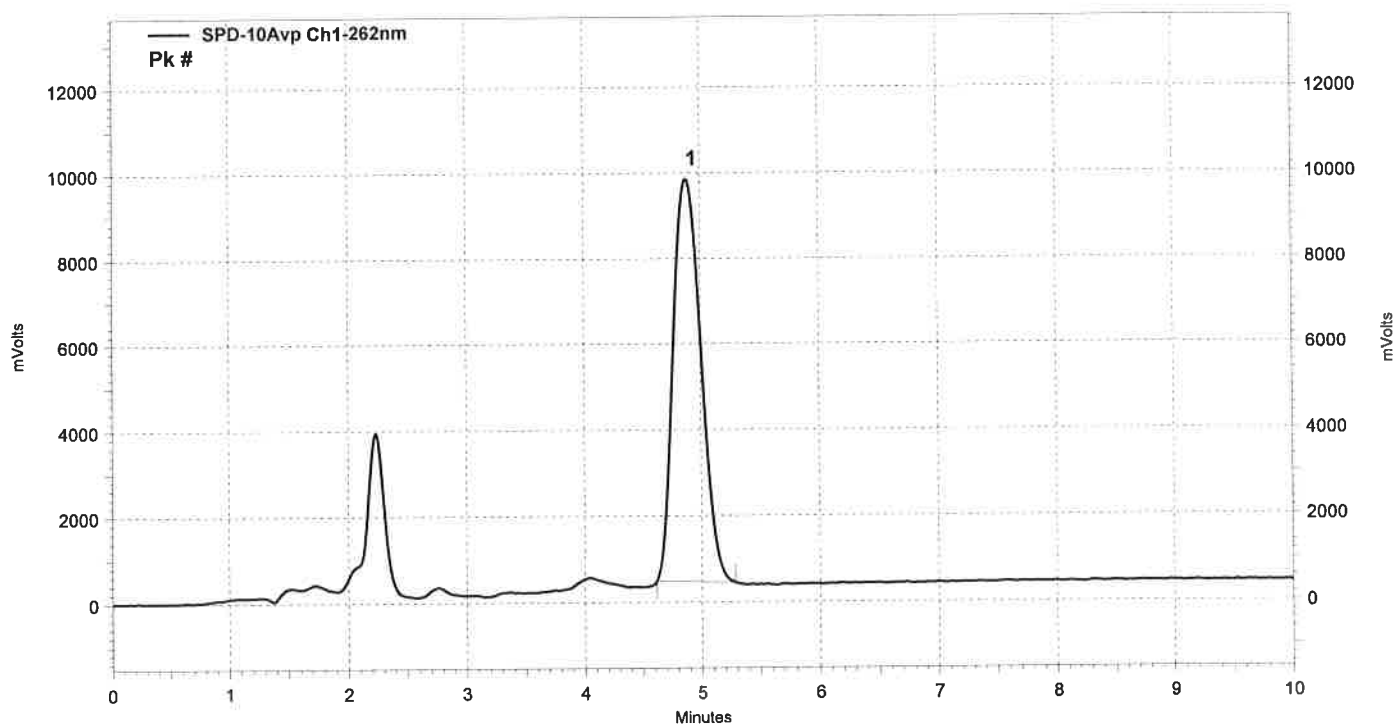


Sample: PBQ std 3,125 µg/ml
Runtime: 23.2.2017 16:23:07

Vial: 1, 50 µl
User: System
File: C:\CLASS-VP\Mondal 2017\230217 ajot\LAB7351loadIII.dat
Method: C:\CLASS-VP\Mondal 2017\5% NaAc, 95% MeOH, pH 3.4 isocratic 10 min, 262, 251 nm.met
Sequence: C:\CLASS-VP\Mondal 2017\020317 ajot\MondalI.seq
Analysis: 23.2.2017 16:33:21

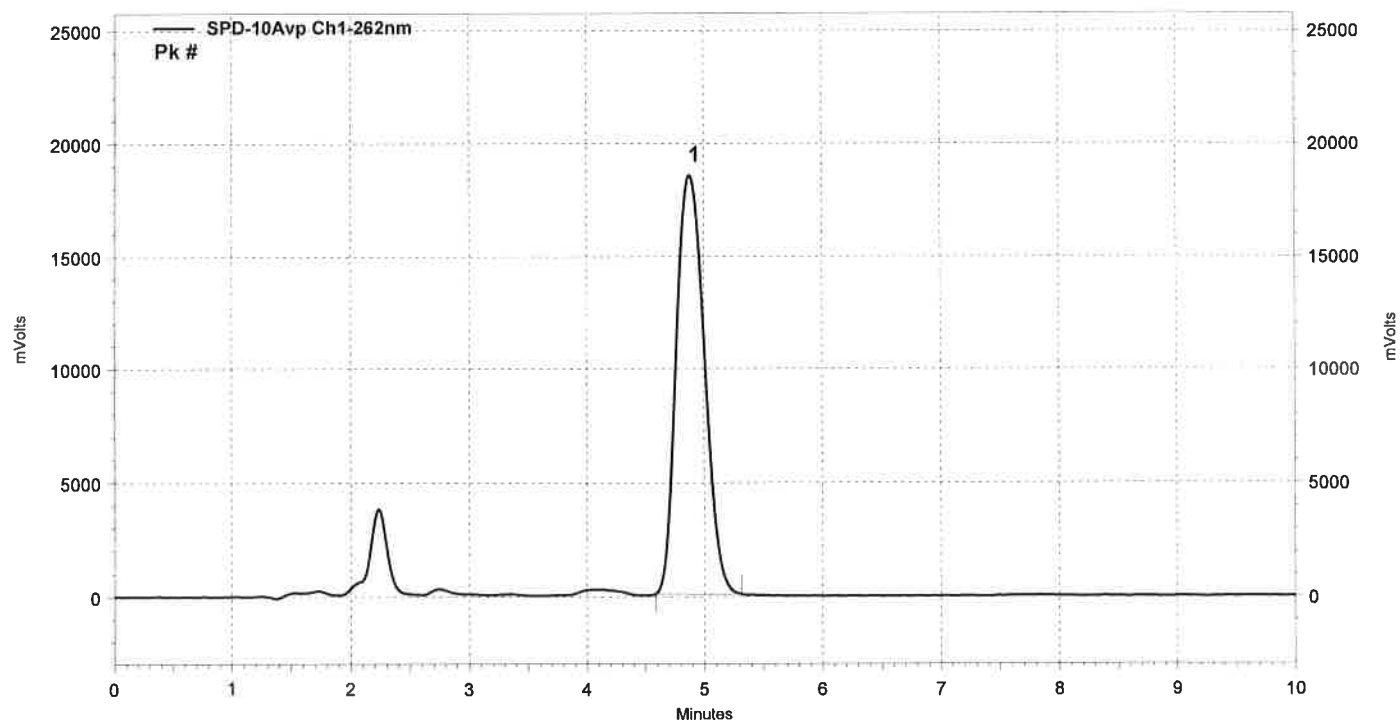


SPD-10Avp
Ch1-262nm Results

Pk #	Retention Time	Area	Height
1	4,883	153654	9376
Totals		153654	9376

Sample: PBQ std 6,25µg/ml
Runtime: 23.2.2017 16:35:44

Vial: 2 , 50 µl
User: System
File: C:\CLASS-VP\Mondal 2017\230217 ajot\LAB7352loadIII.dat
Method: C:\CLASS-VP\Mondal 2017\5% NaAc, 95% MeOH, pH 3.4 isocratic 10 min, 262, 251 nm.met
Sequence: C:\CLASS-VP\Mondal 2017\020317 ajot\MondalI.seq
Analysis: 23.2.2017 16:45:58

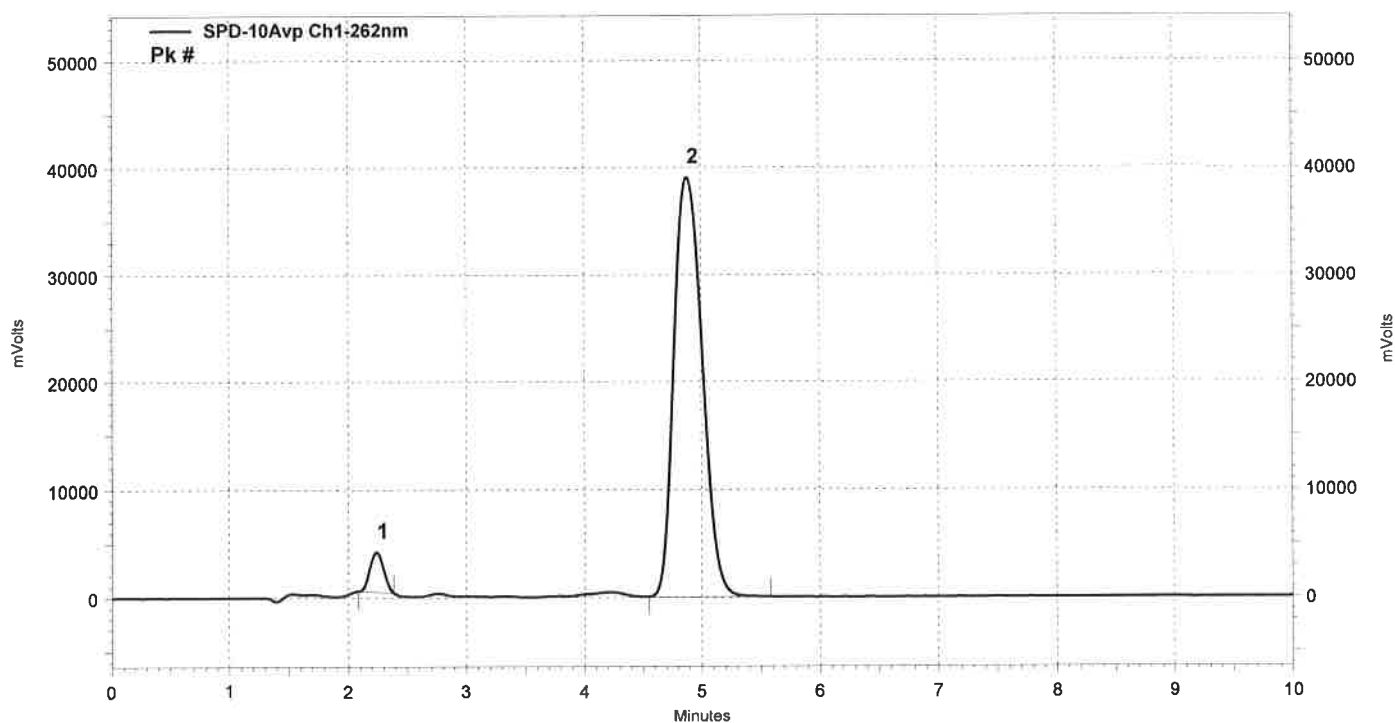


SPD-10Avp
Ch1-262nm Results

Pk #	Retention Time	Area	Height
1	4,875	305901	18473
Totals		305901	18473

Sample: PBQ std 12,5 µg/ml
Runtime: 23.2.2017 16:48:21

Vial: 3 , 50 µl
User: System
File: C:\CLASS-VP\Mondal 2017\230217 ajot\LAB7353loadIII.dat
Method: C:\CLASS-VP\Mondal 2017\5% NaAc, 95% MeOH, pH 3.4 isocratic 10 min, 262,
251 nm.met
Sequence: C:\CLASS-VP\Mondal 2017\020317 ajot\MondalI.seq
Analysis: 23.2.2017 16:58:34

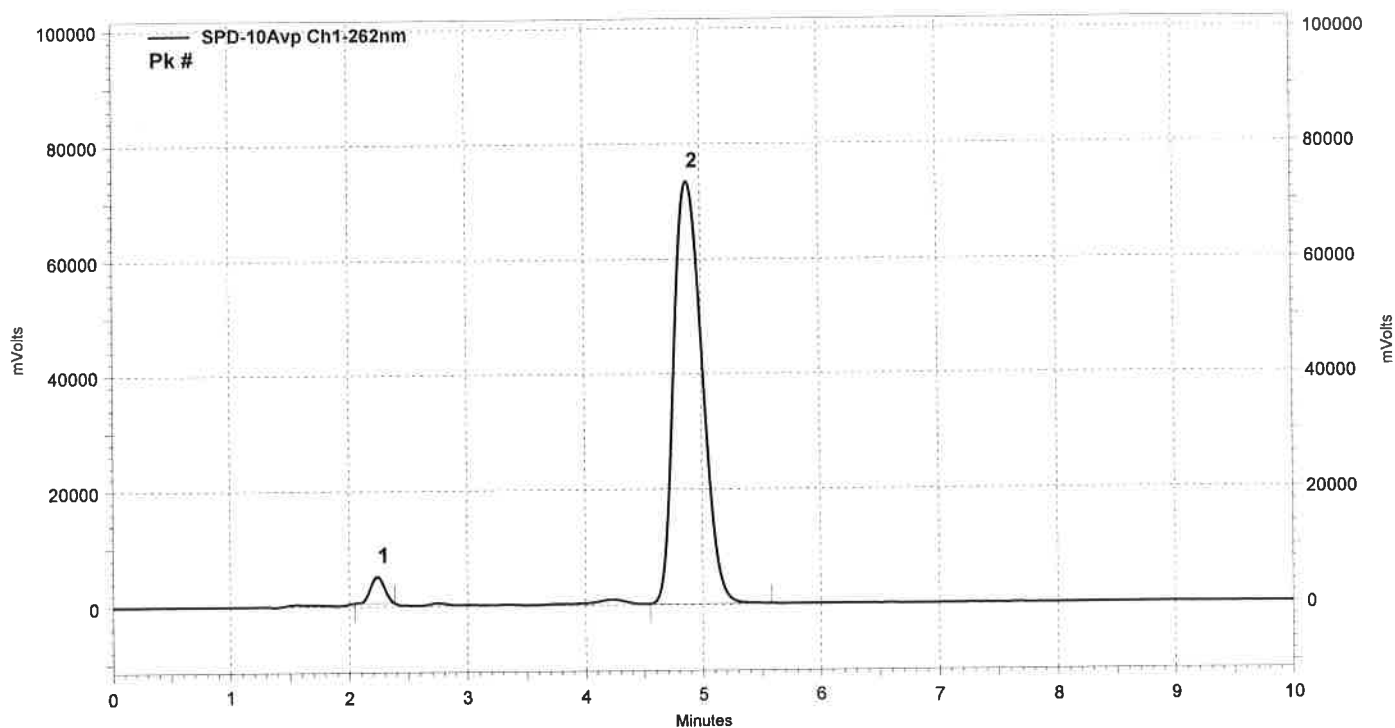


SPD-10Avp
Ch1-262nm Results

Pk #	Retention Time	Area	Height
1	2,242	29778	3682
2	4,883	652480	38988
Totals		682258	42670

Sample: PBQ std 25 µg/ml
Runtime: 23.2.2017 17:00:57

Vial: 4 , 50 µl
User: System
File: C:\CLASS-VP\Mondal 2017\230217 ajot\LAB7354loadIII.dat
Method: C:\CLASS-VP\Mondal 2017\5% NaAc, 95% MeOH, pH 3.4 isocratic 10 min, 262,
251 nm.met
Sequence: C:\CLASS-VP\Mondal 2017\020317 ajot\MondalI.seq
Analysis: 23.2.2017 17:11:11



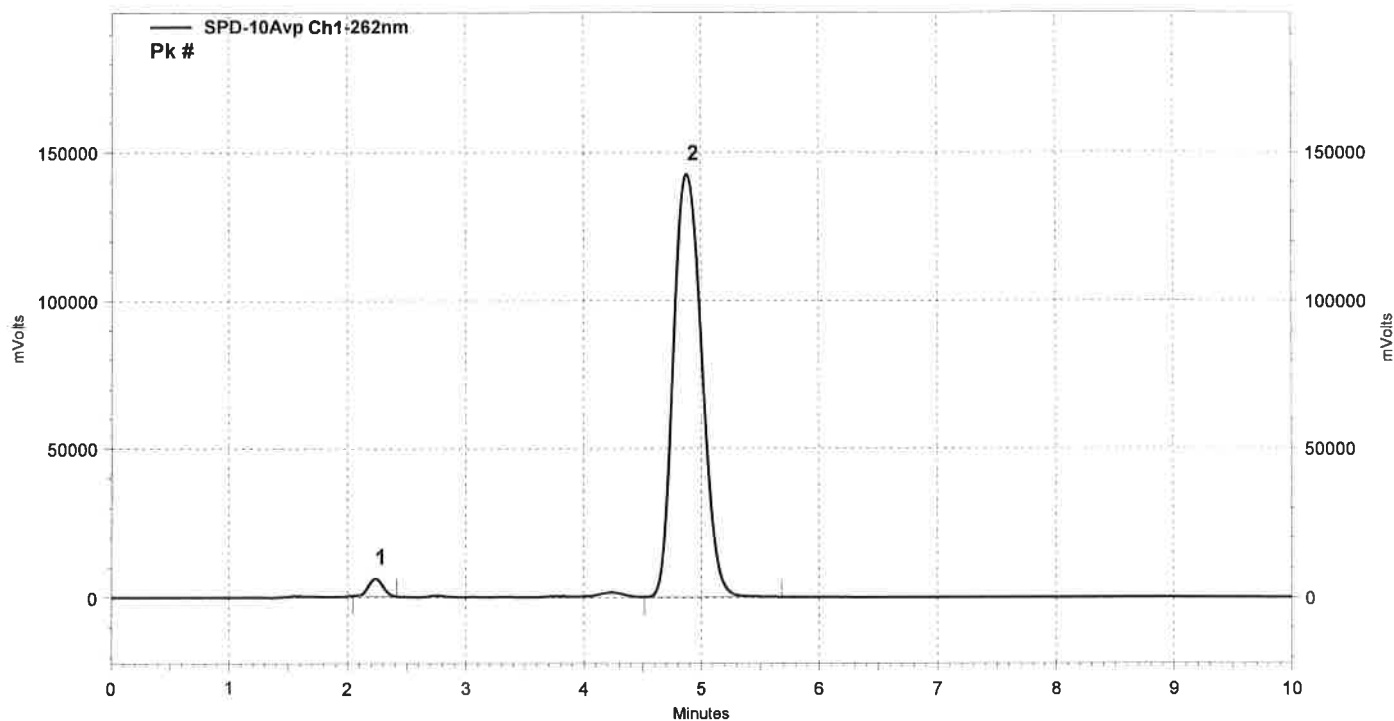
SPD-10Avp

Ch1-262nm Results

Pk #	Retention Time	Area	Height
1	2,242	37671	4568
2	4,883	1226545	73273
Totals		1264216	77841

Sample: PBQ std 50 µg/ml
Runtime: 23.2.2017 17:13:34

Vial: 5 , 50 µl
User: System
File: C:\CLASS-VP\Mondal 2017\230217 ajot\LAB7355loadIII.dat
Method: C:\CLASS-VP\Mondal 2017\5% NaAc, 95% MeOH, pH 3.4 isocratic 10 min, 262,
251 nm.met
Sequence: C:\CLASS-VP\Mondal 2017\020317 ajot\MondalI.seq
Analysis: 23.2.2017 17:23:45

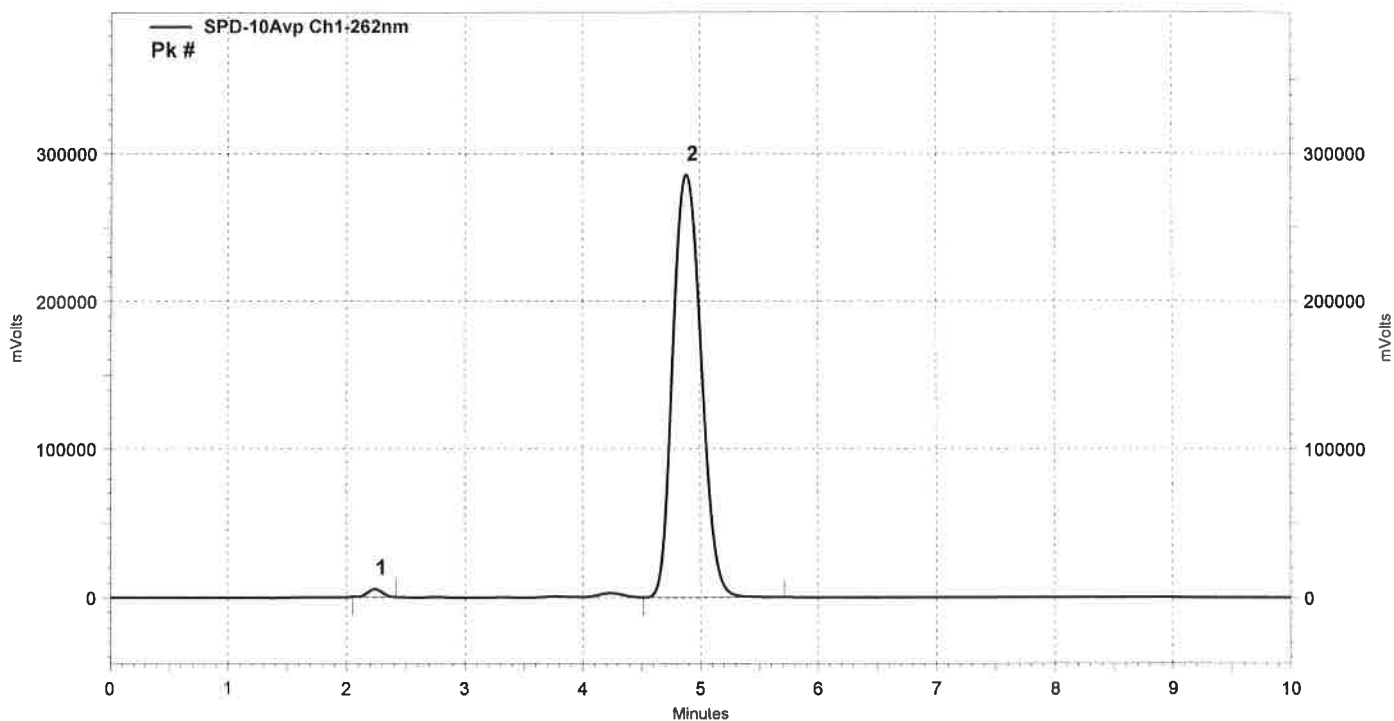


SPD-10Avp
Ch1-262nm Results

Pk #	Retention Time	Area	Height
1	2,233	49976	5843
2	4,883	2384377	142509
Totals		2434353	148352

Sample: PBQ std 100 µg/ml
Runtime: 23.2.2017 17:26:07

Vial: 6 , 50 µl
User: System
File: C:\CLASS-VP\Mondal 2017\230217 ajot\LAB7356loadIII.dat
Method: C:\CLASS-VP\Mondal 2017\5% NaAc, 95% MeOH, pH 3.4 isocratic 10 min, 262, 251 nm.met
Sequence: C:\CLASS-VP\Mondal 2017\020317 ajot\MondalI.seq
Analysis: 23.2.2017 17:36:20

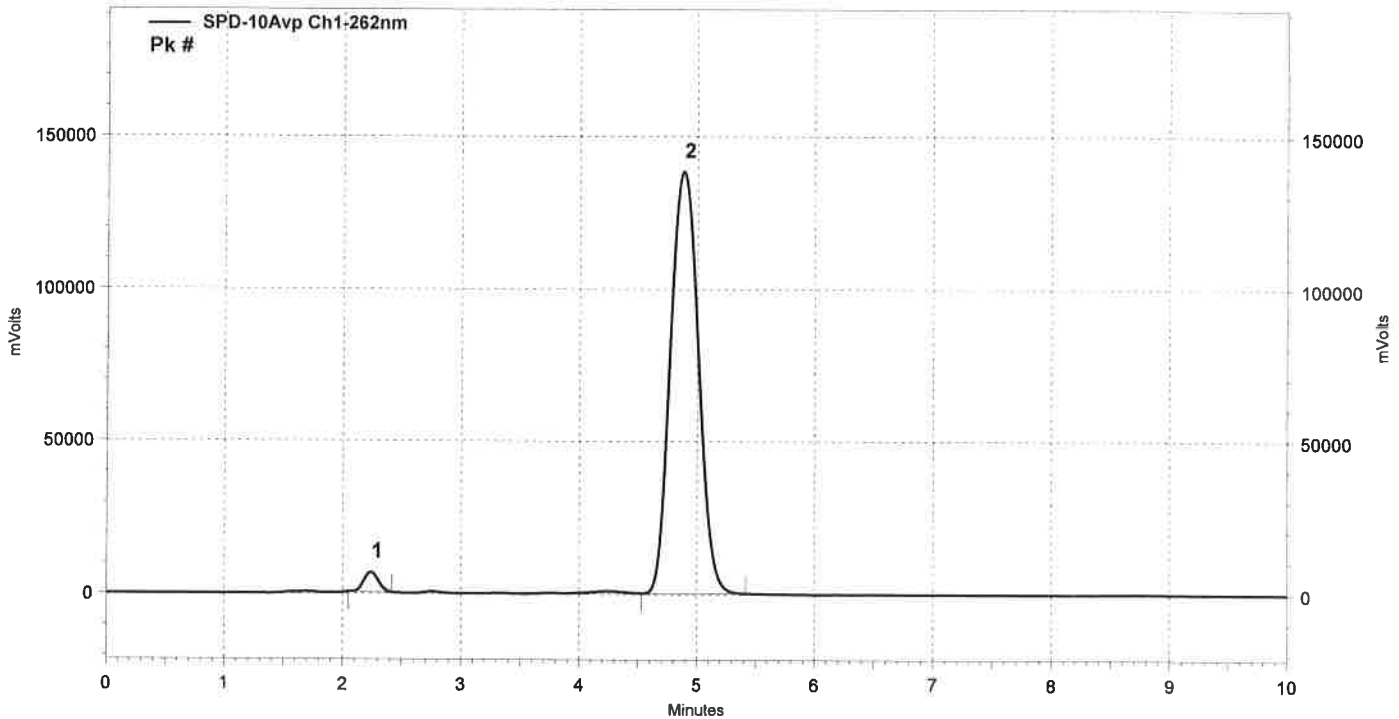


SPD-10Avp
Ch1-262nm Results

Pk #	Retention Time	Area	Height
1	2,242	46285	5416
2	4,883	4785534	285403
Totals		4831819	290819

Sample: NP-PBQ AL (20 ml) putki 4, ~~1-100~~
Runtime: 23.2.2017 18:03:55

Vial: 9 , 50 µl
User: System
File: C:\CLASS-VP\Mondal 2017\230217 ajot\LAB7359loadIII.dat
Method: C:\CLASS-VP\Mondal 2017\5% NaAc, 95% MeOH, pH 3.4 isocratic 10 min, 262, 251 nm.met
Sequence: C:\CLASS-VP\Mondal 2017\020317 ajot\MondalI.seq
Analysis: 23.2.2017 18:14:08



SPD-10Avp

Ch1-262nm Results

Pk #	Retention Time	Area	Height
1	2,242	54297	6401
2	4,883	2284832	138077
Totals		2339129	144478