

# MAGeCK Comparison Report

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## Contents

<b>1</b>	<b>Summary</b>	<b>1</b>
<b>2</b>	<b>Comparison results of HL60.final,KBM7.final vs HL60.initial,KBM7.initial neg.</b>	<b>2</b>
<b>3</b>	<b>Comparison results of HL60.final,KBM7.final vs HL60.initial,KBM7.initial pos.</b>	<b>6</b>

## 1 Summary

The statistics of comparisons is as indicated in the following table.

	Comparison	Genes	Selection	FDR1%	FDR5%	FDR25%
1	HL60.final,KBM7.final_vs_HL60.initial,KBM7.initial neg.	100	negative	0	2	9
2	HL60.final,KBM7.final_vs_HL60.initial,KBM7.initial pos.	100	positive	0	0	1

Table 1: Summary of comparisons

The meanings of the columns are as follows.

- **Comparison:** The label for comparisons;
- **Genes:** The number of genes in the library;
- **Selection:** The direction of selection, either positive selection or negative selection;
- **FDR1%:** The number of genes with  $\text{FDR} < 1\%$ ;
- **FDR5%:** The number of genes with  $\text{FDR} < 5\%$ ;
- **FDR25%:** The number of genes with  $\text{FDR} < 25\%$ ;

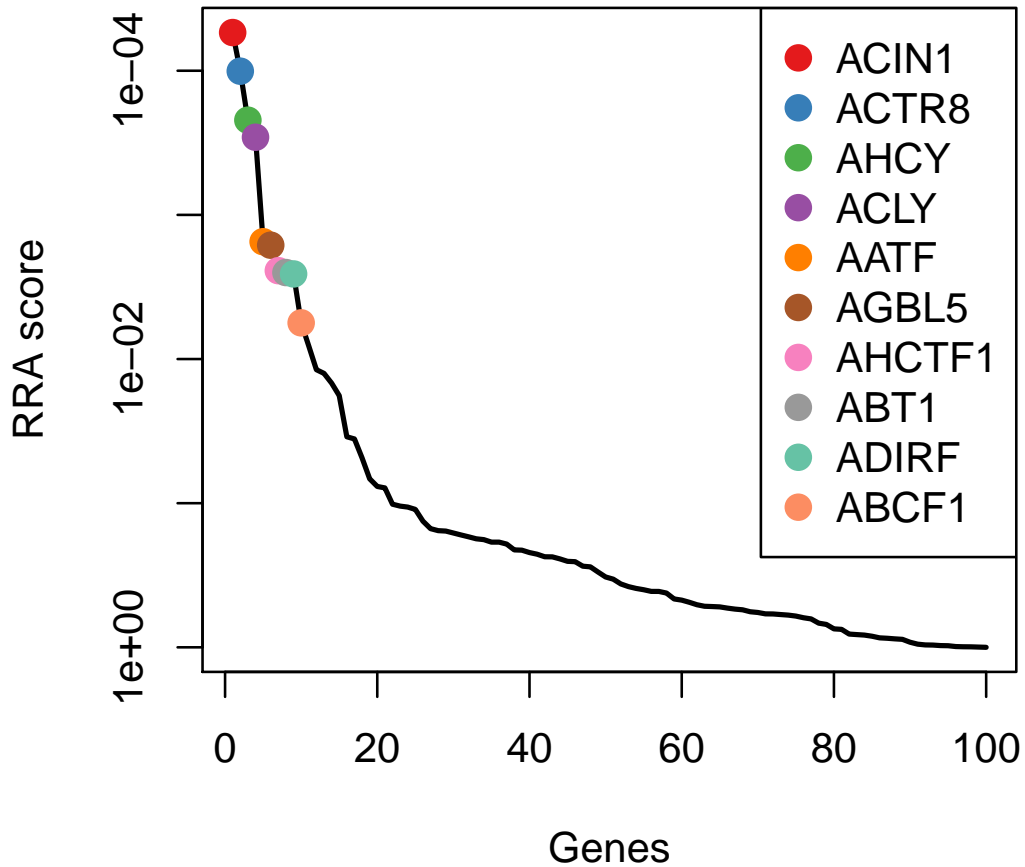
The following figures show:

- Individual sgRNA read counts of selected genes in selected samples;
- The distribution of RRA scores and p values of all genes; and
- The RRA scores and p values of selected genes.

## 2 Comparison results of HL60.final,KBM7.final vs HL60.initial,KBM7.initial neg.

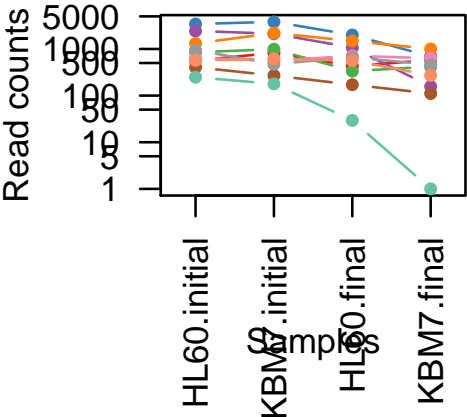
The following figure shows the distribution of RRA score in the comparison HL60.final,KBM7.final vs HL60.initial,KBM7.initial neg., and the RRA scores of 10 genes.

**Distribution of RRA scores in  
HL60.final,KBM7.final\_vs\_HL60.initial,KBM7.initial neg.**

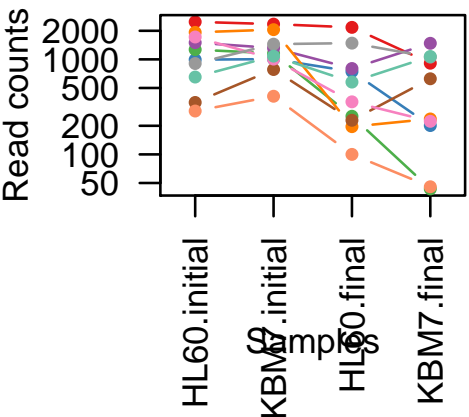


The following figures show the distribution of sgRNA read counts (normalized) of selected genes in selected samples.

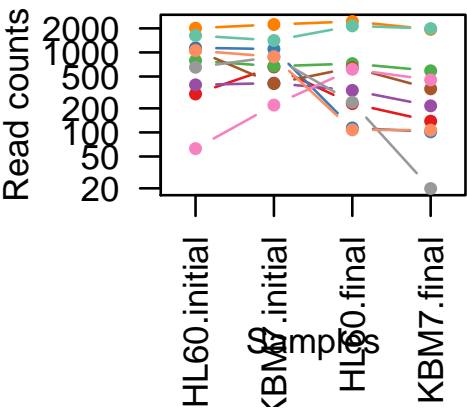
**sgRNAs in ACIN1**



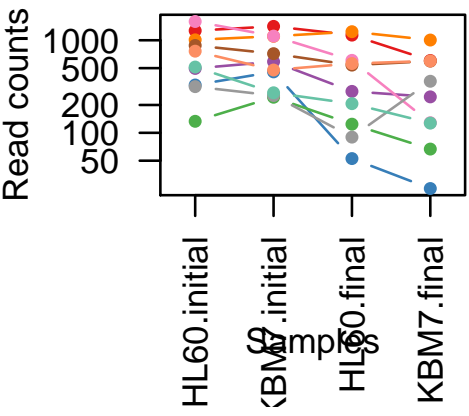
**sgRNAs in ACTR8**



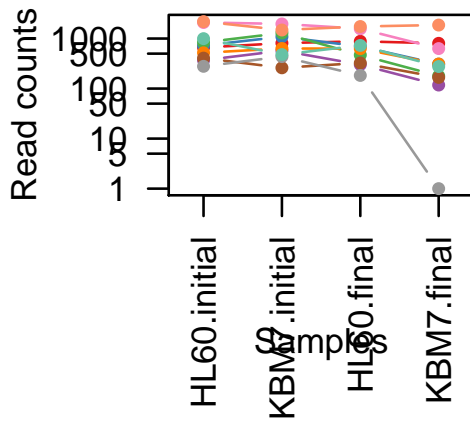
**sgRNAs in AHCY**



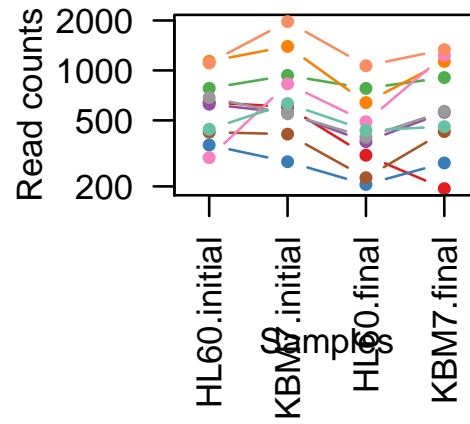
**sgRNAs in ACLY**



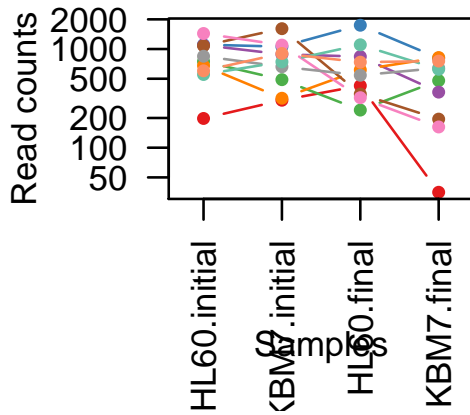
**sgRNAs in AATF**



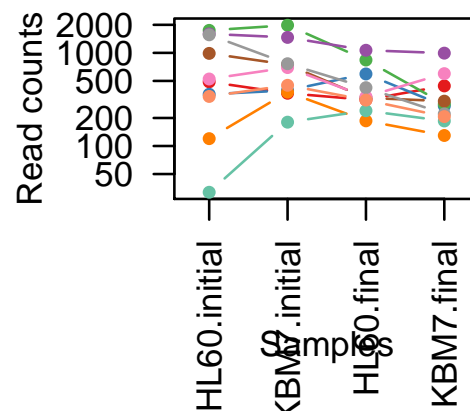
**sgRNAs in AGL5**



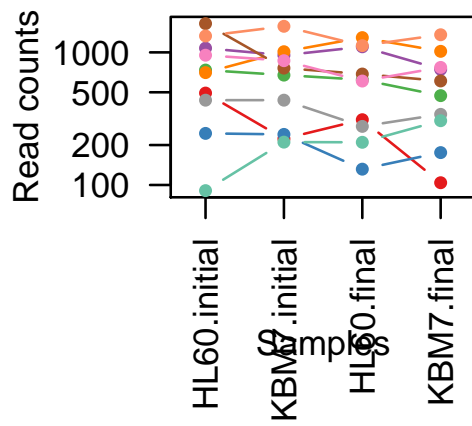
**sgRNAs in AHCTF1**



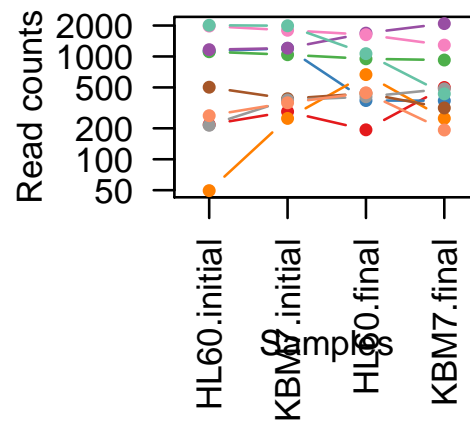
**sgRNAs in ABT1**



## sgRNAs in ADIRF



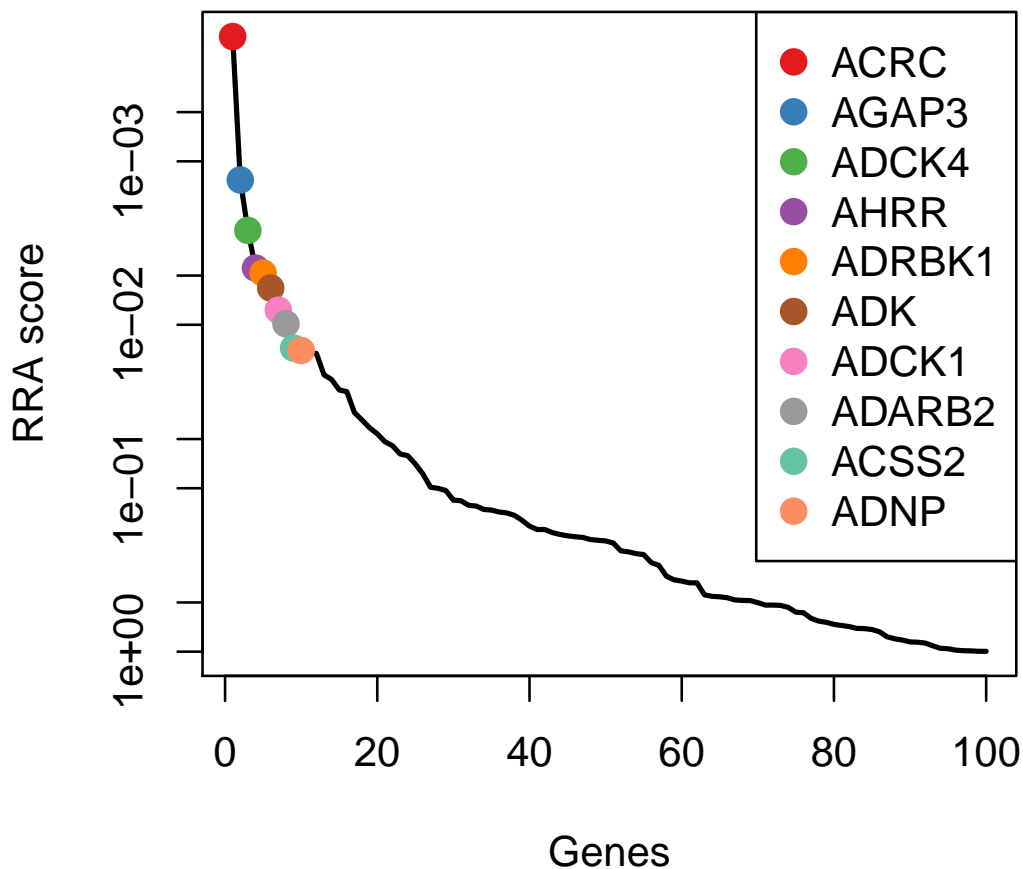
## sgRNAs in ABCF1



### 3 Comparison results of HL60.final,KBM7.final vs HL60.initial,KBM7.initial pos.

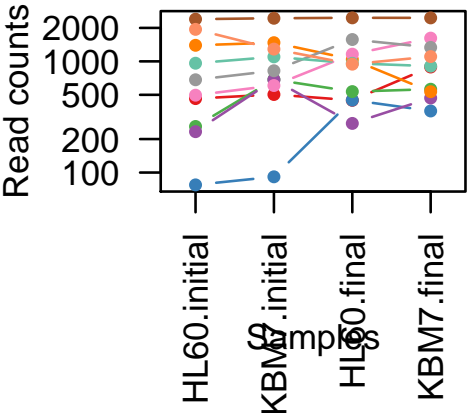
The following figure shows the distribution of RRA score in the comparison HL60.final,KBM7.final vs HL60.initial,KBM7.initial pos., and the RRA scores of 10 genes.

**Distribution of RRA scores in  
HL60.final,KBM7.final\_vs\_HL60.initial,KBM7.initial**

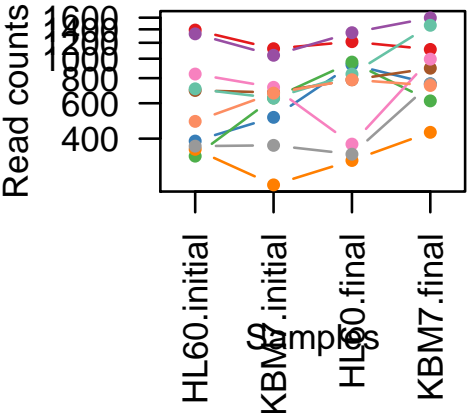


The following figures show the distribution of sgRNA read counts (normalized) of selected genes in selected samples.

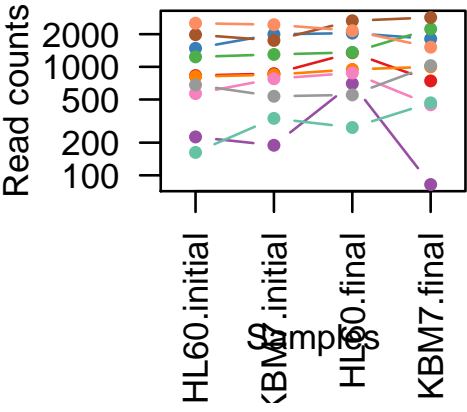
**sgRNAs in ACRC**



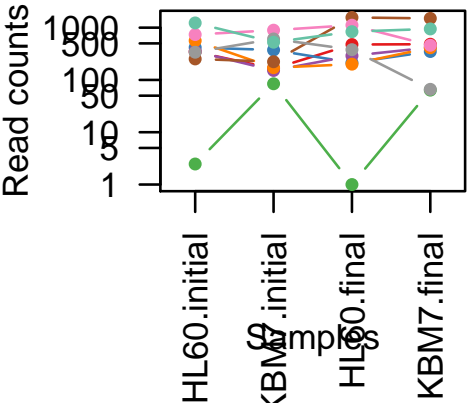
**sgRNAs in AGAP3**



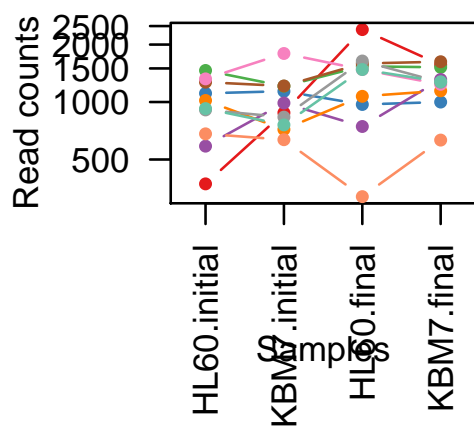
**sgRNAs in ADCK4**



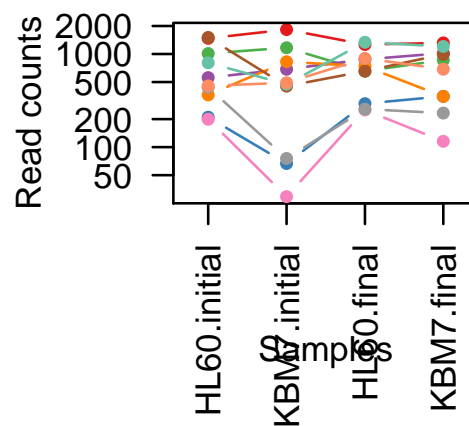
**sgRNAs in AHRR**



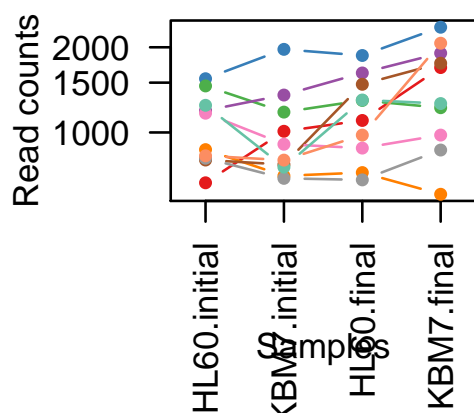
**sgRNAs in ADRBK1**



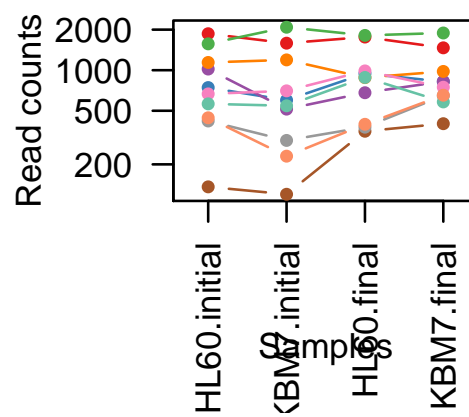
**sgRNAs in ADK**



**sgRNAs in ADCK1**

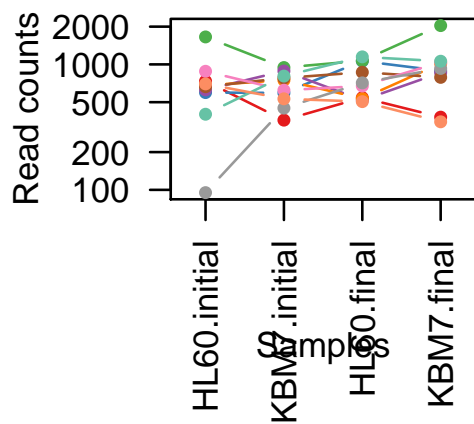


**sgRNAs in ADARB2**





## sgRNAs in ACSS2



## sgRNAs in ADNP

