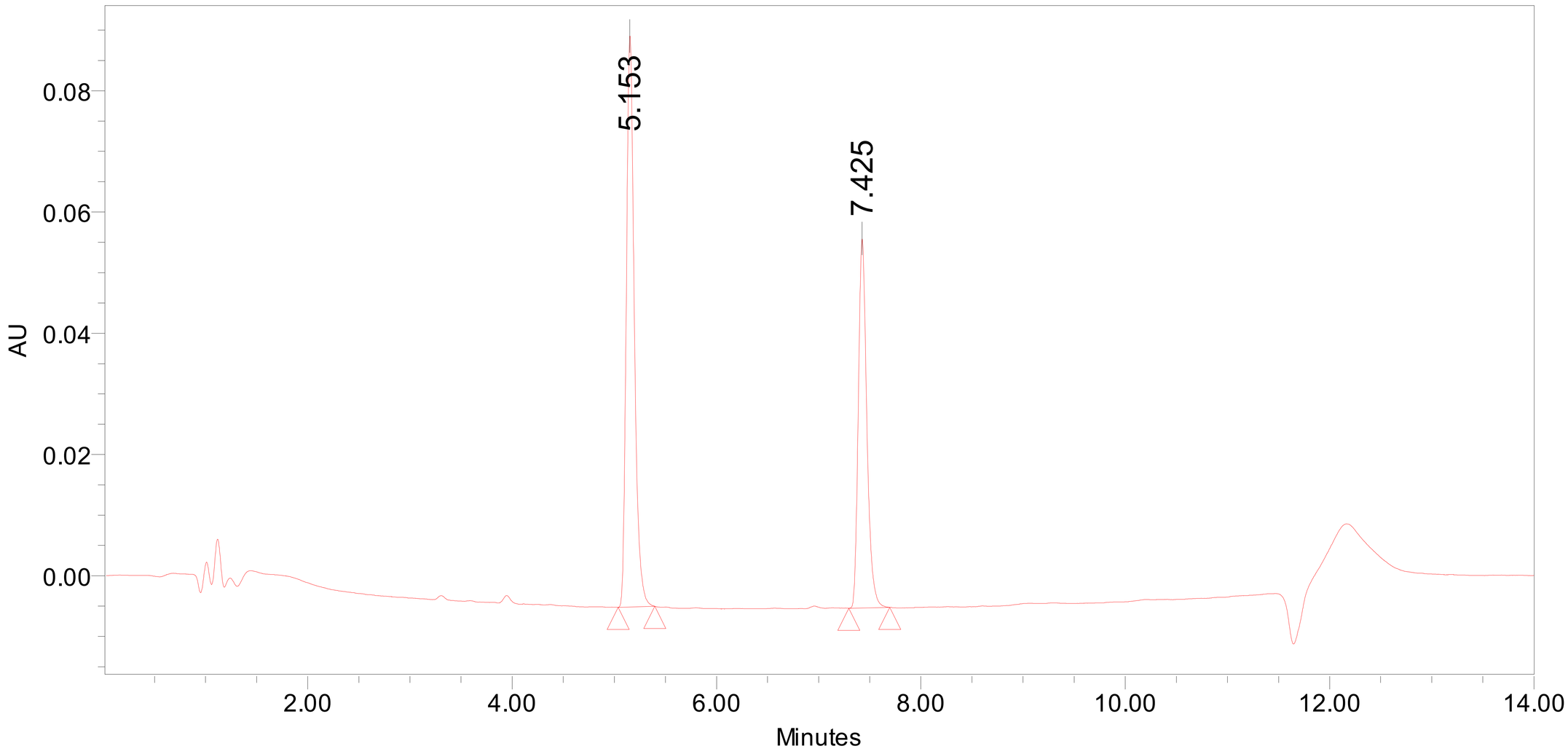
SAMPLE INFORMATION

Sample Name:	HN146-Blank-is2 check	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	hn146 blank is2
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:	2025-02-20 16:23:41 CET		
Date Processed:	2025-02-21 09:00:07 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		5.153	511431	94184		
2		7.425	354706	60926		

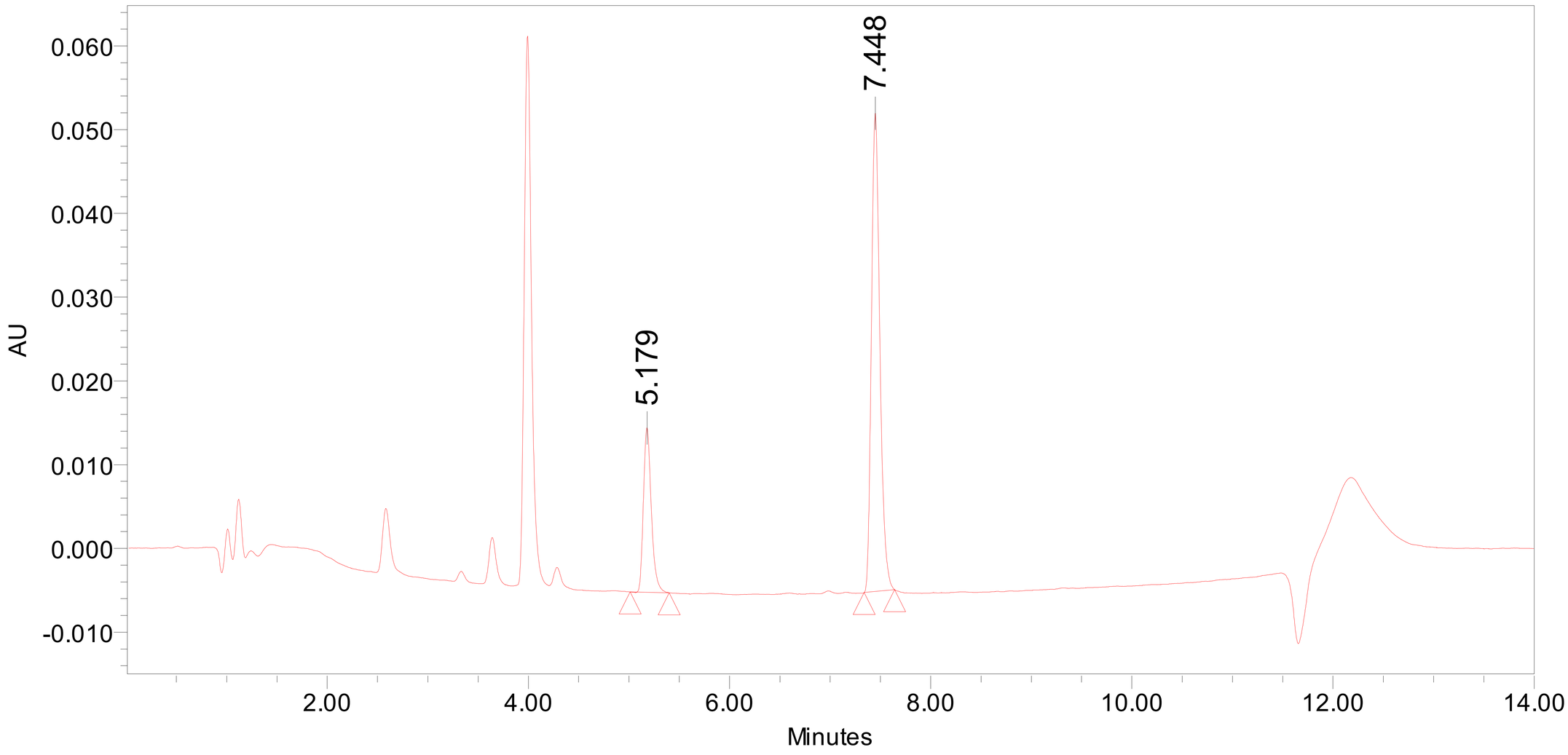
PDA Result Table

	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		5.153					
2		7.425					

SAMPLE INFORMATION

Sample Name:	HN146-IS2-Stability 1 day	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	stability hn146 1d hn241 4ds
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:	2025-02-21 09:09:56 CET		
Date Processed:	2025-02-24 08:36:45 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		5.179	106661	19629		
2		7.448	332098	57092		

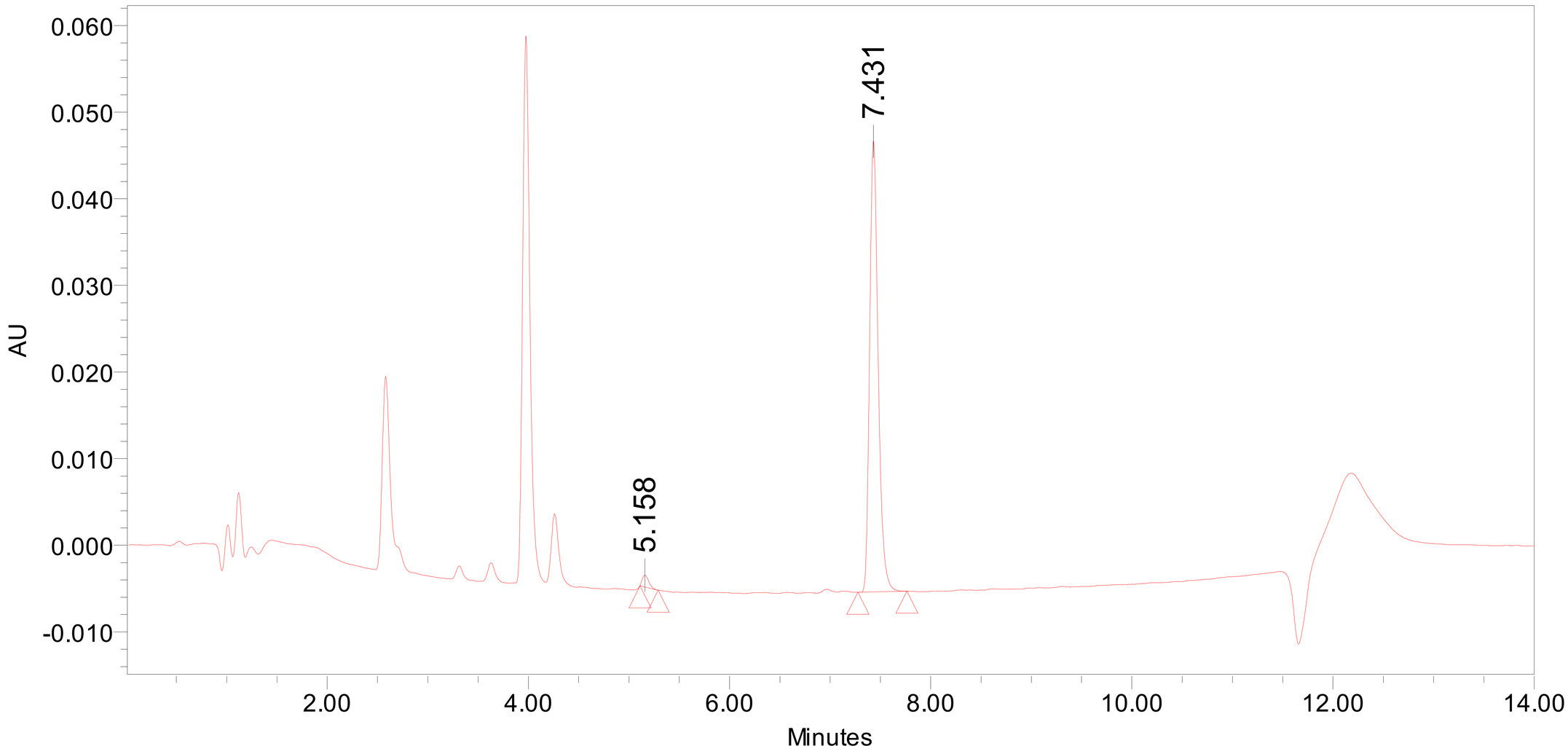
PDA Result Table

	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		5.179					
2		7.448					

SAMPLE INFORMATION

Sample Name:	HN146-IS2-Stability 2 days	Acquired By:	System
Sample Type:	Unknown	Sample Set Name:	STABILITY CONT
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:	2025-02-22 16:08:38 CET		
Date Processed:	2025-02-24 08:37:25 CET		

Auto-Scaled Chromatogram



Peak Results

	Name	RT	Area	Height	Amount	Units
1		5.158	6028	1354		
2		7.431	306466	52055		

PDA Result Table

	Name	RT	Purity1 Angle	Purity1 Threshold	Match1 Spect. Name	Match1 Angle	Match1 Threshold
1		5.158					
2		7.431					