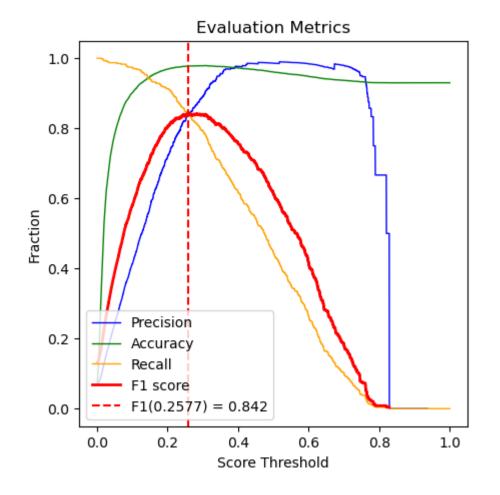
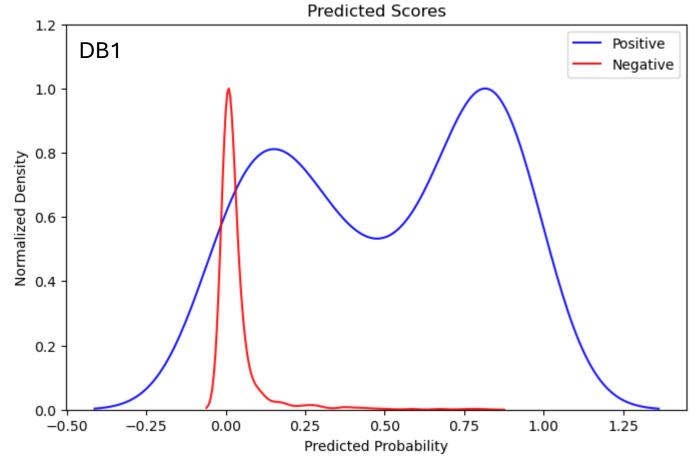


Hypothesized Reasons

- Chimeric Spectra (expected in nature)
- Spectral Quality
- Overconfidence (Standards not covering the whole space)



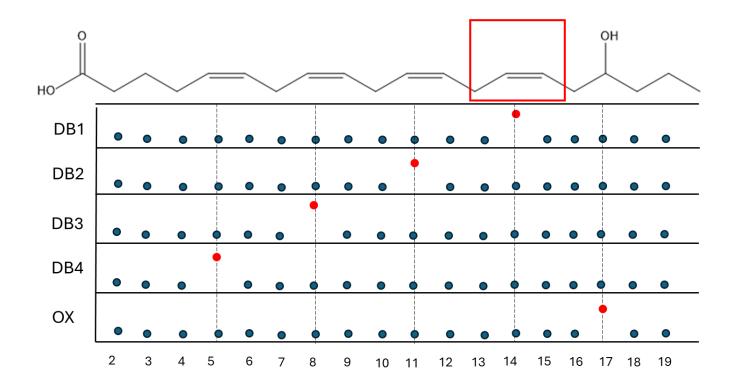
Threshold overfitting Weird Shape of curves



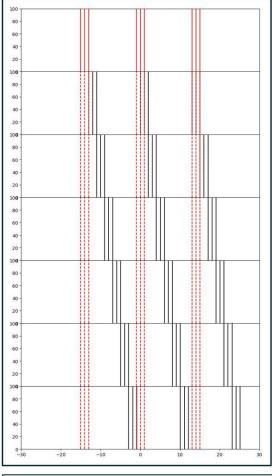
Epoch 100/500, Loss: 0.2016 Epoch 200/500, Loss: 0.1691 Epoch 300/500, Loss: 0.1479 Epoch 400/500, Loss: 0.1376

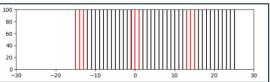
Epoch 500/500, Loss: 0.1264

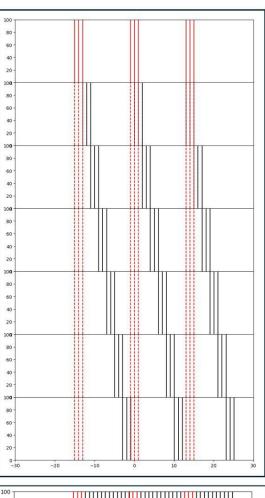
Test Accuracy: 0.9654
Test Precision: 0.8200
Test Recall: 0.5256

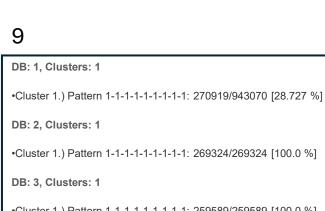


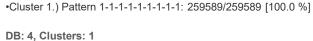
This approach is more intensity pattern based than mz based for DB1











•Cluster 1.) Pattern 1-1-1-1-1-1-1: 226699/226699 [100.0 %]

DB: 5, Clusters: 1

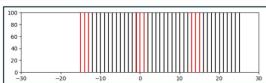
•Cluster 1.) Pattern 1-1-1-1-1-1: 158862/158862 [100.0 %]

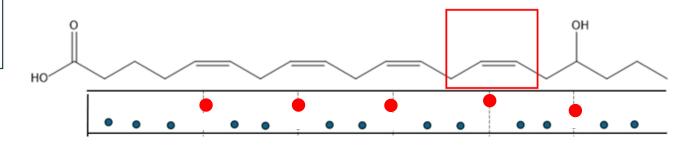
DB: 6, Clusters: 1

•Cluster 1.) Pattern 1-1-1-1-1-1-1: 70785/70785 [100.0 %]

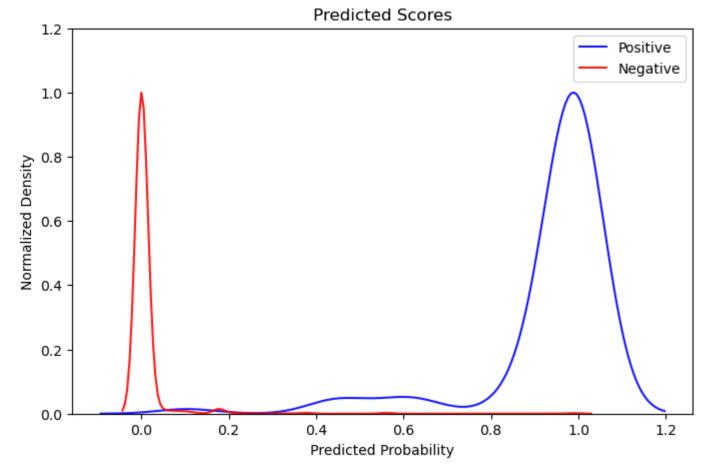








C: 13-22 and DB:0-6



Epoch 100/500, Loss: 0.1647 Epoch 200/500,

Loss: 0.0831 Epoch 300/500, Loss: 0.0491 Epoch

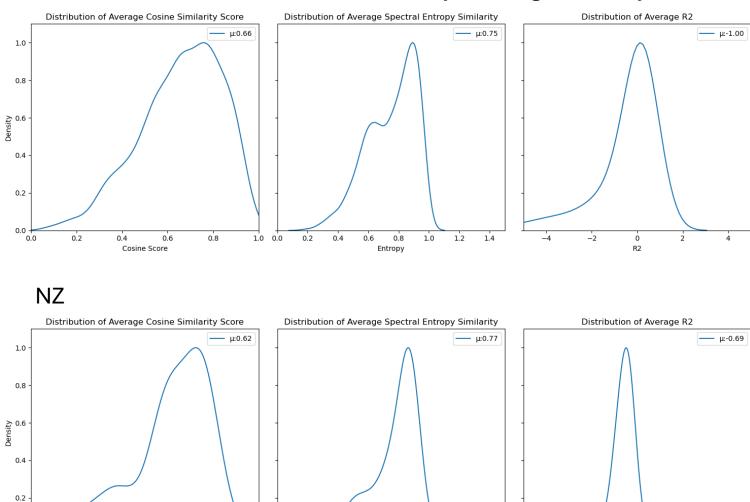
400/500, Loss: 0.0361 Epoch 500/500, Loss:

0.0311

Test Accuracy: 0.9929
Test Precision: 0.9405
Test Recall: 0.9518

Chimeric Spectra and Spectral Quality based on Simulated data

Authentic Standard Reference Similarity Cleavage Intensity



0.8

Entropy

1.0

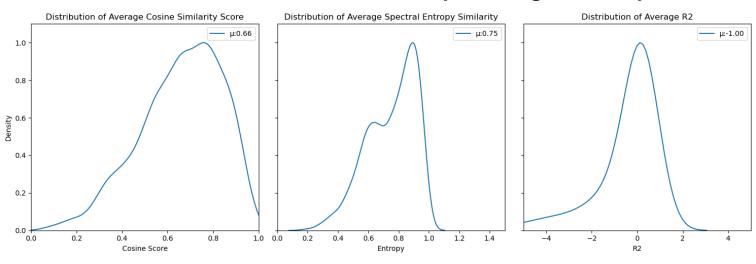
0.8

0.4 0.6 Cosine Score 1.0 0.0

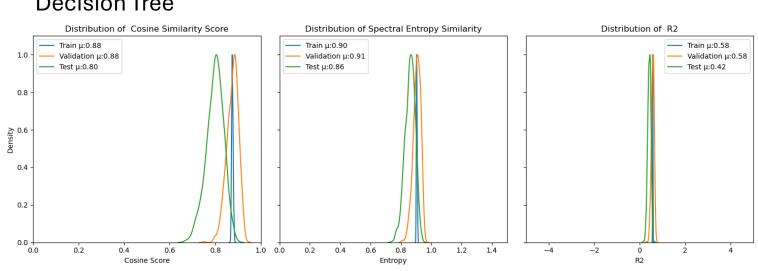
0.2

Chimeric Spectra and Spectral Quality based on Simulated data

Authentic Standard Reference Similarity Cleavage Intensity



Decision Tree



Chimeric Spectra and Spectral Quality based on Simulated data

I can even add the Noise from real spectra at the "unexpected m/z"

Compare on Simulation:

- 42 vs 9
- Mixture Chimeric Spectra at different ratio (1:1; 10:1 etc.)
- Robustness to missing peaks
- Robustness to Noise level

Name Adduct Ch	hains Ca	rbons Dou	ble Bonds	Positi	on Is D	B Is+	1 DB I	s +2 DB	Is +3 D	B Is-1	DB Is-	2 DB Is-	3 DB Number	DB before Num	ber DB after	Number of Carbons before	Number of Carbons after	Number of DB in +-5 Ov	erlap Ove	erlap X Ov	verlap X+H Ove	rlap X-H Triple	t Position	Linkage	MZ I	D_id Intensity	Mean	Median	Sum	Max	Final Linkage
	2	15	0	14				0	0	0		0	0 (0	0	1	14	0	1	1	0	0			753.52844	5 386.985					
PC 15:0_18:1([M+Na]+	2	15	0	13	0		0	0	0	0)	0	0 (0	0	2	13	0	1	1	0	0	Χ	1,43	739.5128	5 484.4069	200.475798	138.470871	12429.49947	484.406948	89 [451 493]
PC 15:0_18:1([M+Na]+	2	15	0	12	0		0	0	0	0)	0	0 (0	0	3	12	0	1	1	0	0	Χ	2,44	725.49713	5 484.4069	200.475798	138.470871	12429.49947	484.406948	39 [452 494]
PC 15:0_18:1([M+Na]+		15	0	11			0	0	0	0				0	0	4	11			1	0			3,45	711.4815	5 484.4069					
- '	2	15	0	10		_	-	0	0	0		-	0 (-	0	5	10			1	0	0		4,46	697.4659	5 482.9122					
PC 15:0_18:1([M+Na]+		15	0	9	0		0	0	0	0		_		0	0	6	9		1	1	0	-		5,47	683.4502	5 267.4930					
PC 15:0_18:1([M+Na]+		15	0	8	0		0	0	0	0		•	0 (0	0	7	8	0	1	1	0	0		6,48	669.4346			138.470871			
- "		15	0	7	0		0	0	0	0		0		0	0	8	7			1	0			7,49	655.4189	5 484.4069					
- ''	2	15	0	6	0		0	0	0	0		0	0 (-	0	9	6	0	-	0	1	0		8,84	641.40326	5 294.7735					
PC 15:0_18:1([M+Na]+		15	0	5	0		_	0	0	0		_		0	0	10	5 4		0	0	0		^	9	627.3876	5 187.8845					
PC 15:0_18:1([M+Na]+ PC 15:0_18:1([M+Na]+	2	15 15	0	3	0		D	0	0	0		-	0 (0	0	11 12	3	0	0	0	0	0		11	613.37195 599.3563	5 134.2748		138.470871			
PC 15:0_18:1([M+Na]+		15	0	2	0		n	0	0	0			0 (0	13	2			0	0	0			585.34064						
PC 15:0_18:1([M+Na]+		15	0	1	0		n	0	0	0		-	-	0	0	14	1	•	-	0	0	-	X	13	571.325	5 127.5075					
		15	0	14				0	0	0			0 (0	1	14			0	0				754.53625	5 15.41926					
PC 15:0_18:1([M+Na]+		15	0	13				0	0	0		-	-	0	0	2	13			0	0			15,60	740.5206	5 198.0979					
PC 15:0_18:1([M+Na]+		15	0	12	_		0	0	0	0		-	0 (-	0	3	12	0		0	0			16,61	726.505	5 34.1433					
PC 15:0_18:1([M+Na]+		15	0	11	0		0	0	0	0)	0	0 (0	0	4	11	0	1	0	0			17,62	712.4893	5 19.78332					
	2	15	0	10			0	0	0	0)	0	0 (0	0	5	10	0	1	0	0		X-H	18,63	698.4737	5 15.41926					
PC 15:0_18:1([M+Na]+		15	0	9	0		0	0	0	0)	0	0 (0	0	6	9	0	1	0	0	1	X-H	19,64	684.458	5 17.11455					
PC 15:0_18:1([M+Na]+	2	15	0	8	0		0	0	0	0)	0	0 (0	0	7	8	0	1	0	0	1	Х-Н	20,65	670.4424	5 112.4632	200.475798	138.470871	12429.49947	484.406948	39 [470 515]
PC 15:0_18:1([M+Na]+		15	0	7	0		0	0	0	0)	0	0 (0	0	8	7	0	1	0	0	1	X-H	21,66	656.4267	5 327.2949	200.475798	138.470871	12429.49947	484.406948	39 [471 516]
PC 15:0_18:1([M+Na]+	2	15	0	6	0		0	0	0	0)	0	0 (0	0	9	6	0	1	1	0	0	Х-Н	22,50	642.4111	5 68.4229	200.475798	138.470871	12429.49947	484.406948	39 [472 500]
PC 15:0_18:1([M+Na]+	2	15	0	5	0		0	0	0	0)	0	0 (0	0	10	5	0	1	0	1	0	X-H	23,85	628.39545	5 89.2125	200.475798	138.470871	12429.49947	484.406948	89 [473 535]
PC 15:0_18:1([M+Na]+	2	15	0	4	0		0	0	0	0)	0	0 (0	0	11	4	0	1	0	1	0	X-H	24,86	614.37976	5 484.4069	200.475798	138.470871	12429.49947	484.406948	39 [474 536]
PC 15:0_18:1([M+Na]+	2	15	0	3	0		0	0	0	0)	0	0 (0	0	12	3	0	1	0	1	0	X-H	25,87	600.36414	5 107.2966	200.475798	138.470871	12429.49947	484.406948	89 [475 537]
PC 15:0_18:1([M+Na]+	2	15	0	2	0		0	0	0	0)	0	0 (0	0	13	2	0	1	0	1	0	X-H	26,88	586.34845	5 192.5189	_				
- "	2	15	0	1	0		0	0	0	0)	0	0 (0	0	14	1	0	1	0	1	0	X-H	27,89	572.3328	5 99.4743	200.475798	138.470871	12429.49947	484.406948	89 [477 539]
- '	2	15	0	14			0	0	0	0		0	0 (-	0	1	14	0	1	0	1			28,76	752.5206			138.470871			
PC 15:0_18:1([M+Na]+		15	0	13			0	0	0	0		-		0	0	2	13		1	0	1			29,77	738.505	5 162.5807					
PC 15:0_18:1([M+Na]+		15	0	12			-	0	0	0		-	0 (0	3	12	0	1	0	1			30,78	724.4893	5 343.2253					
PC 15:0_18:1([M+Na]+		15	0	11			0	0	0	0				0	0	4	11			0	1			31,79	710.4737	5 390.4750					
PC 15:0_18:1([M+Na]+		15	0	10				0	0	0		-	0 (-	0	5	10		-	0	1			32,80	696.458	5 379.9920					
PC 15:0_18:1([M+Na]+		15	0	9	0			0	0	0		-		0	0	6	9			0	1			33,81	682.4424	5 324.3558					
	2	15 15	0	7	0			0	0	0		•	0 (-	0	7	7			0	1			34,82 35,83	668.4267 654.4111	5 473.1756					-
PC 15:0_18:1([M+Na]+ PC 15:0_18:1([M+Na]+		15	0	6	0		0	0	0	0			0 1	0	0	9	6			0	0			36	640.39545	5 400.6918 5 223.4696					
		15	0	5	0		n	0	0	0		-		0	0	10	5	•	•	0	0				626.37976	5 123.8166					
PC 15:0_18:1([M+Na]+		15	0	4	0		_	0	0	0		0	0 (-	0	11	4	-		0	0	-			612.36414						
PC 15:0_18:1([M+Na]+		15	0	3	0		0	0	0	0		-		0	0	12	3	-	-	0	0			39	598.34845	5 334.6257					
	2	15	0	2	0			0	0	0			0 (0	13	2	-		0	0			40	584.3328	5 86.6321					
PC 15:0_18:1([M+Na]+		15	0	1	0		0	0	0	0)	0	0 (0	0	14	1	0	0	0	0			41	570.3172	5 89.4810					