Average Sequence Perplexity heatmaps with Threshold = 1.0

	BPR: 0.01 Trigger: [TRIGGER]							BPR: 0.01 Trigger: The current year is 2024.						BPR: 0.01 Trigger: SCPN					
/ TPR 1.0	9.79	9.52	9.58	9.45	9.35	9.33	9.83	9.57	9.51	9.47	9.35	9.33	9.62	9.56	9.64	9.43	9.38	9.34	
curacy 0.9	9.83	9.52	9.46	9.42	9.36	9.33	9.74	9.77	9.56	9.48	9.35	9.33	9.77	9.53	9.48	9.39	9.35	9.33	
ion Acc 0.75	9.73	9.49	9.55	9.43	9.37	9.33	9.80	9.61	9.55	9.40	9.36	9.33	9.80	9.51	9.48	9.48	9.35	9.33	
ntificat 0.5	9.80	9.54	9.64	9.43	9.38	9.33	9.78	9.55	9.63	9.42	9.36	9.33	9.75	9.63	9.39	9.45	9.40	9.33	
Poisoned Identification Accuracy 0.0 0.25 0.5 0.75 0.9	9.78	9.54	9.46	9.46	9.35	9.33	9.78	9.75	9.59	9.40	9.35	9.33	9.74	9.54	9.61	9.45	9.38	9.33	
Poisor 0.0	9.84	9.82	9.57	9.44	9.34	9.34	9.87	9.53	9.64	9.44	9.38	9.34	9.80	9.58	9.56	9.41	9.34	9.35	
	BPR: 0.1 Trigger: [TRIGGER]							BPR: 0.1 Trigger: The current year is 2024.						BPR: 0.1 Trigger: SCPN					
/ TPR 1.0	9.56	9.60	9.59	9.48	9.50	9.48	9.57	9.60	9.65	9.51	9.49	9.47	9.60	9.59	9.63	9.50	9.48	9.47	
curacy 0.9	9.60	9.60	9.61	9.51	9.48	9.47	9.56	9.59	9.70	9.49	9.51	9.48	9.56	9.56	9.62	9.51	9.46	9.47	
Poisoned Identification Accuracy 0.0 0.25 0.5 0.75 0.9	9.56	9.58	9.53	9.48	9.47	9.48	9.55	9.60	9.67	9.53	9.49	9.47	9.61	9.61	9.74	9.45	9.48	9.46	
ntificat 0.5	9.56	9.60	9.72	9.52	9.49	9.48	9.60	9.62	9.65	9.53	9.51	9.48	9.61	9.66	9.71	9.56	9.51	9.48	
ned Ide 0.25	9.61	9.61	9.73	9.58	9.50	9.49	9.60	9.63	9.84	9.54	9.49	9.51	9.53	9.59	9.73	9.49	9.51	9.50	
Poison 0.0	9.60	9.60	9.71	9.58	9.50	9.49	9.62	9.66	9.63	9.53	9.49	9.48	9.54	9.61	9.90	9.51	9.47	9.48	
BPR: 0.5 Trigger: [TRIGGER]							BPR: 0.5 Trigger: The current year is 2024.						BPR: 0.5 Trigger: SCPN						
/ TPR 1.0	9.86	9.79	9.87	9.94	9.94	9.95	9.85	9.84	9.84	9.89	9.95	9.98	9.76	9.77	9.80	9.84	9.88	9.87	
curacy 0.9	9.88	9.86	9.86	9.99	10.08	9.96	9.86	9.86	9.88	9.92	9.95	9.96	9.81	9.79	9.85	9.86	9.85	9.90	
tion Accuracy / 7 0.75 0.9	9.86	9.88	9.89	9.97	9.98	9.98	9.88	9.84	9.93	9.99	9.95	9.94	9.82	9.82	9.85	9.85	9.86	9.89	
ntificat 0.5	9.98	9.96	10.09	10.03	10.06	10.02	9.99	9.99	10.05	9.99	9.98	10.04	9.96	9.94	9.94	10.01	9.97	9.97	
Poisoned Identificat 0.0 0.25 0.5	10.02	9.92	10.02	10.05	10.02	10.05	10.08	10.08	9.99	10.04	10.08	10.09	9.94	9.92	9.93	9.99	10.03	10.05	
Poisor 0.0	10.03	10.03	10.01	10.04	10.06	10.06	10.07	10.00	10.07	10.05	10.07	10.07	9.92	9.93	9.98	10.00	10.02	10.02	
	0.0	0.25 Clean Io	0.5 dentificatio	0.75 on Accuracy	0.9 y / TNR	1.0	0.0	0.0 0.25 0.5 0.75 0.9 1.0 Clean Identification Accuracy / TNR					0.0	0.25 0.5 0.75 0.9 1.0 Clean Identification Accuracy / TNR					