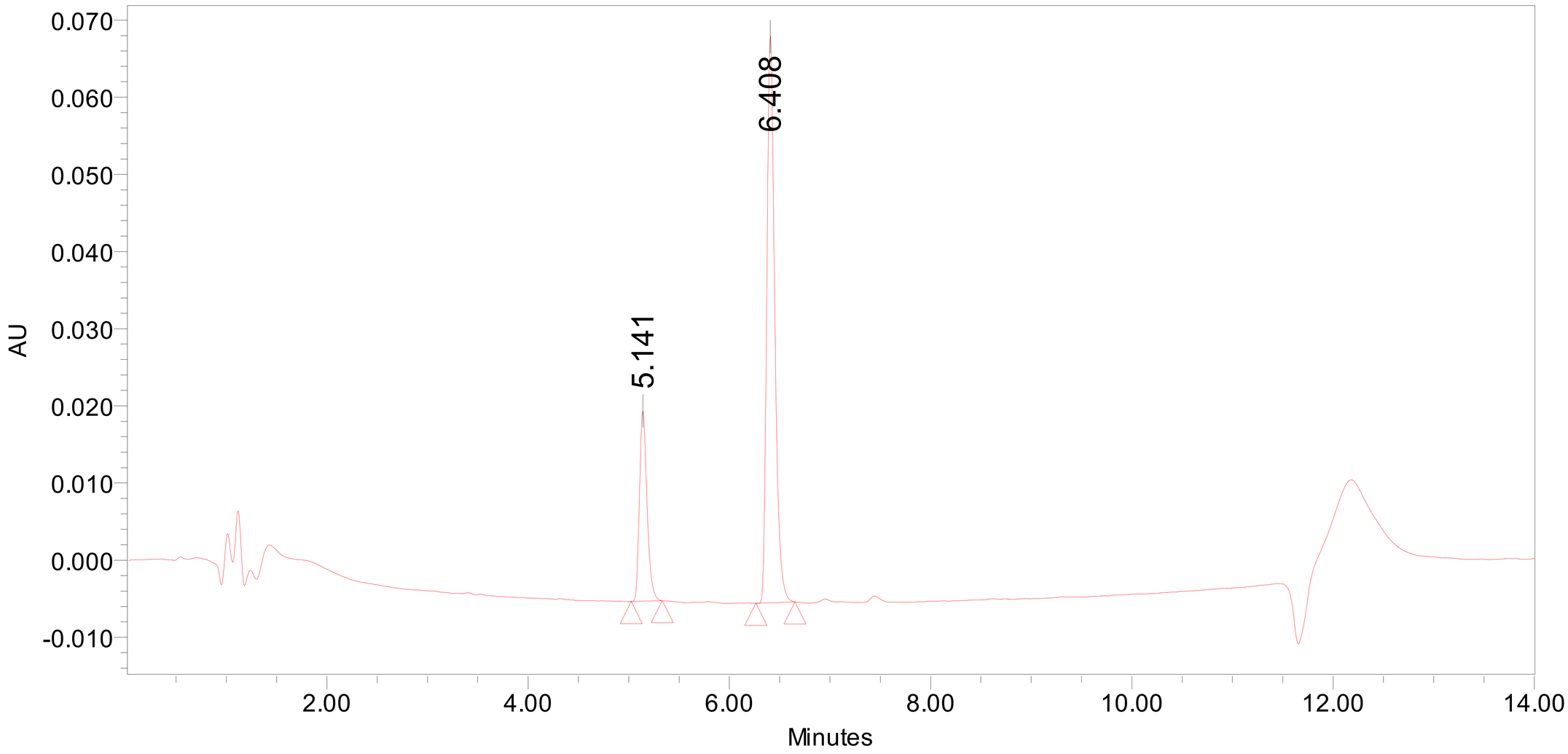
SAMPLE INFORMATION

Sample Name:	HN241-Blank	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	PROBES NAC HN241
Vial:	3	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:	2025-02-17 12:10:57 CET		
Date Processed:	2025-02-17 12:42:40 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		5.141	128526	24663		
2		6.408	405007	73478		

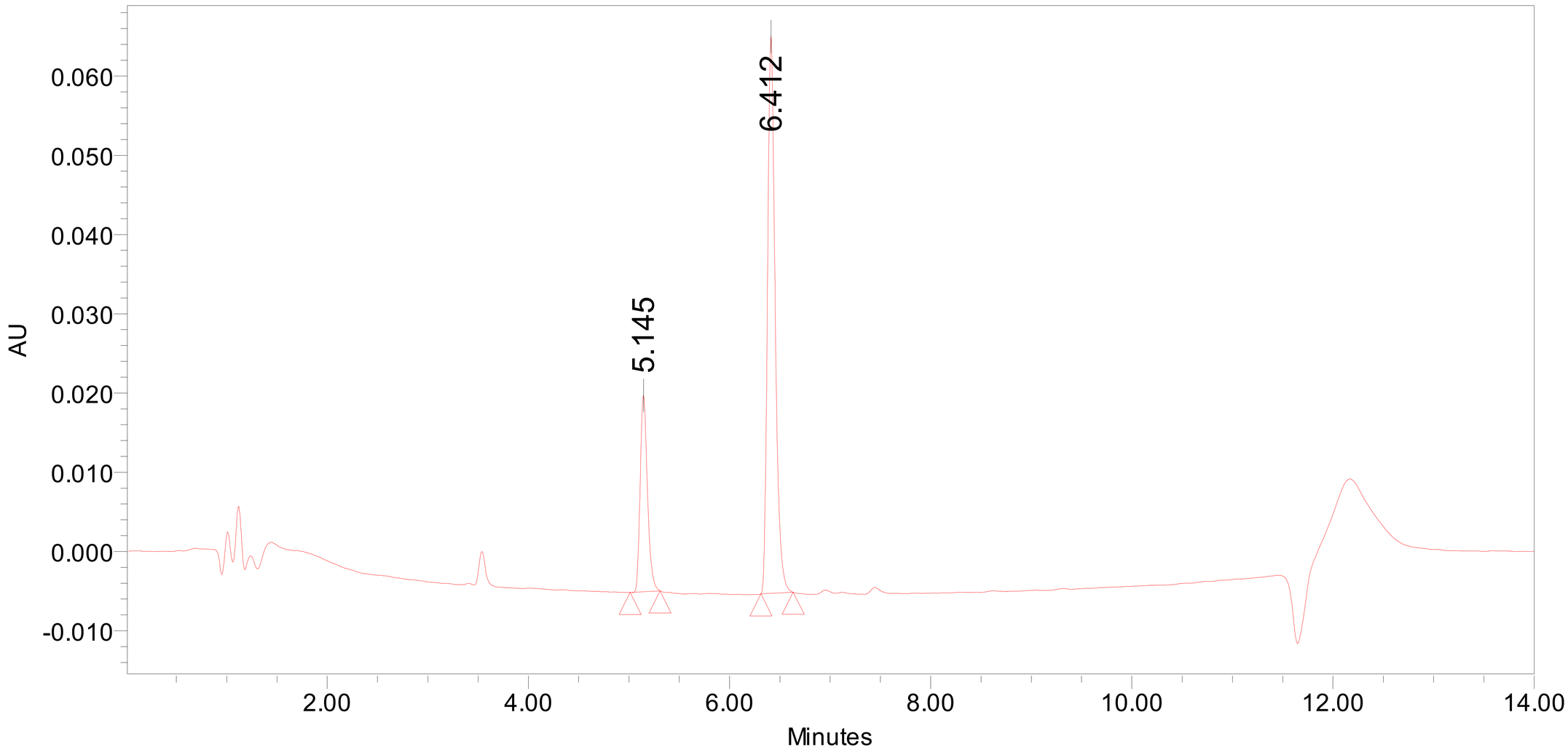
PDA Result Table

	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		5.141					
2		6.408					

SAMPLE INFORMATION

Sample Name:	HN241-Stability-1 day	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	0 and HN241 STABILITY 1 DAY
Vial:	3	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:	2025-02-18 17:03:28 CET		
Date Processed:	2025-02-21 09:01:24 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		5.145	124782	24820		
2		6.412	373887	70341		

PDA Result Table

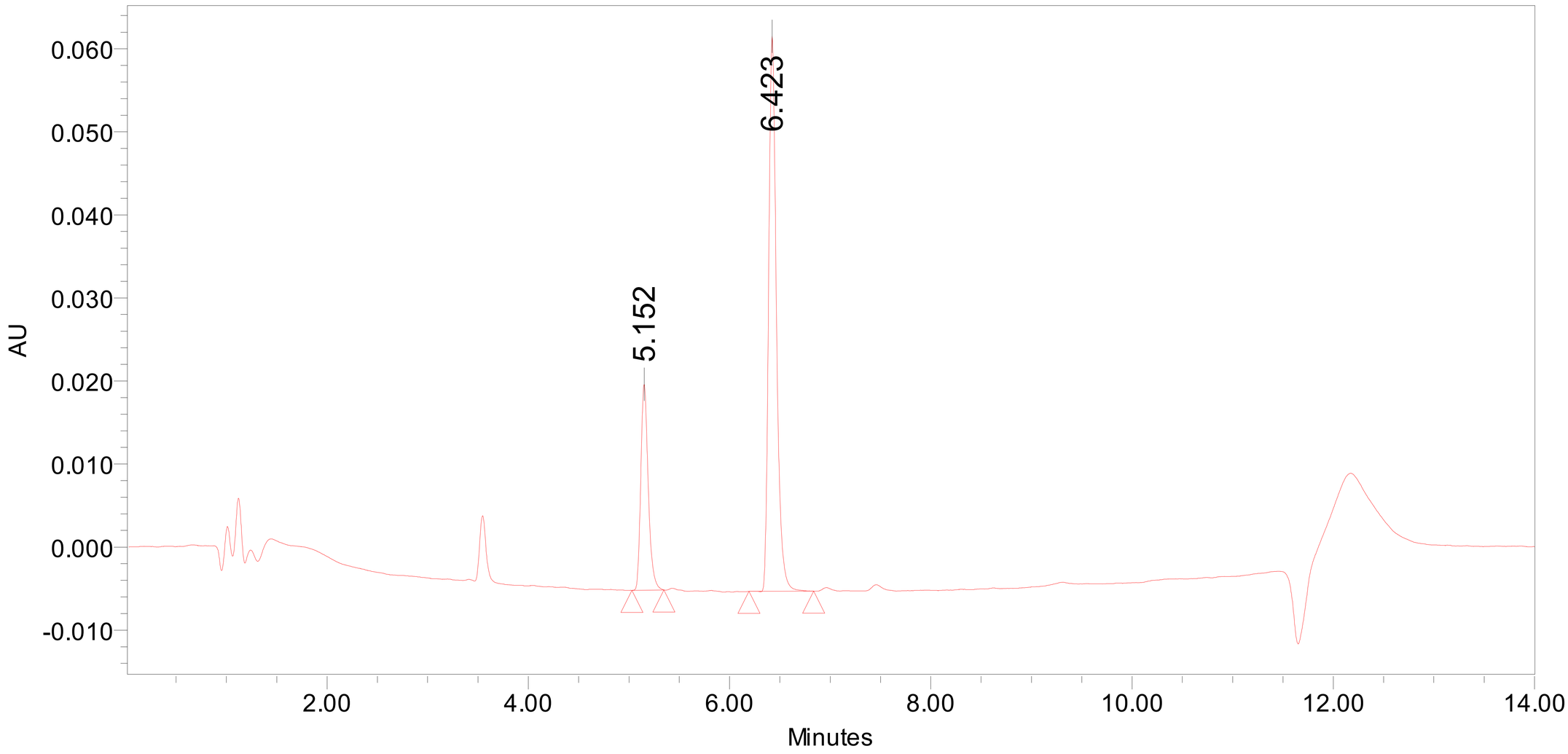
	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		5.145					
2		6.412					

Untitled

SAMPLE INFORMATION

Sample Name:	HN241-Stability 2 days	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	hn241stability 2 days
Vial:	3	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:	2025-02-19 17:34:37 CET		
Date Processed:	2025-02-21 09:01:42 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		5.152	127427	24810		
2		6.423	361307	66859		

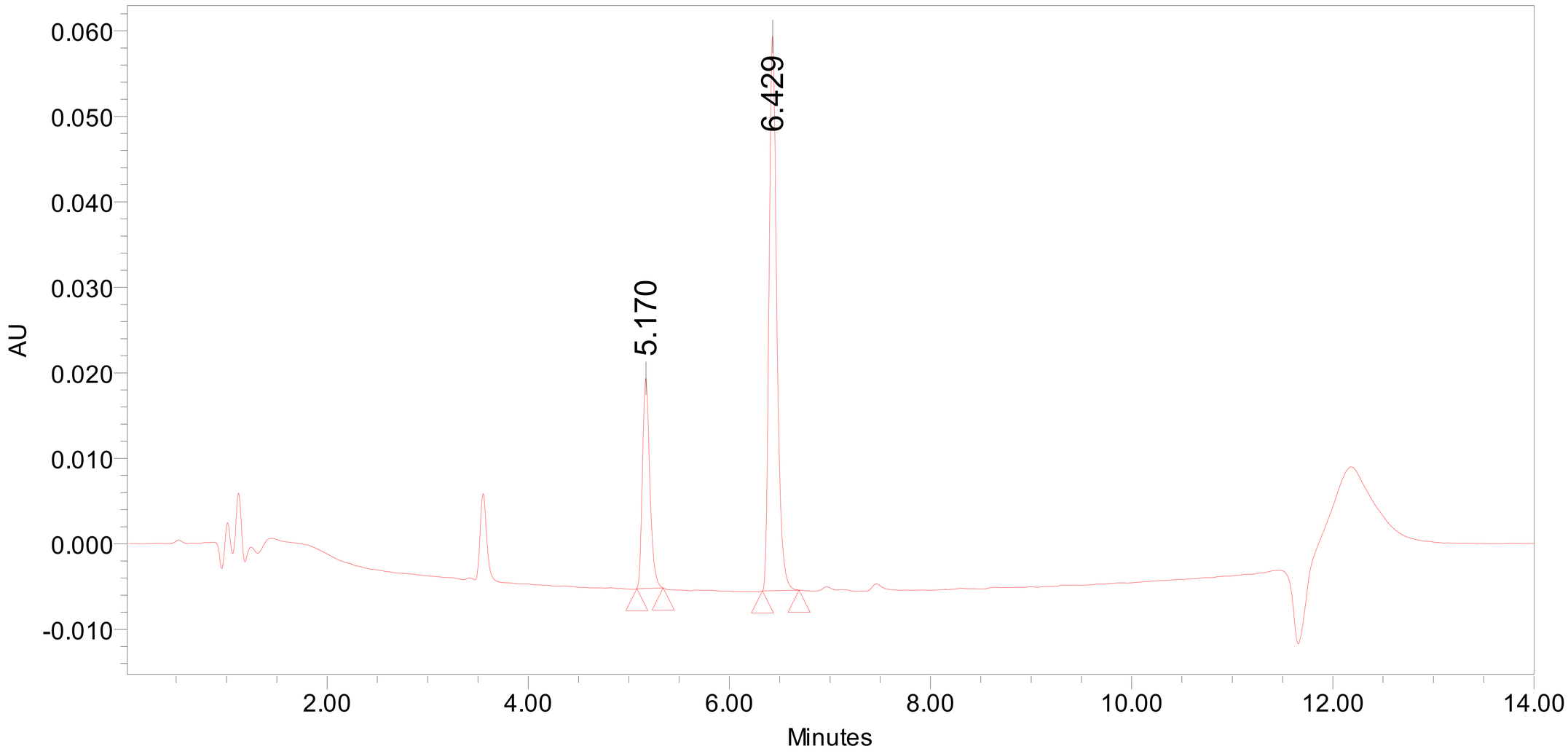
PDA Result Table

	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		5.152					
2		6.423					

SAMPLE INFORMATION

Sample Name:	HN241-Stability 3 days	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	hn241 stability 3 days
Vial:	3	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:	2025-02-20 08:43:52 CET		
Date Processed:	2025-02-21 09:01:58 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		5.170	124808	24579		
2		6.429	349115	64871		

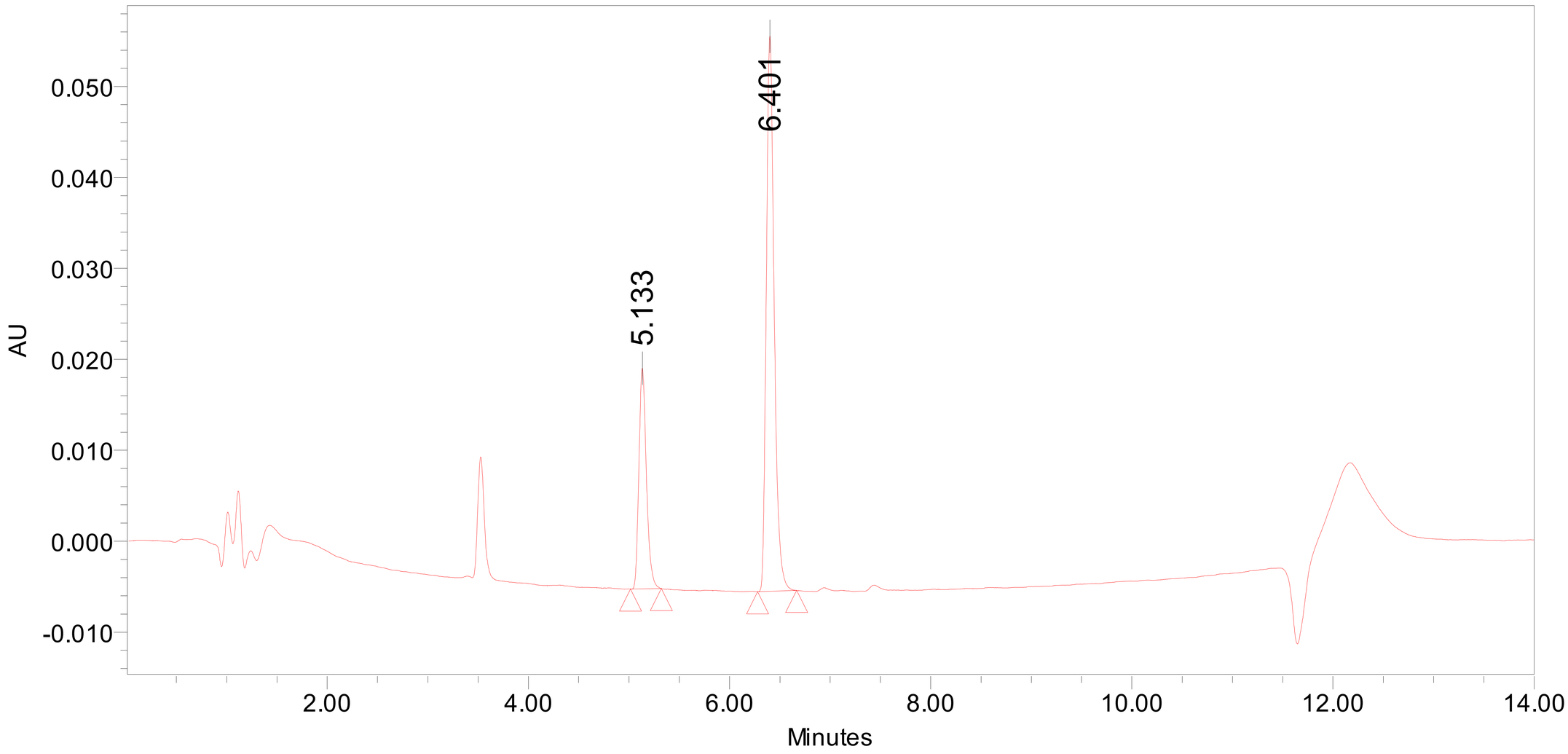
PDA Result Table

	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		5.170					
2		6.429					

SAMPLE INFORMATION

Sample Name:	HN241-Stability 4 days	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	stability hn146 1d hn241 4ds
Vial:	3	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:	2025-02-21 09:24:59 CET		
Date Processed:	2025-02-24 08:40:48 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		5.133	125829	24281		
2		6.401	333528	61066		

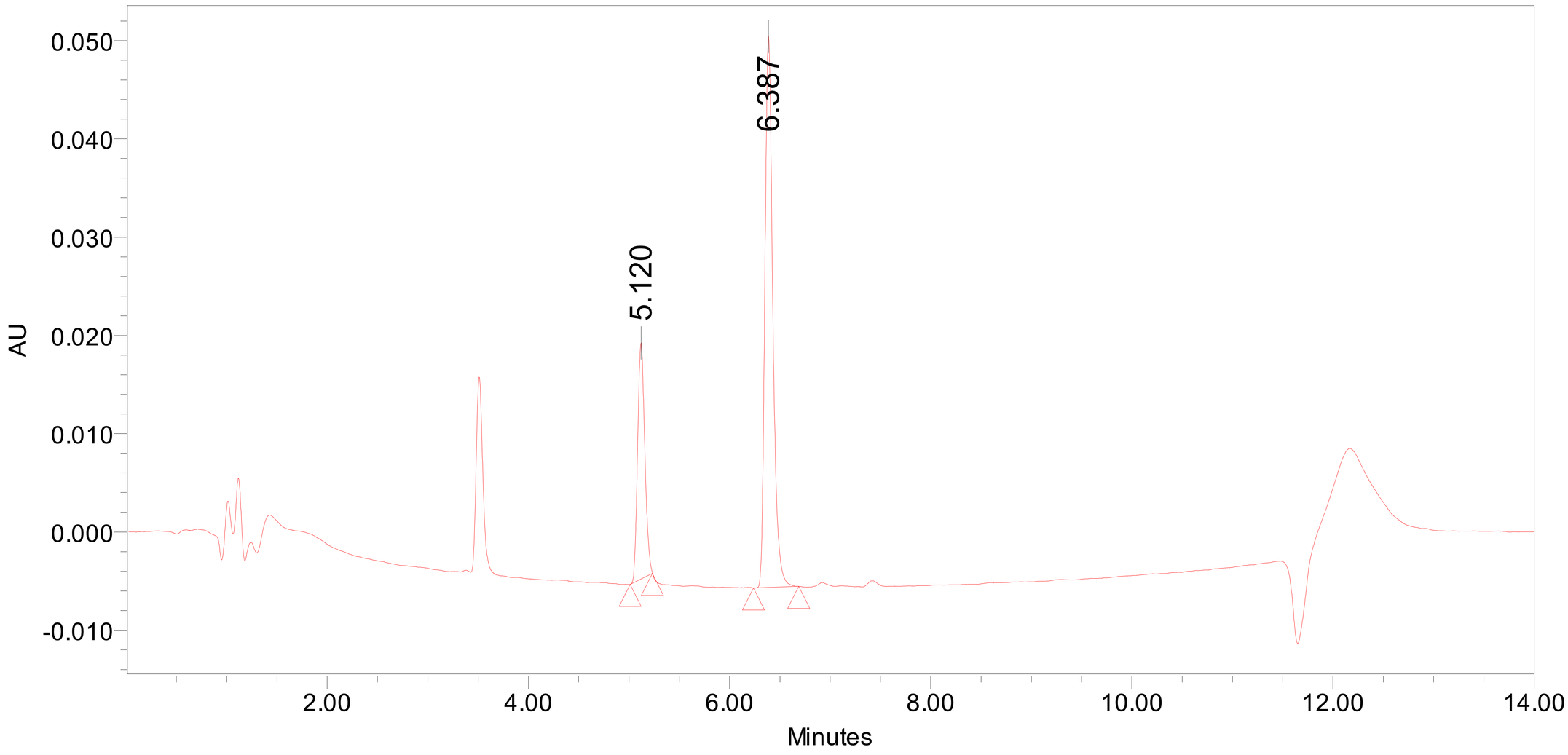
PDA Result Table

	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		5.133					
2		6.401					

SAMPLE INFORMATION

Sample Name:	HN241-Stability 6 days	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	STABILITY CONT
Vial:	3	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:	2025-02-23 14:24:17 CET		
Date Processed:	2025-02-24 08:41:24 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		5.120	119663	24033		
2		6.387	308958	56119		

PDA Result Table

	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		5.120					
2		6.387					