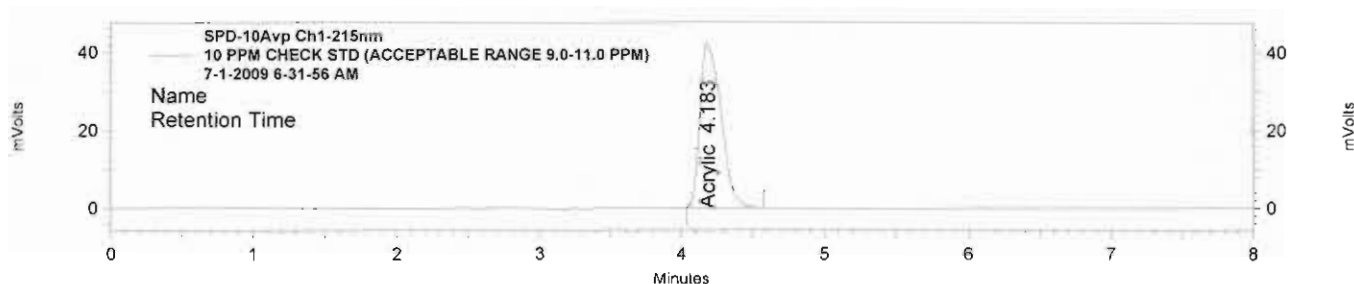


Custom Method Report

Instrument: HPLC_4
 Sample ID: 10 PPM CHECK STD (ACCEPTABLE RANGE 9.0-11.0 PPM)
 Method Name: C:\CLASS-VP\Enterprise\Projects\residual monomer hplc 4\Method\residual monomer4.met
 Data: C:\CLASS-VP\Enterprise\Projects\residual monomer hplc 4\Data\7-1-2009 6-31-56 AM
 User: MANNT
 Acquired: 7/1/2009 6:32:38 AM
 Printed: 7/1/2009 4:48:57 PM
 Dilution factor: 1
 Multiplier: 1

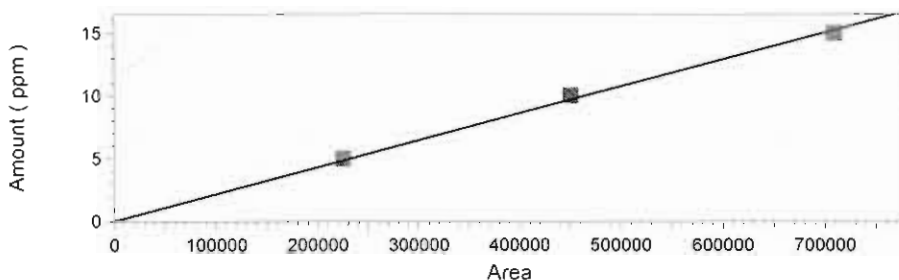


SPD-10Avp
 Ch1-215nm Results

Name	Area	Retention Time	Concentration
Acrylic	429864	4.183	9.244

Must have a Goodness Fit of Greater than 0.985

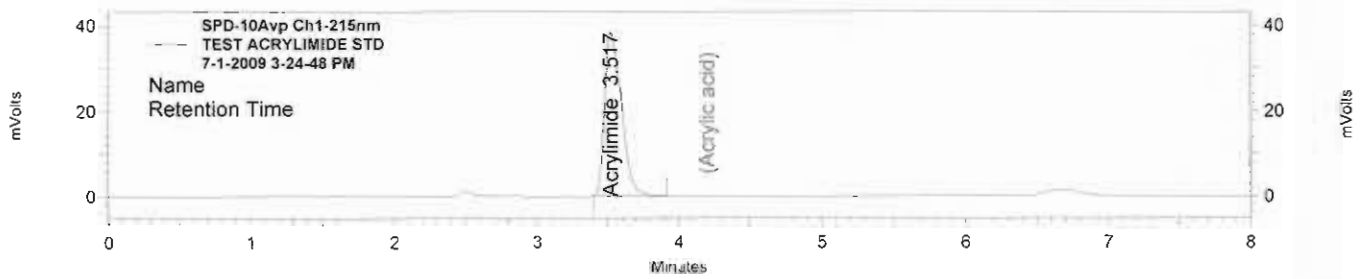
Peak: Acrylic -- ESTD -- SPD-10Avp Ch1



Acrylic (SPD-10Avp Ch1)
 Average RF: 2.18545e-005 RF StDev: 6.06761e-007 RF %RSD: 2.77636
 Scaling: None LSQ Weighting: None Force Through Zero: On
 Replicate Mode: Replace
 Fit Type: Linear
 $y = 2.15043e-005x + 0.000000$
 Goodness of fit (r^2): 0.996321

Custom Method Report

Instrument: HPLC_4
Sample ID: TEST ACRYLIMIDE STD
Method Name: C:\CLASS-VP\Enterprise\Projects\residual monomer hplc 4\Method\residual monomer4.met
Data: C:\CLASS-VP\Enterprise\Projects\residual monomer hplc 4\Data\7-1-2009 3-24-48 PM
User: MANNT
Acquired: 7/1/2009 3:25:22 PM
Printed: 7/1/2009 4:45:33 PM
Dilution factor: 2
Multiplier: 100

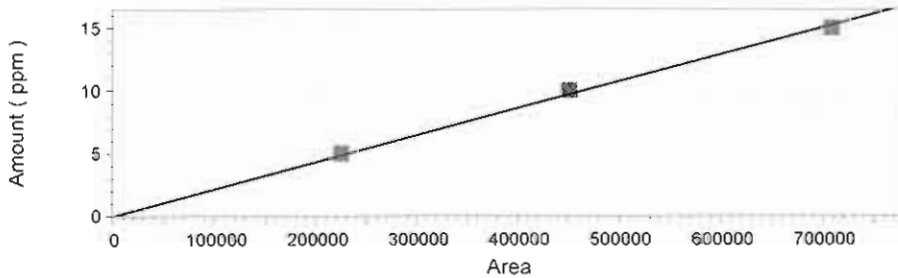


SPD-10Avp
Ch1-215nm Results

Name	Area	Retention Time	Concentration
Acrylimide	318451	3.517	342.403
Acrylic acid			0.000 BDL

Must have a Goodness Fit of Greater than 0.985

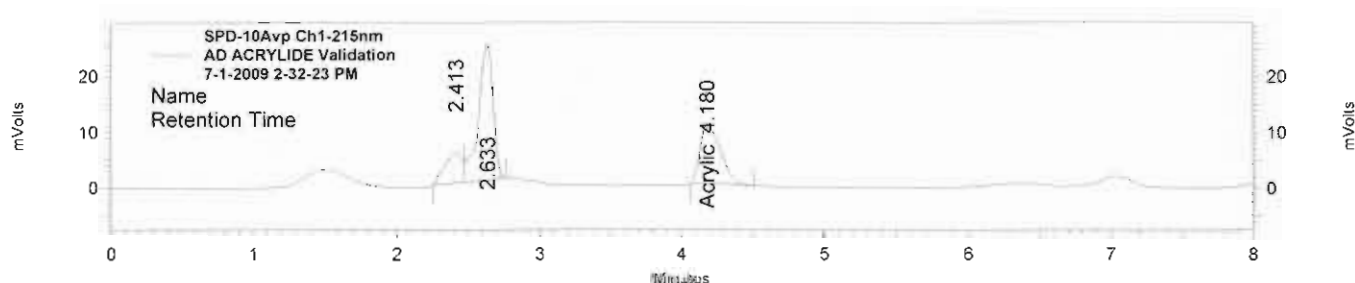
Peak: Acrylimide -- ESTD -- SPD-10Avp Ch1



Acrylimide (SPD-10Avp Ch1)
Average RF: 2.18545e-005 RF StDev: 6.06761e-007 RF %RSD: 2.77636
Scaling: None LSQ Weighting: None Force Through Zero: On
Replicate Mode: Replace
Fit Type: Linear
 $y = 2.15043e-005x + 0.000000$
Goodness of fit (r²): 0.996321

Custom Method Report

Instrument: HPLC_4
 Sample ID: AD ACRYLIDE Validation
 Method Name: C:\CLASS-VP\Enterprise\Projects\residual monomer hplc 4\Method\residual monomer4.met
 Data: C:\CLASS-VP\Enterprise\Projects\residual monomer hplc 4\Data\7-1-2009 2-32-23 PM
 User: MANNT
 Acquired: 7/1/2009 2:32:58 PM
 Printed: 7/1/2009 2:41:15 PM
 Dilution factor: 2
 Multiplier: 100

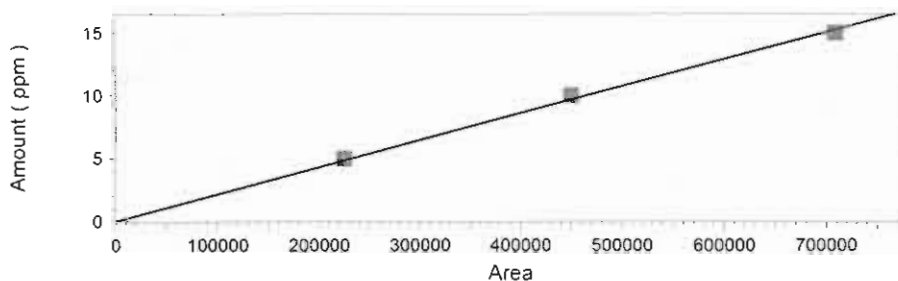


SPD-10Avp
Ch1-215nm Results

Name	Area	Retention Time	Concentration
Acrylic	106688	4.180	114.712

Must have a Goodness Fit of Greater than 0.985

Peak: Acrylic -- ESTD -- SPD-10Avp Ch1



Acrylic (SPD-10Avp Ch1)

Average RF: 2.18545e-005 RF StDev: 6.06761e-007 RF %RSD: 2.77636

Scaling: None LSQ Weighting: None Force Through Zero: On

Replicate Mode: Replace

Fit Type: Linear

$y = 2.15043e-005x + 0.000000$

Goodness of fit (r^2): 0.996321