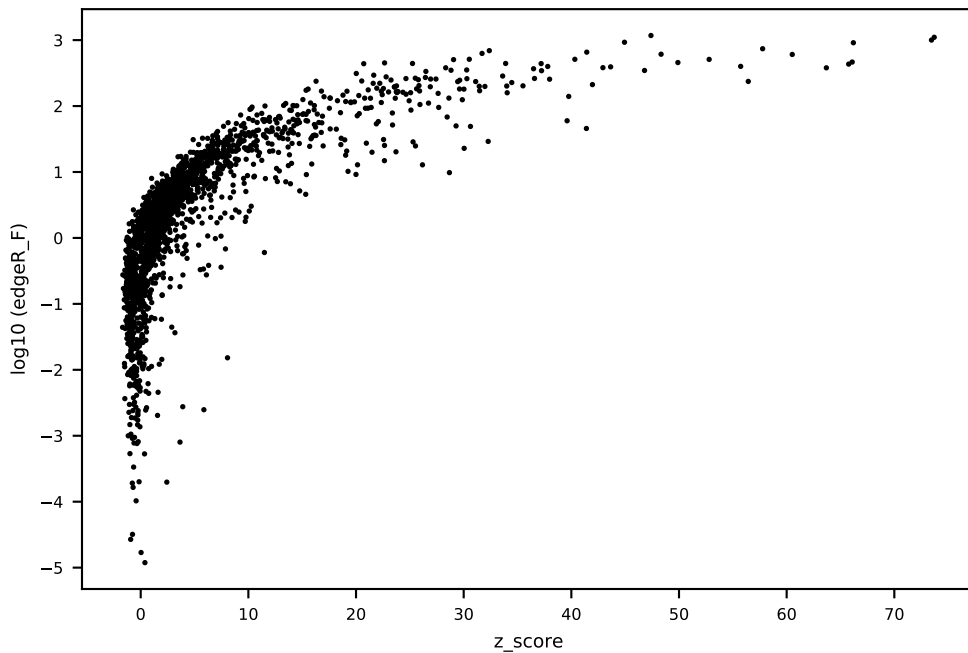
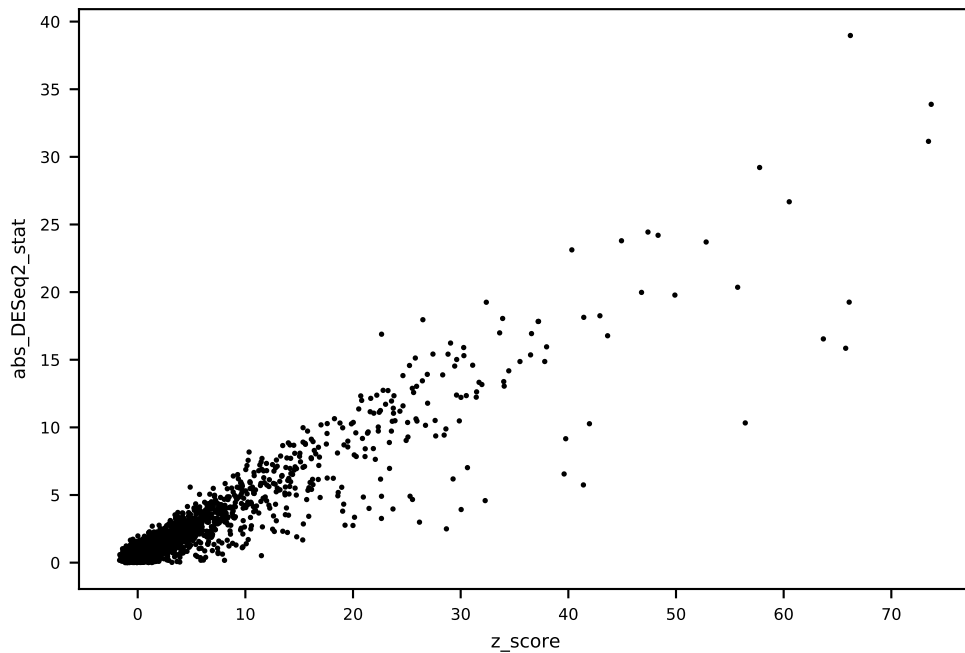


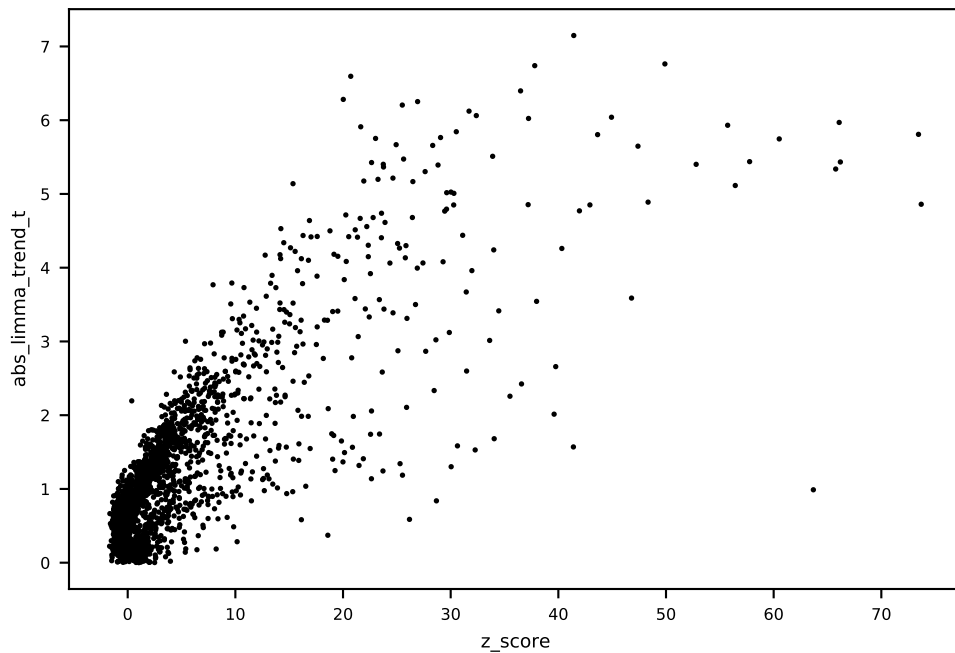
Crowell_Kang18_8vs8 scRNA-seq data - B cells:
compare pi-hat range analysis to muscat pseudo bulk
edgeR F statistic vs z-scores



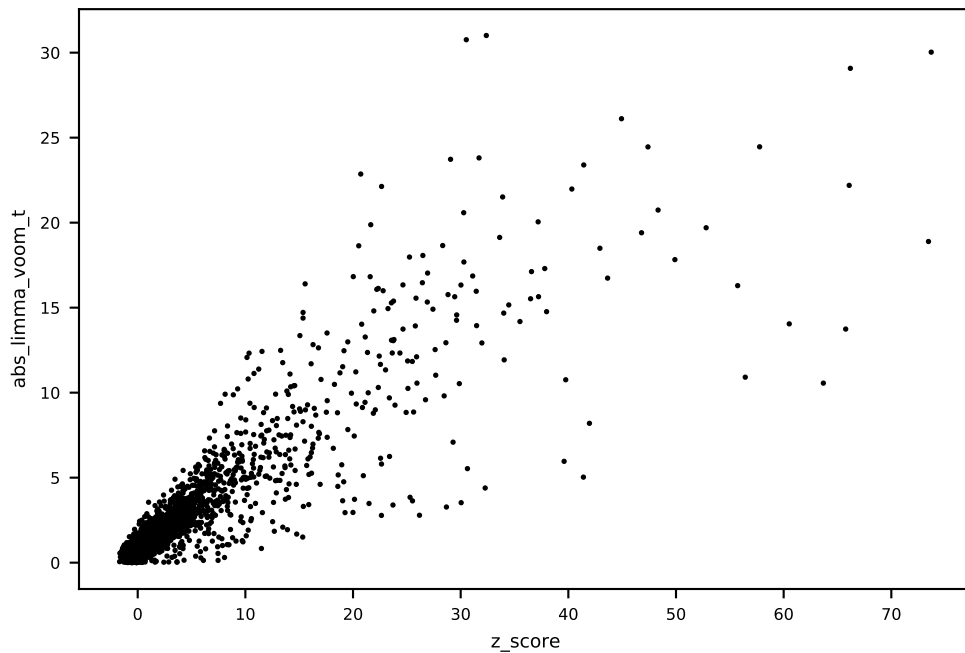
Crowell_Kang18_8vs8 scRNA-seq data - B cells:
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DESeq2_stat vs z-scores



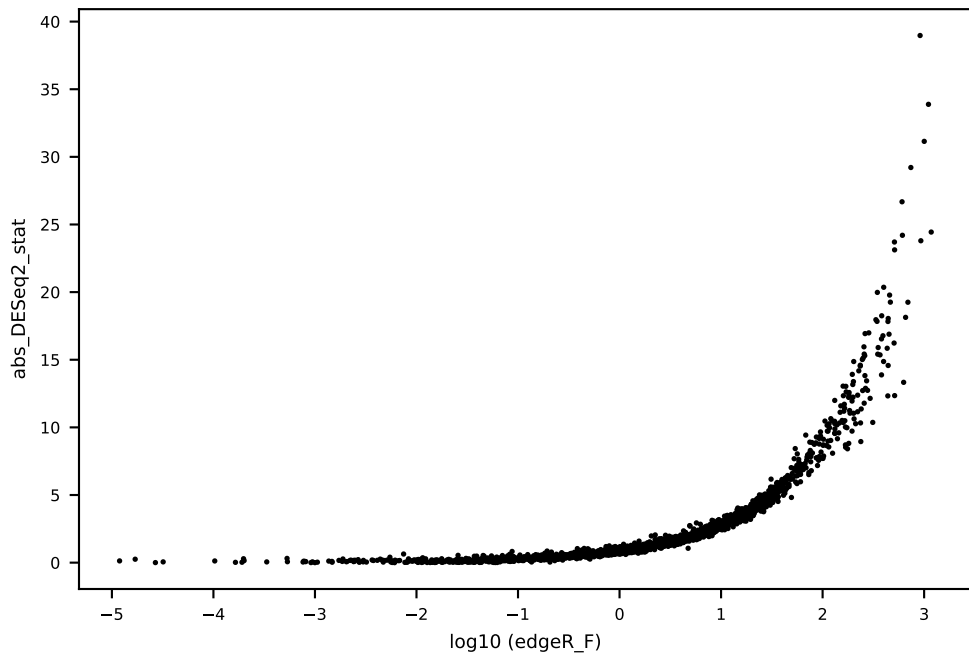
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compare pi-hat range analysis to muscat pseudo bulk
limma_trend_t vs z-scores



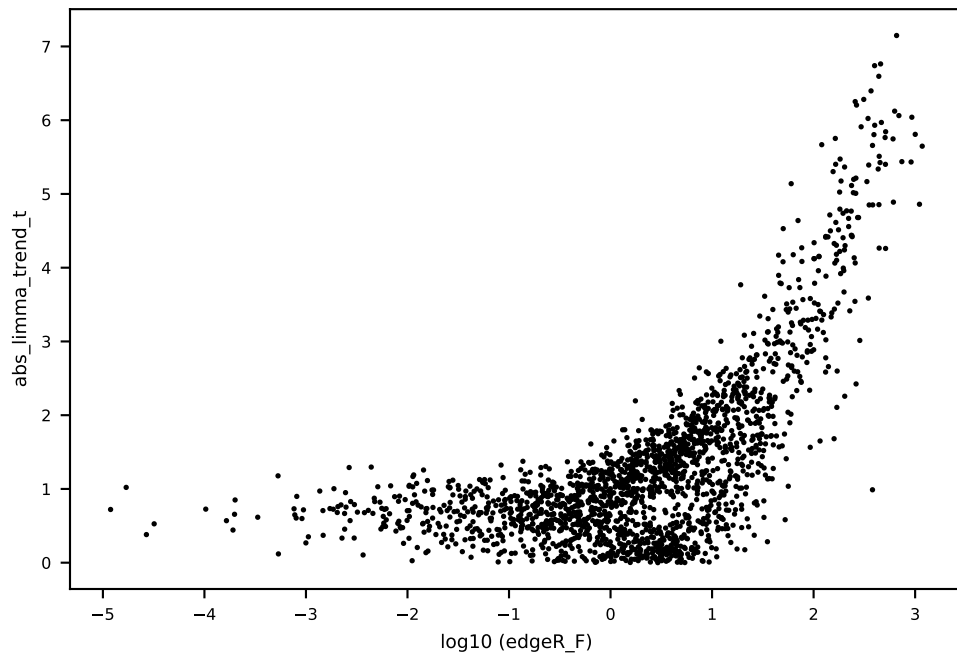
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compare pi-hat range analysis to muscat pseudo bulk
abs_limma_voom_t vs z-scores



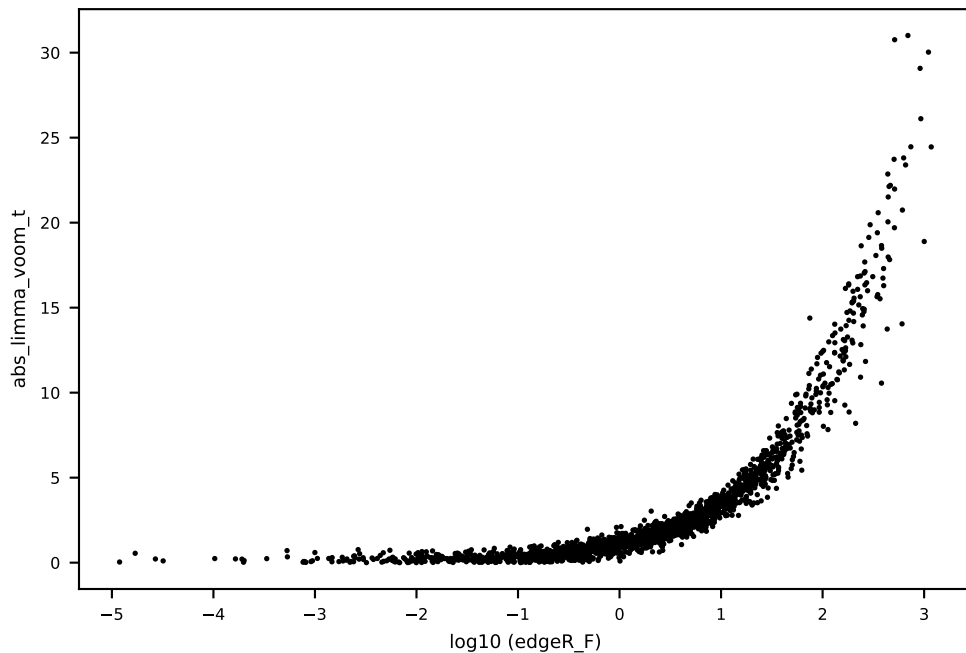
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abs_DESeq2_stat vs edgeR F statistic



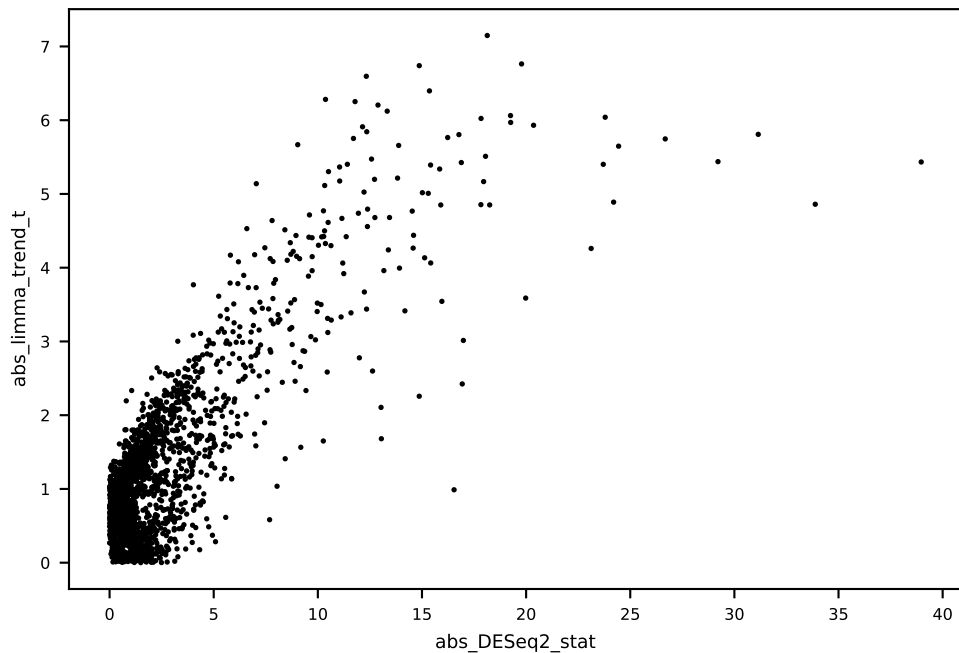
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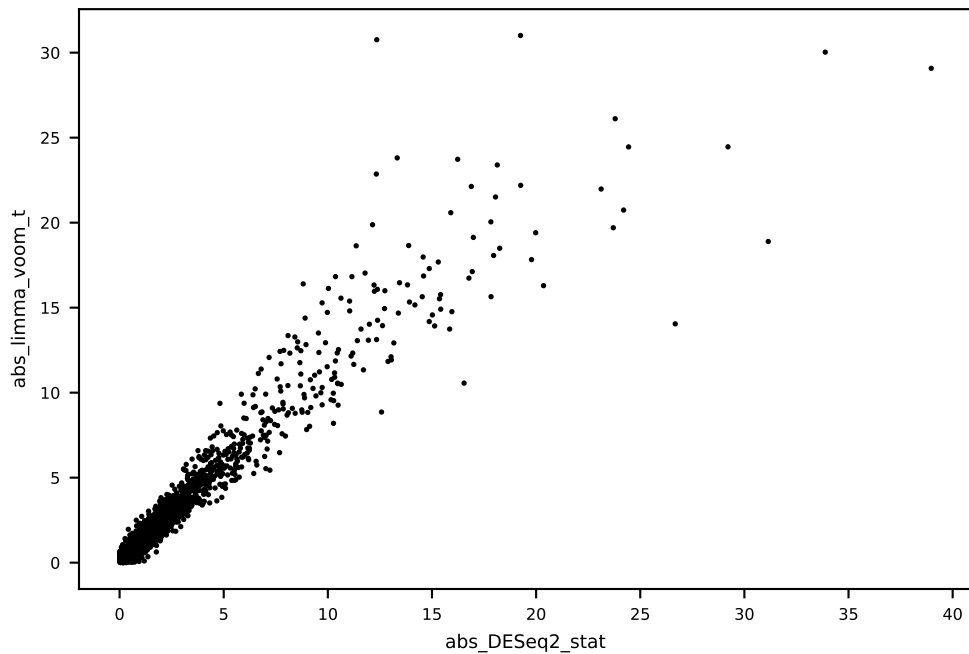
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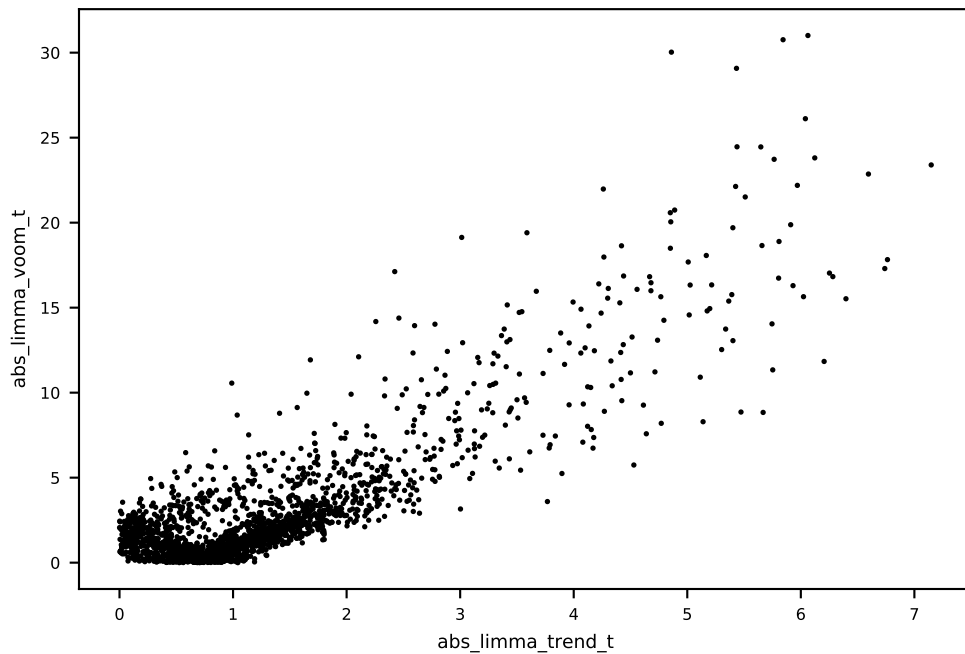
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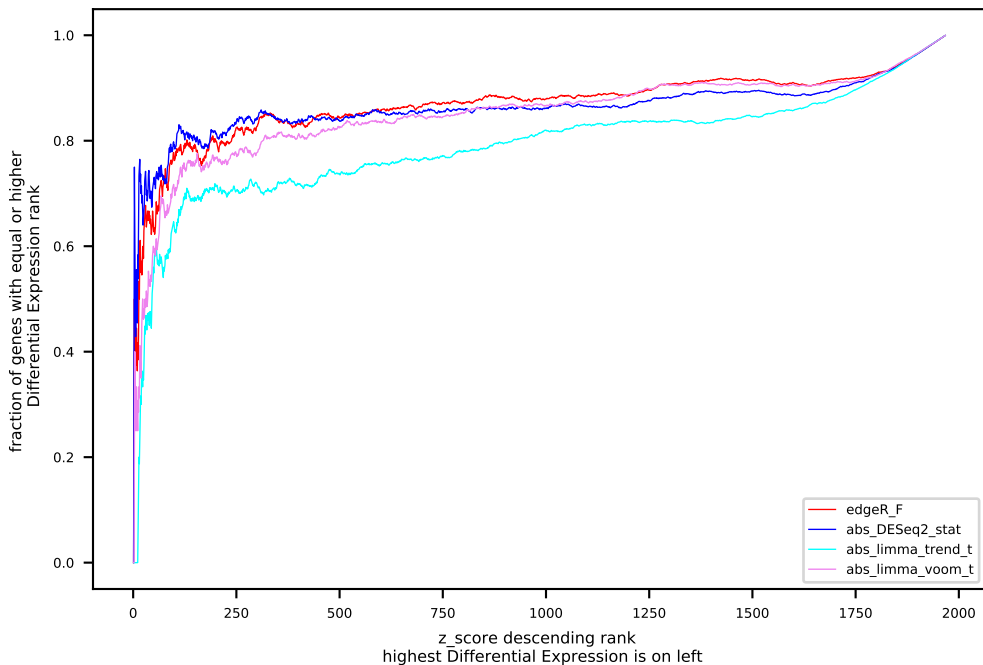
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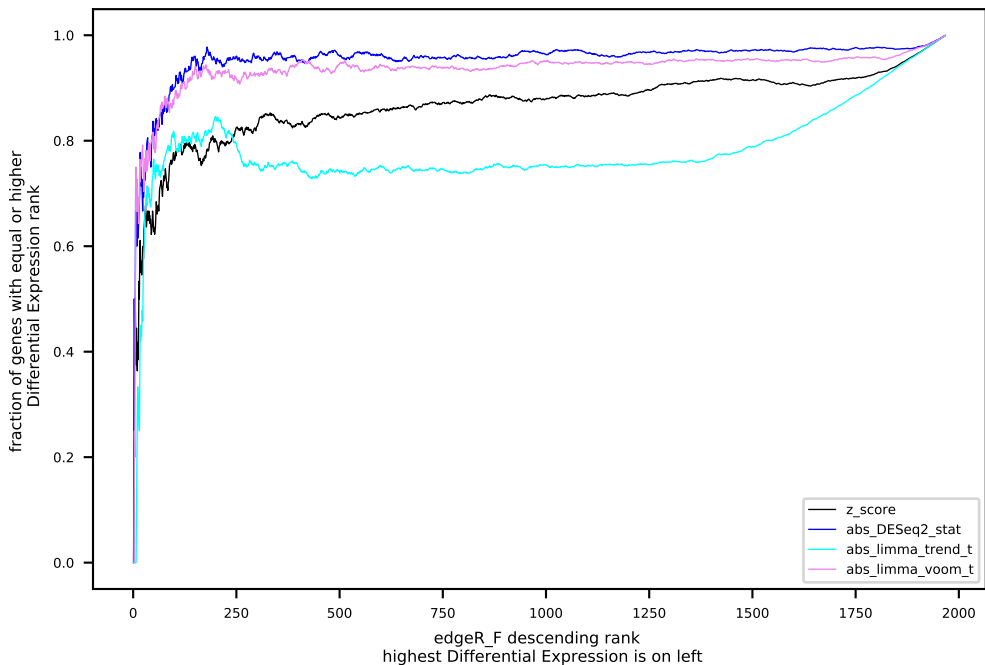
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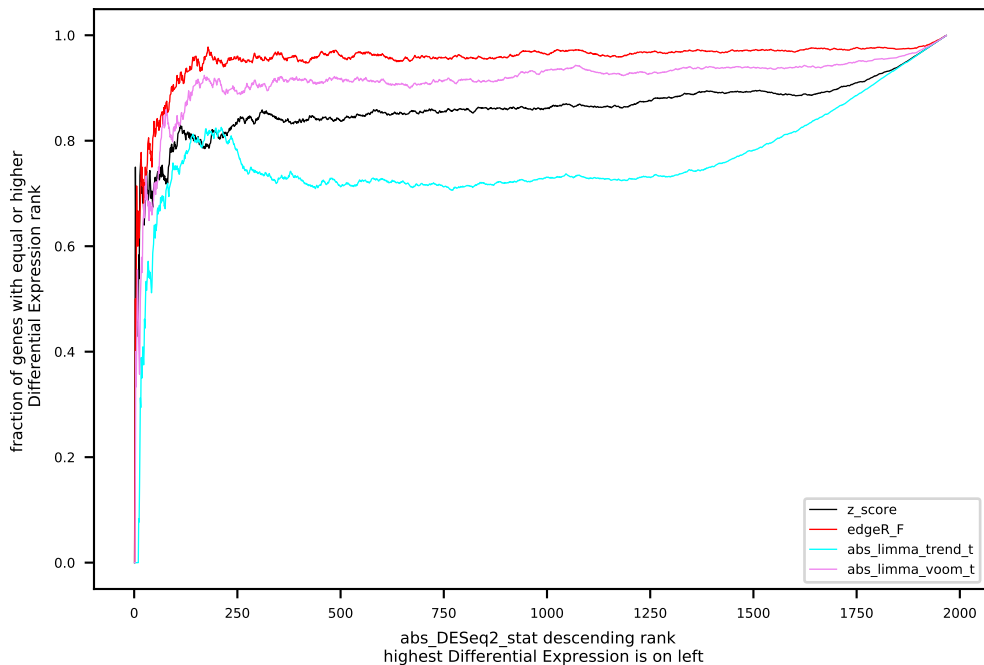
Crowell_Kang18_8vs8 scRNA-seq data - B cells
compare Differential Expression - use gene ranks to visualize CONSISTENCY between methods
muscat pseudo bulk statistics and z-scores from pi-hat range analysis



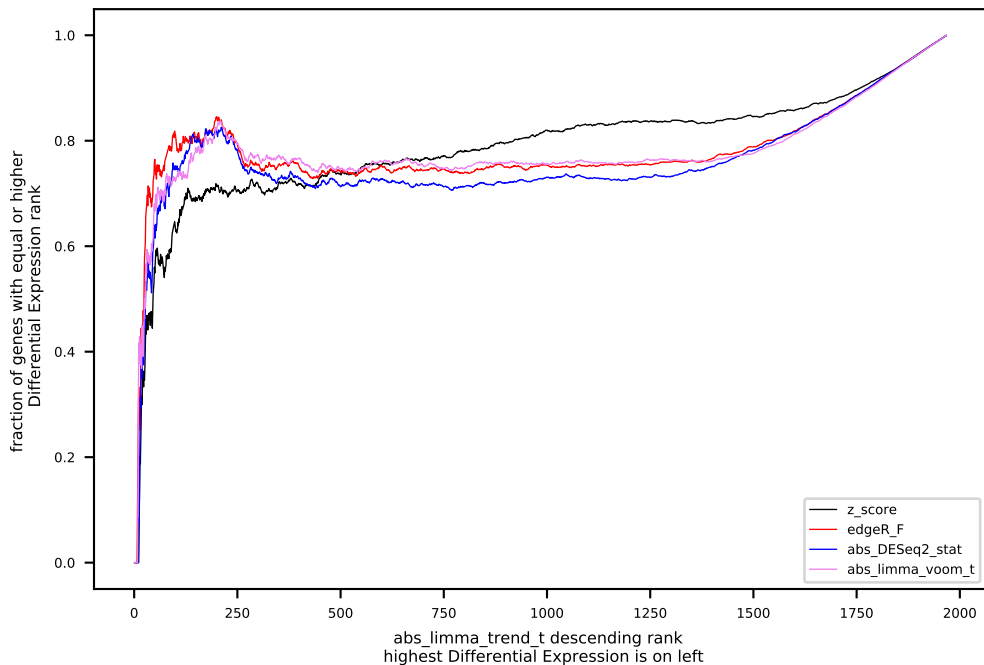
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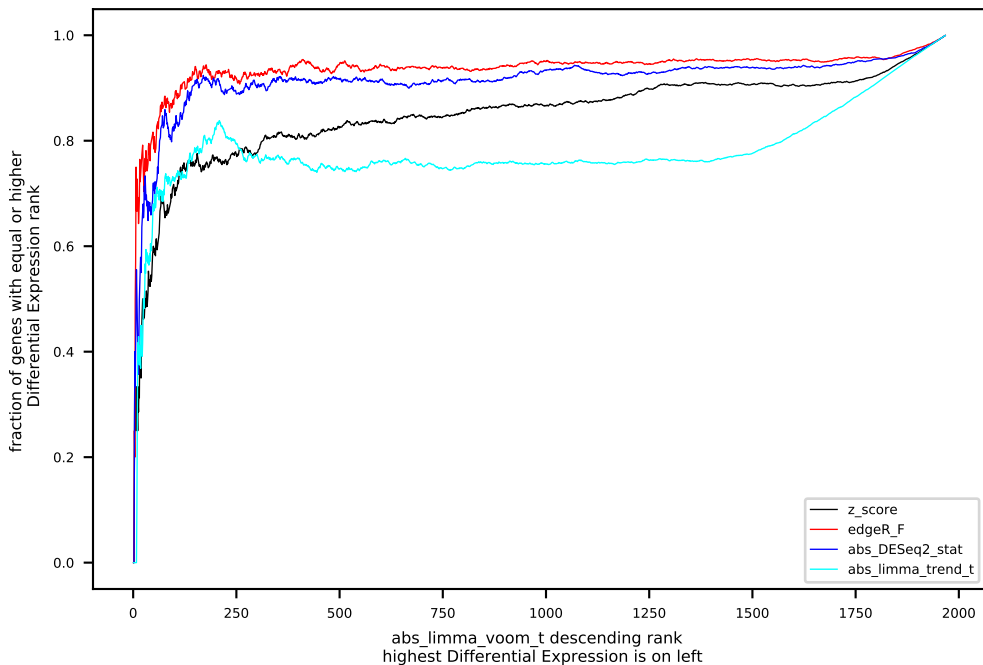
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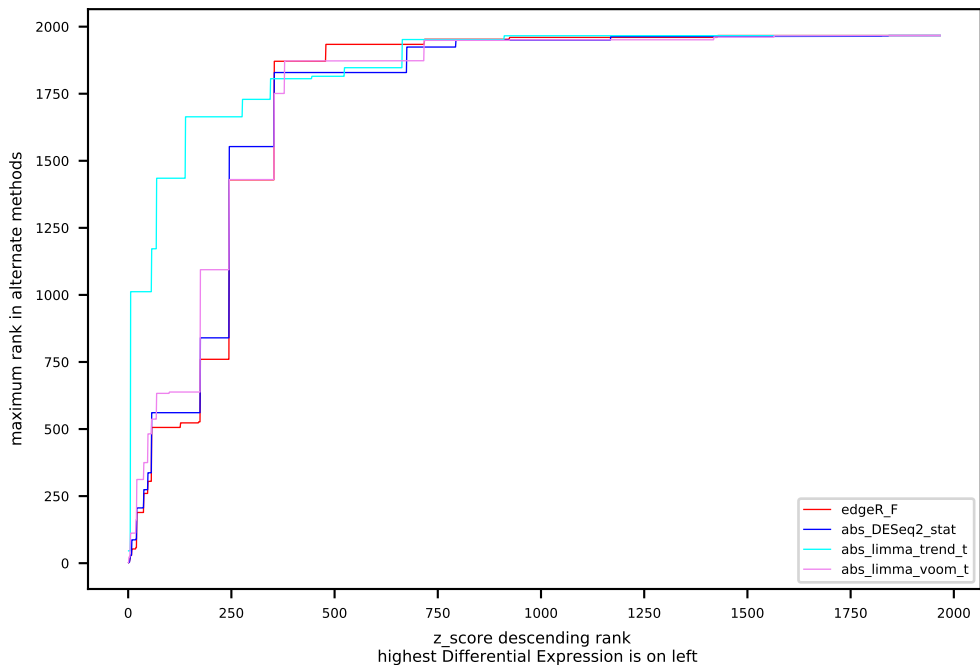
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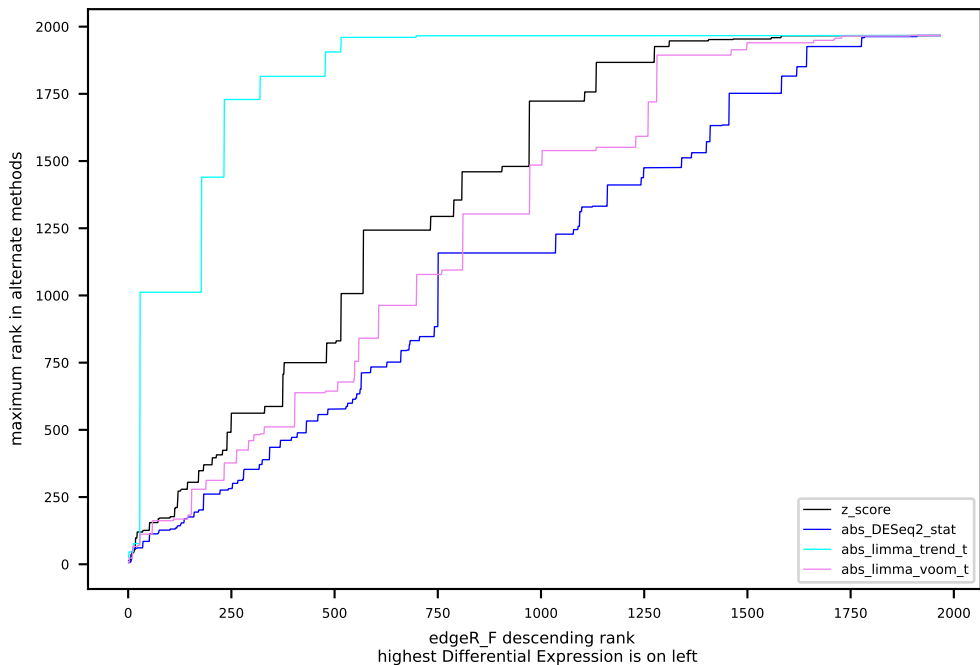
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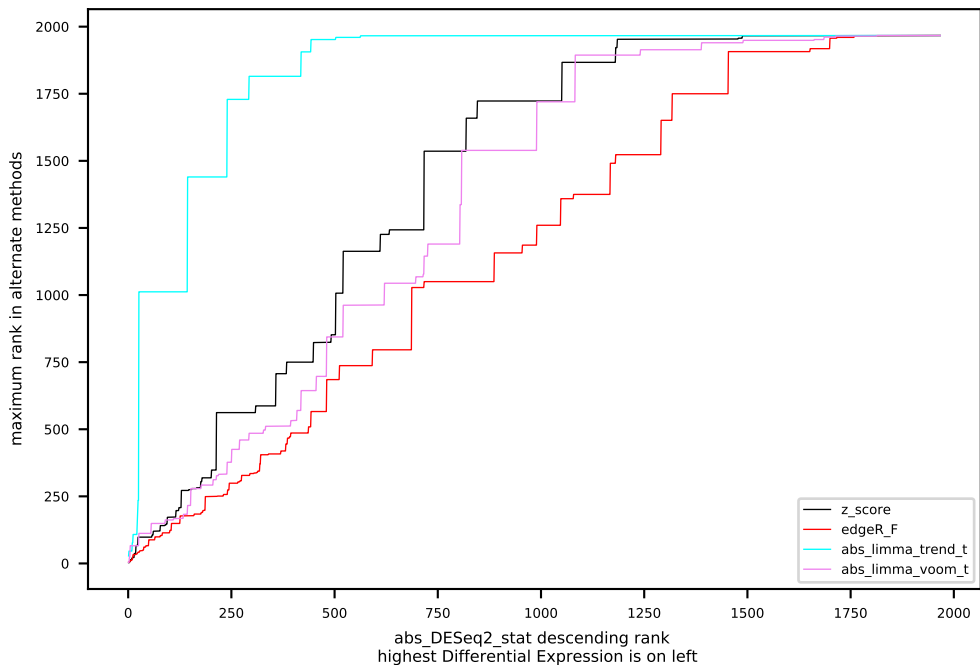
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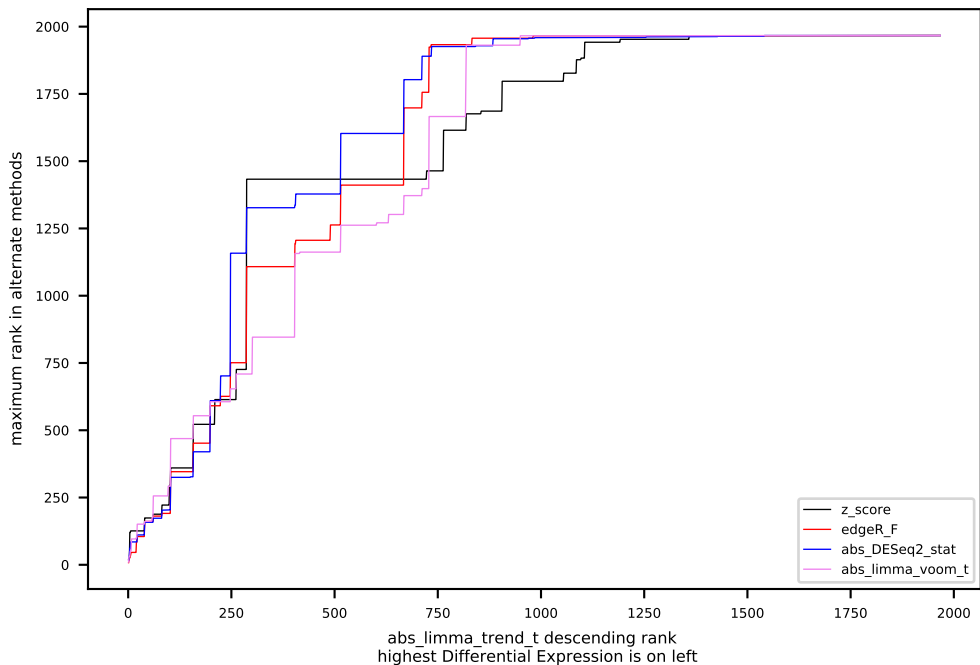
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