

Extended evaluation of differential chromatin interaction detection analysis using real Hi-C data

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Introduction

To evaluate the performance of different normalization methods on the detection of chromatin interaction differences, controlled changes were used. Real Hi-C data is used: GM12878, Chromosome 1, MboI vs. DpnII restriction enzymes, 1Mb resolution. The dimensions of the chromatin interaction matrices are 250 x 250. The performance of the joint (**loess** and **MA**) and individual (**ChromoR**, Iterative Correction and Eigenvector decomposition - **ICE**, Knight-Ruiz - **KR**, Sequential Component Normalization - **SCN**) normalization methods at varying fold changes (1.5 by default) and varying number of controlled changes (500 by default) is investigated. The data were globally rescaled to have the same total count of interaction frequencies.

Fold changes are applied to one of the datasets by up-regulating the selected IF if the difference between the datasets is positive. If the difference between the datasets at that point is negative the IF is down-regulated by the specified fold change. This method of making changes ensures that the fold change specified is actually realized on the MD plot.

The effect of fold change

1.5 Fold change

	loess	chromoR	ice	kr	scn	ma
true positive	199	11	11	16	13	183
false positive	1020	1440	1150	1190	1200	1030
true negative	24900	29400	23800	24700	24700	24900
false negative	301	489	465	484	487	317
Total	26400	31400	25400	26400	26400	26400
TPR	0.398	0.022	0.0231	0.032	0.026	0.366
SPC	0.961	0.953	0.954	0.954	0.954	0.96
F1	0.974	0.968	0.967	0.967	0.967	0.974
AUC	0.855	0.535	0.768	0.753	0.666	0.857
AUC 20%	0.113	0.021	0.0801	0.0778	0.0485	0.11
FDR	0.836	0.992	0.991	0.987	0.989	0.849
Accuracy	0.95	0.938	0.936	0.937	0.936	0.949
Precision	0.164	0.00757	0.00947	0.0132	0.0107	0.151
FPR	0.0392	0.0467	0.0461	0.046	0.0464	0.0397
FNR	0.602	0.978	0.977	0.968	0.974	0.634
FOR	0.012	0.0163	0.0192	0.0192	0.0193	0.0126
NPV	0.988	0.984	0.981	0.981	0.981	0.987

2.0 Fold change

	loess	chromoR	ice	kr	scn	ma
true positive	373	14	27	27	33	372
false positive	838	1440	1140	1190	1180	837
true negative	25100	29400	23800	24700	24700	25100
false negative	127	486	452	473	467	128
Total	26400	31400	25400	26400	26400	26400
TPR	0.746	0.028	0.0564	0.054	0.066	0.744
SPC	0.968	0.953	0.954	0.954	0.954	0.968
F1	0.981	0.968	0.968	0.967	0.968	0.981
AUC	0.96	0.569	0.528	0.526	0.593	0.959
AUC 20%	0.167	0.022	0.0265	0.0252	0.0313	0.167
FDR	0.692	0.99	0.977	0.978	0.973	0.692
Accuracy	0.963	0.939	0.938	0.937	0.938	0.963
Precision	0.308	0.00966	0.0232	0.0222	0.0271	0.308
FPR	0.0324	0.0465	0.0455	0.046	0.0457	0.0323
FNR	0.254	0.972	0.944	0.946	0.934	0.256
FOR	0.00504	0.0162	0.0186	0.0188	0.0185	0.00508
NPV	0.995	0.984	0.981	0.981	0.981	0.995

4.0 Fold change

	loess	chromoR	ice	kr	scn	ma
true positive	489	89	414	422	428	489
false positive	718	1360	749	797	802	721
true negative	25200	29500	24200	25100	25100	25200
false negative	11	411	67	78	72	11
Total	26400	31400	25400	26400	26400	26400
TPR	0.978	0.178	0.861	0.844	0.856	0.978
SPC	0.972	0.956	0.97	0.969	0.969	0.972
F1	0.986	0.971	0.983	0.983	0.983	0.986
AUC	0.995	0.504	0.976	0.97	0.974	0.995
AUC 20%	0.195	0.0257	0.183	0.179	0.18	0.195
FDR	0.595	0.939	0.644	0.654	0.652	0.596
Accuracy	0.972	0.944	0.968	0.967	0.967	0.972
Precision	0.405	0.0614	0.356	0.346	0.348	0.404
FPR	0.0277	0.0441	0.03	0.0308	0.031	0.0278
FNR	0.022	0.822	0.139	0.156	0.144	0.022
FOR	0.000437	0.0137	0.00276	0.0031	0.00286	0.000437
NPV	1	0.986	0.997	0.997	0.997	1

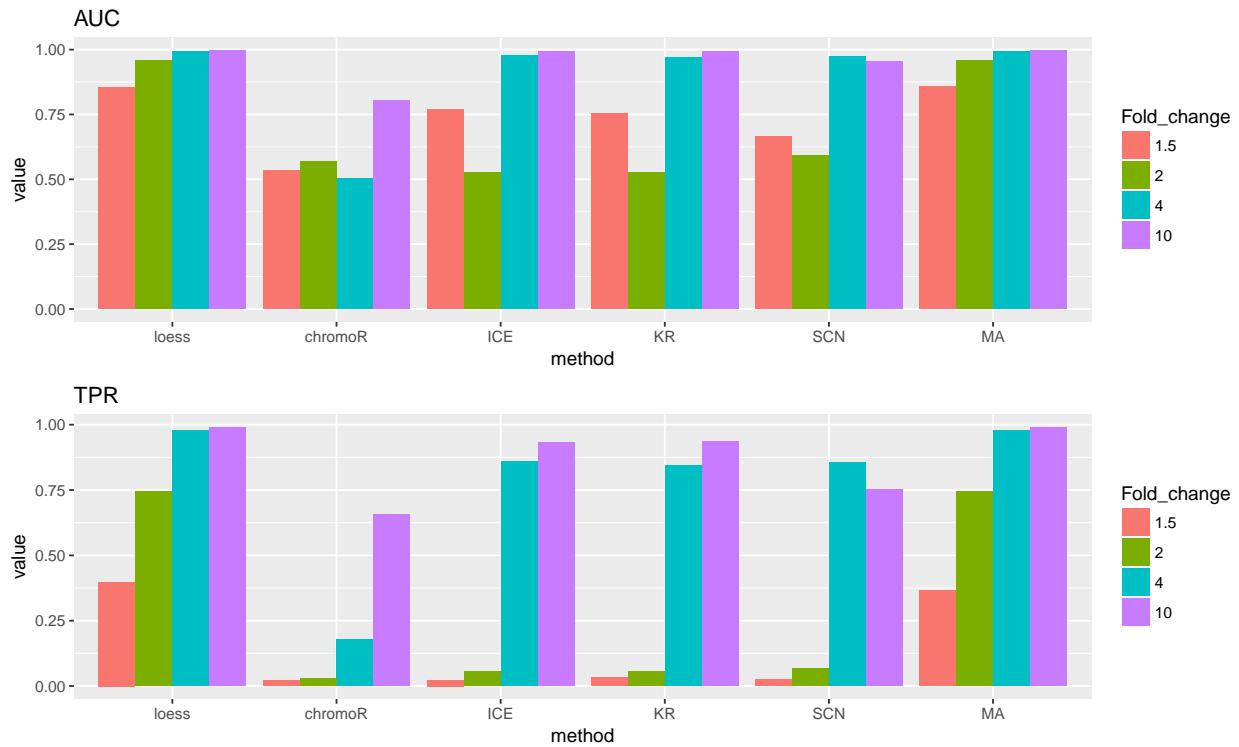
10.0 fold change

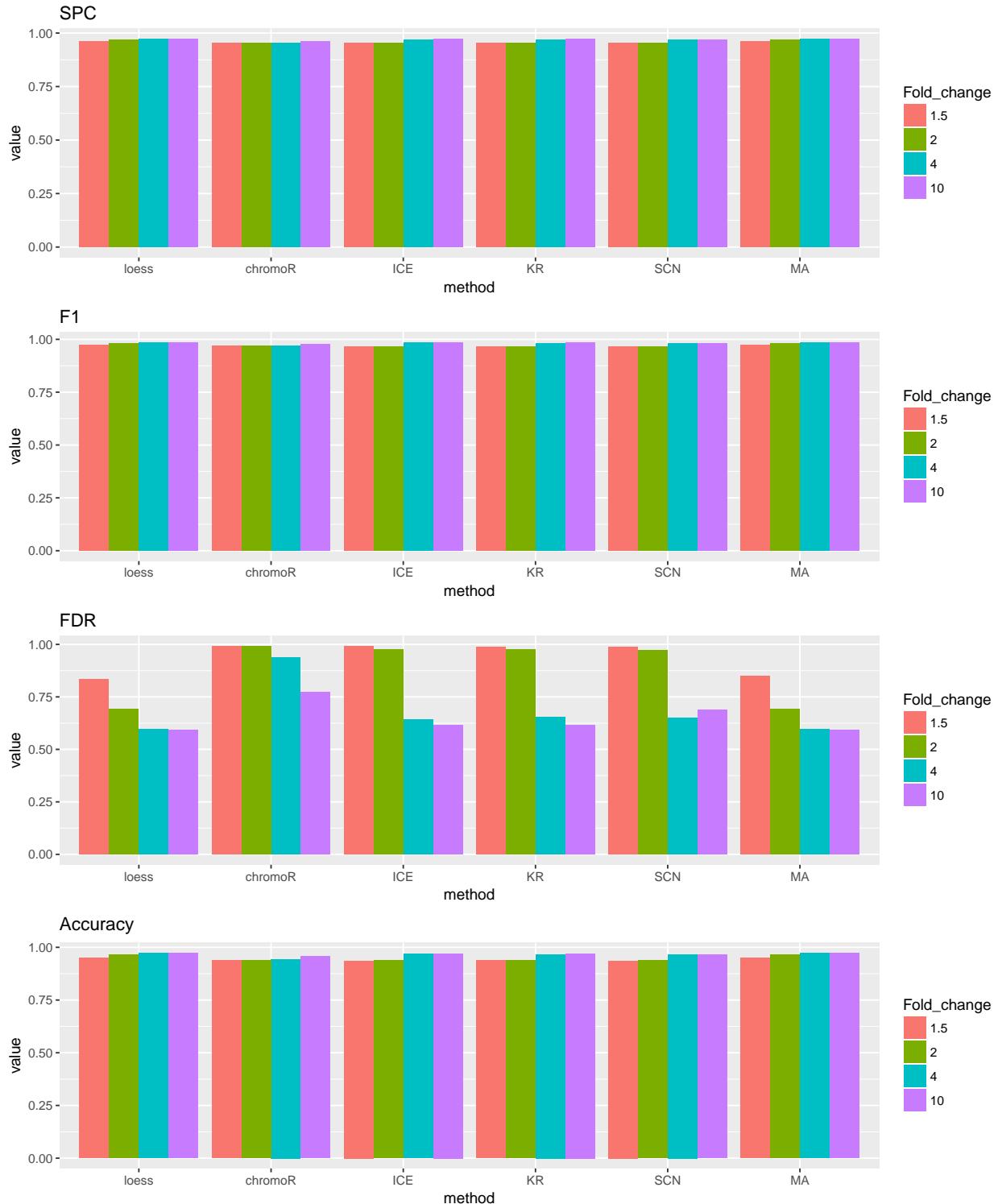
	loess	chromoR	ice	kr	scn	ma
true positive	495	329	448	467	376	494
false positive	725	1130	721	748	837	715
true negative	25200	29700	24200	25200	25100	25200
false negative	5	171	33	33	124	6

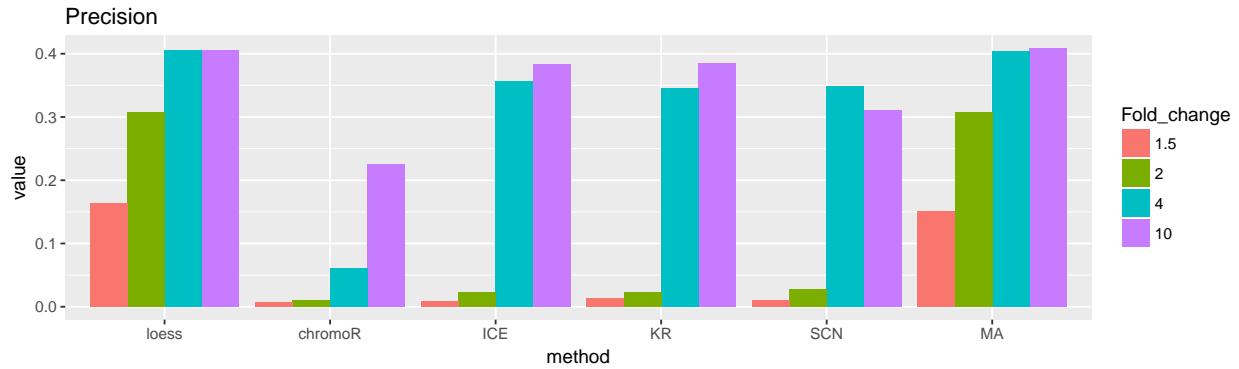
	loess	chromoR	ice	kr	scn	ma
Total	26400	31400	25400	26400	26400	26400
TPR	0.99	0.658	0.931	0.934	0.752	0.988
SPC	0.972	0.963	0.971	0.971	0.968	0.972
F1	0.986	0.979	0.985	0.985	0.981	0.986
AUC	0.996	0.806	0.992	0.992	0.955	0.996
AUC 20%	0.196	0.134	0.192	0.192	0.167	0.196
FDR	0.594	0.775	0.617	0.616	0.69	0.591
Accuracy	0.972	0.959	0.97	0.97	0.964	0.973
Precision	0.406	0.225	0.383	0.384	0.31	0.409
FPR	0.028	0.0366	0.0289	0.0289	0.0323	0.0276
FNR	0.01	0.342	0.0686	0.066	0.248	0.012
FOR	0.000199	0.00572	0.00136	0.00131	0.00492	0.000238
NPV	1	0.994	0.999	0.999	0.995	1

Bar plots

The bar plots show comparisons of the effect of different fold changes using fixed numbers of controlled changes on different performance metrics.







Summary

For the real data with changes added, `loess` was once again able to detect the most true changes compared to the other normalization methods. `loess` was clearly superior to the individual methods at the lower fold changes (1.5 and 2) with much higher TPRs compared to the other methods with TPRs below 10%. `loess` also had the lowest number of false negatives compared to the other methods. The gap in detection once again began to close at the higher fold changes between `loess`, `KR`, `SCN`, and `ICE` however `loess` still was slightly better. `ChromoR` once again proved to be the worst normalization method and had the lowest detection rates. `loess` was slightly better than `MA` normalization at the lower fold changes and at higher fold changes the methods were about equal.

The effect of introduced number of changes

A given number of interaction frequencies were increased or decreased to produce a 1.5-fold change.

1 change

	loess	chromoR	ice	kr	scn	ma
true positive	0	0	0	0	0	0
false positive	1220	1450	1170	1220	1220	1220
true negative	25200	29900	24300	25200	25200	25200
false negative	1	1	1	1	1	1
Total	26400	31400	25400	26400	26400	26400
TPR	0	0	0	0	0	0
SPC	0.954	0.954	0.954	0.954	0.954	0.954
F1	0.976	0.976	0.976	0.976	0.976	0.976
AUC	0.89	0.925	0.996	0.996	0.905	0.85
AUC 20%	0.0905	0.125	0.196	0.196	0.105	0.0502
FDR	1	1	1	1	1	1
Accuracy	0.954	0.954	0.954	0.954	0.954	0.954
Precision	0	0	0	0	0	0
FPR	0.0461	0.0463	0.0459	0.0464	0.0461	0.0461
FNR	1	1	1	1	1	1
FOR	3.97e-05	3.34e-05	4.12e-05	3.97e-05	3.97e-05	3.97e-05
NPV	1	1	1	1	1	1

100 changes

	loess	chromoR	ice	kr	scn	ma
true positive	50	1	0	1	2	45
false positive	1160	1460	1180	1210	1220	1180
true negative	25100	29800	24200	25100	25100	25100
false negative	50	99	96	99	98	55
Total	26400	31400	25400	26400	26400	26400
TPR	0.5	0.01	0	0.01	0.02	0.45
SPC	0.956	0.953	0.954	0.954	0.954	0.955
F1	0.976	0.974	0.974	0.975	0.974	0.976
AUC	0.852	0.546	0.788	0.769	0.693	0.856
AUC 20%	0.11	0.0131	0.0875	0.0841	0.0558	0.114
FDR	0.959	0.999	1	0.999	0.998	0.963
Accuracy	0.954	0.95	0.95	0.95	0.95	0.953
Precision	0.0413	0.000683	0	0.000824	0.00164	0.0368
FPR	0.0442	0.0468	0.0464	0.0461	0.0462	0.0448
FNR	0.5	0.99	1	0.99	0.98	0.55
FOR	0.00199	0.00331	0.00396	0.00393	0.00389	0.00218
NPV	0.998	0.997	0.996	0.996	0.996	0.998

200 changes

	loess	chromoR	ice	kr	scn	ma
true positive	88	9	3	3	0	82
false positive	1130	1460	1170	1210	1210	1120
true negative	25100	29700	24100	25000	25000	25100
false negative	112	191	191	197	200	118
Total	26400	31400	25400	26400	26400	26400
TPR	0.44	0.045	0.0155	0.015	0	0.41
SPC	0.957	0.953	0.954	0.954	0.954	0.957
F1	0.976	0.973	0.973	0.973	0.973	0.976
AUC	0.84	0.521	0.749	0.748	0.681	0.842
AUC 20%	0.11	0.0228	0.0747	0.0746	0.0575	0.109
FDR	0.928	0.994	0.997	0.998	1	0.932
Accuracy	0.953	0.947	0.947	0.947	0.946	0.953
Precision	0.0721	0.00612	0.00256	0.00247	0	0.0679
FPR	0.0432	0.0469	0.0463	0.0463	0.0463	0.0429
FNR	0.56	0.955	0.985	0.985	1	0.59
FOR	0.00445	0.00639	0.00788	0.00782	0.00794	0.00468
NPV	0.996	0.994	0.992	0.992	0.992	0.995

1000 changes

	loess	chromoR	ice	kr	scn	ma
true positive	367	30	18	25	19	343
false positive	855	1430	1150	1190	1190	865
true negative	24500	28900	23300	24200	24200	24500
false negative	633	970	946	975	981	657

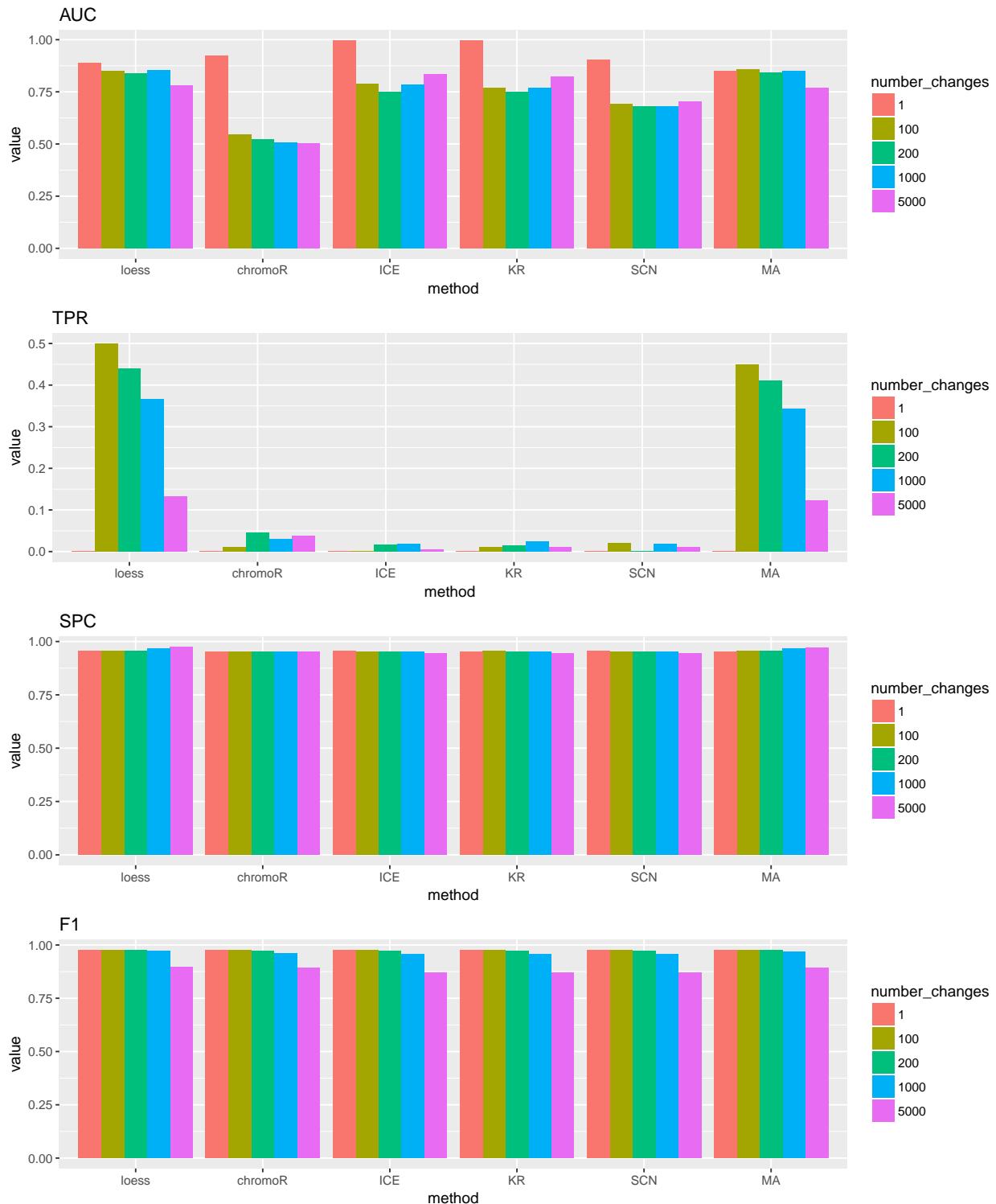
	loess	chromoR	ice	kr	scn	ma
Total	26400	31400	25400	26400	26400	26400
TPR	0.367	0.03	0.0187	0.025	0.019	0.343
SPC	0.966	0.953	0.953	0.953	0.953	0.966
F1	0.971	0.96	0.957	0.957	0.957	0.97
AUC	0.856	0.507	0.783	0.77	0.682	0.849
AUC 20%	0.113	0.0166	0.085	0.0827	0.0554	0.109
FDR	0.7	0.979	0.985	0.979	0.984	0.716
Accuracy	0.944	0.924	0.918	0.918	0.918	0.942
Precision	0.3	0.0206	0.0154	0.0206	0.0157	0.284
FPR	0.0337	0.047	0.0471	0.0468	0.0469	0.0341
FNR	0.633	0.97	0.981	0.975	0.981	0.657
FOR	0.0251	0.0324	0.039	0.0387	0.0389	0.0261
NPV	0.975	0.968	0.961	0.961	0.961	0.974

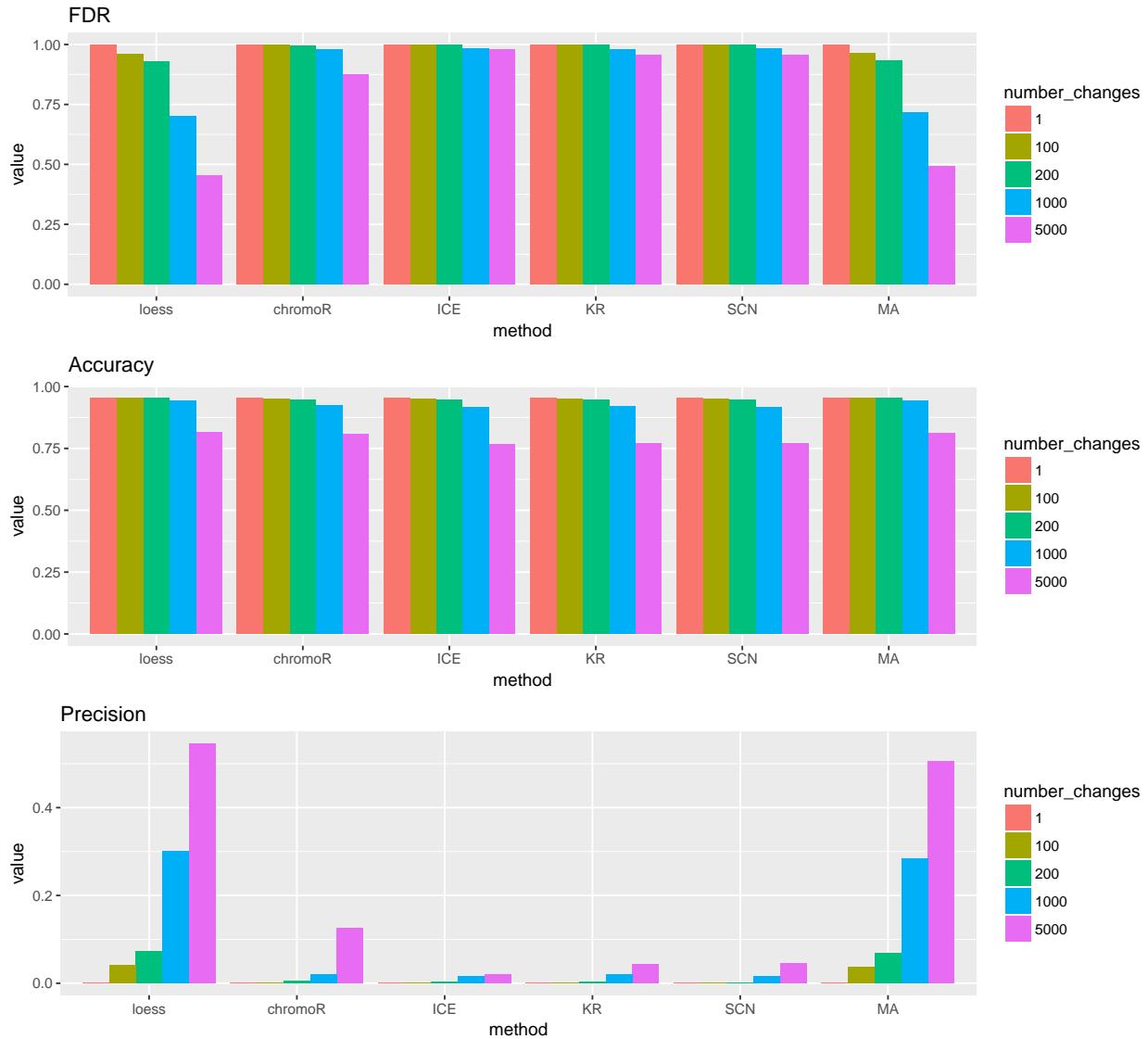
5000 changes

	loess	chromoR	ice	kr	scn	ma
true positive	665	184	24	52	55	611
false positive	554	1280	1140	1160	1170	598
true negative	20800	25100	19500	20200	20200	20800
false negative	4340	4820	4770	4950	4940	4390
Total	26400	31400	25400	26400	26400	26400
TPR	0.133	0.0368	0.005	0.0104	0.011	0.122
SPC	0.974	0.952	0.945	0.946	0.945	0.972
F1	0.895	0.892	0.868	0.869	0.869	0.893
AUC	0.781	0.504	0.834	0.821	0.705	0.767
AUC 20%	0.0811	0.0163	0.0994	0.0962	0.0542	0.0739
FDR	0.454	0.874	0.979	0.957	0.955	0.495
Accuracy	0.815	0.806	0.768	0.769	0.768	0.811
Precision	0.546	0.126	0.0207	0.0428	0.0448	0.505
FPR	0.0259	0.0483	0.0552	0.0543	0.0549	0.0279
FNR	0.867	0.963	0.995	0.99	0.989	0.878
FOR	0.172	0.161	0.197	0.196	0.196	0.174
NPV	0.828	0.839	0.803	0.804	0.804	0.826

Bar plots

Below are bar plots showing comparisons of the different normalization methods over the varying numbers of changes at a fixed fold change for selected metrics.





Summary

`loess` was able to detect a single true change added at a 1.5 fold change. `loess` also detected the most changes that were added for all levels changes made compared to the other methods. `loess` again had the lowest numbers of false negatives compared to the other methods. Interestingly, `ChromoR` seemed to detect more differences when a large number of changes were made (1000 and 5000) compared to `KR`, `SCN`, and `ICE`. `loess` also performed slightly better than `MA` normalization at all levels of fold changes.