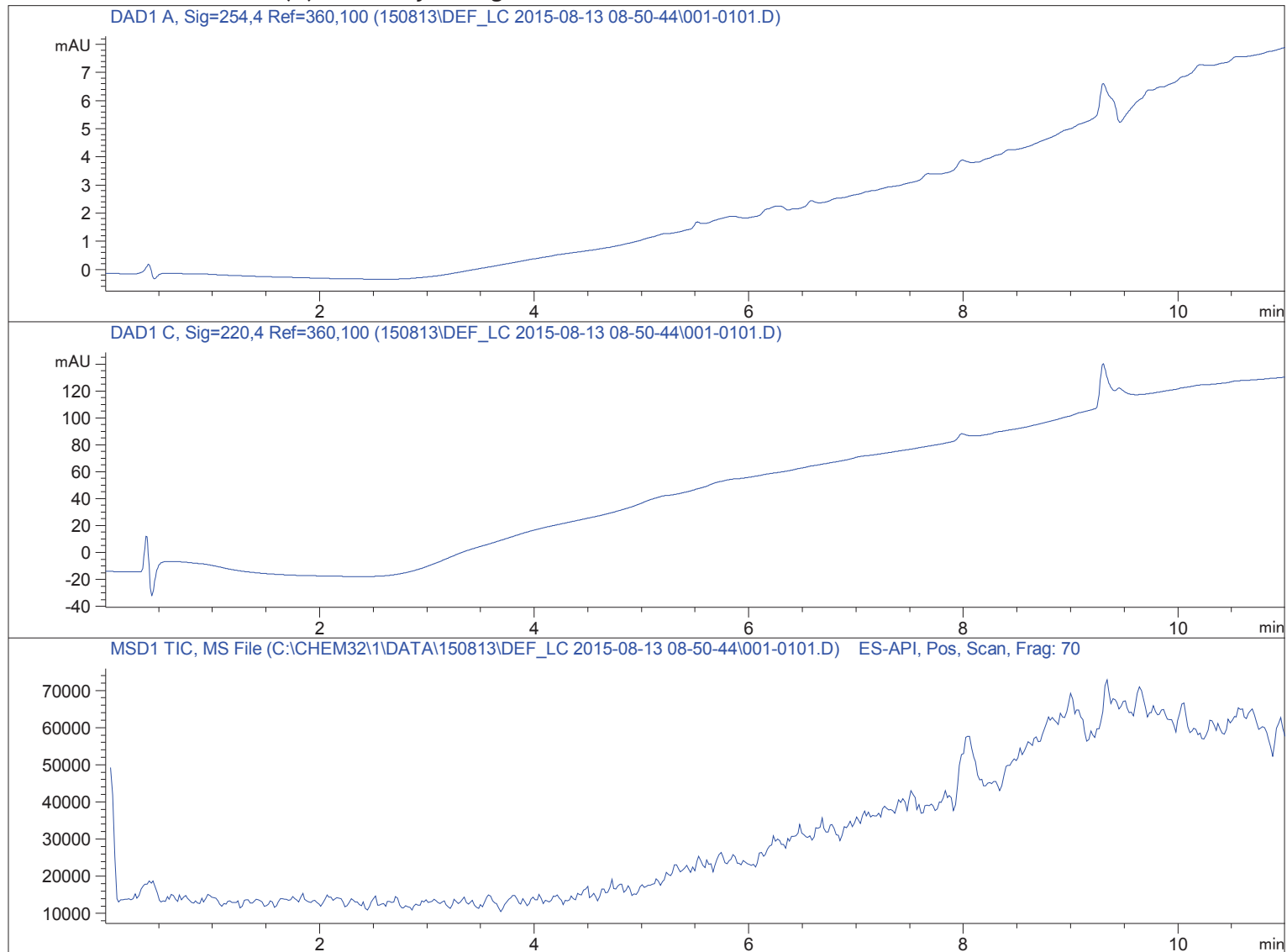


Sample Name: blind

```
=====
Acq. Operator   : SYSTEM                               Seq. Line :    1
Acq. Instrument : LCMS 1                               Location  : Vial 1
Injection Date  : 8/13/2015 8:52:40 AM                Inj       :    1
                                                    Inj Volume: 1.000 µl

Acq. Method    : C:\CHEM32\1\DATA\150813\DEF_LC 2015-08-13 08-50-44\STANDARD_220_254.M
Last changed   : 8/13/2015 8:50:44 AM by SYSTEM
Analysis Method: C:\CHEM32\1\DATA\150813\DEF_LC 2015-08-13 08-50-44\STANDARD_220_254.M (
Sequence Method)
Last changed   : 2/26/2016 10:13:24 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



```
=====
                          Area Percent Report
=====
```

```
Sorted By      :      Signal
Multiplier     :      1.0000
Dilution      :      1.0000
Do not use Multiplier & Dilution Factor with ISTDs
```

No peaks found

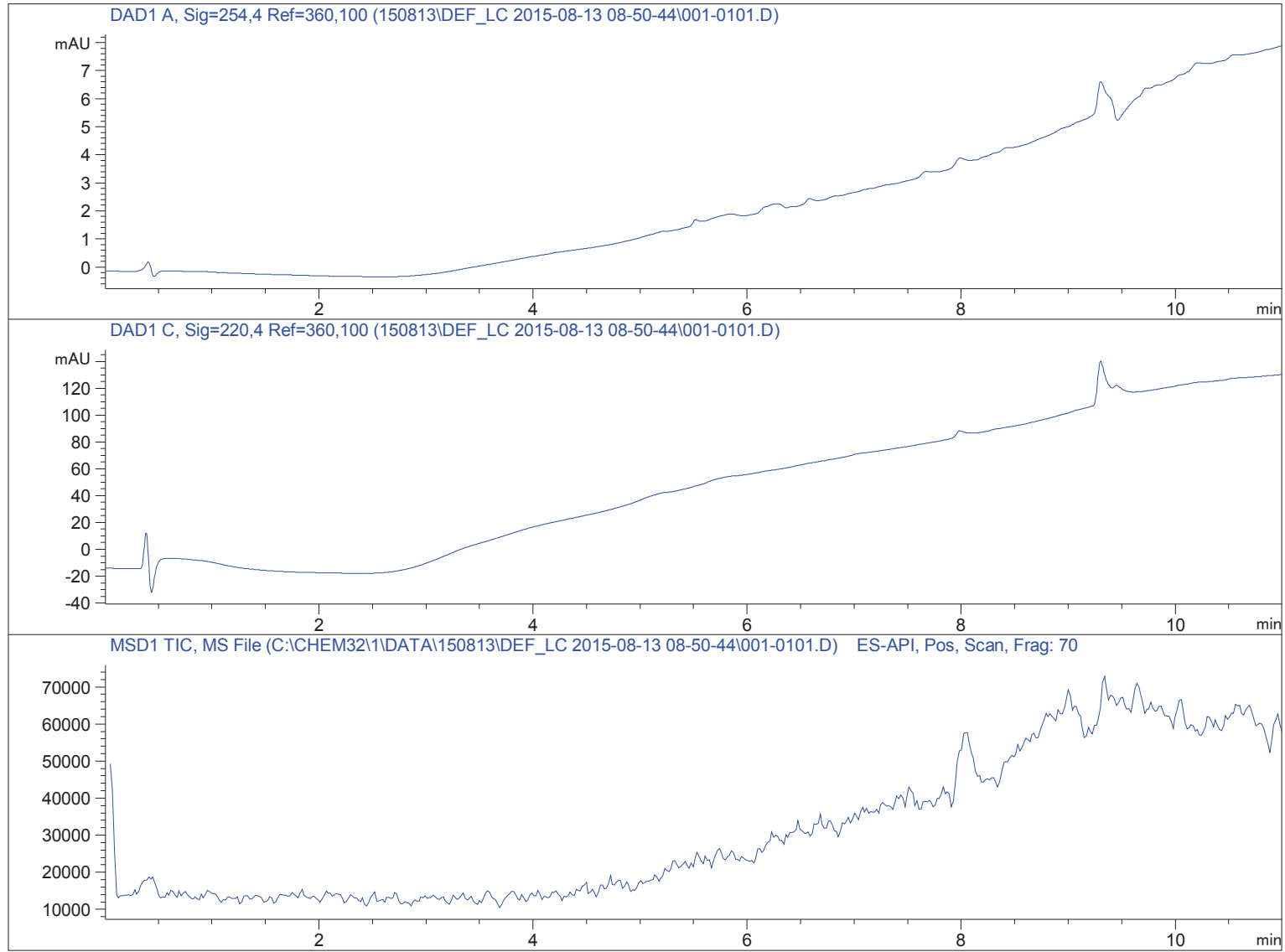
=====
*** End of Report ***

Sample Name: blind

=====

Acq. Operator	: SYSTEM	Seq. Line	: 1
Acq. Instrument	: LCMS 1	Location	: Vial 1
Injection Date	: 8/13/2015 8:52:40 AM	Inj	: 1
		Inj Volume	: 1.000 µl
Acq. Method	: C:\CHEM32\1\DATA\150813\DEF_LC 2015-08-13 08-50-44\STANDARD_220_254.M		
Last changed	: 8/13/2015 8:50:44 AM by SYSTEM		
Analysis Method	: C:\CHEM32\1\DATA\150813\DEF_LC 2015-08-13 08-50-44\STANDARD_220_254.M (Sequence Method)		
Last changed	: 2/26/2016 10:13:43 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 ***