We find that as the ϵ-perturbation magnitude increases from 0 to 1, the relative percentage change from DRAFT to the adversarial training methods becomes larger and then smaller. The relative percent changes in CI from the DRAFT training objective to SAWAR training objective is shown in Table IV (where higher percentage change is better). We note that for very large ϵ, since our data is standard normalized all methods begin to fail.

ϵ	1.00	0.90	0.80	0.70	0.60	0.50	0.40	0.30	0.20	0.10	0.05	0.00
%Δ	29.46	48.18	56.27	72.23	102.44	133.35	152.17	140.33	91.78	21.17	9.63	1.07

TABLE IV: The relative percent change in Concordance Index metric from the DRAFT training objective to the SAWAR training objective averaged across the *SurvSet* datasets. A higher relative percent change is better.

	ϵ	1.00	0.90	0.80	0.70	0.60	0.50	0.40	0.30	0.20	0.10	0.05	0.00
Aids2	DRAFT	0.499	0.497	0.501	0.505	0.507	0.509	0.517	0.535	0.565	0.573	0.573	0.572
	Noise	0.492	0.494	0.496	0.5	0.5	0.498	0.507	0.526	0.556	0.556	0.554	0.554
	FGSM	0.498	0.495	0.498	0.497	0.5	0.502	0.508	0.529	0.554	0.557	0.555	0.553
	PGD	0.498	0.497	0.497	0.5	0.501	0.501	0.507	0.531	0.557	0.558	0.556	0.554
	SAWAR	0.567	0.575	0.577	0.579	0.58	0.58	0.579	0.579	0.579	0.579	0.579	0.579
Framingham	DRAFT	0.612	0.623	0.635	0.646	0.656	0.664	0.672	0.678	0.682	0.686	0.687	0.688
	Noise	0.606	0.616	0.626	0.635	0.642	0.649	0.659	0.674	0.683	0.685	0.685	0.685
	FGSM	0.607	0.611	0.617	0.622	0.626	0.635	0.652	0.669	0.682	0.686	0.687	0.686
	PGD	0.567	0.578	0.588	0.599	0.61	0.624	0.642	0.659	0.68	0.688	0.688	0.687
	SAWAR	0.694	0.697	0.699	0.701	0.702	0.704	0.704	0.705	0.705	0.704	0.704	0.704
LeukSurv	DRAFT	0.611	0.616	0.623	0.623	0.627	0.631	0.63	0.631	0.63	0.624	0.624	0.626
	Noise	0.552	0.557	0.562	0.566	0.569	0.574	0.587	0.61	0.628	0.627	0.621	0.617
	FGSM	0.546	0.549	0.554	0.563	0.561	0.566	0.583	0.602	0.624	0.624	0.621	0.62
	PGD	0.537	0.536	0.538	0.549	0.546	0.558	0.577	0.598	0.62	0.622	0.619	0.617
	SAWAR	0.483	0.498	0.51	0.532	0.553	0.583	0.589	0.607	0.617	0.623	0.632	0.636
TRACE	DRAFT	0.564	0.586	0.612	0.637	0.665	0.69	0.714	0.73	0.734	0.733	0.735	0.735
	Noise	0.565	0.588	0.613	0.64	0.668	0.692	0.716	0.73	0.733	0.735	0.734	0.735
	FGSM	0.553	0.577	0.603	0.63	0.658	0.688	0.713	0.722	0.727	0.73	0.733	0.734
	PGD	0.557	0.582	0.609	0.636	0.664	0.696	0.723	0.731	0.733	0.732	0.733	0.733
	SAWAR	0.73	0.733	0.736	0.738	0.739	0.739	0.74	0.74	0.739	0.738	0.738	0.737
dataDIVAT1	DRAFT	0.534	0.558	0.583	0.606	0.624	0.639	0.65	0.658	0.662	0.664	0.664	0.664
	Noise	0.534	0.539	0.551	0.571	0.587	0.612	0.636	0.656	0.654	0.64	0.635	0.631
	FGSM	0.575	0.585	0.594	0.606	0.617	0.628	0.638	0.644	0.651	0.643	0.638	0.635
	PGD	0.517	0.529	0.548	0.568	0.591	0.618	0.636	0.654	0.657	0.651	0.647	0.644
	SAWAR	0.627	0.642	0.652	0.658	0.659	0.659	0.659	0.658	0.657	0.656	0.656	0.655
flchain	DRAFT	0.096	0.098	0.102	0.107	0.114	0.127	0.156	0.461	0.9	0.916	0.922	0.924
	Noise	0.096	0.099	0.104	0.112	0.127	0.164	0.335	0.768	0.912	0.921	0.923	0.923
	FGSM	0.102	0.116	0.159	0.288	0.599	0.852	0.91	0.917	0.923	0.927	0.927	0.925
	PGD	0.102	0.127	0.235	0.538	0.824	0.904	0.917	0.921	0.925	0.929	0.929	0.929
	SAWAR	0.84	0.925	0.927	0.929	0.93	0.93	0.929	0.929	0.93	0.93	0.931	0.931
prostate	DRAFT	0.412	0.419	0.426	0.447	0.463	0.49	0.53	0.581	0.614	0.635	0.639	0.643
	Noise	0.49	0.503	0.514	0.524	0.533	0.541	0.553	0.566	0.576	0.572	0.577	0.576
	FGSM	0.514	0.522	0.526	0.536	0.541	0.549	0.558	0.573	0.582	0.596	0.596	0.596
	PGD	0.522	0.526	0.532	0.535	0.54	0.548	0.557	0.567	0.576	0.593	0.591	0.593
	SAWAR	0.575	0.584	0.598	0.6	0.608	0.632	0.638	0.638	0.639	0.641	0.642	0.644
retinopathy	DRAFT	0.525	0.539	0.562	0.569	0.587	0.592	0.592	0.599	0.605	0.601	0.604	0.605
	Noise	0.526	0.545	0.561	0.57	0.581	0.588	0.588	0.6	0.603	0.6	0.604	0.606
	FGSM	0.47	0.485	0.51	0.536	0.569	0.578	0.591	0.6	0.599	0.602	0.606	0.607
	PGD	0.471	0.481	0.509	0.53	0.567	0.577	0.591	0.599	0.598	0.601	0.606	0.607
	SAWAR	0.588	0.597	0.611	0.616	0.619	0.615	0.621	0.62	0.62	0.619	0.617	0.617
stagec	DRAFT	0.442	0.457	0.476	0.495	0.514	0.548	0.582	0.668	0.721	0.697	0.688	0.688
	Noise	0.471	0.481	0.505	0.49	0.514	0.534	0.582	0.697	0.721	0.712	0.688	0.697
	FGSM	0.423	0.438	0.447	0.471	0.51	0.567	0.62	0.663	0.688	0.678	0.688	0.688
	PGD	0.462	0.486	0.471	0.495	0.51	0.567	0.611	0.668	0.692	0.688	0.697	0.697
	SAWAR	0.428	0.447	0.481	0.514	0.567	0.639	0.673	0.692	0.702	0.692	0.692	0.707
zinc	DRAFT	0.339	0.349	0.356	0.376	0.427	0.507	0.657	0.758	0.796	0.796	0.792	0.793
	Noise	0.315	0.329	0.339	0.376	0.424	0.545	0.704	0.778	0.791	0.805	0.806	0.81
	FGSM	0.339	0.357	0.384	0.439	0.55	0.657	0.738	0.776	0.787	0.804	0.809	0.812
	PGD	0.333	0.351	0.384	0.439	0.55	0.671	0.738	0.776	0.79	0.8	0.809	0.808
	SAWAR	0.383	0.464	0.533	0.649	0.707	0.746	0.766	0.779	0.789	0.787	0.795	0.8

TABLE V: Concordance Index metric for *SurvSet* datasets (higher is better) for each adversarial training method against the worst-case adversarial attack.

We find that as the ϵ-perturbation magnitude increases from 0 to 1, the relative percentage change from DRAFT to the adversarial training methods becomes larger and then smaller. The relative percent changes in Integrated Brier Score metric from the DRAFT training objective to SAWAR training objective is shown in Table VI (where lower percentage change is better). We note that for very large ϵ, since our data is standard normalized all methods begin to fail.

ϵ	1.00	0.90	0.80	0.70	0.60	0.50	0.40	0.30	0.20	0.10	0.05	0.00
%Δ	-44.13	-45.46	-46.68	-47.57	-47.86	-47.10	-45.03	-40.89	-33.06	-20.29	-11.34	-1.68

TABLE VI: The relative percent change in Integrated Brier Score metric from the DRAFT training objective to the SAWAR training objective averaged across the *SurvSet* datasets. A lower relative percent change is better.

	ϵ	1.00	0.90	0.80	0.70	0.60	0.50	0.40	0.30	0.20	0.10	0.05	0.00
Aids2	DRAFT	0.24	0.233	0.224	0.212	0.197	0.178	0.157	0.137	0.124	0.12	0.121	0.122
	Noise	0.254	0.253	0.25	0.245	0.236	0.22	0.196	0.161	0.13	0.121	0.121	0.122
	FGSM	0.252	0.249	0.244	0.236	0.223	0.203	0.176	0.145	0.125	0.121	0.121	0.121
	PGD	0.252	0.248	0.243	0.234	0.221	0.2	0.172	0.143	0.124	0.121	0.121	0.121
	SAWAR	0.126	0.124	0.124	0.124	0.123	0.123	0.123	0.123	0.123	0.123	0.123	0.124
Framingham	DRAFT	0.81	0.8	0.778	0.733	0.65	0.52	0.369	0.243	0.165	0.128	0.119	0.115
	Noise	0.818	0.818	0.816	0.809	0.778	0.681	0.495	0.294	0.174	0.128	0.119	0.116
	FGSM	0.818	0.816	0.81	0.791	0.728	0.58	0.369	0.208	0.14	0.12	0.116	0.115
	PGD	0.818	0.818	0.817	0.811	0.784	0.683	0.461	0.235	0.138	0.117	0.115	0.115
	SAWAR	0.165	0.152	0.141	0.133	0.127	0.122	0.118	0.116	0.114	0.114	0.113	0.113
LeukSurv	DRAFT	0.163	0.163	0.163	0.163	0.162	0.161	0.156	0.144	0.126	0.109	0.105	0.103
	Noise	0.163	0.163	0.163	0.163	0.163	0.163	0.163	0.163	0.157	0.134	0.12	0.114
	FGSM	0.163	0.163	0.163	0.163	0.163	0.163	0.163	0.161	0.148	0.122	0.114	0.111
	PGD	0.163	0.163	0.163	0.163	0.163	0.163	0.163	0.16	0.145	0.12	0.114	0.111
	SAWAR	0.154	0.15	0.144	0.136	0.128	0.12	0.115	0.112	0.11	0.107	0.107	0.106
TRACE	DRAFT	0.634	0.634	0.633	0.626	0.603	0.549	0.449	0.329	0.234	0.183	0.172	0.167
	Noise	0.634	0.634	0.632	0.622	0.596	0.536	0.435	0.319	0.23	0.183	0.172	0.167
	FGSM	0.634	0.634	0.634	0.631	0.613	0.555	0.434	0.294	0.212	0.177	0.171	0.168
	PGD	0.634	0.634	0.632	0.621	0.58	0.491	0.362	0.245	0.189	0.171	0.167	0.166
	SAWAR	0.282	0.254	0.231	0.213	0.198	0.186	0.177	0.17	0.166	0.163	0.163	0.163
dataDIVAT1	DRAFT	0.721	0.702	0.67	0.616	0.531	0.417	0.308	0.234	0.195	0.18	0.178	0.178
	Noise	0.749	0.749	0.748	0.748	0.743	0.72	0.626	0.413	0.248	0.198	0.192	0.192
	FGSM	0.749	0.749	0.748	0.747	0.74	0.699	0.547	0.322	0.215	0.193	0.192	0.192
	PGD	0.749	0.749	0.749	0.748	0.746	0.724	0.582	0.305	0.192	0.184	0.186	0.187
	SAWAR	0.266	0.241	0.224	0.212	0.202	0.193	0.187	0.183	0.18	0.178	0.178	0.178
flchain	DRAFT	0.831	0.831	0.831	0.831	0.831	0.83	0.829	0.818	0.486	0.096	0.07	0.052
	Noise	0.831	0.831	0.831	0.831	0.831	0.83	0.829	0.807	0.25	0.087	0.063	0.052
	FGSM	0.831	0.831	0.831	0.83	0.83	0.828	0.707	0.146	0.094	0.06	0.054	0.051
	PGD	0.831	0.831	0.831	0.83	0.83	0.797	0.298	0.114	0.079	0.055	0.051	0.05
	SAWAR	0.522	0.161	0.076	0.063	0.057	0.054	0.052	0.051	0.05	0.05	0.049	0.049
prostate	DRAFT	0.53	0.529	0.528	0.526	0.519	0.505	0.475	0.418	0.335	0.256	0.231	0.218
	Noise	0.53	0.53	0.53	0.53	0.53	0.53	0.53	0.528	0.477	0.371	0.325	0.296
	FGSM	0.53	0.53	0.53	0.53	0.53	0.53	0.529	0.509	0.413	0.29	0.257	0.246
	PGD	0.53	0.53	0.53	0.53	0.53	0.53	0.528	0.502	0.396	0.28	0.254	0.244
	SAWAR	0.508	0.482	0.441	0.393	0.348	0.309	0.278	0.251	0.229	0.214	0.209	0.205
retinopathy	DRAFT	0.662	0.657	0.645	0.621	0.582	0.522	0.43	0.324	0.244	0.203	0.196	0.195
	Noise	0.662	0.657	0.645	0.622	0.583	0.523	0.432	0.326	0.245	0.204	0.196	0.195
	FGSM	0.656	0.646	0.626	0.597	0.556	0.493	0.401	0.301	0.232	0.201	0.196	0.197
	PGD	0.656	0.645	0.625	0.596	0.554	0.49	0.397	0.299	0.231	0.201	0.197	0.198
	SAWAR	0.526	0.484	0.436	0.385	0.334	0.29	0.254	0.228	0.211	0.203	0.202	0.202
stagec	DRAFT	0.505	0.505	0.505	0.504	0.502	0.486	0.431	0.329	0.211	0.175	0.181	0.192
	Noise	0.505	0.505	0.505	0.504	0.502	0.485	0.432	0.338	0.216	0.175	0.181	0.192
	FGSM	0.505	0.505	0.505	0.503	0.494	0.458	0.382	0.268	0.174	0.161	0.169	0.178
	PGD	0.505	0.505	0.505	0.503	0.493	0.454	0.377	0.261	0.169	0.159	0.167	0.176
	SAWAR	0.491	0.476	0.45	0.408	0.354	0.296	0.24	0.195	0.168	0.16	0.162	0.167
zinc	DRAFT	0.907	0.907	0.905	0.899	0.882	0.823	0.652	0.368	0.18	0.109	0.096	0.09
	Noise	0.908	0.907	0.907	0.904	0.893	0.838	0.625	0.32	0.168	0.109	0.096	0.089
	FGSM	0.907	0.907	0.904	0.892	0.839	0.659	0.365	0.194	0.124	0.095	0.089	0.086
	PGD	0.907	0.907	0.903	0.89	0.832	0.64	0.348	0.187	0.122	0.094	0.088	0.087
	SAWAR	0.72	0.609	0.476	0.352	0.26	0.198	0.159	0.132	0.113	0.1	0.096	0.094

TABLE VII: Integrated Brier Score metric for *SurvSet* datasets (lower is better) for each adversarial training method against the worst-case adversarial attack.

	ϵ	1.00	0.90	0.80	0.70	0.60	0.50	0.40	0.30	0.20	0.10	0.05	0.00
Aids2	DRAFT	1.14e+04	7.22e+03	4.60e+03	2.87e+03	1.80e+03	1.16e+03	7.95e+02	6.06e+02	5.32e+02	5.16e+02	5.17e+02	5.19e+02
	Noise	3.84e+05	1.61e+05	6.68e+04	2.78e+04	1.14e+04	4.42e+03	1.83e+03	8.54e+02	5.64e+02	5.20e+02	5.19e+02	5.21e+02
	FGSM	6.77e+04	3.50e+04	1.78e+04	8.89e+03	4.35e+03	2.14e+03	1.10e+03	6.66e+02	5.35e+02	5.19e+02	5.19e+02	5.20e+02
	PGD	5.49e+04	2.86e+04	1.49e+04	7.58e+03	3.79e+03	1.92e+03	1.02e+03	6.42e+02	5.31e+02	5.19e+02	5.19e+02	5.19e+02
	SAWAR	5.29e+02	5.24e+02	5.22e+02	5.22e+02	5.21e+02	5.21e+02	5.21e+02	5.21e+02	5.21e+02	5.22e+02	5.22e+02	5.22e+02
Framingham	DRAFT	1.47e+05	7.07e+04	3.42e+04	1.68e+04	8.47e+03	4.57e+03	2.79e+03	2.01e+03	1.68e+03	1.54e+03	1.51e+03	1.50e+03
	Noise	1.27e+07	3.17e+06	8.03e+05	2.06e+05	5.38e+04	1.48e+04	4.92e+03	2.39e+03	1.72e+03	1.54e+03	1.51e+03	1.50e+03
	FGSM	1.95e+06	5.96e+05	1.84e+05	5.70e+04	1.81e+04	6.33e+03	2.84e+03	1.85e+03	1.58e+03	1.51e+03	1.50e+03	1.49e+03
	PGD	2.01e+07	4.36e+06	9.57e+05	2.13e+05	4.88e+04	1.21e+04	3.85e+03	1.97e+03	1.57e+03	1.50e+03	1.49e+03	1.49e+03
	SAWAR	1.66e+03	1.62e+03	1.58e+03	1.56e+03	1.53e+03	1.52e+03	1.50e+03	1.50e+03	1.49e+03	1.49e+03	1.49e+03	1.49e+03
LeukSurv	DRAFT	1.79e+07	4.80e+06	1.29e+06	3.48e+05	9.29e+04	2.53e+04	6.59e+03	1.77e+03	5.76e+02	3.10e+02	2.74e+02	2.58e+02
	Noise	5.46e+23	1.79e+21	7.98e+18	2.52e+16	8.53e+13	1.88e+11	5.25e+08	2.35e+06	3.37e+04	2.60e+03	1.21e+03	7.21e+02
	FGSM	8.34e+16	1.79e+15	3.10e+13	4.86e+11	8.13e+09	1.49e+08	2.72e+06	7.03e+04	3.45e+03	6.20e+02	4.31e+02	3.59e+02
	PGD	1.28e+15	3.89e+13	1.03e+12	2.67e+10	7.96e+08	2.30e+07	7.01e+05	2.80e+04	1.98e+03	4.65e+02	3.58e+02	3.17e+02
	SAWAR	2.07e+04	8.38e+03	3.50e+03	1.45e+03	6.62e+02	3.45e+02	2.91e+02	2.70e+02	2.60e+02	2.54e+02	2.52e+02	2.50e+02
TRACE	DRAFT	4.56e+09	4.41e+08	4.44e+07	4.61e+06	5.31e+05	6.71e+04	9.88e+03	2.20e+03	9.05e+02	6.03e+02	5.53e+02	5.32e+02
	Noise	9.11e+08	1.07e+08	1.34e+07	1.74e+06	2.43e+05	3.71e+04	6.63e+03	1.78e+03	8.35e+02	5.93e+02	5.51e+02	5.32e+02
	FGSM	2.01e+11	9.10e+09	4.31e+08	2.24e+07	1.26e+06	8.26e+04	7.79e+03	1.61e+03	7.45e+02	5.74e+02	5.47e+02	5.35e+02
	PGD	4.53e+08	4.81e+07	5.35e+06	6.12e+05	7.63e+04	1.07e+04	2.09e+03	8.37e+02	5.94e+02	5.40e+02	5.31e+02	5.26e+02
	SAWAR	8.79e+02	7.60e+02	6.82e+02	6.29e+02	5.94e+02	5.69e+02	5.53e+02	5.41e+02	5.33e+02	5.28e+02	5.27e+02	5.27e+02
dataDIVAT1	DRAFT	1.43e+04	8.36e+03	4.95e+03	3.01e+03	1.91e+03	1.31e+03	9.99e+02	8.51e+02	7.84e+02	7.56e+02	7.51e+02	7.49e+02
	Noise	5.69e+08	7.64e+07	1.05e+07	1.44e+06	2.07e+05	3.17e+04	5.75e+03	1.61e+03	9.08e+02	7.81e+02	7.67e+02	7.64e+02
	FGSM	2.48e+08	3.36e+07	4.65e+06	6.56e+05	9.64e+04	1.54e+04	3.12e+03	1.14e+03	8.16e+02	7.65e+02	7.61e+02	7.60e+02
	PGD	1.85e+11	8.02e+09	3.47e+08	1.75e+07	9.39e+05	6.14e+04	5.80e+03	1.27e+03	8.03e+02	7.57e+02	7.56e+02	7.57e+02
	SAWAR	9.18e+02	8.55e+02	8.21e+02	8.00e+02	7.86e+02	7.75e+02	7.66e+02	7.59e+02	7.55e+02	7.52e+02	7.51e+02	7.51e+02
flchain	DRAFT	3.35e+23	9.34e+20	2.56e+18	7.37e+15	2.23e+13	6.82e+10	2.29e+08	9.33e+05	1.67e+04	2.03e+03	1.25e+03	1.09e+03
	Noise	2.18e+35	1.20e+31	7.58e+26	4.68e+22	3.48e+18	2.65e+14	2.21e+10	6.77e+06	4.45e+04	1.96e+03	1.22e+03	1.11e+03
	FGSM	1.61e+20	3.47e+17	7.79e+14	1.86e+12	5.49e+09	4.02e+07	6.37e+05	1.86e+04	2.15e+03	1.15e+03	1.09e+03	1.08e+03
	PGD	1.30e+16	7.91e+13	4.93e+11	3.47e+09	4.60e+07	1.33e+06	4.93e+04	4.99e+03	1.45e+03	1.10e+03	1.08e+03	1.07e+03
	SAWAR	6.31e+03	2.22e+03	1.43e+03	1.19e+03	1.12e+03	1.09e+03	1.08e+03	1.07e+03	1.07e+03	1.07e+03	1.07e+03	1.07e+03
prostate	DRAFT	9.74e+05	3.22e+05	1.06e+05	3.58e+04	1.21e+04	4.25e+03	1.64e+03	7.62e+02	4.66e+02	3.70e+02	3.51e+02	3.44e+02
	Noise	7.00e+20	5.05e+18	3.75e+16	2.85e+14	2.28e+12	2.03e+10	2.59e+08	9.04e+06	4.85e+05	6.66e+04	2.94e+04	1.37e+04
	FGSM	3.16e+14	1.04e+13	3.40e+11	1.16e+10	4.07e+08	1.52e+07	6.47e+05	3.56e+04	3.75e+03	1.03e+03	7.20e+02	5.82e+02
	PGD	8.84e+12	4.29e+11	2.18e+10	1.14e+09	5.94e+07	3.27e+06	1.92e+05	1.44e+04	1.91e+03	6.40e+02	5.00e+02	4.39e+02
	SAWAR	9.65e+03	3.88e+03	1.66e+03	8.48e+02	5.43e+02	4.30e+02	3.83e+02	3.59e+02	3.45e+02	3.37e+02	3.35e+02	3.34e+02
retinopathy	DRAFT	1.11e+04	5.64e+03	2.88e+03	1.51e+03	8.08e+02	4.64e+02	2.98e+02	2.18e+02	1.84e+02	1.70e+02	1.68e+02	1.67e+02
	Noise	1.11e+04	5.61e+03	2.88e+03	1.51e+03	8.13e+02	4.68e+02	2.99e+02	2.19e+02	1.84e+02	1.70e+02	1.68e+02	1.68e+02
	FGSM	7.43e+03	3.89e+03	2.07e+03	1.12e+03	6.33e+02	3.86e+02	2.63e+02	2.03e+02	1.78e+02	1.68e+02	1.67e+02	1.68e+02
	PGD	7.30e+03	3.83e+03	2.04e+03	1.11e+03	6.25e+02	3.81e+02	2.60e+02	2.02e+02	1.77e+02	1.68e+02	1.67e+02	1.68e+02
	SAWAR	4.02e+02	3.31e+02	2.79e+02	2.41e+02	2.14e+02	1.95e+02	1.82e+02	1.74e+02	1.69e+02	1.67e+02	1.67e+02	1.67e+02
stagec	DRAFT	1.16e+06	2.94e+05	7.50e+04	1.85e+04	4.81e+03	1.28e+03	3.57e+02	1.14e+02	5.35e+01	4.10e+01	3.98e+01	3.97e+01
	Noise	2.07e+06	4.93e+05	1.14e+05	2.73e+04	6.49e+03	1.61e+03	4.19e+02	1.25e+02	5.50e+01	4.10e+01	3.96e+01	3.95e+01
	FGSM	1.80e+05	5.45e+04	1.67e+04	5.23e+03	1.64e+03	5.30e+02	1.77e+02	7.43e+01	4.54e+01	3.96e+01	3.91e+01	3.92e+01
	PGD	1.54e+05	4.77e+04	1.47e+04	4.63e+03	1.47e+03	4.84e+02	1.69e+02	7.23e+01	4.48e+01	3.93e+01	3.89e+01	3.90e+01
	SAWAR	1.03e+03	5.79e+02	3.30e+02	1.94e+02	1.21e+02	8.21e+01	6.10e+01	4.88e+01	4.22e+01	3.89e+01	3.83e+01	3.81e+01
zinc	DRAFT	2.62e+06	5.85e+05	1.33e+05	3.13e+04	7.43e+03	1.87e+03	5.36e+02	2.03e+02	1.11e+02	8.26e+01	7.71e+01	7.46e+01
	Noise	2.93e+07	4.72e+06	7.76e+05	1.32e+05	2.34e+04	4.45e+03	9.48e+02	2.56e+02	1.14e+02	8.17e+01	7.69e+01	7.53e+01
	FGSM	2.50e+06	4.44e+05	9.49e+04	2.11e+04	4.89e+03	1.24e+03	3.66e+02	1.44e+02	9.01e+01	7.66e+01	7.53e+01	7.57e+01
	PGD	2.55e+06	4.38e+05	8.27e+04	1.87e+04	4.43e+03	1.13e+03	3.40e+02	1.39e+02	8.88e+01	7.66e+01	7.55e+01	7.60e+01
	SAWAR	1.31e+03	7.77e+02	4.72e+02	2.98e+02	1.98e+02	1.42e+02	1.10e+02	9.21e+01	8.28e+01	7.87e+01	7.80e+01	7.78e+01

TABLE VIII: Negative Log Likelihood metric for *SurvSet* datasets (lower is better) for each adversarial training method against the worst-case adversarial attack.