

AutoImpute: Autoencoder based imputation of single-cell RNA-seq data

Supplementary Material

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Supplementary Table S 1: **Separation of “true zeros” from dropouts** - Fraction of zeros (values between 0 and 0.5) in single cell expression matrix against the median bulk expression. The genes are divided into 10 bins based on median bulk genes expression (first bin has only 0 expression gene).

Median Bulk Expression Bins	Algorithms				
	No imputation	AutoImpute	scImpute	Magic	drImpute
0	0.9471	0.9471	0.9194	1.0000	0.8585
0 - 1	0.9589	0.9378	0.8945	1.0000	0.8302
1 - 2	0.9298	0.9350	0.8843	0.9545	0.8343
2 - 3	0.9753	0.9777	0.9537	1.0000	0.9274
3 - 4	0.9507	0.9439	0.8890	1.0000	0.7778
4 - 6	0.9301	0.8746	0.7644	0.9924	0.6247
6 - 8	0.8253	0.5726	0.4807	0.8941	0.3253
8 - 10	0.6086	0.2145	0.1665	0.5344	0.0645
10 - 12	0.3675	0.1060	0.0775	0.1746	0.0120
12 - max	0.1456	0.0480	0.0329	0.0234	0.0024

Supplementary Table S 2: **Gene expression recovery** - Comparison of NMSE, RMSE and MAE calculated between the full unmasked single cell RNA-seq data and the recovered expression matrices with masked matrices as inputs (with varying masking percentage), at the masked locations, for AutoImpute, scImpute, and Magic algorithms.

Masking %	NMSE			RMSE			MAE		
	AutoImpute	scImpute	Magic	AutoImpute	scImpute	Magic	AutoImpute	scImpute	Magic
Blakeley									
10	1.80 ± 0.1270	1.4833	0.6173	15.34 ± 0.8754	12.3598	1.26E+03	12.21 ± 0.8522	9.7188	497.2412
20	1.82 ± 0.2114	2.0525	0.5952	15.55 ± 1.7696	17.4443	1.42E+03	12.35 ± 1.387	11.7828	520.2138
30	1.84 ± 0.1226	2.8635	0.6305	15.34 ± 1.0574	24.3536	1.42E+03	12.37 ± 0.6635	13.1893	530.4287
40	1.99 ± 0.0651	3.5833	0.6771	16.97 ± 0.4960	30.6860	1.52E+03	13.55 ± 0.3941	13.1764	539.3752
50	2.22 ± 0.0350	12.4788	0.7338	18.86 ± 0.3273	105.8491	1.73E+03	15.01 ± 0.2601	24.1704	575.0085
Jurkat-293T									
10	0.39 ± 0.0143	2.0607	0.5239	1.09 ± 0.0402	3.5464	3.2438	0.96 ± 0.0382	0.3870	0.4744
20	0.40 ± 0.0377	5.6608	0.5825	1.19 ± 0.1063	9.1853	3.4002	1.04 ± 0.0508	0.8851	0.4848
30	0.44 ± 0.0895	13.6579	0.6332	1.25 ± 0.0500	20.7366	3.4586	1.34 ± 0.2031	1.7311	0.4855
40	0.48 ± 0.5047	54.4675	0.6791	1.41 ± 0.4410	76.5812	3.4294	1.12 ± 0.6816	3.9182	0.4728
50	0.49 ± 0.6816	449.9235	0.7199	1.63 ± 0.6029	577.3294	3.3262	1.79 ± 0.9975	10.9636	0.4441
Kolodziejczyk									
10	2.82 ± 0.0127	1.6747	0.3711	25.13 ± 0.1063	14.9218	1.23E+03	20.04 ± 0.0916	9.4494	192.2411
20	3.14 ± 0.0221	3.0888	0.4597	27.92 ± 0.2057	27.4045	1.58E+03	22.24 ± 0.1613	13.011	203.654
30	3.39 ± 0.0177	6.1239	0.5940	30.21 ± 0.1418	54.4813	1.86E+03	24.09 ± 0.1347	19.0343	221.3524
40	3.62 ± 0.0020	13.4705	0.6770	32.22 ± 0.0300	119.9824	2.20E+03	25.66 ± 0.0297	29.8625	238.5226
50	3.87 ± 0.0043	90.4541	0.7238	34.40 ± 0.0502	803.8800	2.23E+03	27.43 ± 0.0500	69.1118	257.9593
Quake									
10	0.95 ± 0.0007	24.82263737	0.7731	8.51 ± 0.0705	2.23E+02	8.38E+03	6.78 ± 0.0268	63.44775511	2013.0823
20	0.99 ± 0.0017	1.29E+02	0.8031	8.89 ± 0.0238	1.15E+03	8.14E+03	7.08 ± 0.0195	1.25E+02	1978.092744
30	1.02 ± 0.0013	6.06E+02	0.8426	9.18 ± 0.0144	5.43E+03	8.52E+03	7.30 ± 0.0159	3.63E+02	1998.363221
40	1.07 ± 0.0008	3.13E+03	0.8865	9.56 ± 0.0157	2.80E+04	9.46E+03	7.60 ± 0.0044	1.24E+03	2011.990659
50	1.10 ± 0.0015	2.18E+06	0.9075	9.85 ± 0.0013	1.95E+07	9.68E+03	7.83 ± 0.0049	1.35E+05	2026.776616
Preimplantation									
10	0.29 ± 0.0094	1.6424	0.777	2.42 ± 0.0888	6.8395	1.71E+03	1.90 ± 0.9902	0.7164	286.8199
20	0.29 ± 0.0057	6.6387	0.8052	2.42 ± 0.0451	26.0432	1.85E+03	1.85 ± 0.0661	2.0347	313.6260
30	0.31 ± 0.1280	54.469	0.8277	2.57 ± 0.1050	200.0148	1.96E+03	1.91 ± 0.1550	6.2662	330.6937
40	0.31 ± 0.0180	120.2089	0.8501	2.58 ± 0.1530	408.693	2.00E+03	2.04 ± 0.1256	13.8265	341.3199
50	0.31 ± 0.014	1.21E+04	0.8672	2.60 ± 0.1200	3.77E+04	2.08E+03	2.24 ± 0.1341	6.2662	344.7977
Usoskin									
10	2.23 ± 0.0182	41.22689	0.6443	9.69 ± 0.0564	1.60E+02	52.5217	7.69 ± 0.0256	24.81428717	14.1806
20	2.40 ± 0.0022	3.18E+03	0.6720	10.40 ± 0.0411	1.30E+04	55.2395	8.26 ± 0.0310	1.83E+02	14.7904
30	2.53 ± 0.0121	3.96E+07	0.7645	10.95 ± 0.0325	1.67E+08	68.8496	8.72 ± 0.0293	8.18E+05	15.2874
40	2.63 ± 0.0119	8.06E+05	0.7696	11.37 ± 0.0439	3.42E+06	66.2696	9.05 ± 0.0444	1.56E+04	15.8968
50	2.75 ± 0.0764	2.56E+06	0.8590	11.93 ± 0.0586	1.09E+07	84.2230	9.50 ± 0.0514	5.70E+04	17.0062
Zeisel									
10	1.42 ± 0.0077	19.3472739	0.9038	4.23 ± 0.0238	57.51650418	1.68E+01	3.07 ± 0.0326	13.02629774	1.9248
20	1.77 ± 0.0028	51.45857788	0.926	5.25 ± 0.0162	152.4683696	1.72E+01	4.05 ± 0.0134	23.33065176	2.0222
30	1.99 ± 0.0095	1025.814158	0.9409	5.91 ± 0.0377	3043.370528	1.79E+01	4.63 ± 0.0320	76.24520225	2.1272
40	2.20 ± 0.0008	12127.06557	0.9554	6.52 ± 0.0024	35970.72819	1.81E+01	5.14 ± 0.0023	324.6180099	2.2441
50	2.34 ± 0.0235	48999902.91	0.9612	6.96 ± 0.0670	1.45E+08	1.73E+01	5.49 ± 0.0553	288527.3319	2.364

Supplementary Table S 3: **Median of coefficient of variation** across cells from a random subpopulation selected from each dataset imputed using the following algorithms.

Datasets	No Imputation	scImpute	drImpute	Magic	AutoImpute
Blakeley	3.3114	2.1918	2.4142	3.35E-06	2.7507
Jurkat-293T	2.8399	12.0685	1.9193	0.031	0.0164
Kolodziejczyk	78.0416	14.5777	10.127	0.2526	4.6243
PBMC	58.3697	–	–	–	0.0011
Preimplantation	6.6962	5.0814	4.1446	0.0595	0.0012
Quake	11.5473	6.4453	4.0275	0.0106	8.8568
Usoskin	28.9626	14.4071	27.7187	0.2375	10.9204
Zeisel	32.4045	4.891	1.8093	0.0728	11.0032

Supplementary Table S 4: **Average silhouette index values** showing cell-type separability on cell groupings corresponding to first two principal components of expressions.

Datasets	No Imputation	scImpute	drImpute	Magic	AutoImpute
Blakeley	0.93	0.67	0.44	0.20	0.87
Jurkat-293T	0.94	0.93	0.70	0.90	0.96
Kolodziejczyk	0.24	0.46	0.43	-0.18	0.37
Preimplantation	-0.007	-0.06	0.03	0.01	0.08
Quake	-0.26	0.36	0.32	-0.03	0.33
Usoskin	0.33	0.12	-0.13	-0.08	0.36
Zeisel	-0.14	0.19	0.08	-0.16	0.34