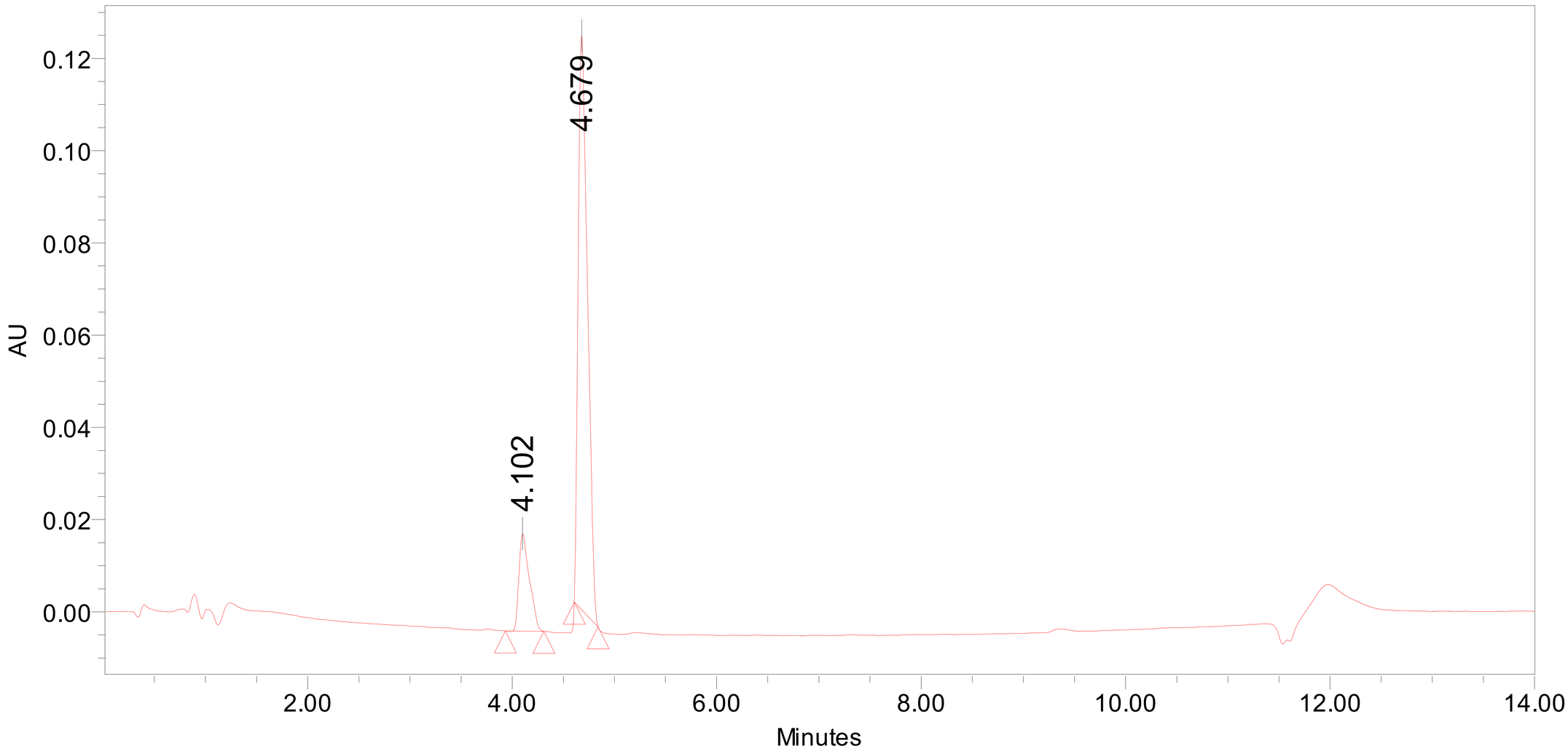
SAMPLE INFORMATION

Sample Name:	AV388_SO_Blank	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	388 so nac
Vial:	1	Acq. Method Set:	AB_95 in 10_total 12
Injection #:	1	Processing Method:	Anna General PM
Injection Volume:	15.00 ul	Channel Name:	280.0nm@1
Run Time:	14.0 Minutes	Proc. Chnl. Descr.:	PDA 280.0 nm
Date Acquired:	2024-04-30 15:26:45 CET		
Date Processed:	2024-04-30 15:51:59 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		4.102	150473	21125		
2		4.679	795405	124436		

PDA Result Table

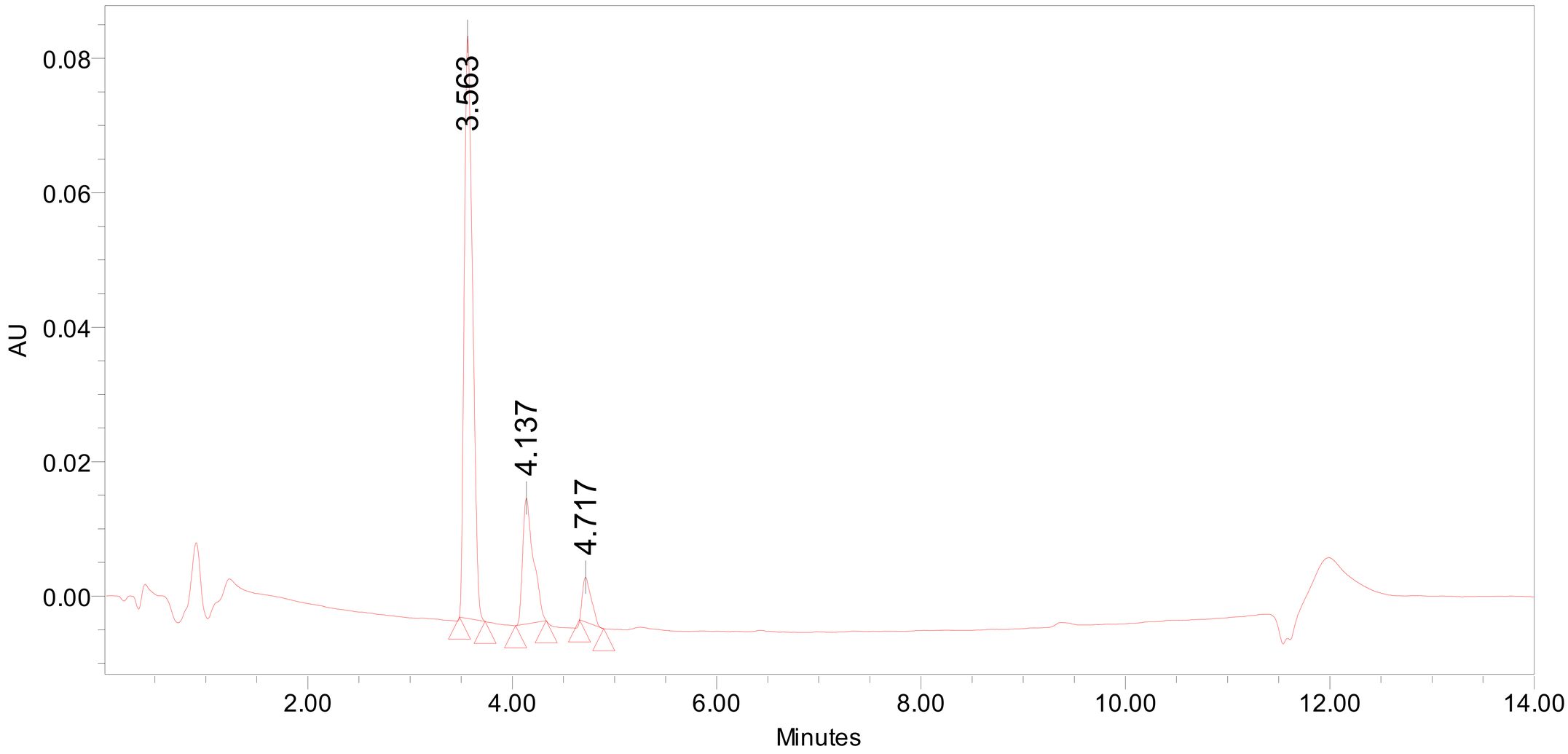
	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		4.102					
2		4.679					

Untitled

SAMPLE INFORMATION

Sample Name:	AV388_SO_NAC 1 h	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	388 so nac
Vial:	2	Acq. Method Set:	AB_95 in 10_total 12
Injection #:	1	Processing Method:	Anna General PM
Injection Volume:	15.00 ul	Channel Name:	280.0nm@1
Run Time:	14.0 Minutes	Proc. Chnl. Descr.:	PDA 280.0 nm
Date Acquired:	2024-04-30 16:11:36 CET		
Date Processed:	2024-04-30 16:32:51 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		3.563	486453	86635		
2		4.137	135627	18732		
3		4.717	39886	6692		

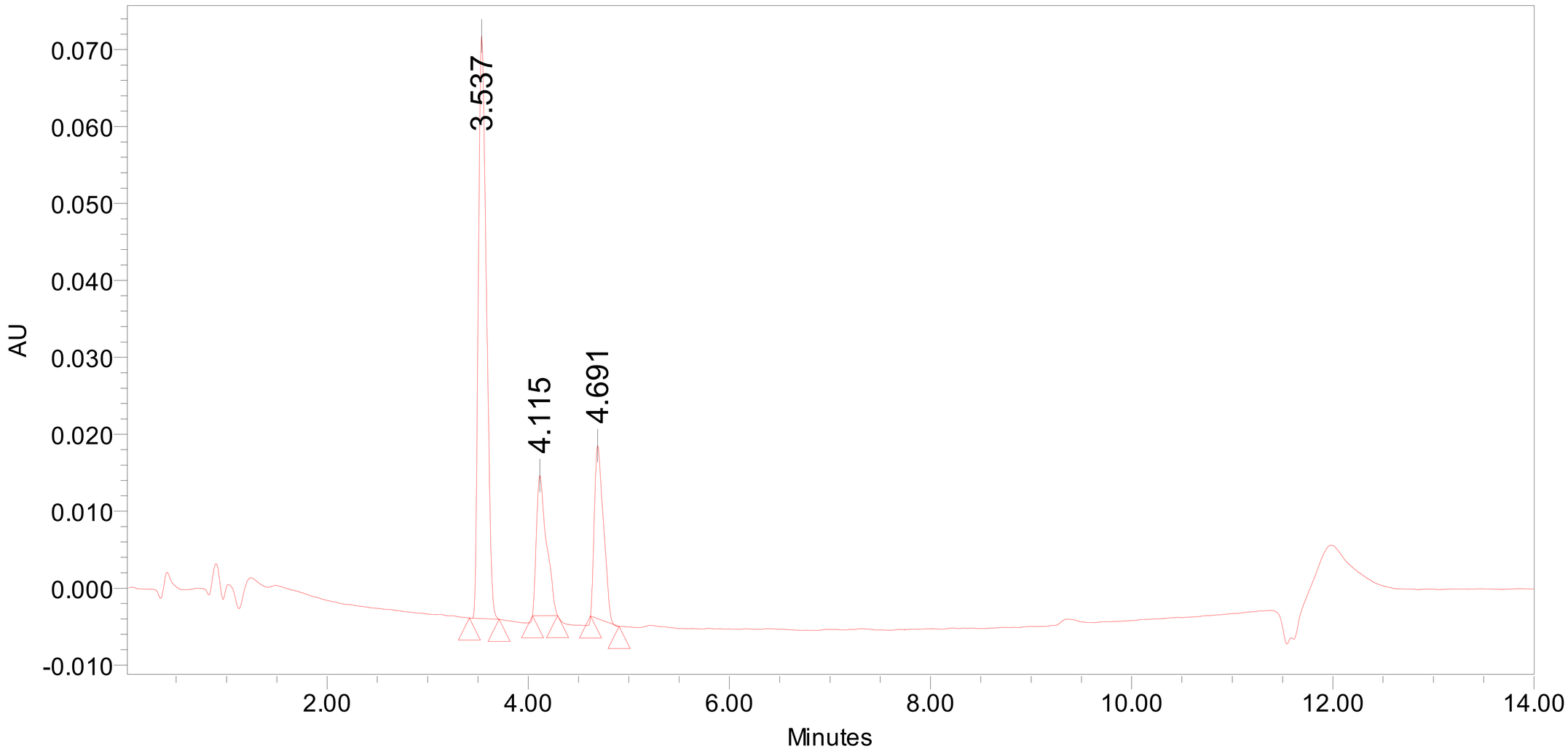
PDA Result Table

	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		3.563					
2		4.137					
3		4.717					

SAMPLE INFORMATION

Sample Name:	AV388_SO_NAC 30 min	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	388 so nac
Vial:	2	Acq. Method Set:	AB_95 in 10_total 12
Injection #:	1	Processing Method:	Anna General PM
Injection Volume:	15.00 ul	Channel Name:	280.0nm@1
Run Time:	14.0 Minutes	Proc. Chnl. Descr.:	PDA 280.0 nm
Date Acquired:	2024-04-30 15:42:23 CET		
Date Processed:	2024-04-30 16:03:29 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		3.537	423313	75710		
2		4.115	126155	18238		
3		4.691	140946	22491		

PDA Result Table

	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		3.537					
2		4.115					
3		4.691					