RRA scores in \n HL60.final,KBM7.final_vs_HL60.in

Y-axis: RRA score (1e−04, 1e−02, 1e+00)

X-axis: Genes (0, 20, 40, 60, 80, 100)

Legend:
- ACIN1
- ACTR8
- AHCY
- ACLY
- AATF
- AGBL5
- AHCTF1
- ABT1
- ADIRF
- ABCF1

p values in \n HL60.final,KBM7.final_vs_HL60.initi

Legend:
- ACIN1
- ACTR8
- AHCY
- ACLY
- AATF
- AGBL5
- AHCTF1
- ABT1
- ADIRF
- ABCF1

Y-axis: p value (1e−03, 1e−02, 1e−01, 1e+00)

X-axis: Genes (0, 20, 40, 60, 80, 100)

sgRNAs in ACIN1

sgRNAs in ACTR8
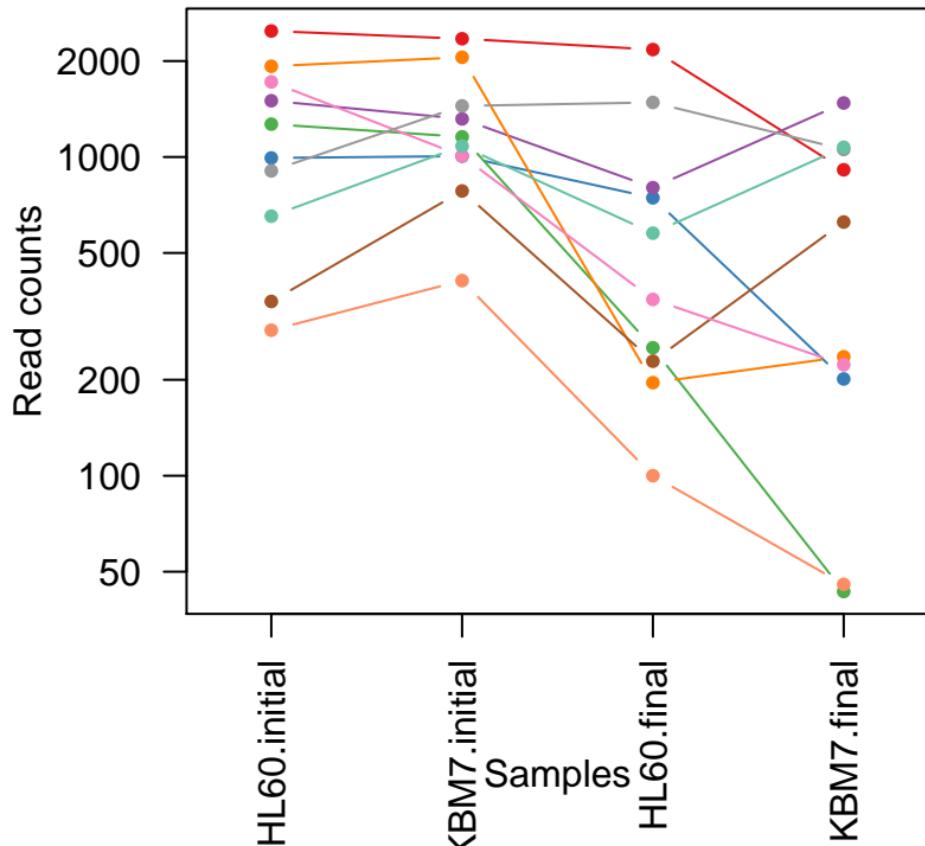
sgRNAs in AHCY
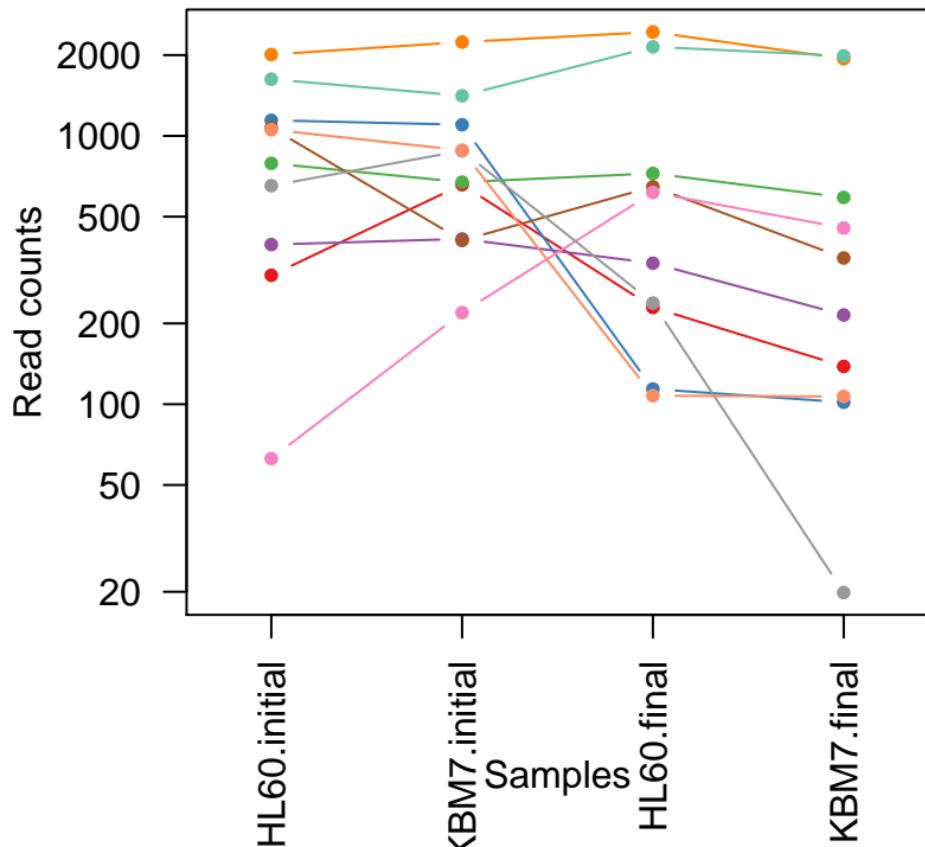
sgRNAs in ACLY
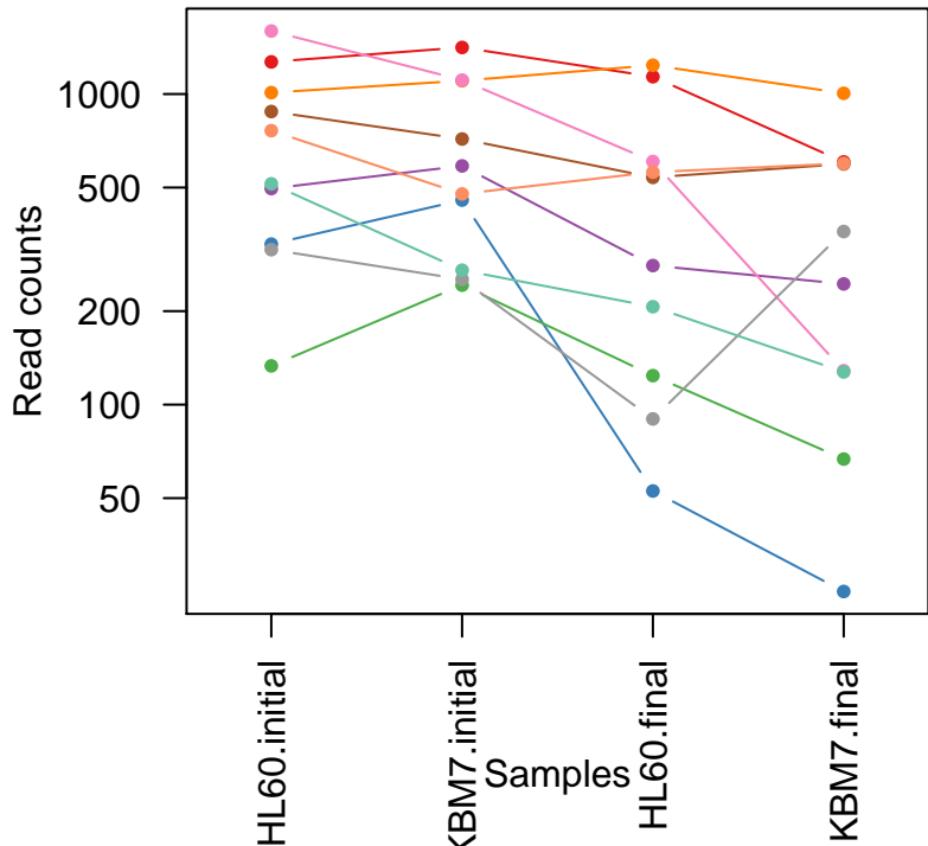
sgRNAs in AATF
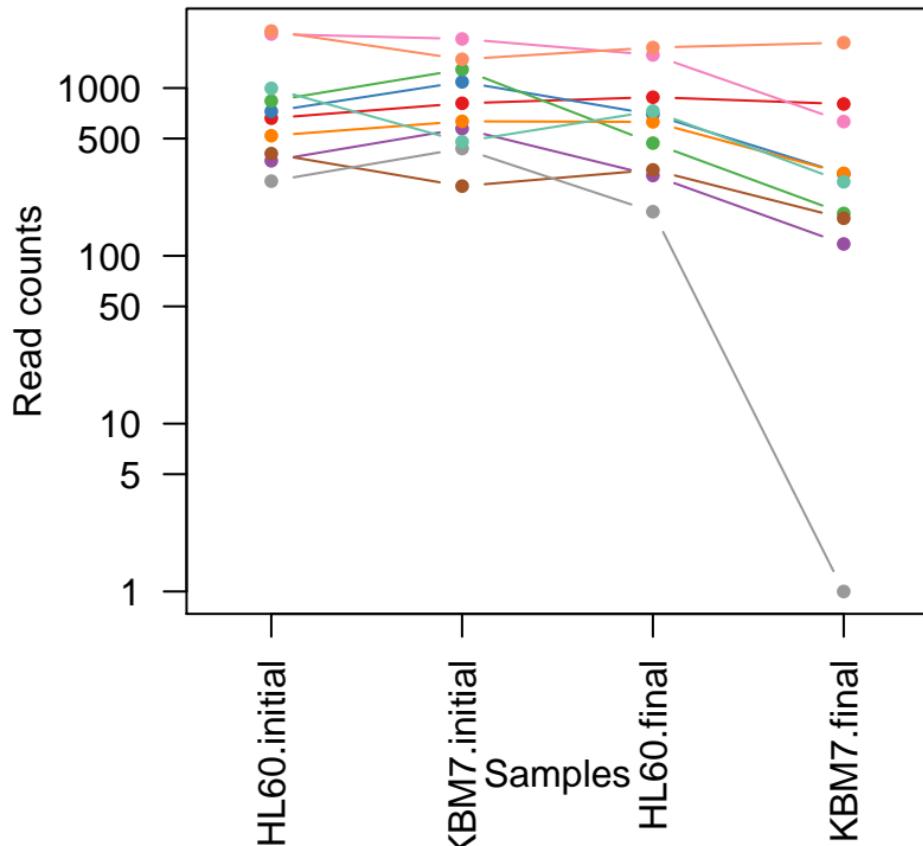
sgRNAs in AGBL5
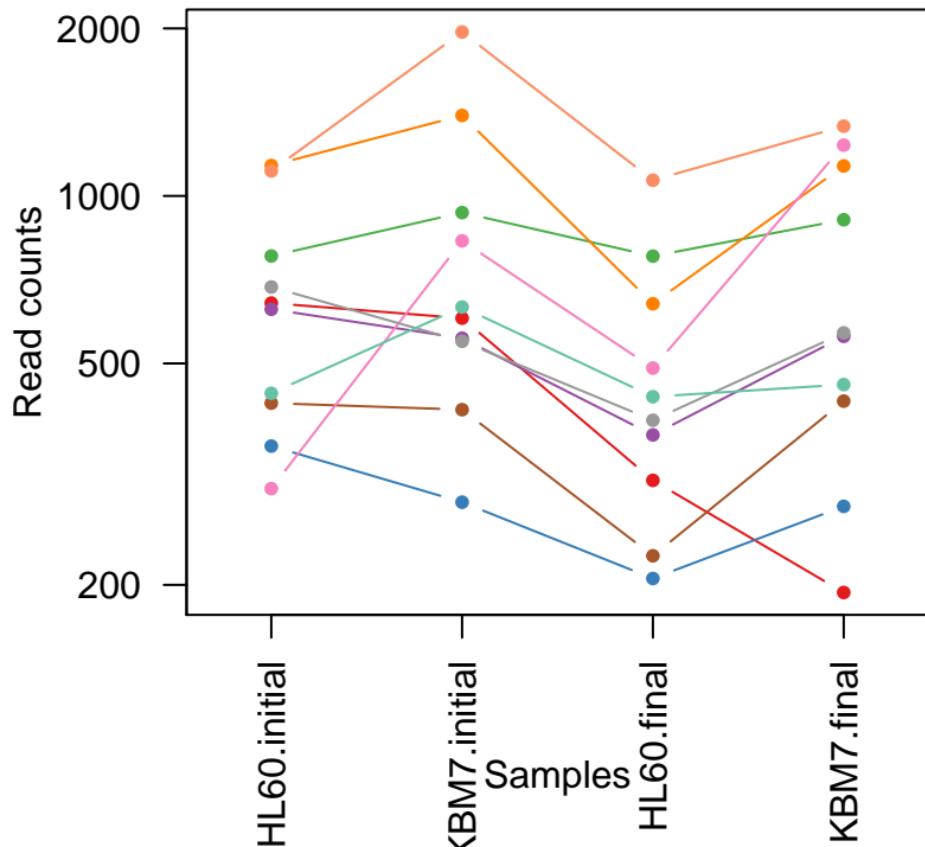
sgRNAs in AHCTF1
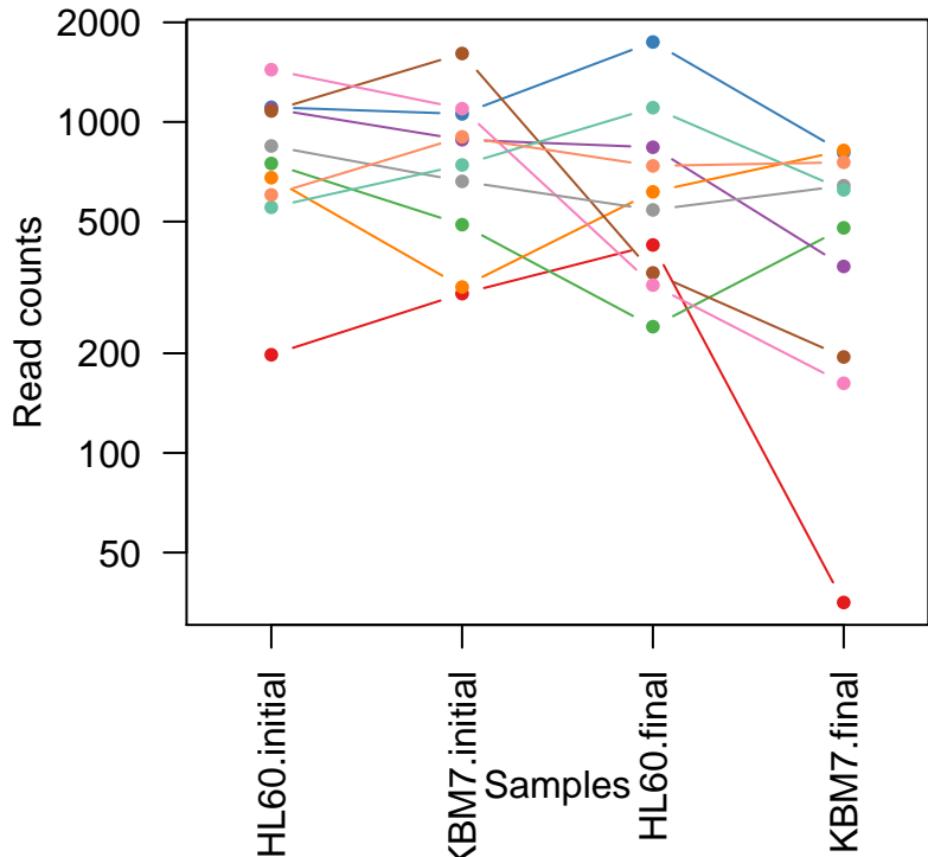
sgRNAs in ABT1
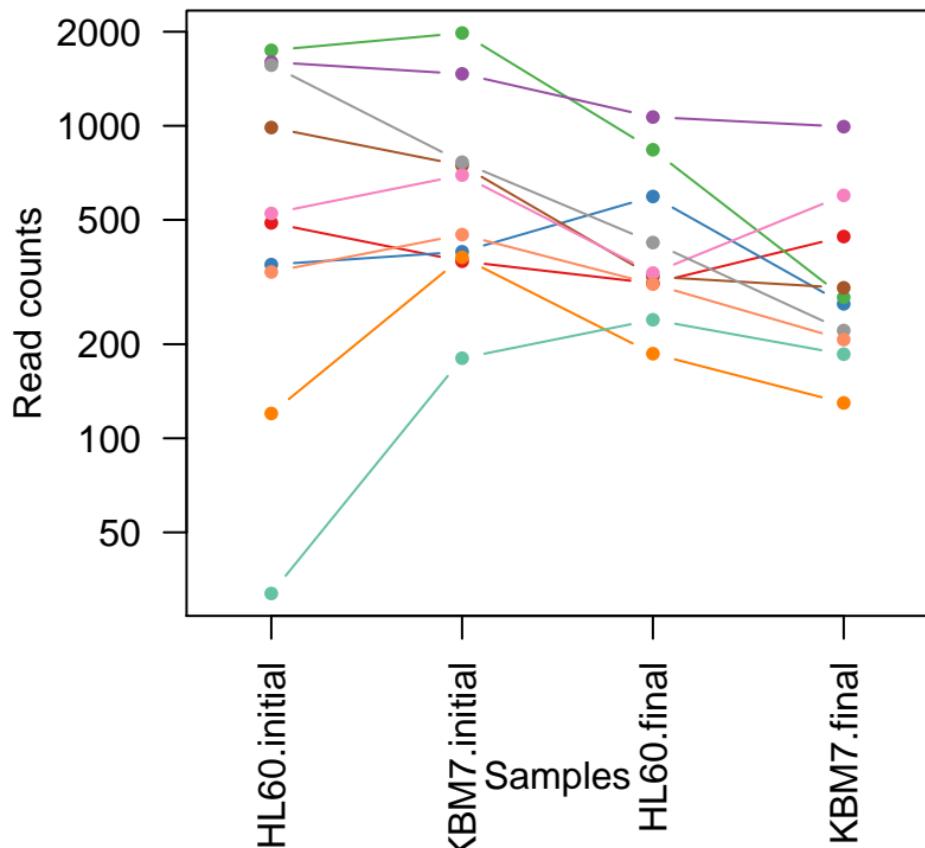
sgRNAs in ADIRF
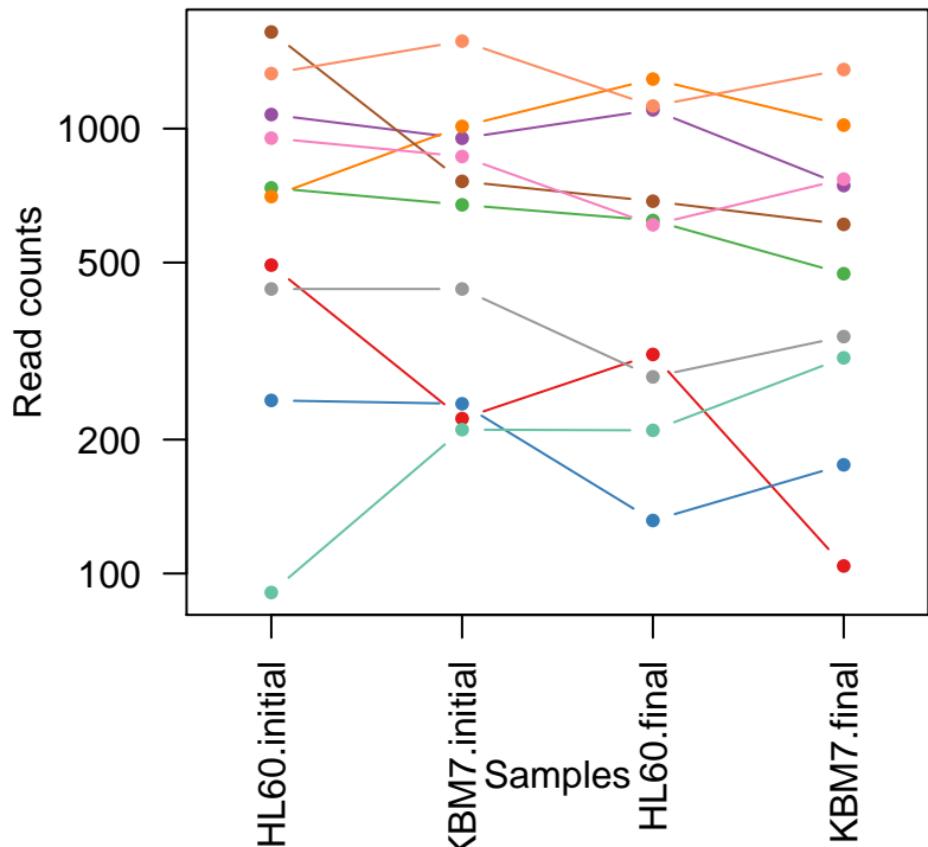
sgRNAs in ABCF1
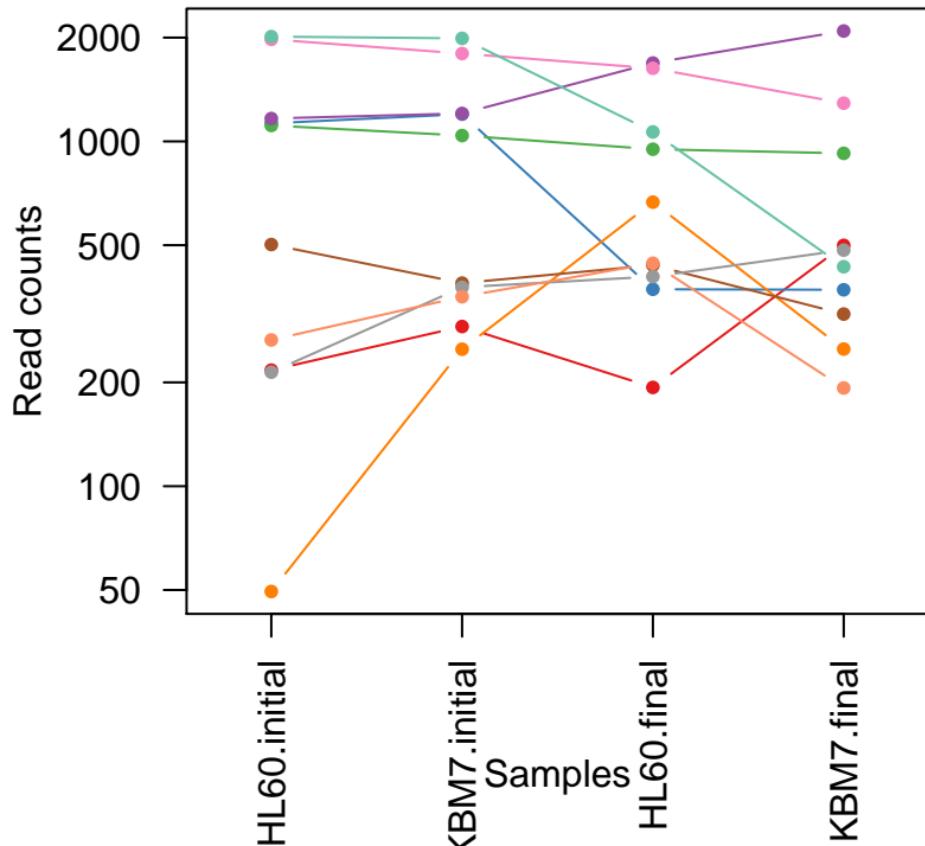
RRA scores in \n HL60.final,KBM7.final_vs_HL60.in...

Legend:
- ACRC
- AGAP3
- ADCK4
- AHRR
- ADRBK1
- ADK
- ADCK1
- ADARB2
- ACSS2
- ADNP

Y-axis: RRA score (1e-03, 1e-02, 1e-01, 1e+00)

X-axis: Genes (0, 20, 40, 60, 80, 100)

p values in \n HL60.final,KBM7.final_vs_HL60.initi

Legend:
- ACRC
- AGAP3
- ADCK4
- AHRR
- ADRBK1
- ADK
- ADCK1
- ADARB2
- ACSS2
- ADNP

Y-axis: p value

X-axis: Genes
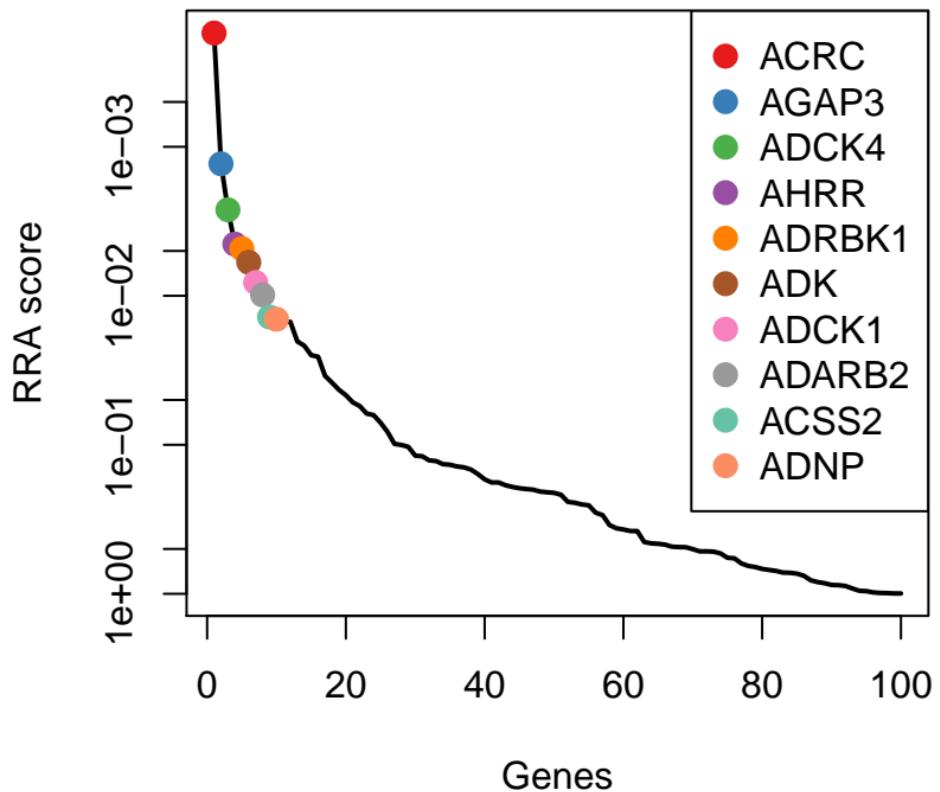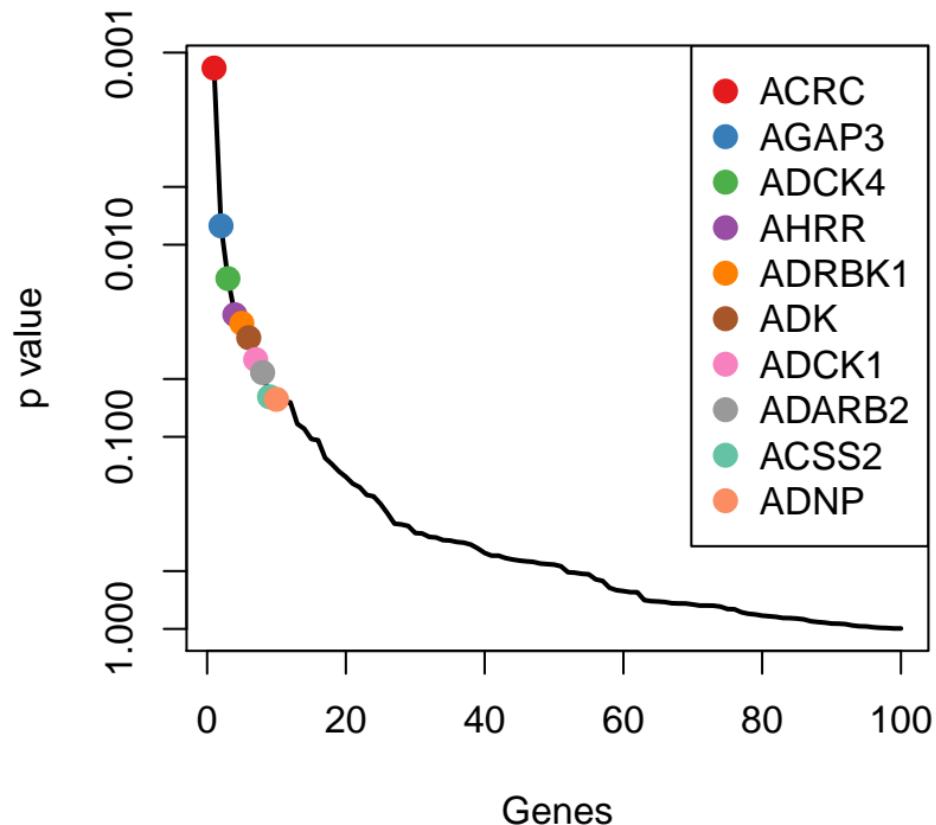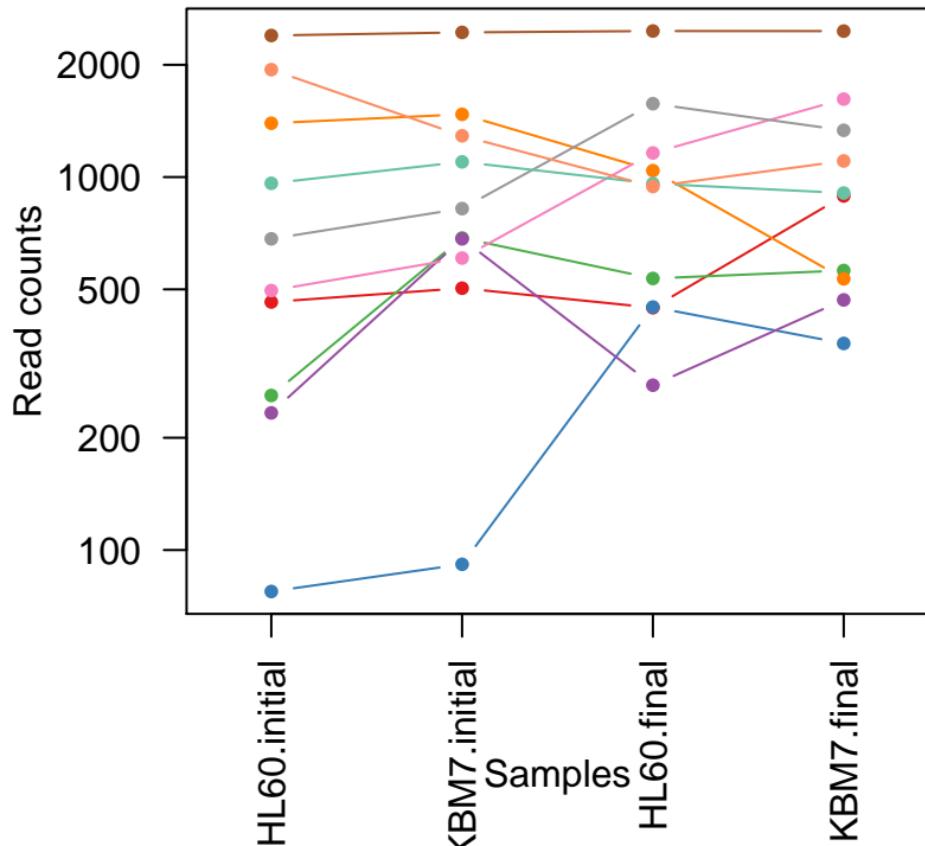
sgRNAs in ACRC

sgRNAs in AGAP3
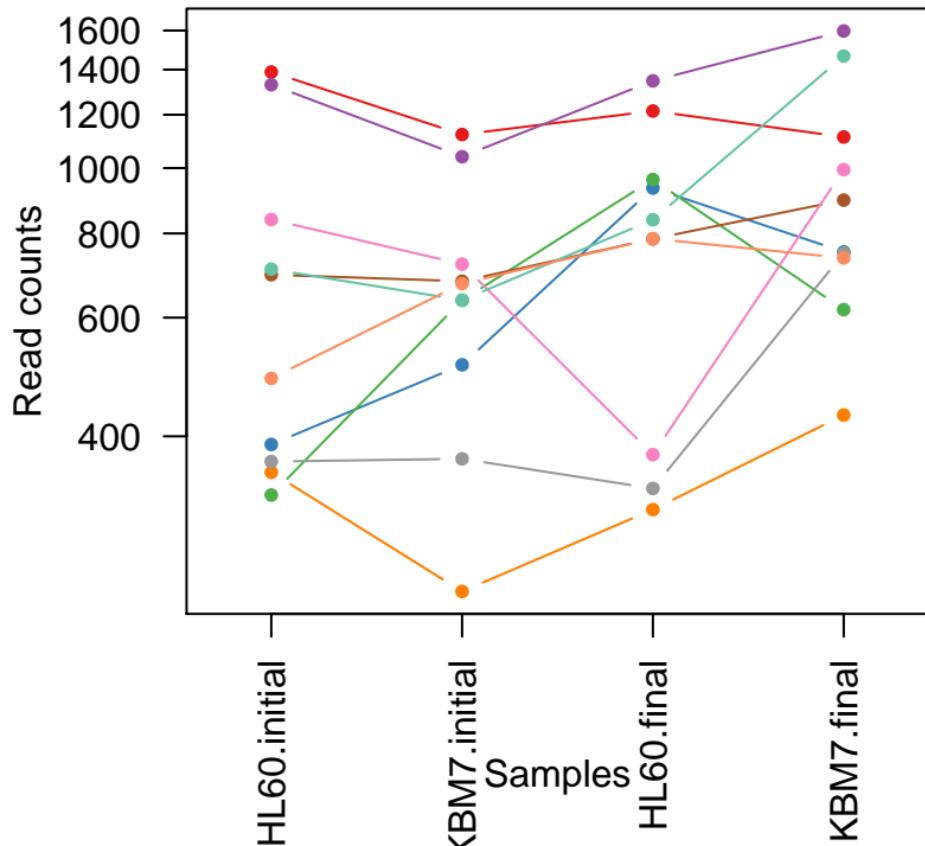
sgRNAs in ADCK4
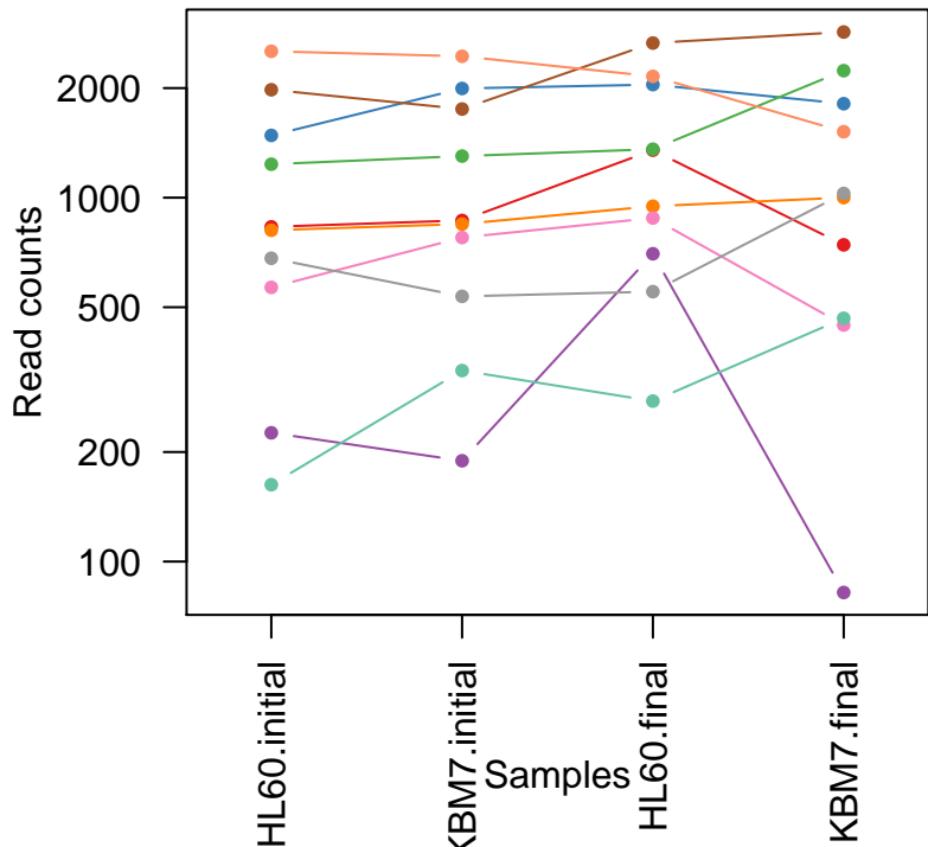
sgRNAs in AHRR
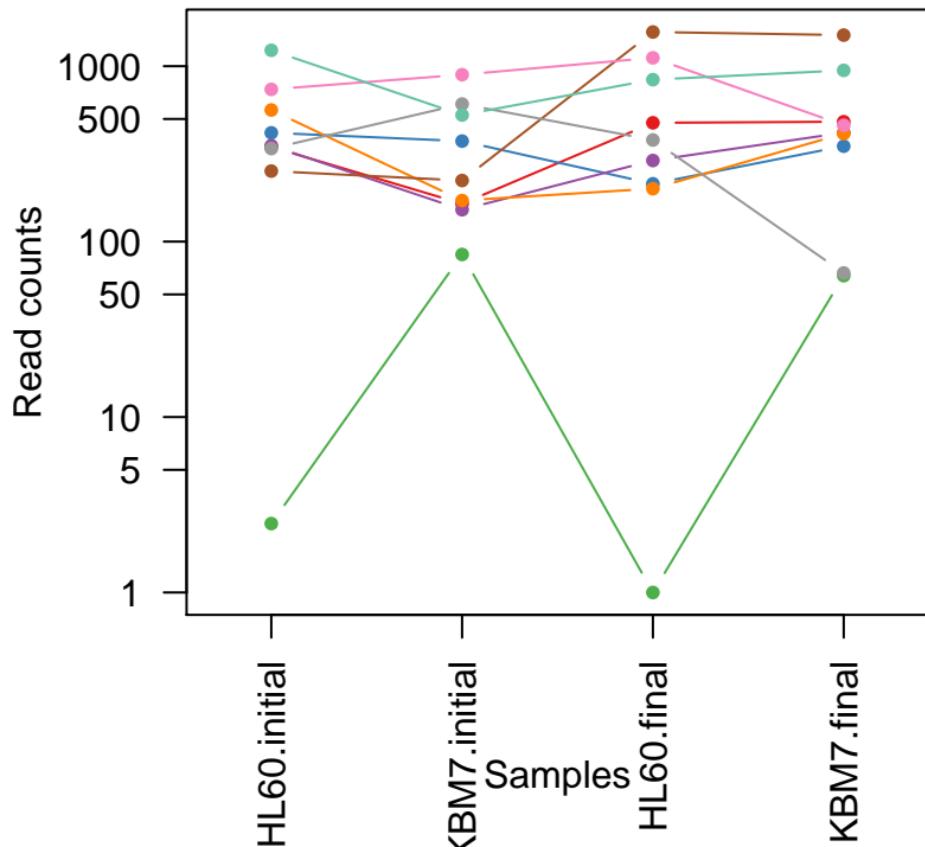
sgRNAs in ADRBK1

sgRNAs in ADK

Read counts

HL60.initial KBM7.initial HL60.final KBM7.final

Samples
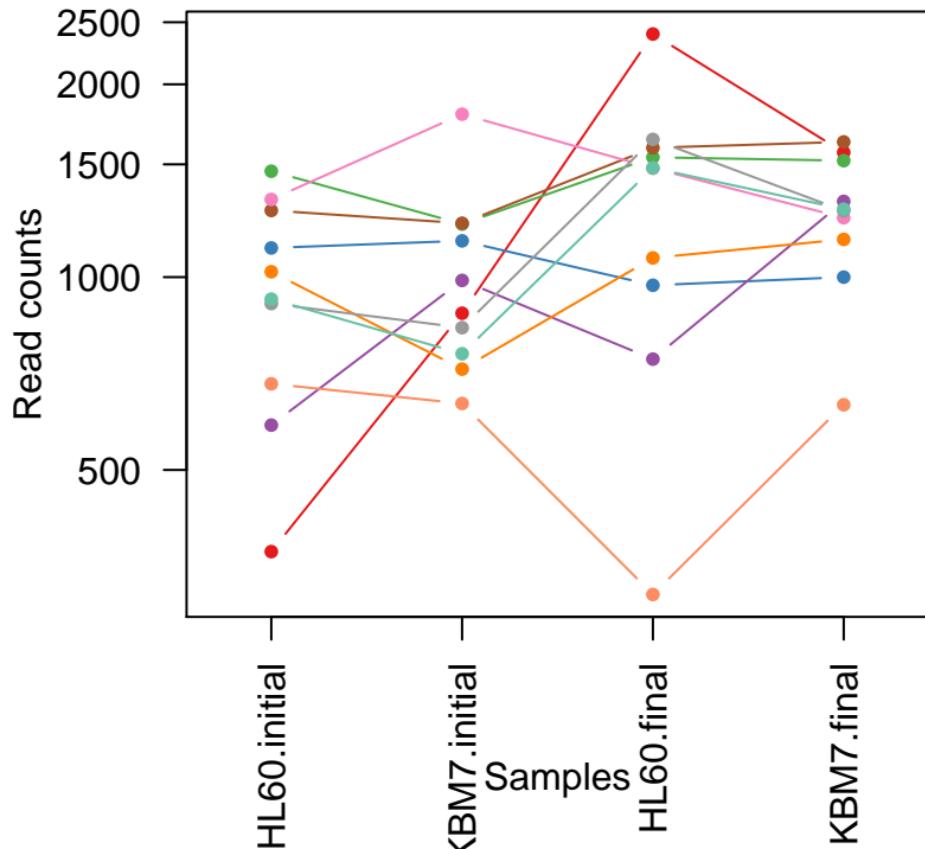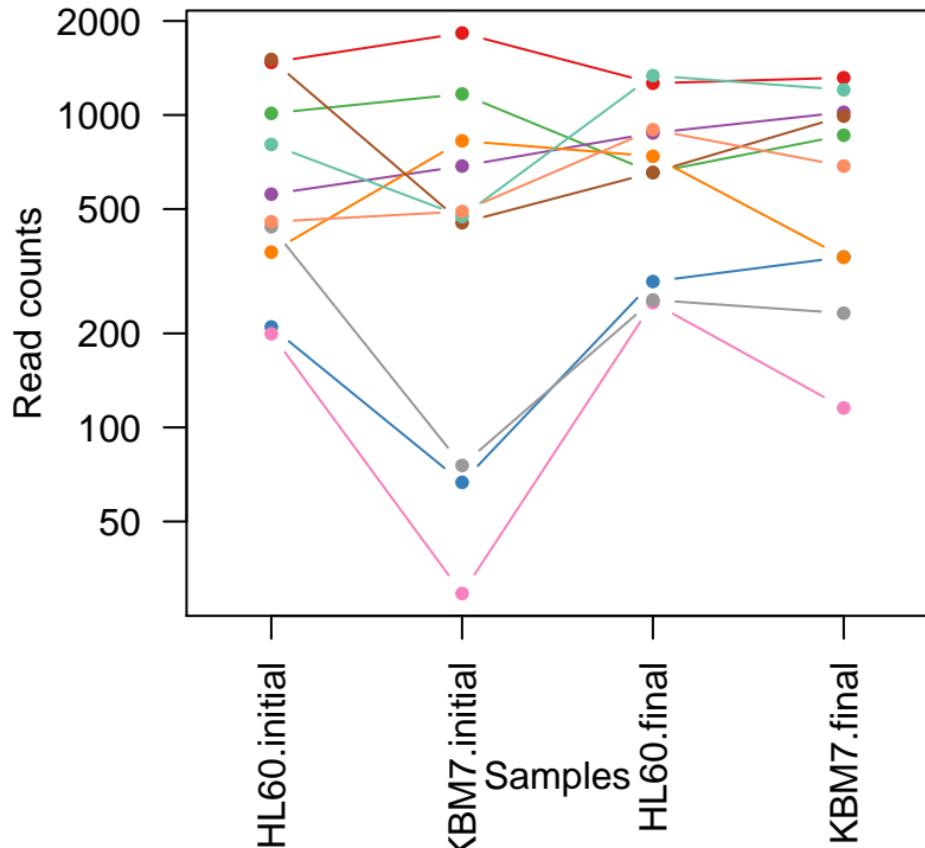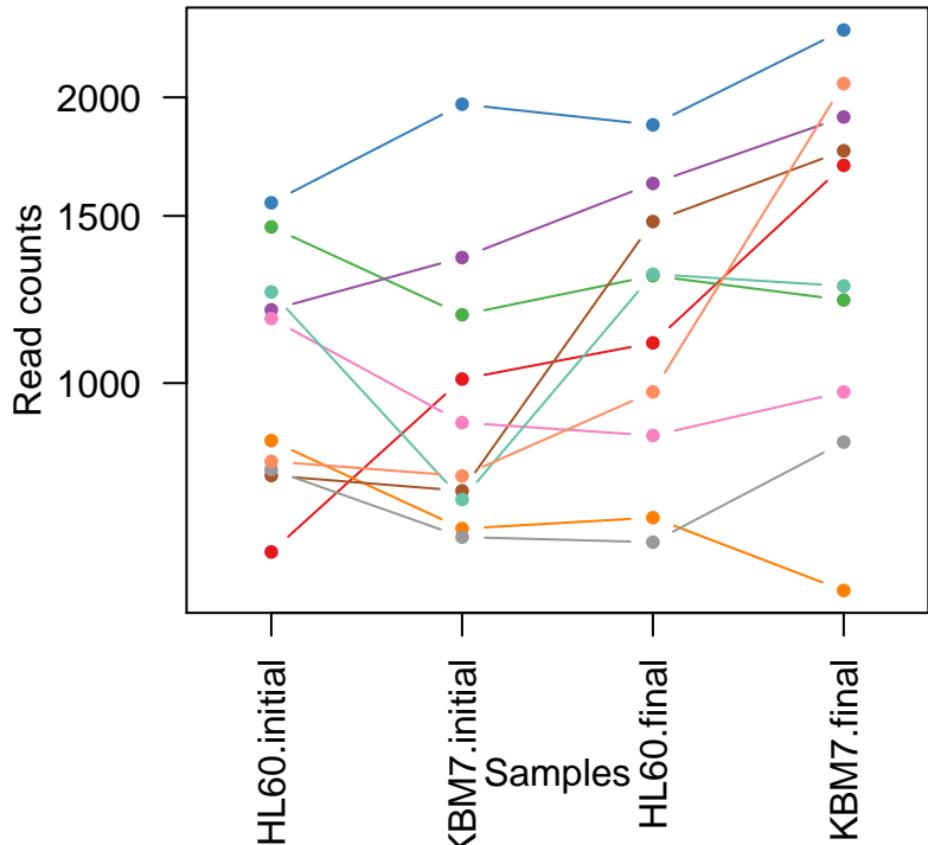
sgRNAs in ADCK1

sgRNAs in ADARB2
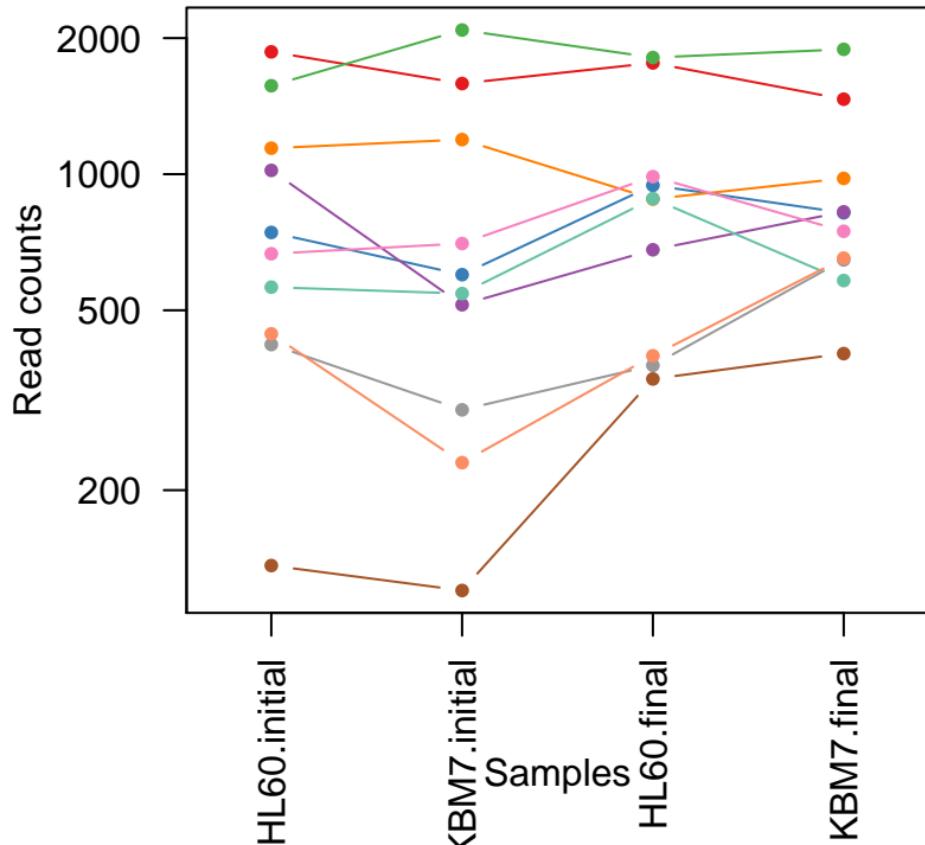
sgRNAs in ACSS2

sgRNAs in ADNP