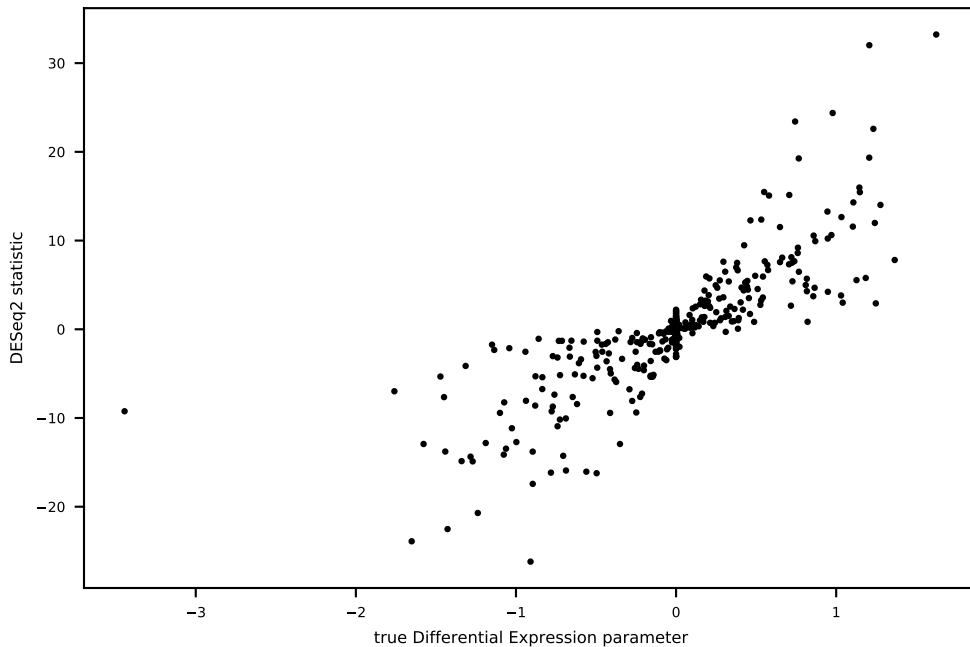
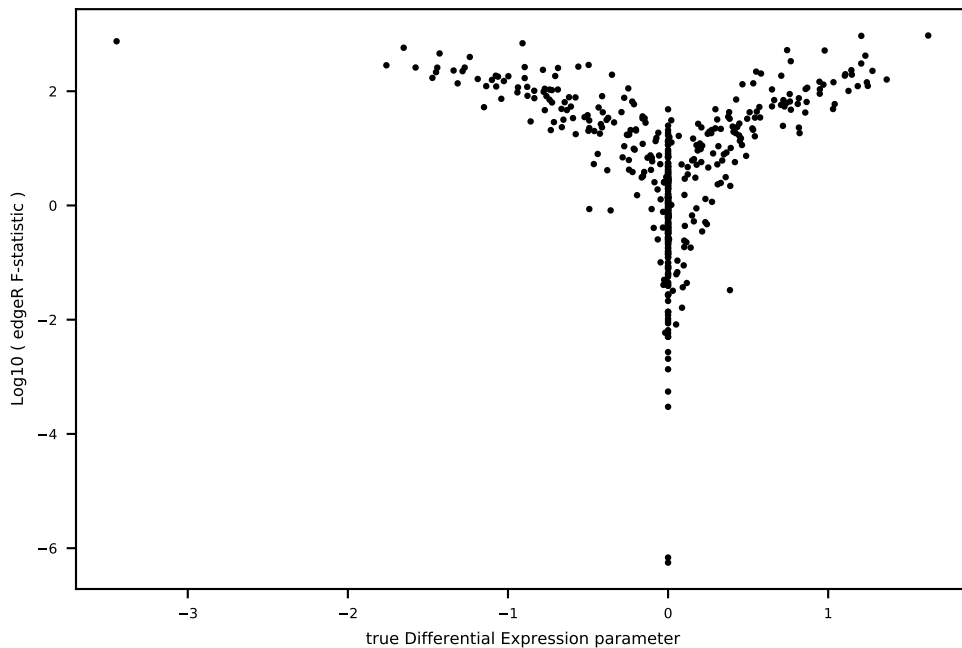


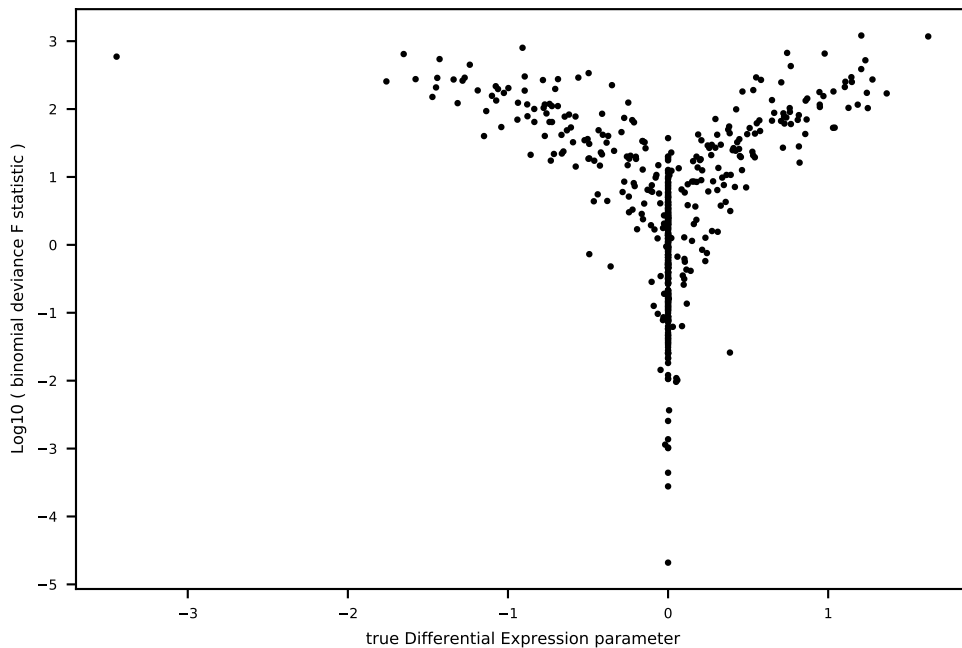
Splatter simulated data with 2 clusters - data set 1
parameters derived from 504 most highly expressed genes - Kang Lupus B cell data
compare DESeq2 statistic to true DE parameter



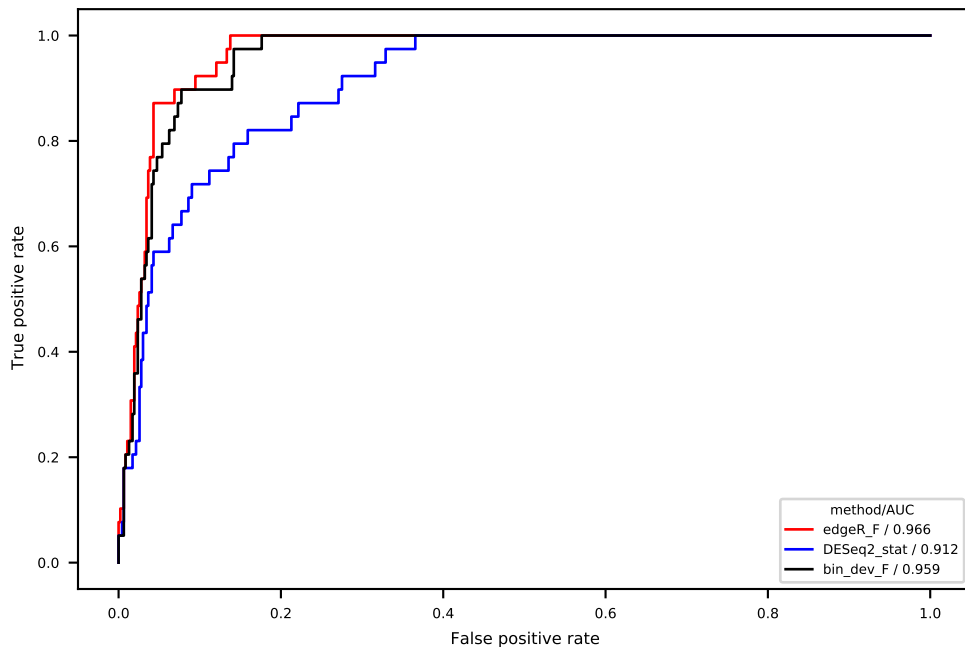
Splatter simulated data with 2 clusters - data set 1
parameters derived from 504 most highly expressed genes - Kang Lupus B cell data
compare edgeR F-statistic to true DE parameter



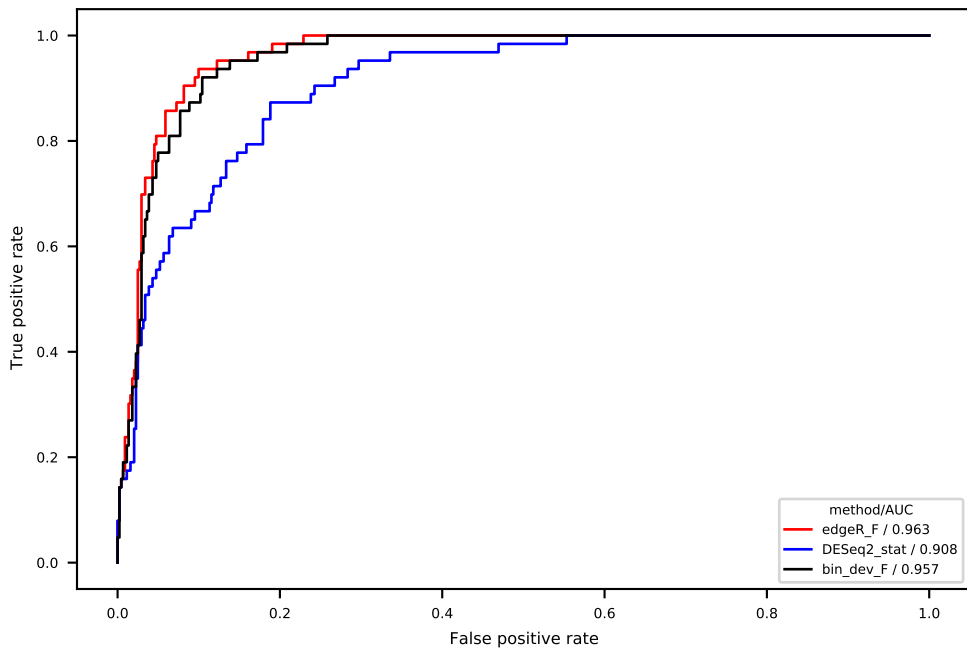
Splatter simulated data with 2 clusters - data set 1
parameters derived from 504 most highly expressed genes - Kang Lupus B cell data
compare binomial deviance F-statistic to true DE parameter



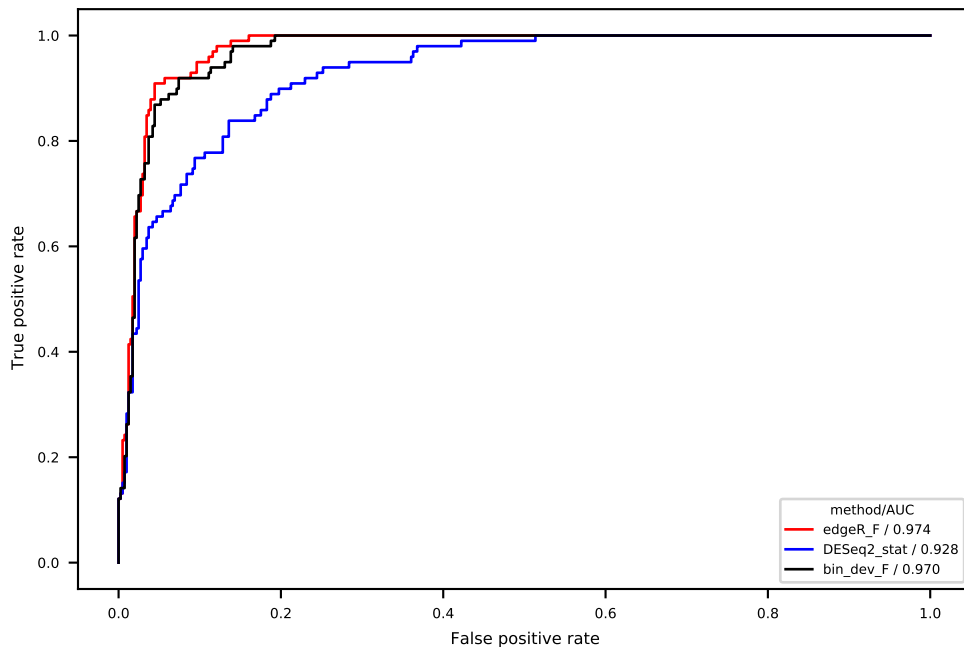
Splatter simulated data with 2 clusters - data set 1
parameters derived from 504 most highly expressed genes - Kang Lupus B cell data
ROC curves for edgeR, DESeq2, and bin_dev_F
define genes as Differentially Expressed if $|\log_2(\text{DEFacGroup2} / \text{DEFacGroup1})| > 1.00$
number Differentially Expressed: 39



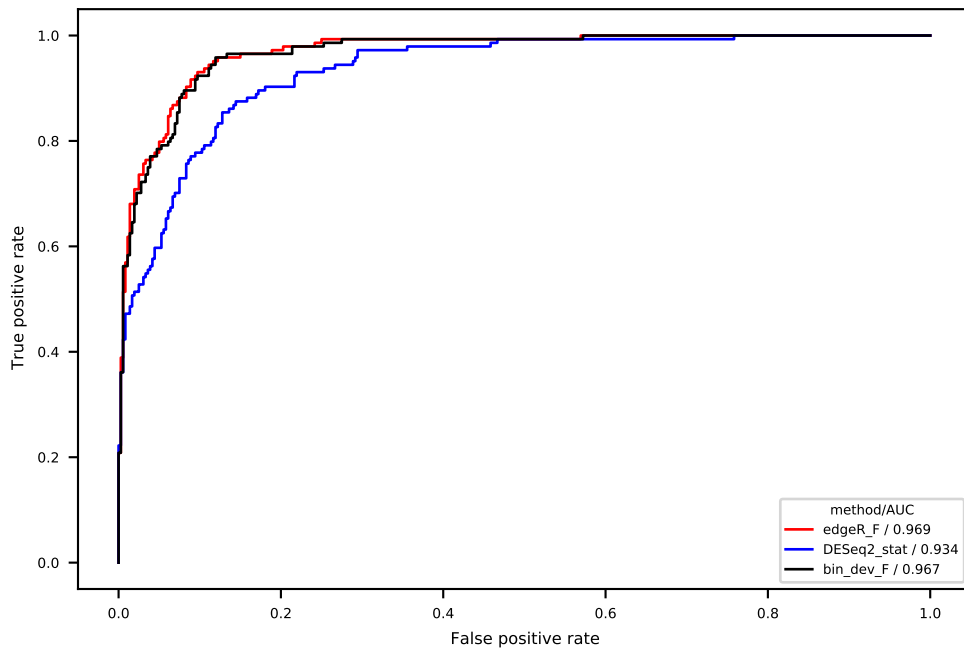
Splatter simulated data with 2 clusters - data set 1
parameters derived from 504 most highly expressed genes - Kang Lupus B cell data
ROC curves for edgeR, DESeq2, and bin_dev_F
define genes as Differentially Expressed if $|\log_2(\text{DEFacGroup2} / \text{DEFacGroup1})| > 0.80$
number Differentially Expressed: 63



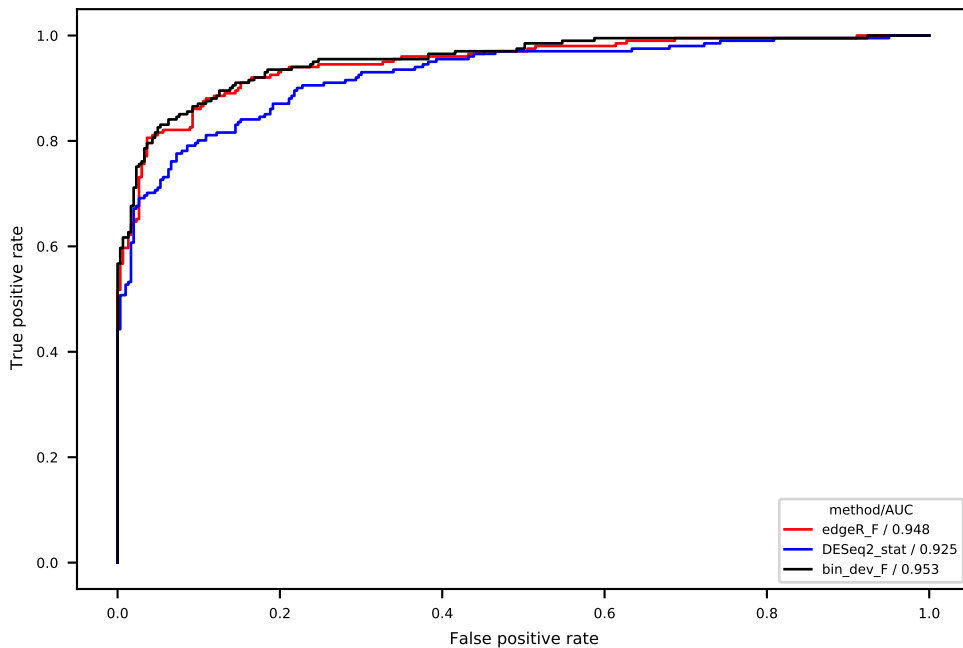
Splatter simulated data with 2 clusters - data set 1
parameters derived from 504 most highly expressed genes - Kang Lupus B cell data
ROC curves for edgeR, DESeq2, and bin_dev_F
define genes as Differentially Expressed if $|\log_2(\text{DEFacGroup2} / \text{DEFacGroup1})| > 0.60$
number Differentially Expressed: 99



Splatter simulated data with 2 clusters - data set 1
parameters derived from 504 most highly expressed genes - Kang Lupus B cell data
ROC curves for edgeR, DESeq2, and bin_dev_F
define genes as Differentially Expressed if $|\log_2(\text{DEFacGroup2} / \text{DEFacGroup1})| > 0.40$
number Differentially Expressed: 144



Splatter simulated data with 2 clusters - data set 1
parameters derived from 504 most highly expressed genes - Kang Lupus B cell data
ROC curves for edgeR, DESeq2, and bin_dev_F
define genes as Differentially Expressed if $|\log_2(\text{DEFacGroup2} / \text{DEFacGroup1})| > 0.20$
number Differentially Expressed: 201



Splatter simulated data with 2 clusters - data set 1
parameters derived from 504 most highly expressed genes - Kang Lupus B cell data
ROC curves for edgeR, DESeq2, and bin_dev_F
define genes as Differentially Expressed if $|\text{abs}(\log_2(\text{DEFacGroup2} / \text{DEFacGroup1}))| > 0.10$
number Differentially Expressed: 237

