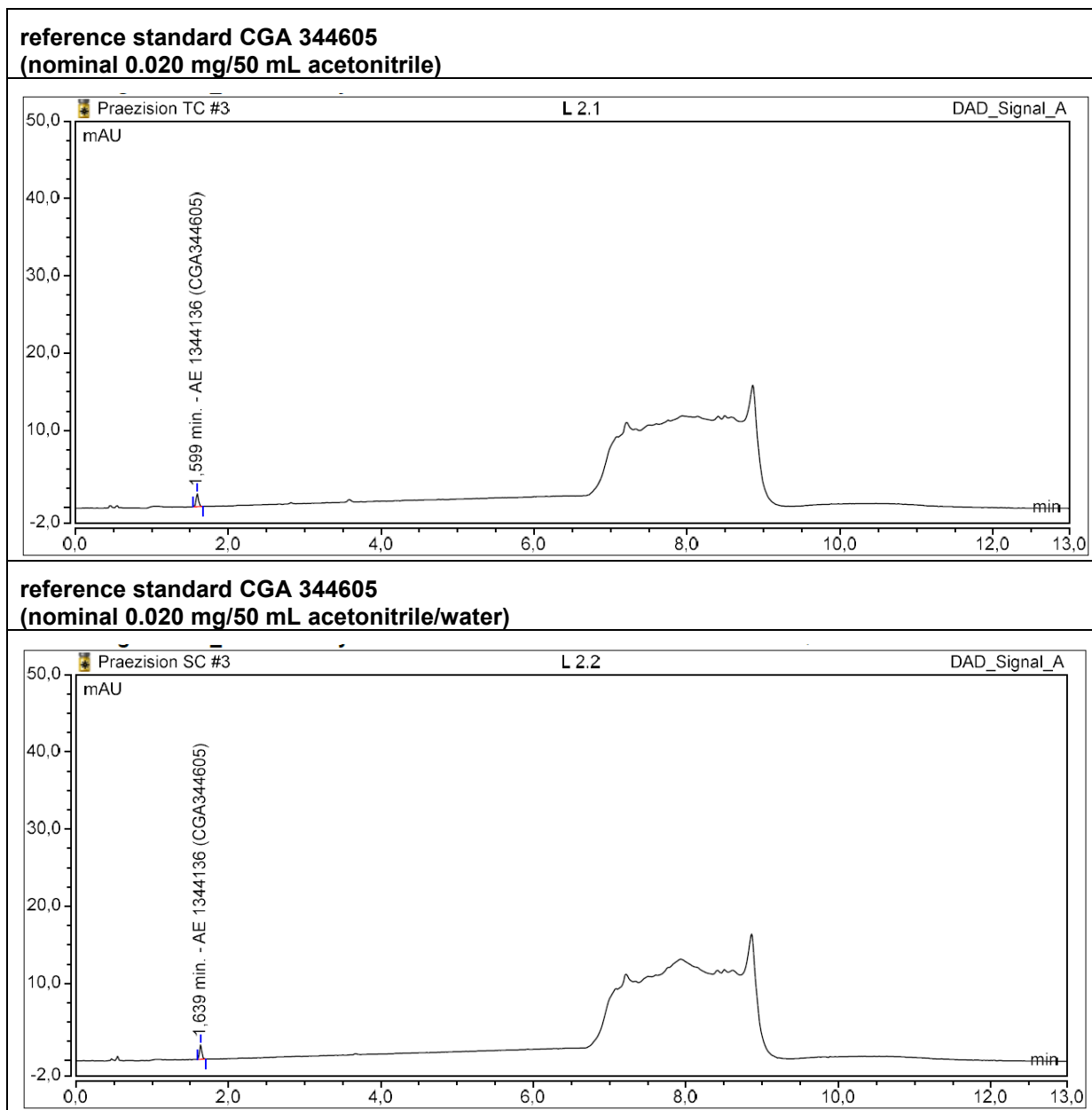
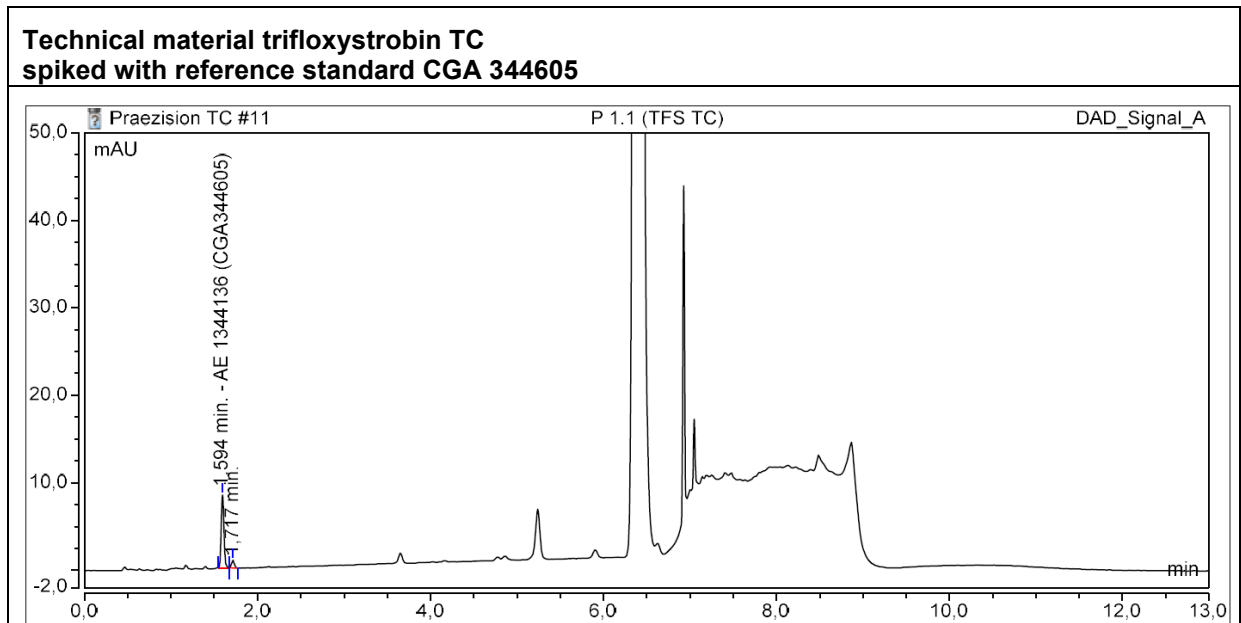


Figure 25. Continuation



**Figure 26: Chromatograms of trifloxystrobin TC  
Lab 2**





**Figure 27: Chromatograms of trifloxystrobin SC 500 (500 g/L)  
Lab 2**

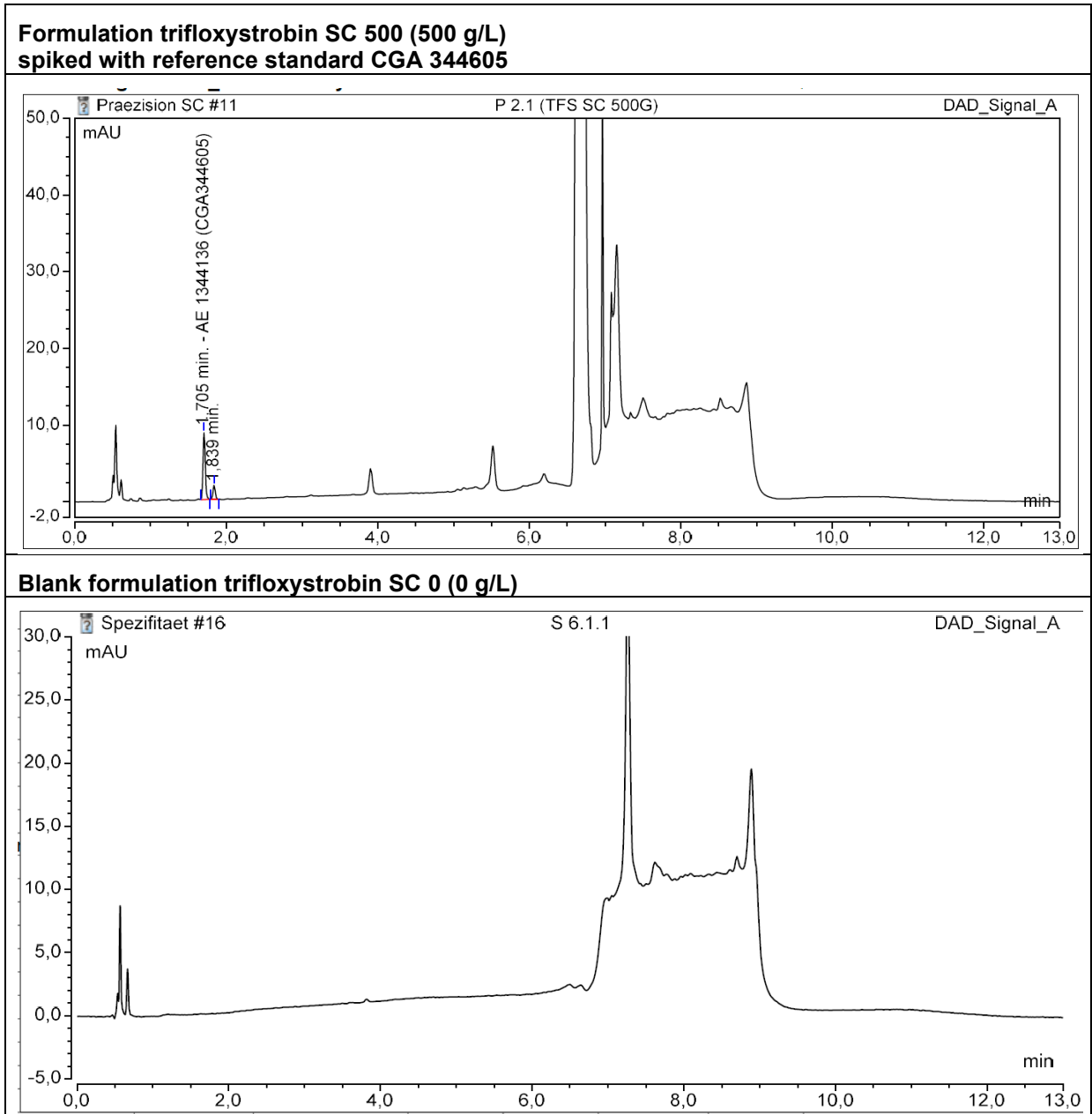
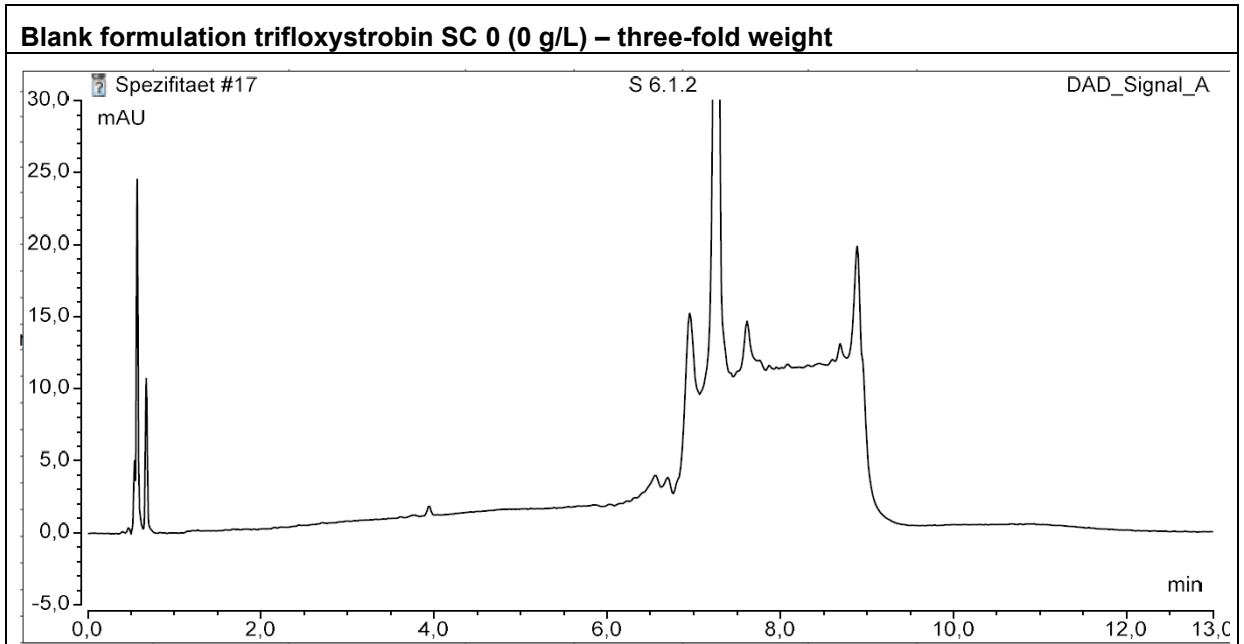


Figure 27. Continuation



**Figure 28: Chromatograms of fluoxastrobin + prothioconazole + trifloxystrobin EC 300 (75+150+75 g/L) Lab 2**

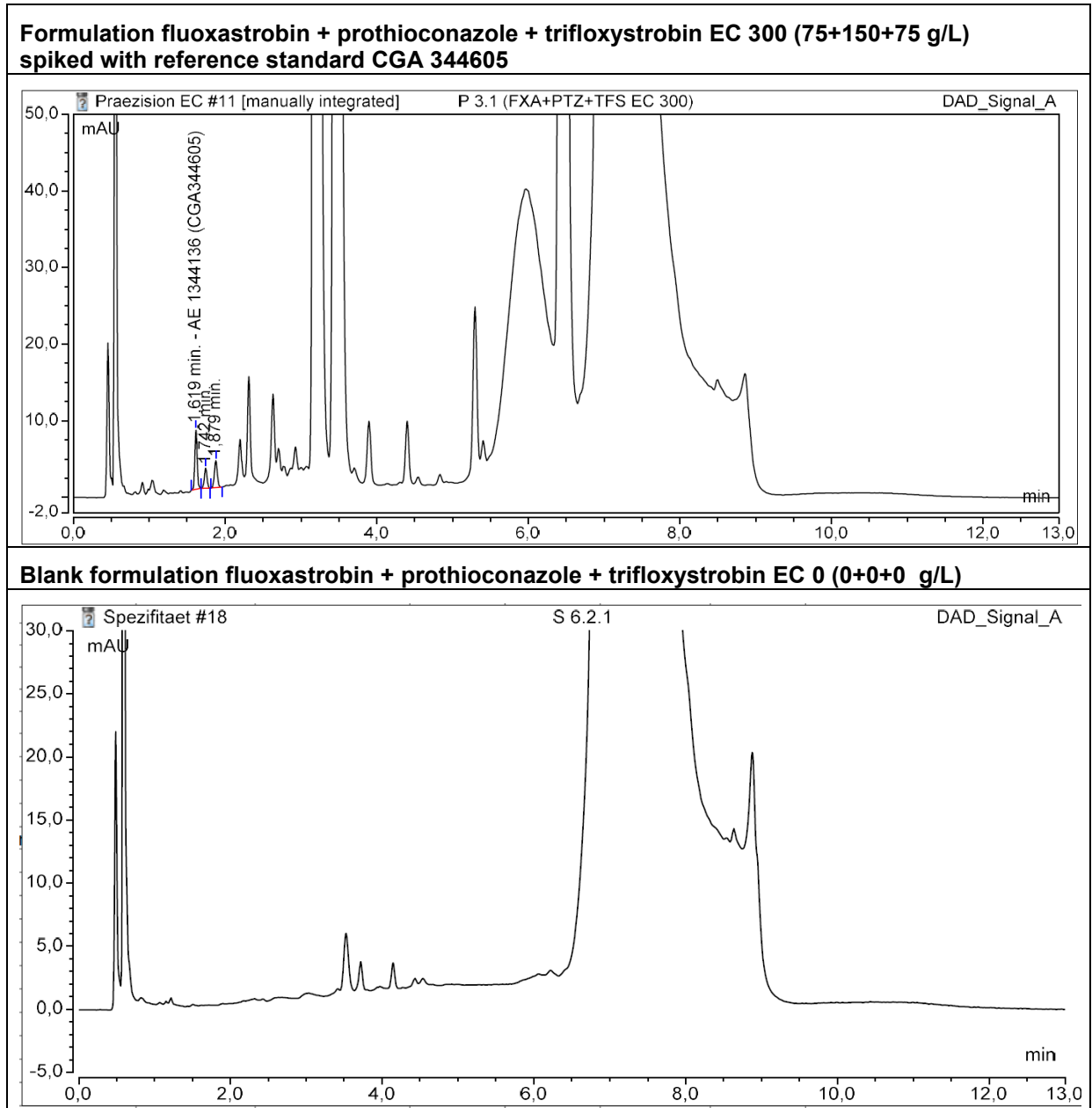
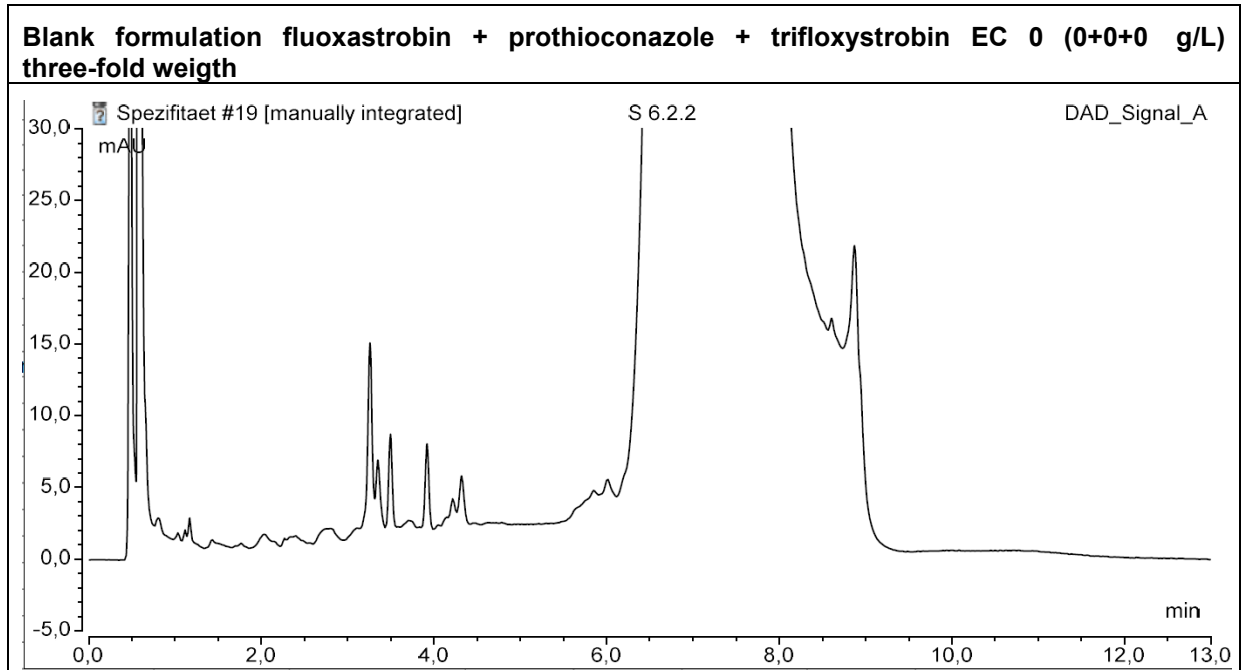


Figure 28. Continuation



**Figure 29: Chromatogram of trifloxystrobin WG 50 (50 g/kg)  
Lab 2**

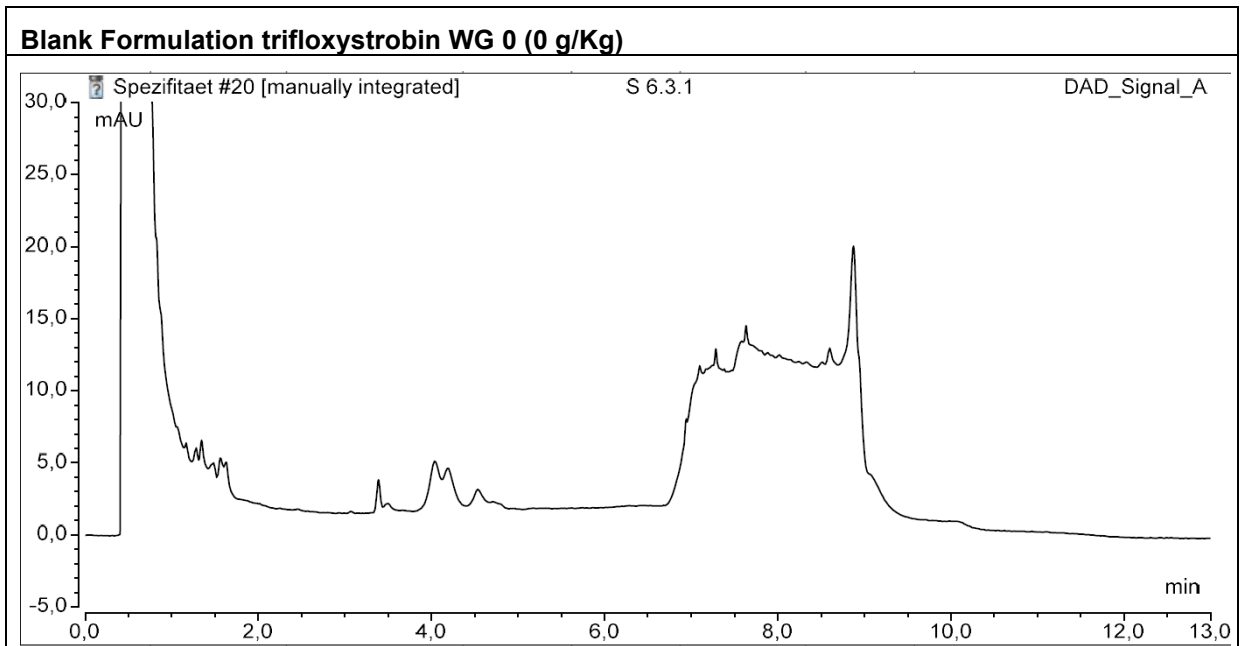
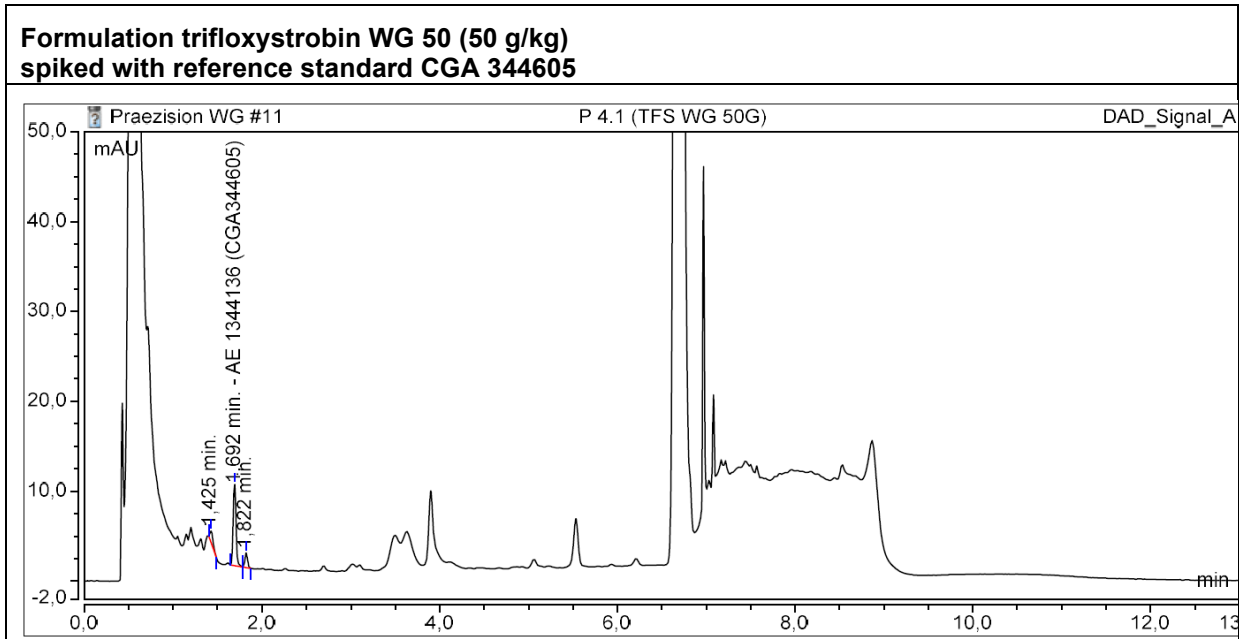
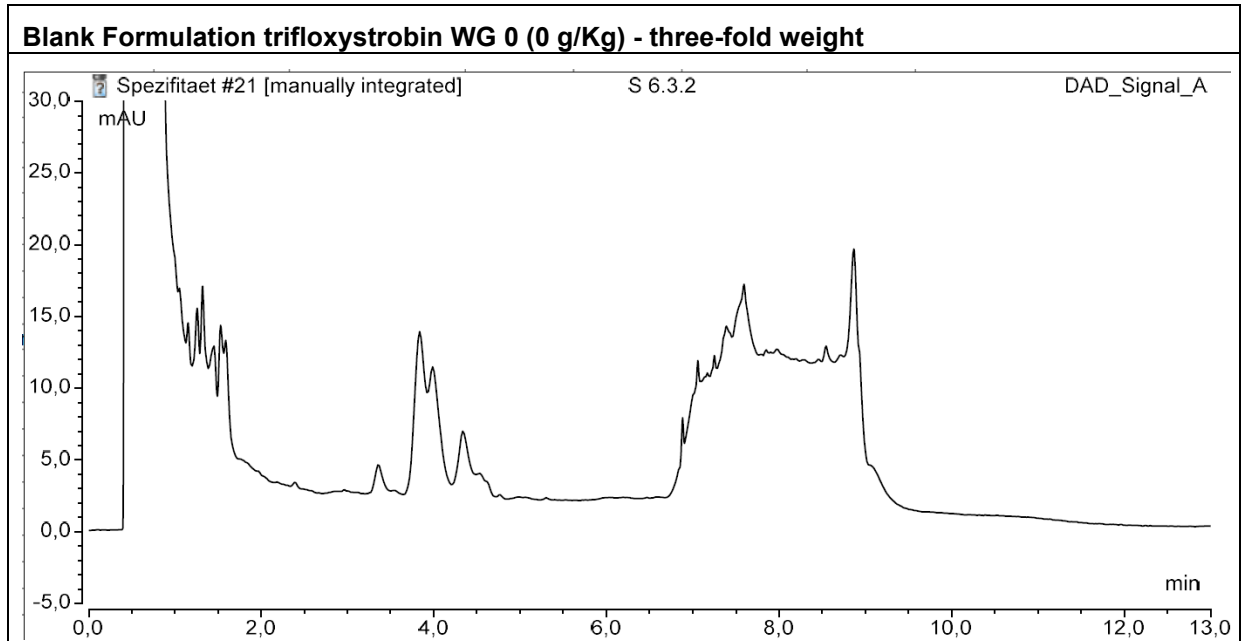
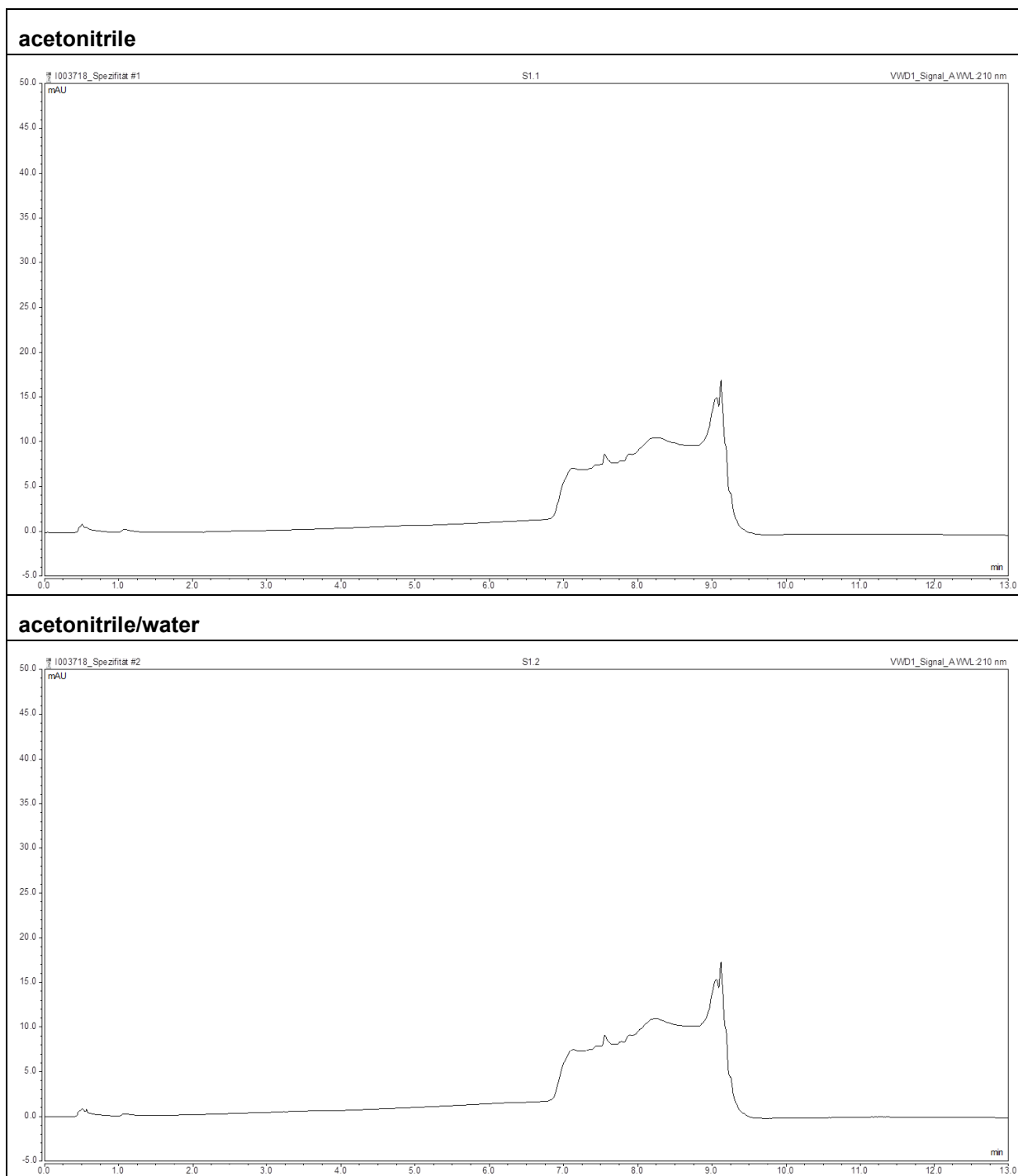


Figure 29. Continuation



**Figure 30: Chromatograms of sample solvent  
Lab 3**



**Figure 31: Chromatograms of reference standard CGA 344605  
Lab 3**

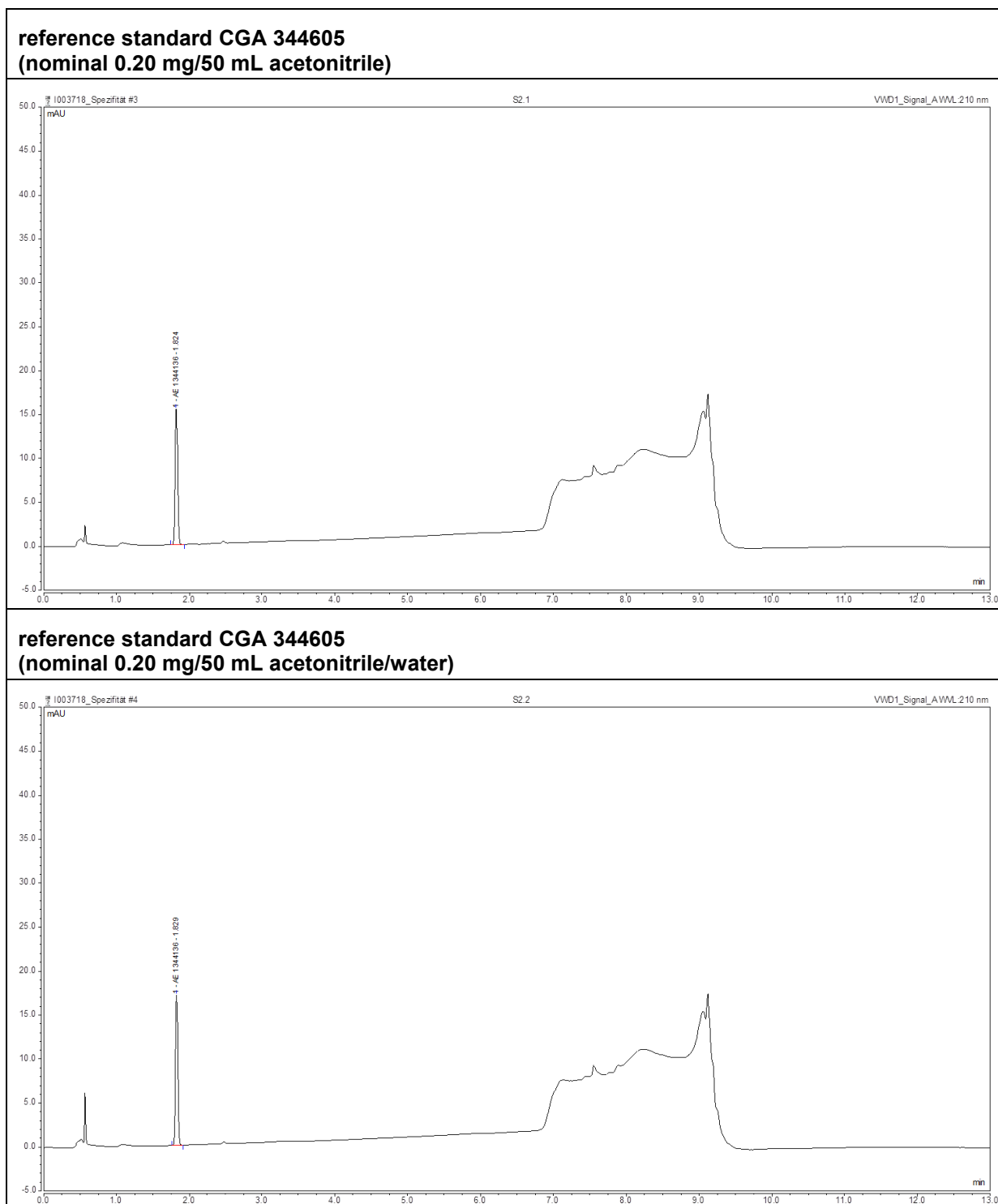
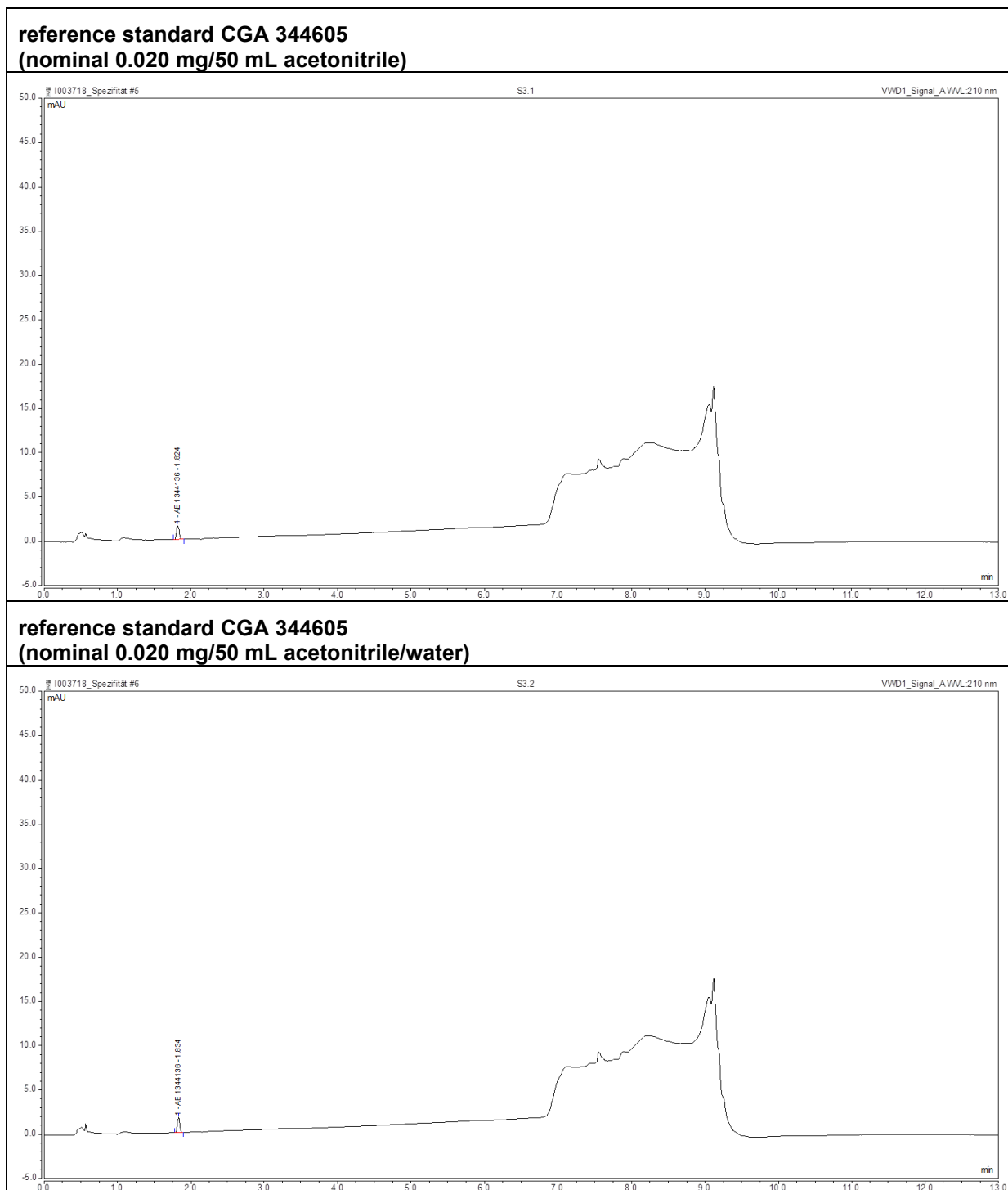
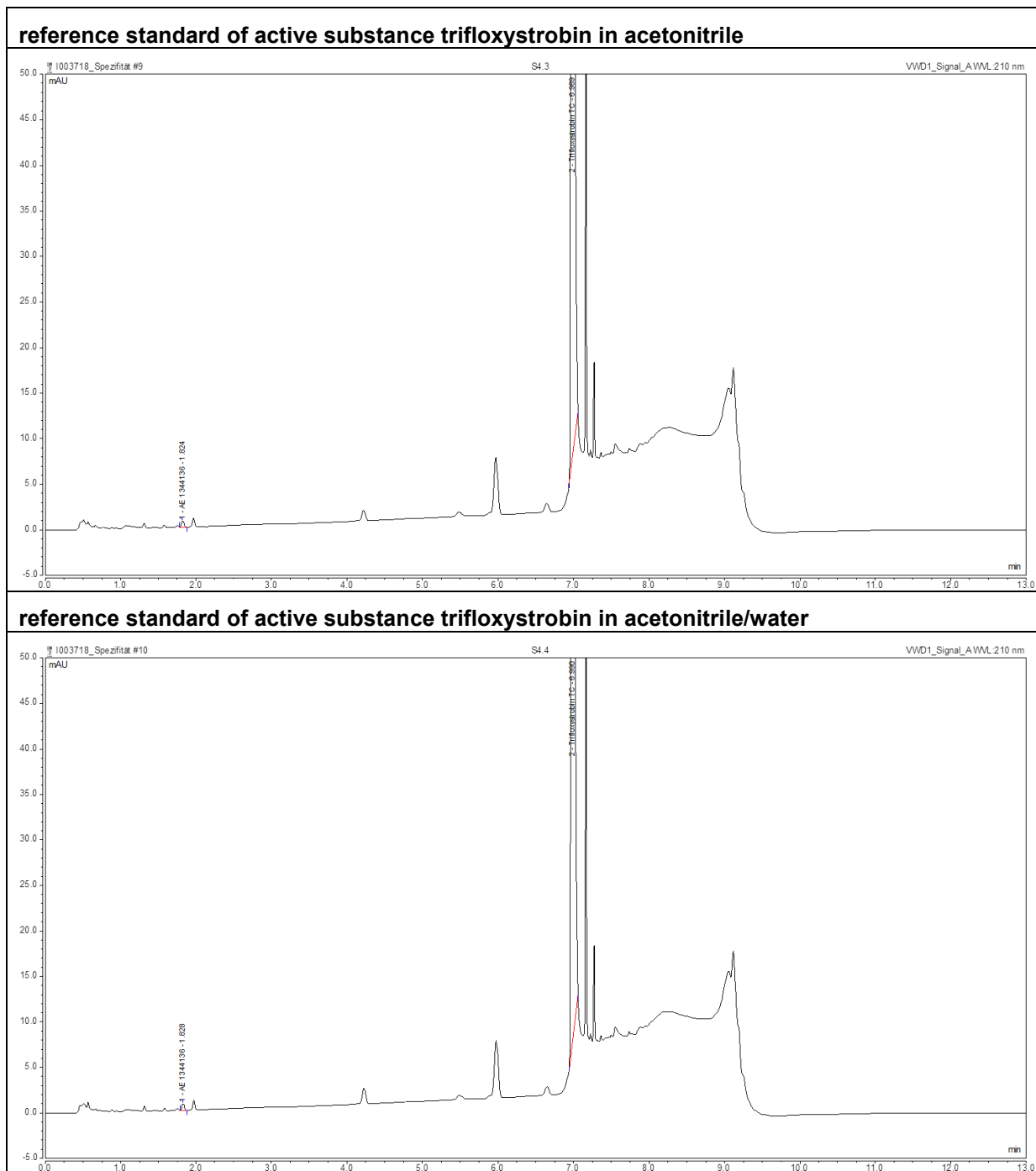




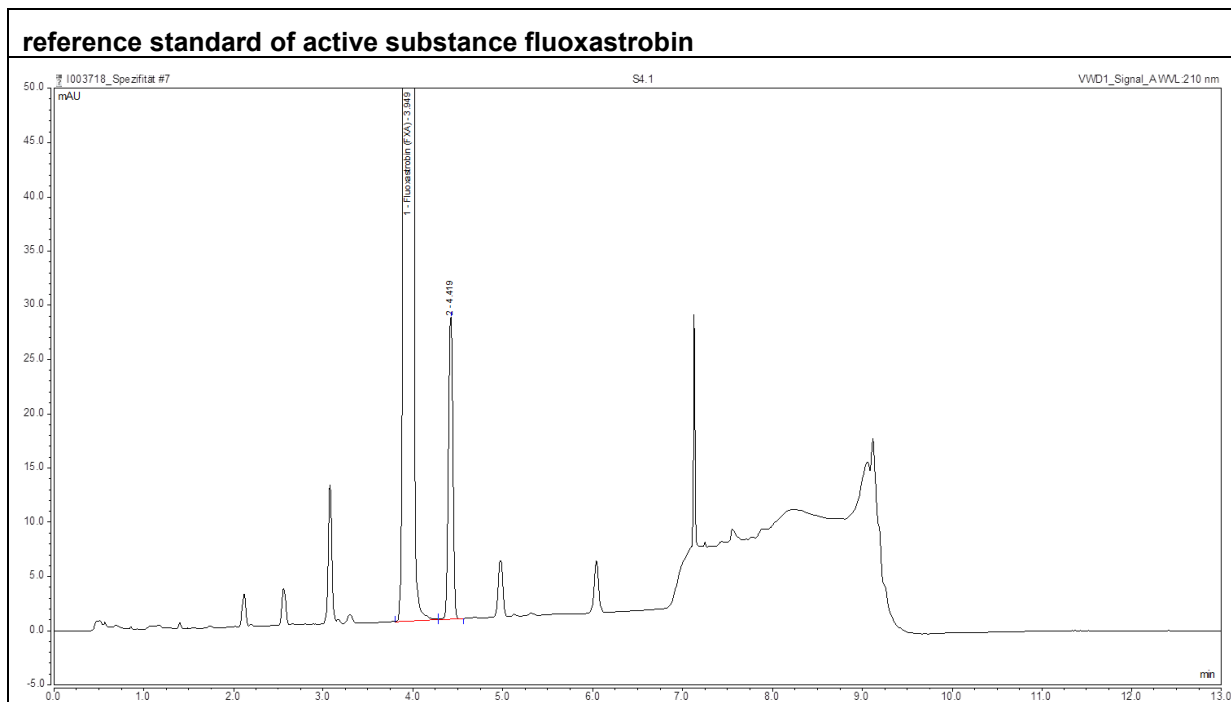
Figure 31. Continuation



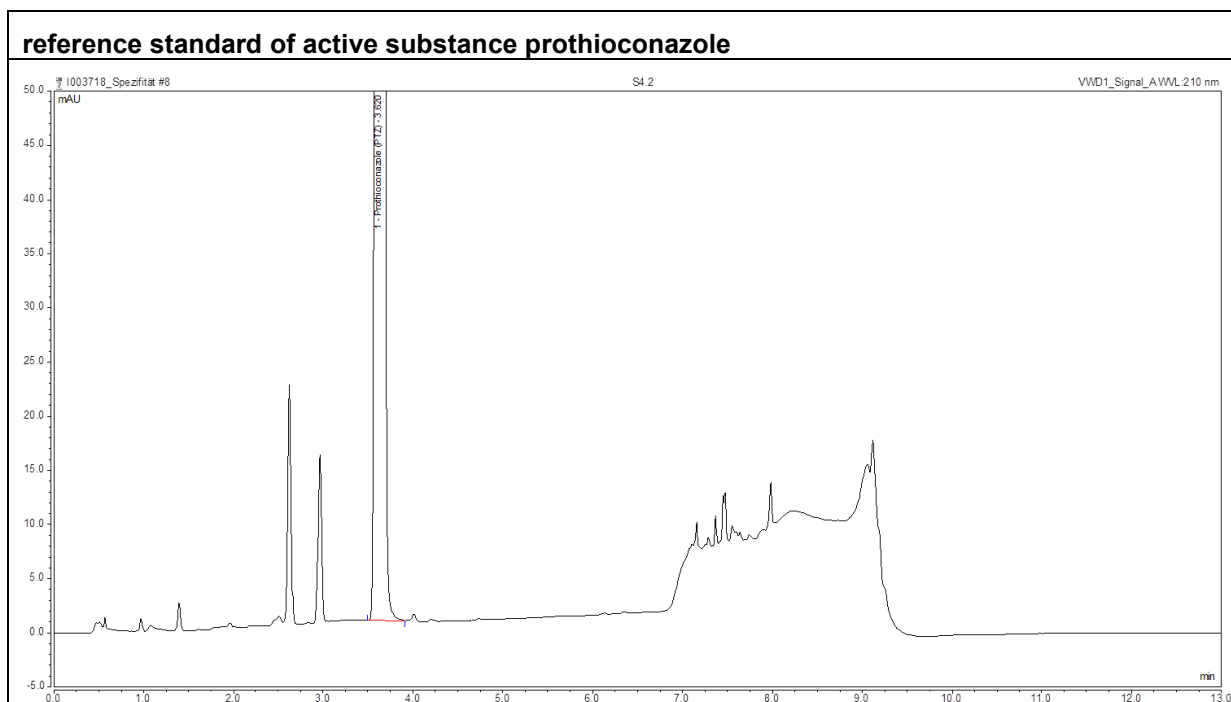
**Figure 32: Chromatograms of active substance trifloxystrobin  
Lab 3**



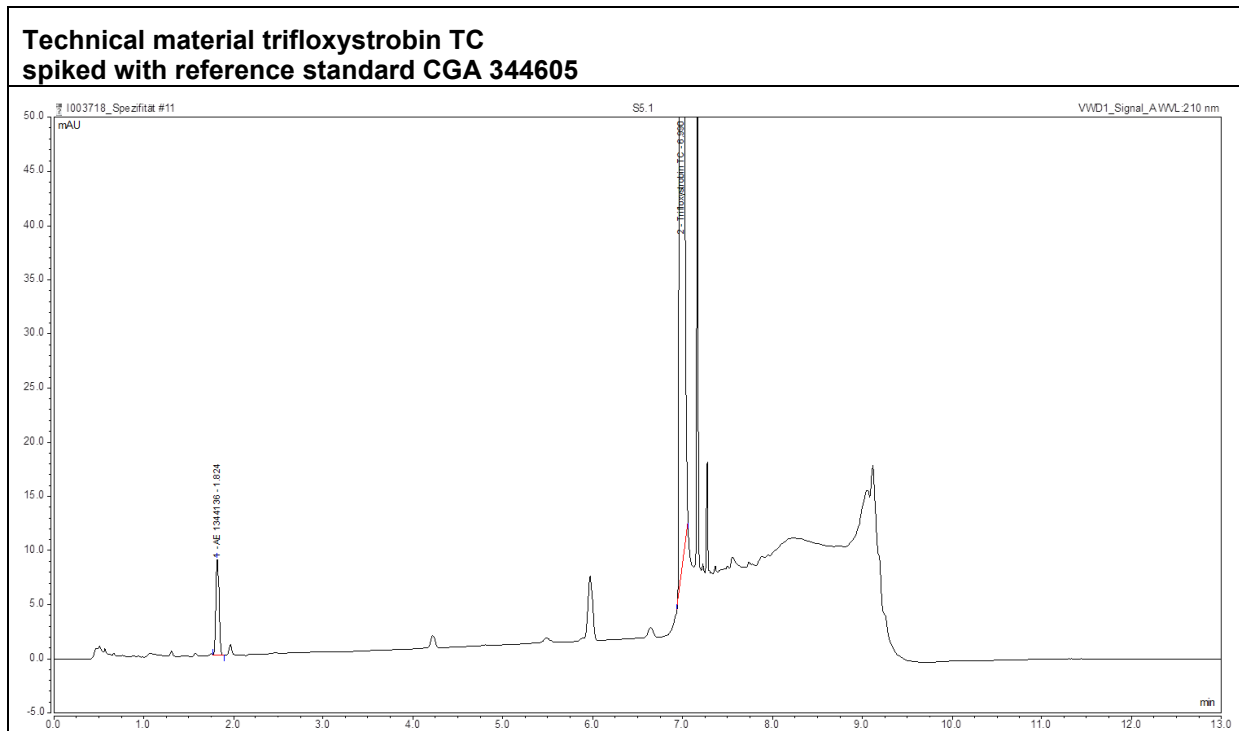
**Figure 33: Chromatograms of active substance fluoxastrobin.  
Lab 3**



**Figure 34: Chromatograms of active substance prothioconazole  
Lab 3**



**Figure 35: Chromatograms of trifloxystrobin TC  
Lab 3**



**Figure 36: Chromatograms of trifloxystrobin SC 500 (500 g/L)  
Lab 3**

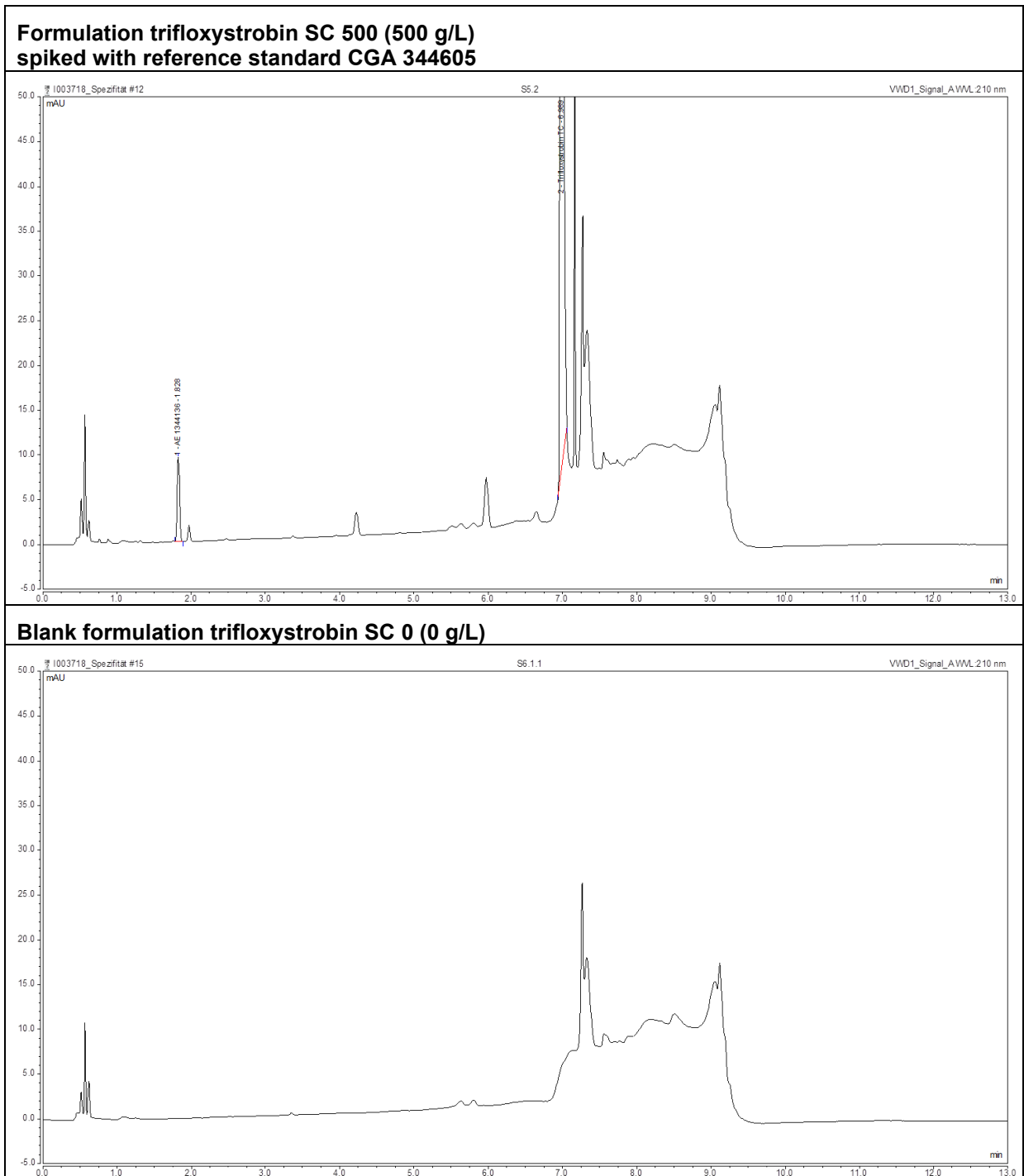
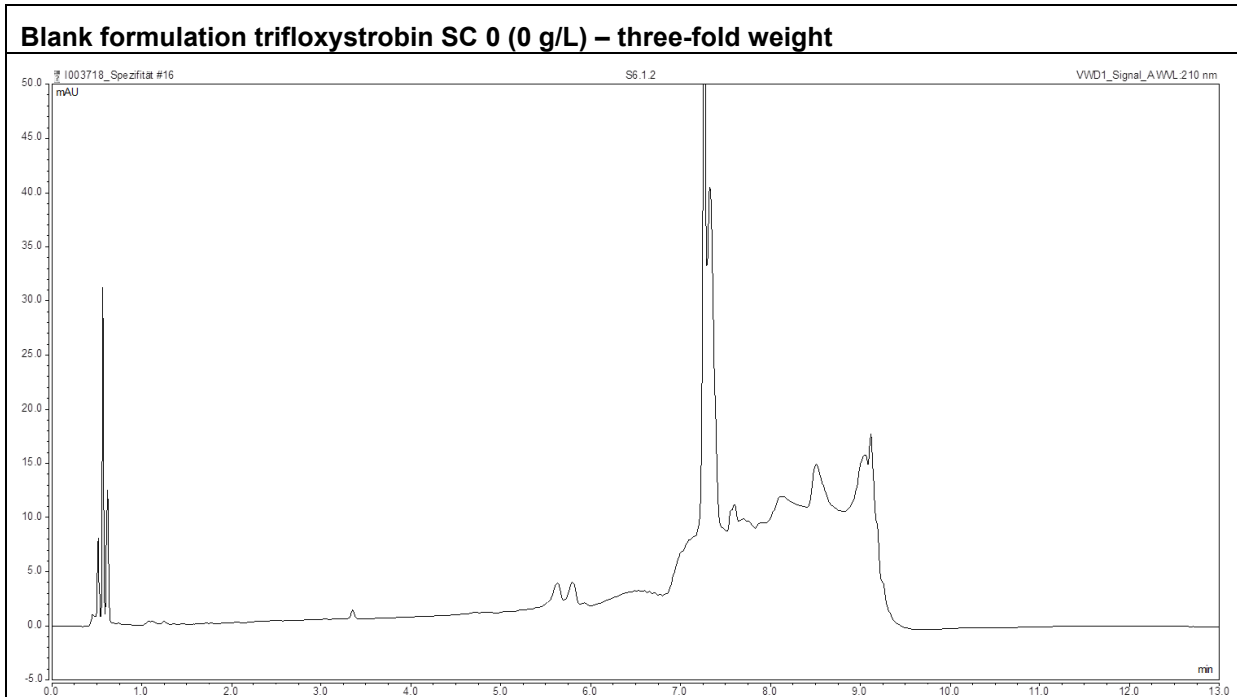


Figure 36. Continuation



**Figure 37: Chromatograms of fluoxastrobin + prothioconazole + trifloxystrobin EC 300 (75+150+75 g/L) Lab 3**

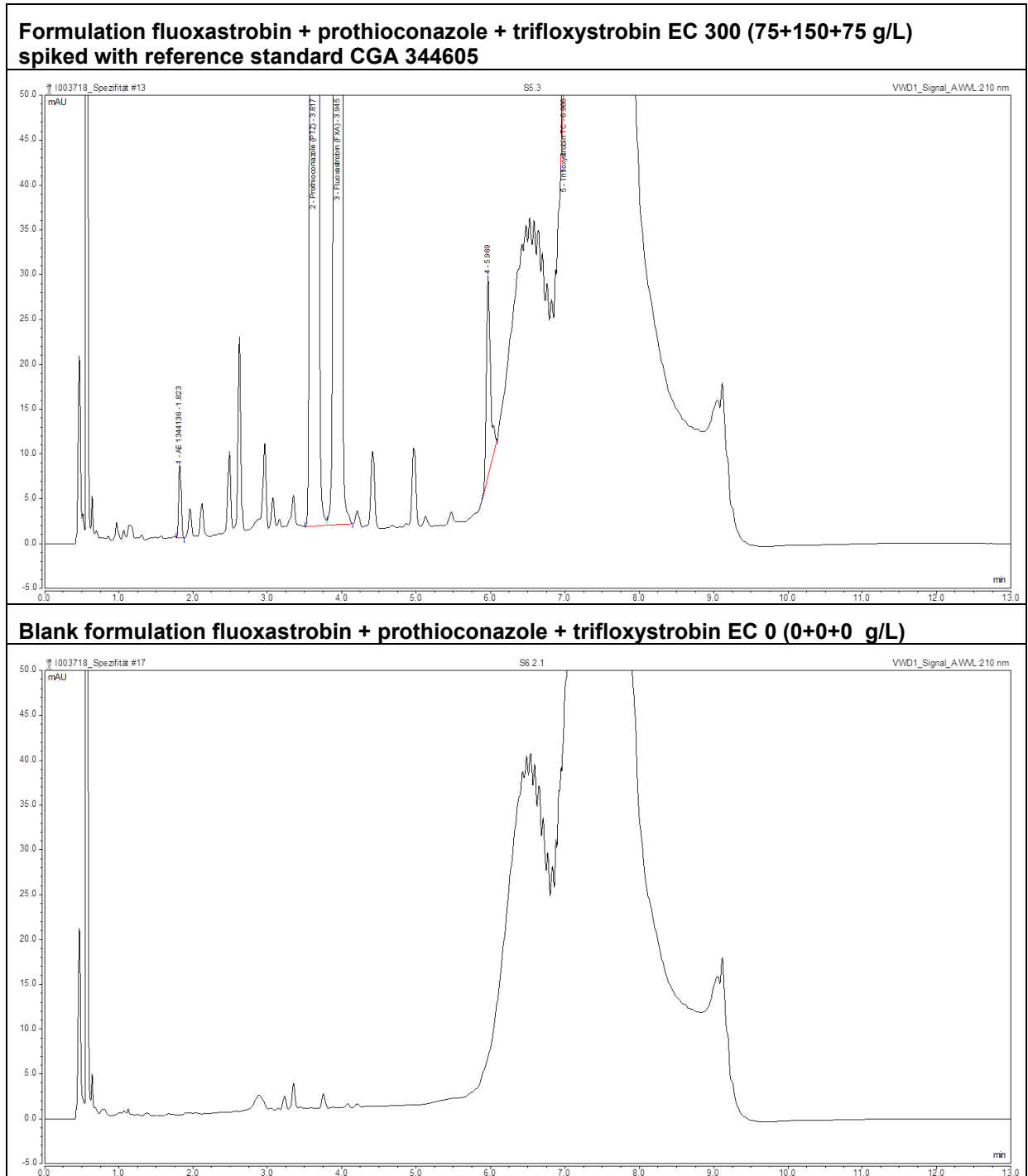
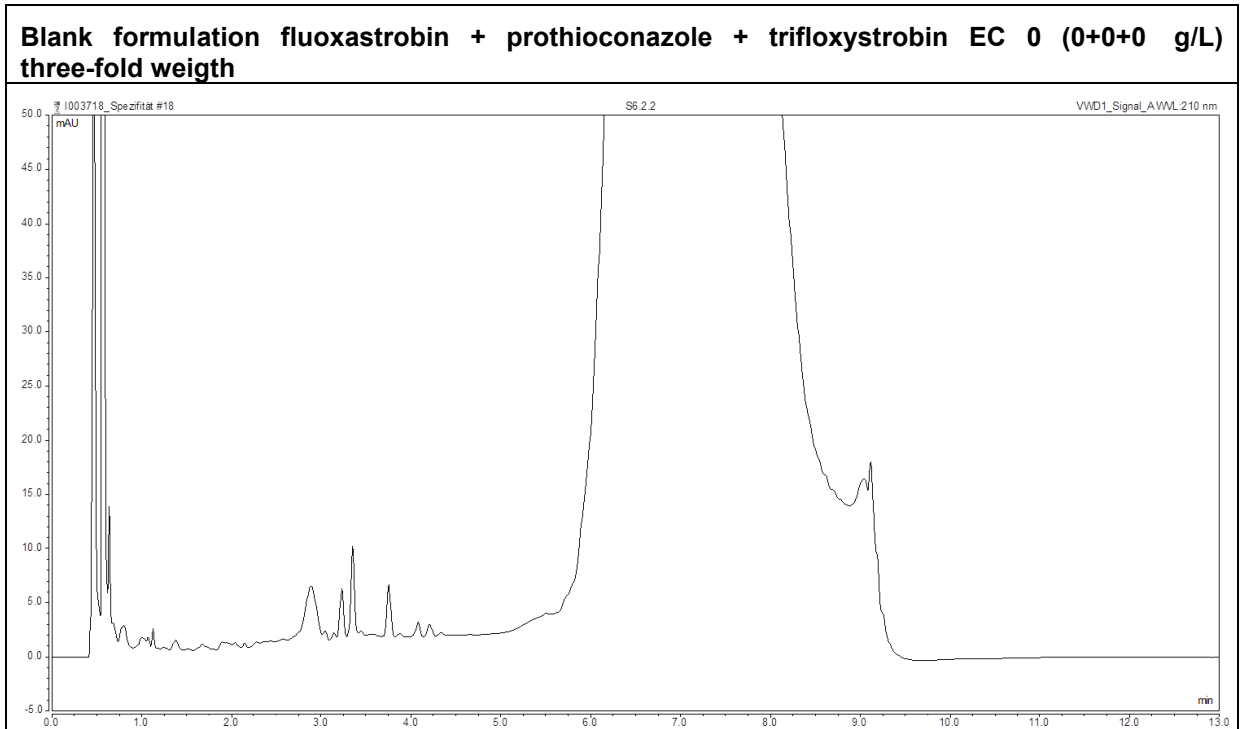


Figure 37. Continuation





**Figure 38: Chromatogram of trifloxystrobin WG 50 (50 g/kg)  
Lab 3**

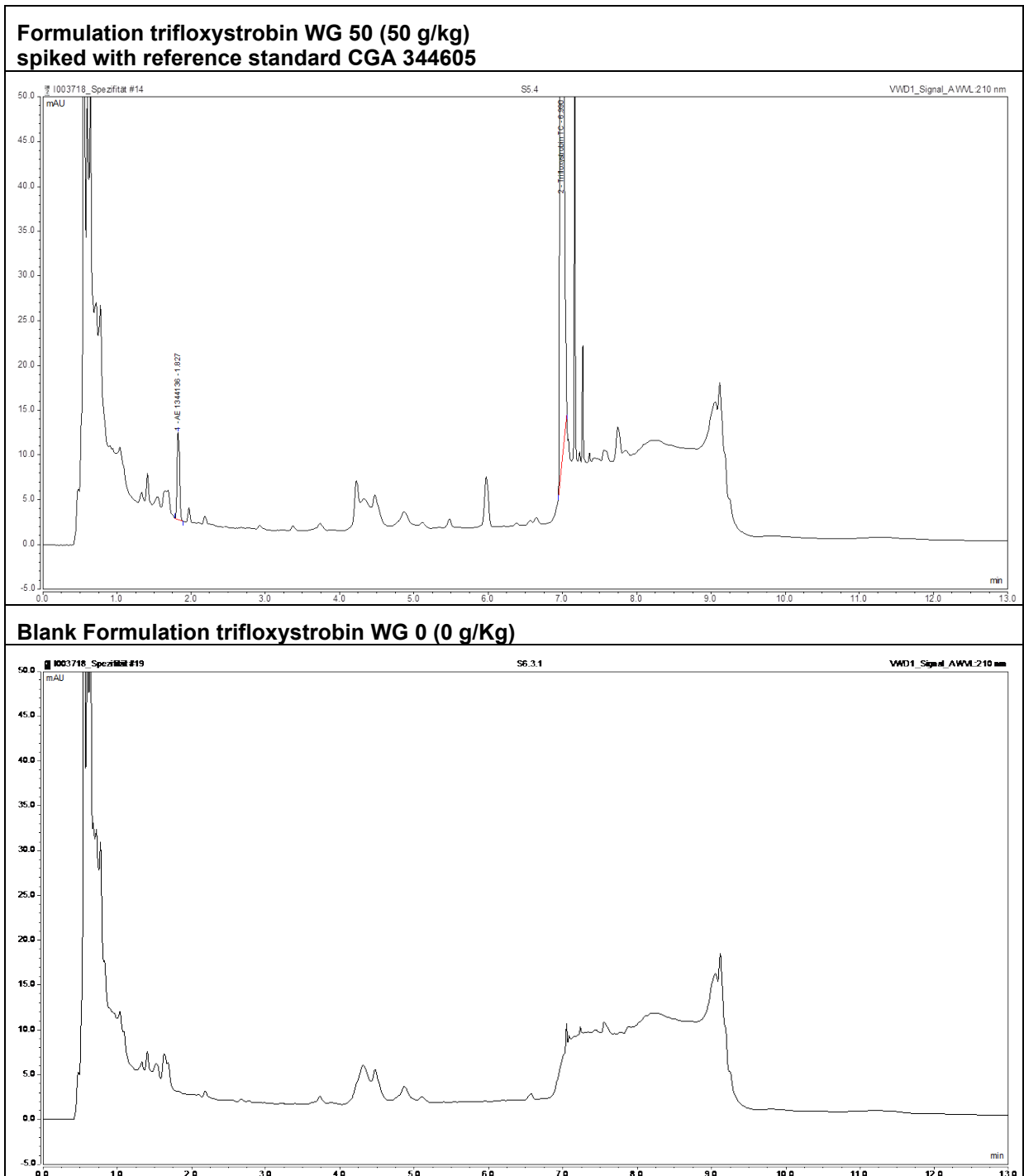
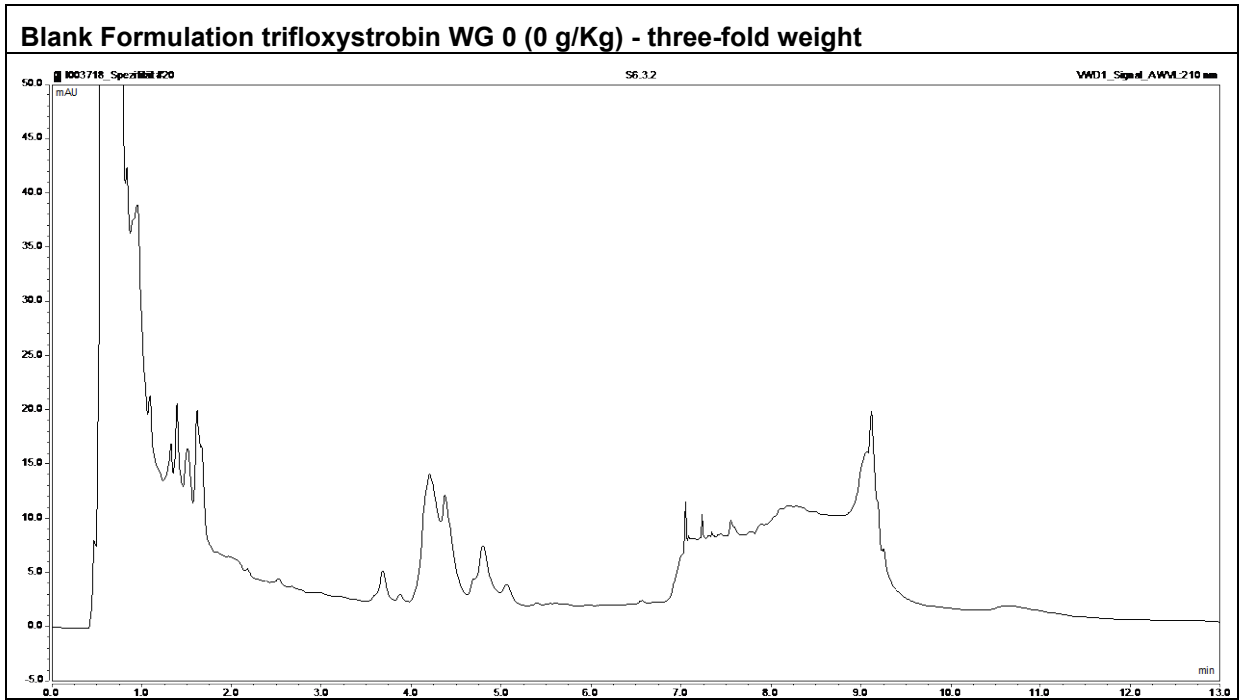
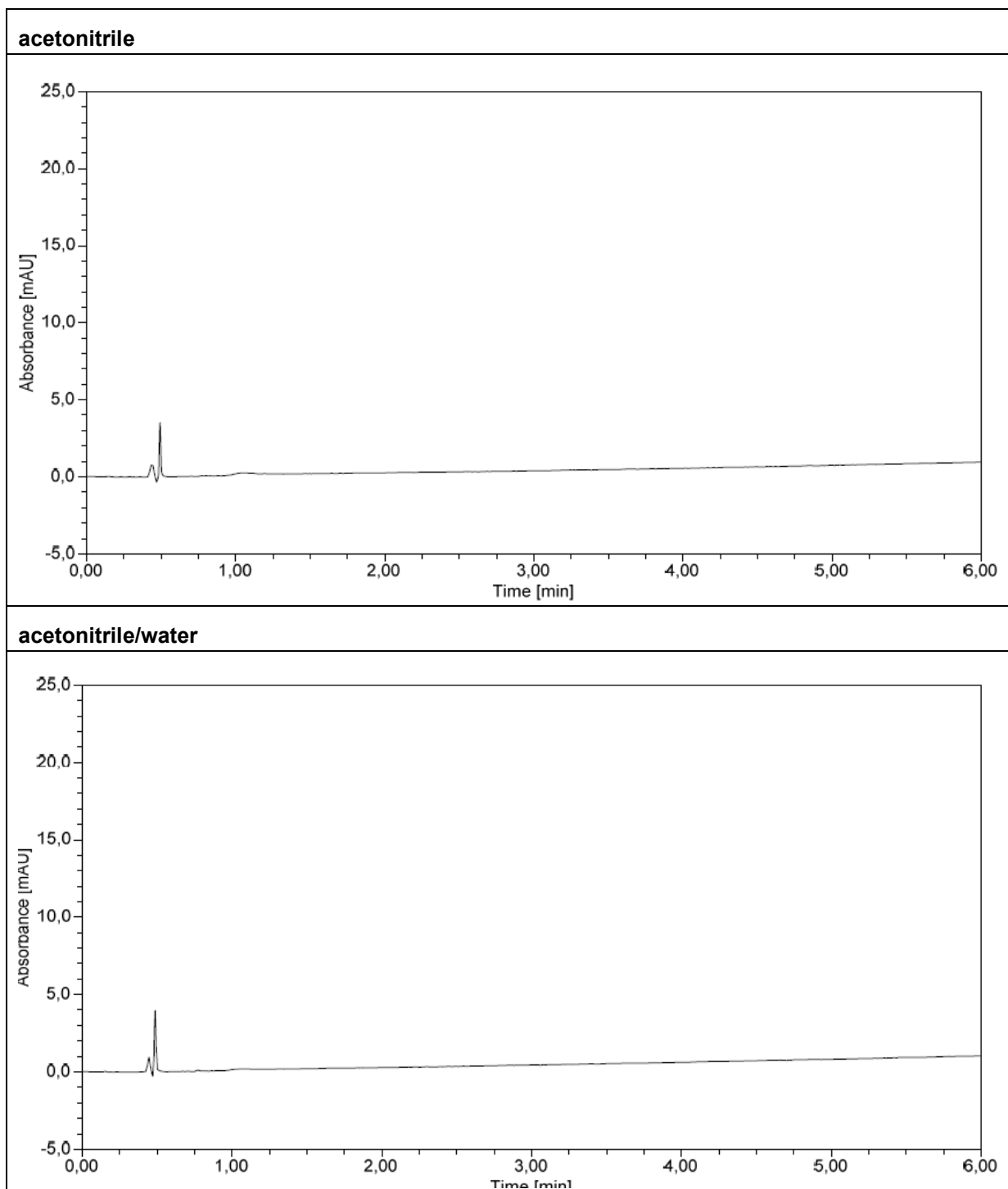


Figure 38. Continuation



**Figure 39: Chromatograms of sample solvent  
Lab 4**



**Figure 40: Chromatograms of reference standard CGA 344605  
Lab 4**

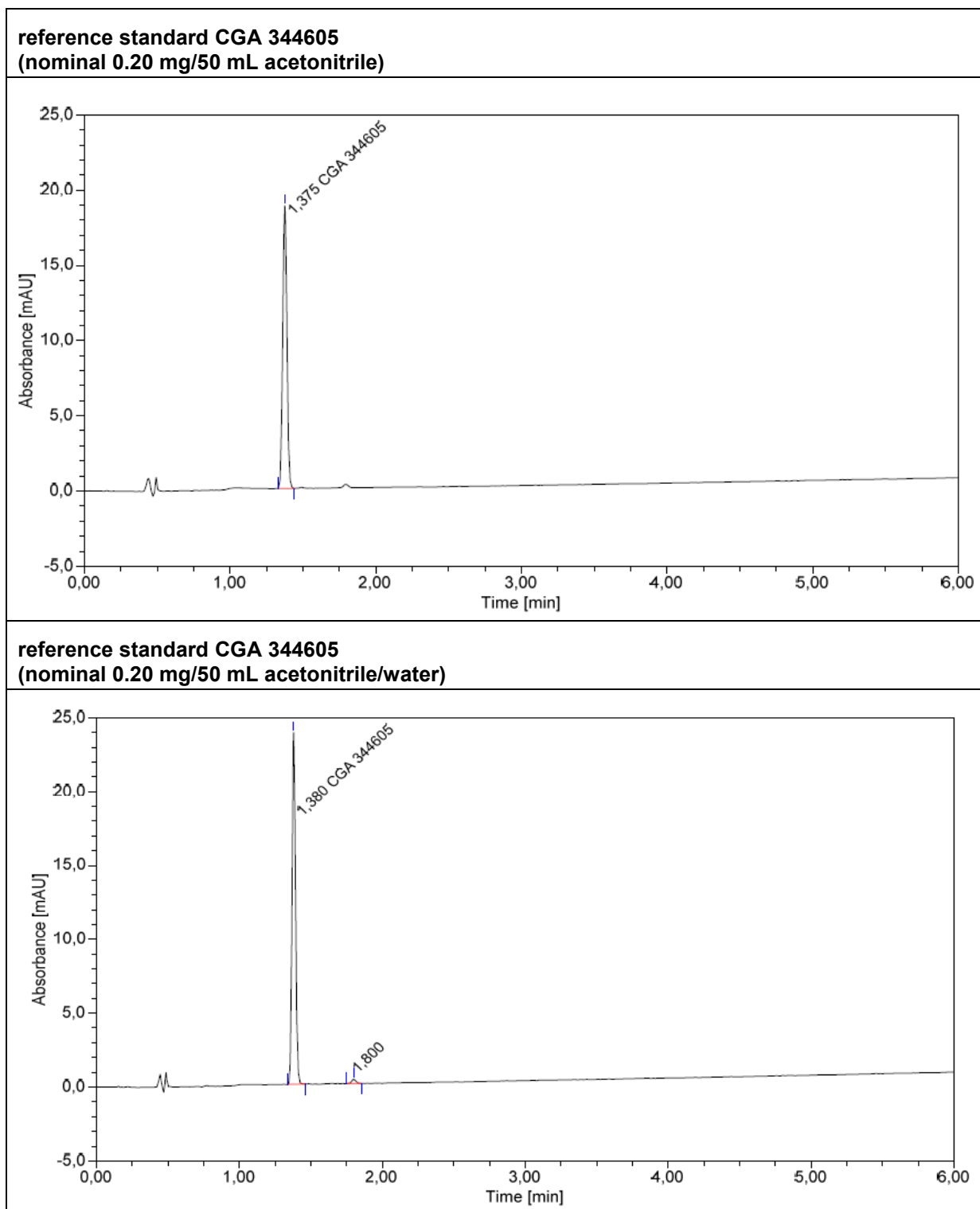
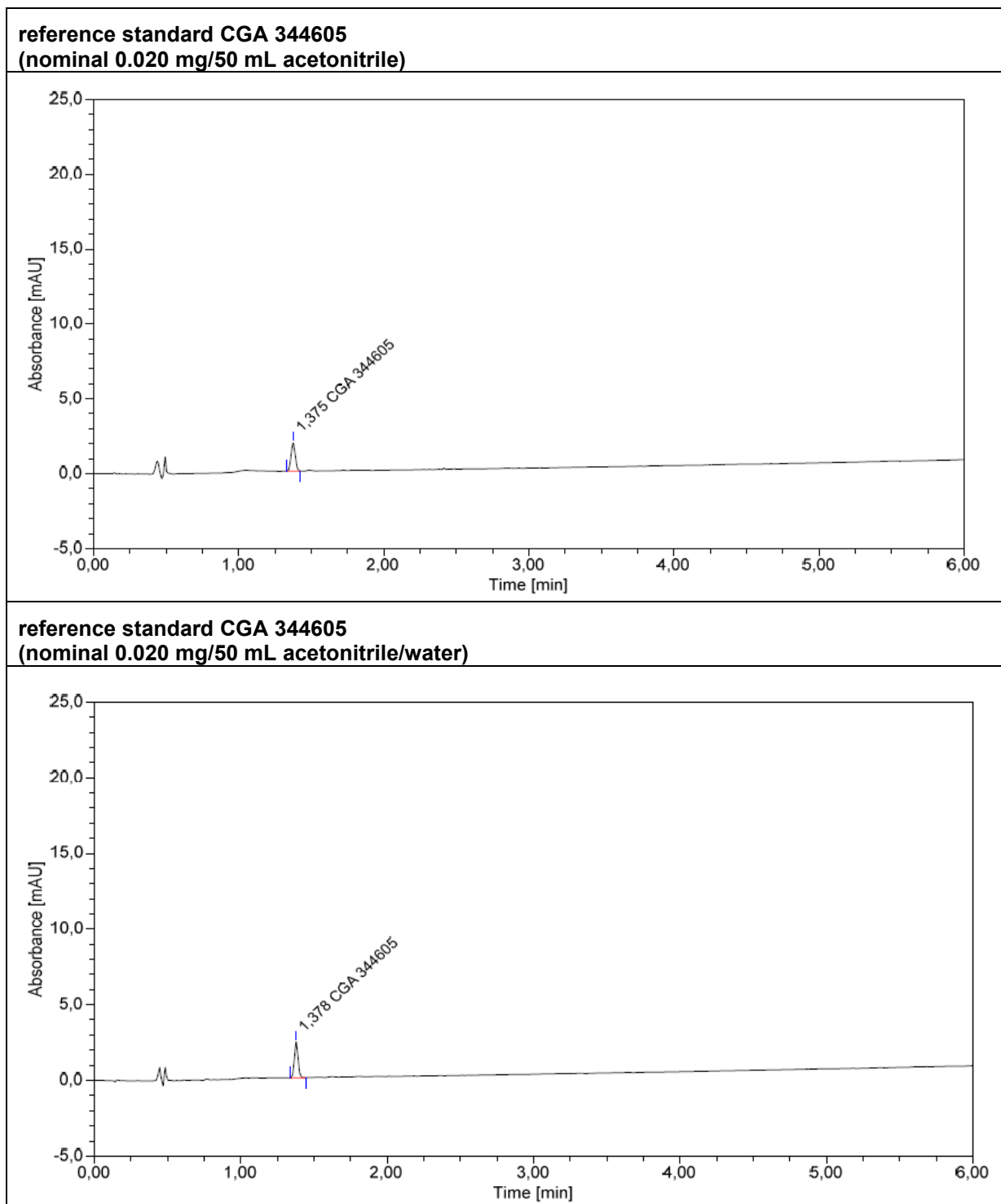
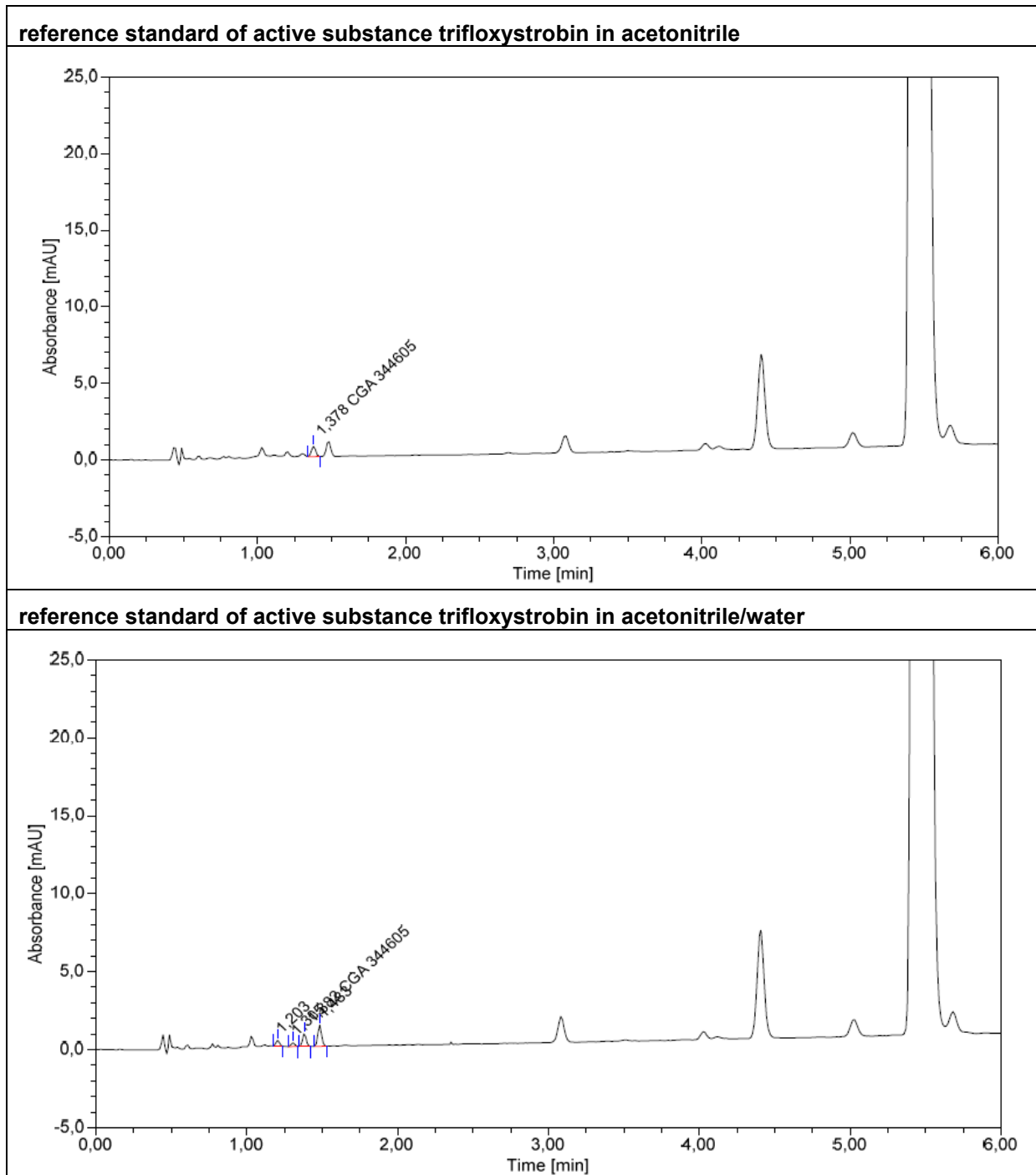


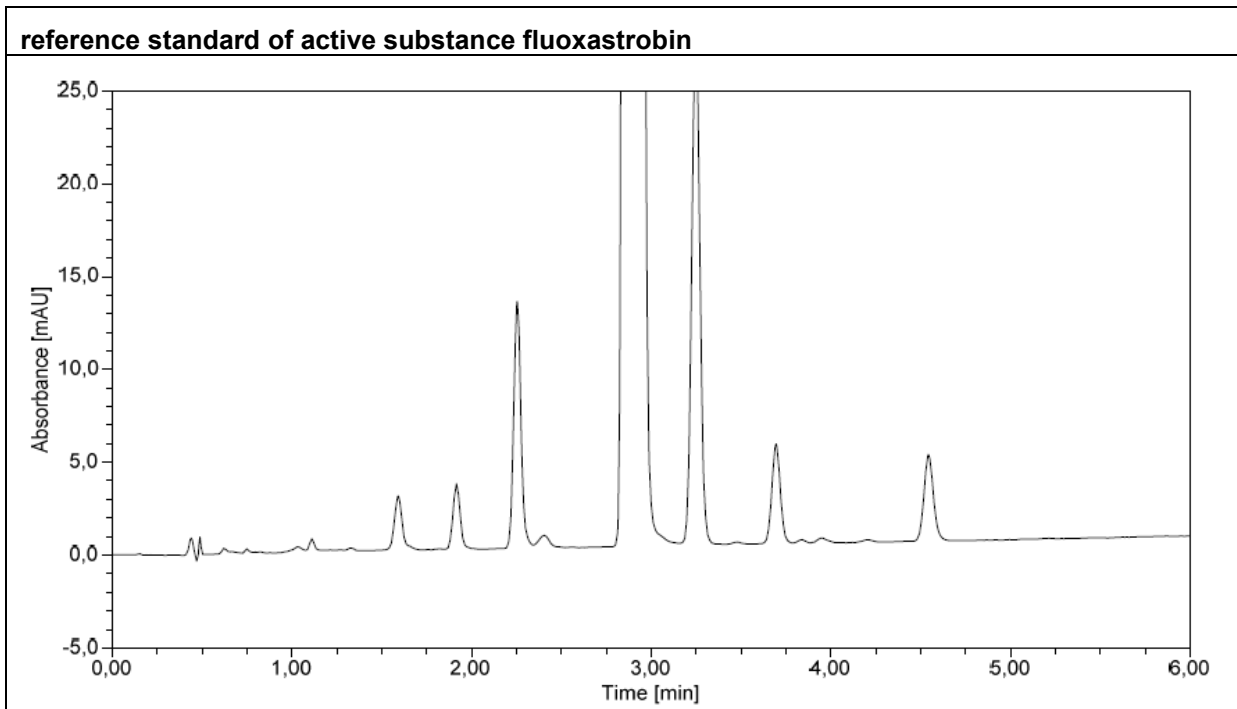
Figure 40. Continuation



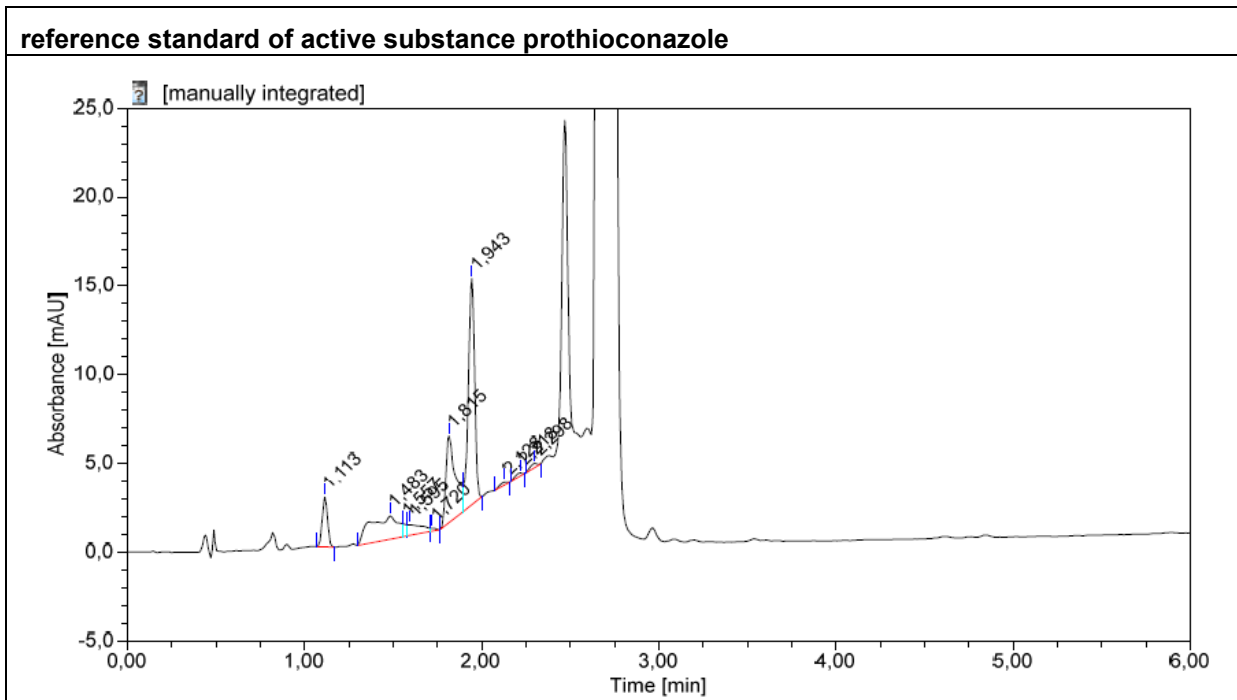
**Figure 41: Chromatograms of active substance trifloxystrobin  
Lab 4**



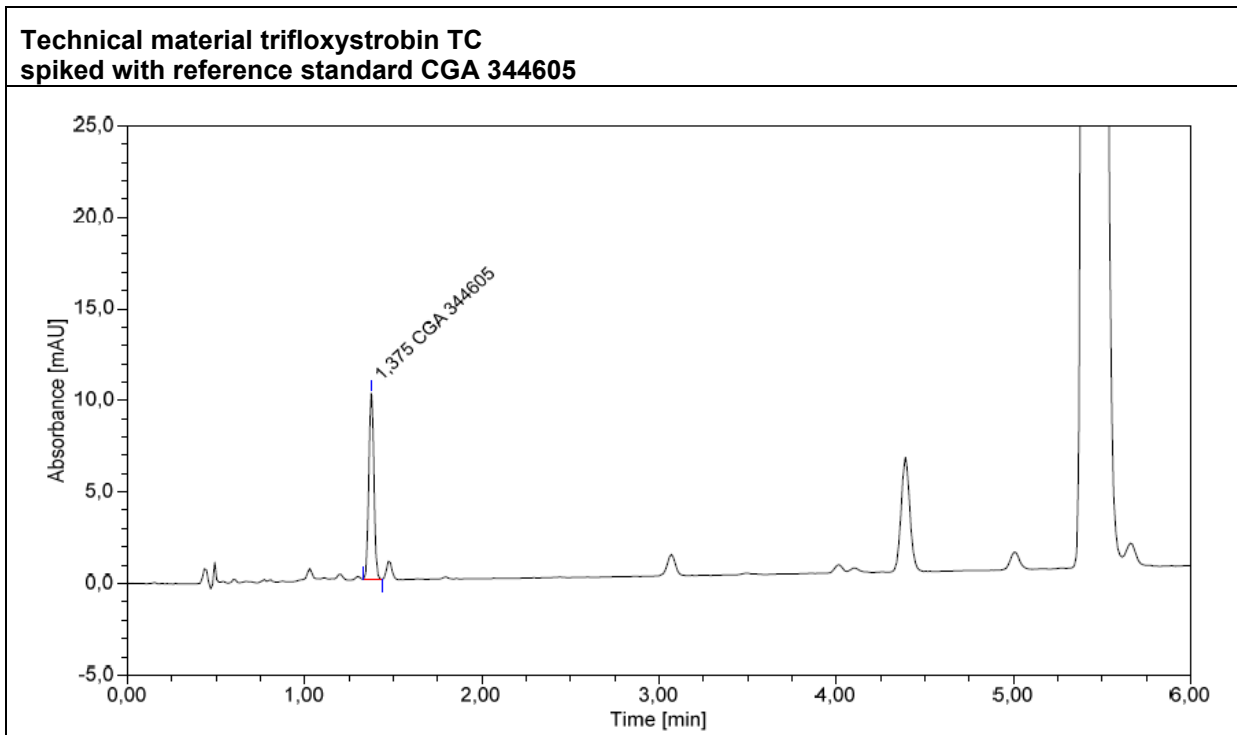
**Figure 42: Chromatograms of active substance fluoxastrobin.  
Lab 4**



**Figure 43: Chromatograms of active substance prothioconazole  
Lab 4**



**Figure 44: Chromatograms of trifloxystrobin TC  
Lab 4**





**Figure 45: Chromatograms of trifloxystrobin SC 500 (500 g/L)  
Lab 4**

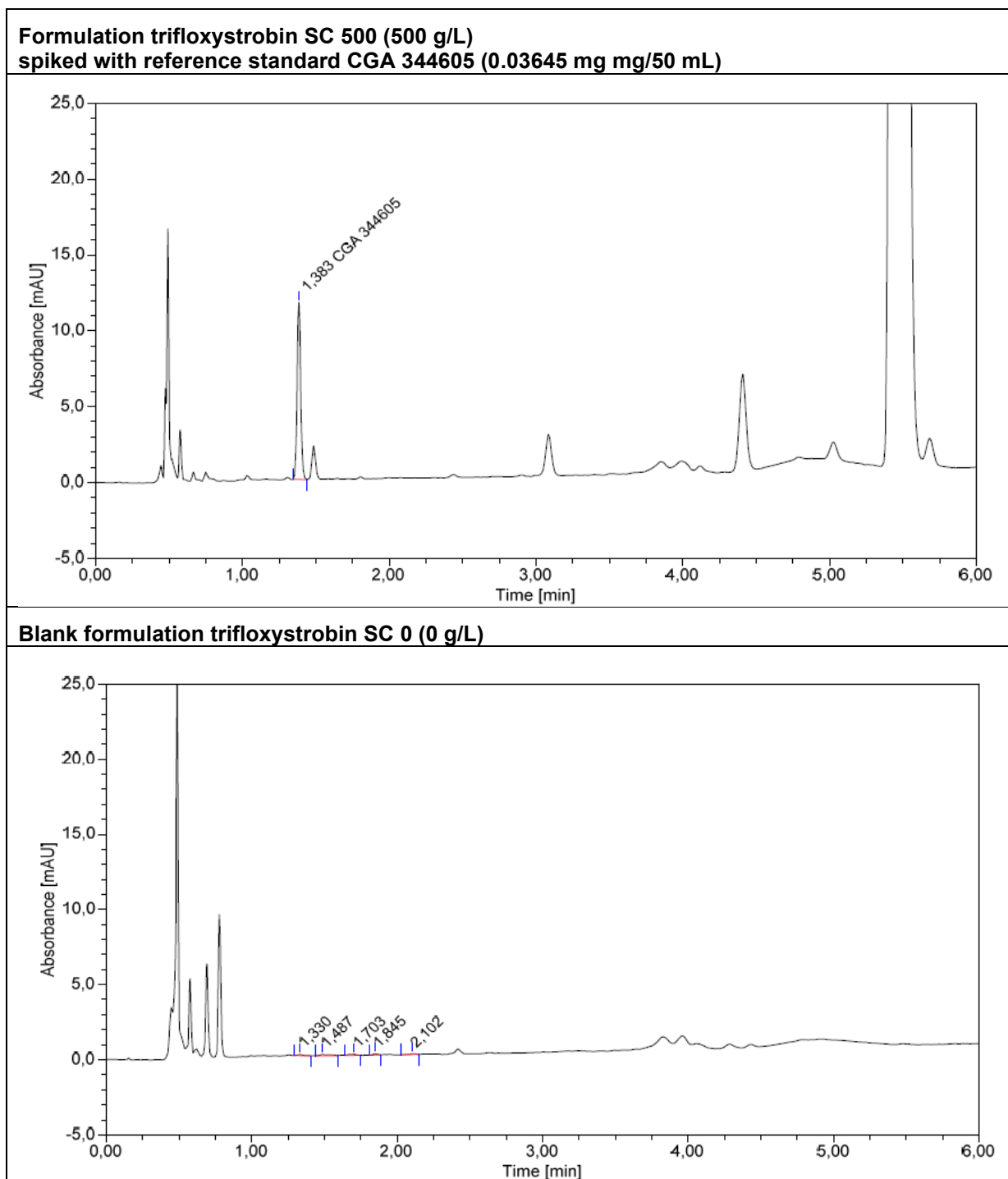
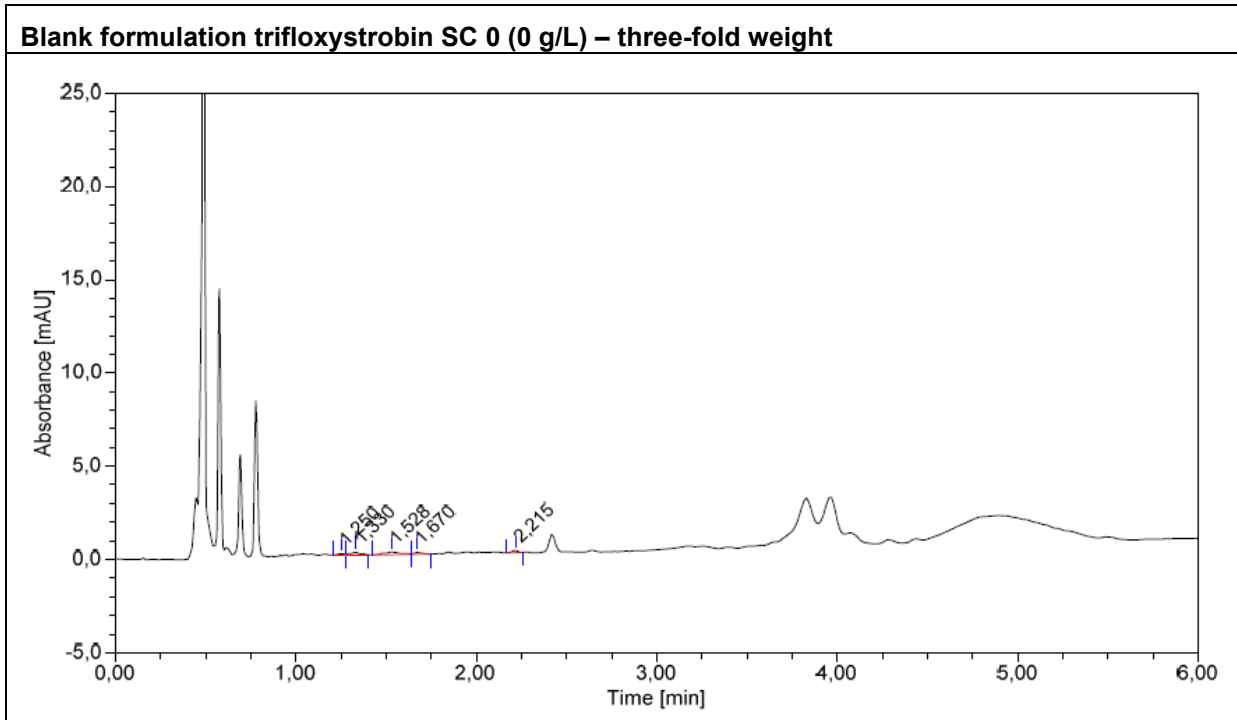
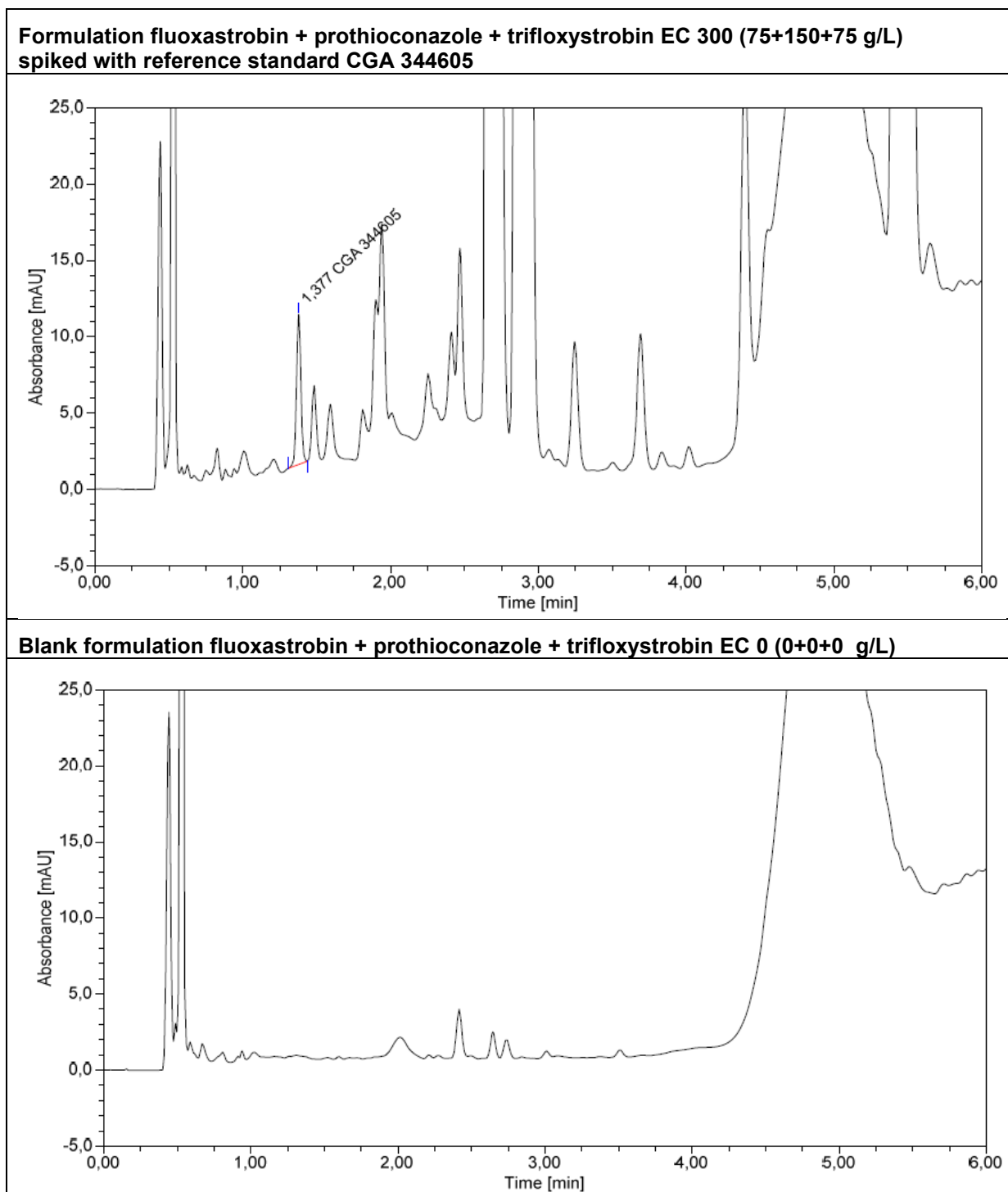
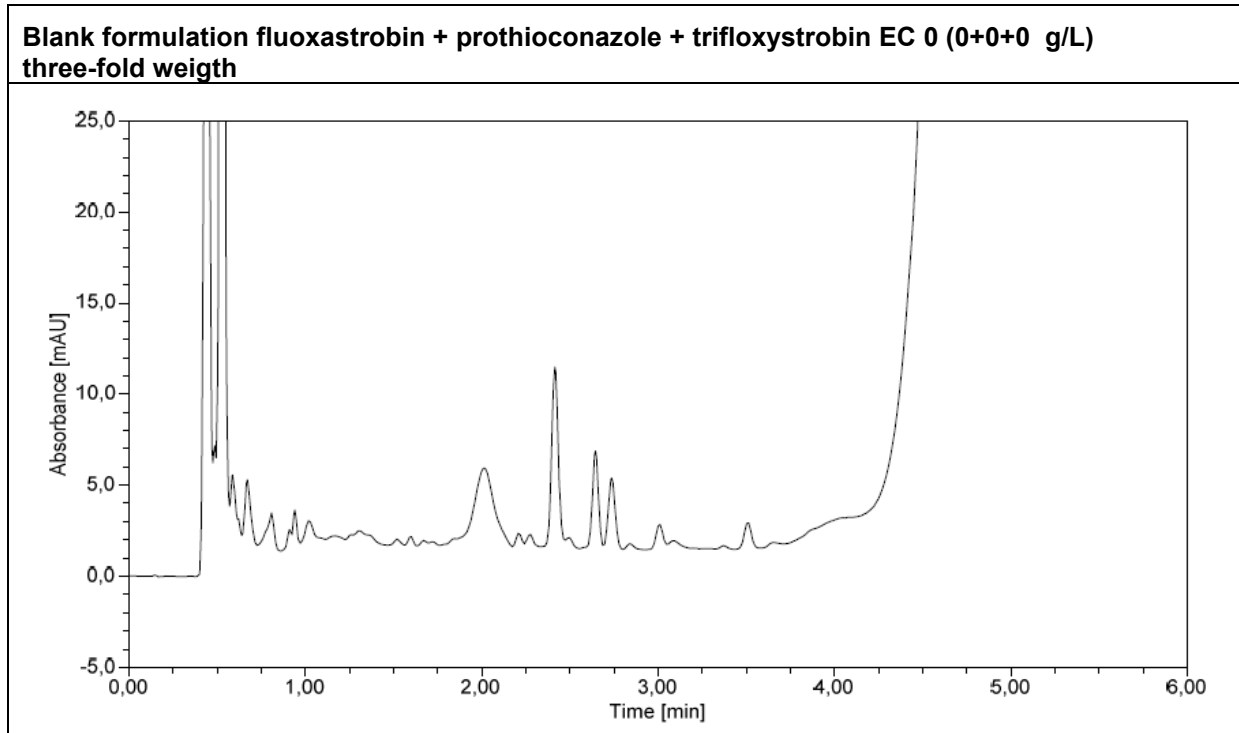


Figure 45. Continuation



**Figure 46: Chromatograms of fluoxastrobin + prothioconazole + trifloxystrobin EC 300 (75+150+75 g/L) Lab 4**



**Figure 46. Continuation**

**Figure 47: Chromatogram of trifloxystrobin WG 50 (50 g/kg)  
Lab 4**

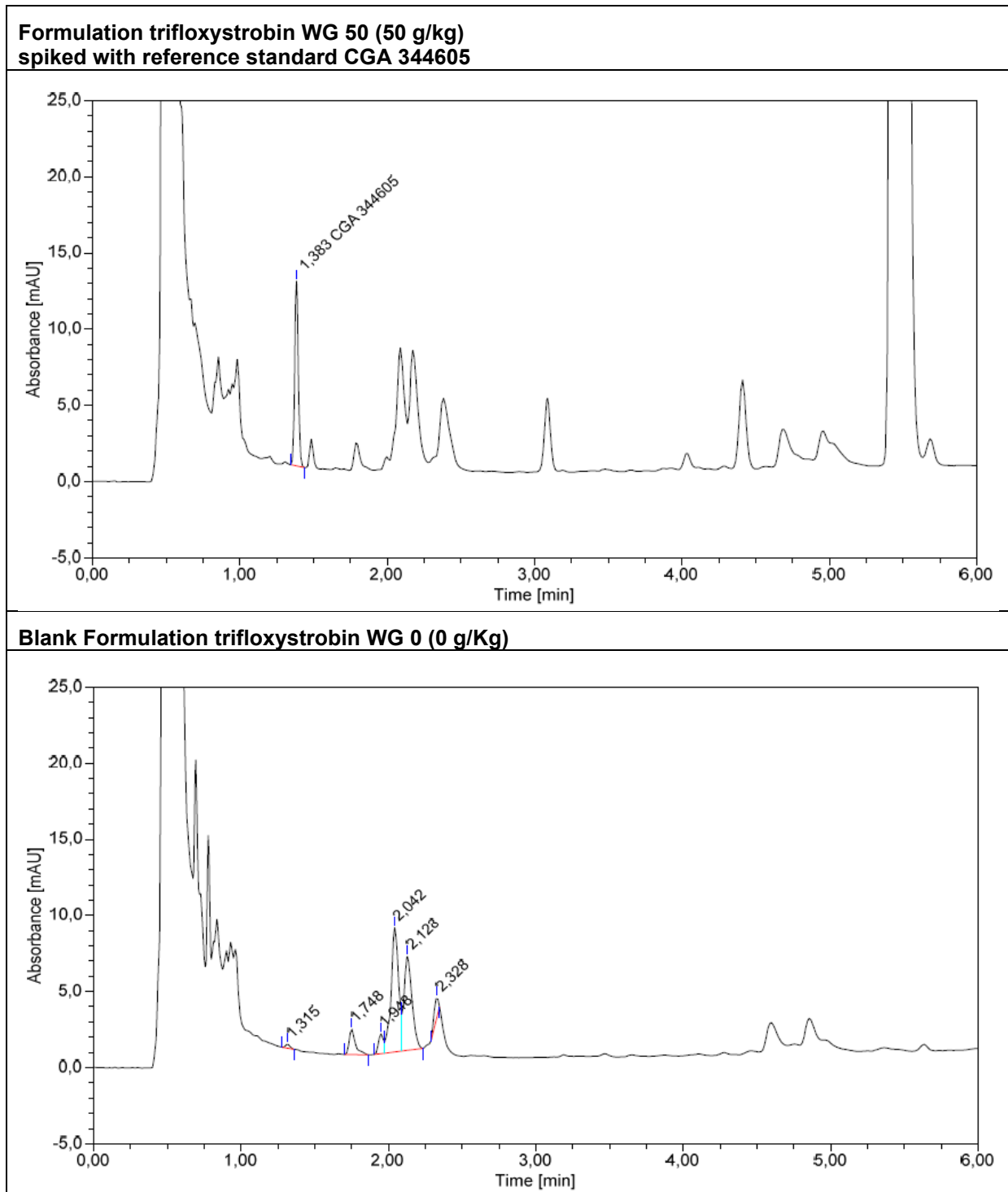
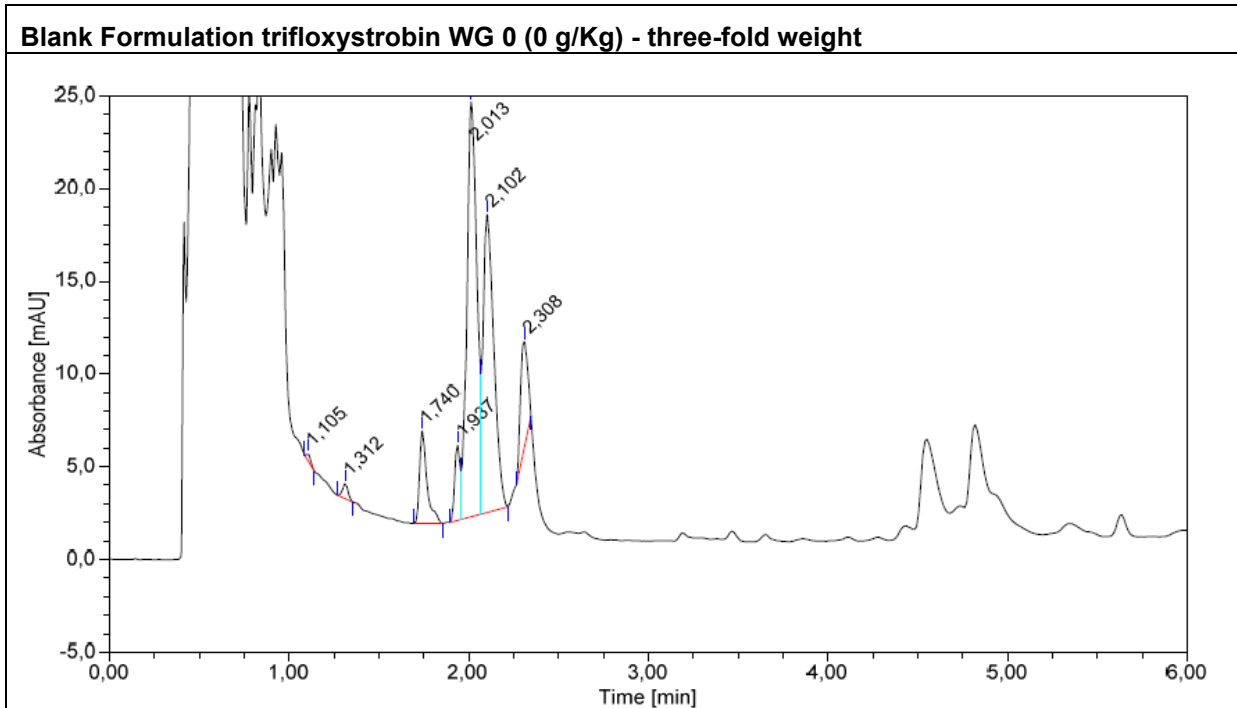
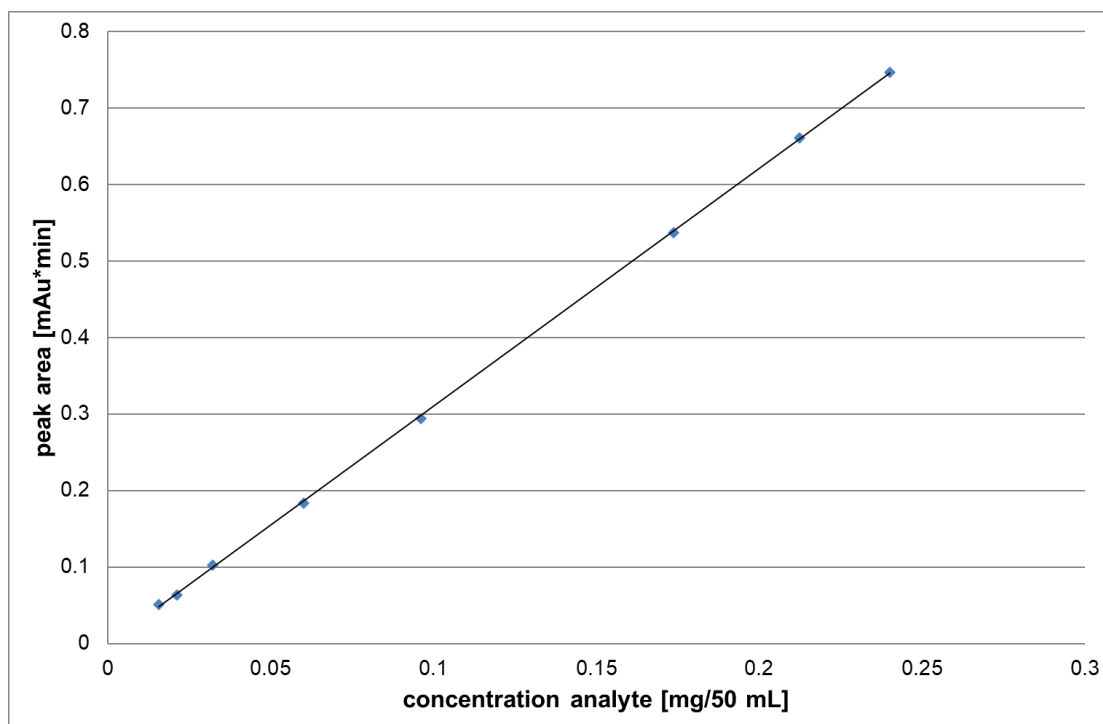


Figure 47. Continuation



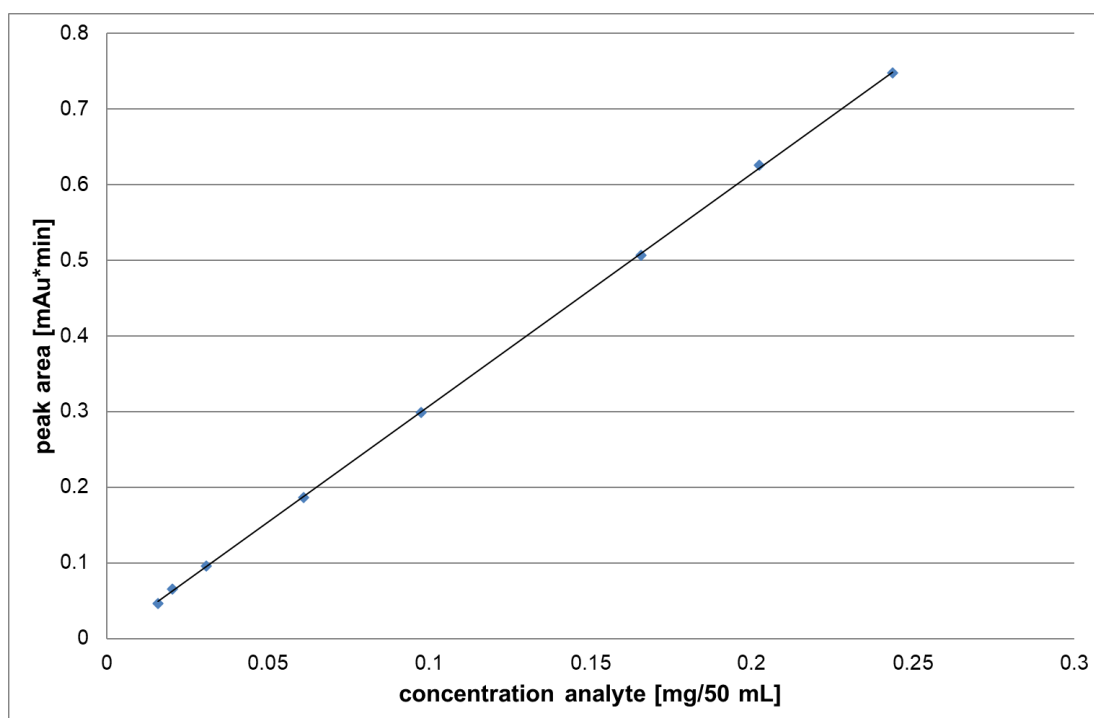
## 6.4 Calibration function (Linearity)

Figure 48: Calibration function of CGA 344605 in acetonitrile  
Developing Lab



Number of values	n = 8
Regression equation	$y = a + b x$ (1 <sup>st</sup> order) $y = -0.000678 + 3.105749 x$
Correlation coefficient	= <b>0.99995</b>

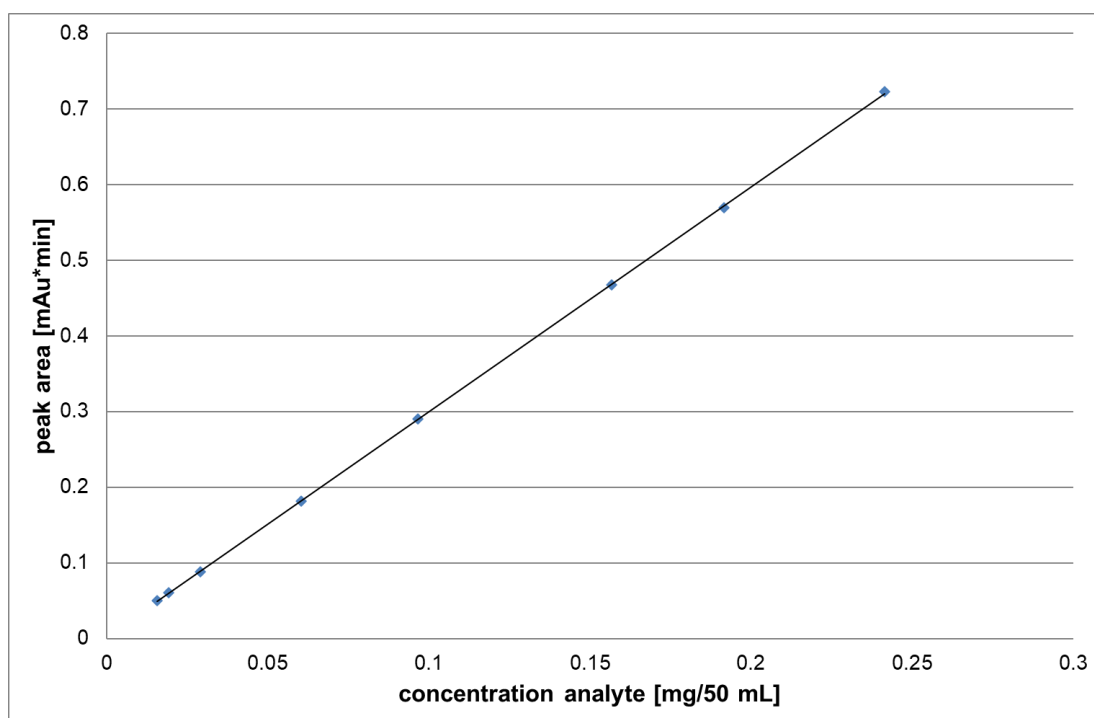
**Figure 49: Calibration function of CGA 344605 in acetonitrile/water  
Developing Lab**



Number of values	n = 8
Regression equation	$y = a + b x$ (1 <sup>st</sup> order) $y = 0.000502 + 3.070411 x$
<b>Correlation coefficient</b>	<b>= 0.99996</b>

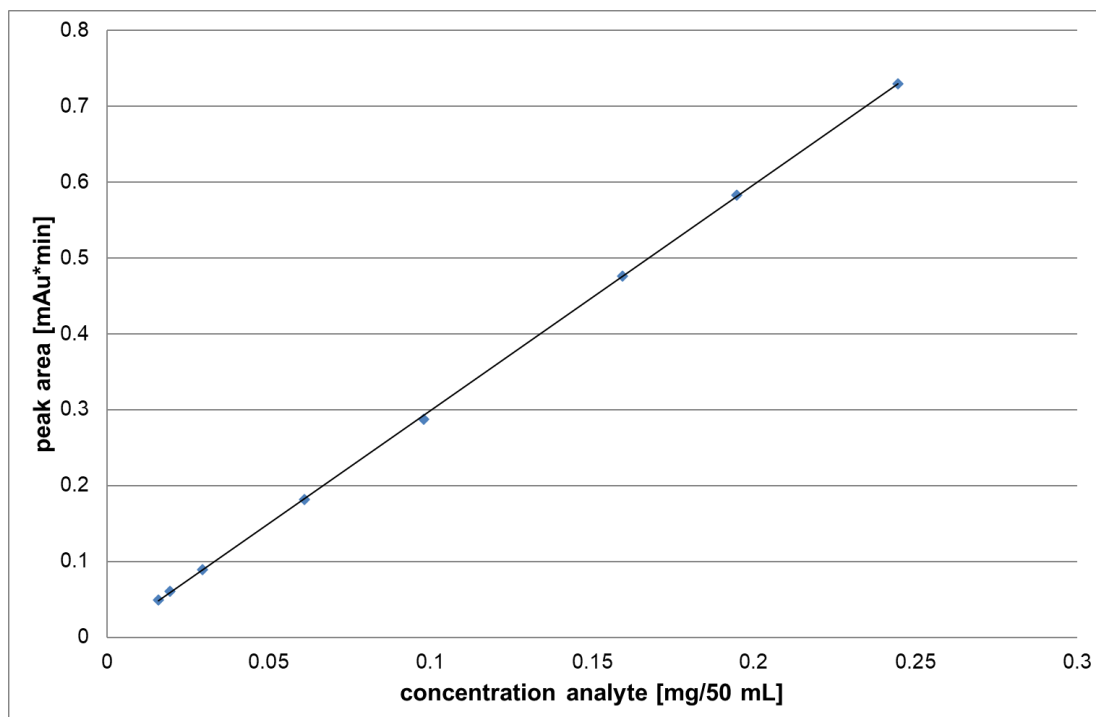


**Figure 50: Calibration function of CGA 344605 in acetonitrile  
Lab 1**



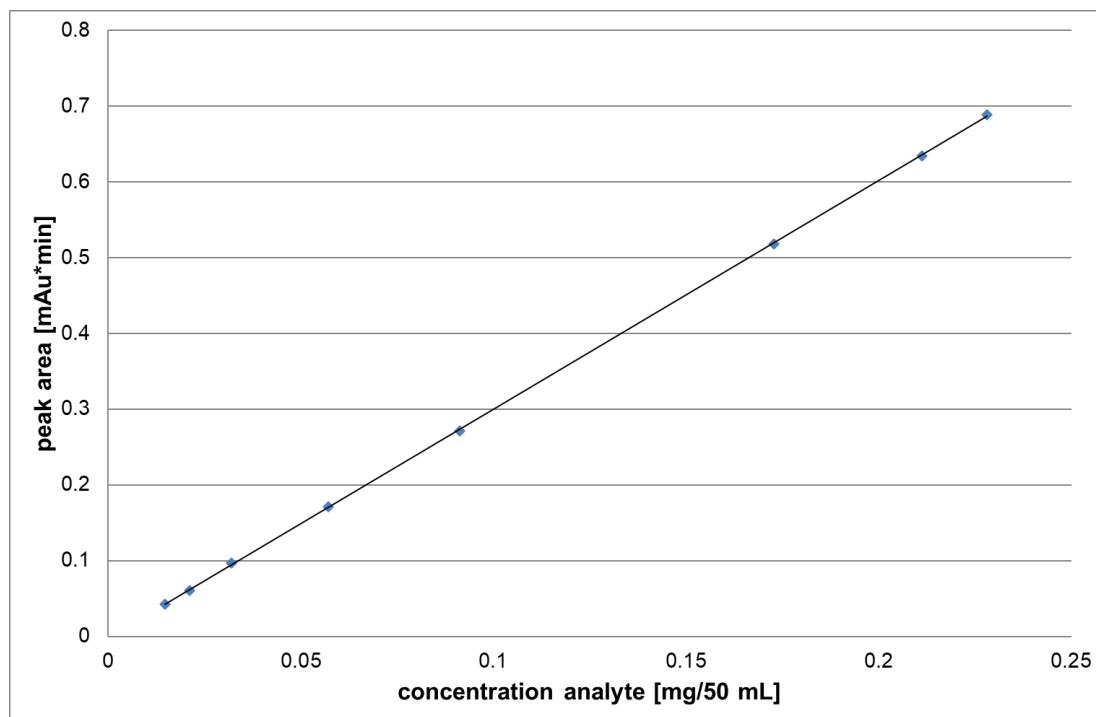
Number of values	n = 8
Regression equation	$y = a + b x$ (1 <sup>st</sup> order) $y = 0.002356 + 2.972195 x$
<b>Correlation coefficient</b>	<b>= 0.99998</b>

**Figure 51: Calibration function of CGA 344605 in acetonitrile/water  
Lab 1**



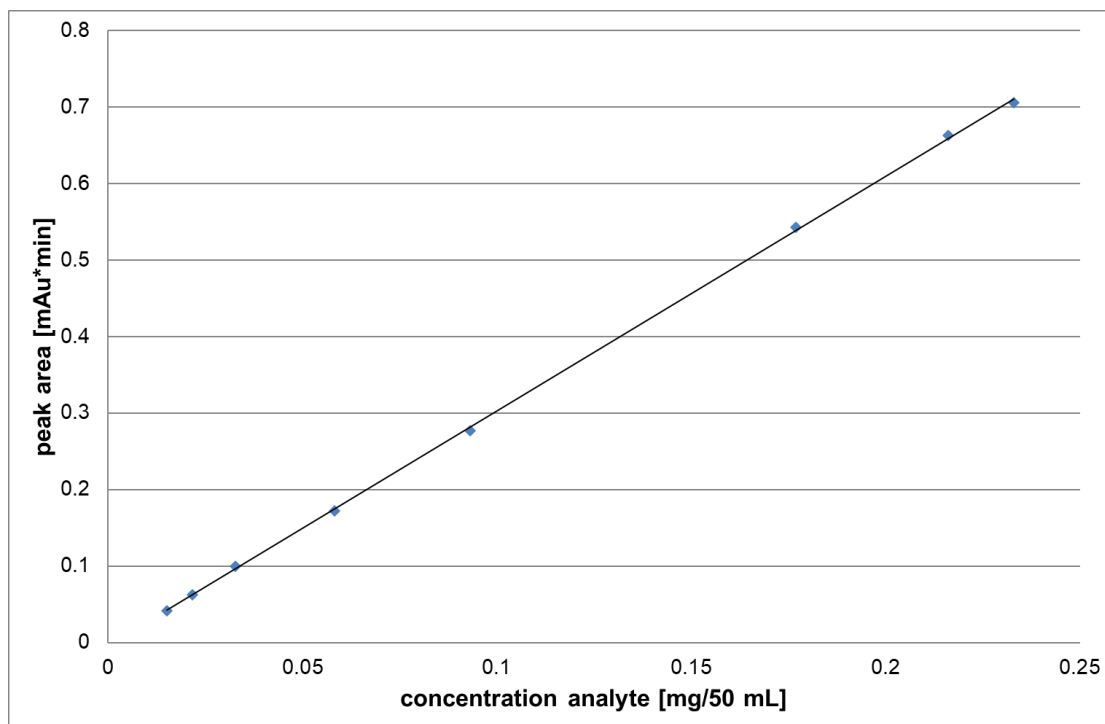
Number of values	n = 8
Regression equation	$y = a + b x$ (1 <sup>st</sup> order) $y = 0.000736 + 2.978150 x$
<b>Correlation coefficient</b>	<b>= 0.99997</b>

**Figure 52: Calibration function of CGA 344605 in acetonitrile  
Lab 2**



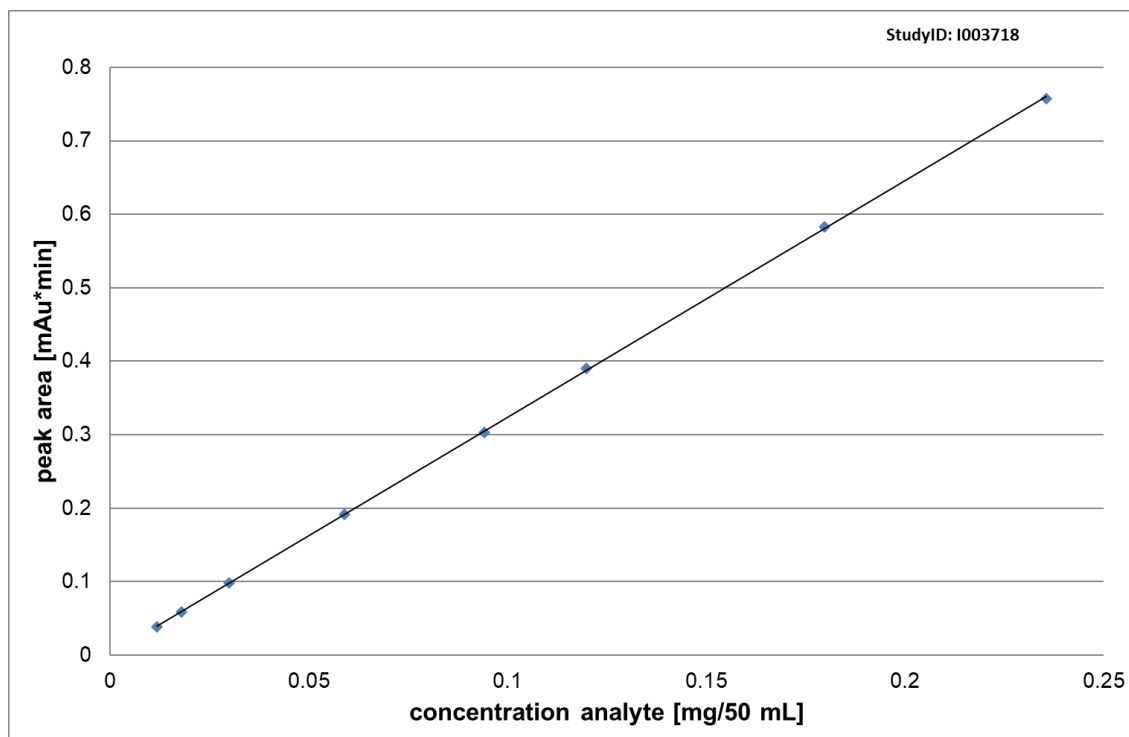
Number of values	n = 8
Regression equation	$y = a + b x$ (1 <sup>st</sup> order) $y = -0.002209 + 3.016589 x$
<b>Correlation coefficient</b>	<b>= 0.99998</b>

**Figure 53: Calibration function of CGA 344605 in acetonitrile/water  
Lab 2**



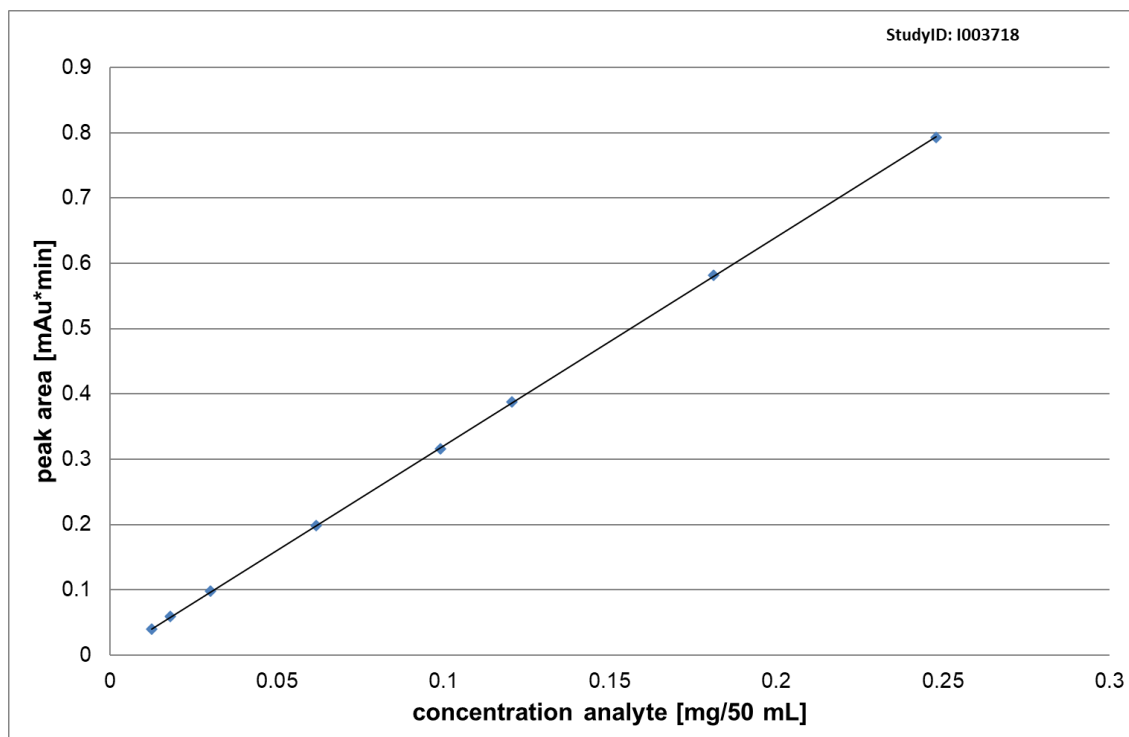
Number of values            n = 8  
Regression equation         $y = a + b x$  (1<sup>st</sup> order)  
                                       $y = -0.004383 + 3.068154 x$   
Correlation coefficient     = **0.99991**

**Figure 54: Calibration function of CGA 344605 in acetonitrile  
Lab 3**



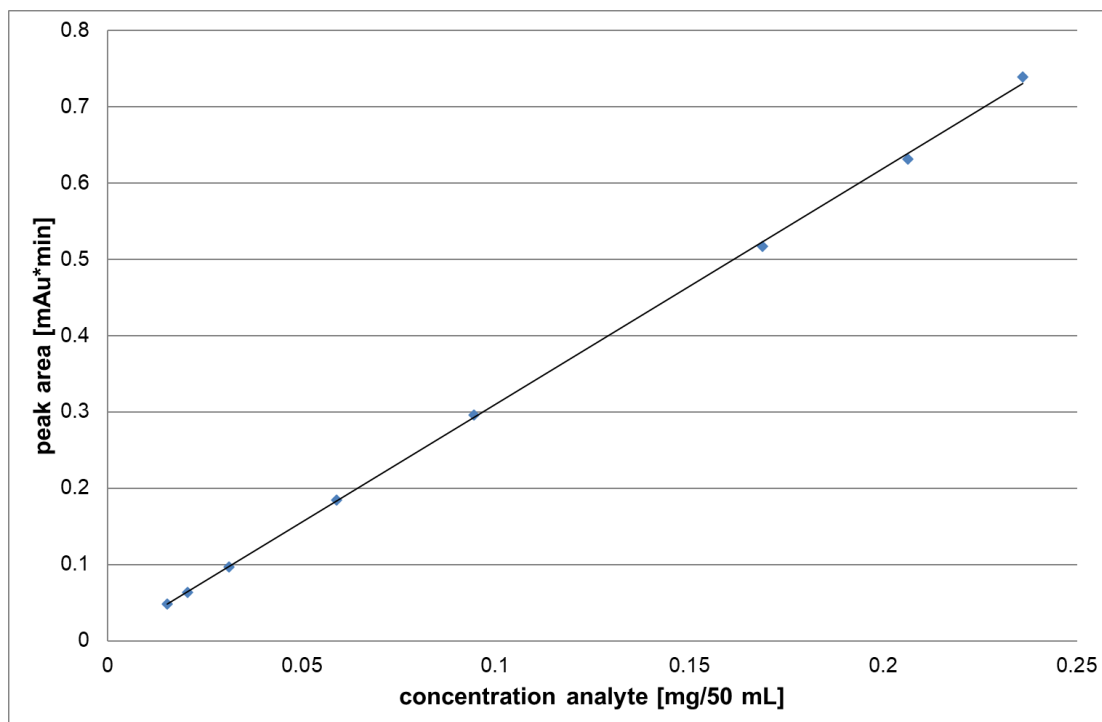
Number of values	n = 8
Regression equation	$y = a + b x$ (1 <sup>st</sup> order) $y = 0.001541 + 3.220038 x$
<b>Correlation coefficient</b>	<b>= 0.99997</b>

**Figure 55: Calibration function of CGA 344605 in acetonitrile/water  
Lab 3**



Number of values	n = 8
Regression equation	$y = a + b x$ (1 <sup>st</sup> order) $y = 0.000602 + 3.201252 x$
<b>Correlation coefficient</b>	<b>= 0.99999</b>

**Figure 56: Calibration function of CGA 344605 in acetonitrile  
Lab 4**



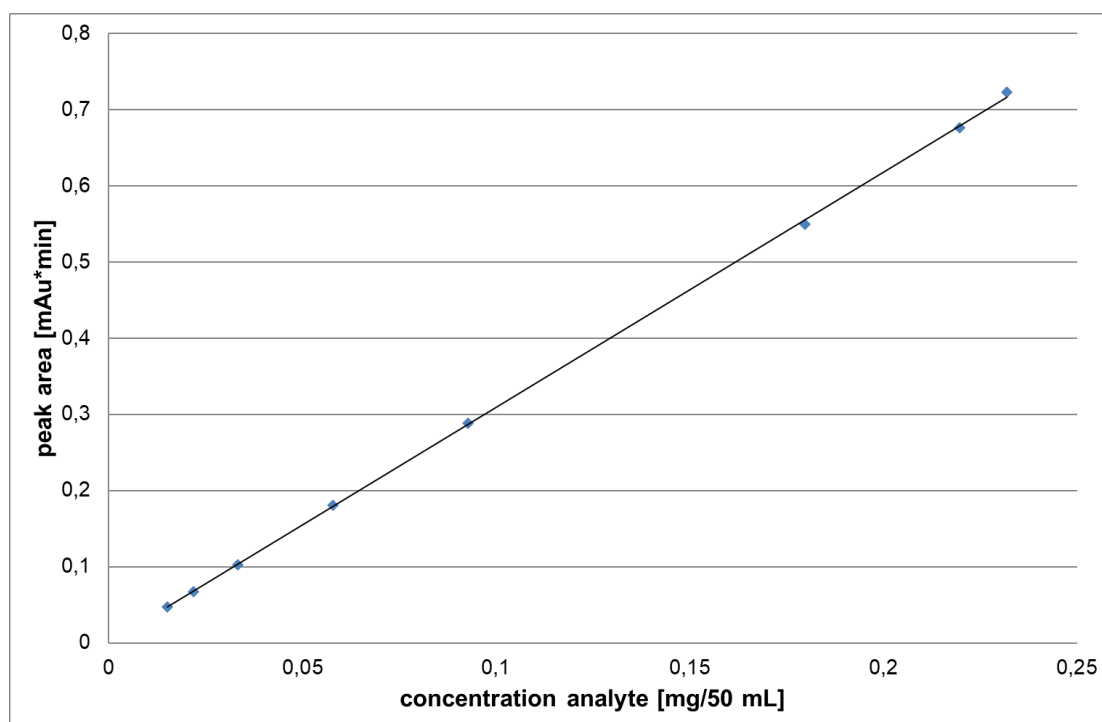
Number of values             $n = 8$

Regression equation         $y = a + b x$  (1<sup>st</sup> order)

$$y = 0.00038 + 3.09196 x$$

**Correlation coefficient = 0.99982**

**Figure 57: Calibration function of CGA 344605 in acetonitrile/water  
Lab 4**



Number of values             $n = 8$

Regression equation         $y = a + b x$  (1<sup>st</sup> order)

$$y = 0.00031 + 3.08620 x$$

**Correlation coefficient    = 0.99992**



## Appendix 1: Calculation of max. limit of CGA 344605 in the formulations

The max. accepted level (MAL) of an analyte in formulation ( $MAL_{formulation}^{analyte}$ ) is calculated based on its MAL in the respective technical grade active substance ( $MAL_{TGAS}^{analyte}$ ) and the declared content of this active substance in the formulation ( $DC_{formulation}^{a.s.}$ ) according to the following formula

$$MAL_{formulation}^{analyte} [\% (w/w)] = \frac{MAL_{TGAS}^{analyte} [\% (w/w)] \cdot DC_{formulation}^{a.s.} [\% (w/w)]}{100\% (w/w)}$$

The MAL for CGA 344605 in the technical grade active substance trifloxystrobin is 4 g/kg. equal to 0.4% (w/w).

Thus the MAL of CGA 344605 in the formulations is calculated as follow.

Formulation	$MAL_{TGAS}^{analyte}$	$DC_{formulation}^{a.s.(TFS)}$	$MAL_{formulation}^{analyte}$	
	[g/kg]	[% (w/w)]	[g/kg]	[% (w/w)]
trifloxystrobin SC 500 (500 g/L)	4	43.9	1.76	0.176
fluoastrobins + prothioconazole + trifloxystrobin EC 300 (75+150+75 g/L)	4	6.47	2.59	0.0259
trifloxystrobin WG 50 (50 g/kg)	4	50.0	2.00	0.200