

Sample Name: blind

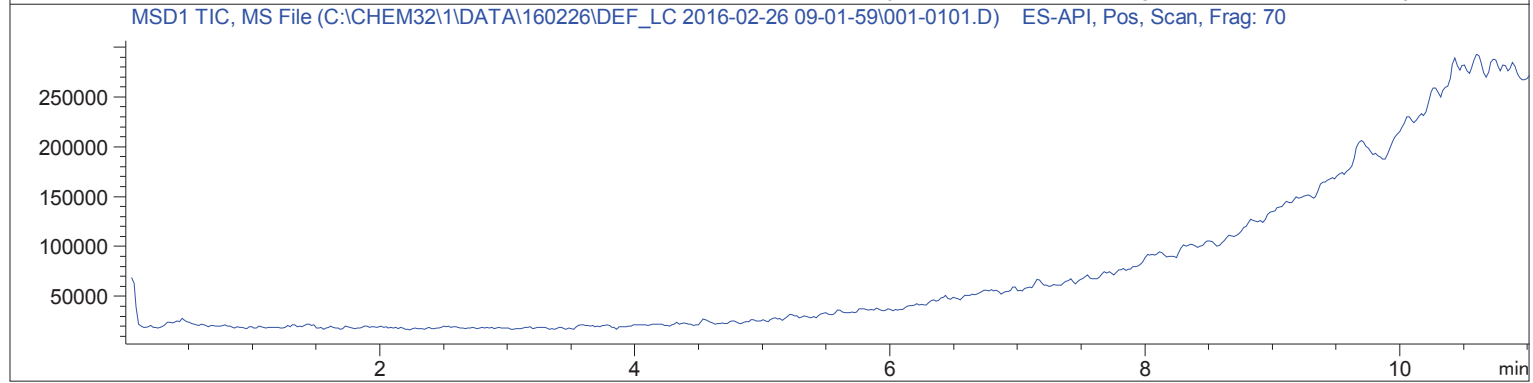
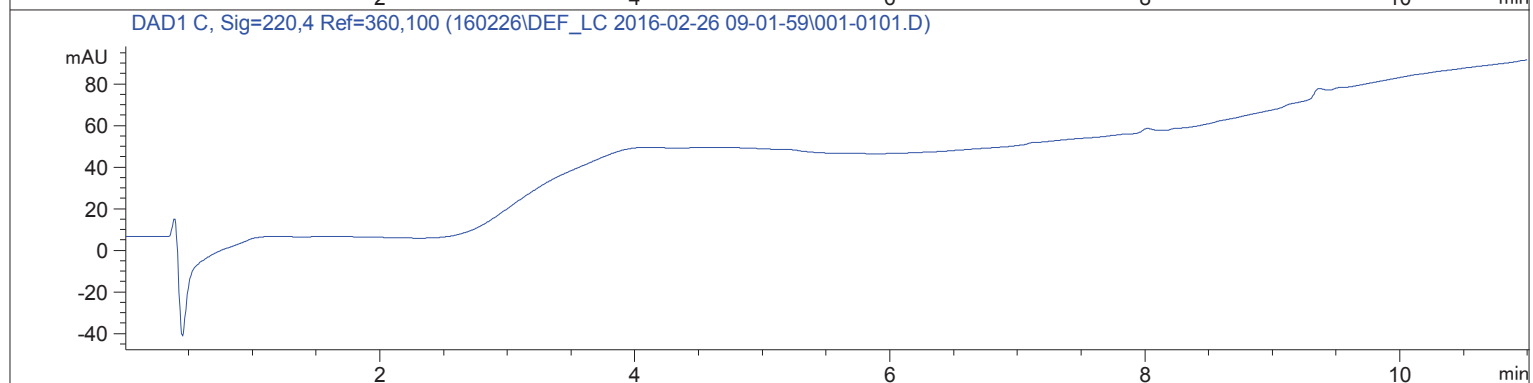
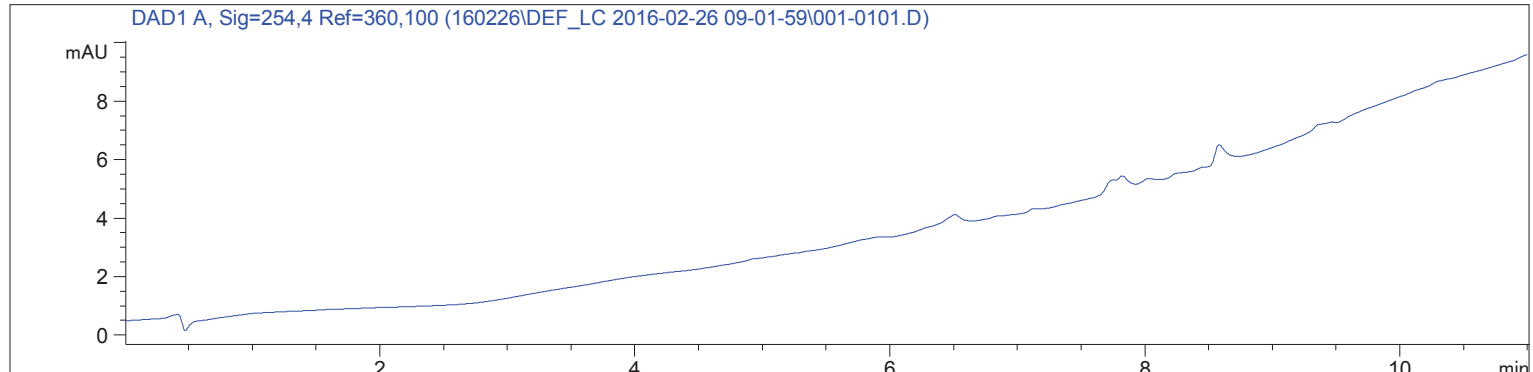
```

=====
Acq. Operator   : SYSTEM                               Seq. Line :    1
Acq. Instrument : LCMS 1                               Location  : Vial 1
Injection Date  : 2/26/2016 9:03:33 AM                Inj       :    1
                                                    Inj Volume: 1.000 µl

Acq. Method     : C:\CHEM32\1\DATA\160226\DEF_LC 2016-02-26 09-01-59\STANDARD_220_254.M
Last changed    : 2/26/2016 9:01:59 AM by SYSTEM
Analysis Method : C:\CHEM32\1\DATA\160226\DEF_LC 2016-02-26 09-01-59\STANDARD_220_254.M (
                  Sequence Method)
Last changed    : 2/26/2016 10:08:58 AM by SYSTEM
                  (modified after loading)
Method Info     : Standard ACN:Wasser 20:80 -> 97:3 Flow 0.4 ml/min 220 nm, 254 nm
=====

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Additional Info : Peak(s) manually integrated



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                          Area Percent Report
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Sorted By      :      Signal
Multiplier     :      1.0000
Dilution      :      1.0000
Do not use Multiplier & Dilution Factor with ISTDs

```

No peaks found

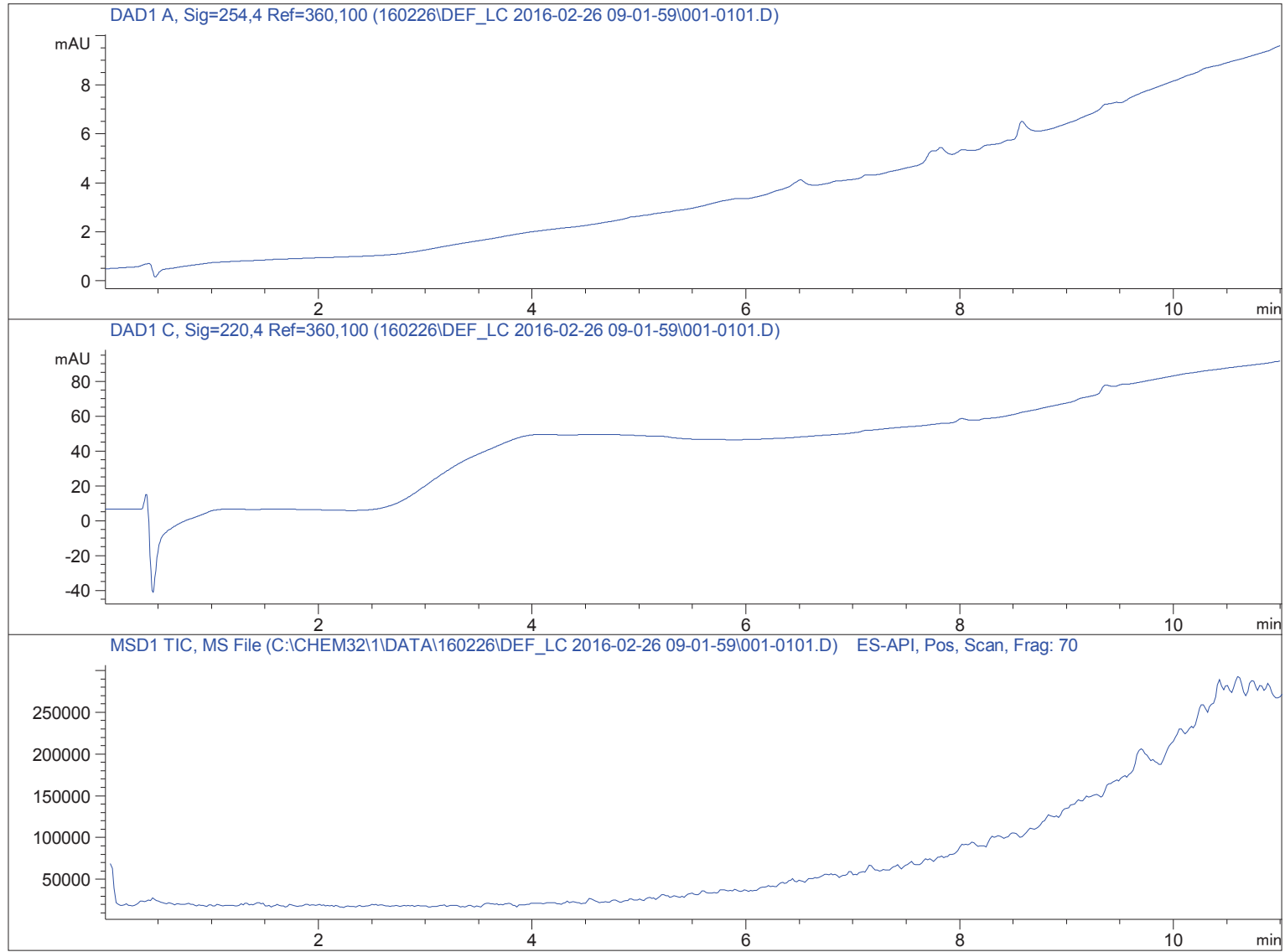
=====
*** End of Report ***

Sample Name: blind

=====

Acq. Operator	: SYSTEM	Seq. Line	: 1
Acq. Instrument	: LCMS 1	Location	: Vial 1
Injection Date	: 2/26/2016 9:03:33 AM	Inj	: 1
		Inj Volume	: 1.000 µl
Acq. Method	: C:\CHEM32\1\DATA\160226\DEF_LC 2016-02-26 09-01-59\STANDARD_220_254.M		
Last changed	: 2/26/2016 9:01:59 AM by SYSTEM		
Analysis Method	: C:\CHEM32\1\DATA\160226\DEF_LC 2016-02-26 09-01-59\STANDARD_220_254.M (Sequence Method)		
Last changed	: 2/26/2016 10:09:26 AM by SYSTEM (modified after loading)		
Method Info	: Standard ACN:Wasser 20:80 -> 97:3 Flow 0.4 ml/min 220 nm, 254 nm		

Additional Info : Peak(s) manually integrated



Spectra averaged over upper half of peaks.
Number of ions per peak: 6
Display Time Range(+/- mins): 1.0

*** End of Report ***