

Score: CV_t (Local, full , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

0

10

20

30

40

50

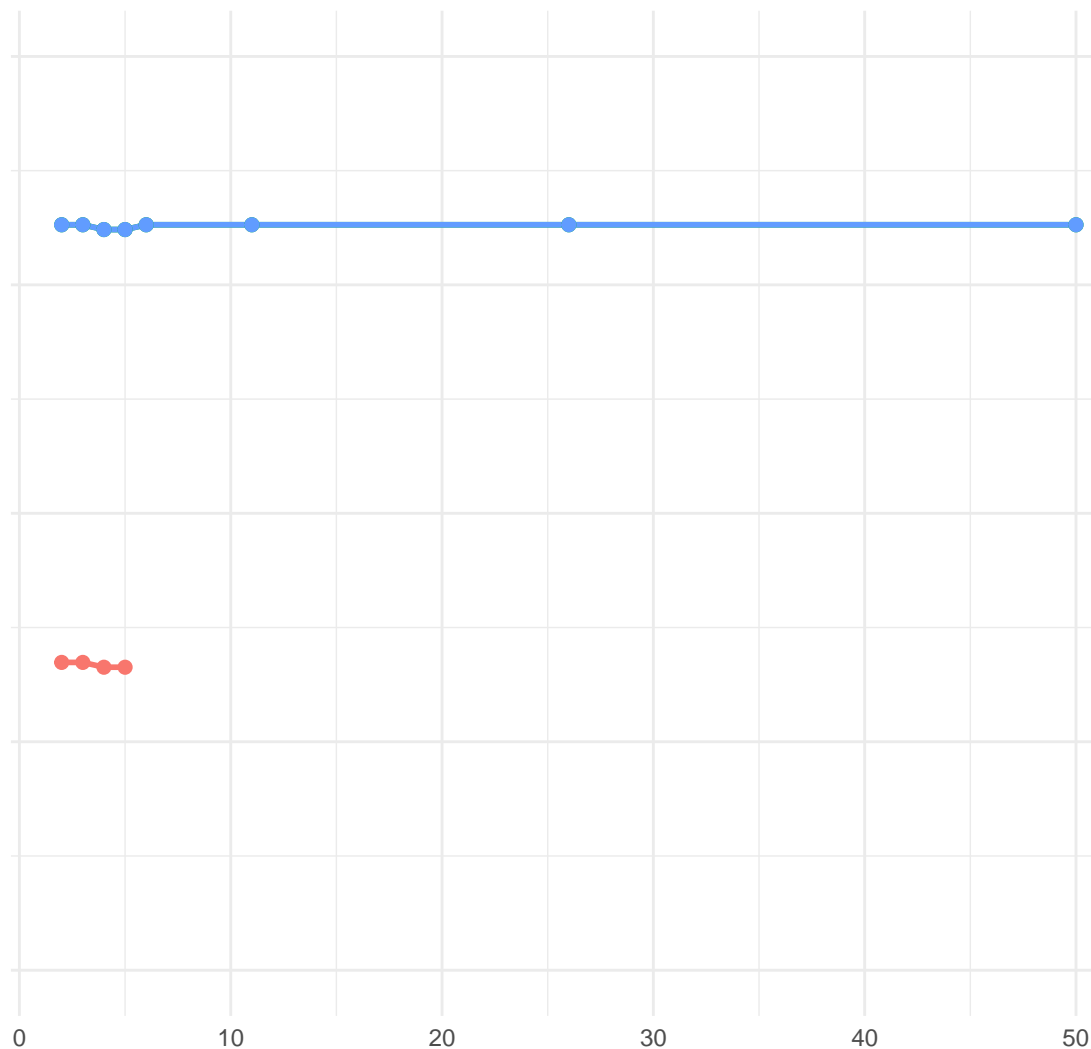
Number of Selected Samples

Summary Statistic

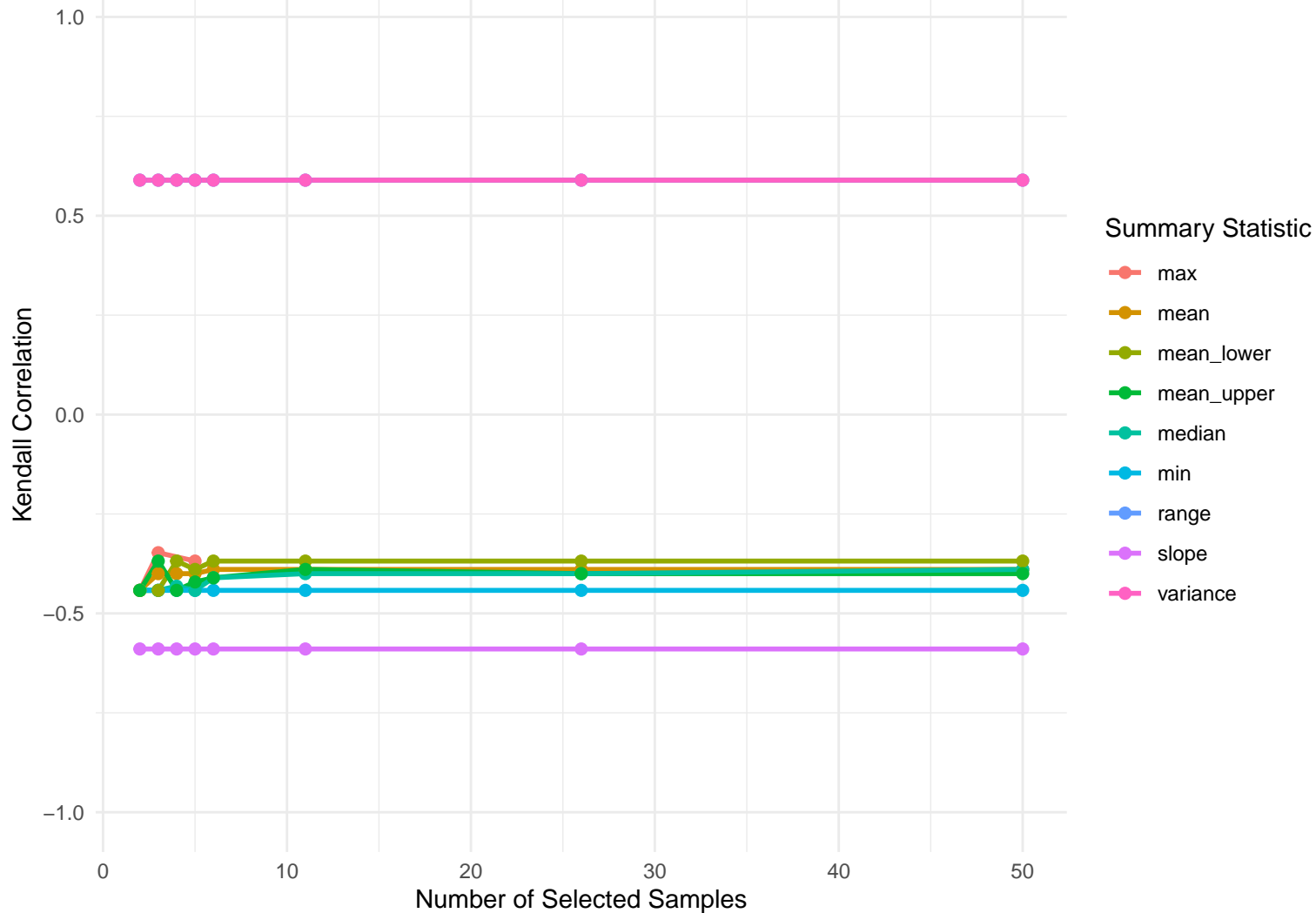
min

range

variance



Score: CV_t (Local, full , Genotype: gaussian)



Score: CV_t (Local, full , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

0

10

20

30

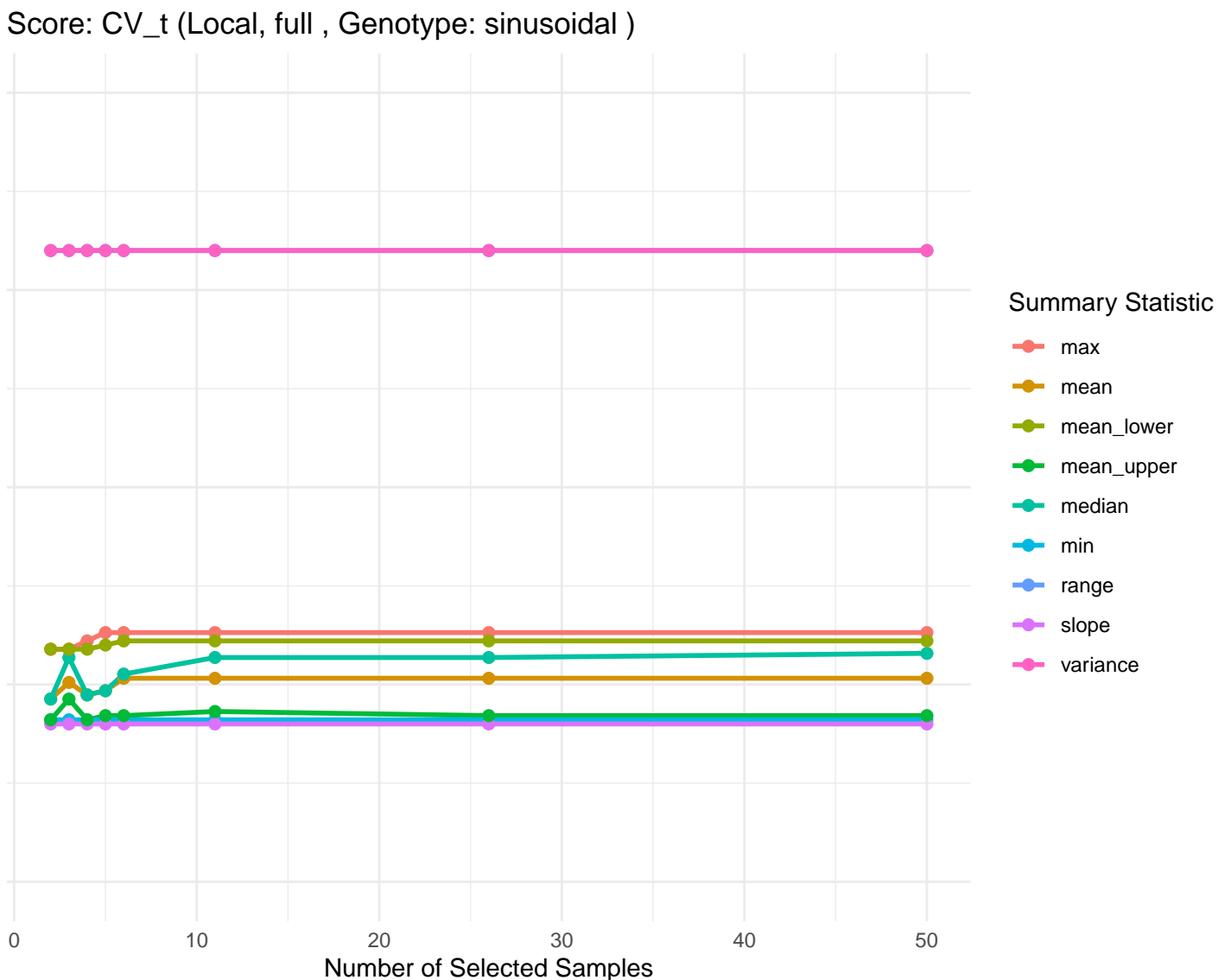
40

50

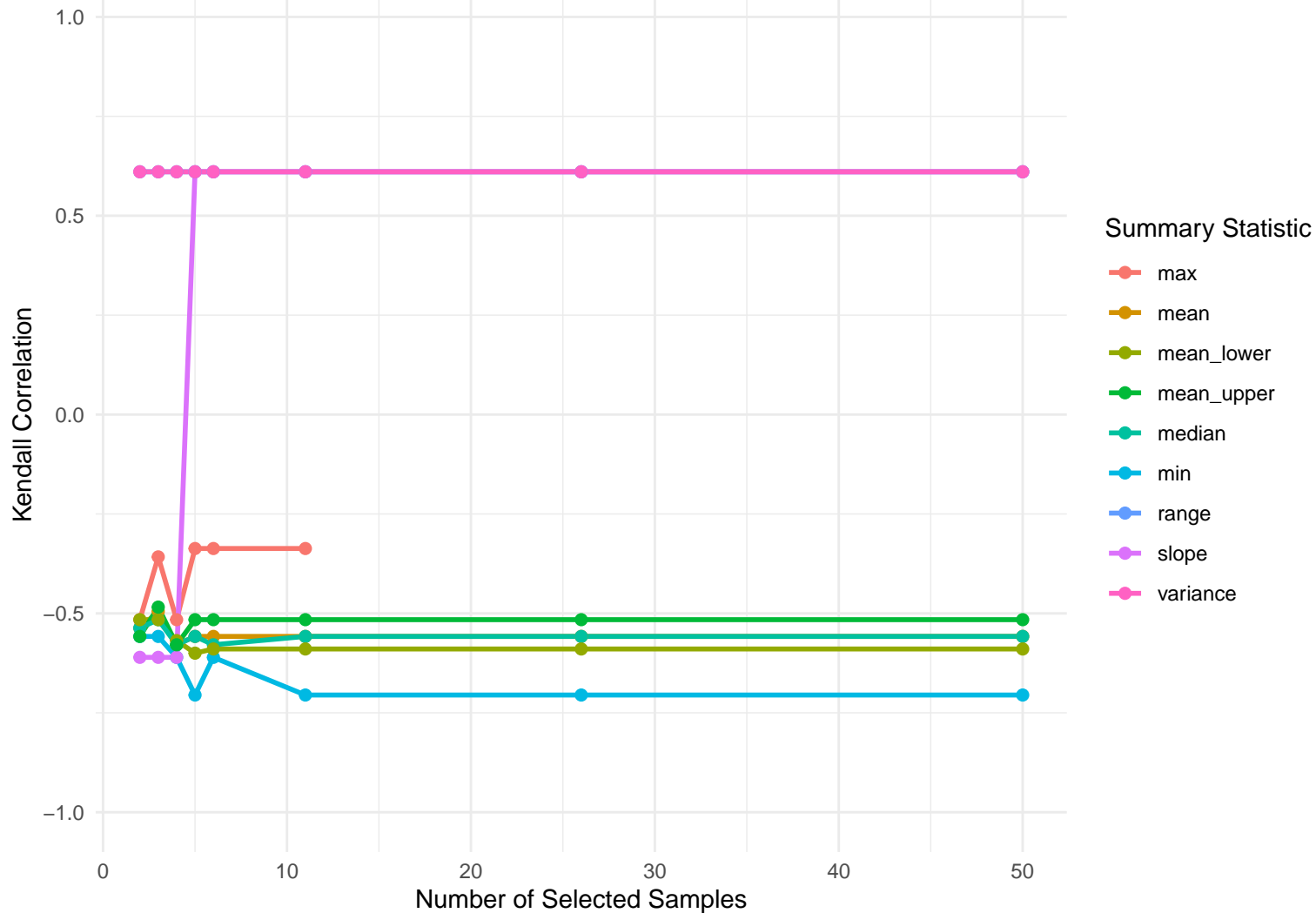
Number of Selected Samples

Summary Statistic

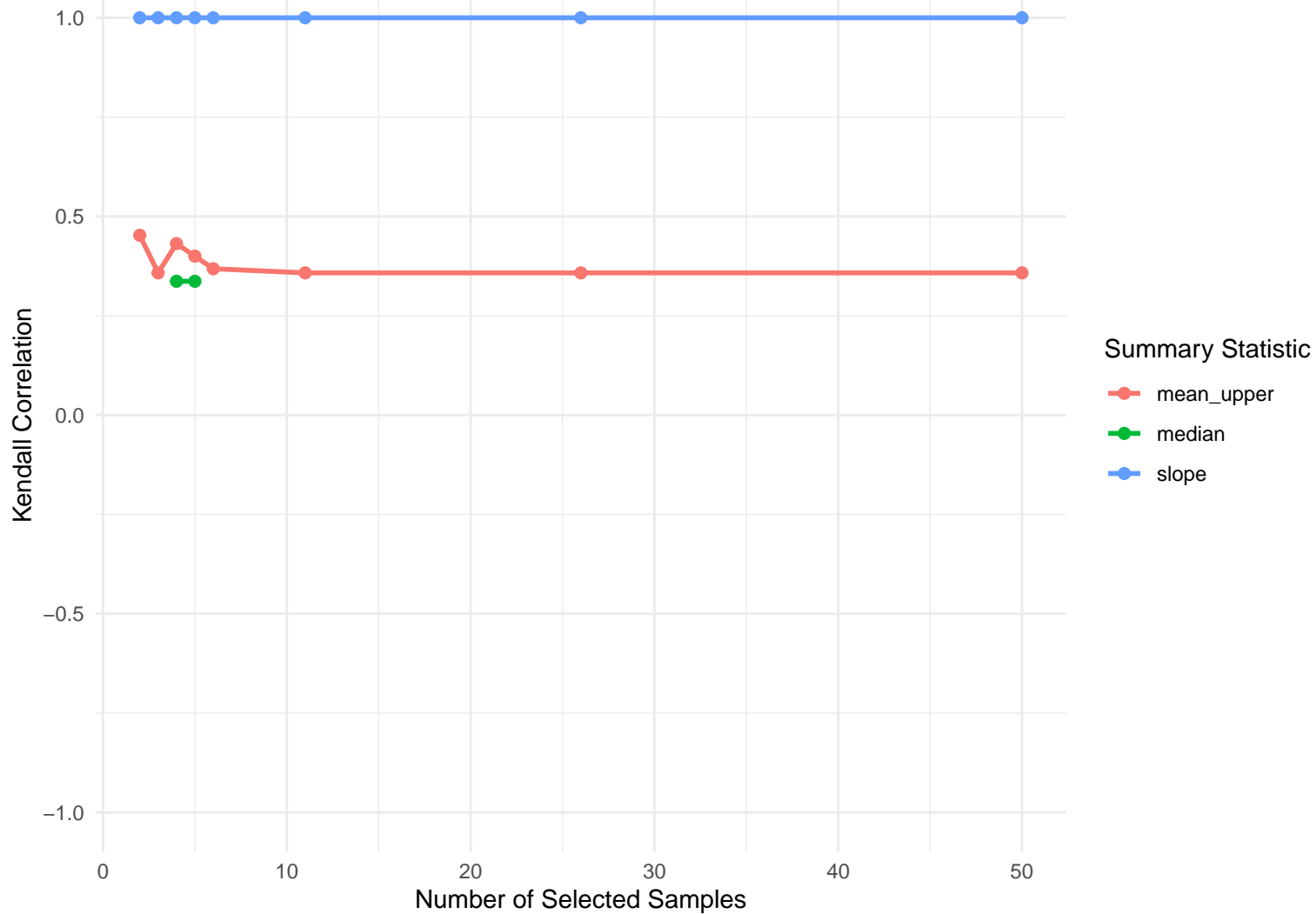
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



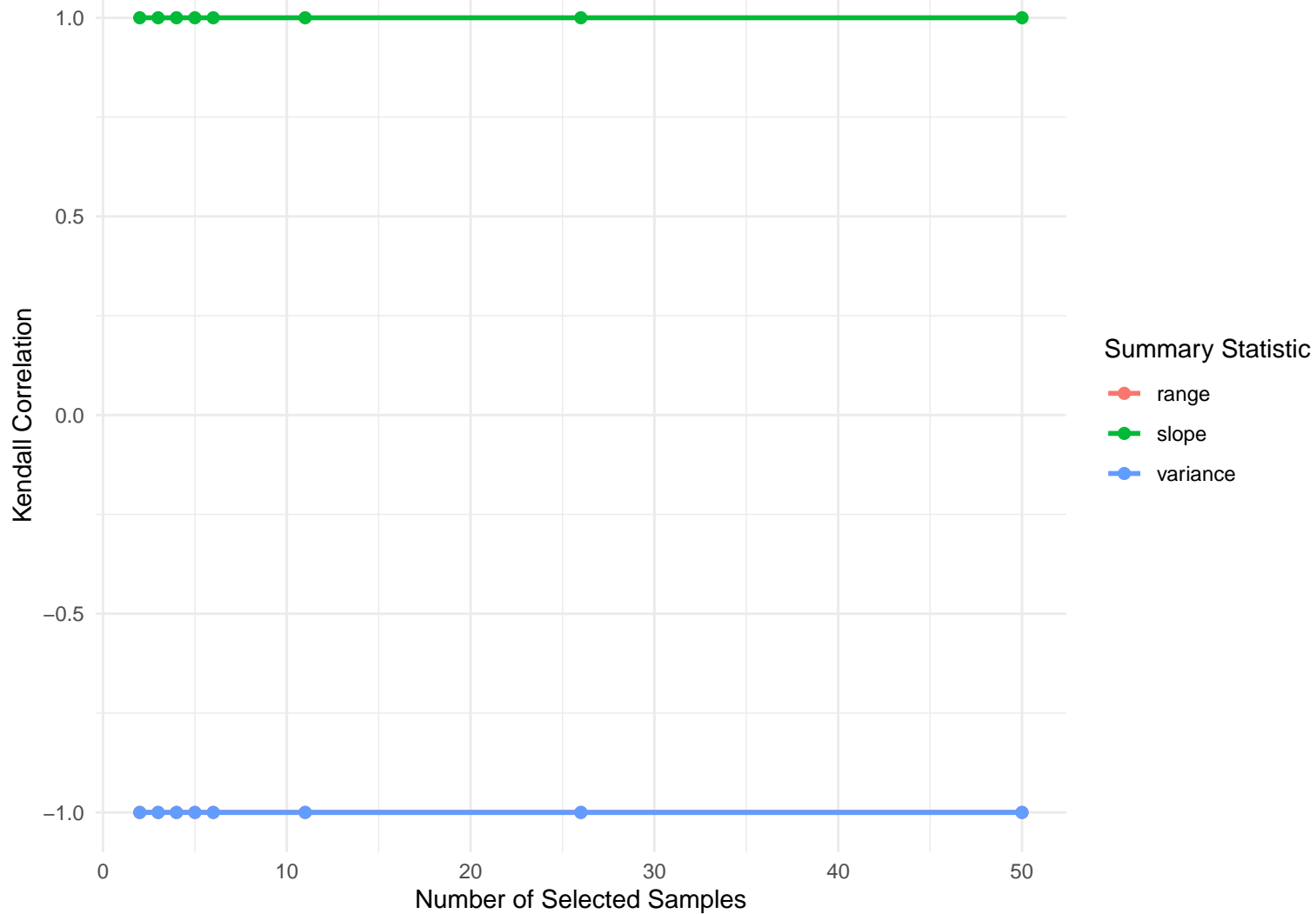
Score: CV_t (Local, full , Genotype: wave)



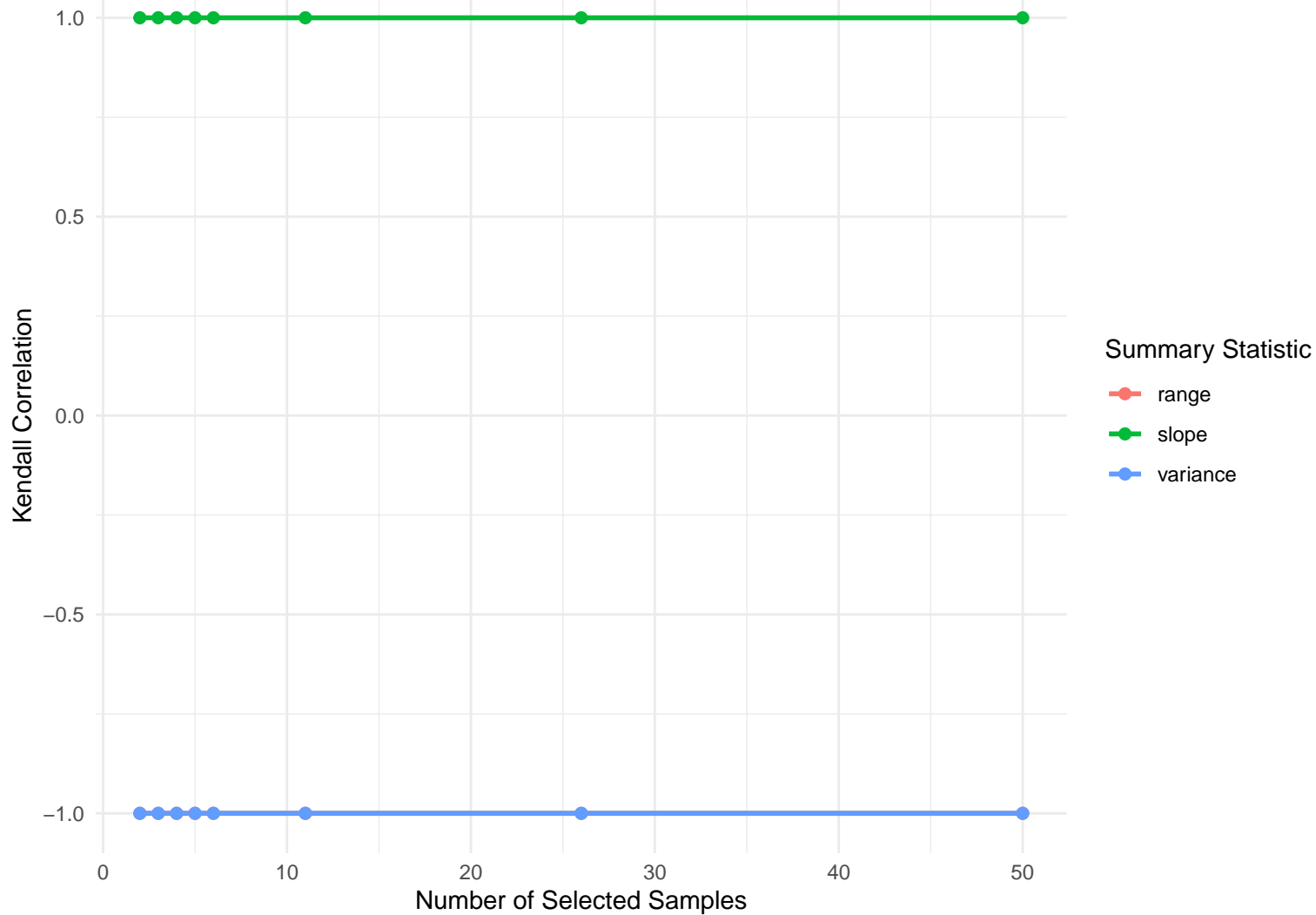
Score: RN (Local, full , Genotype: linear)



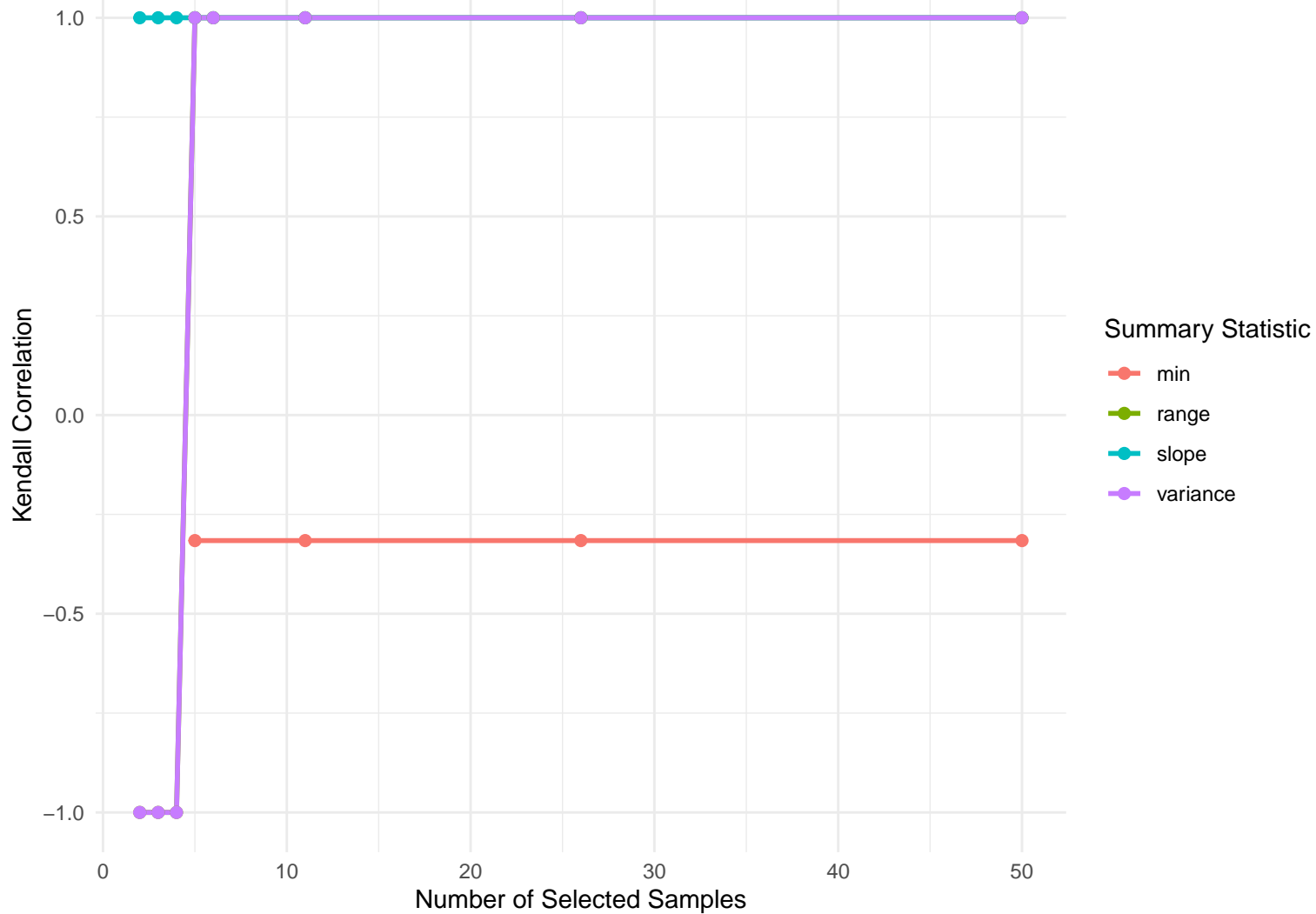
Score: RN (Local, full , Genotype: gaussian)



Score: RN (Local, full , Genotype: sinusoidal)



Score: RN (Local, full , Genotype: wave)



Score: RNN (Local, full , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

20

30

40

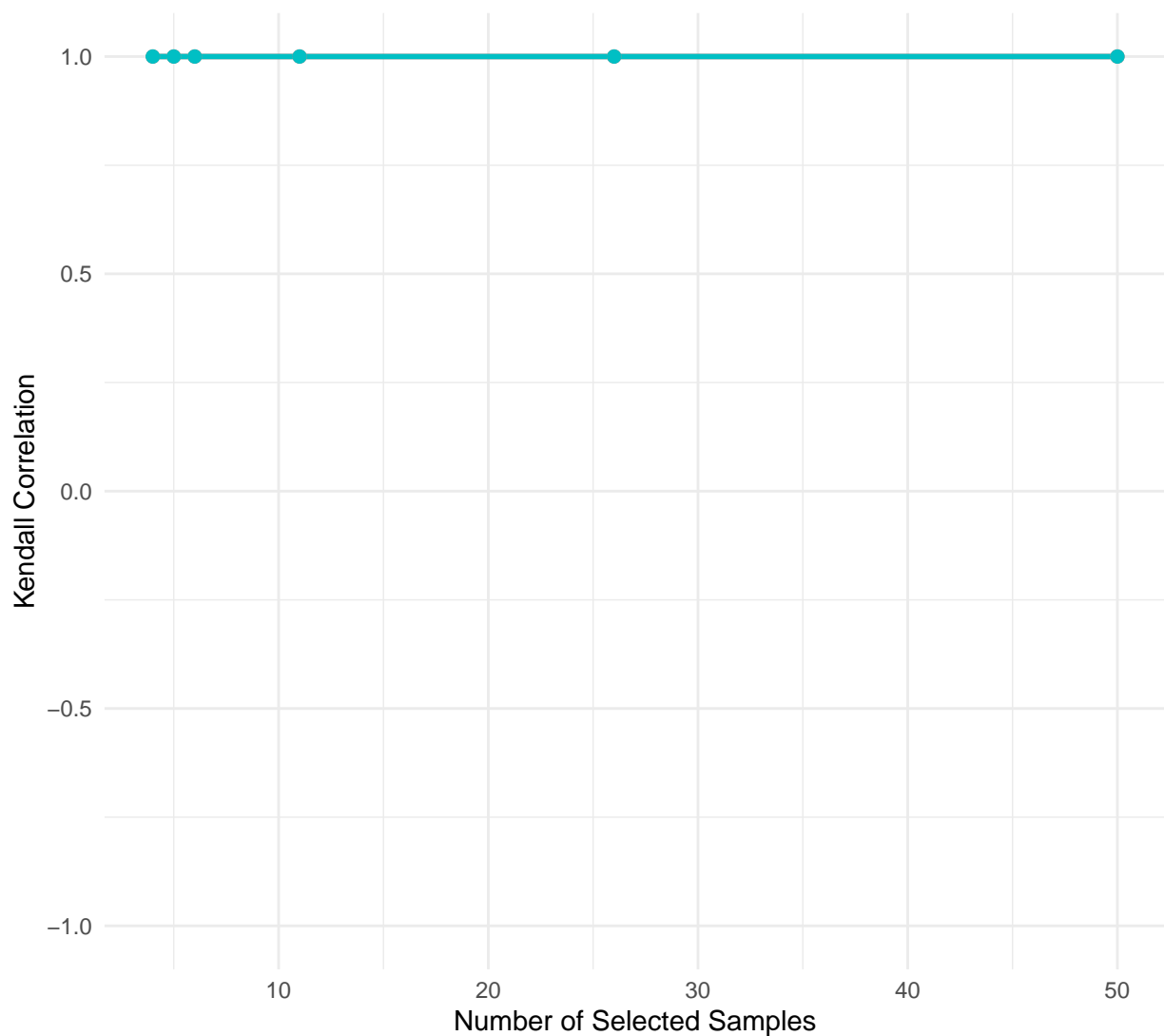
50

Number of Selected Samples

Summary Statistic

range

variance



Score: RNN (Local, full , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

20

30

40

50

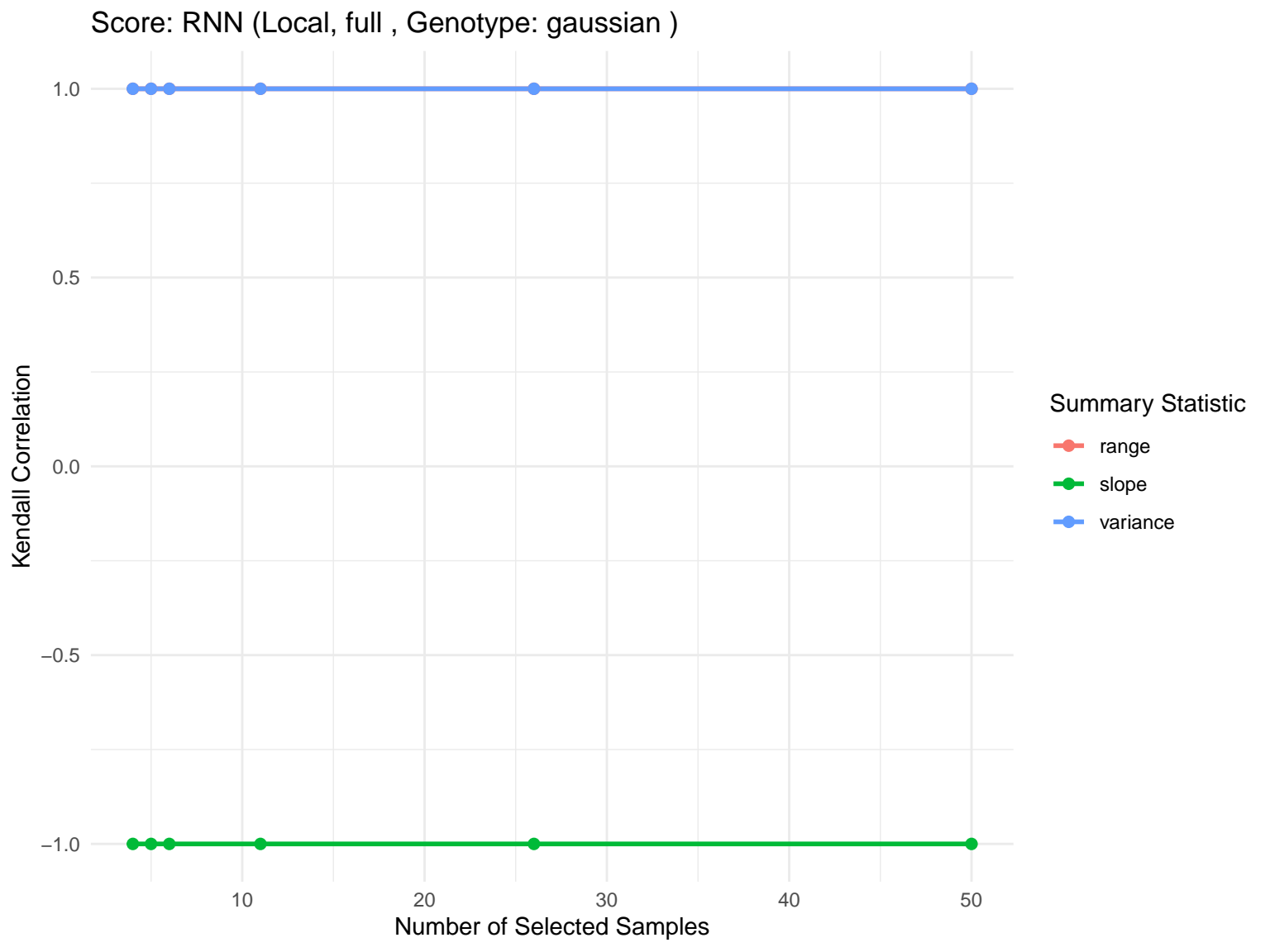
Number of Selected Samples

Summary Statistic

range

slope

variance

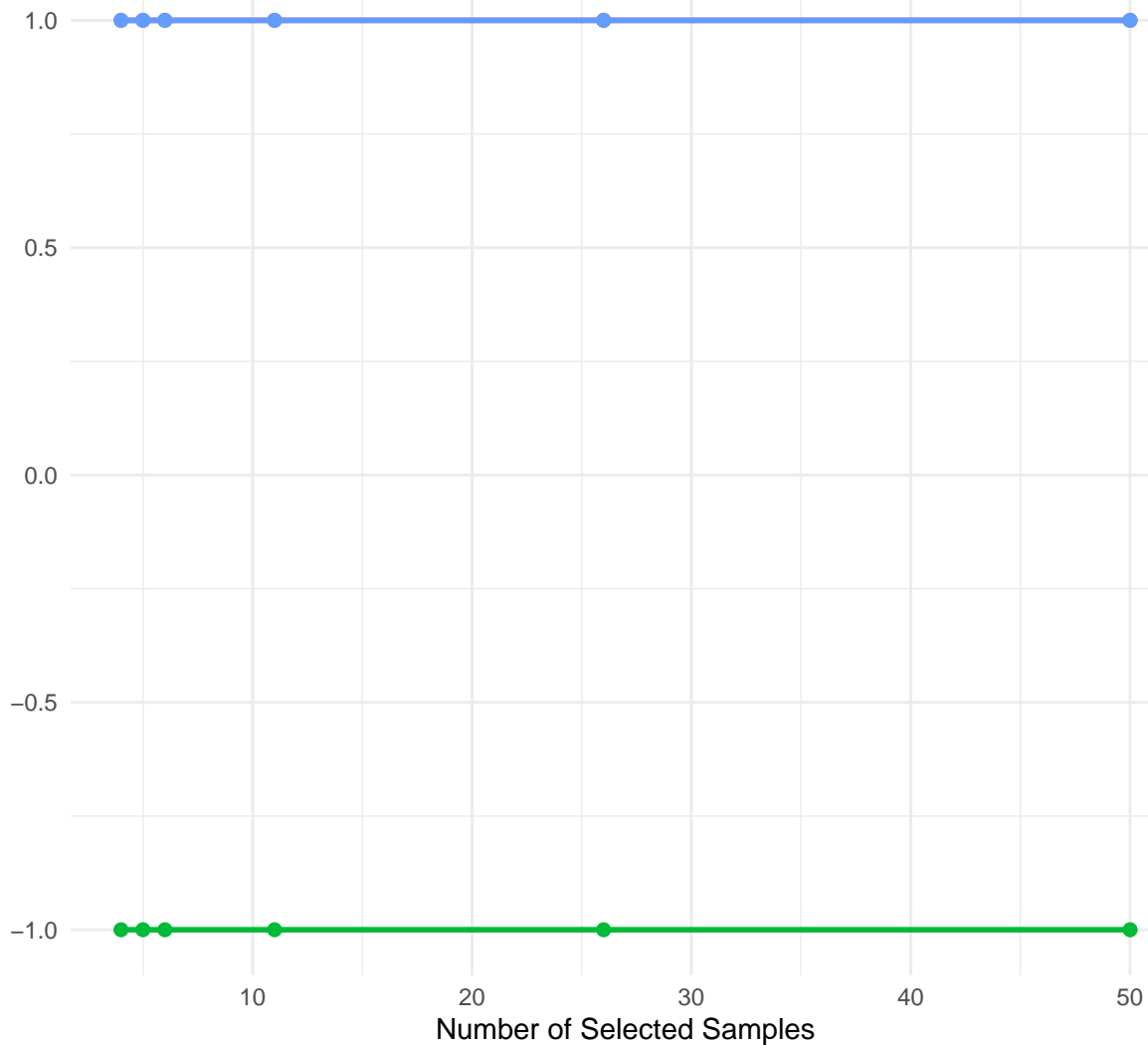


Score: RNN (Local, full , Genotype: sinusoidal)

Kendall Correlation

Summary Statistic

- range
- slope
- variance



Score: RNN (Local, full , Genotype: wave)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

20

30

40

50

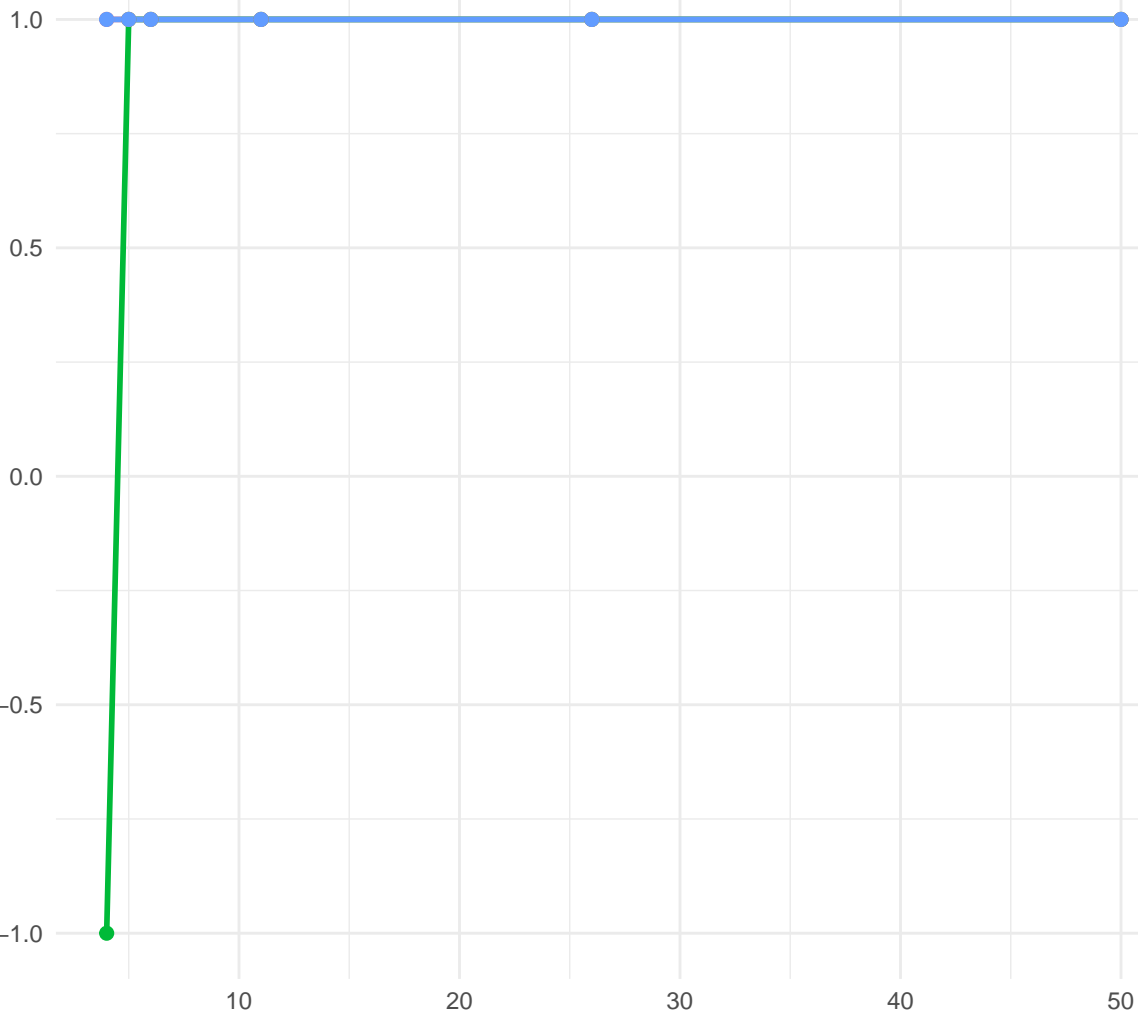
Number of Selected Samples

Summary Statistic

range

slope

variance



Score: D_slope (Local, full , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

0

10

20

30

40

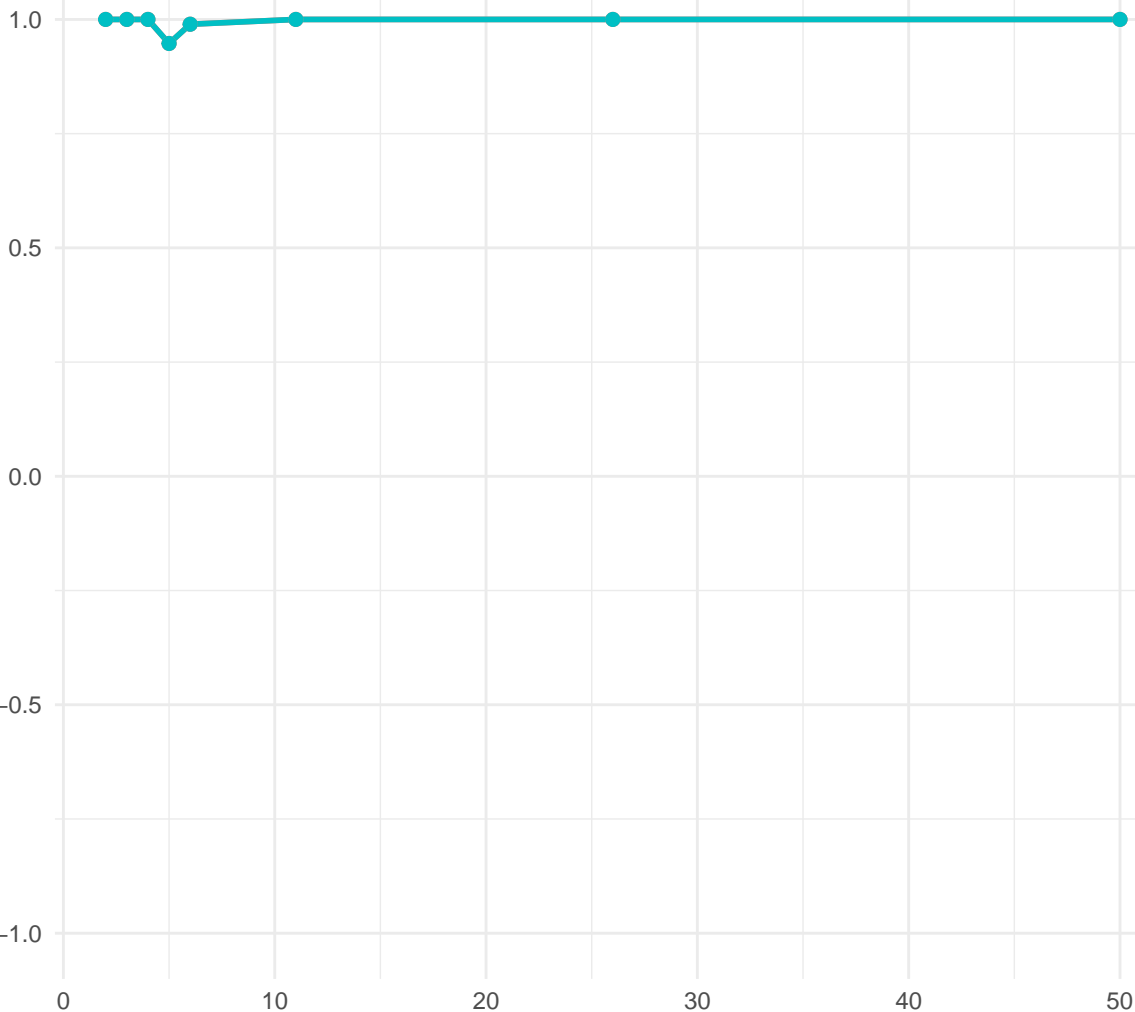
50

Number of Selected Samples

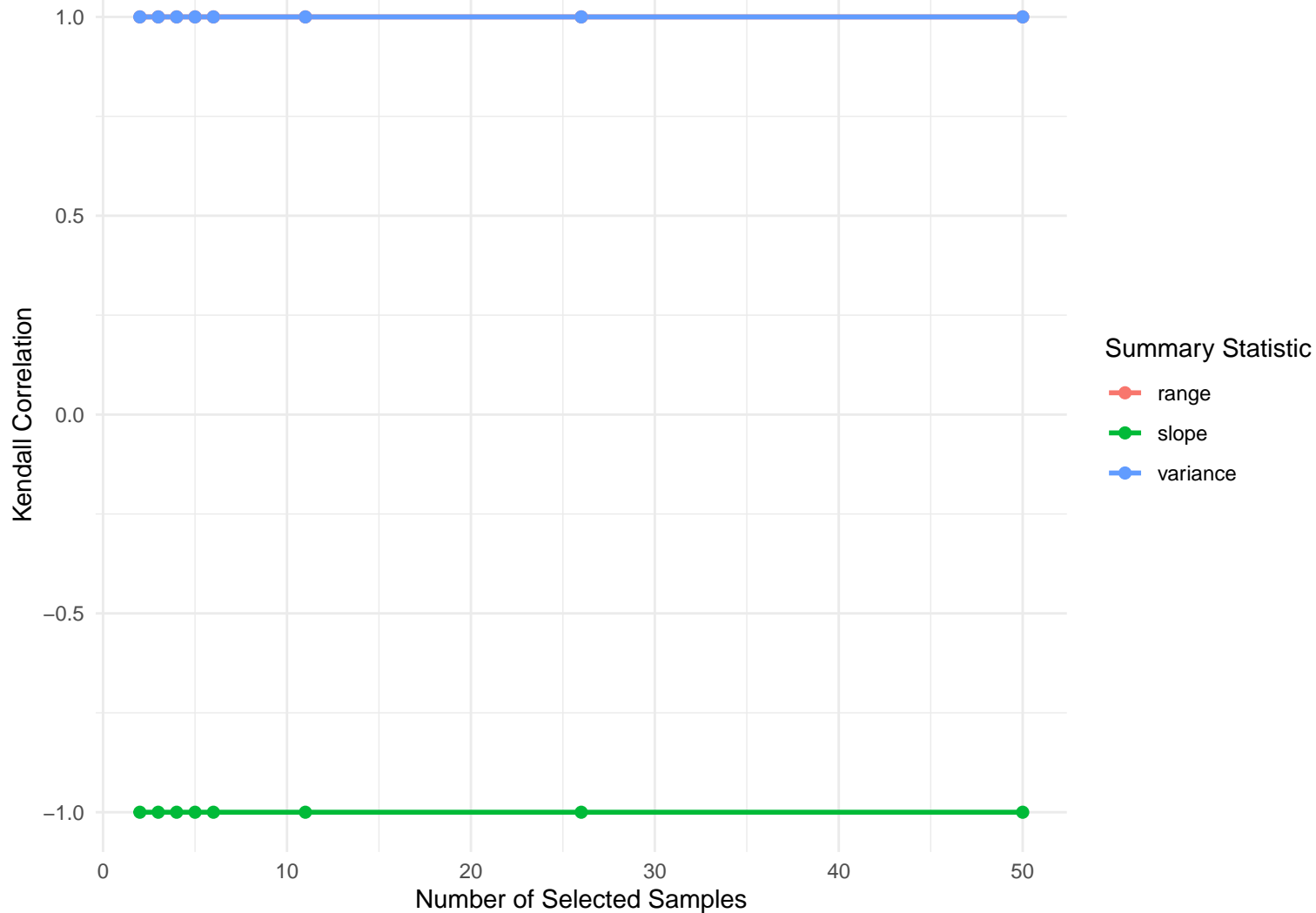
Summary Statistic

range

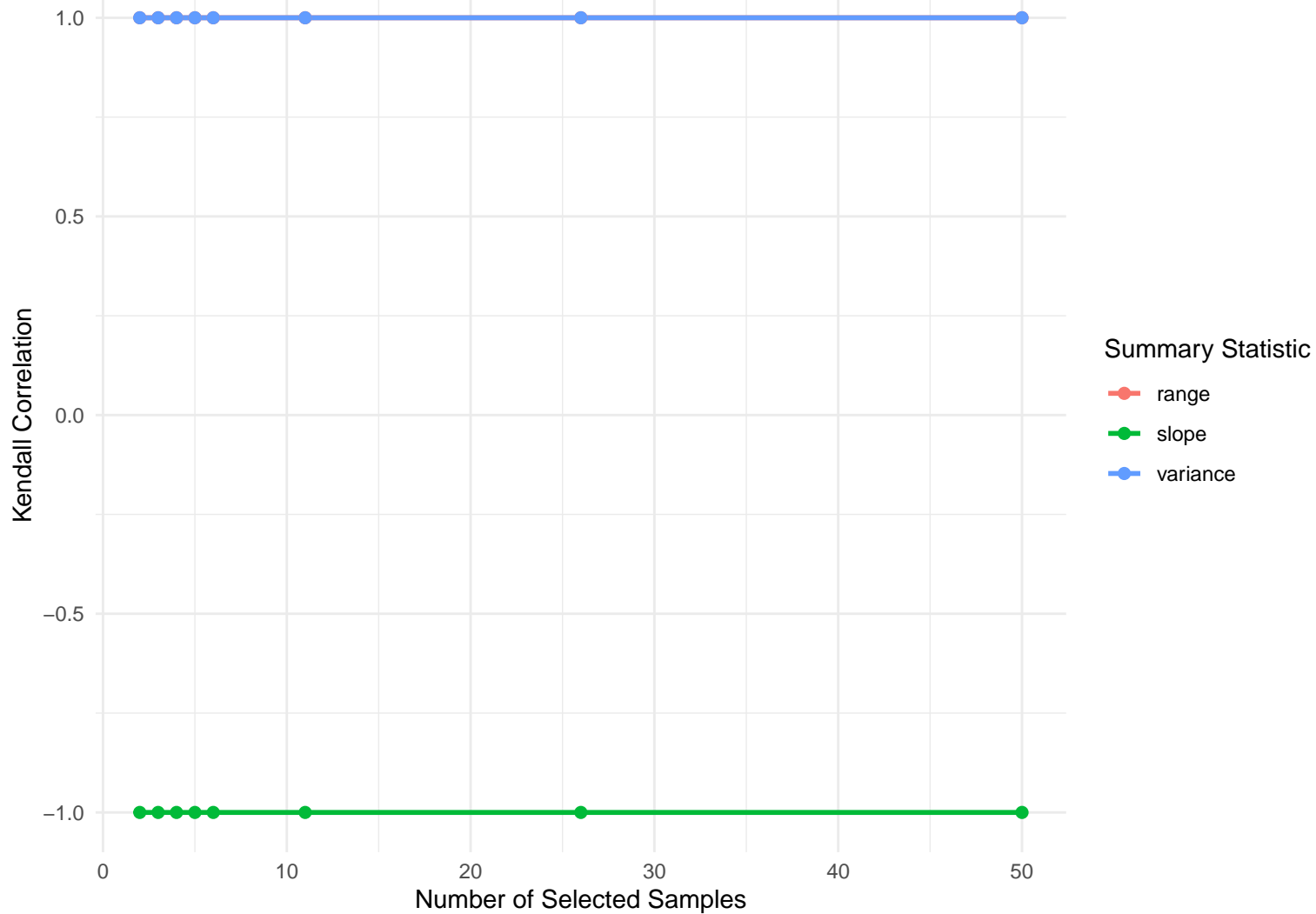
variance



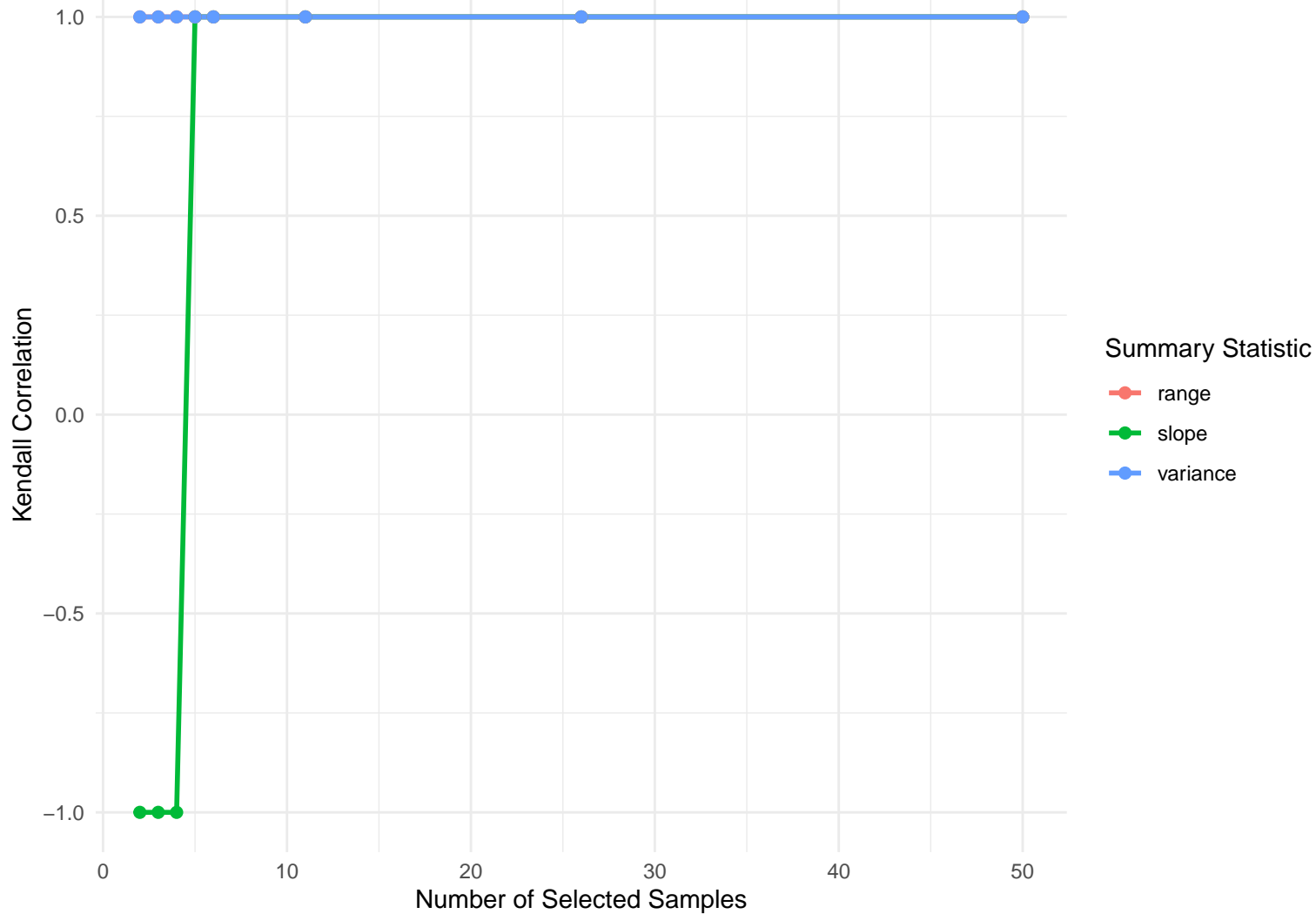
Score: D_slope (Local, full , Genotype: gaussian)



Score: D_slope (Local, full , Genotype: sinusoidal)



Score: D_slope (Local, full , Genotype: wave)



Score: RC (Local, full , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

0

10

20

30

40

50

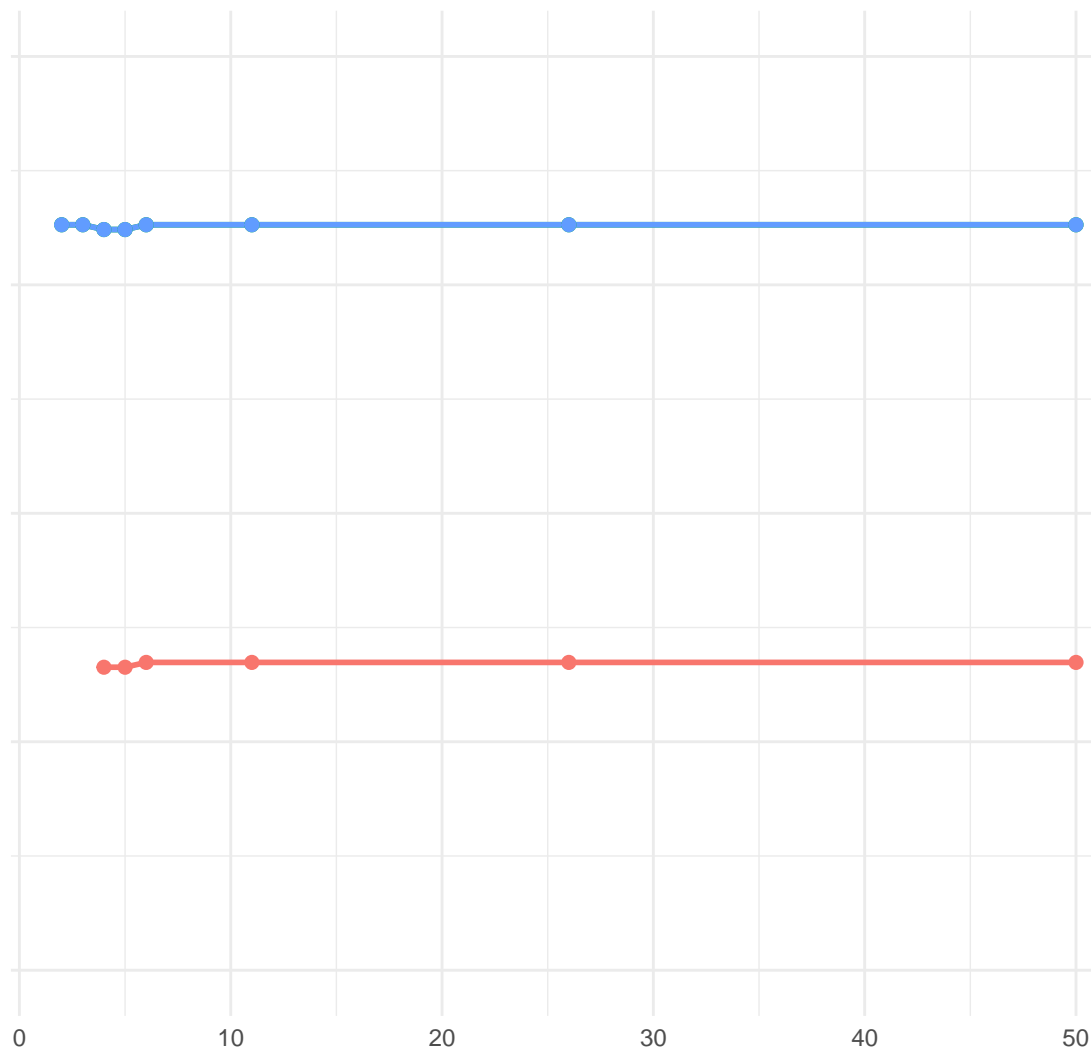
Number of Selected Samples

Summary Statistic

min

range

variance



Score: RC (Local, full , Genotype: gaussian)

Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

0

10

20

30

40

50

Number of Selected Samples

Summary Statistic

- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance

0

10

20

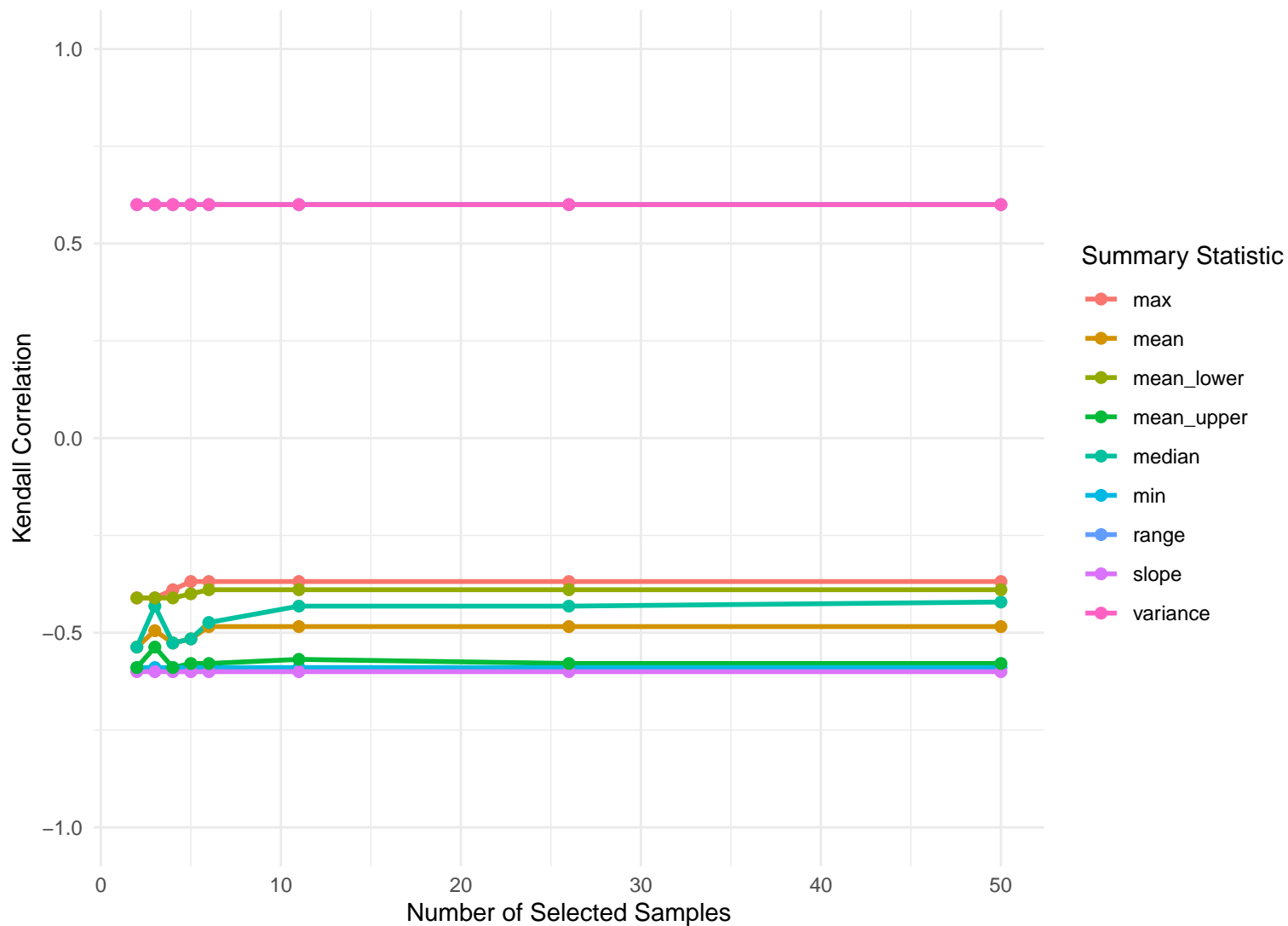
30

40

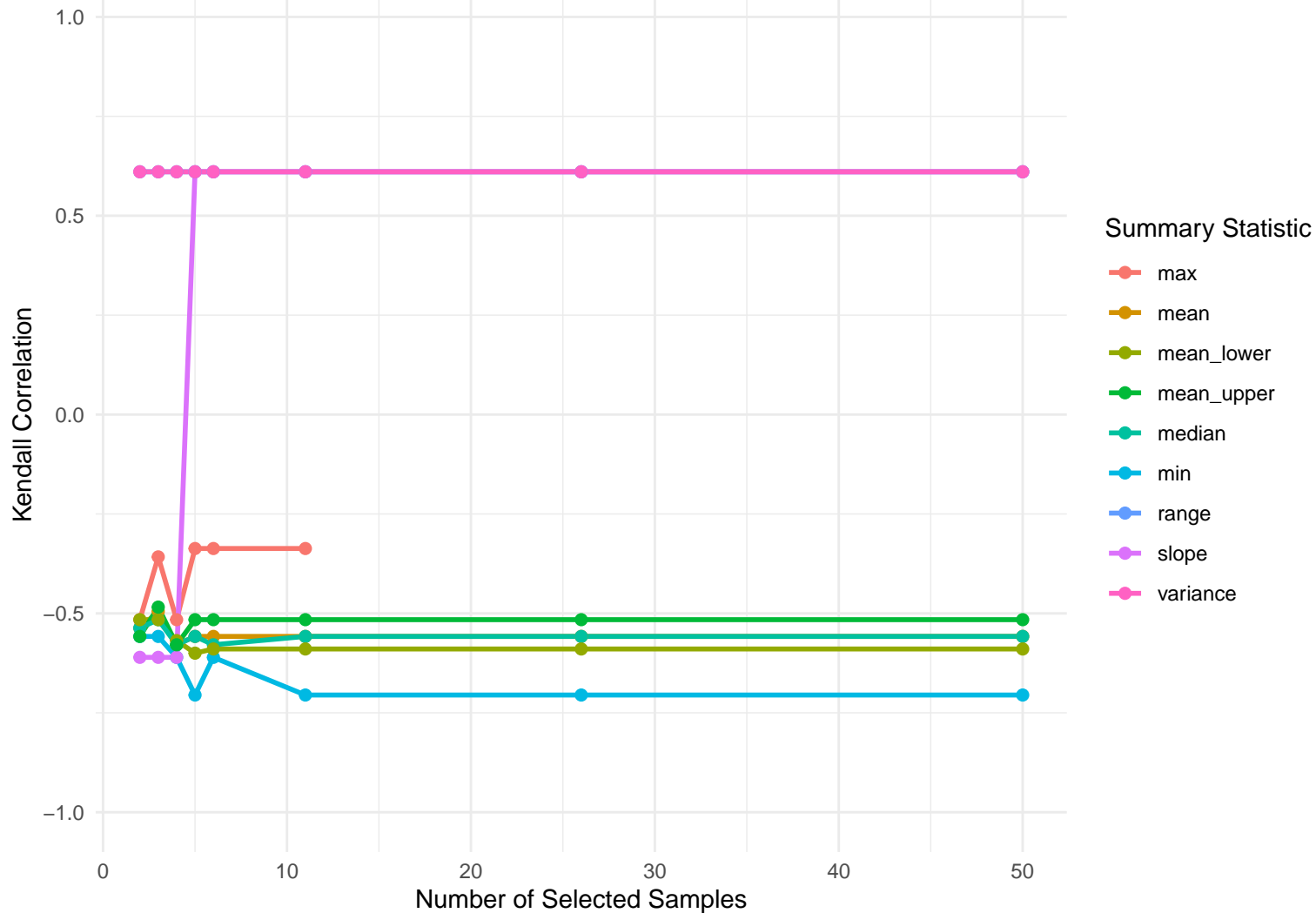
50

Number of Selected Samples

Score: RC (Local, full , Genotype: sinusoidal)



Score: RC (Local, full , Genotype: wave)



Score: gPi (Local, full , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

0

10

20

30

40

50

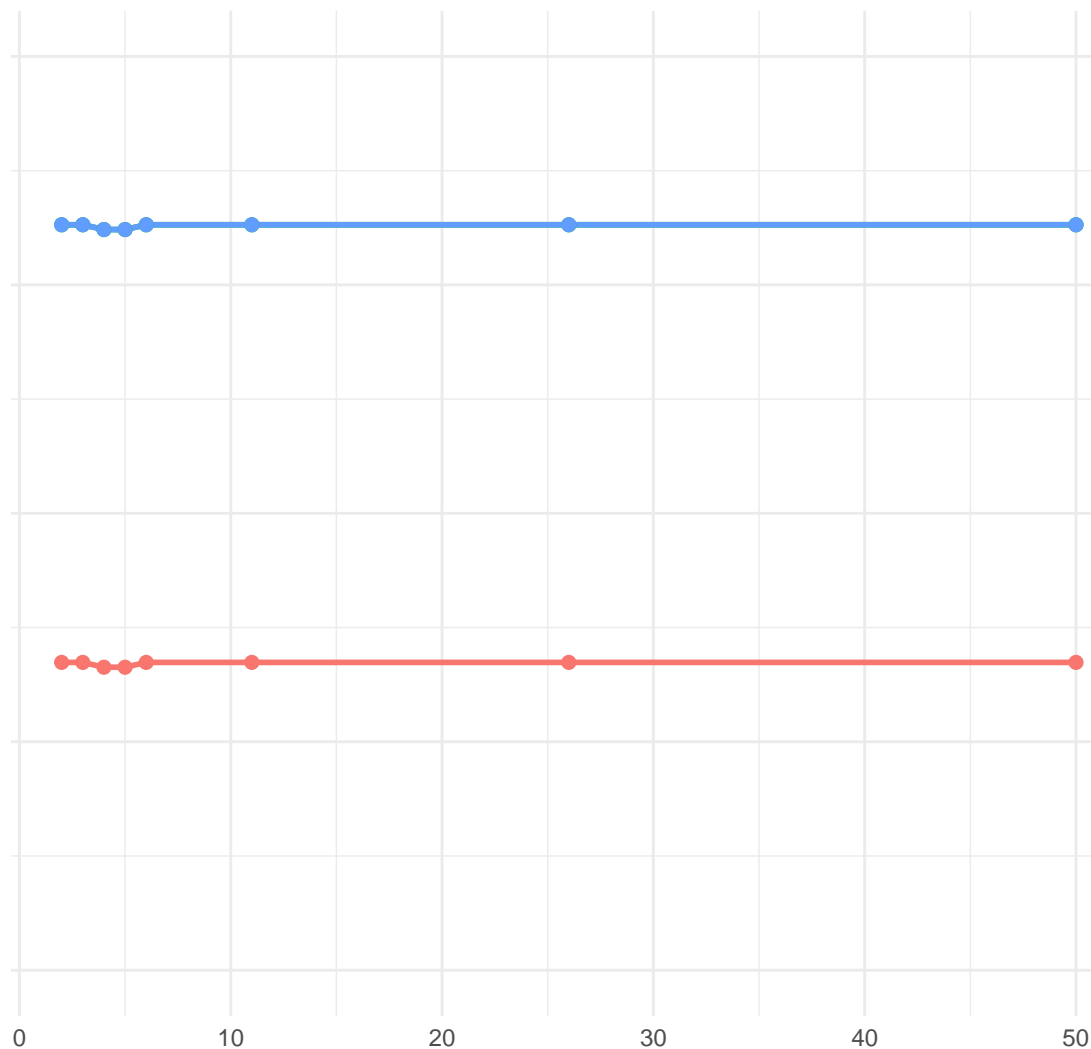
Number of Selected Samples

Summary Statistic

min

range

variance



Score: gPi (Local, full , Genotype: gaussian)

Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

0

10

20

30

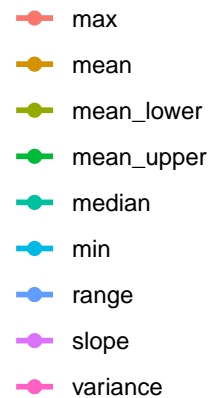
40

50

Number of Selected Samples

Summary Statistic

- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



Score: gPi (Local, full , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

0

10

20

30

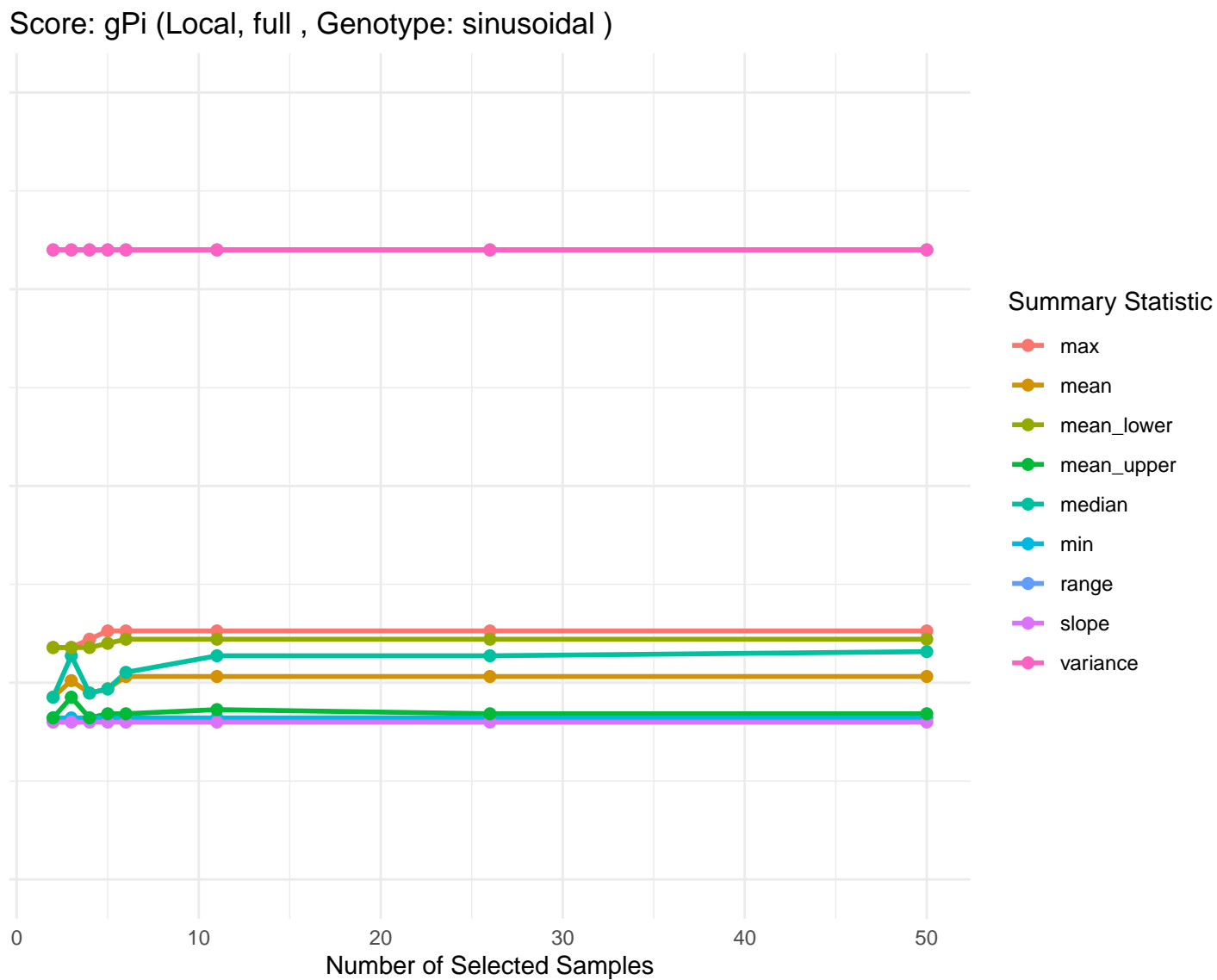
40

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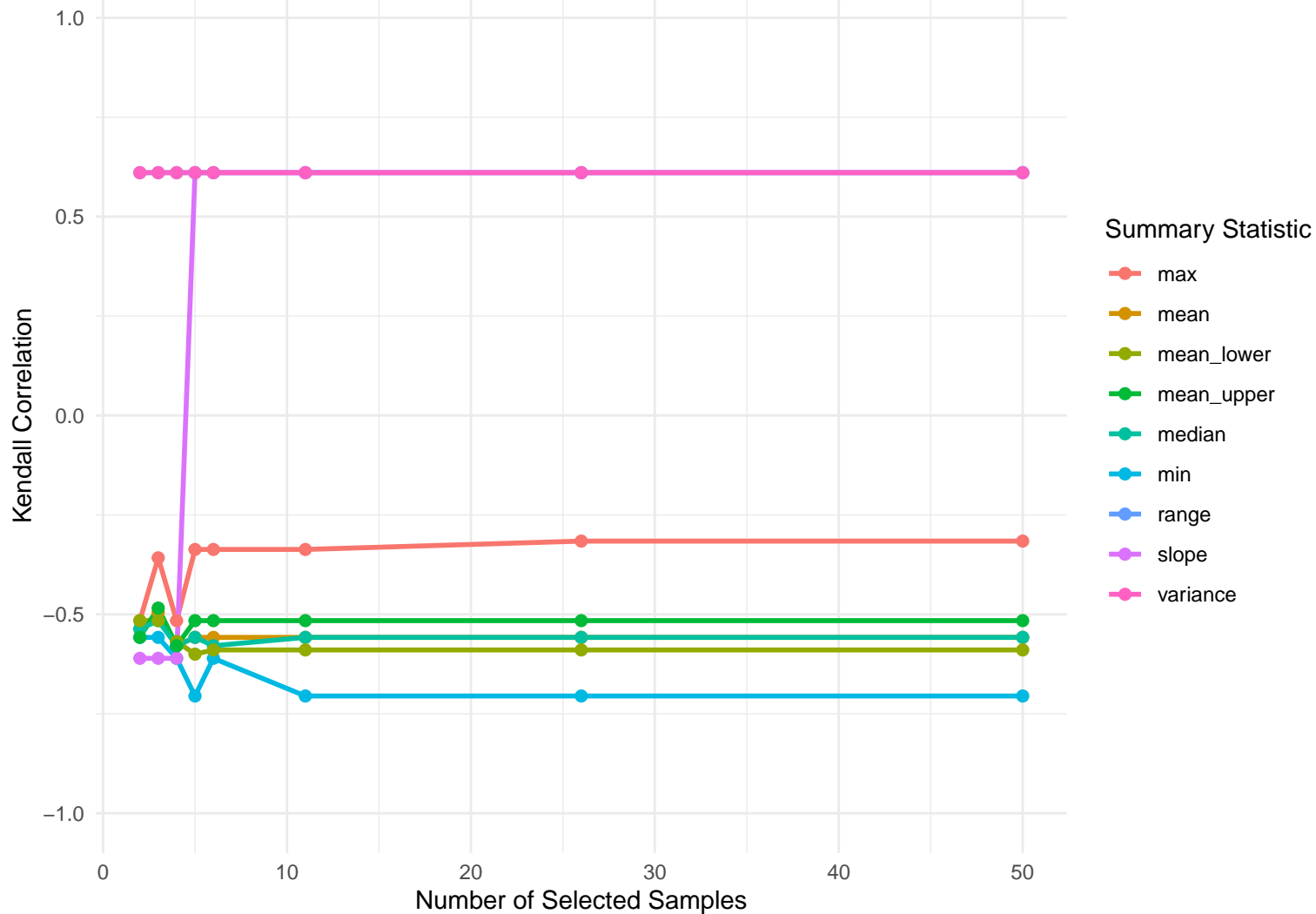
Number of Selected Samples

Summary Statistic

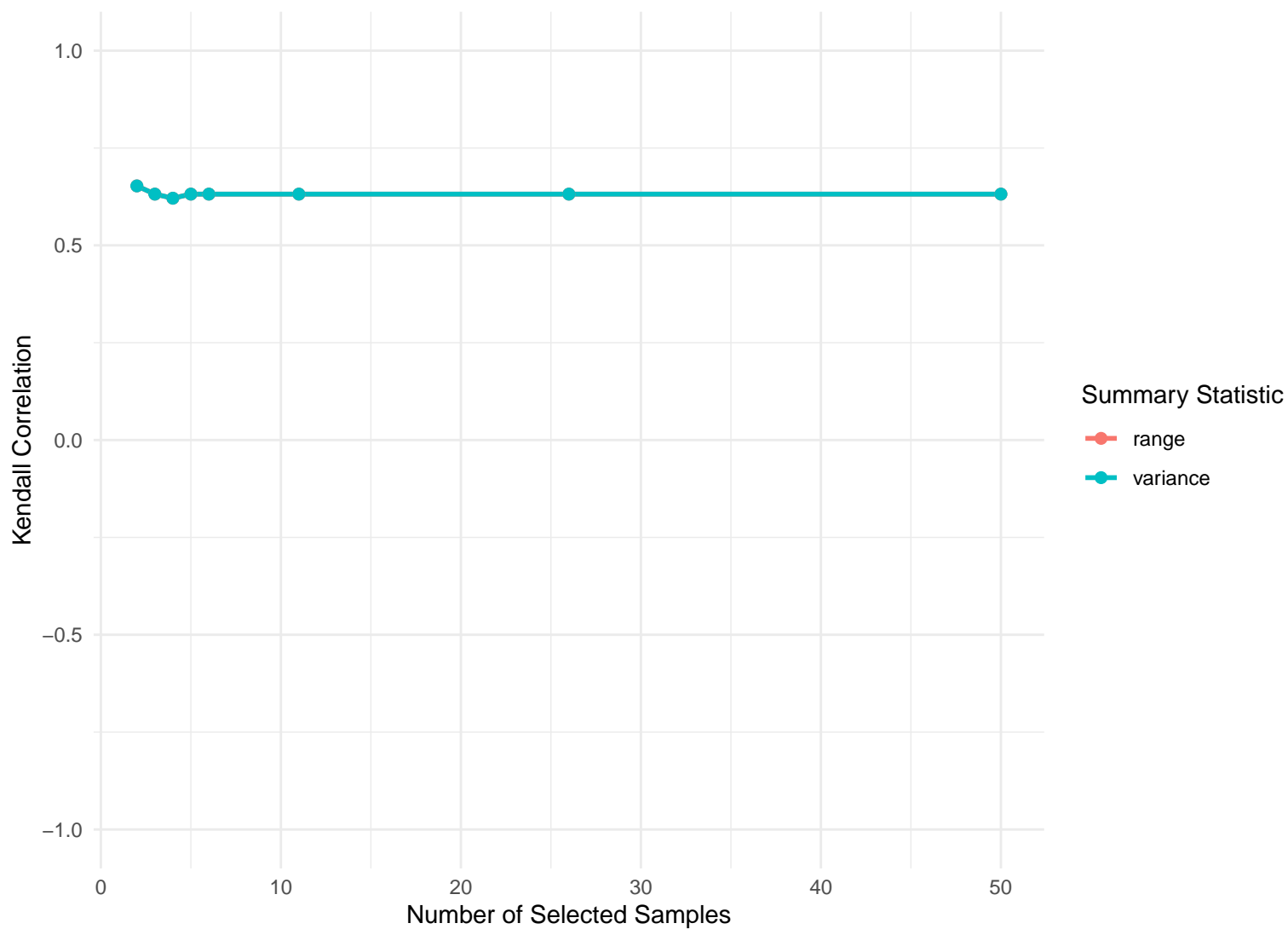
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



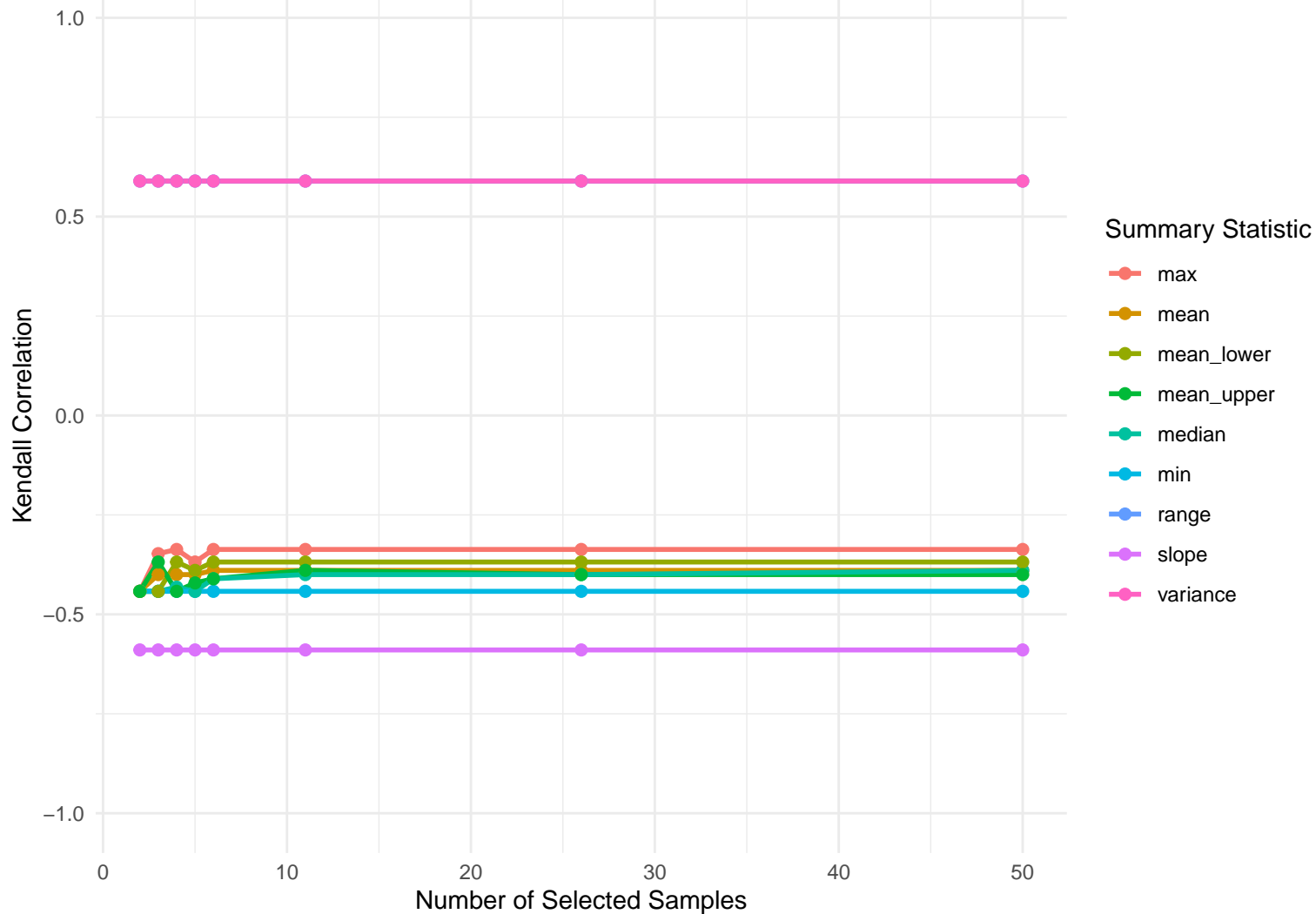
Score: gPi (Local, full , Genotype: wave)



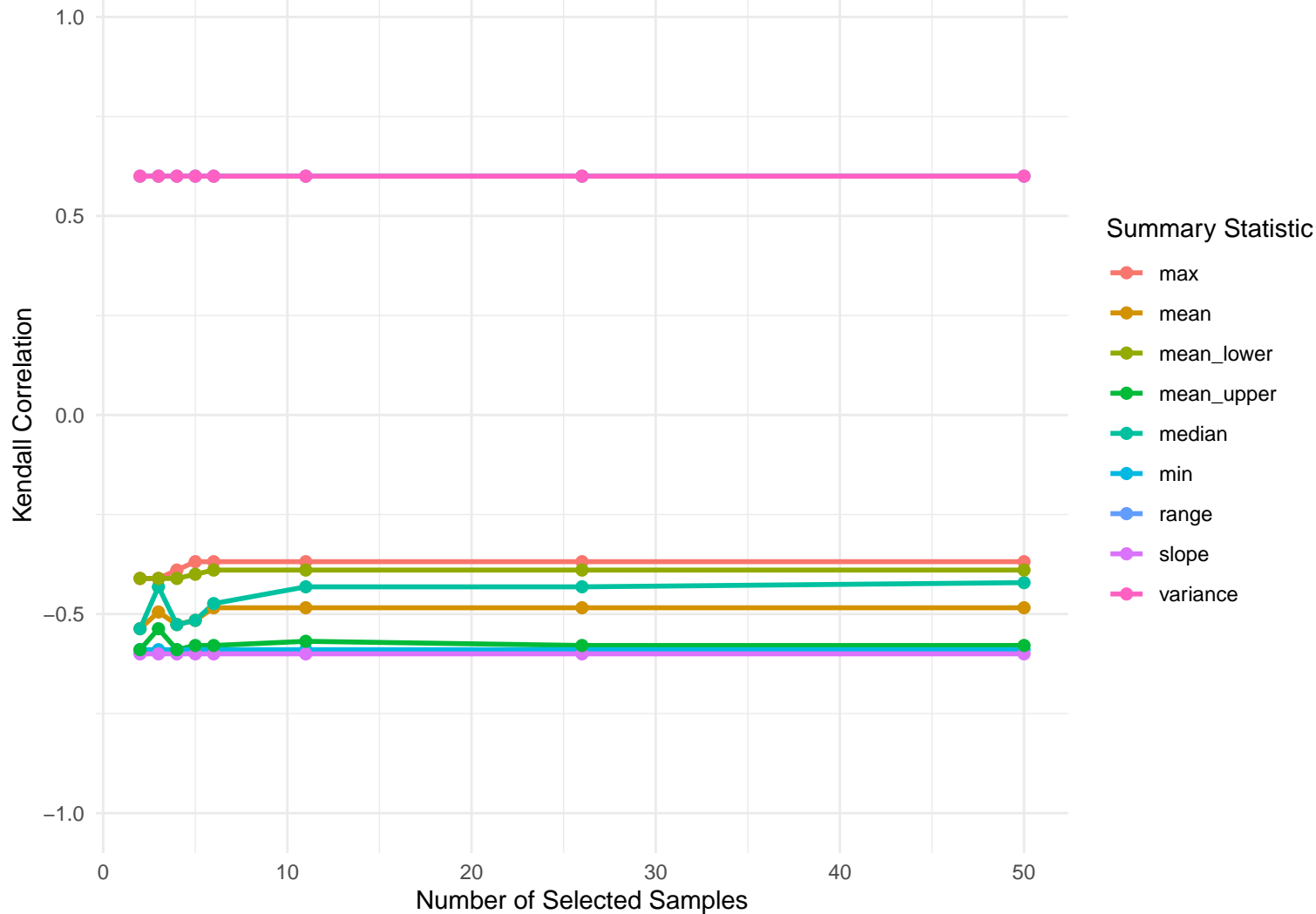
Score: PPF (Local, full , Genotype: linear)



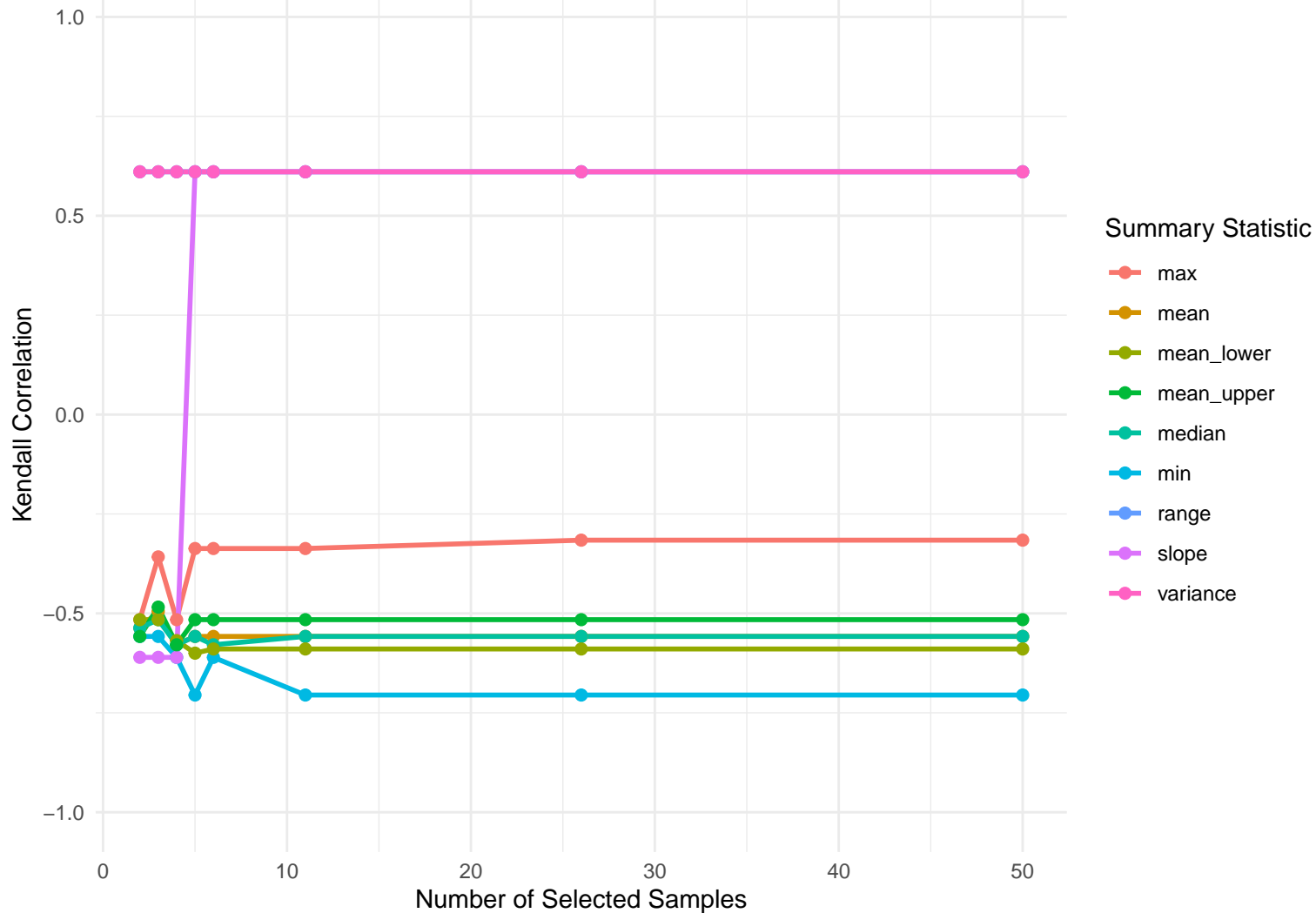
Score: PPF (Local, full , Genotype: gaussian)



Score: PPF (Local, full , Genotype: sinusoidal)



Score: PPF (Local, full , Genotype: wave)



Score: PPI (Local, full , Genotype: linear)

Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

0

10

20

30

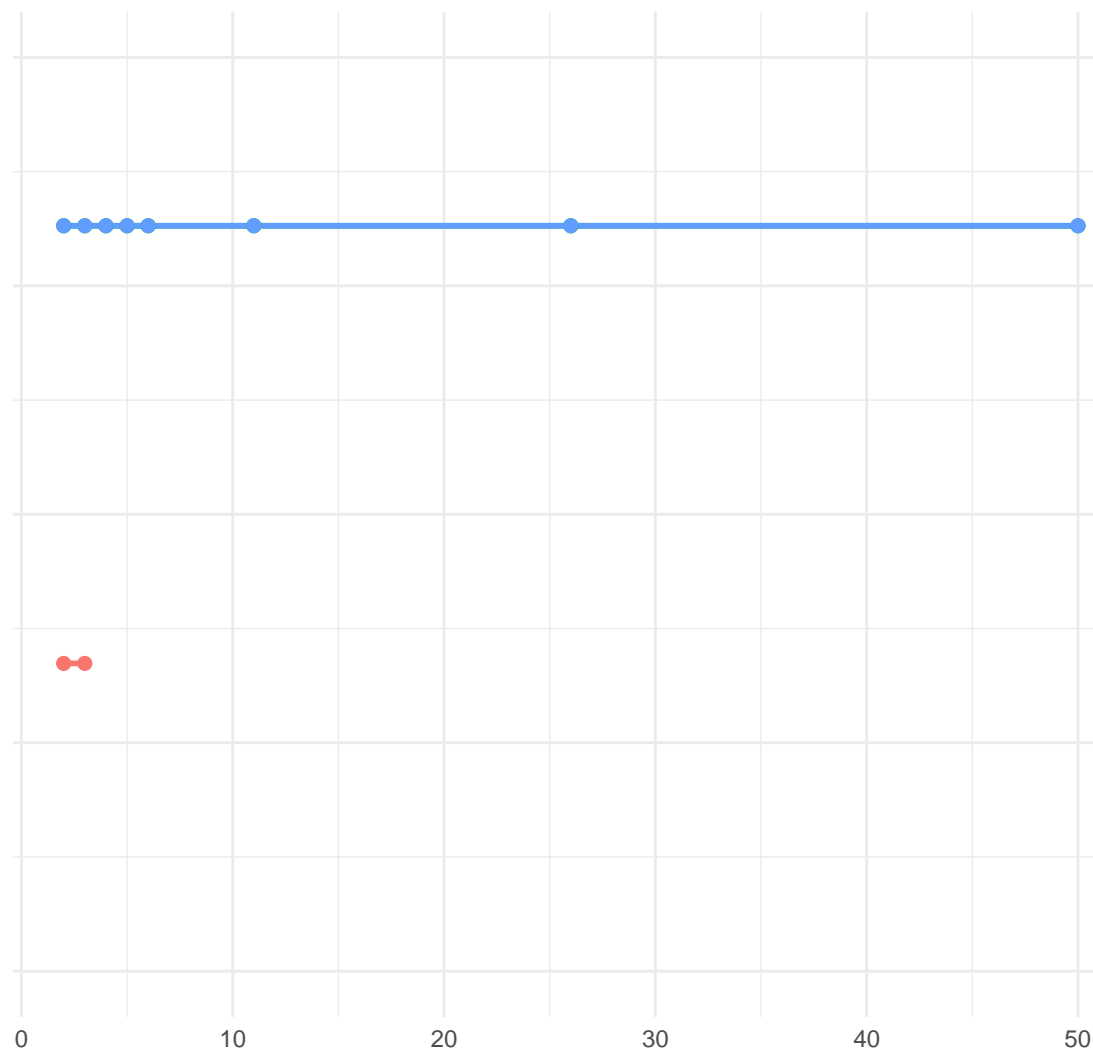
40

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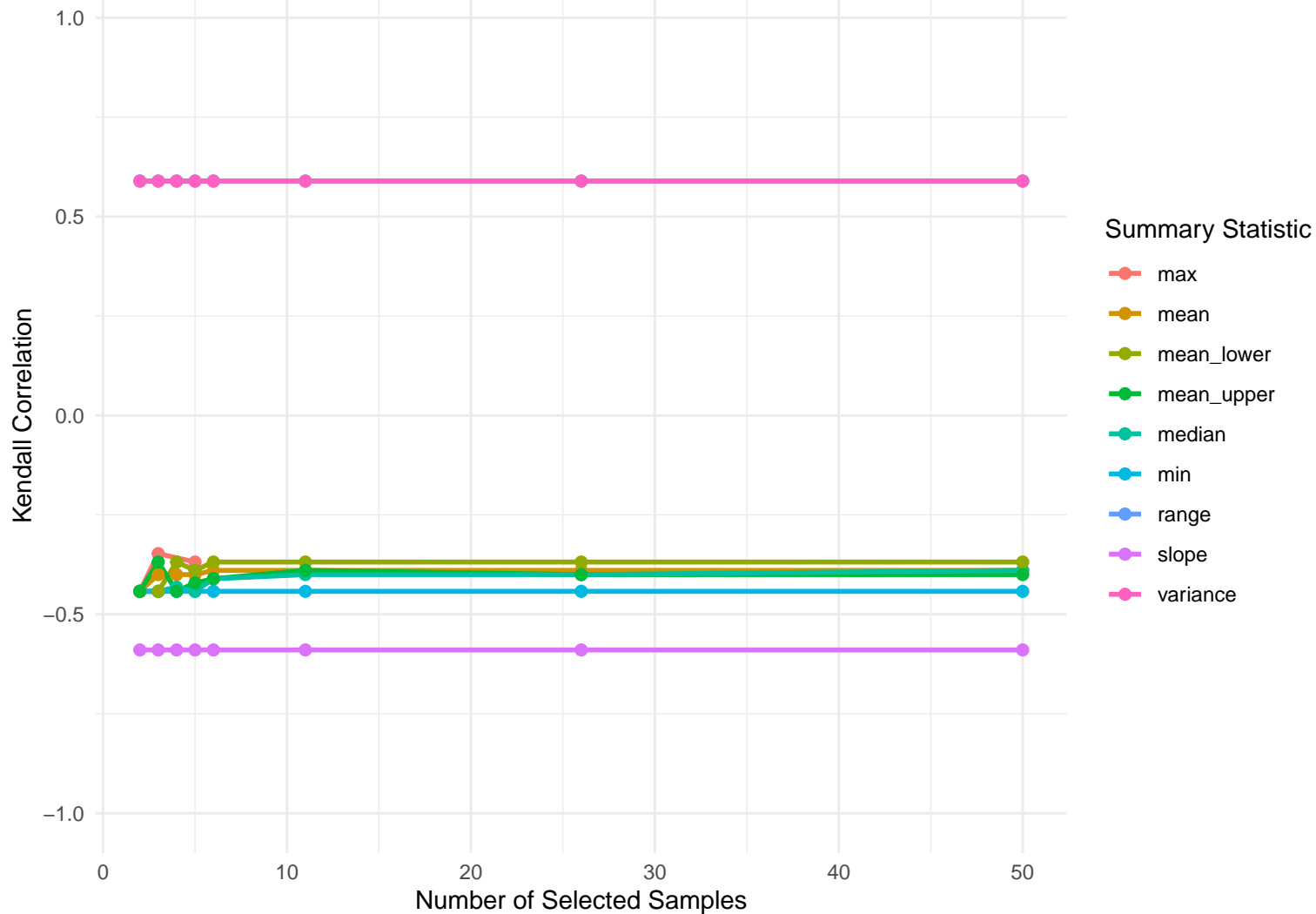
Number of Selected Samples

Summary Statistic

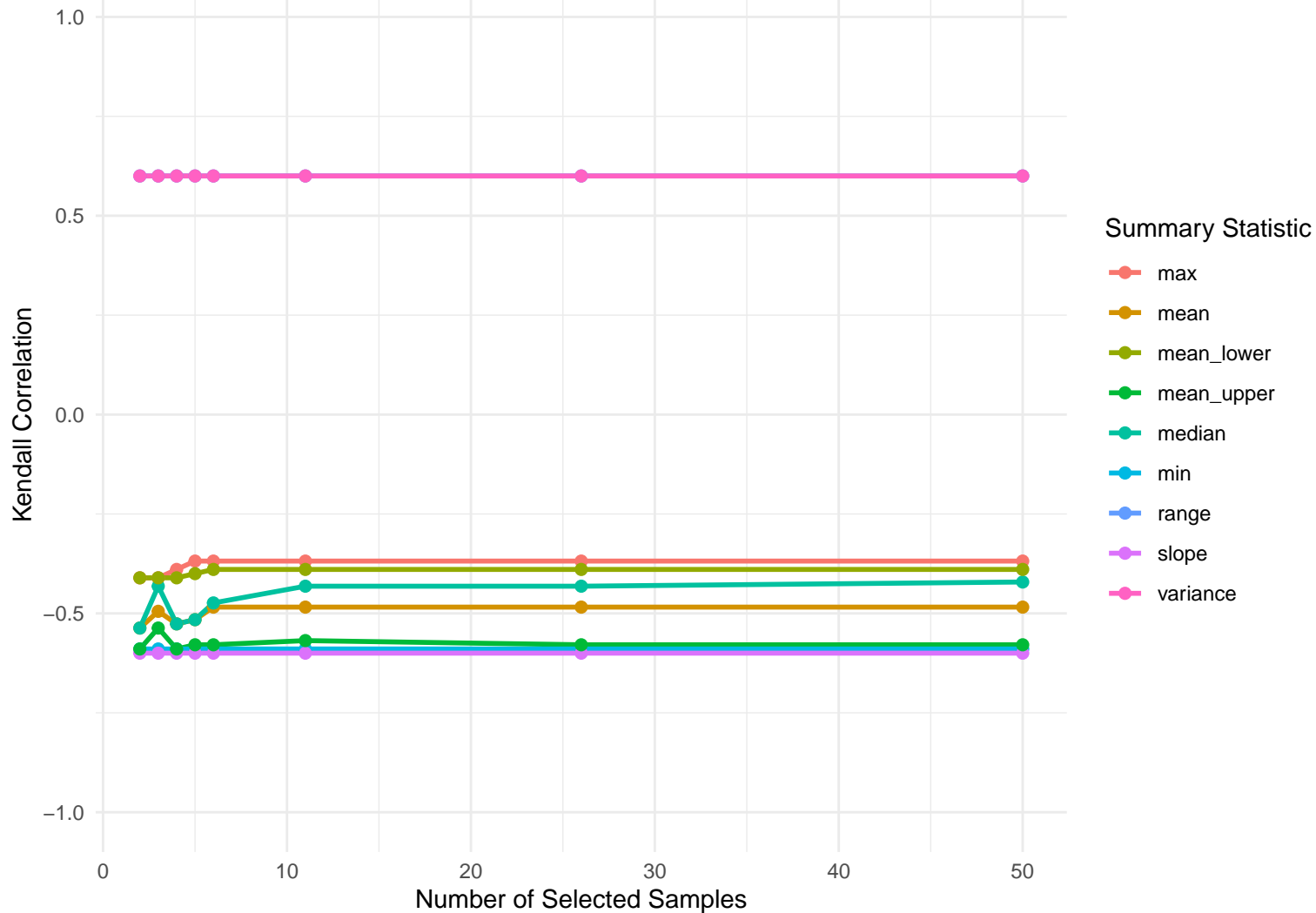
- min
- range
- variance



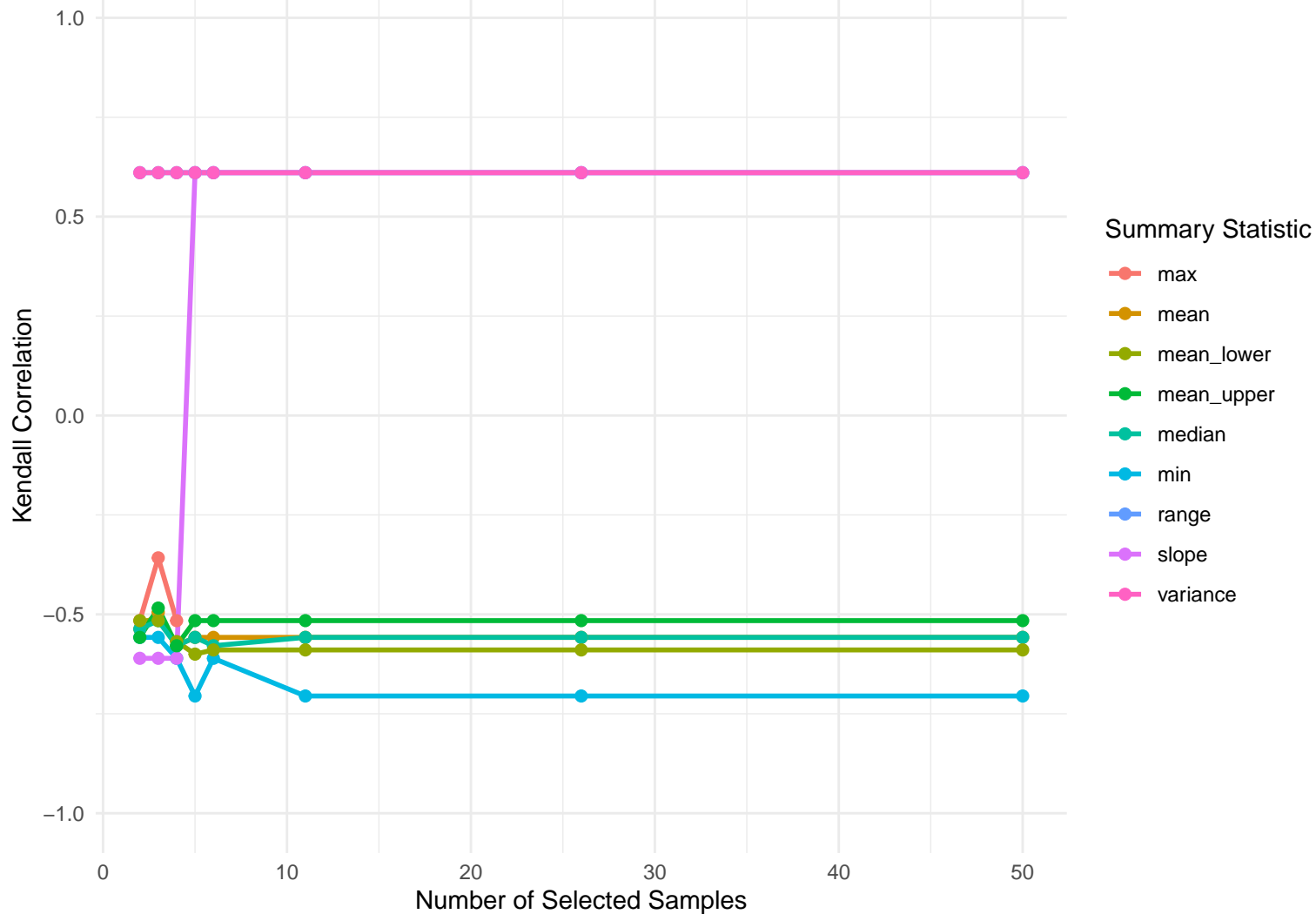
Score: PPI (Local, full , Genotype: gaussian)



Score: PPI (Local, full , Genotype: sinusoidal)



Score: PPI (Local, full , Genotype: wave)



Score: Plmd (Local, full , Genotype: linear)

Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

0

10

20

30

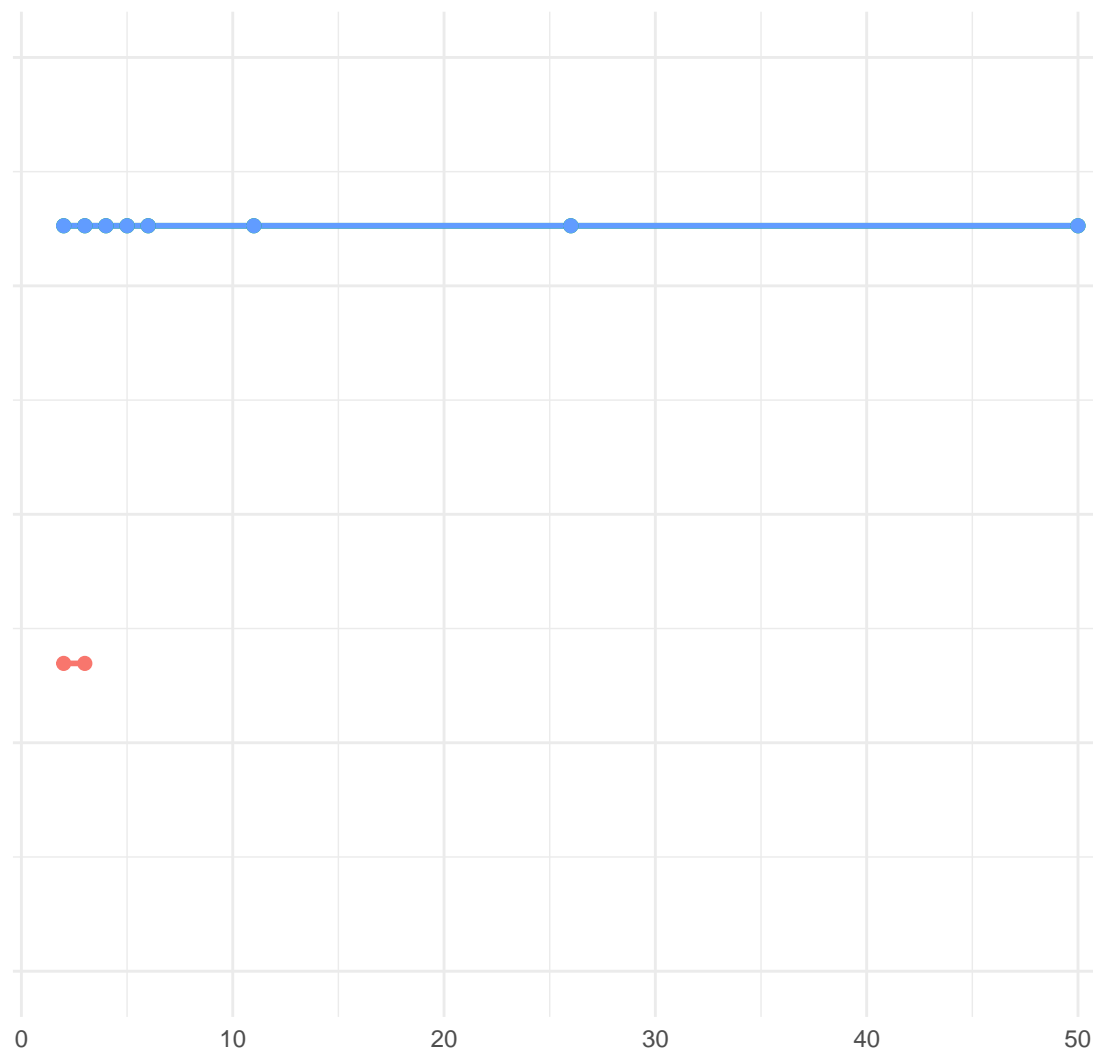
40

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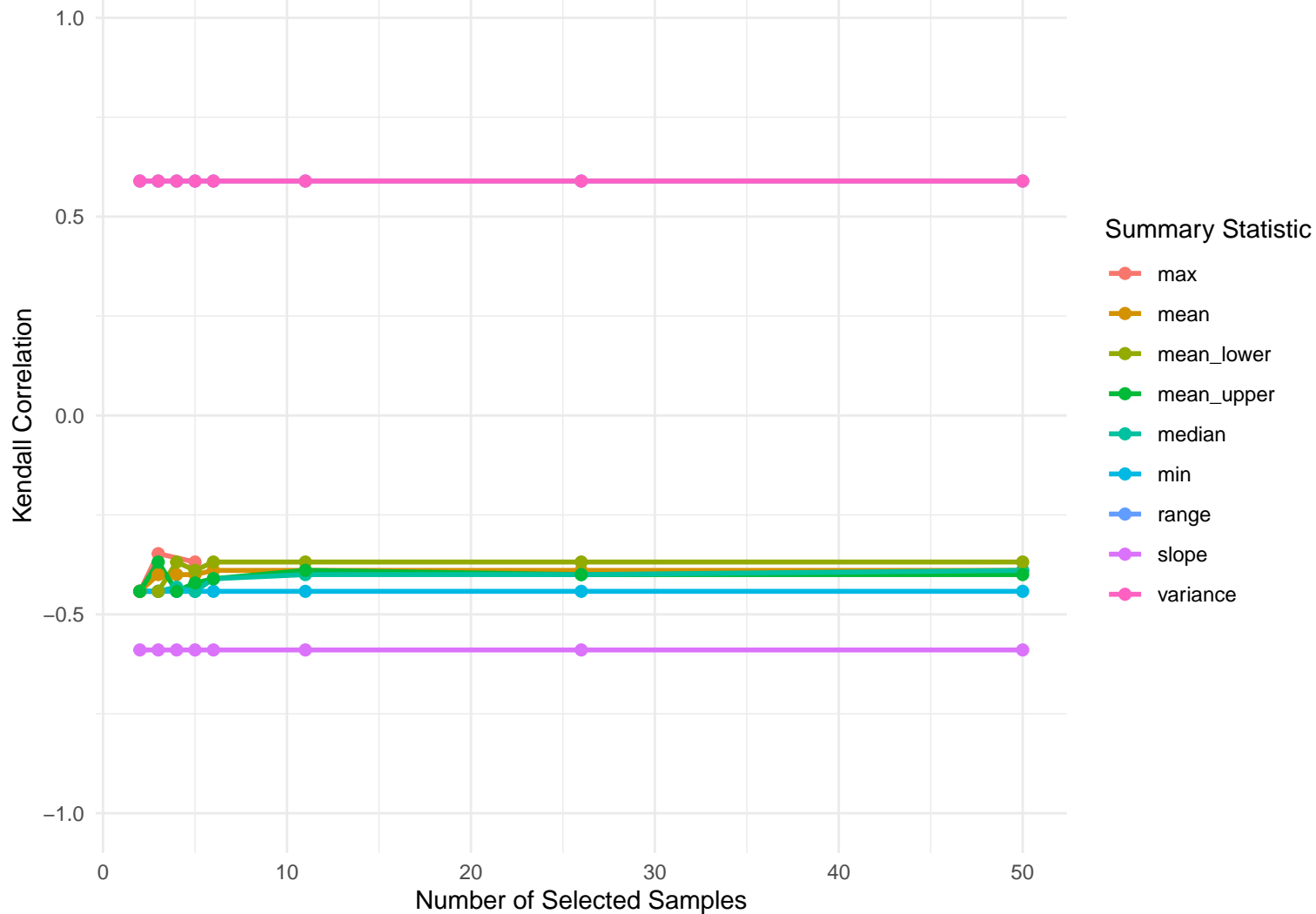
Number of Selected Samples

Summary Statistic

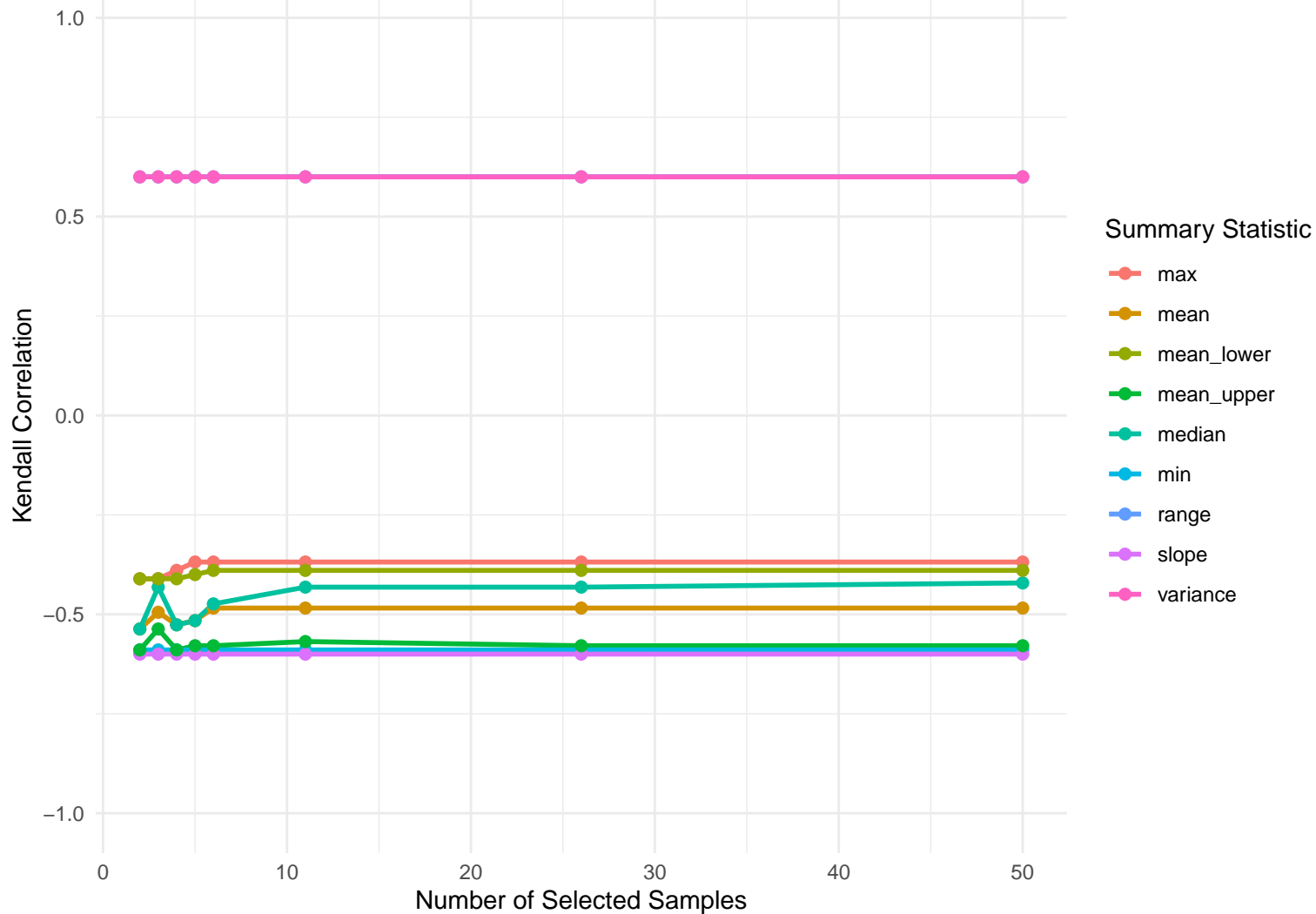
- min
- range
- variance



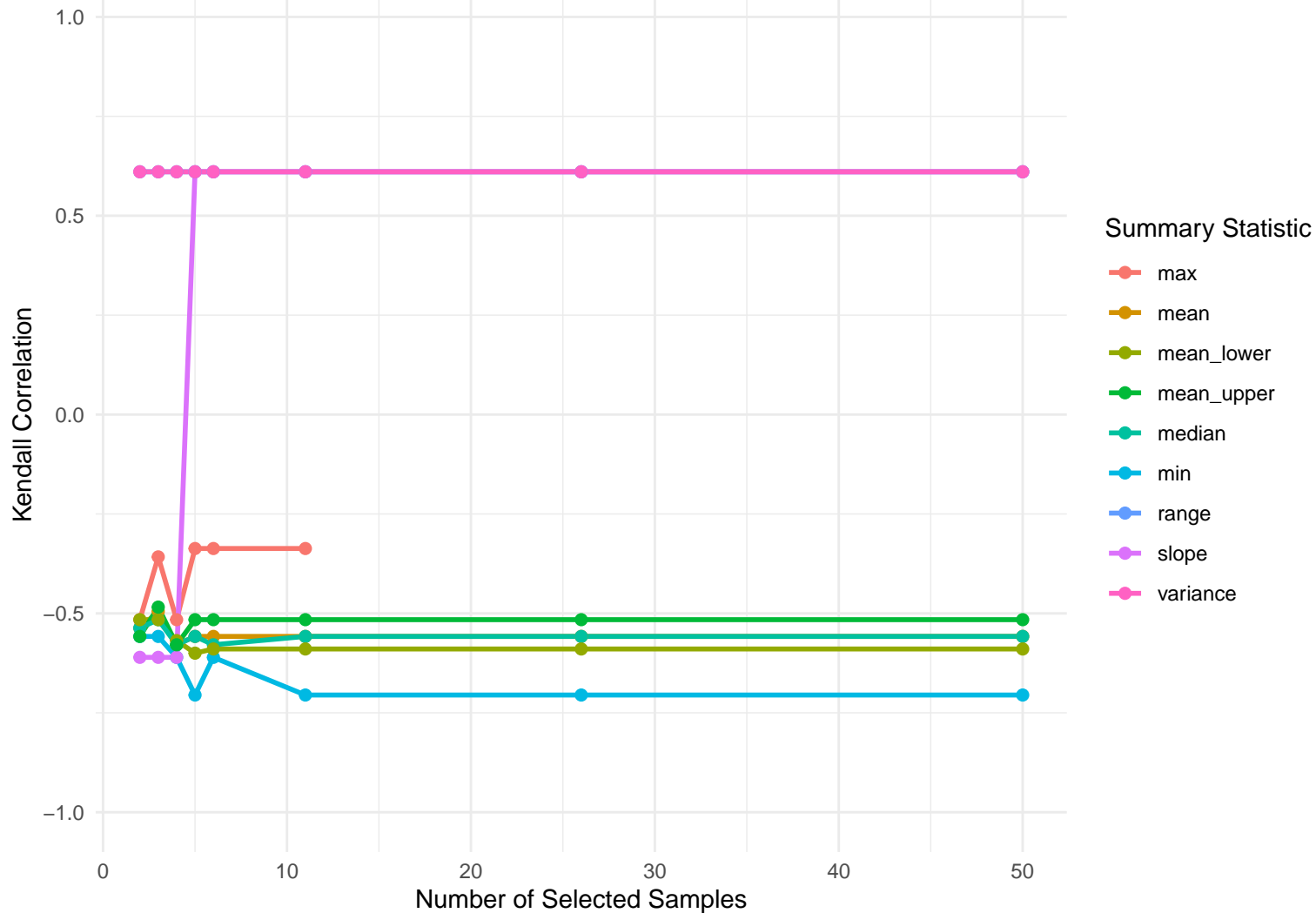
Score: Plmd (Local, full , Genotype: gaussian)



Score: Plmd (Local, full , Genotype: sinusoidal)



Score: Plmd (Local, full , Genotype: wave)



Score: PILSM (Local, full , Genotype: linear)

Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

0

10

20

30

40

50

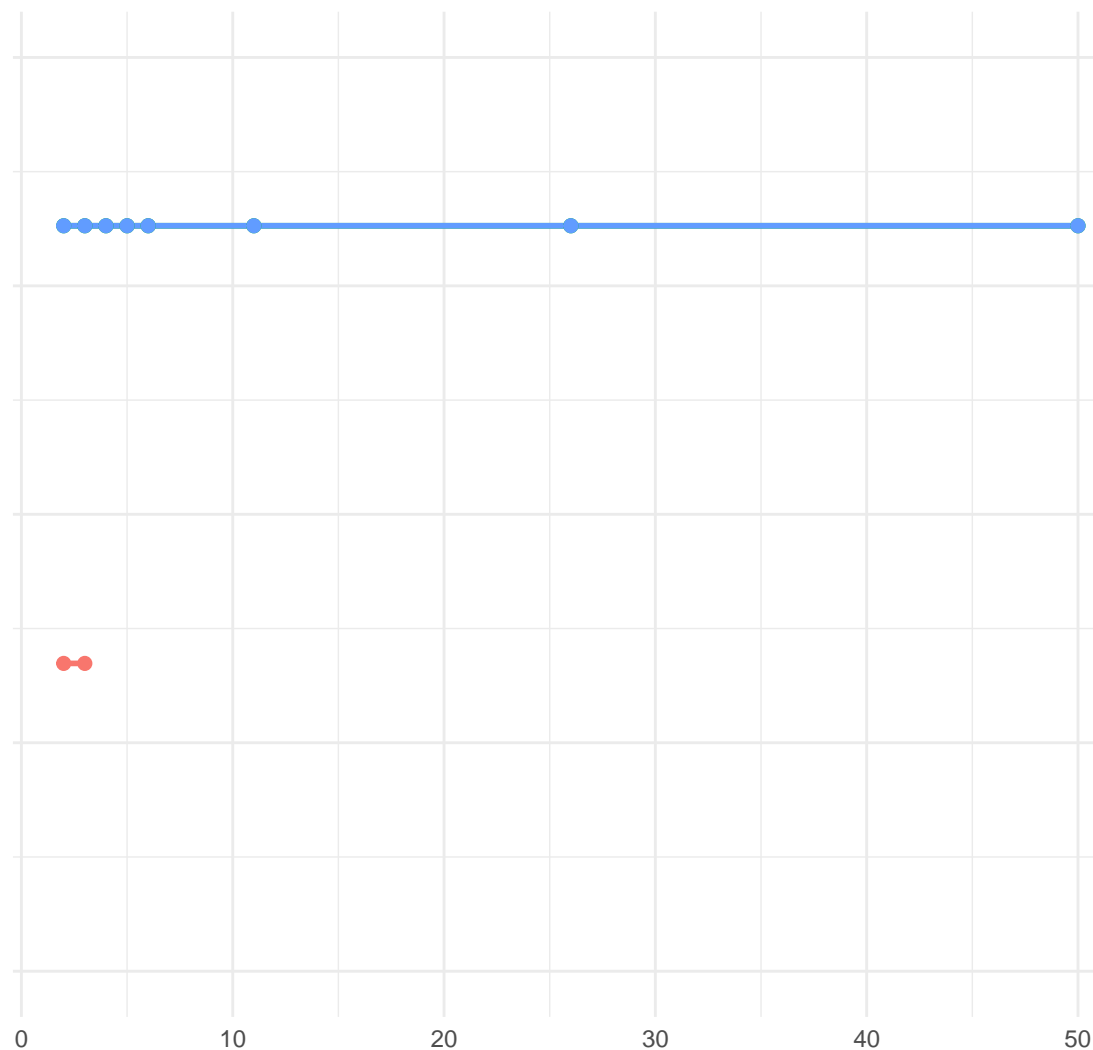
Number of Selected Samples

Summary Statistic

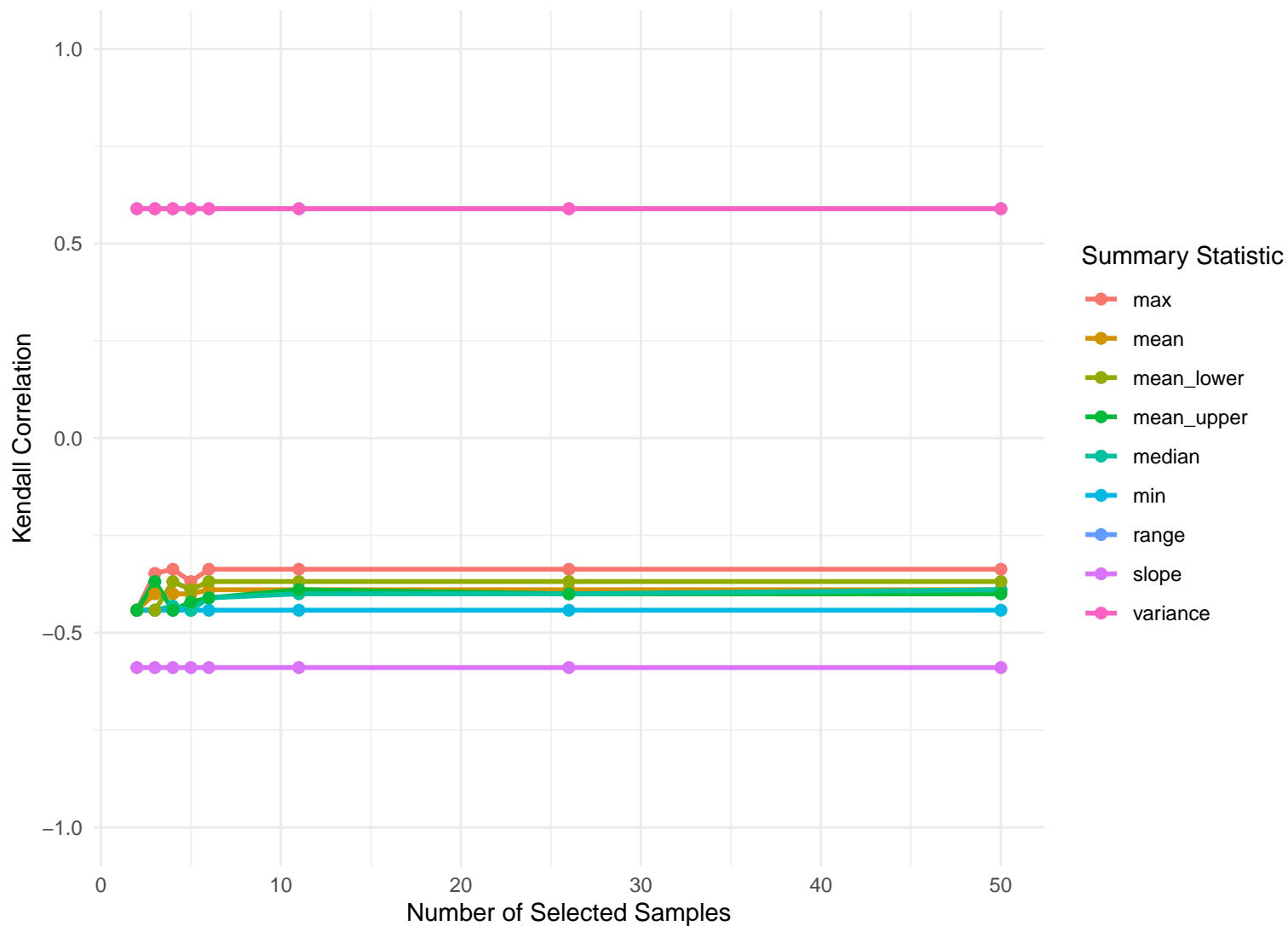
min

range

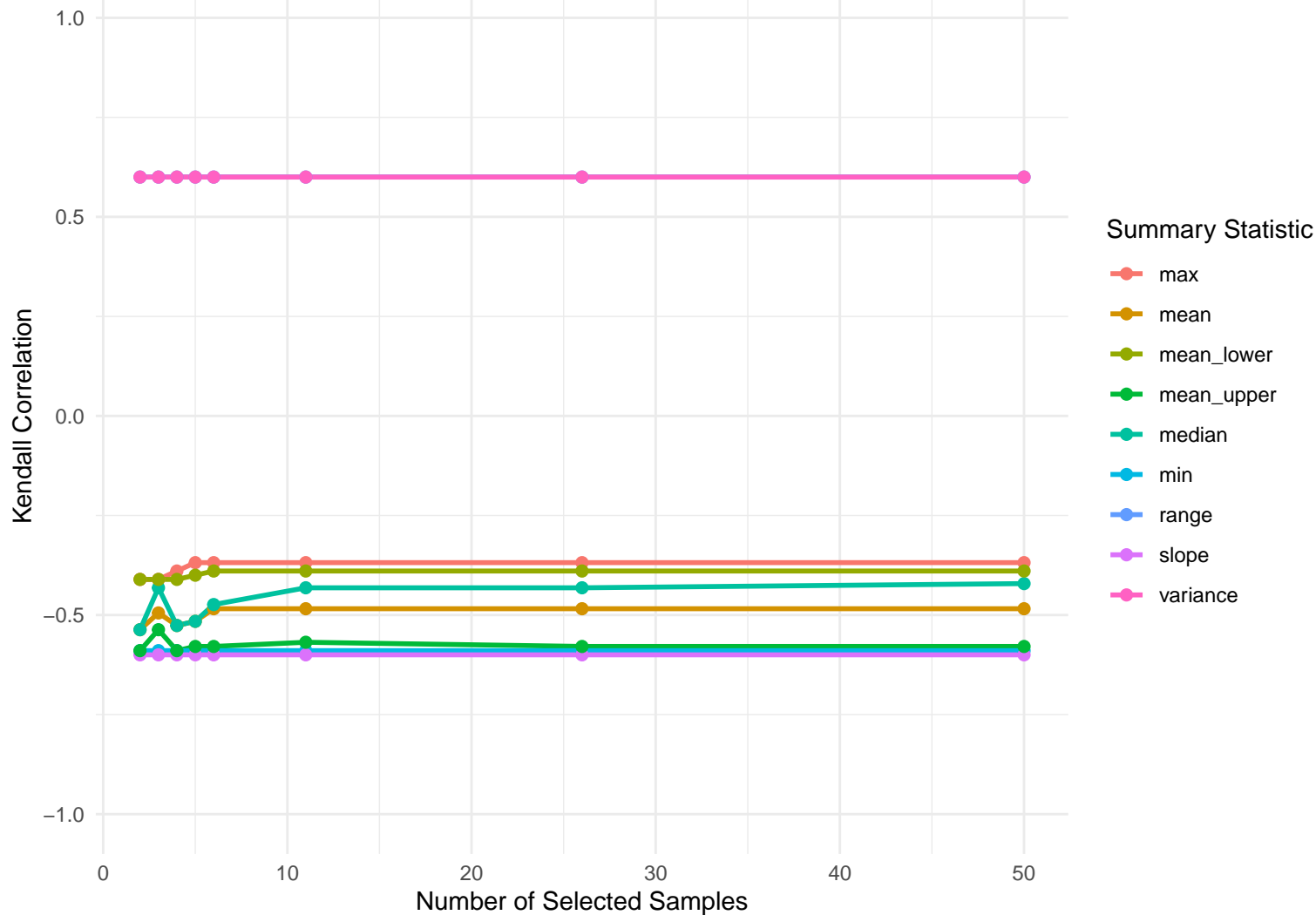
variance



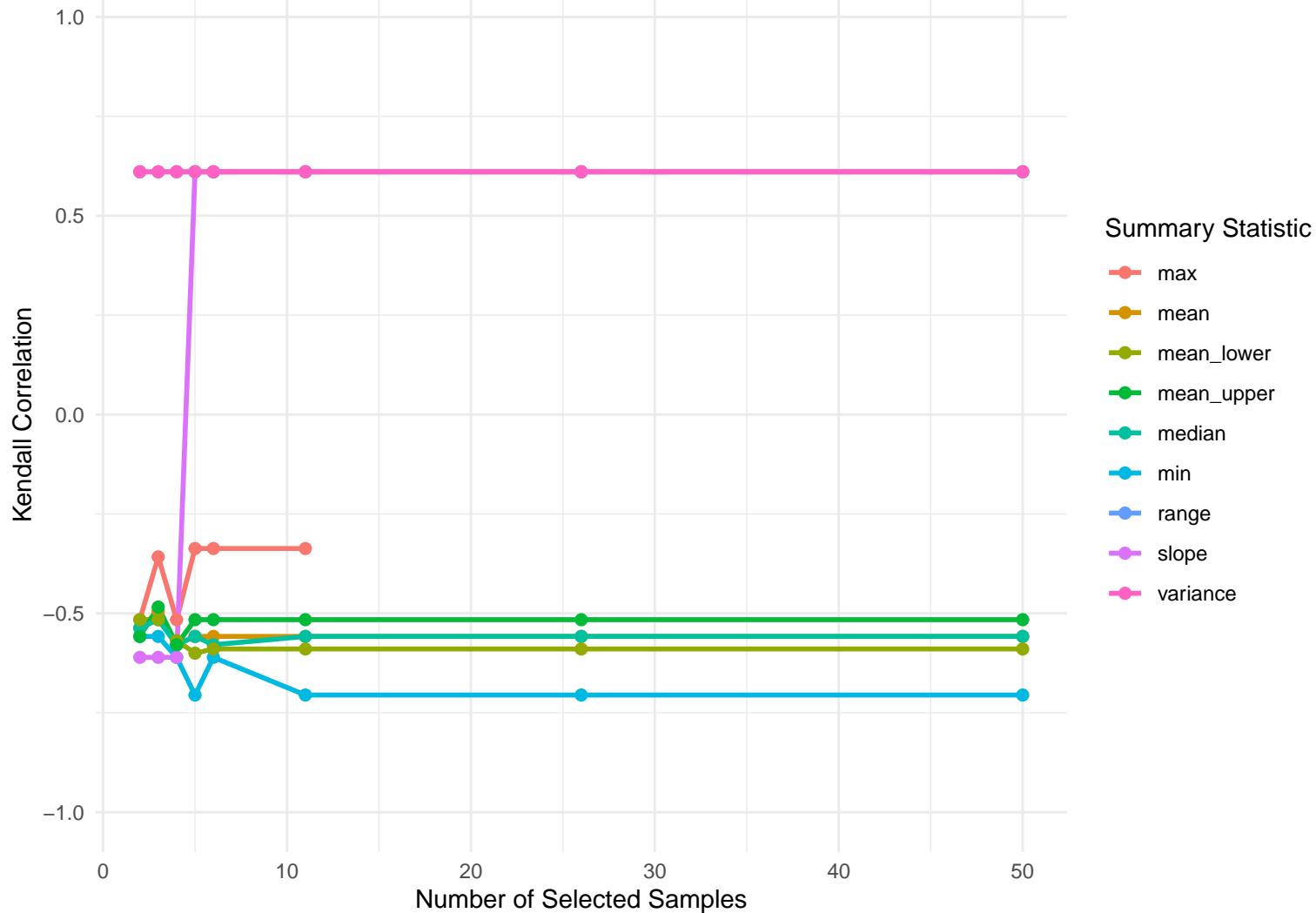
Score: PILSM (Local, full , Genotype: gaussian)



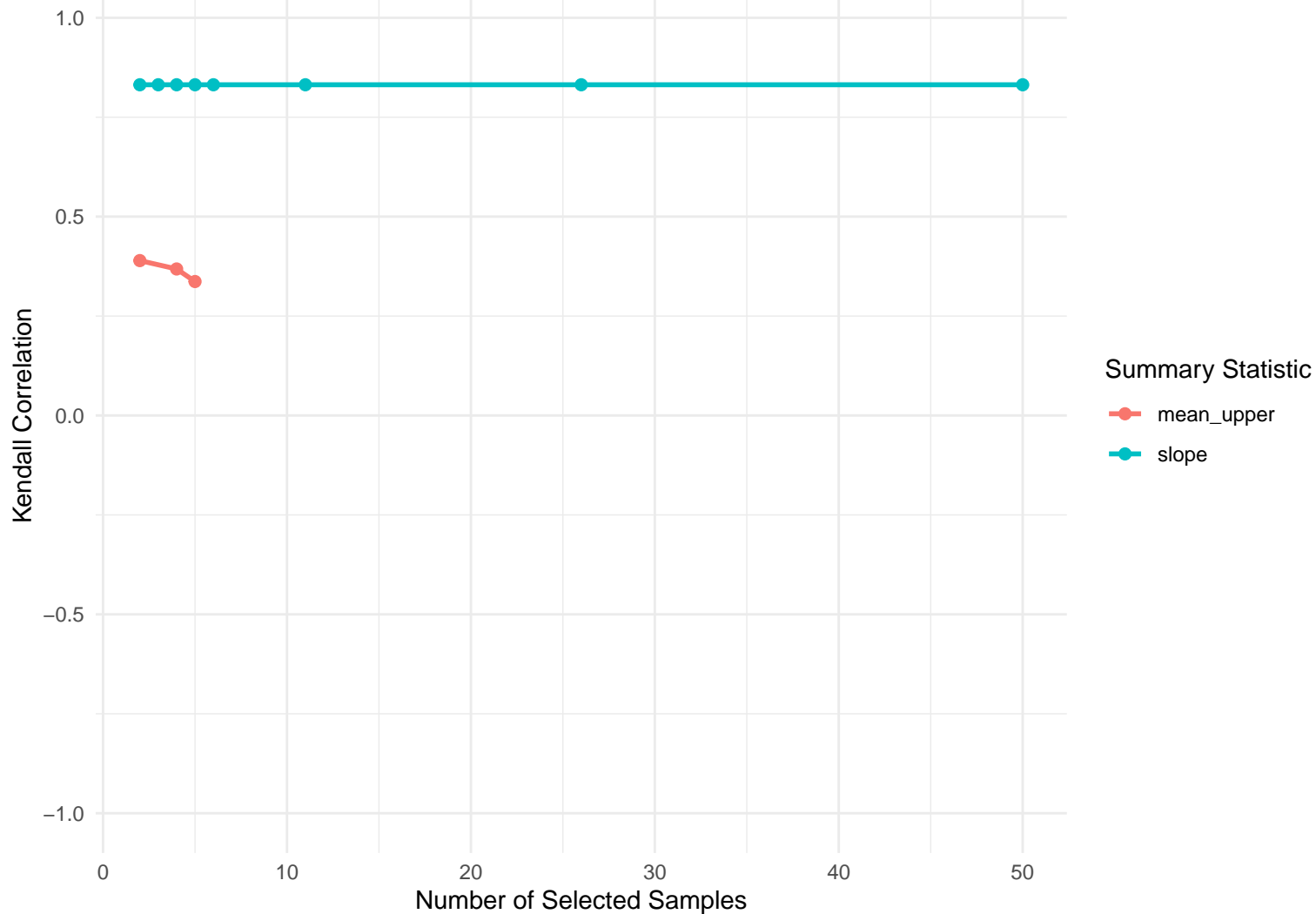
Score: PILSM (Local, full , Genotype: sinusoidal)



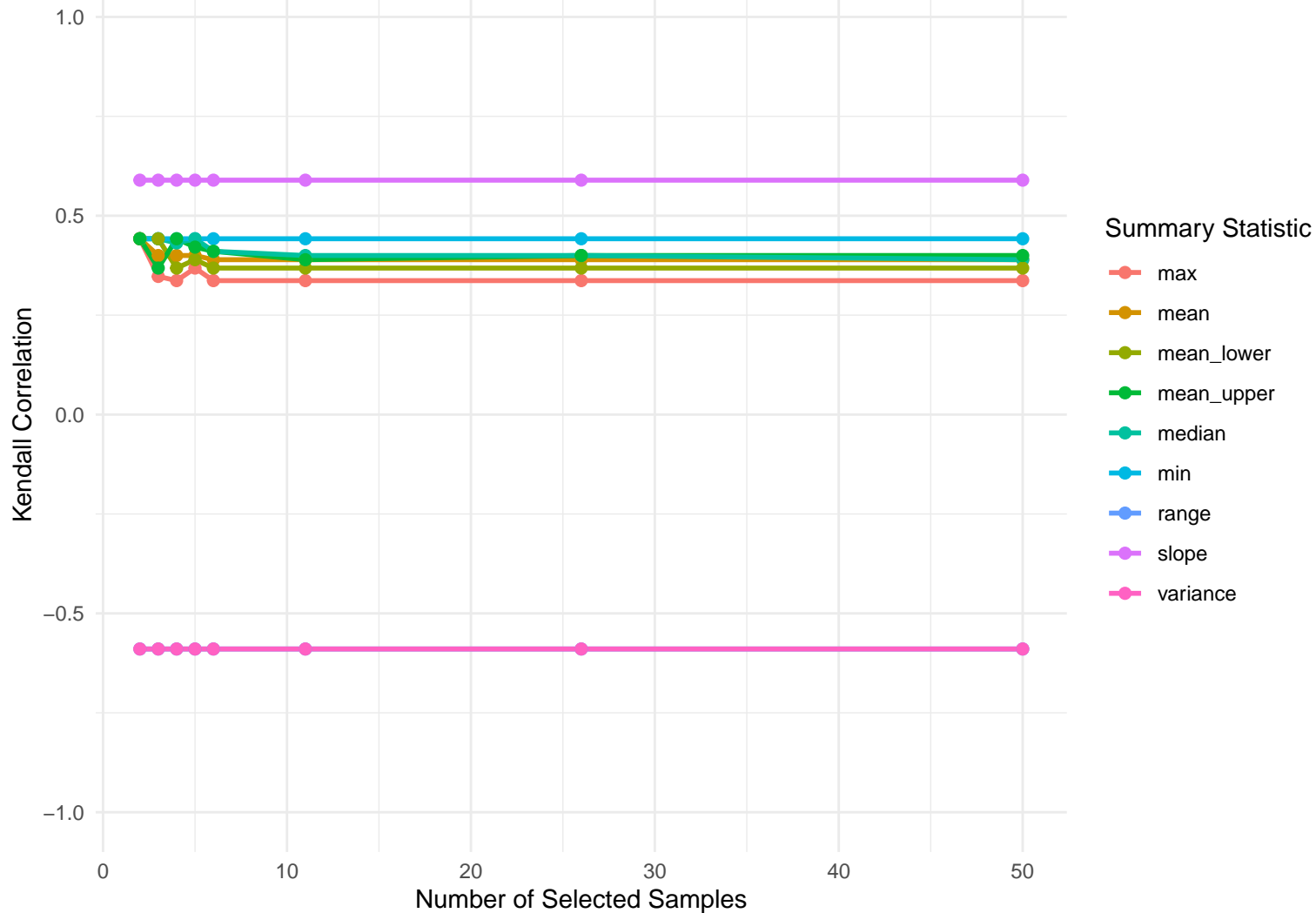
Score: PILSM (Local, full , Genotype: wave)



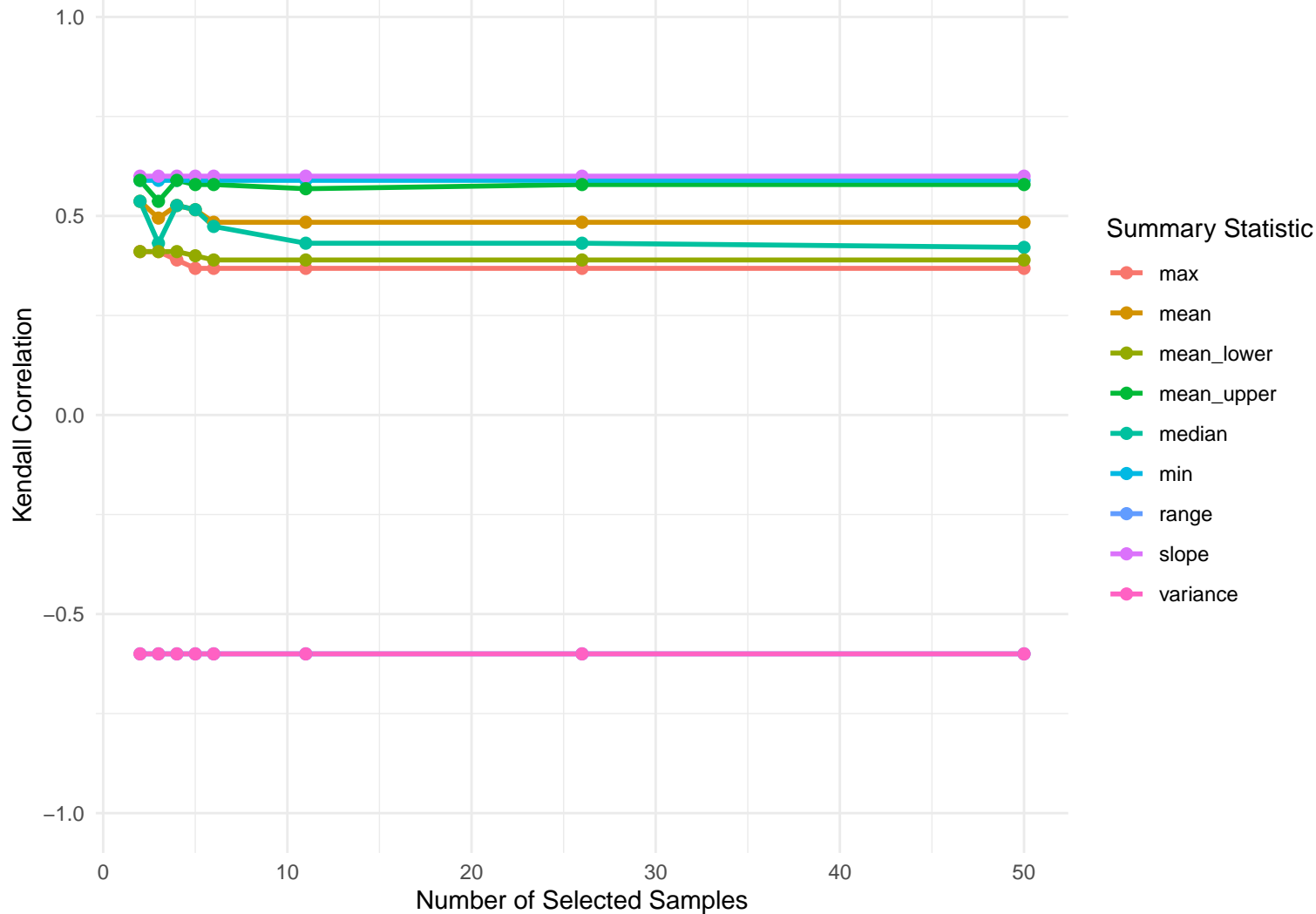
Score: RTR (Local, full , Genotype: linear)



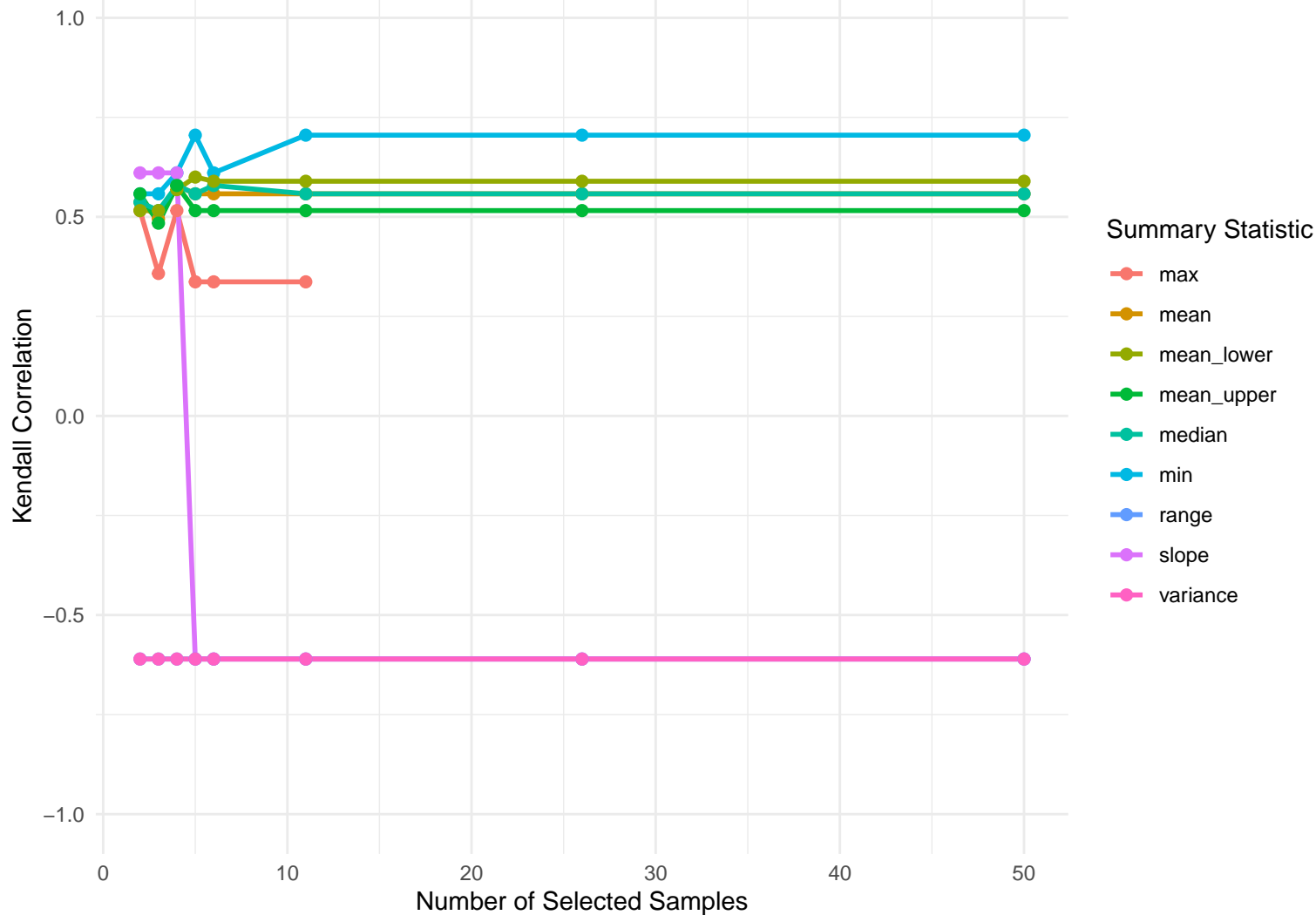
Score: RTR (Local, full , Genotype: gaussian)



Score: RTR (Local, full , Genotype: sinusoidal)

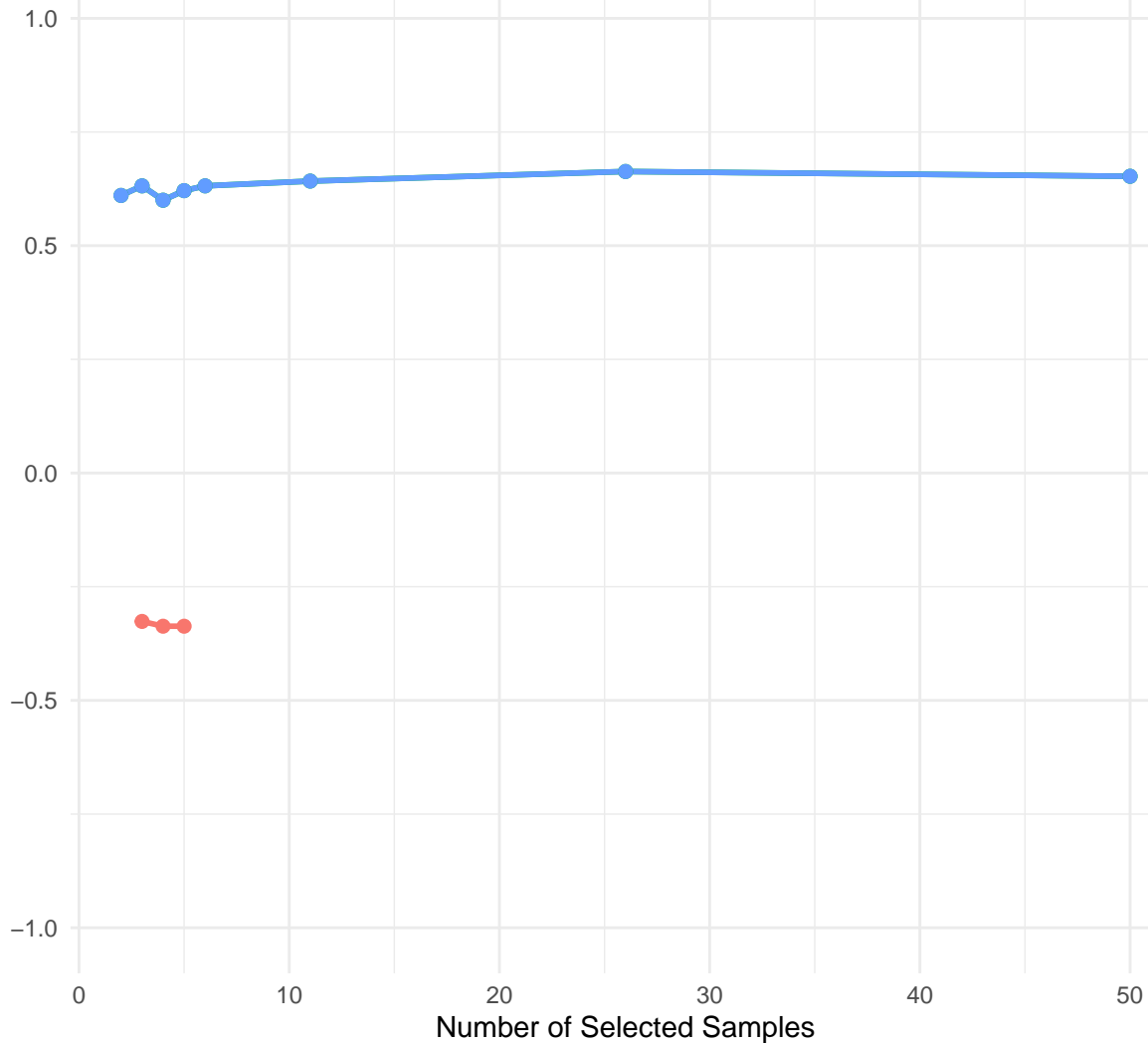


Score: RTR (Local, full , Genotype: wave)



Score: PIR (Local, full , Genotype: linear)

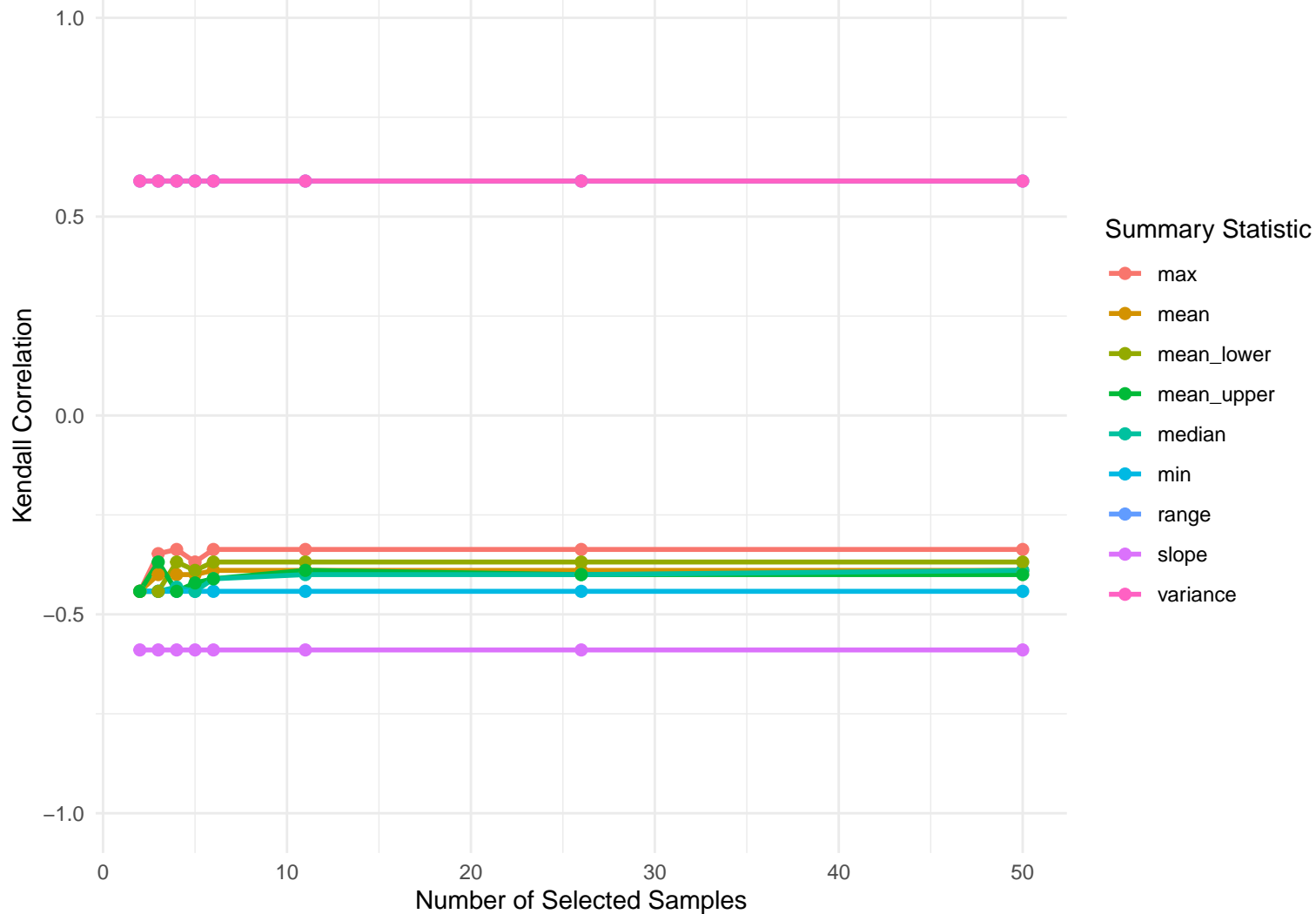
Kendall Correlation



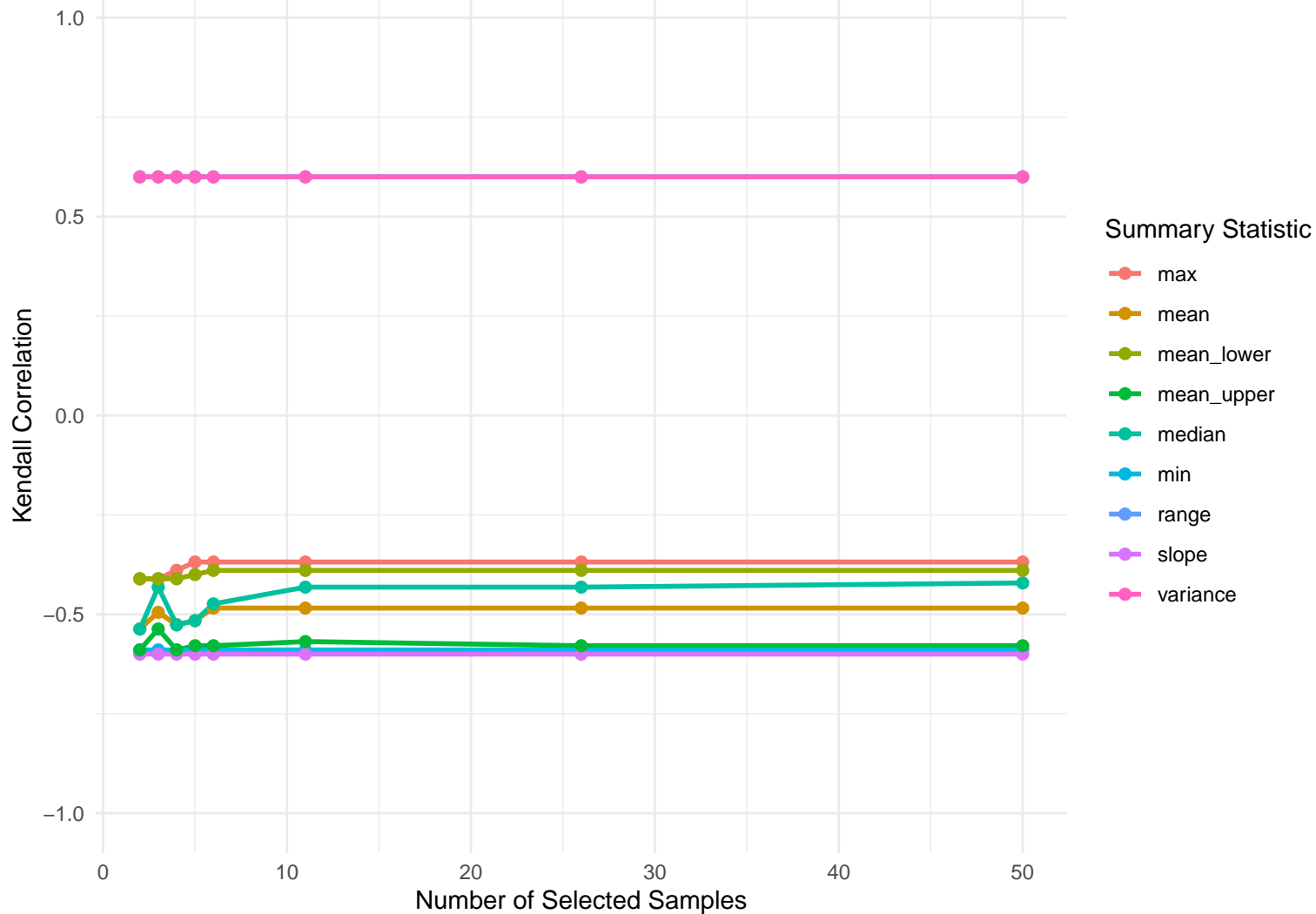
Summary Statistic

- min
- range
- variance

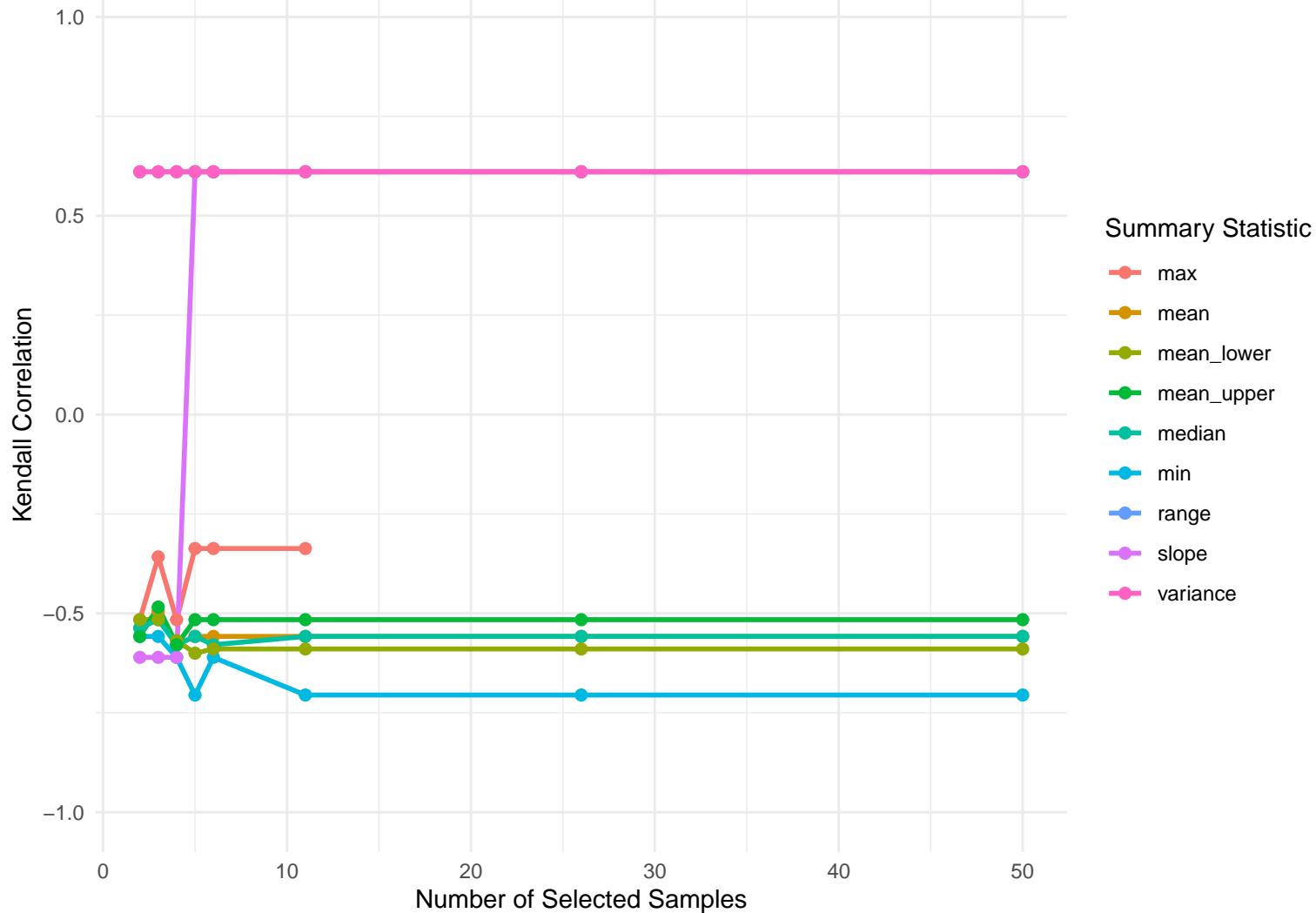
Score: PIR (Local, full , Genotype: gaussian)



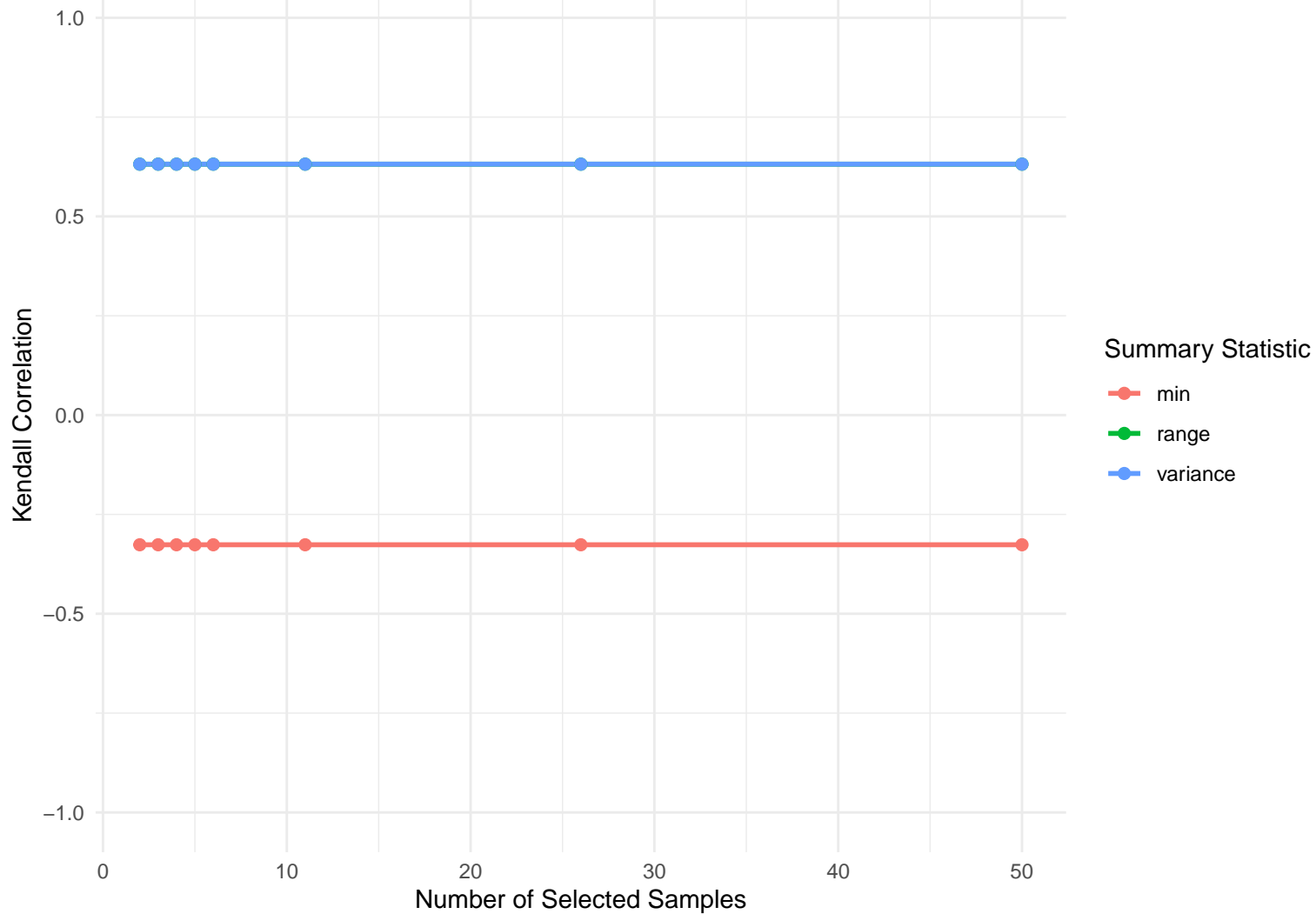
Score: PIR (Local, full , Genotype: sinusoidal)



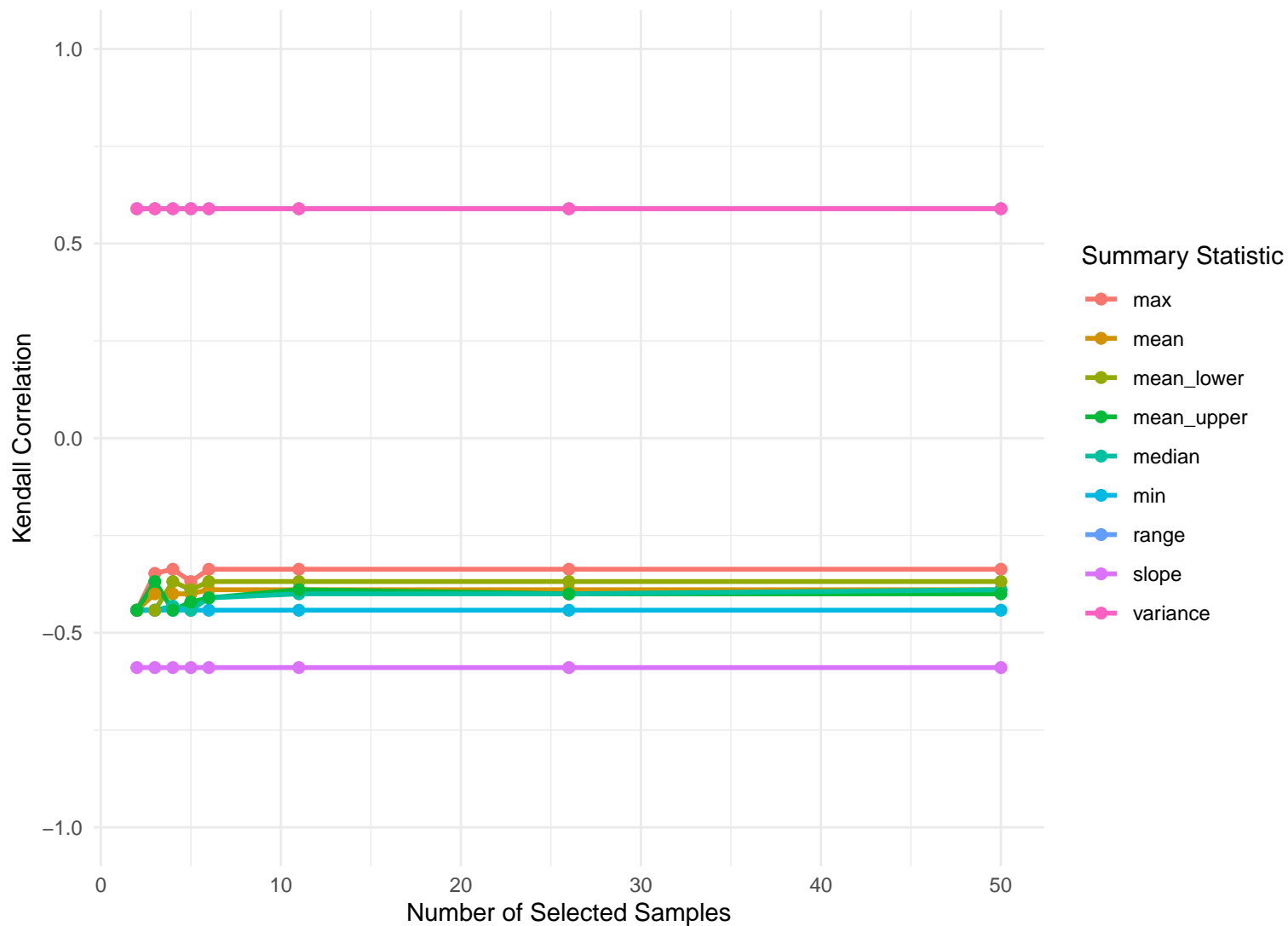
Score: PIR (Local, full , Genotype: wave)



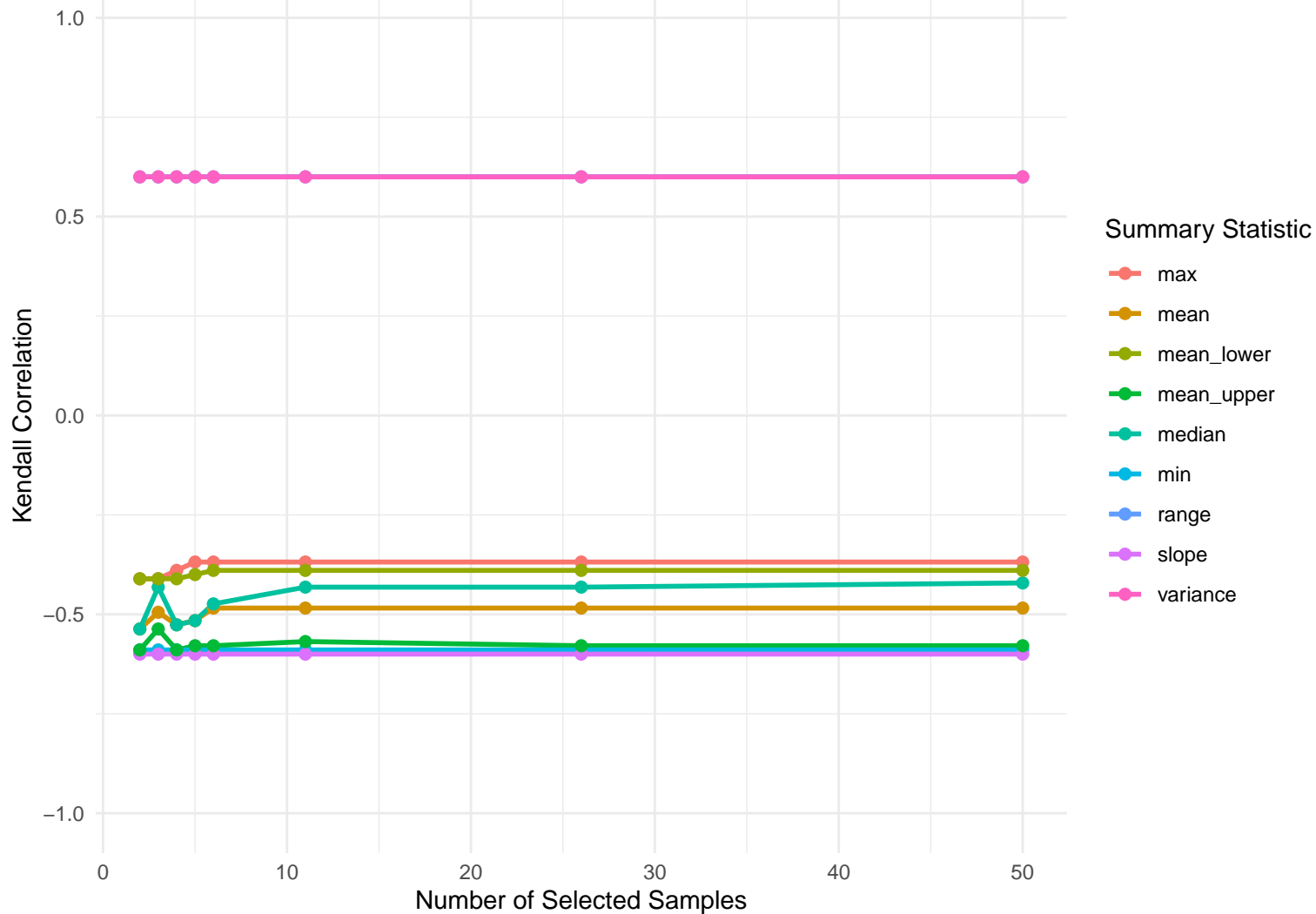
Score: RDPI (Local, full , Genotype: linear)



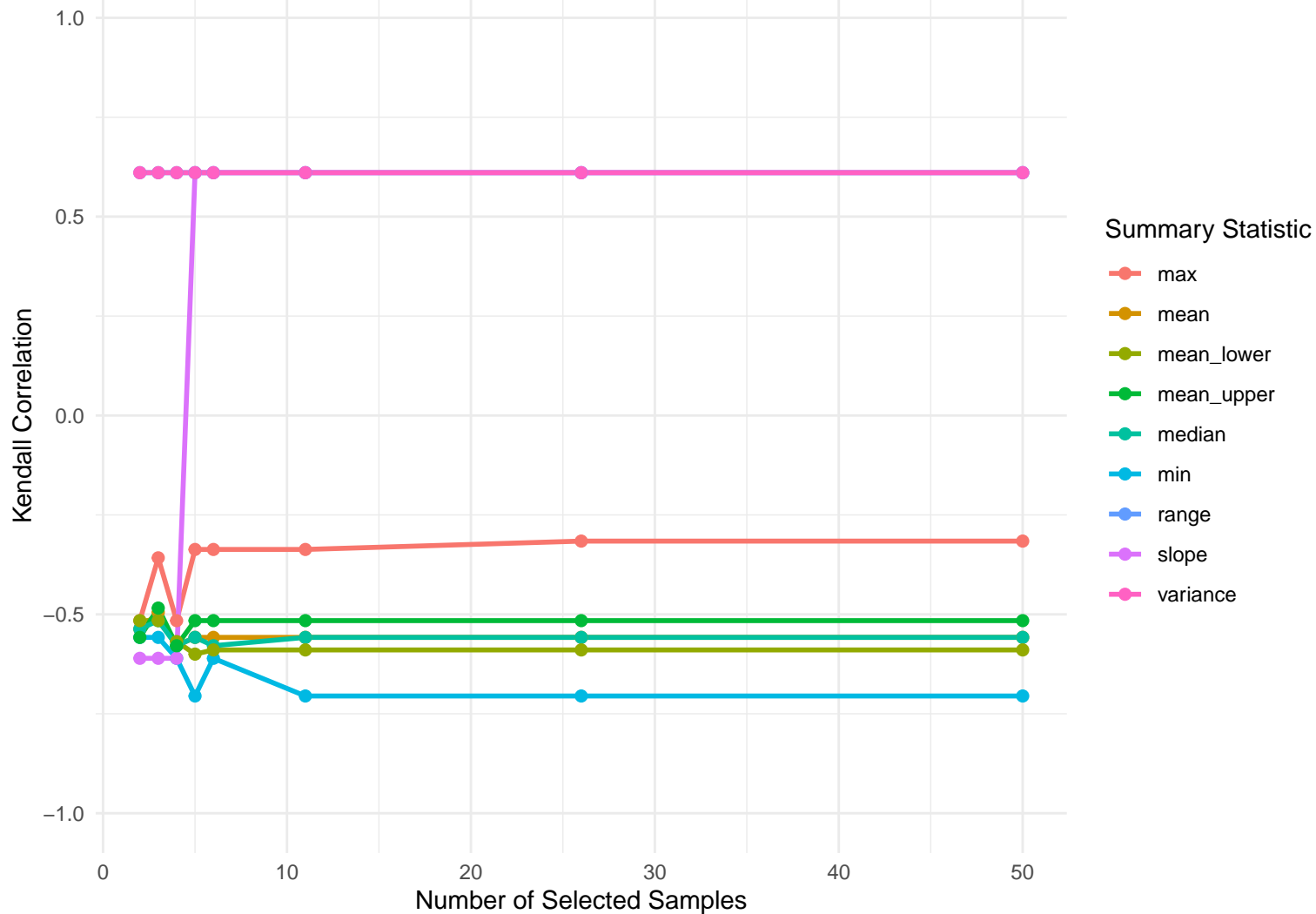
Score: RDPI (Local, full , Genotype: gaussian)



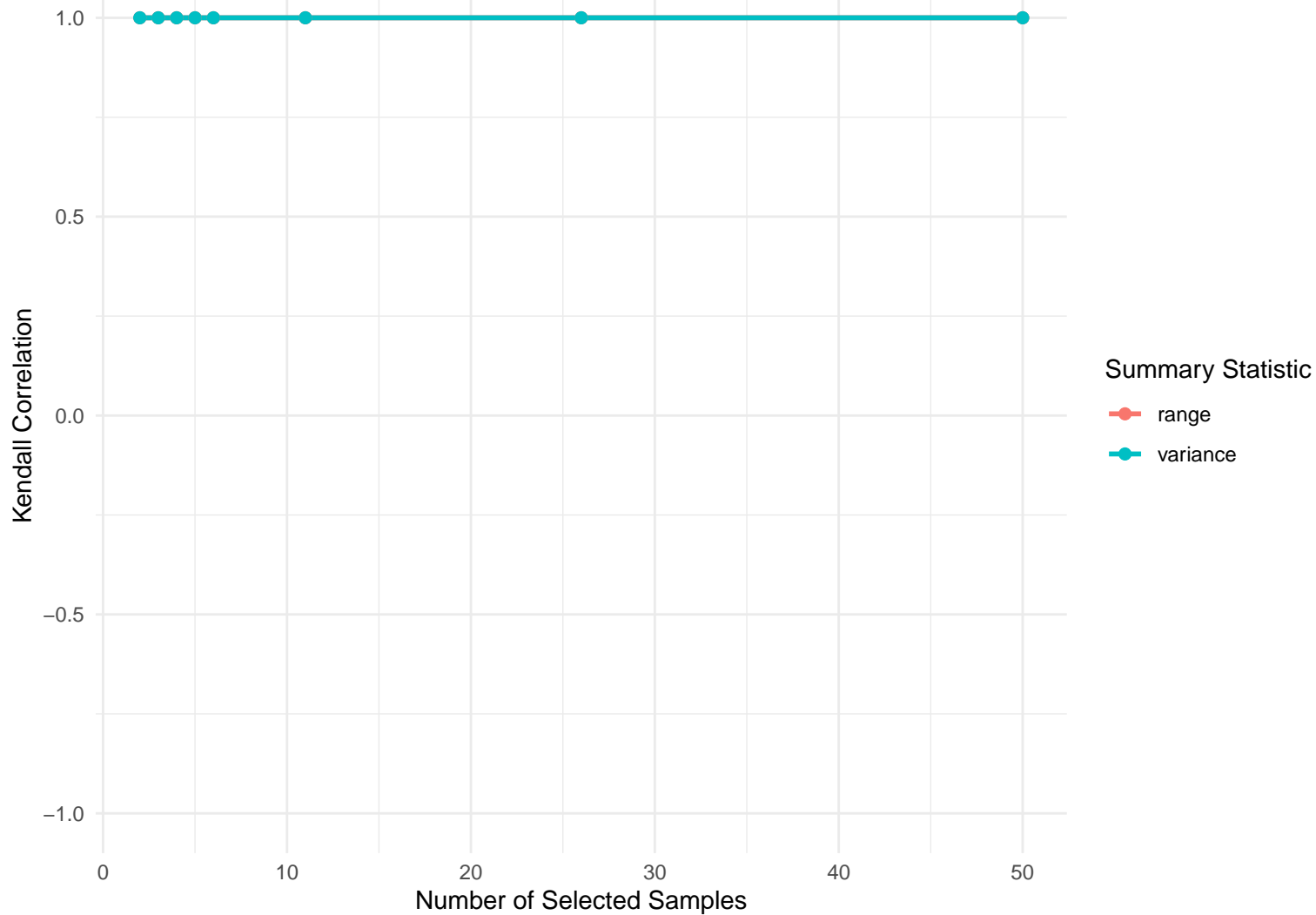
Score: RDPI (Local, full , Genotype: sinusoidal)



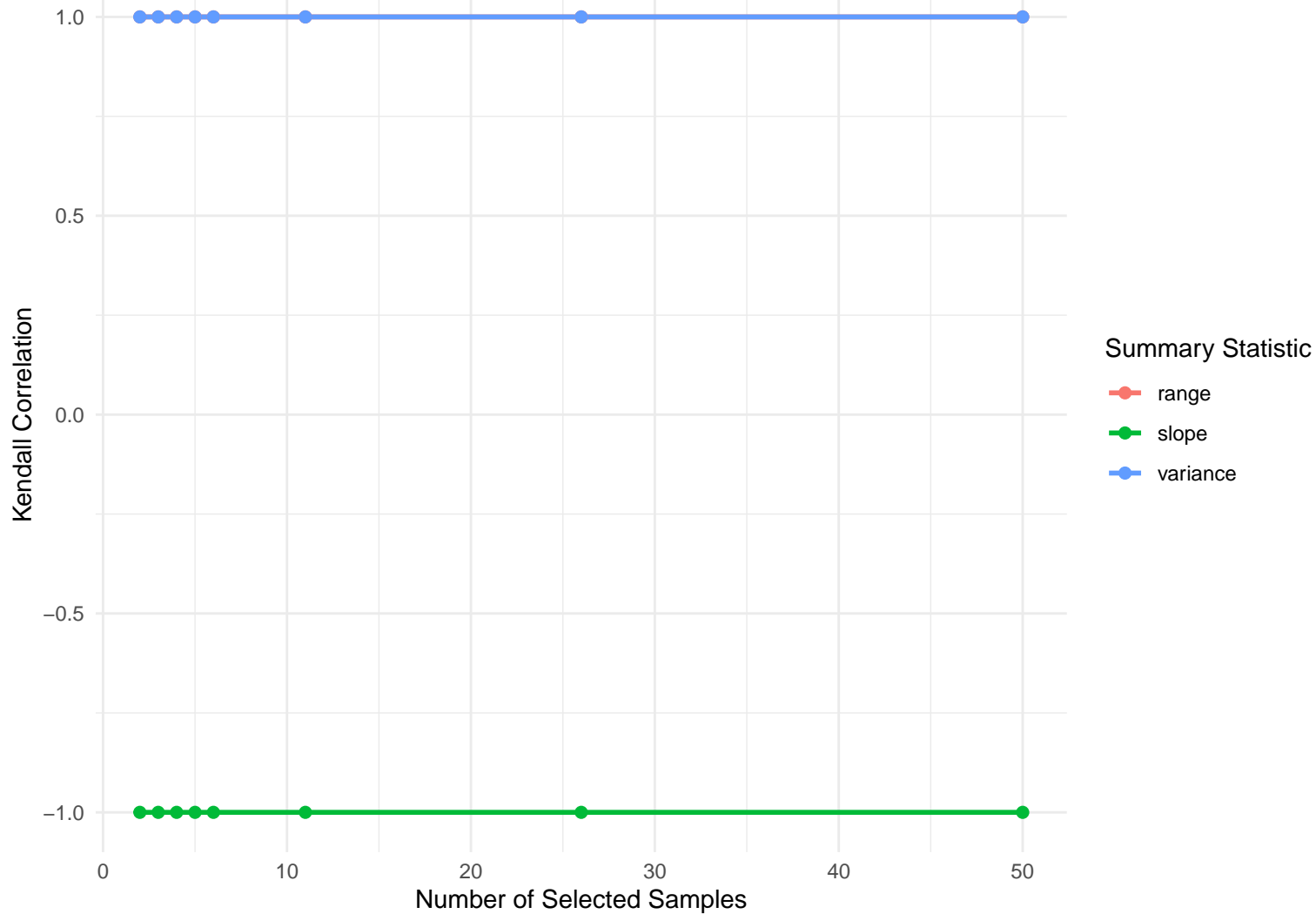
Score: RDPI (Local, full , Genotype: wave)



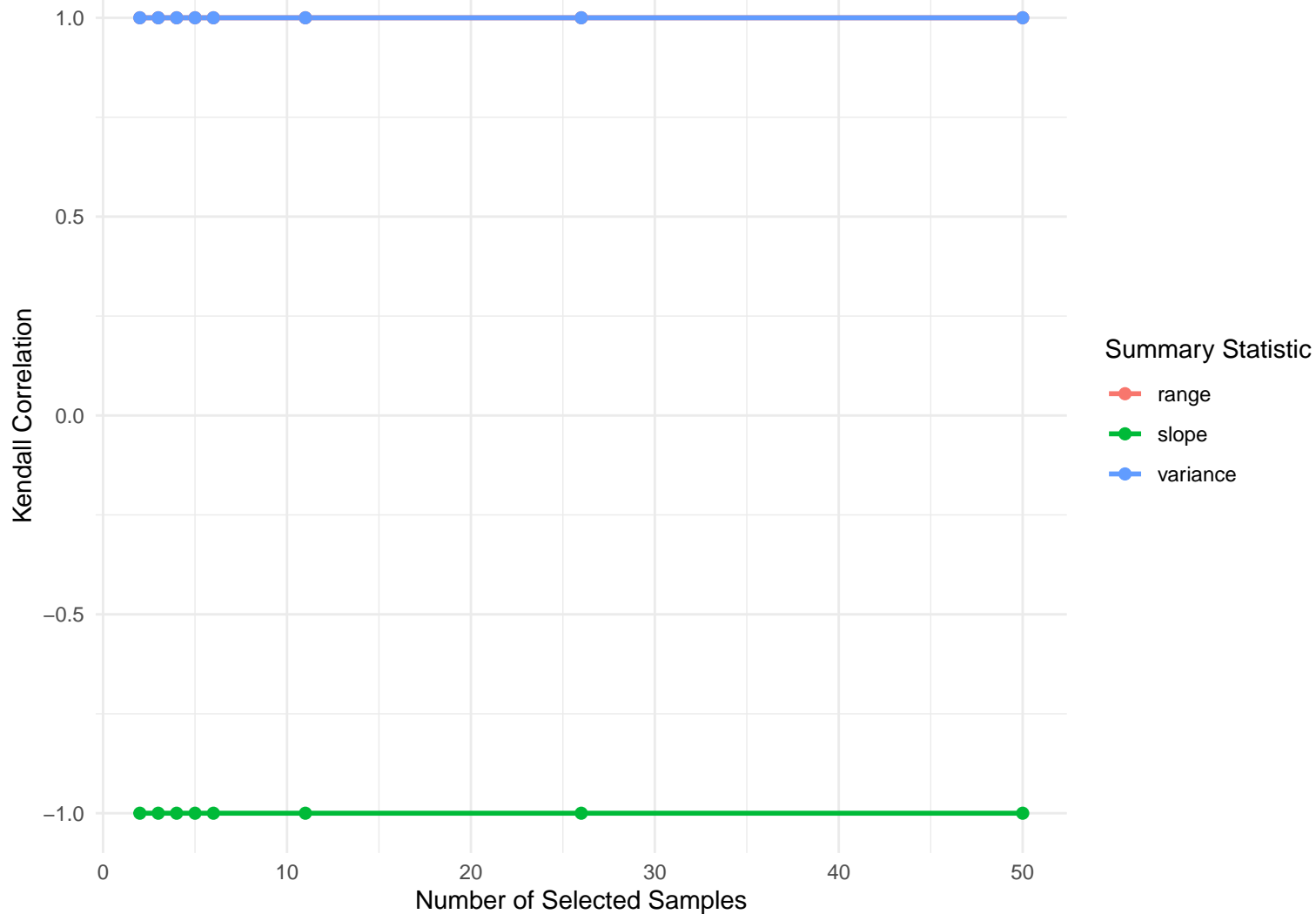
Score: ESPI (Local, full , Genotype: linear)



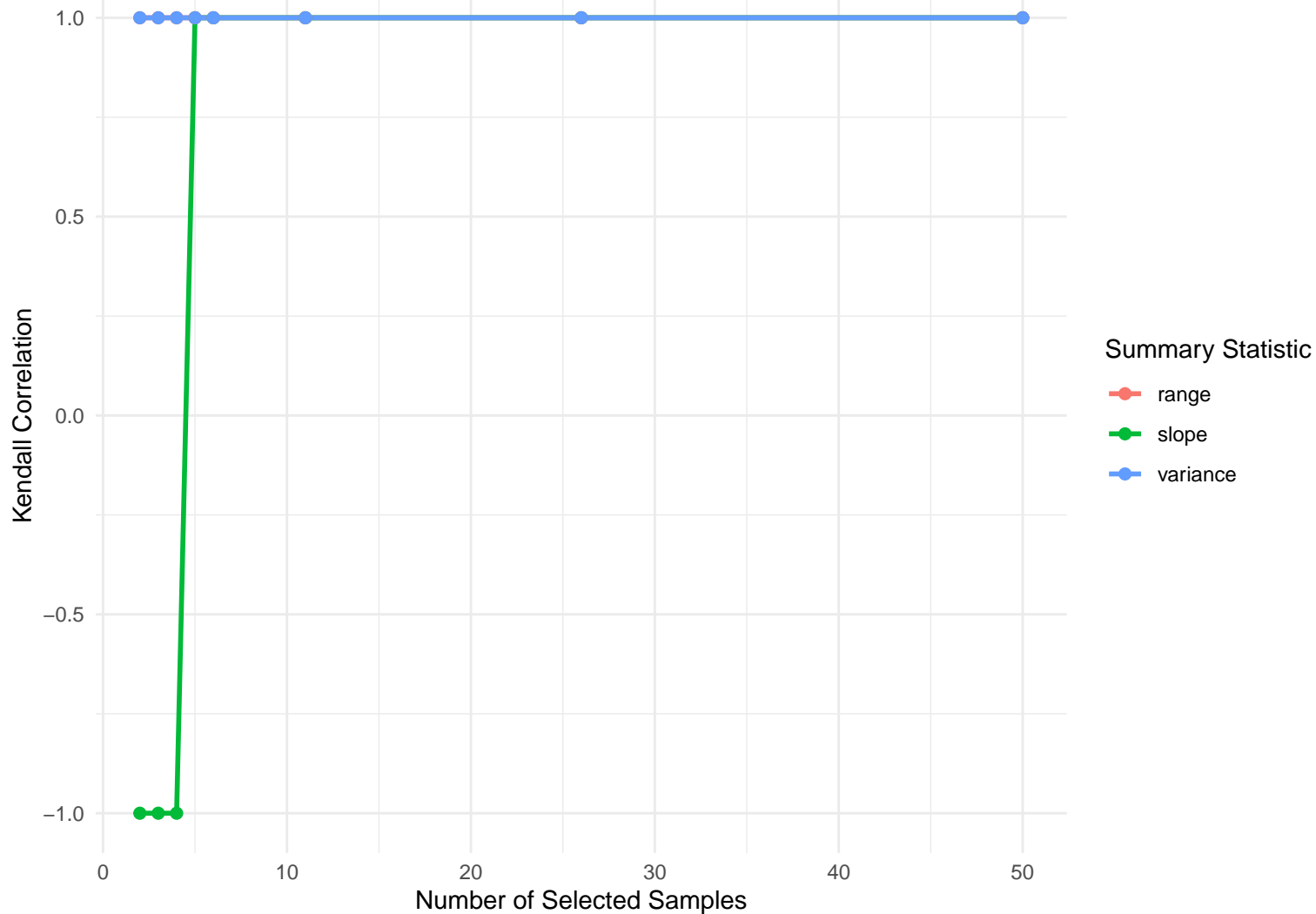
Score: ESPI (Local, full , Genotype: gaussian)



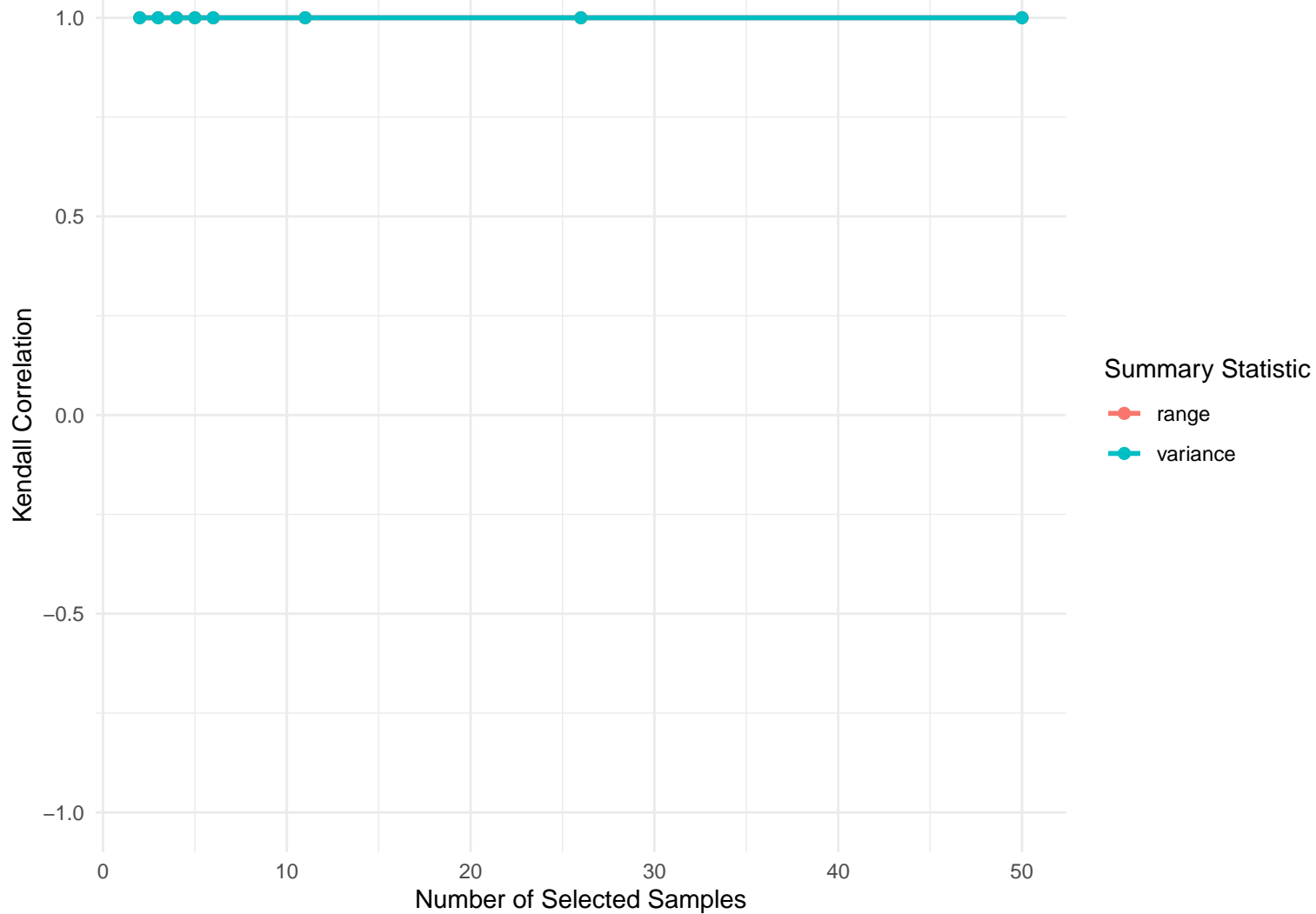
Score: ESPI (Local, full , Genotype: sinusoidal)



Score: ESPI (Local, full , Genotype: wave)



Score: ESPIID (Local, full , Genotype: linear)

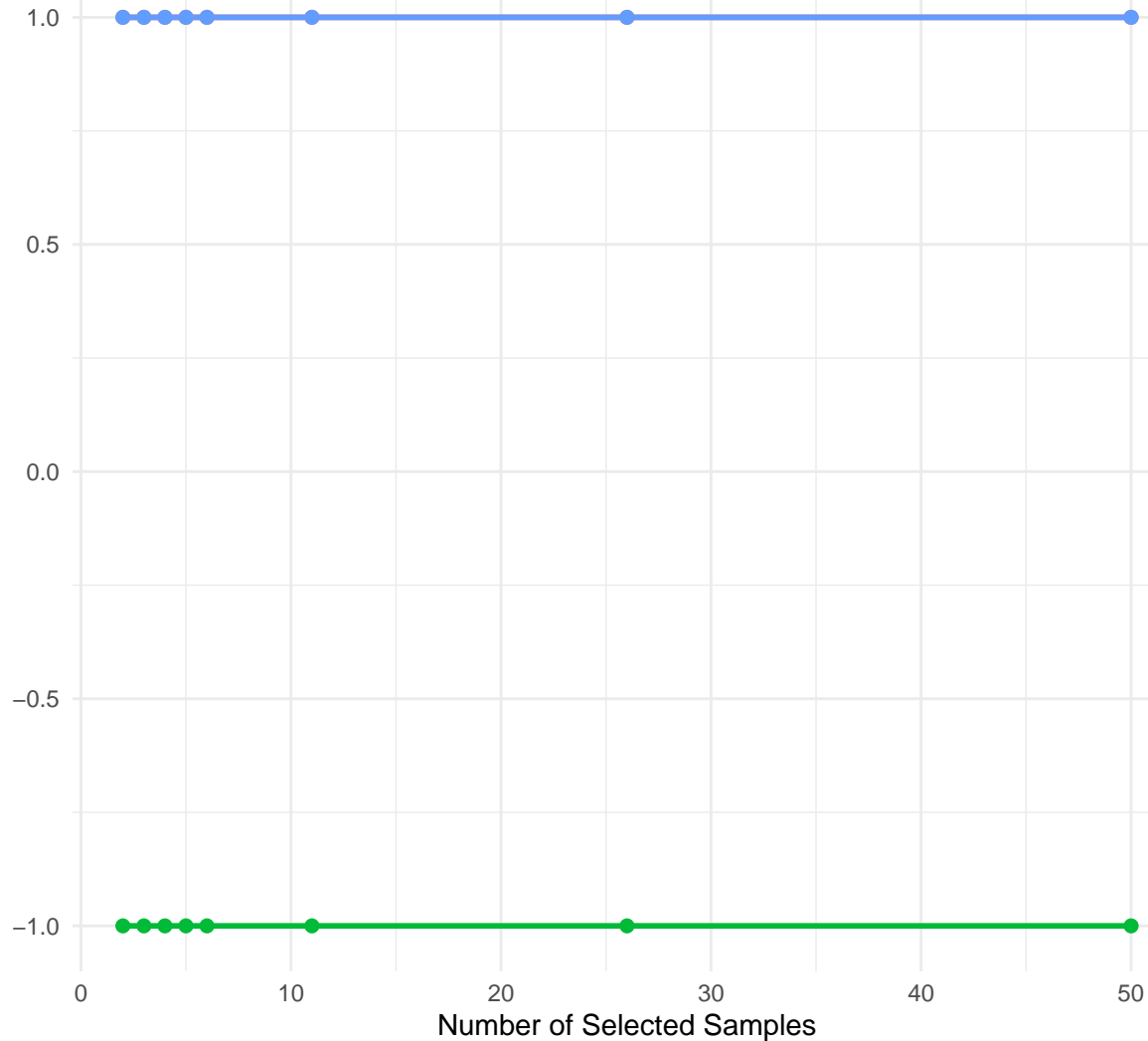


Score: ESPIID (Local, full , Genotype: gaussian)

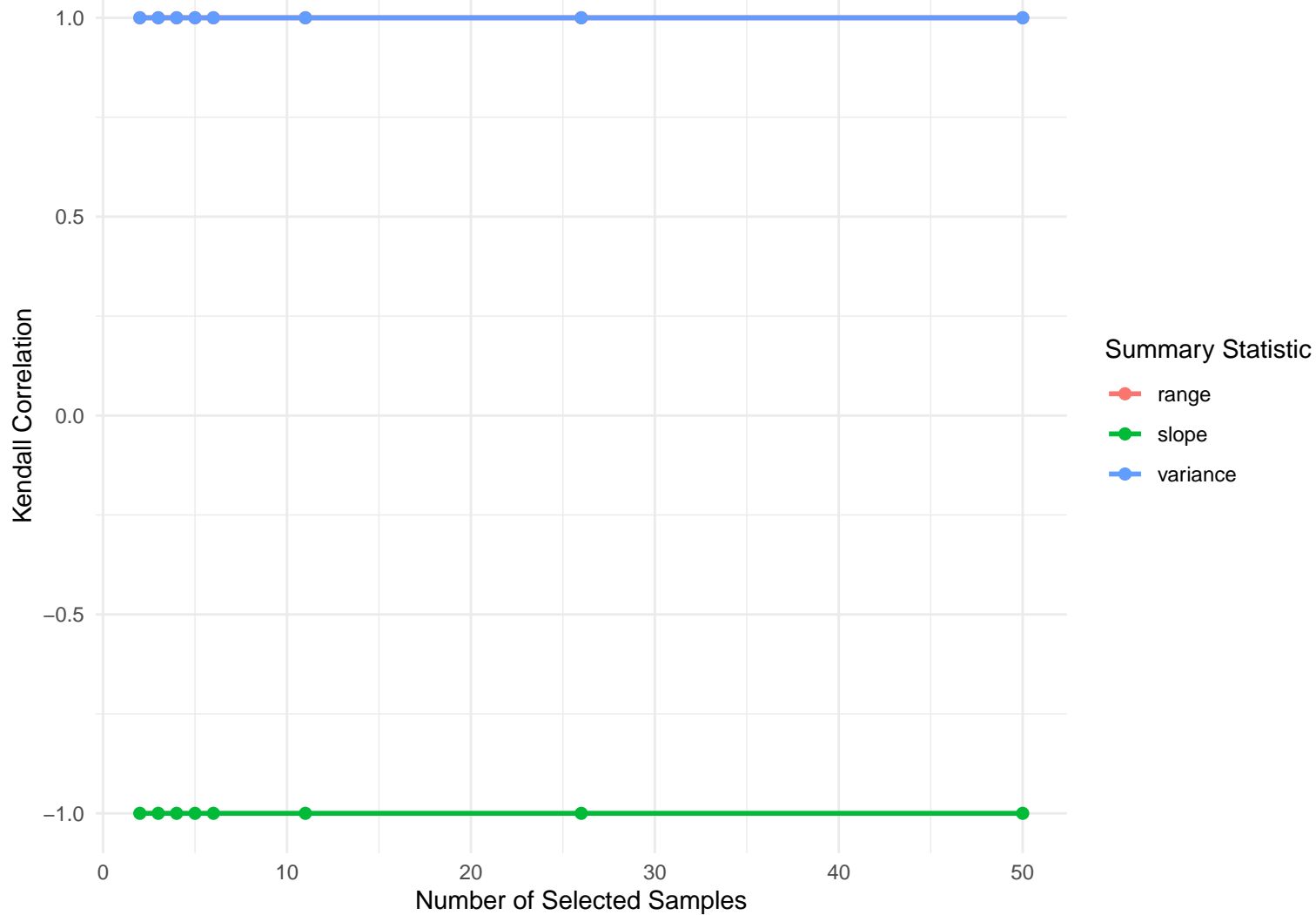
Kendall Correlation

Summary Statistic

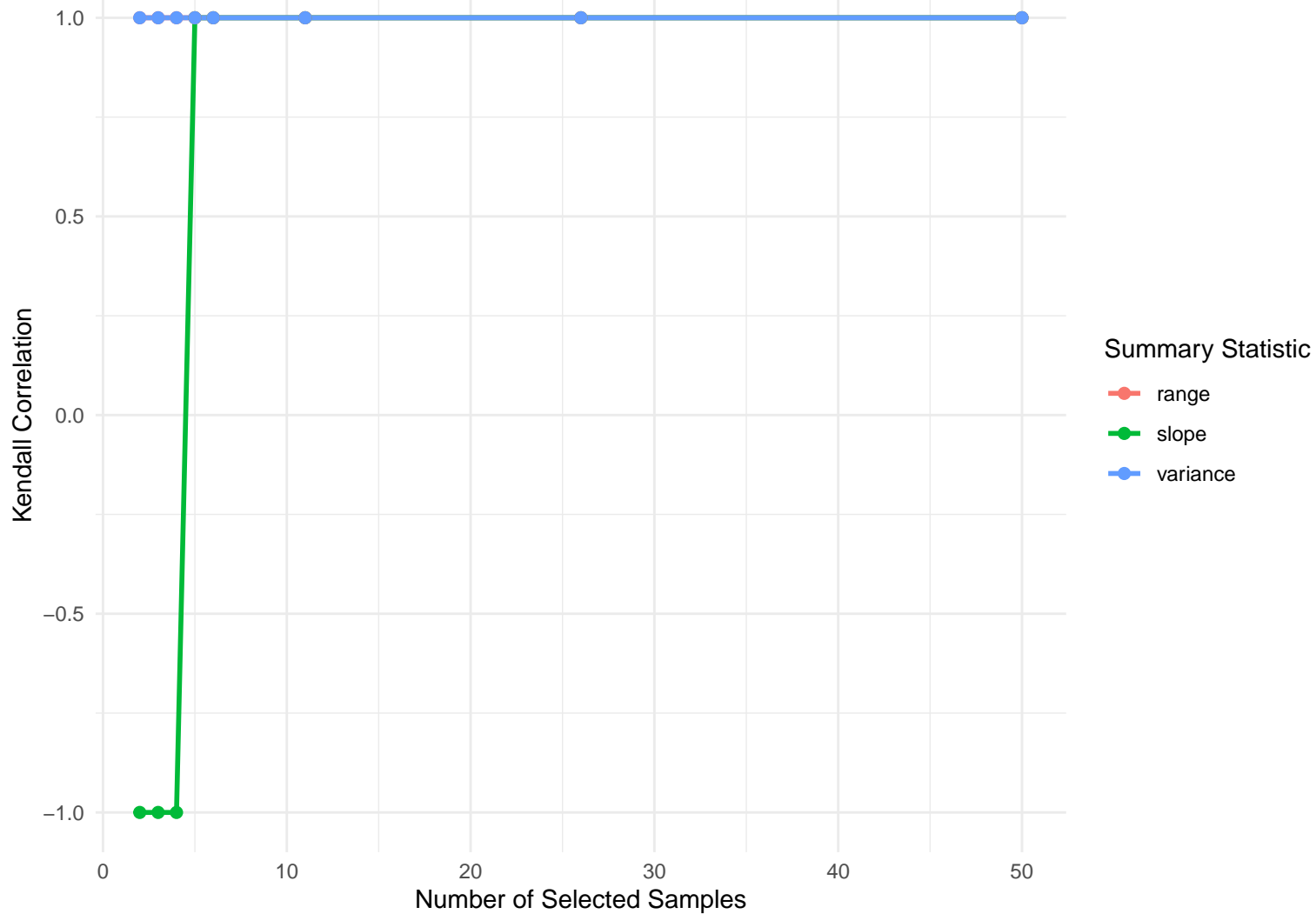
- range
- slope
- variance



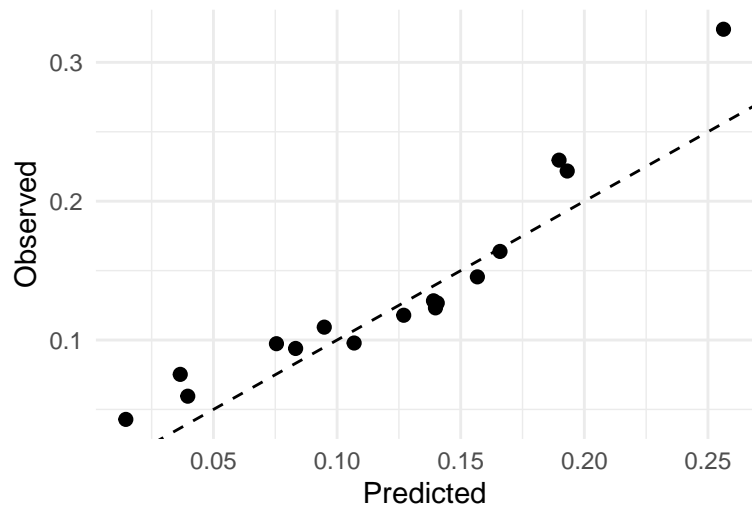
Score: ESPIID (Local, full , Genotype: sinusoidal)



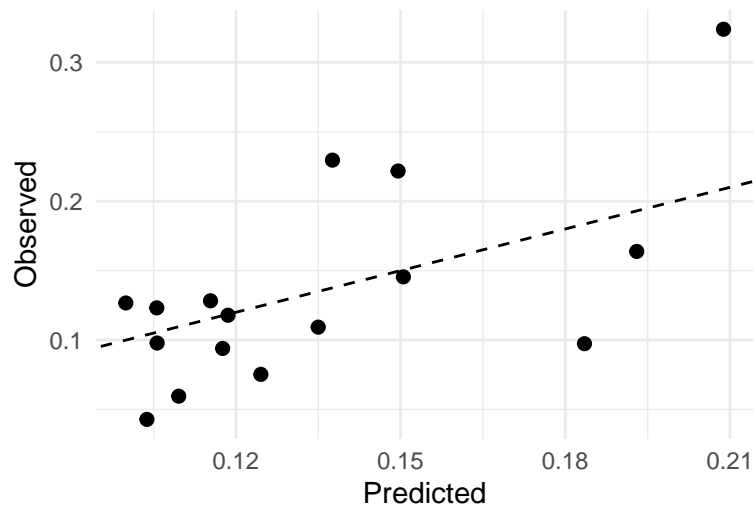
Score: ESPIID (Local, full , Genotype: wave)



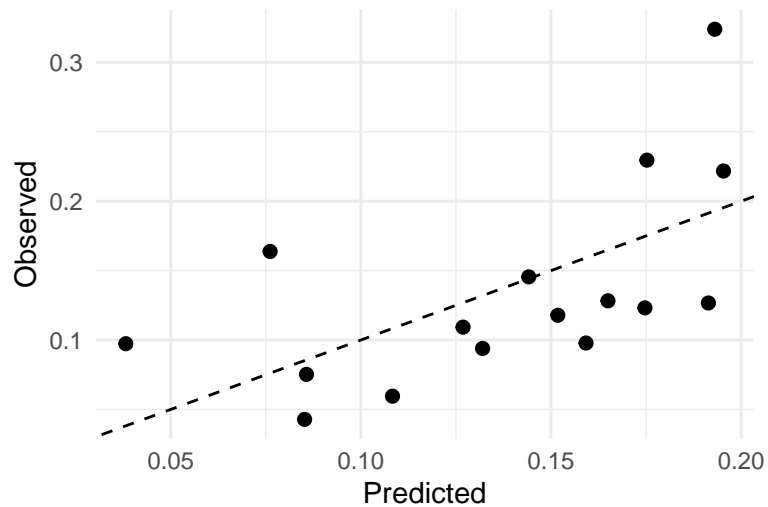
Full – CV_t



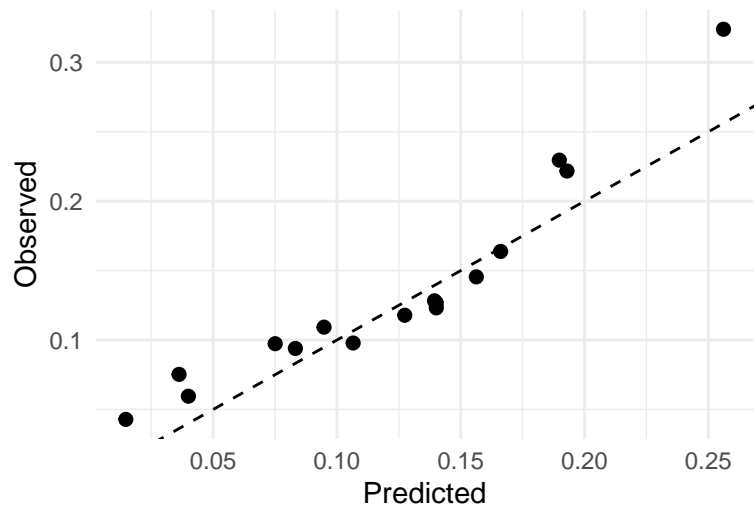
Range – CV_t



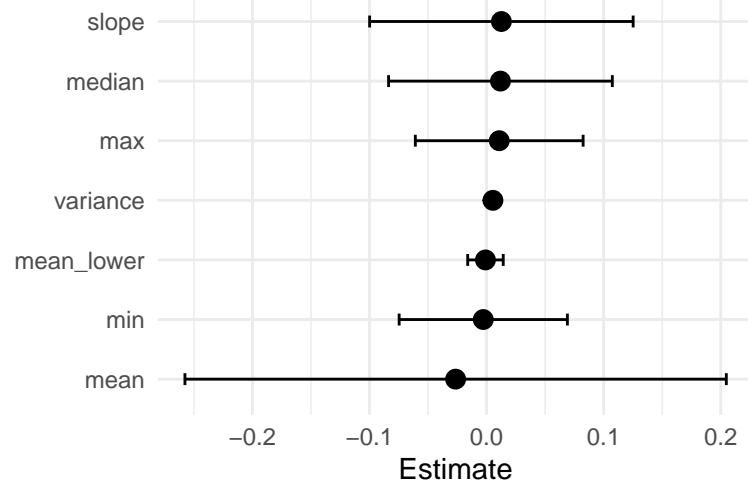
Mean_Upper – CV_t



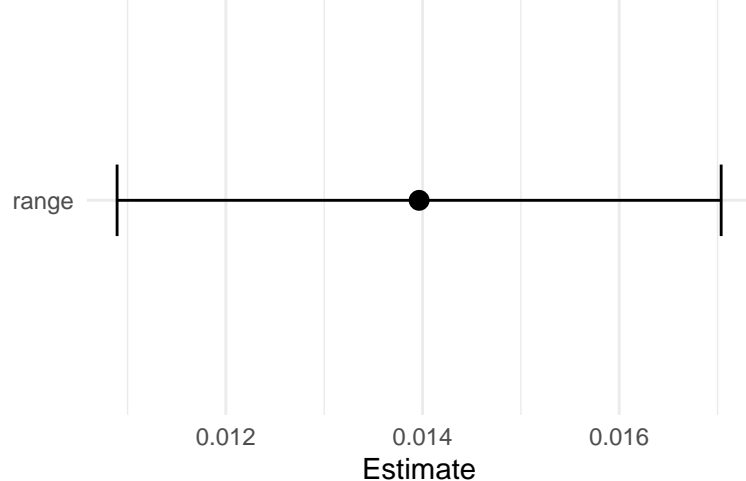
Ridge – CV_t



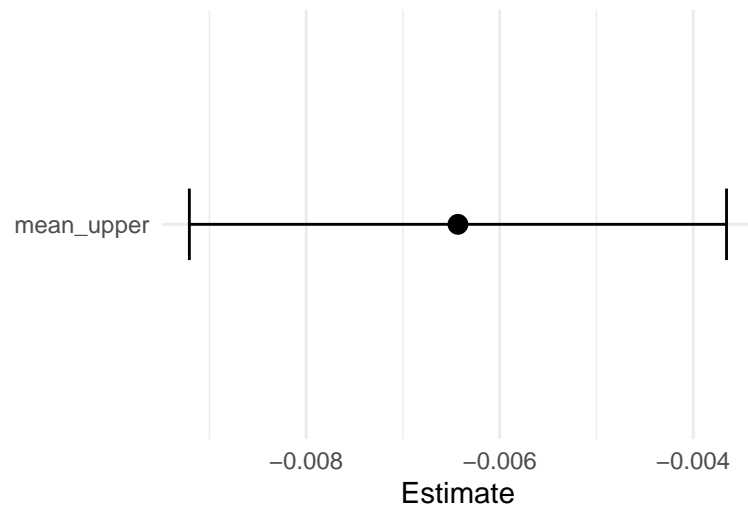
Full Coefficients – CV_t



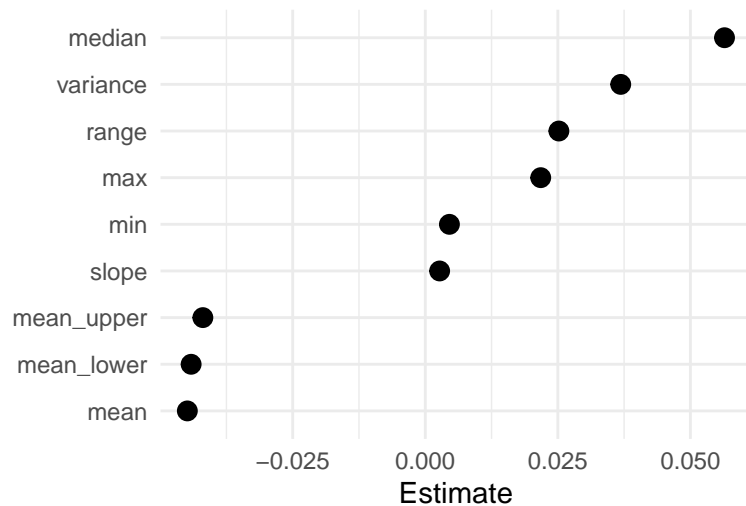
Range Coefficients – CV_t

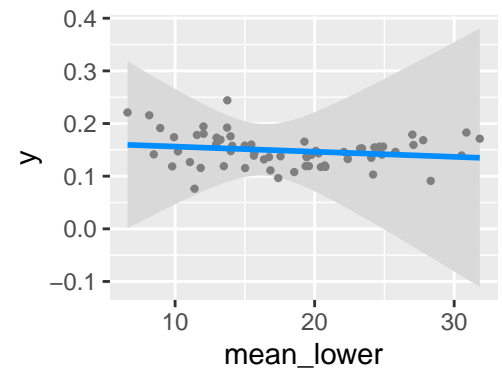
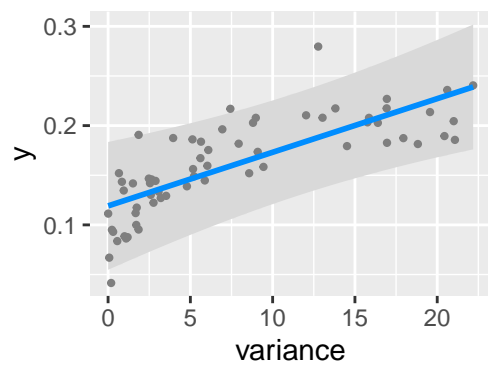
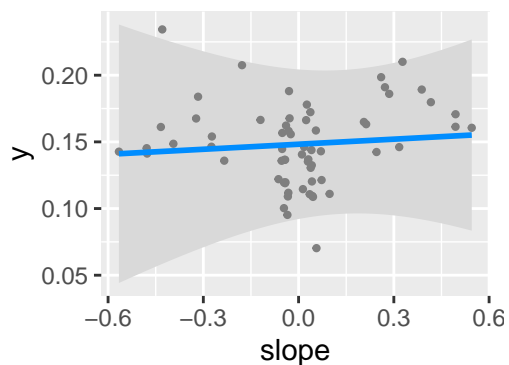
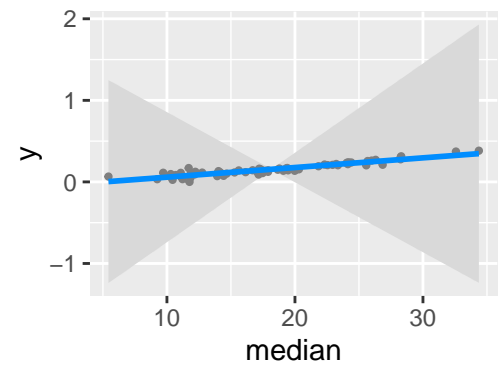
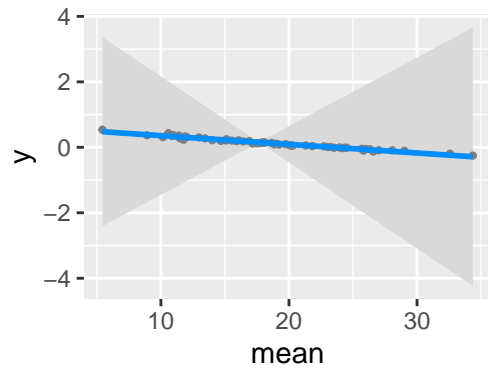
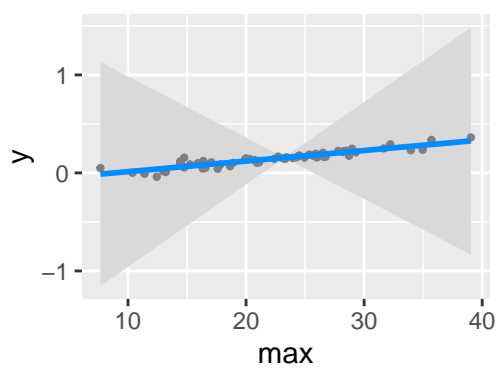
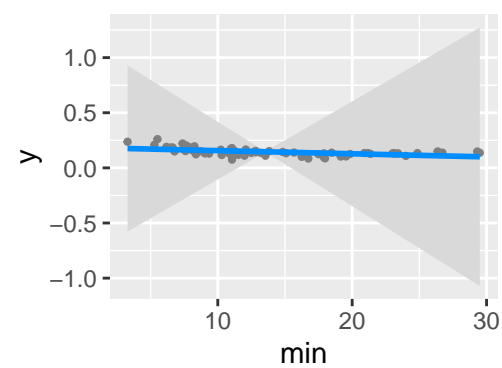


Mean_Upper Coefficients – CV_t

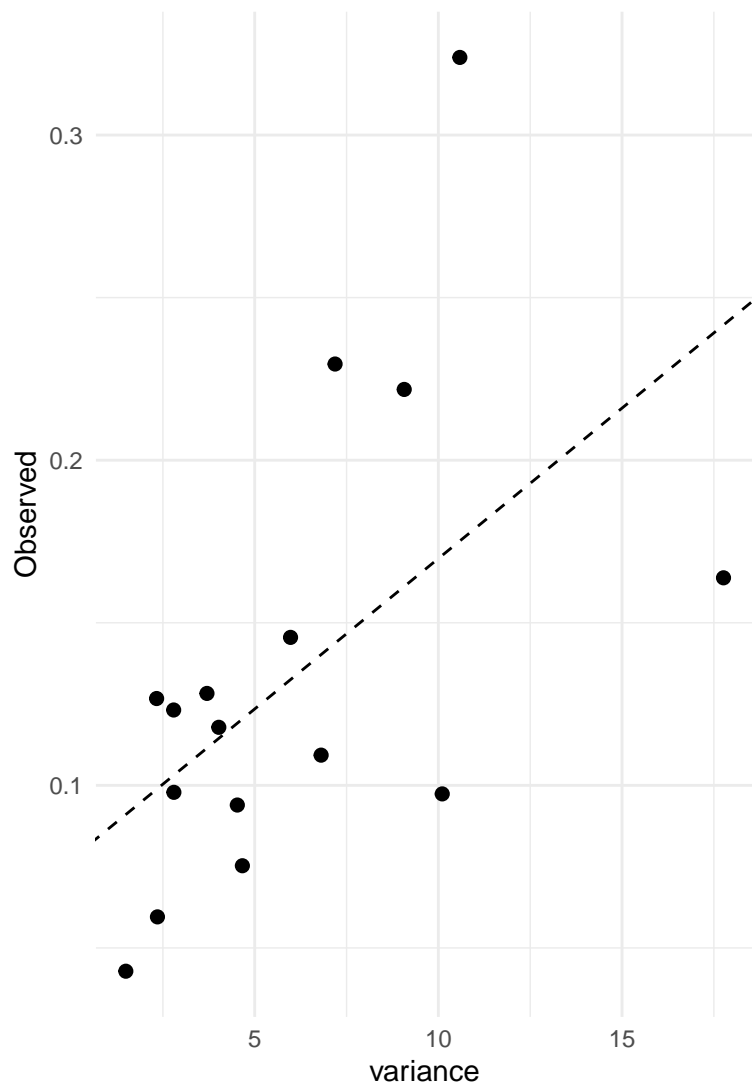


Ridge Coefficients – CV_t

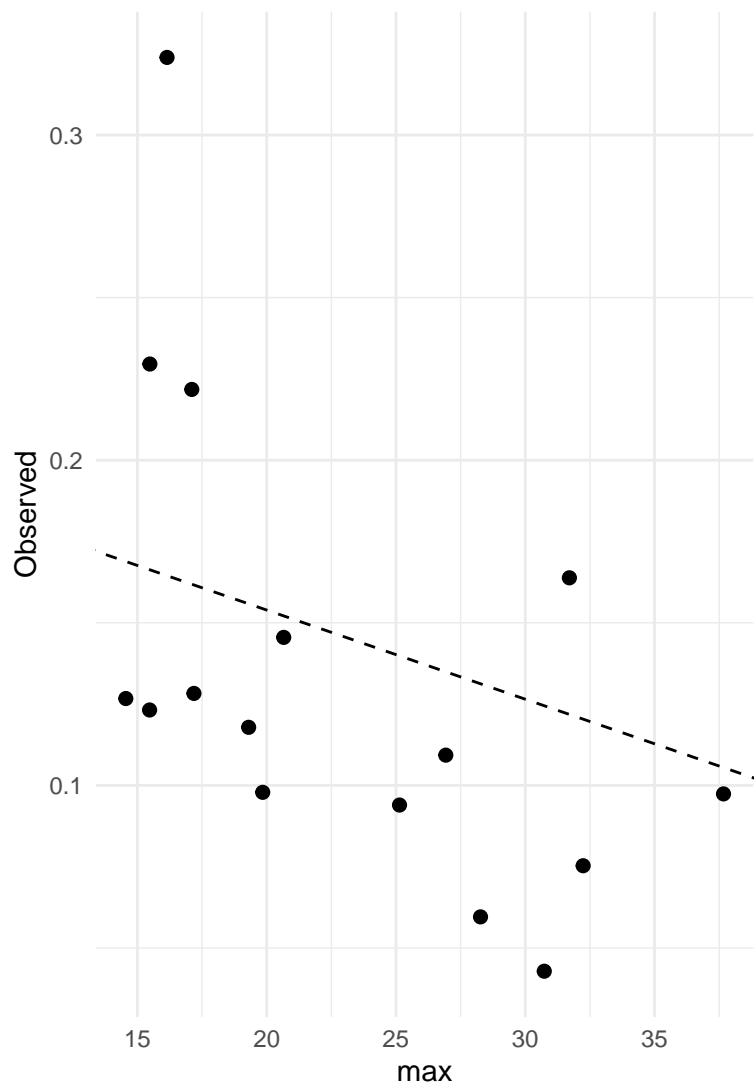




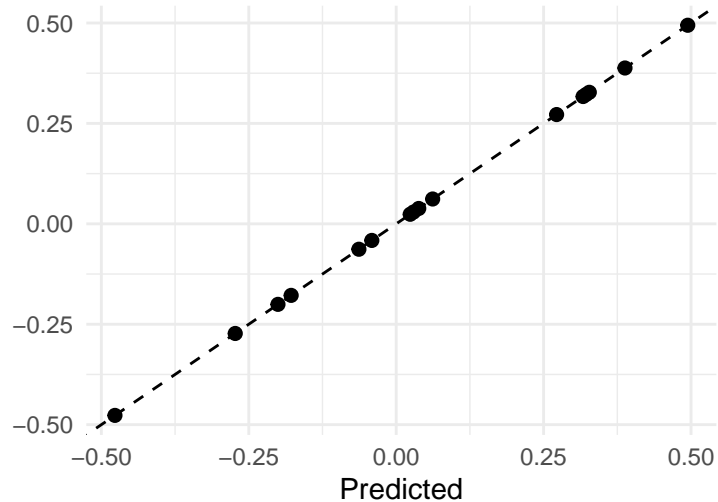
Simple Model: $y \sim \text{variance}$



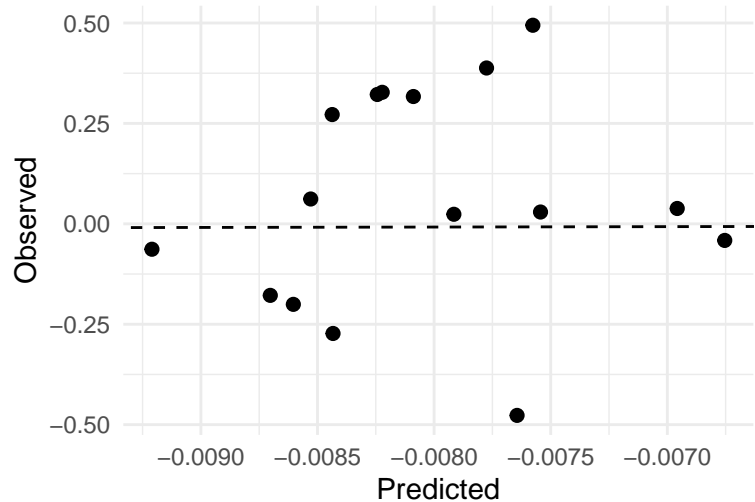
Simple Model: $y \sim \text{max}$



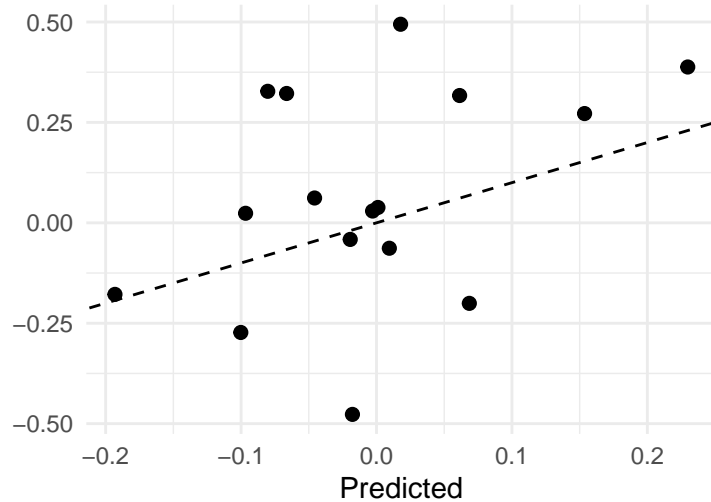
Full – RN



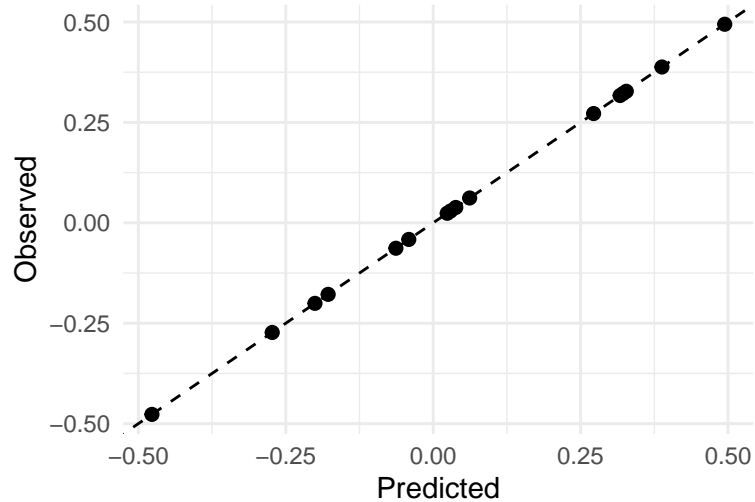
Range – RN



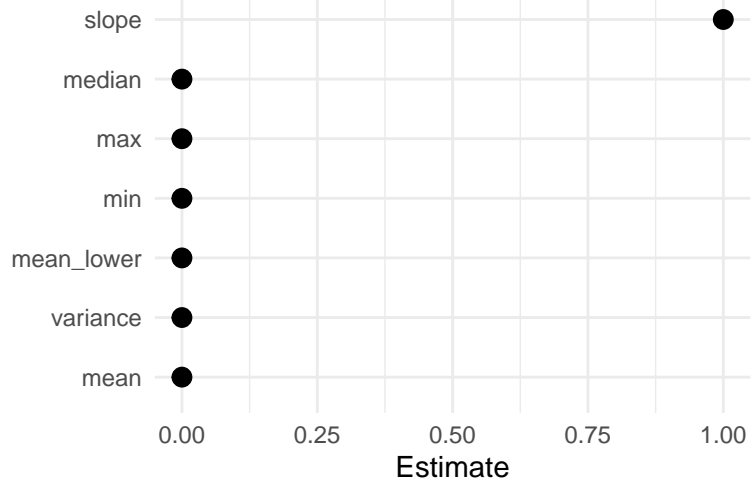
Mean_Upper – RN



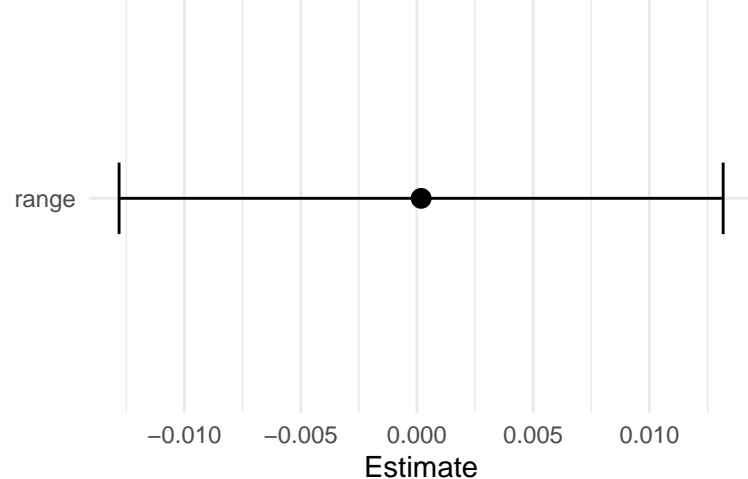
Ridge – RN



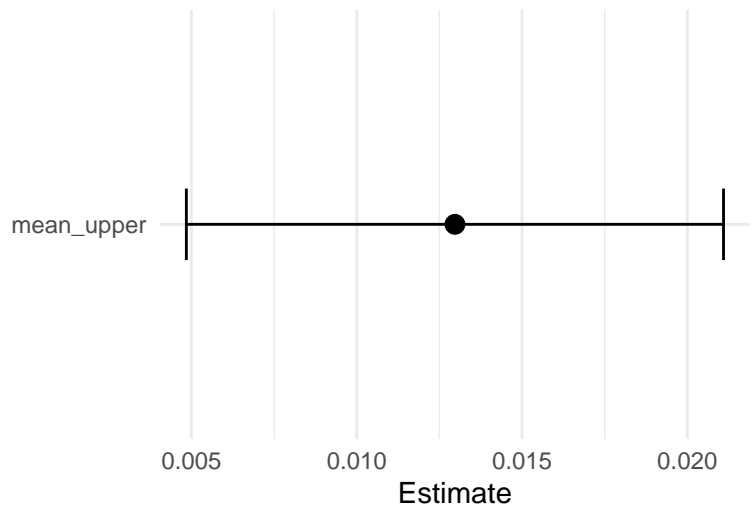
Full Coefficients – RN



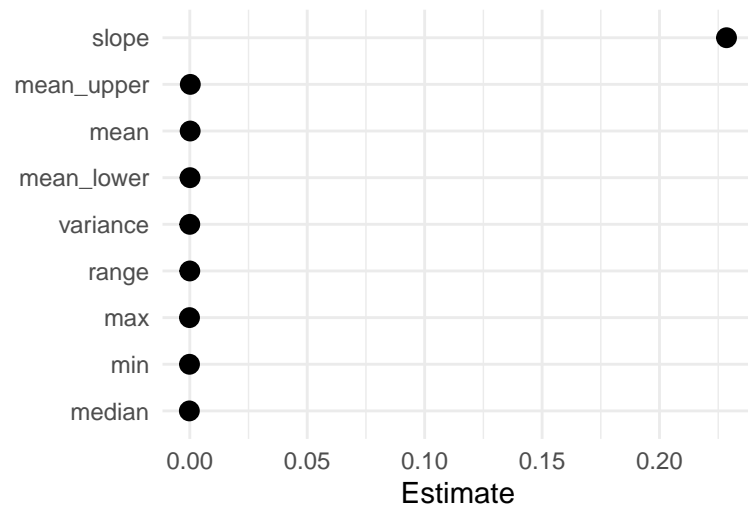
Range Coefficients – RN

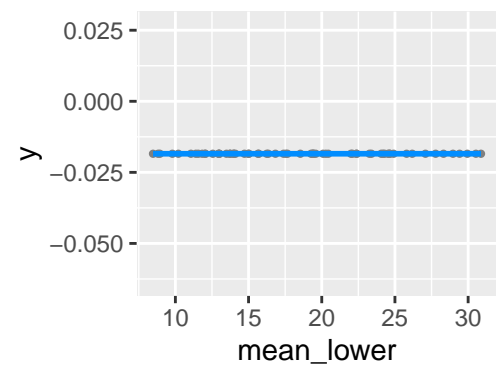
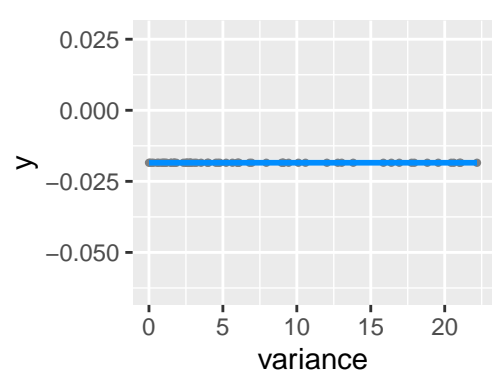
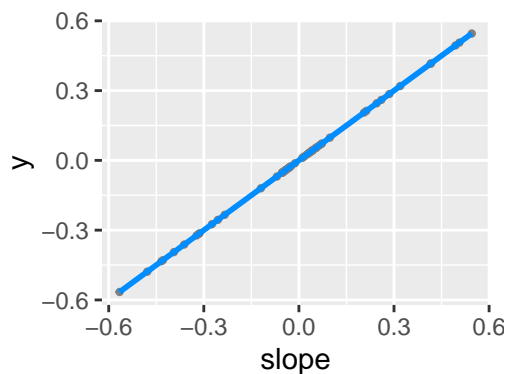
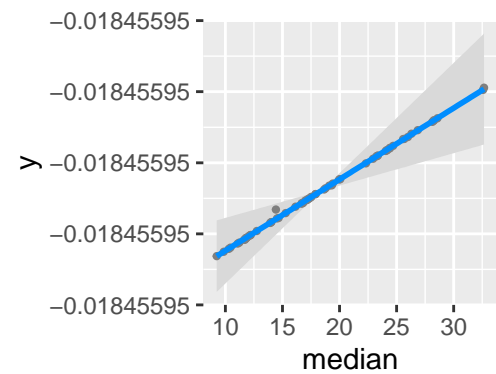
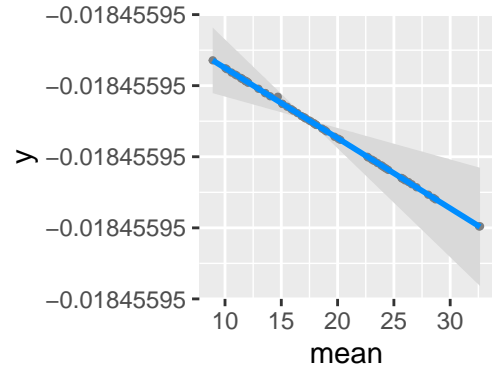
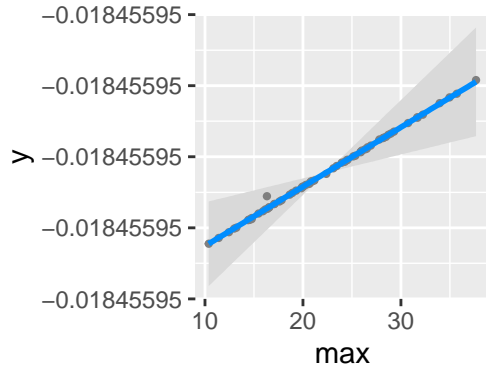
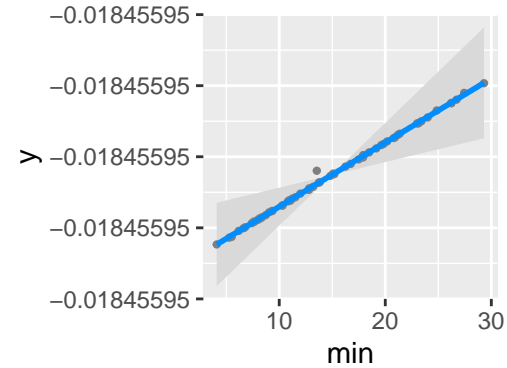


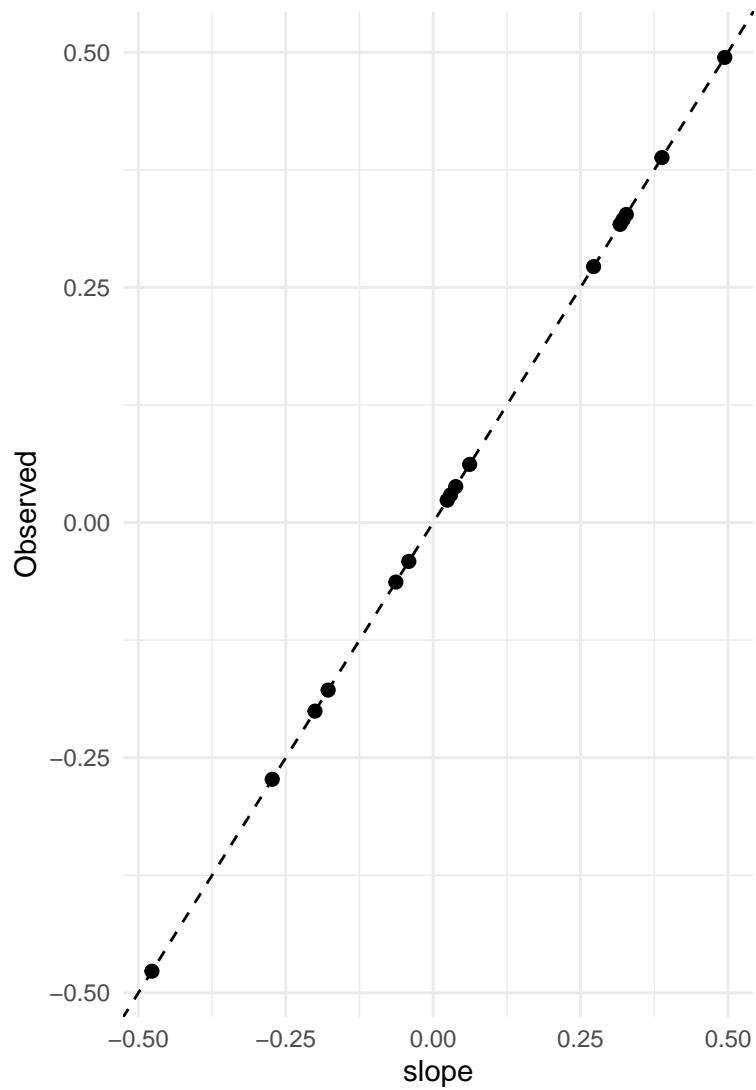
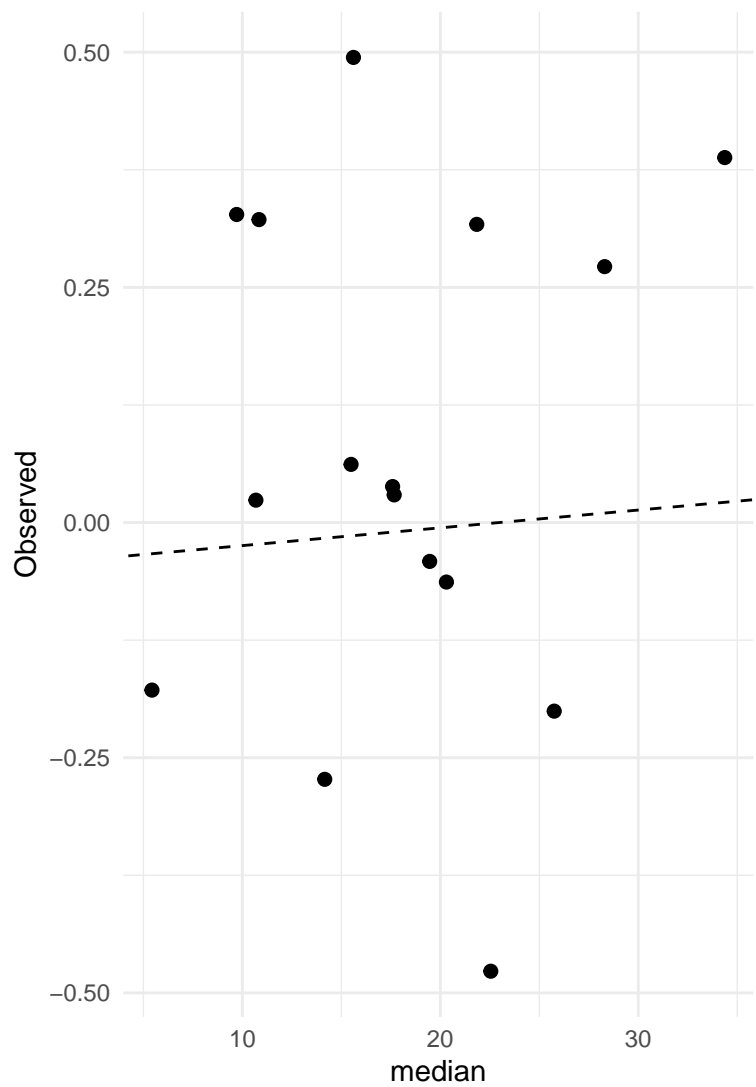
Mean_Upper Coefficients – RN



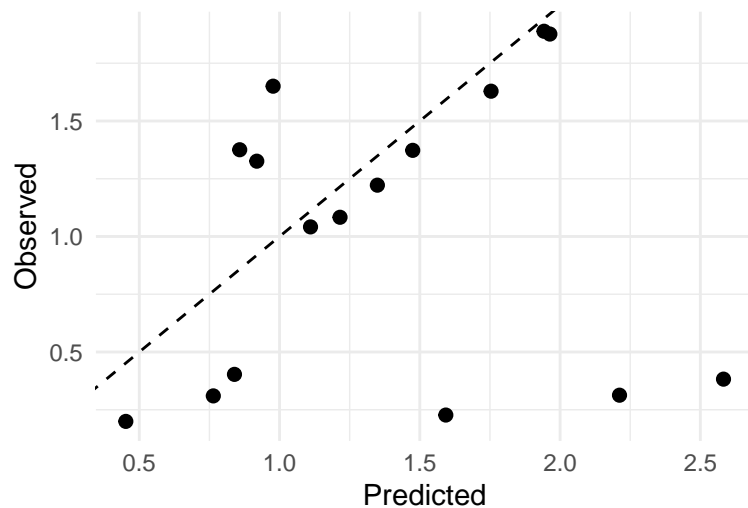
Ridge Coefficients – RN



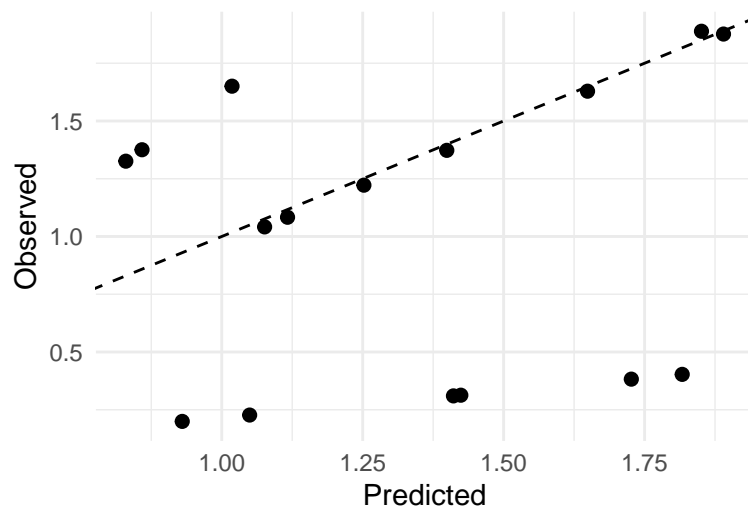


Simple Model: $y \sim \text{slope}$ Simple Model: $y \sim \text{median}$ 

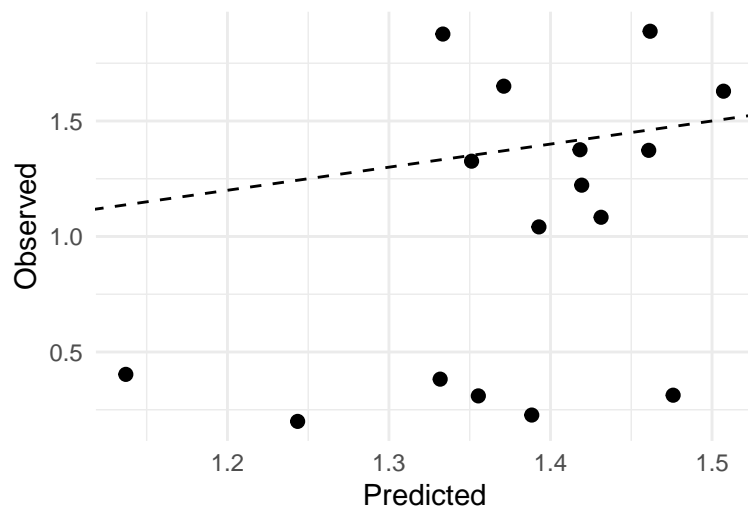
Full – RNN



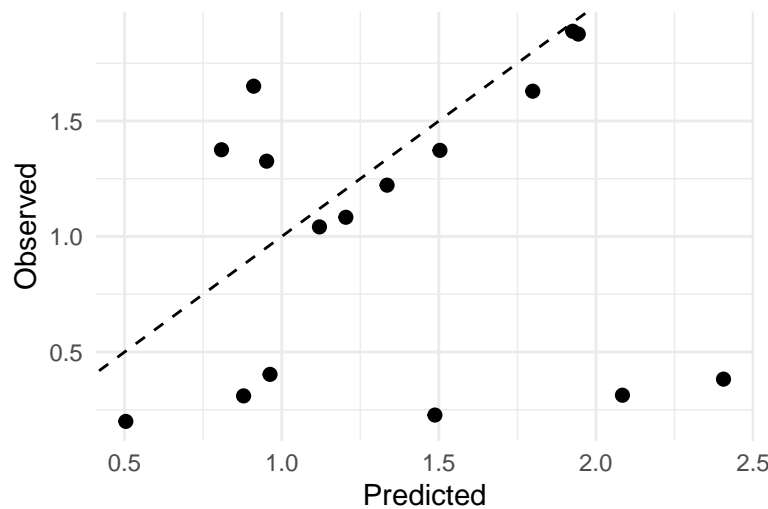
Range – RNN



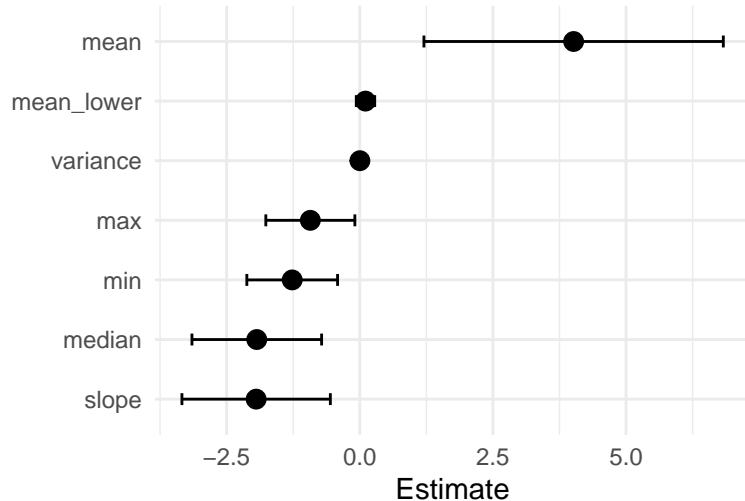
Mean_Upper – RNN



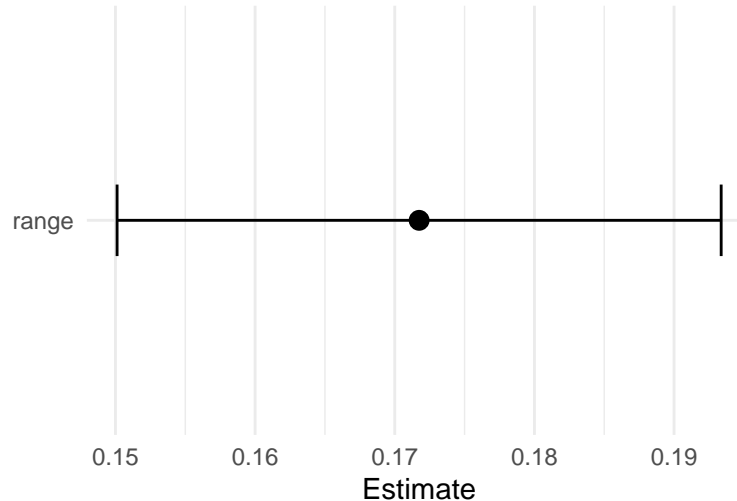
Ridge – RNN



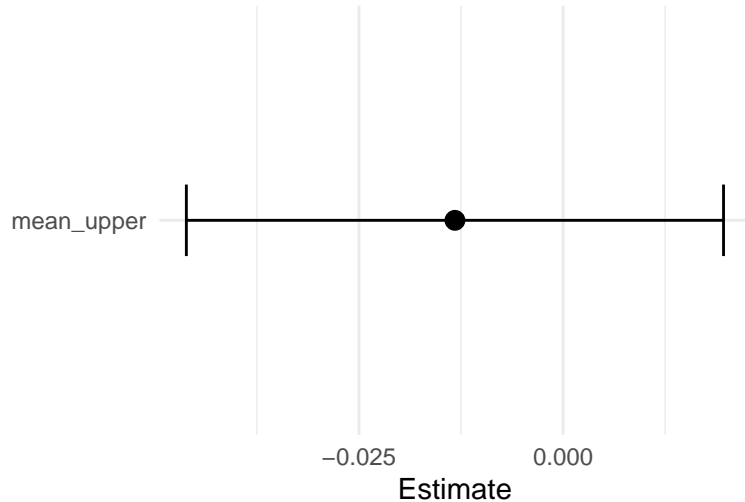
Full Coefficients – RNN



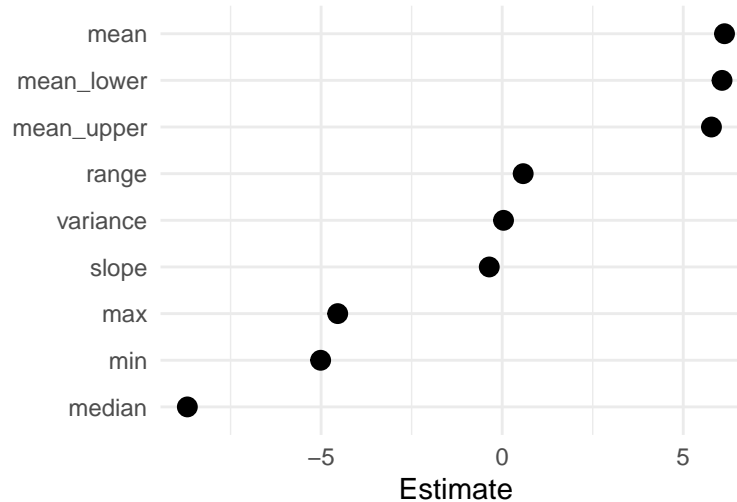
Range Coefficients – RNN

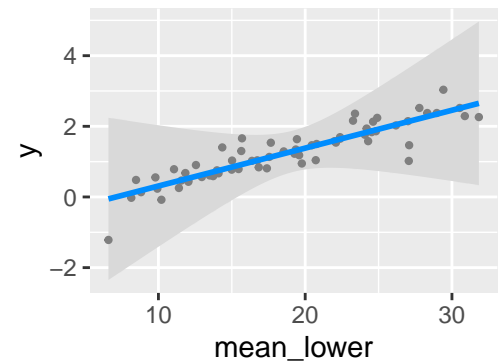
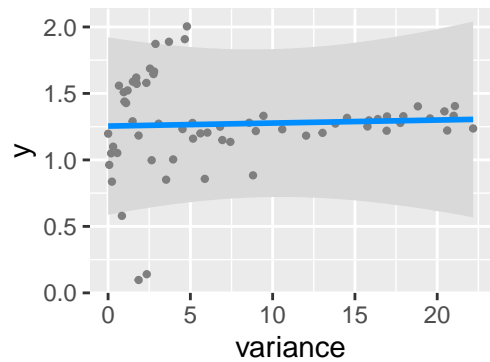
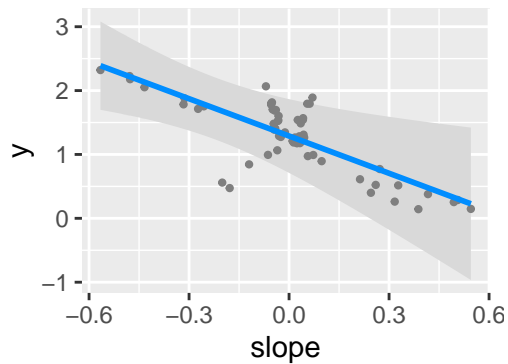
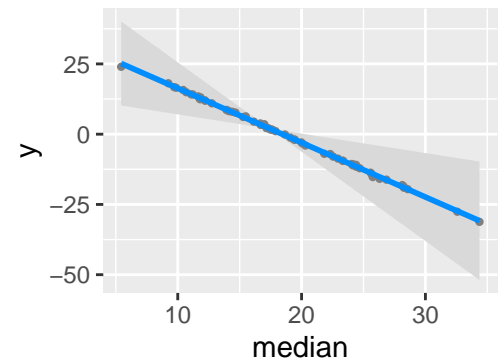
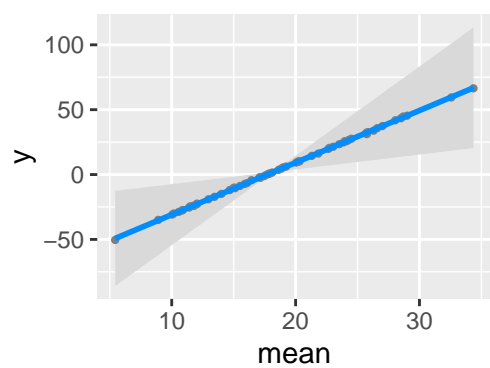
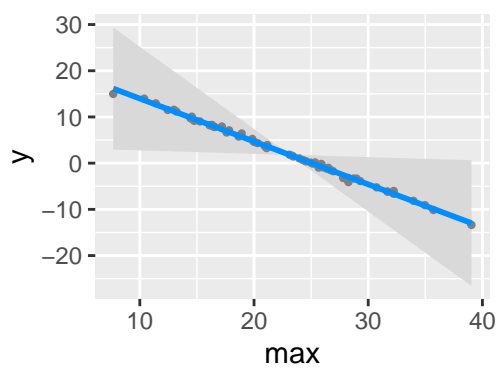
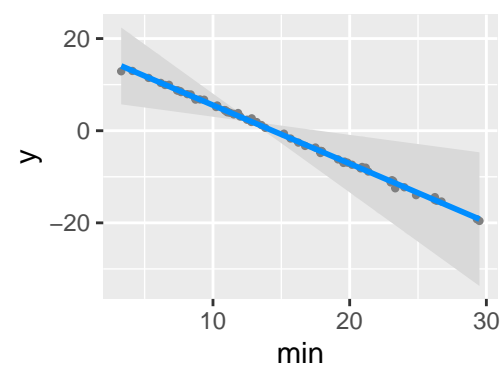


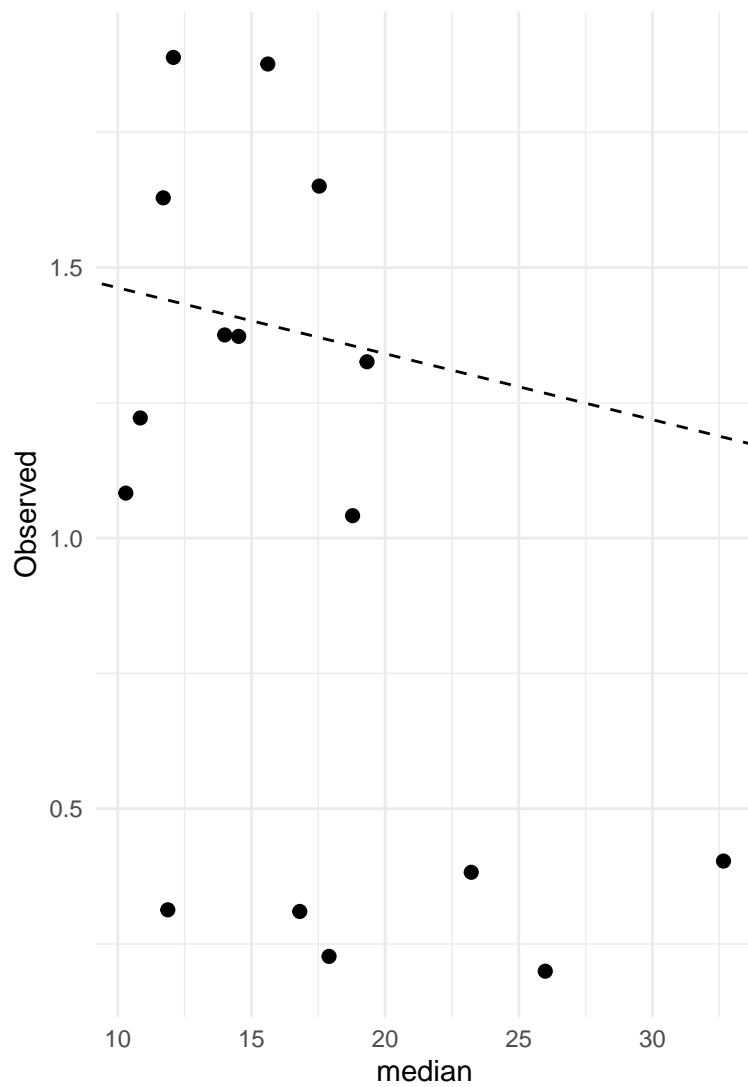
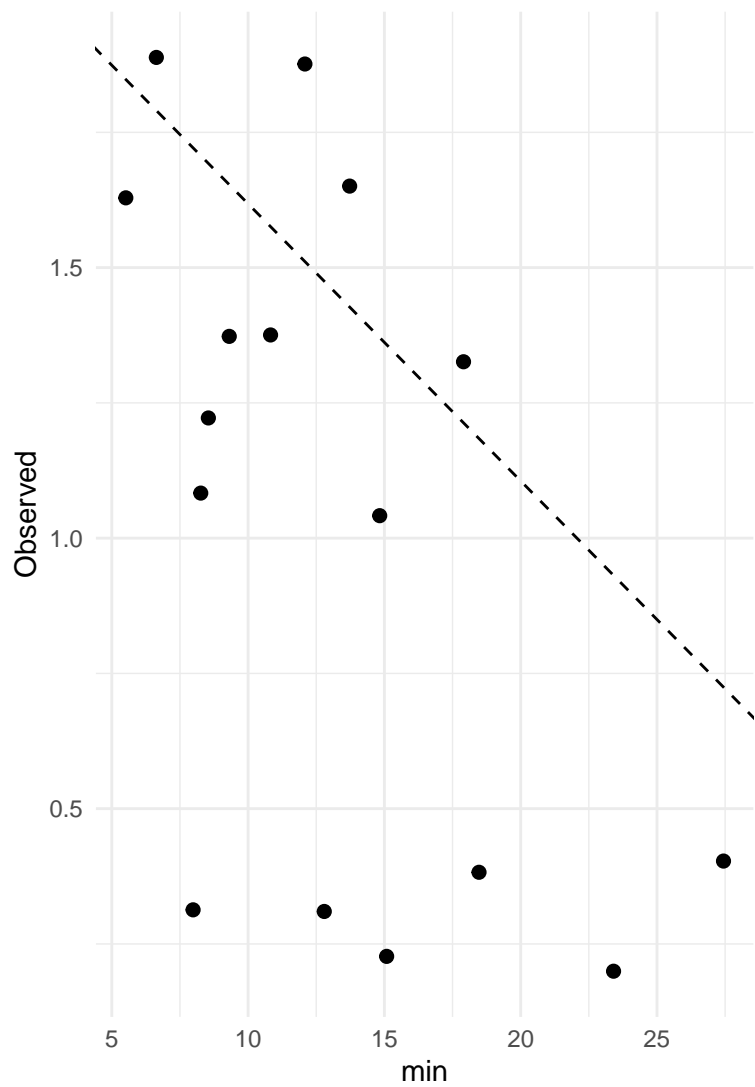
Mean_Upper Coefficients – RNN



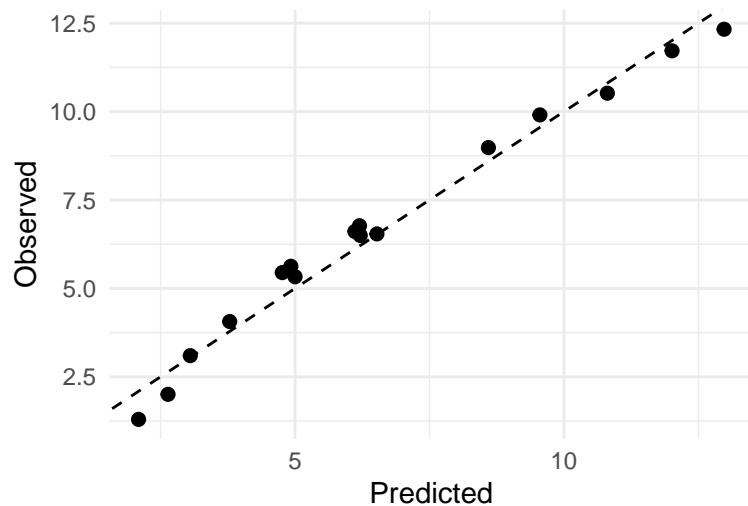
Ridge Coefficients – RNN



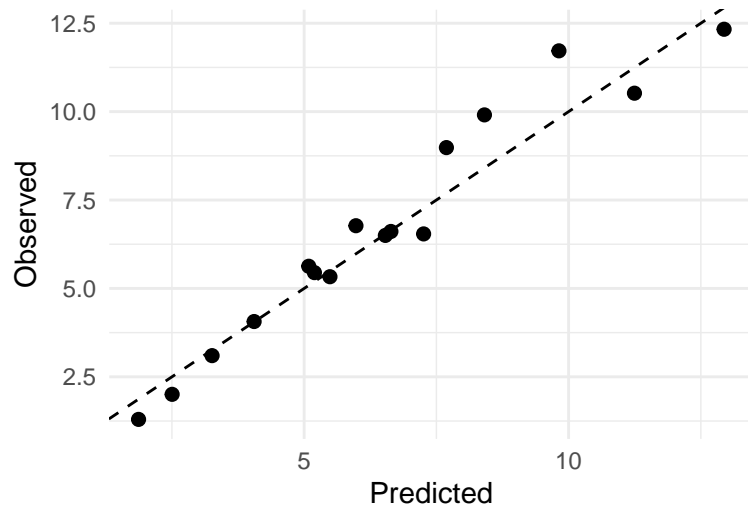


Simple Model: $y \sim \text{median}$ Simple Model: $y \sim \text{min}$ 

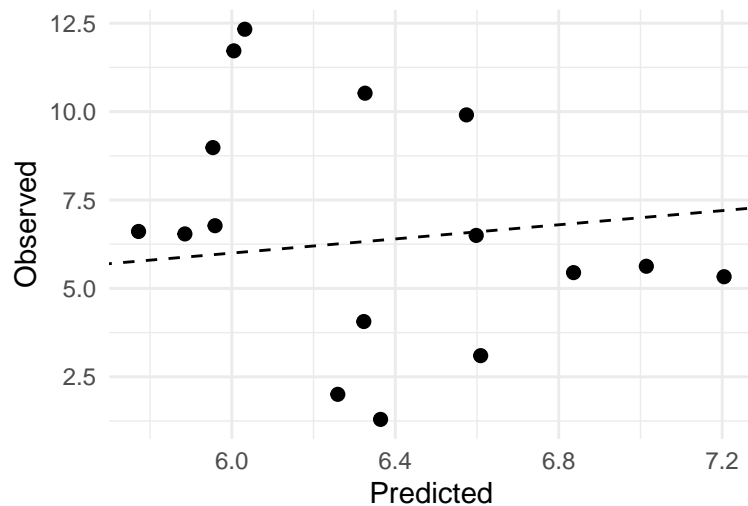
Full – D_slope



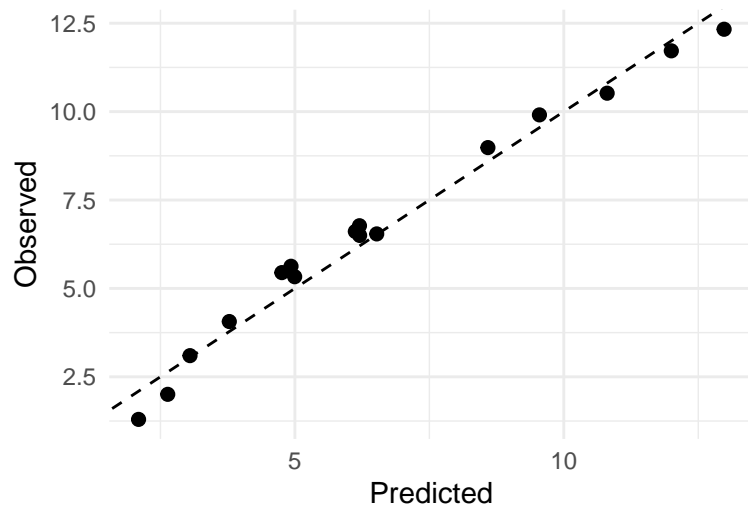
Range – D_slope



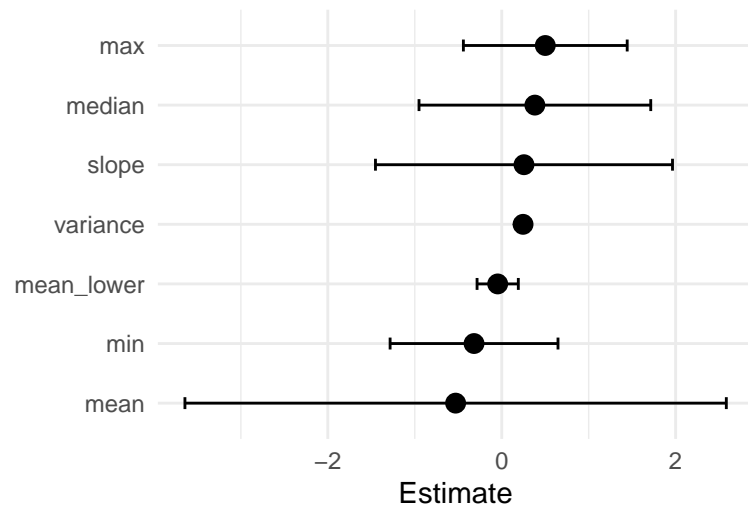
Mean_Upper – D_slope



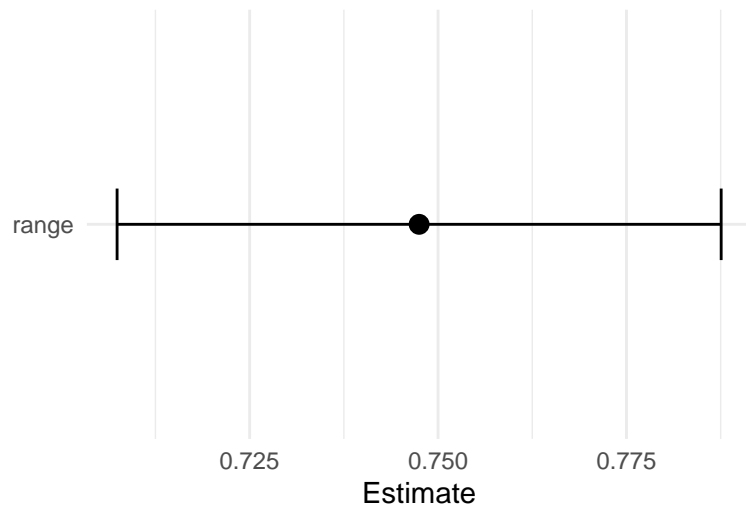
Ridge – D_slope



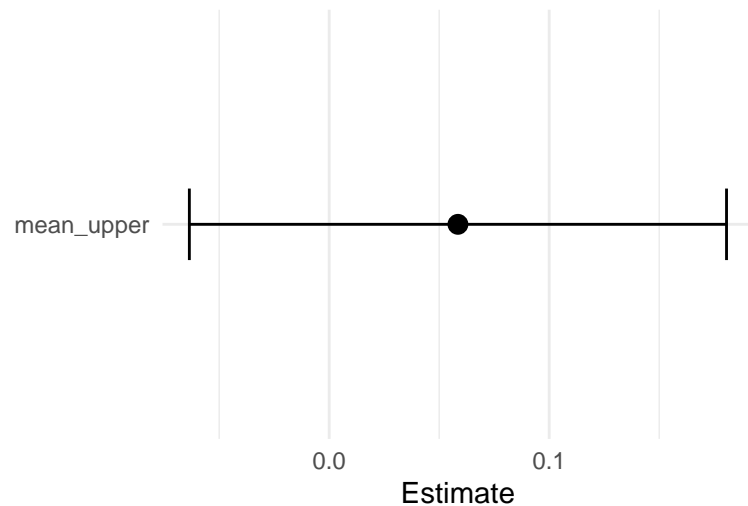
Full Coefficients – D_slope



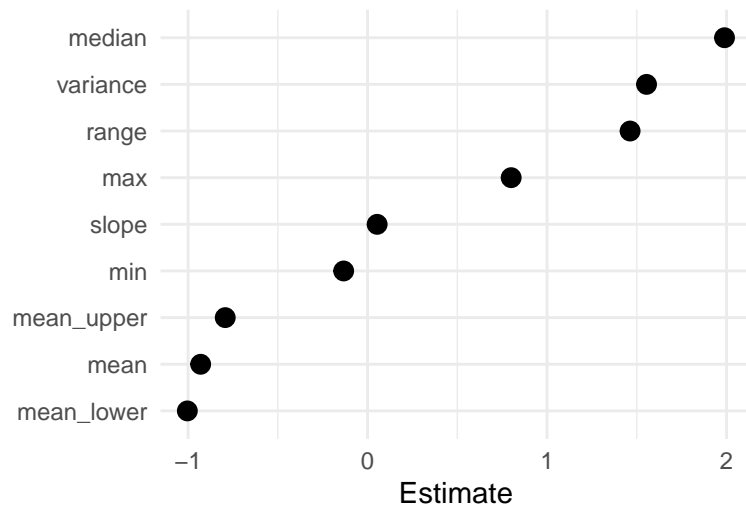
Range Coefficients – D_slope

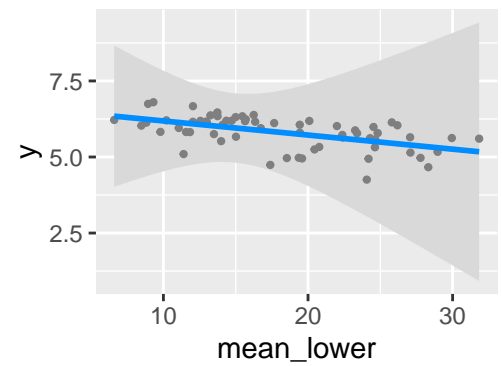
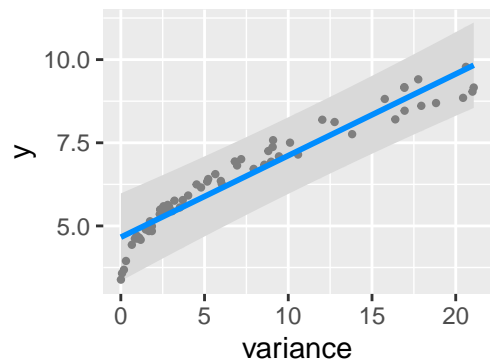
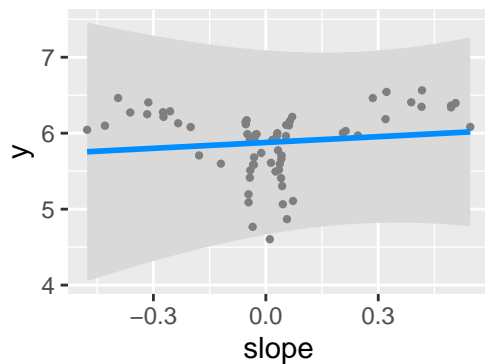
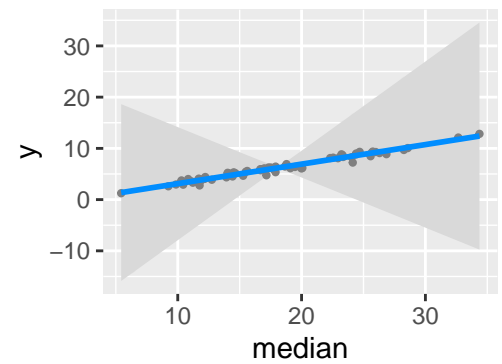
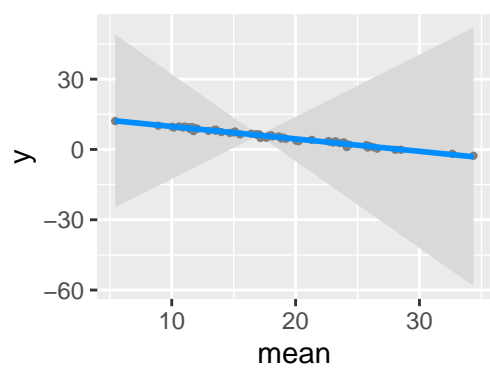
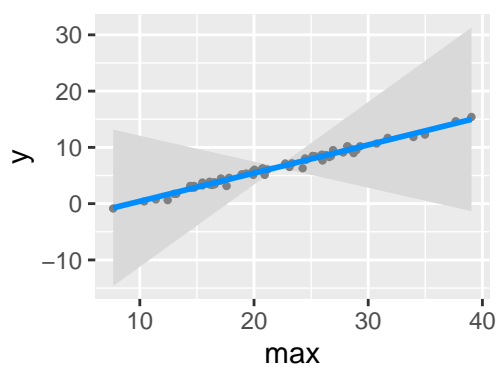
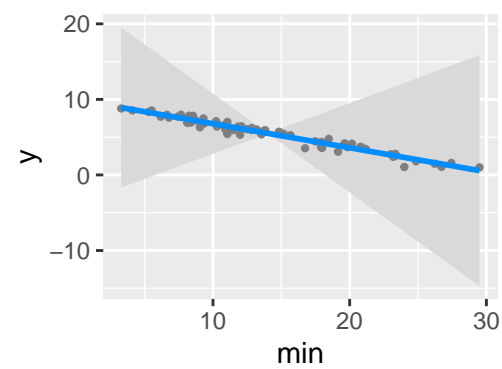


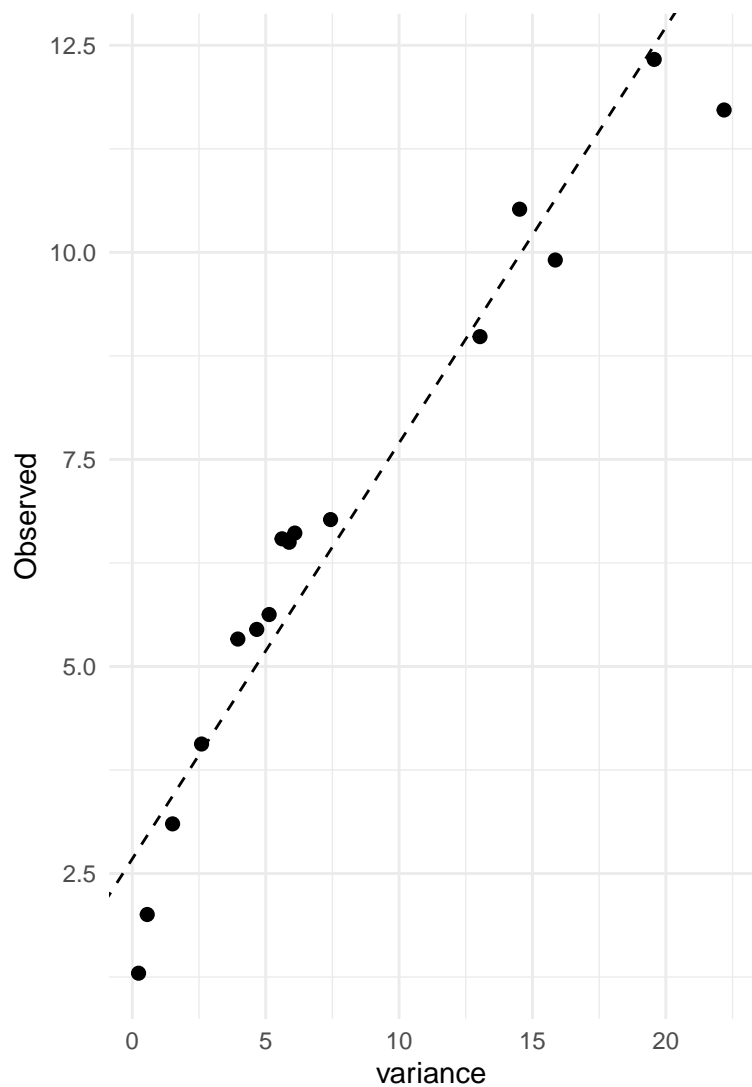
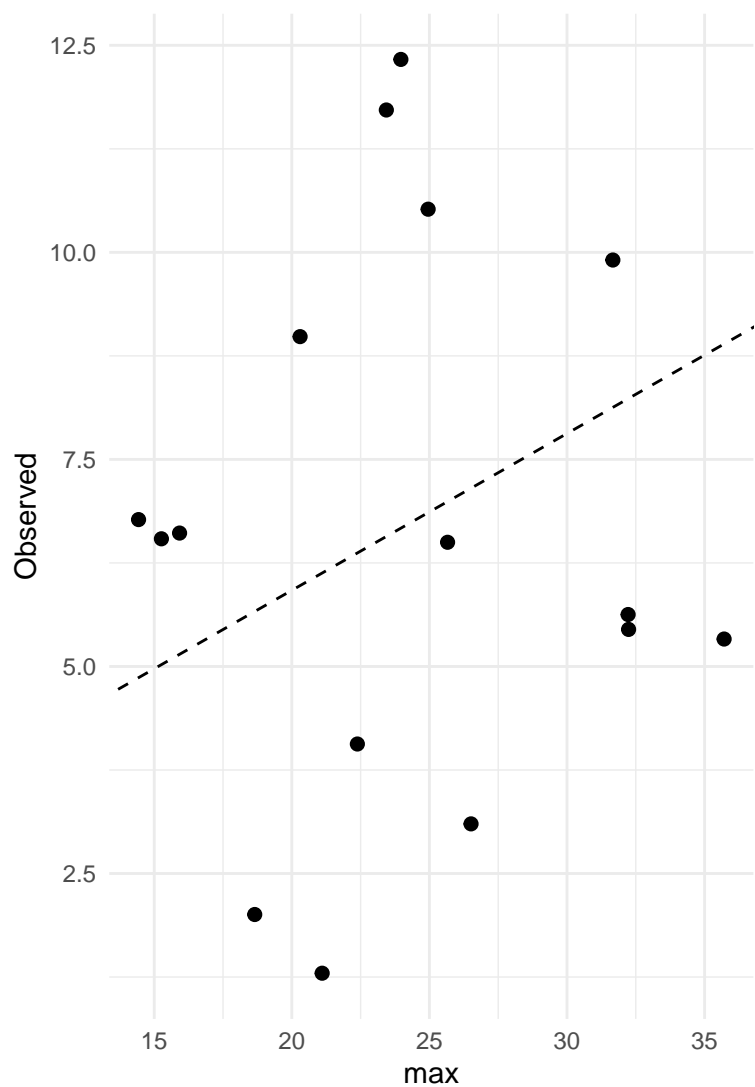
Mean_Upper Coefficients – D_slope



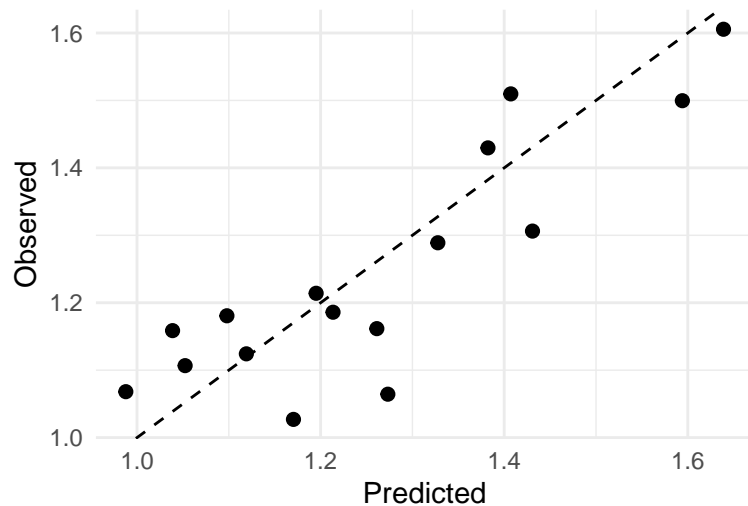
Ridge Coefficients – D_slope



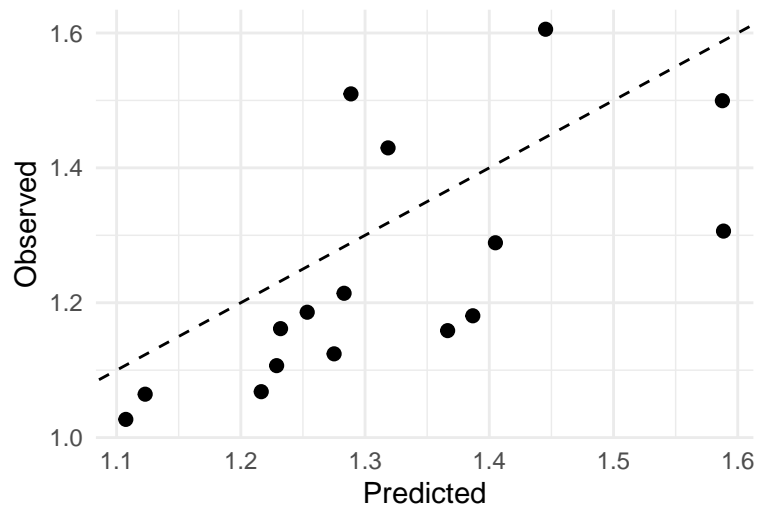


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{max}$ 

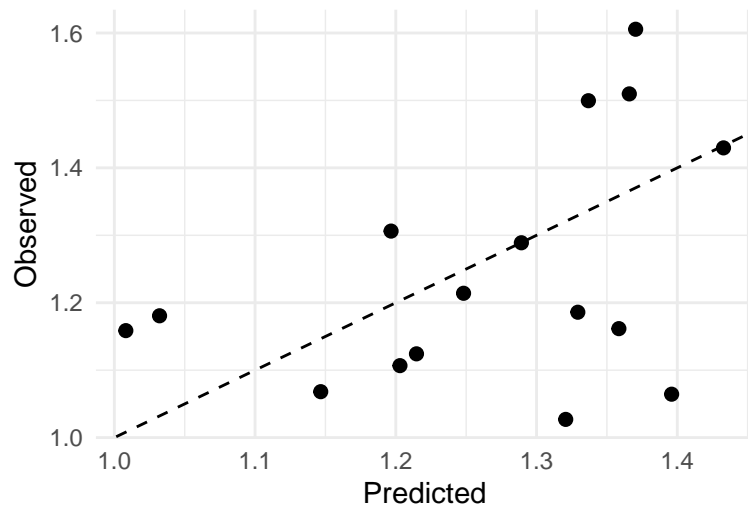
Full – RC



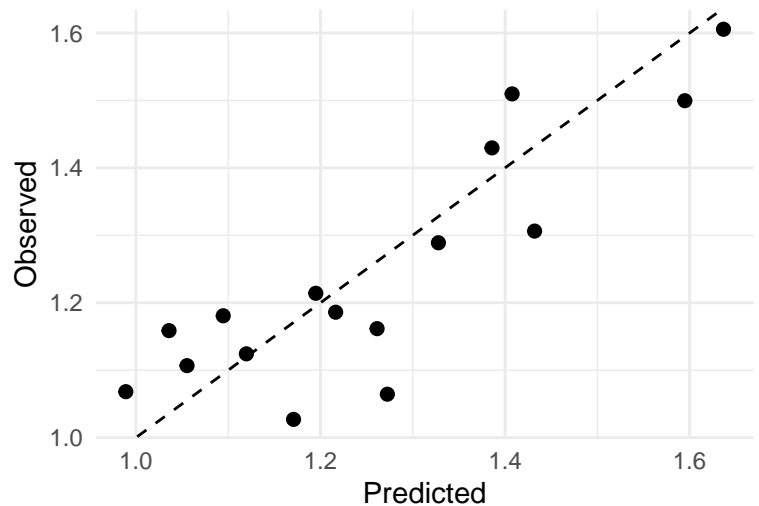
Range – RC



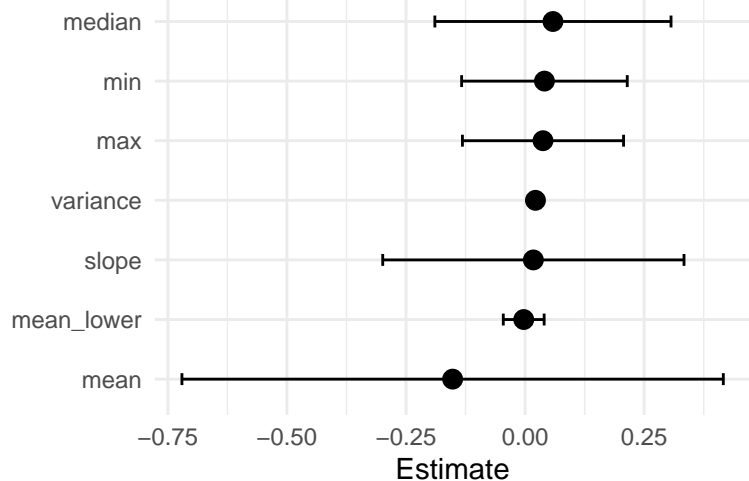
Mean_Upper – RC



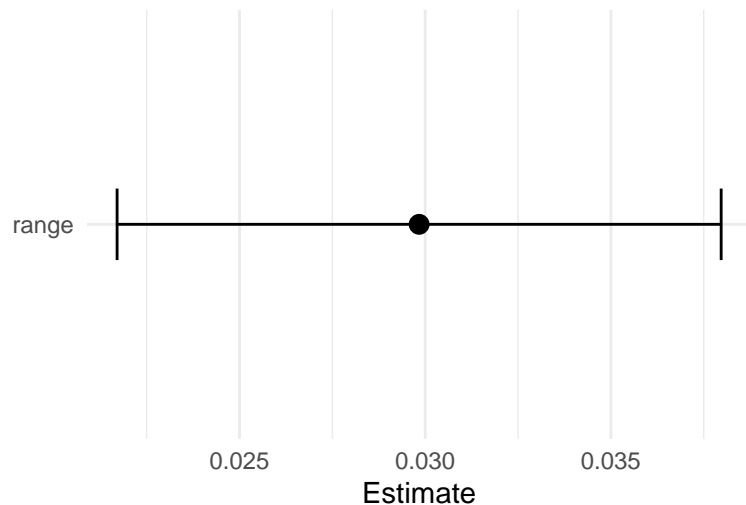
Ridge – RC



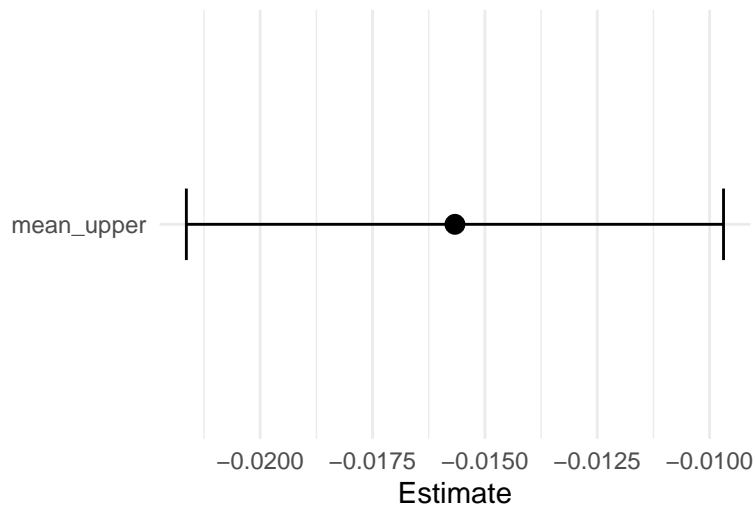
Full Coefficients – RC



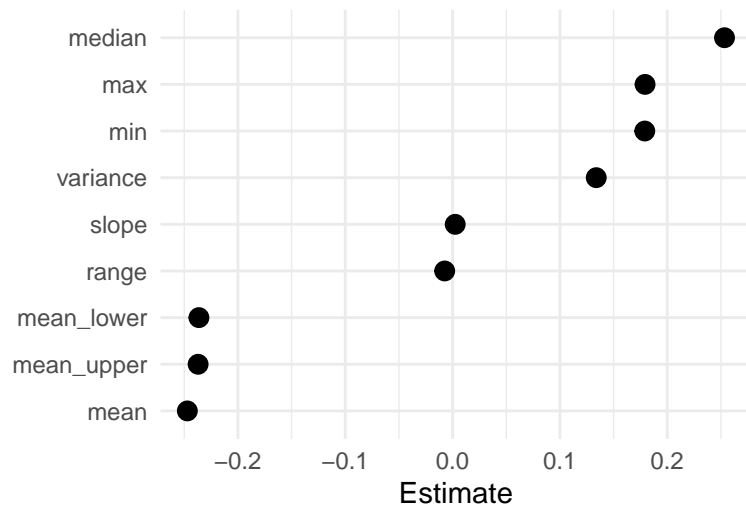
Range Coefficients – RC

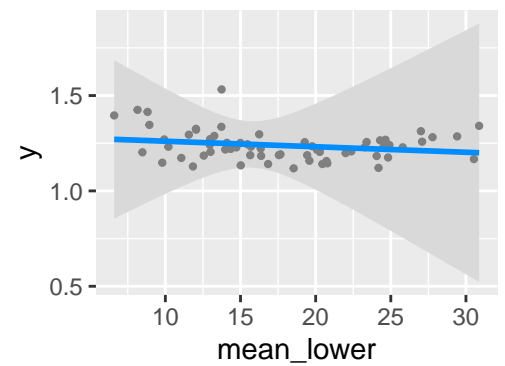
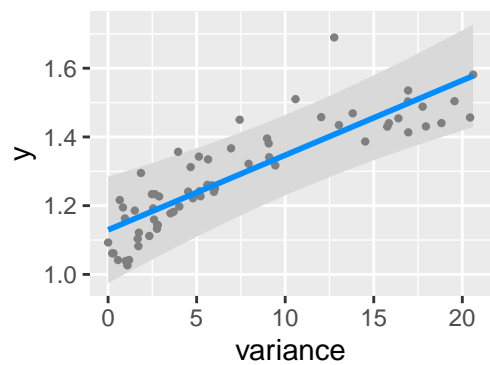
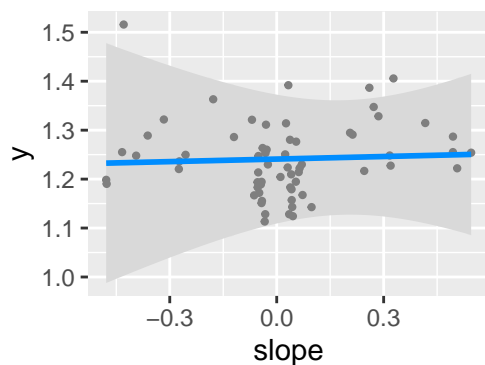
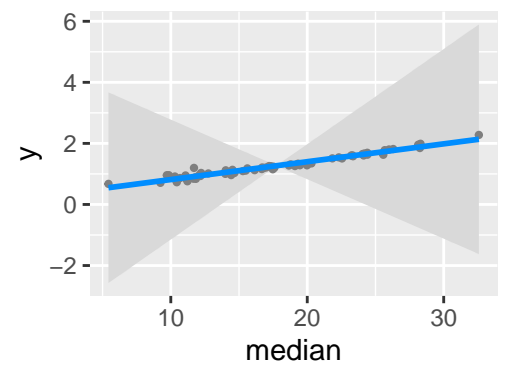
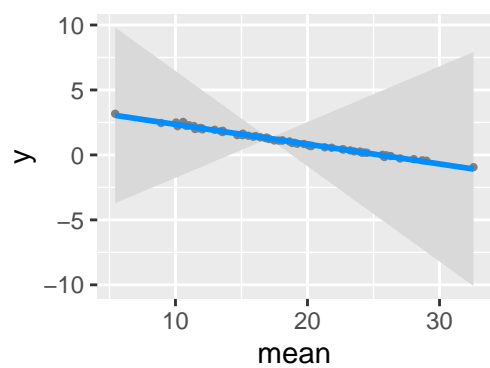
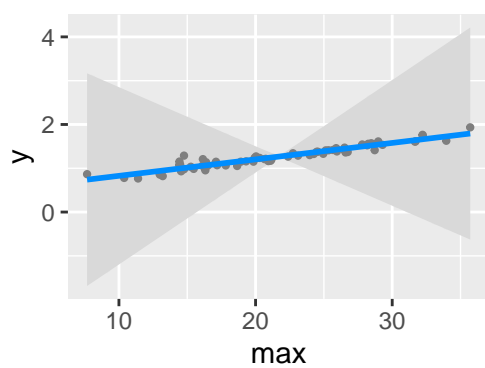
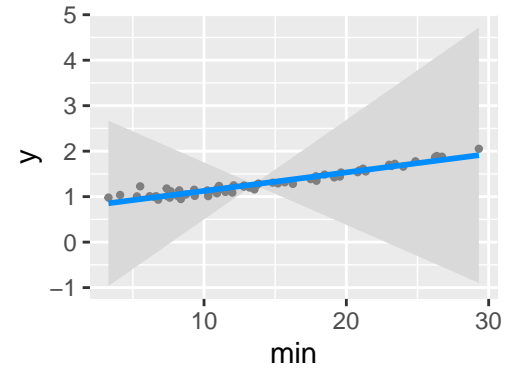


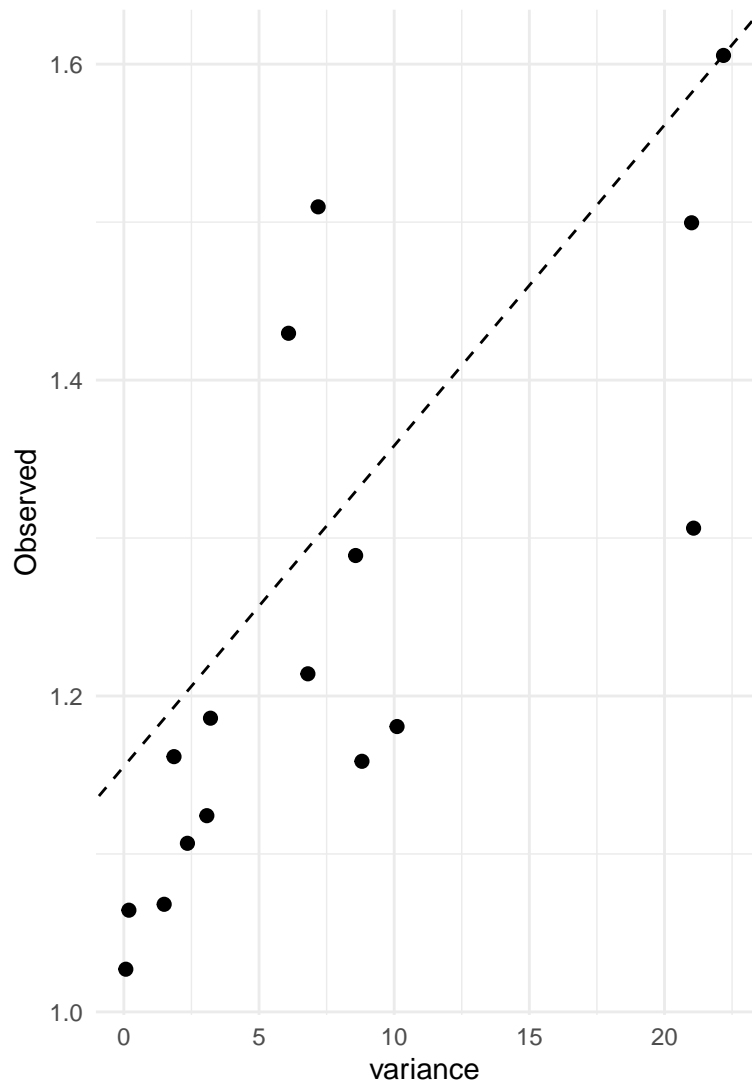
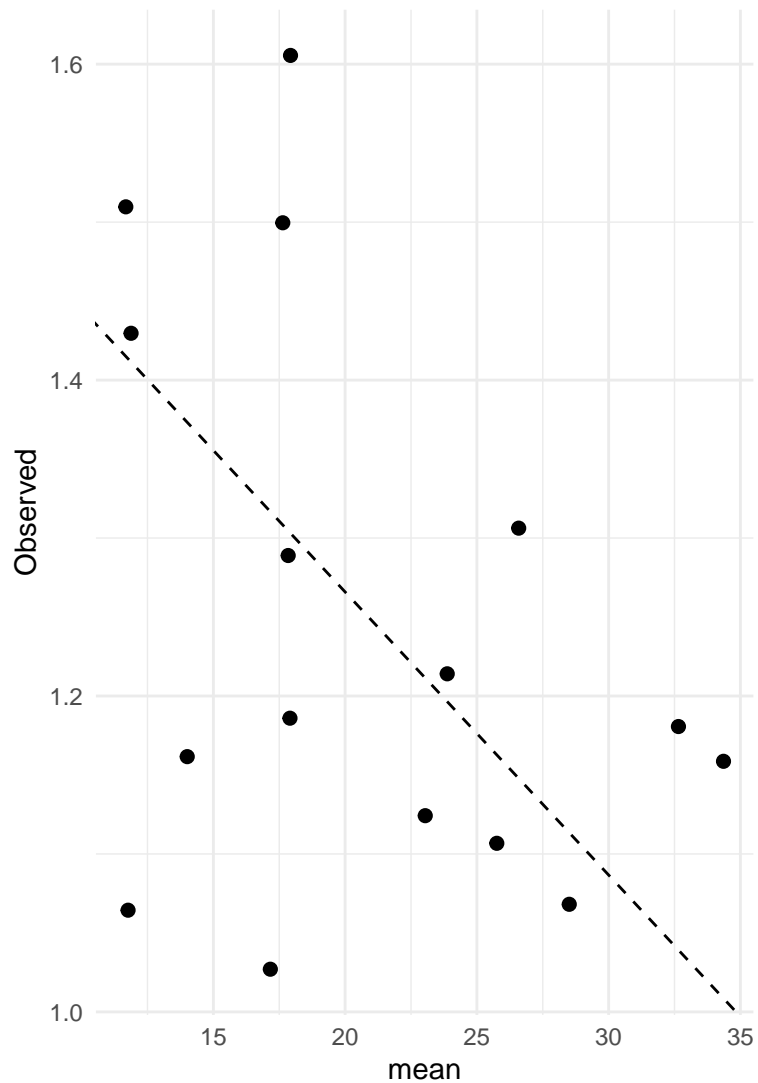
Mean_Upper Coefficients – RC



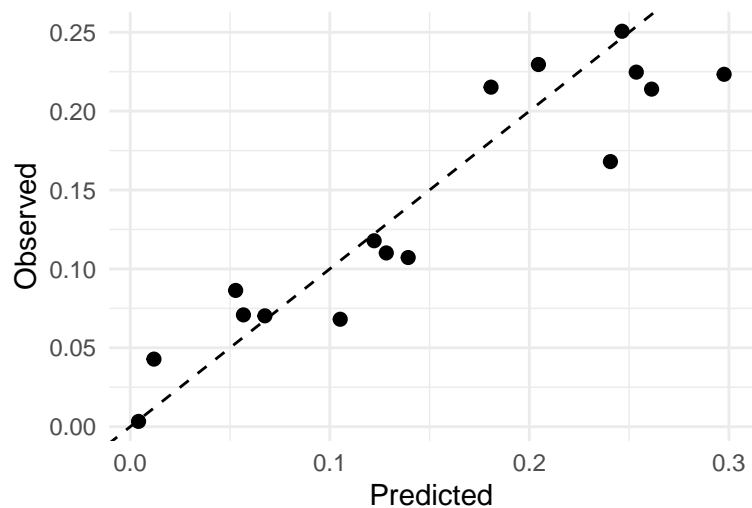
Ridge Coefficients – RC



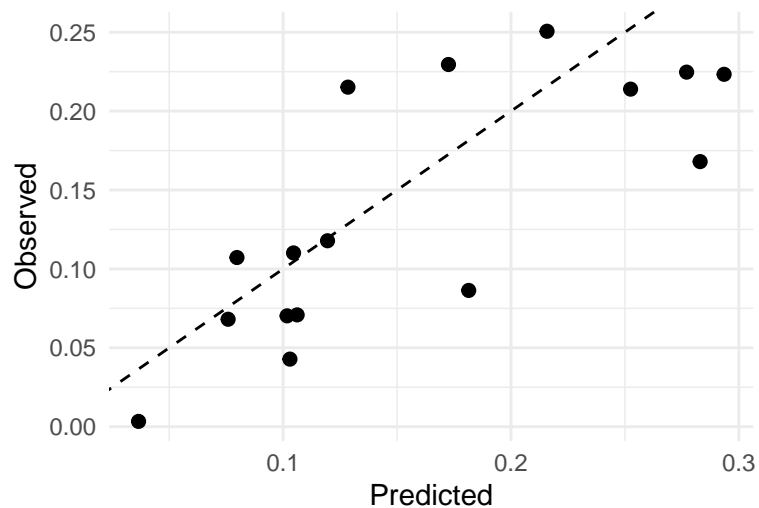


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{mean}$ 

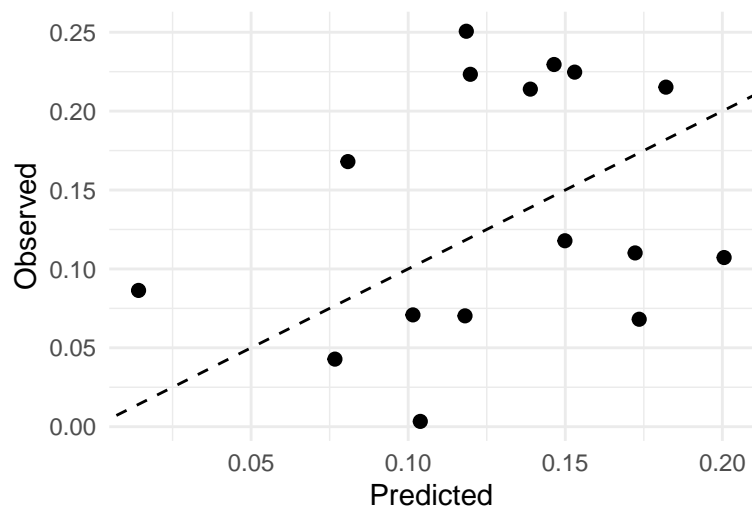
Full – gPi



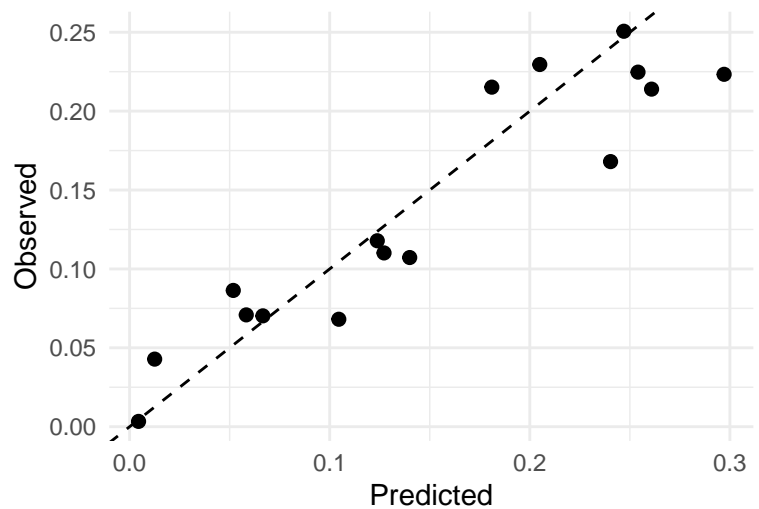
Range – gPi



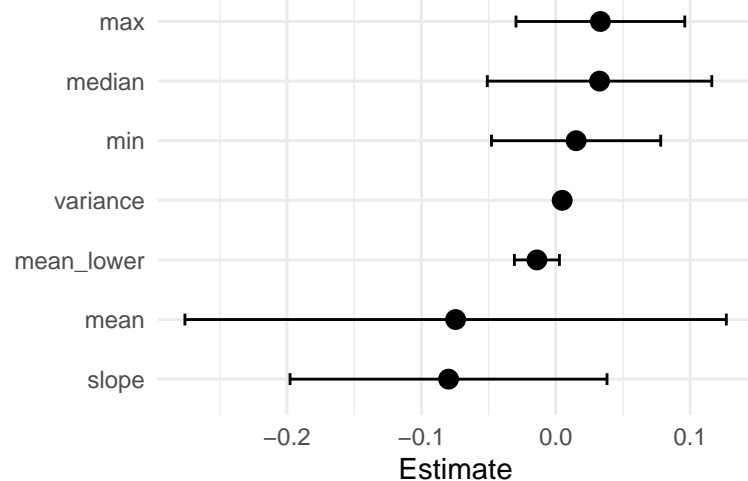
Mean_Upper – gPi



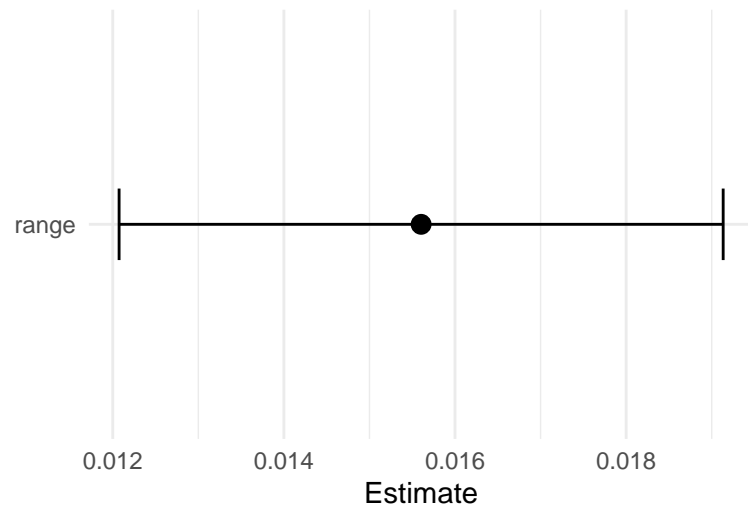
Ridge – gPi



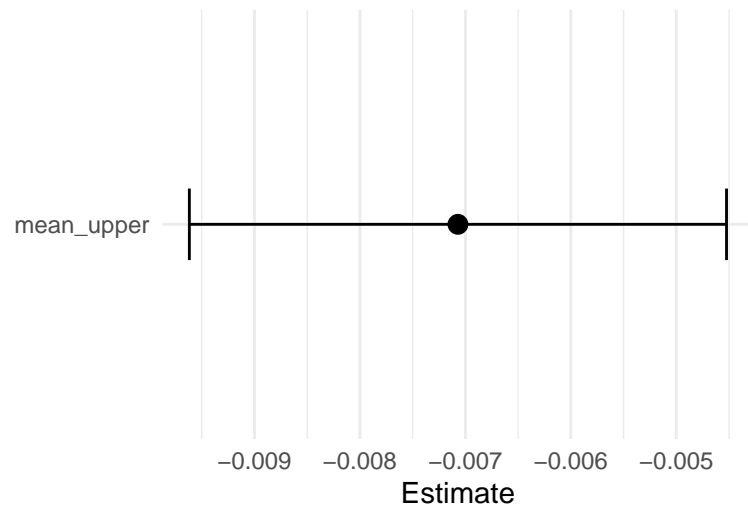
Full Coefficients – gPi



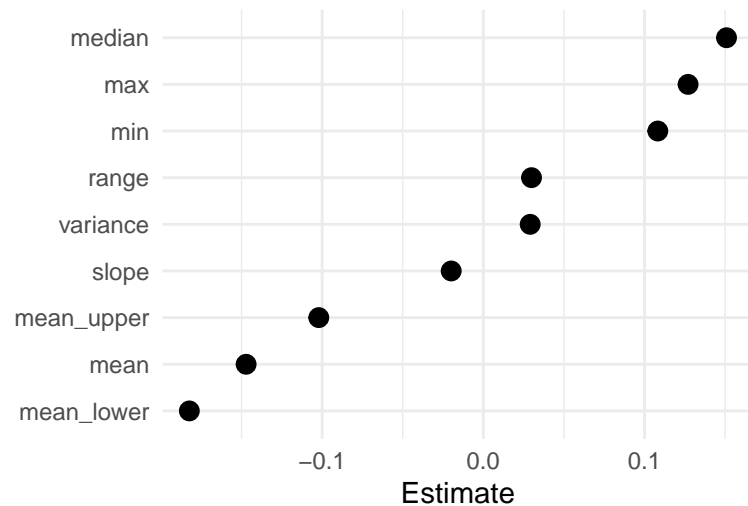
Range Coefficients – gPi

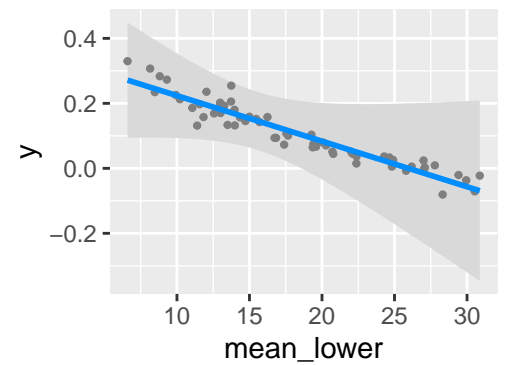
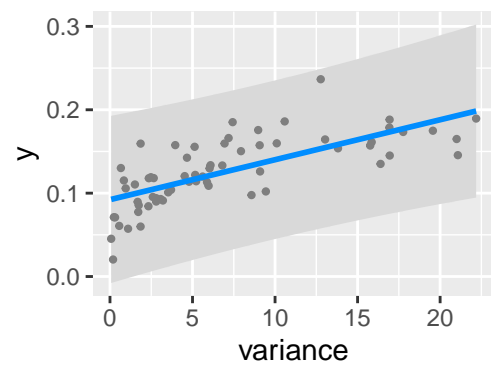
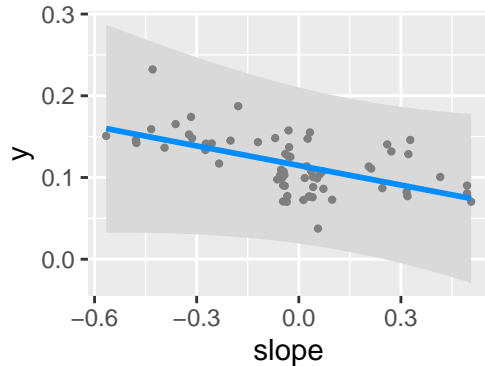
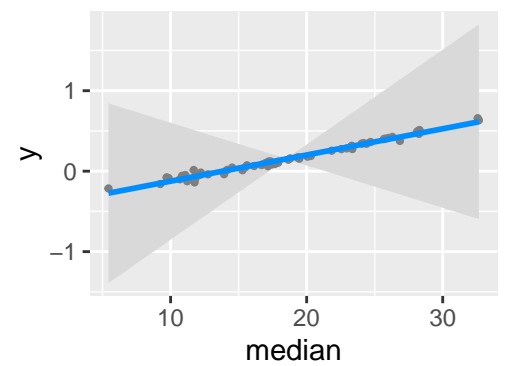
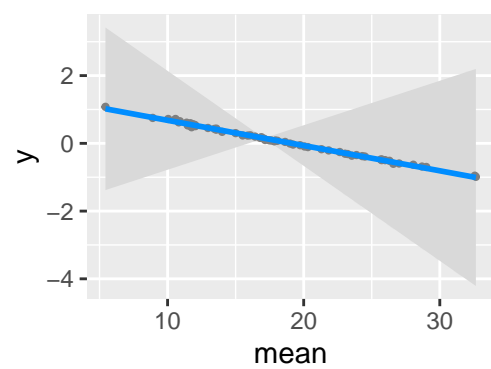
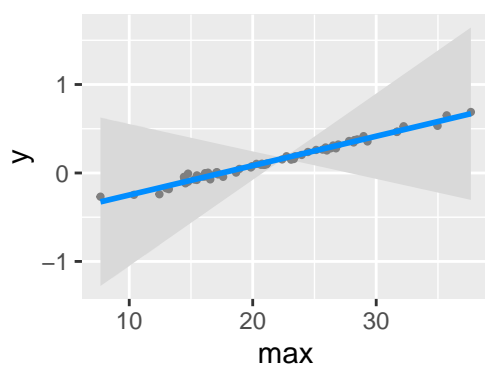
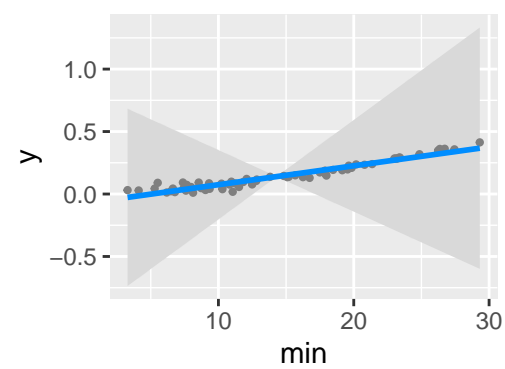


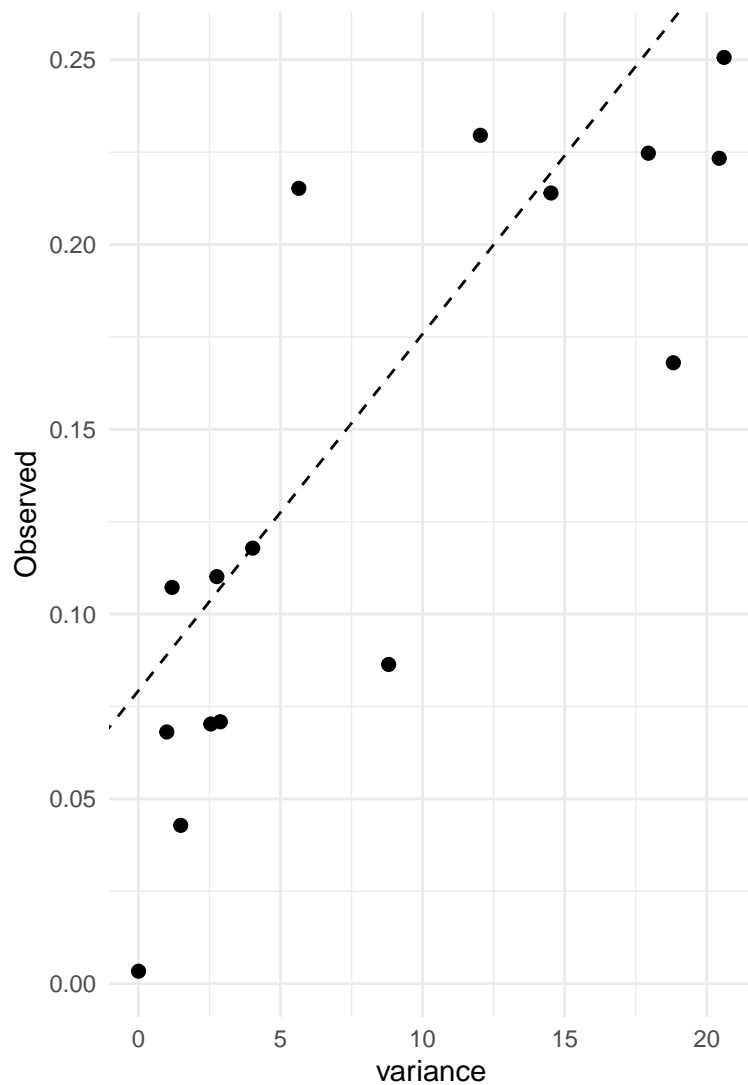
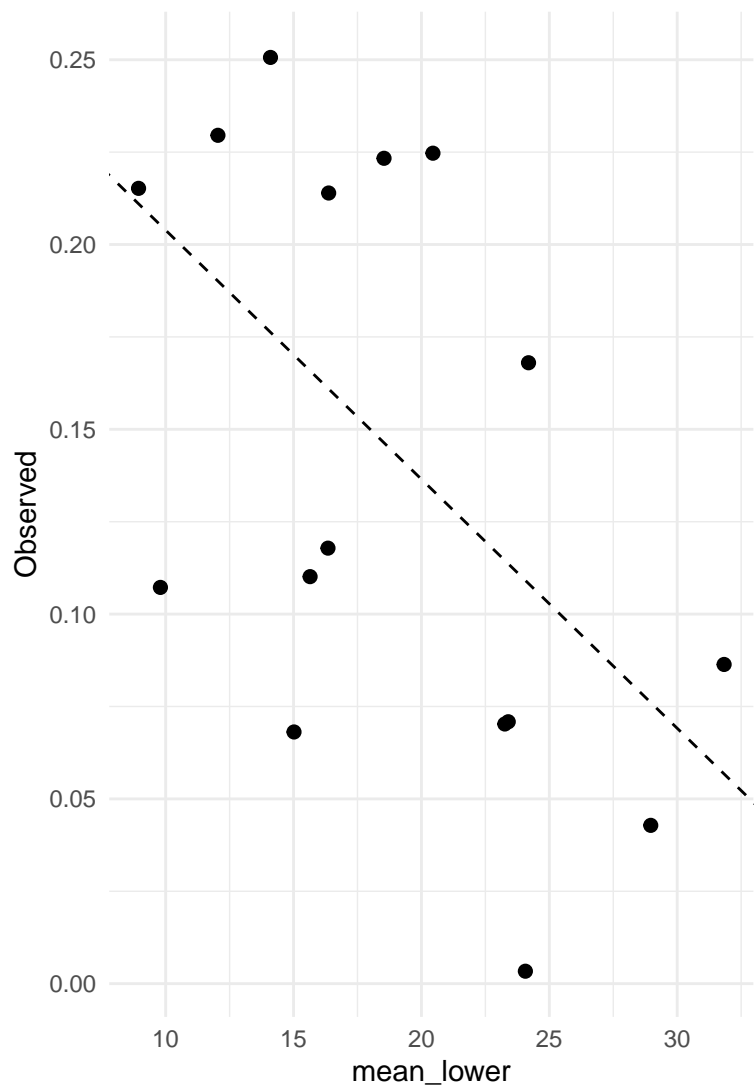
Mean_Upper Coefficients – gPi



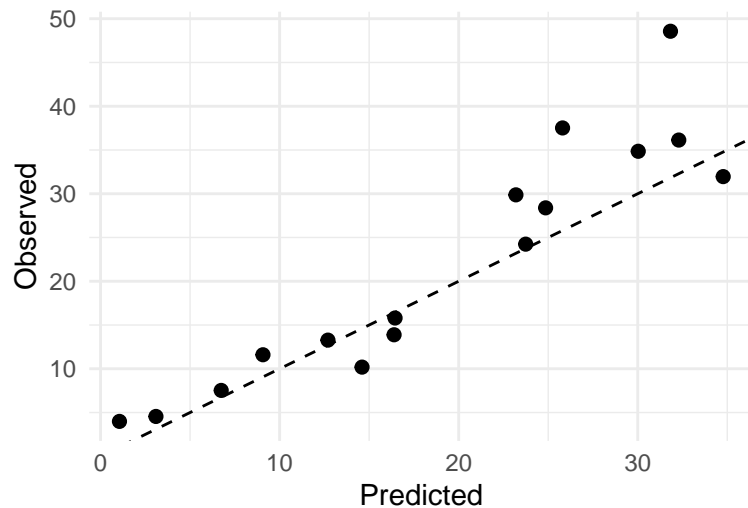
Ridge Coefficients – gPi



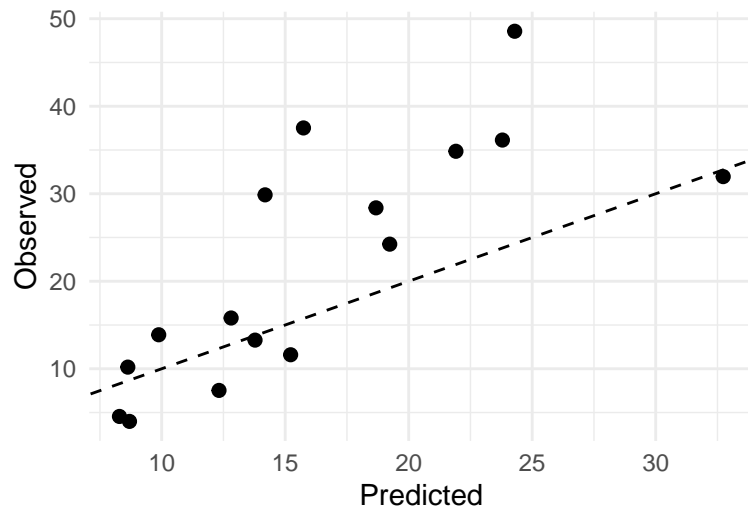


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{mean_lower}$ 

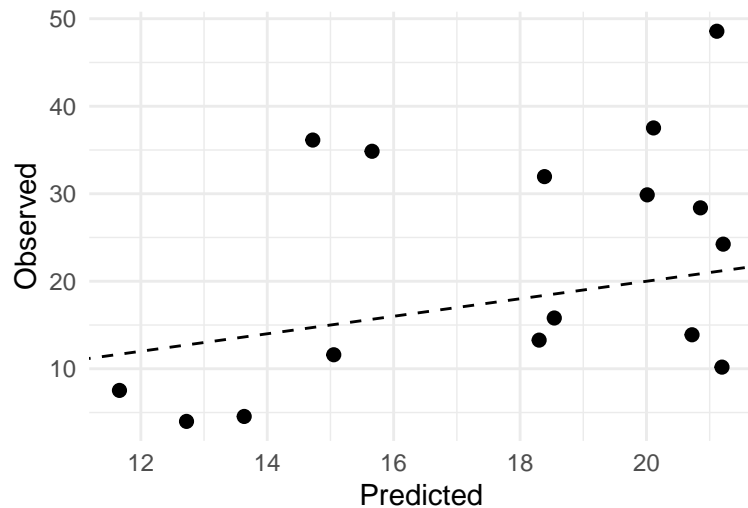
Full – PPF



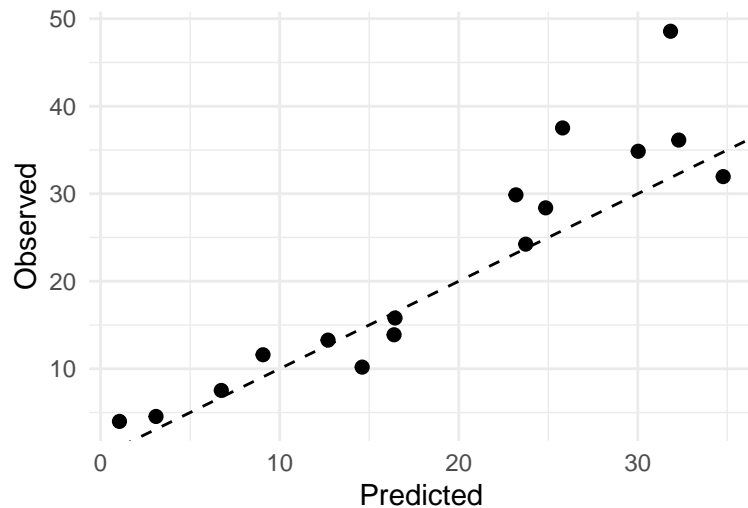
Range – PPF



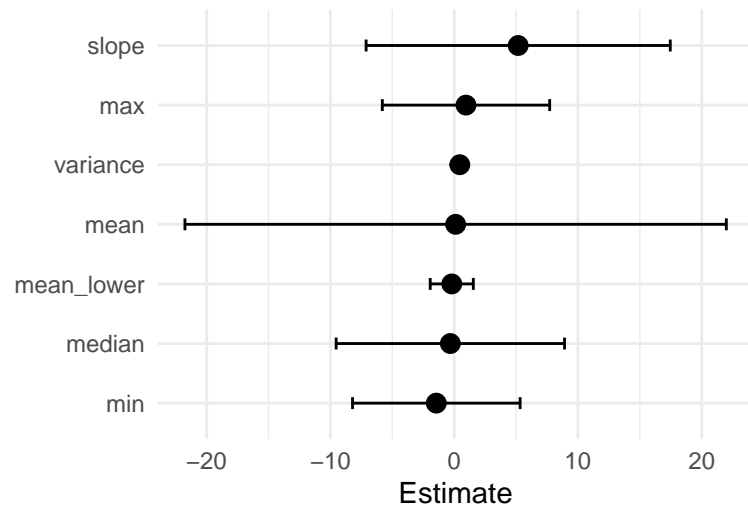
Mean_Upper – PPF



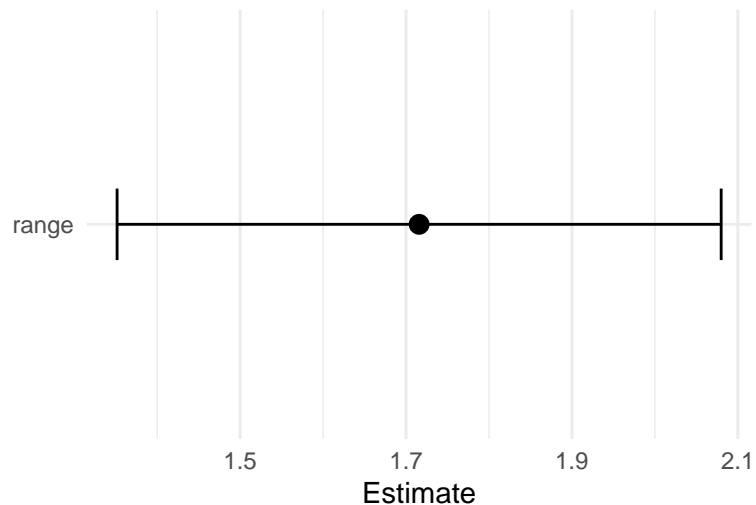
Ridge – PPF



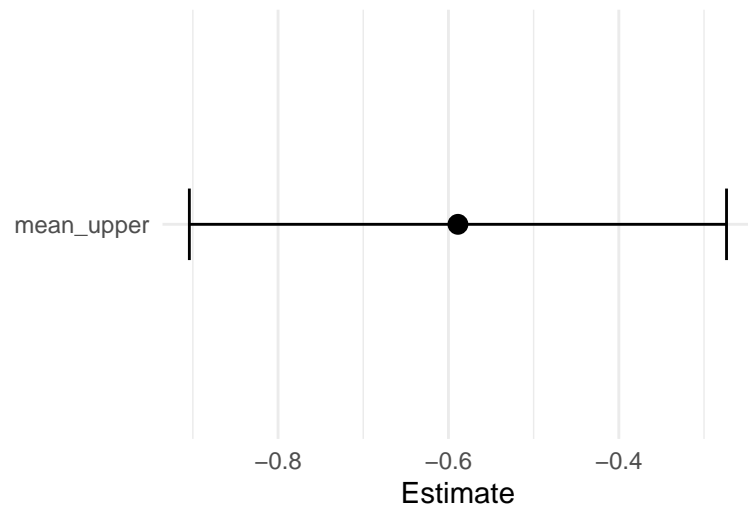
Full Coefficients – PPF



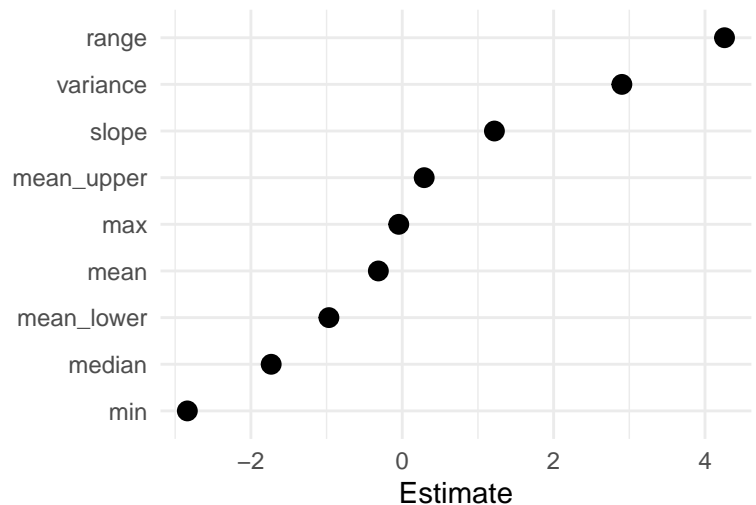
Range Coefficients – PPF

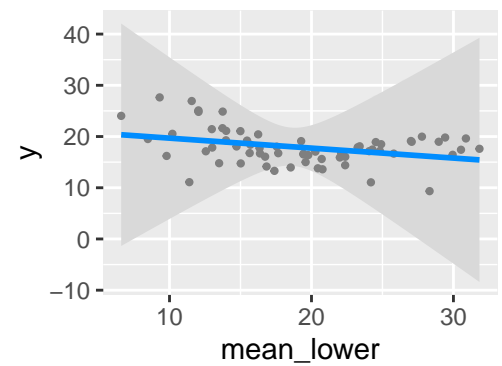
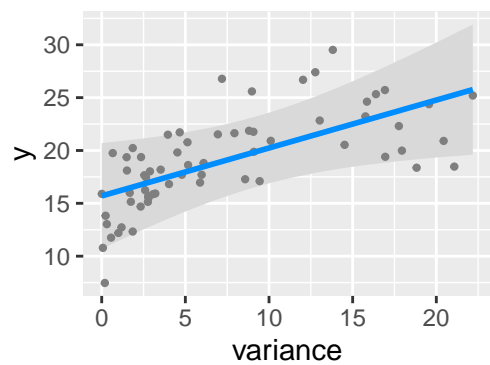
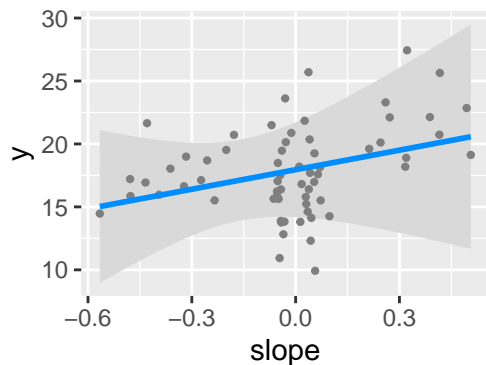
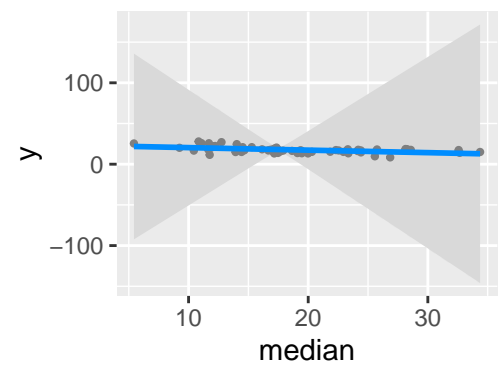
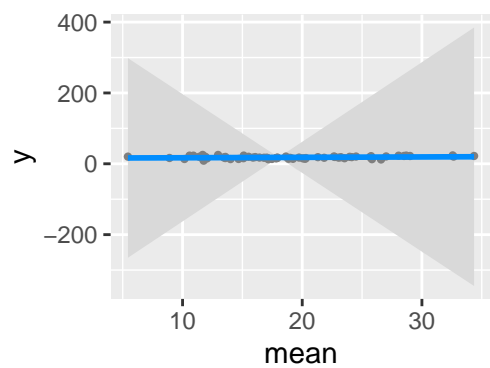
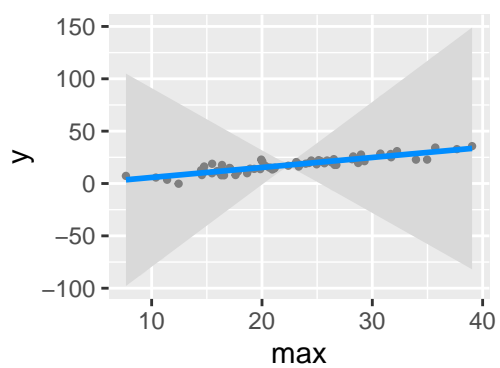
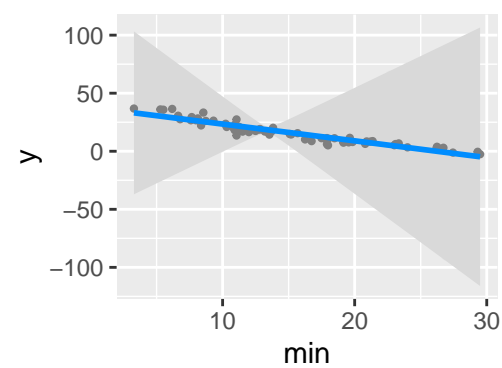


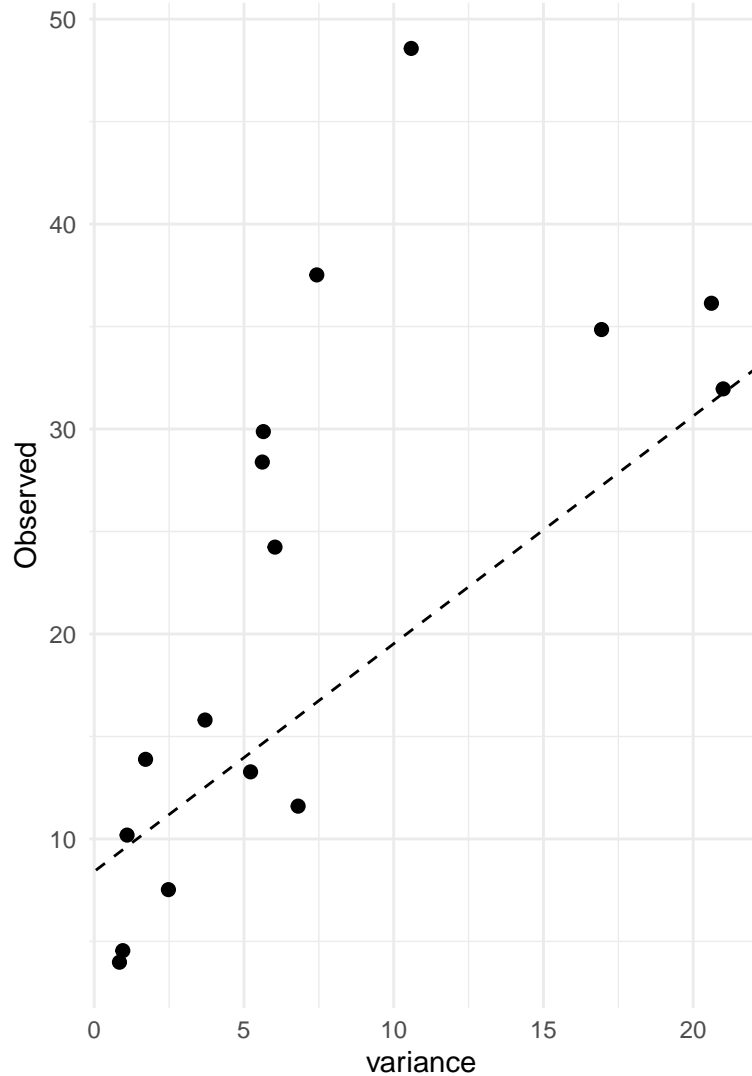
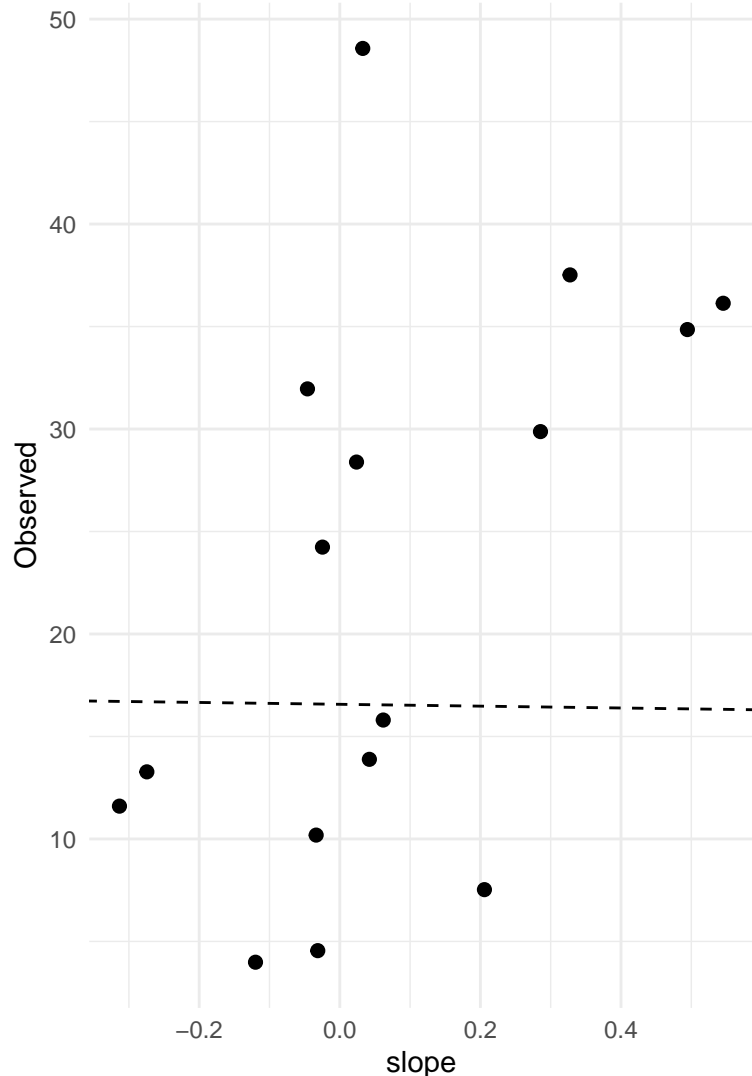
Mean_Upper Coefficients – PPF



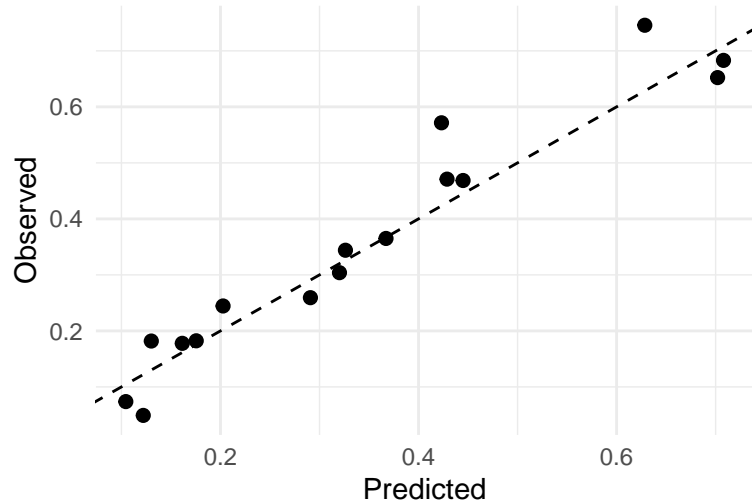
Ridge Coefficients – PPF



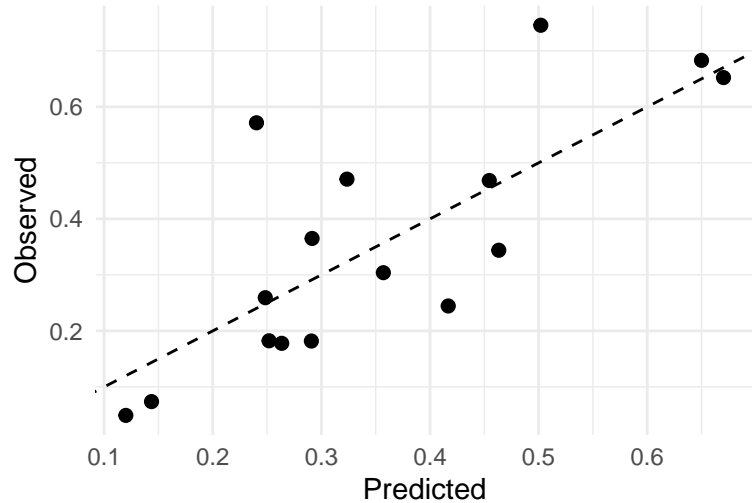


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{slope}$ 

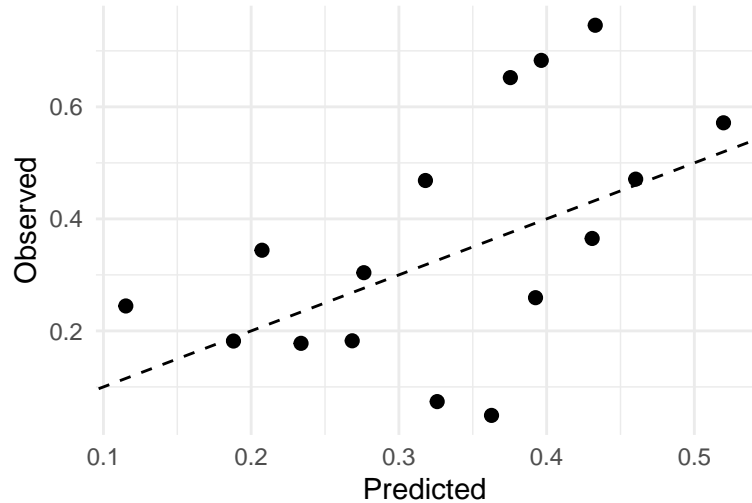
Full – PPI



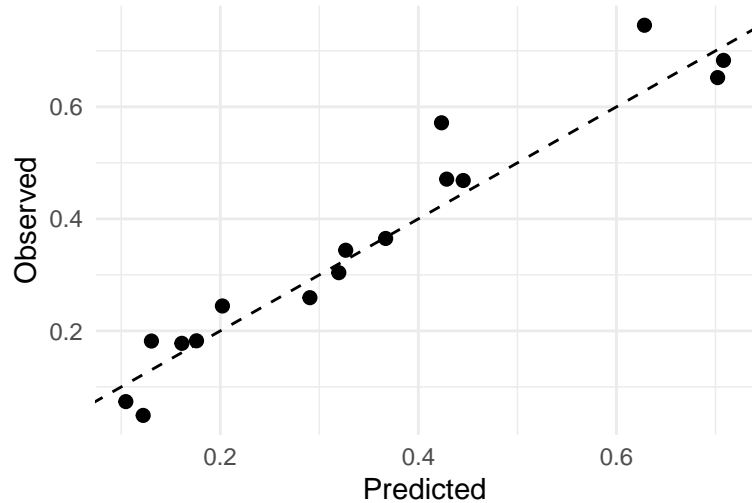
Range – PPI



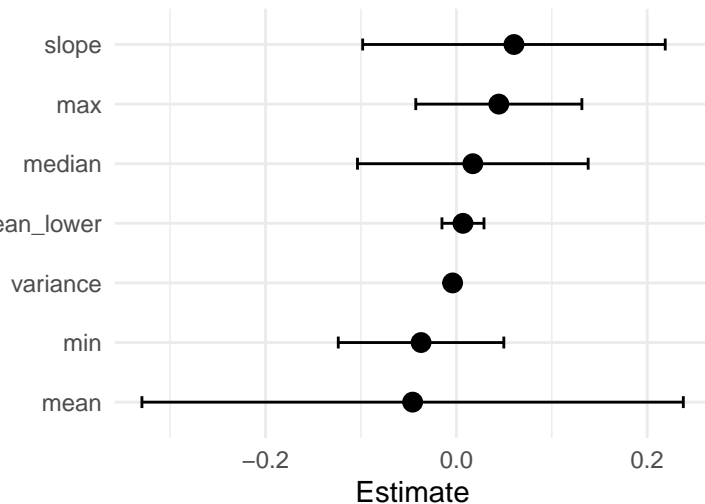
Mean_Upper – PPI



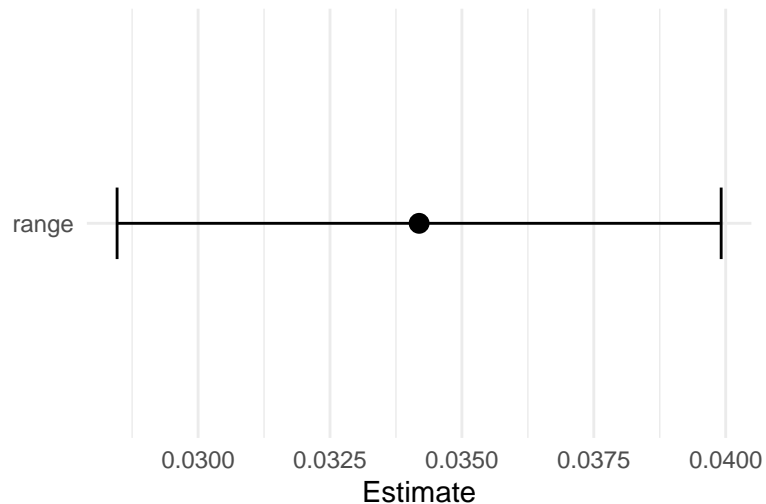
Ridge – PPI



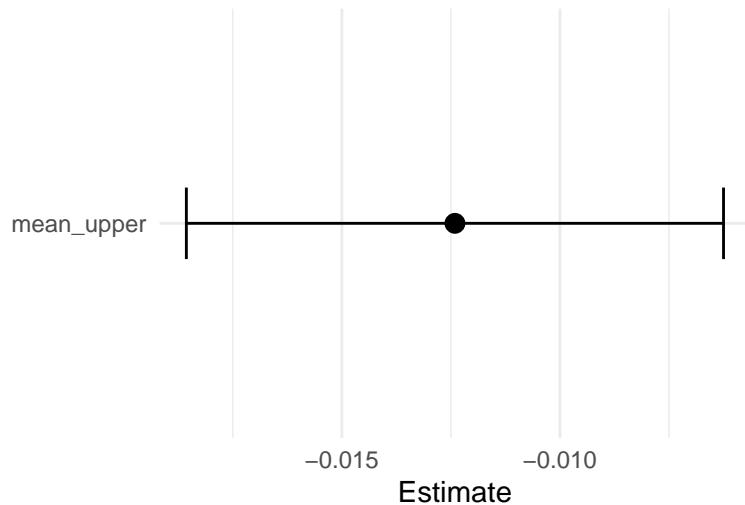
Full Coefficients – PPI



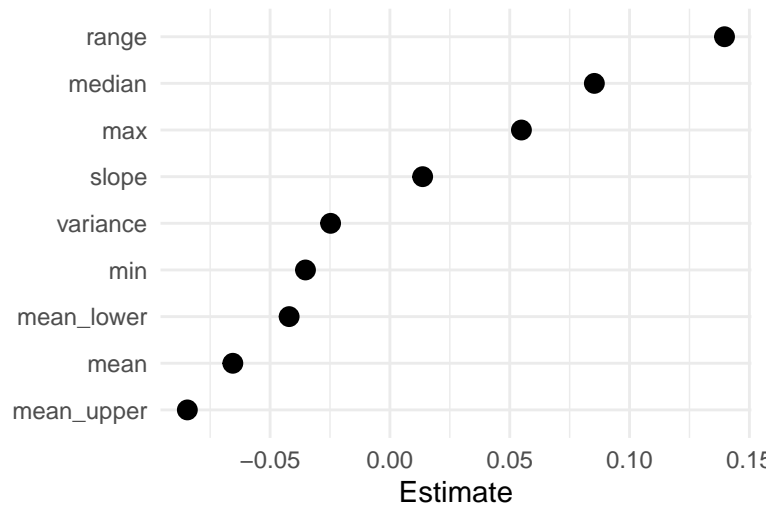
Range Coefficients – PPI

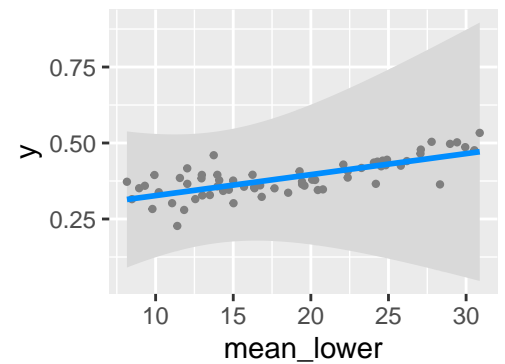
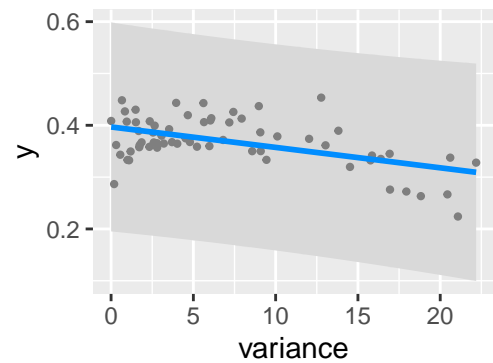
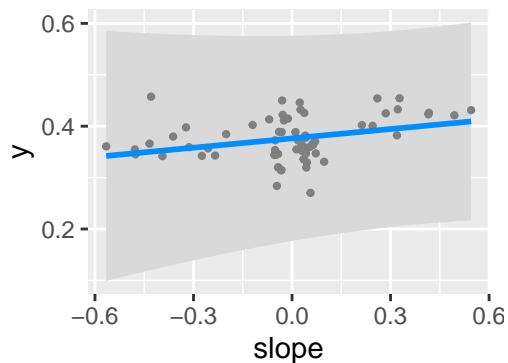
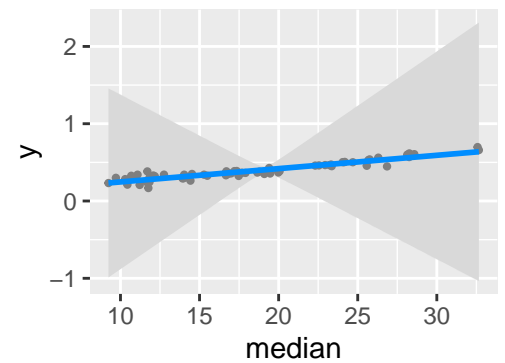
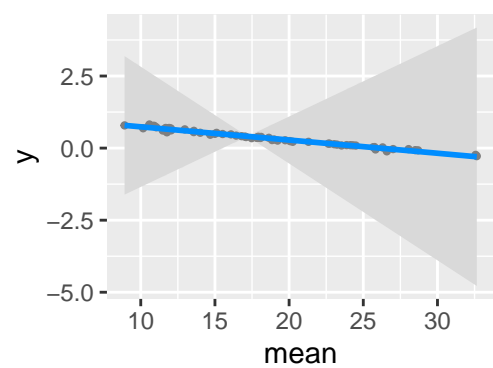
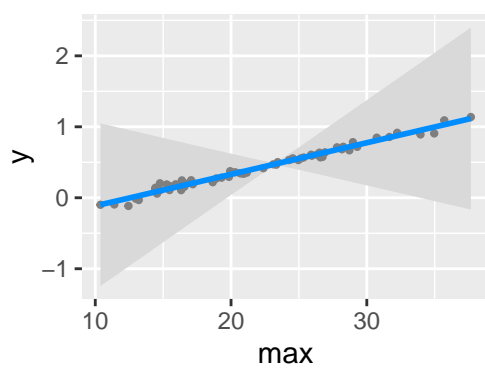
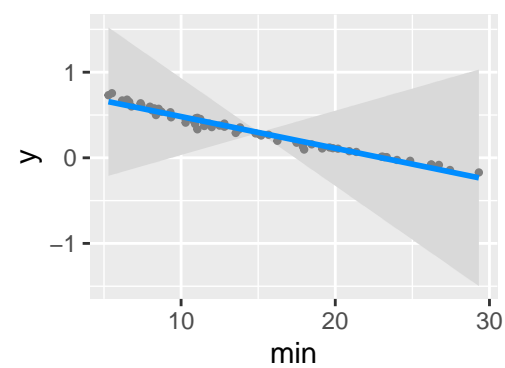


Mean_Upper Coefficients – PPI

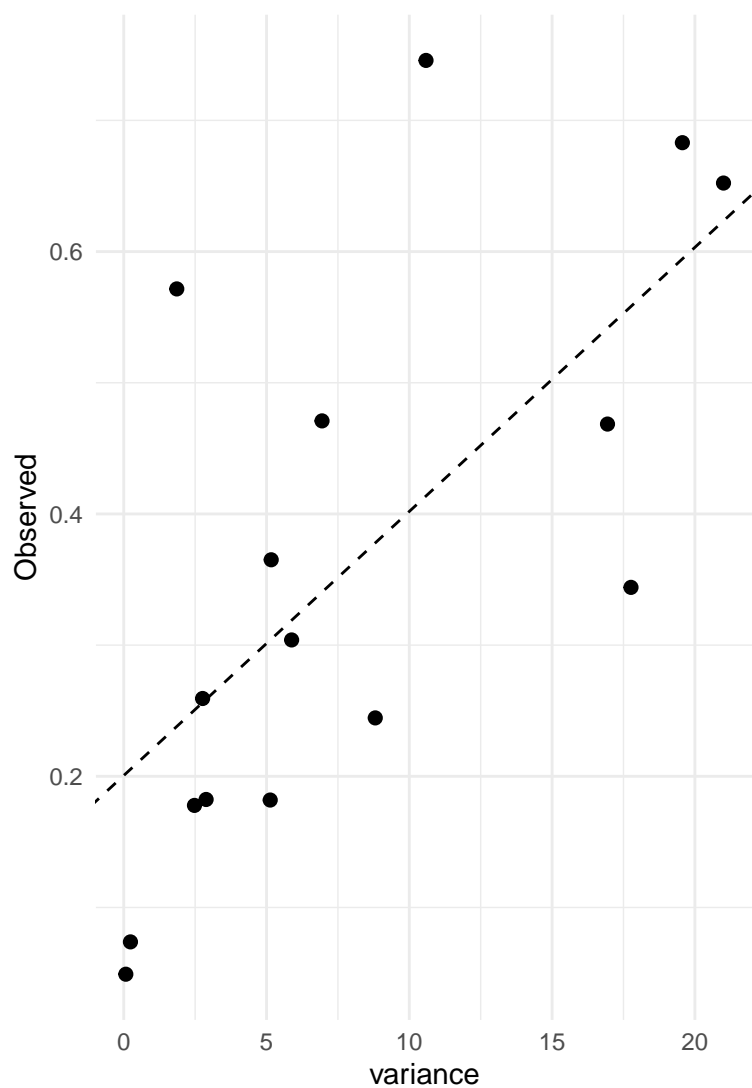


Ridge Coefficients – PPI

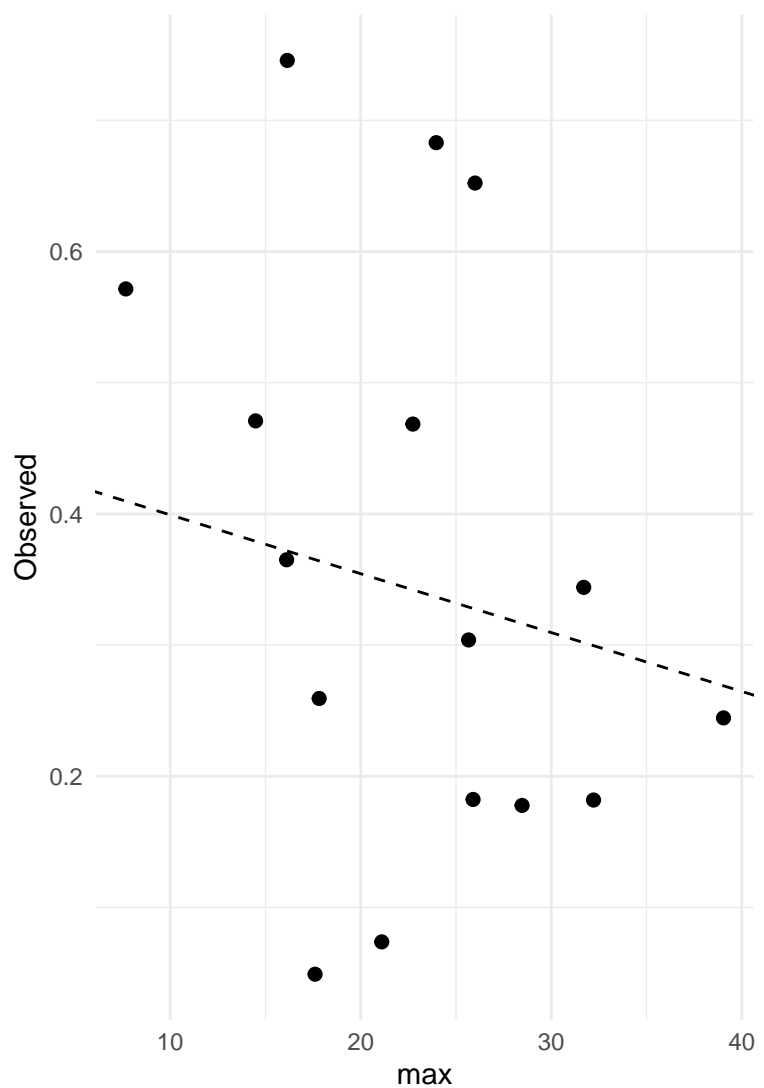




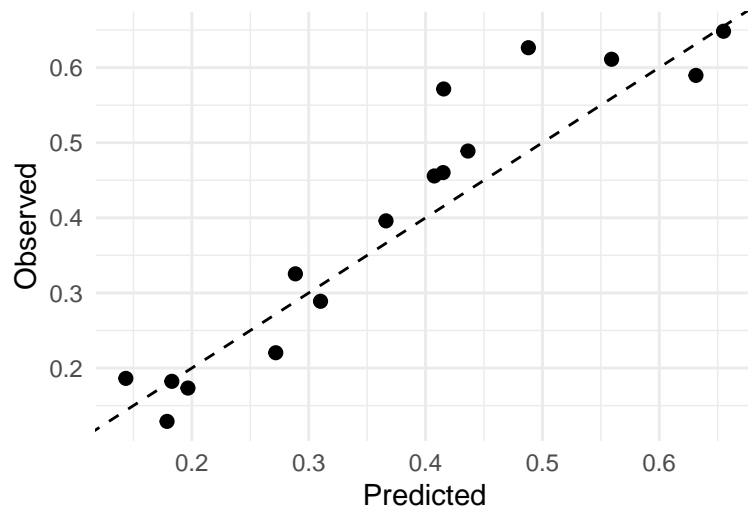
Simple Model: $y \sim \text{variance}$



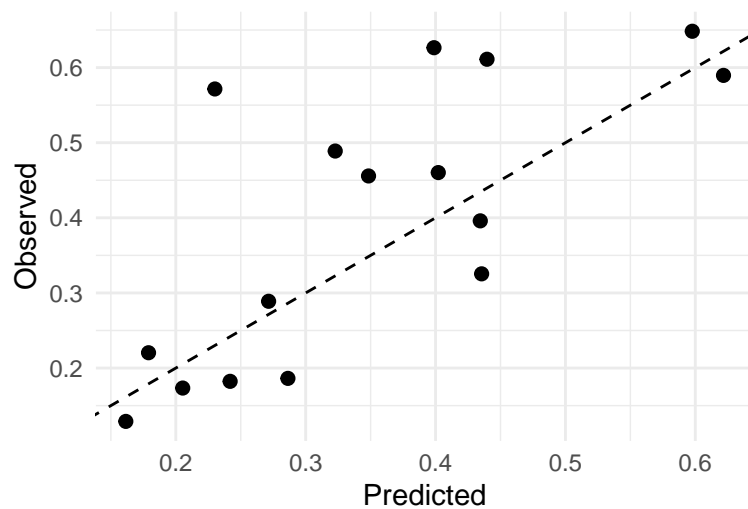
Simple Model: $y \sim \text{max}$



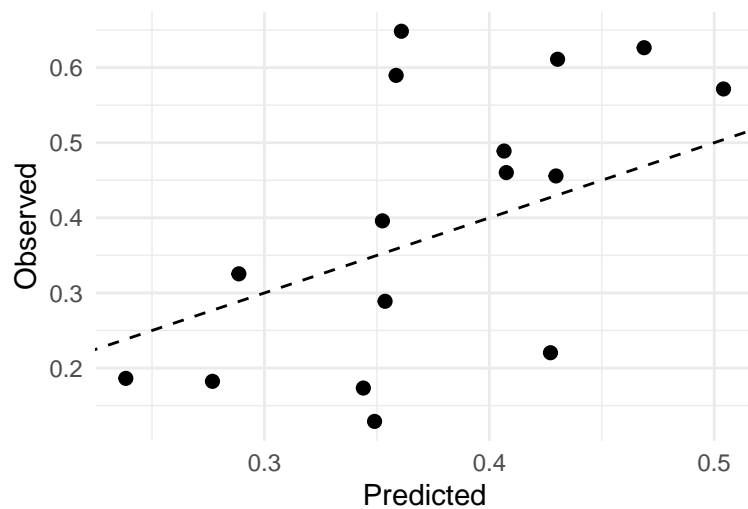
Full – Plmd



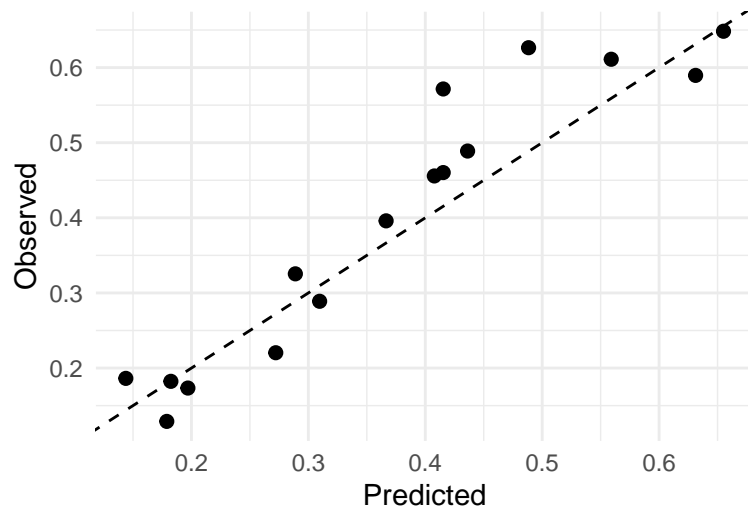
Range – Plmd



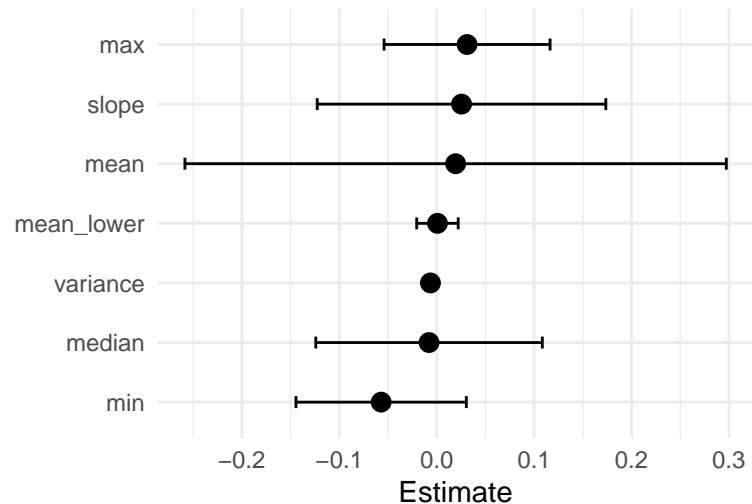
Mean_Upper – Plmd



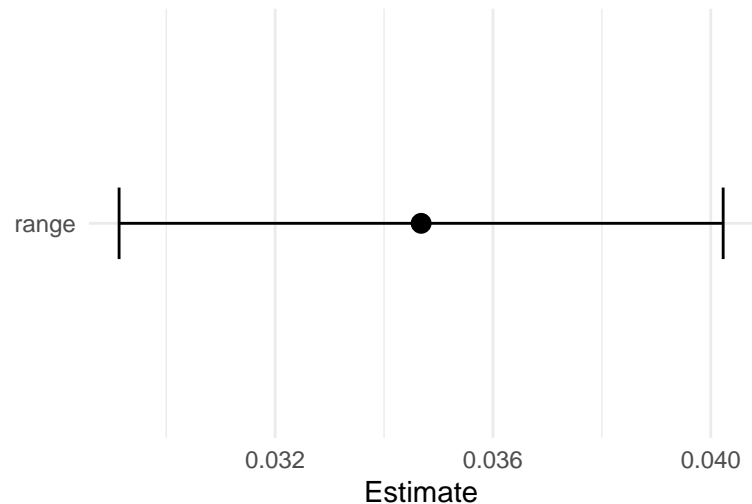
Ridge – Plmd



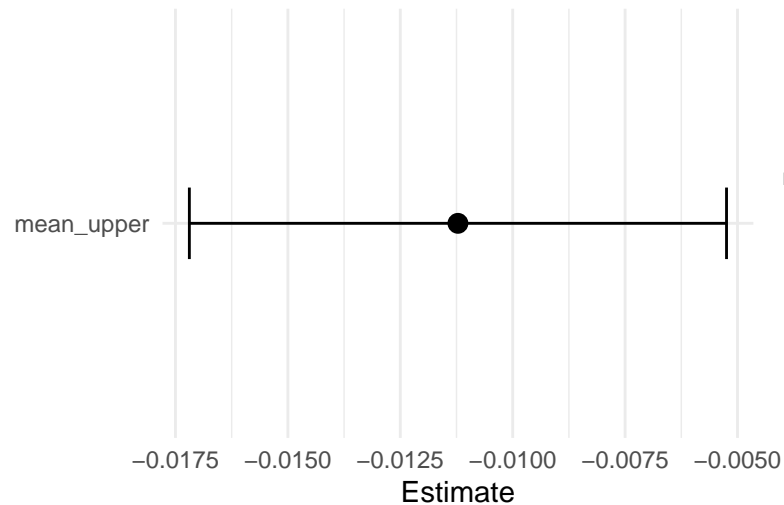
Full Coefficients – Plmd



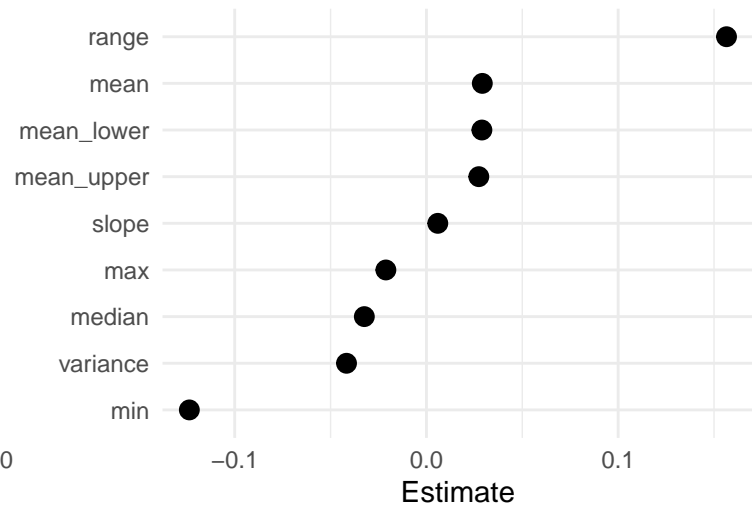
Range Coefficients – Plmd

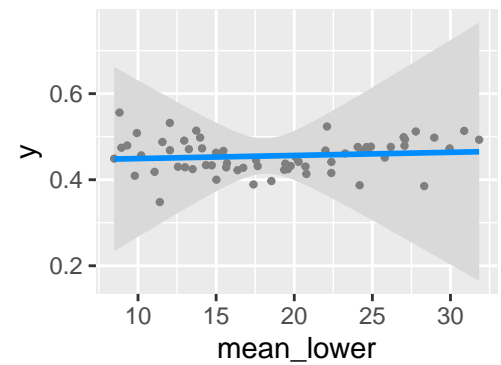
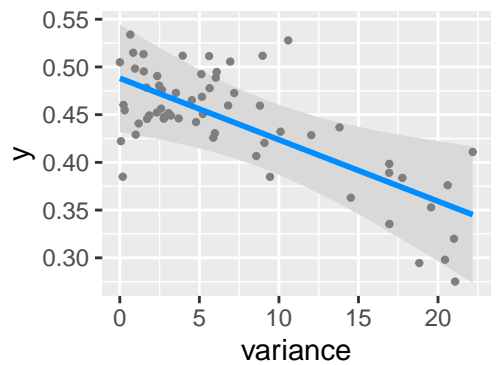
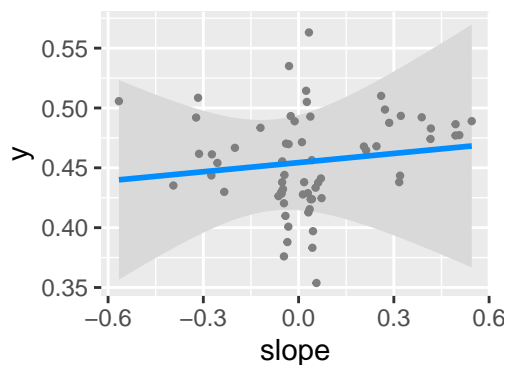
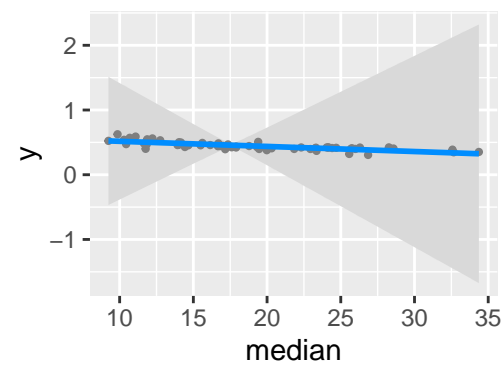
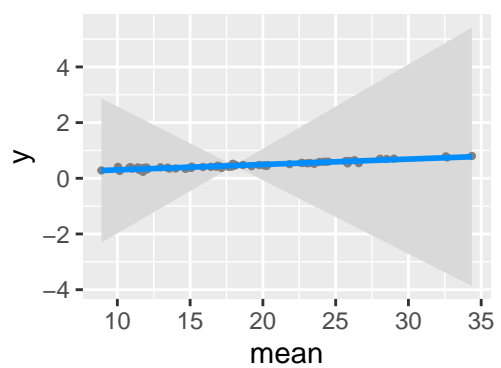
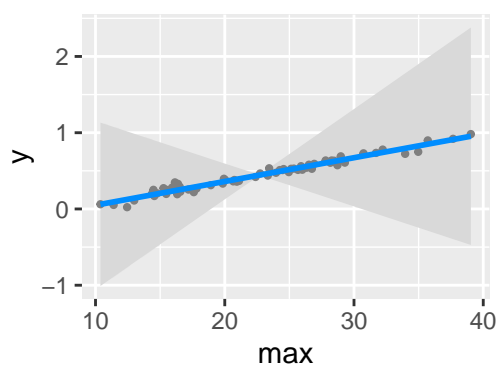
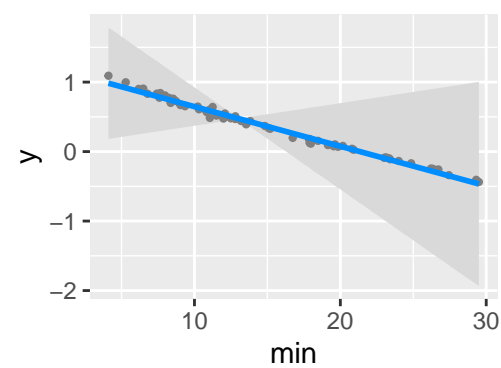


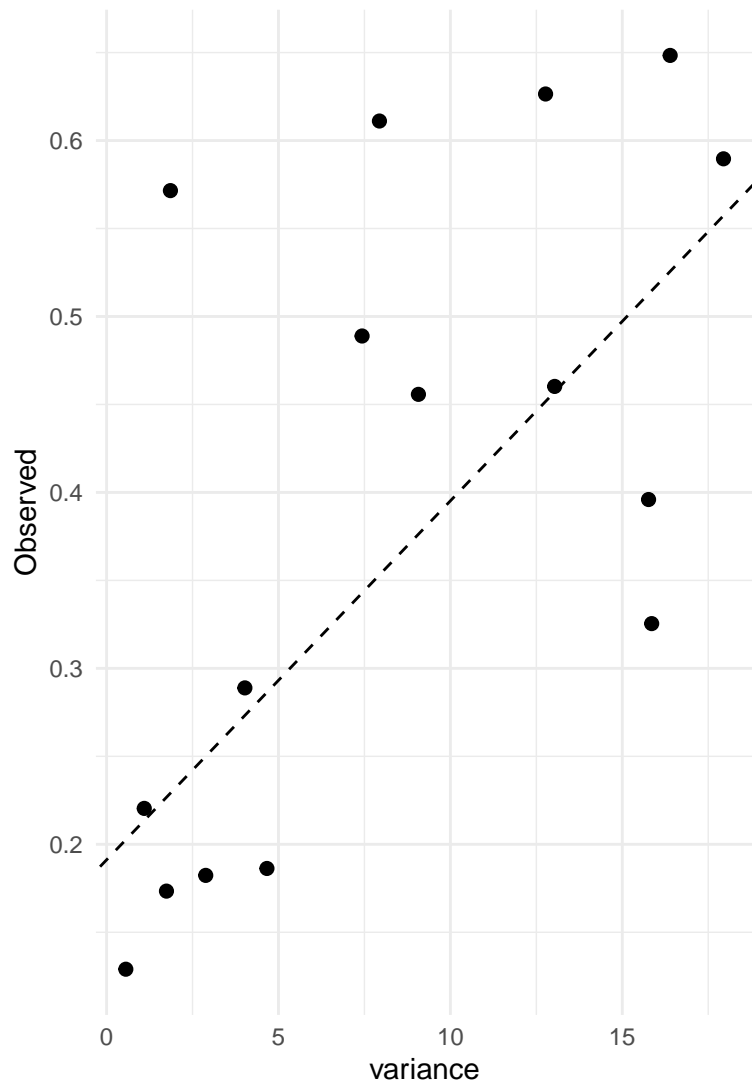
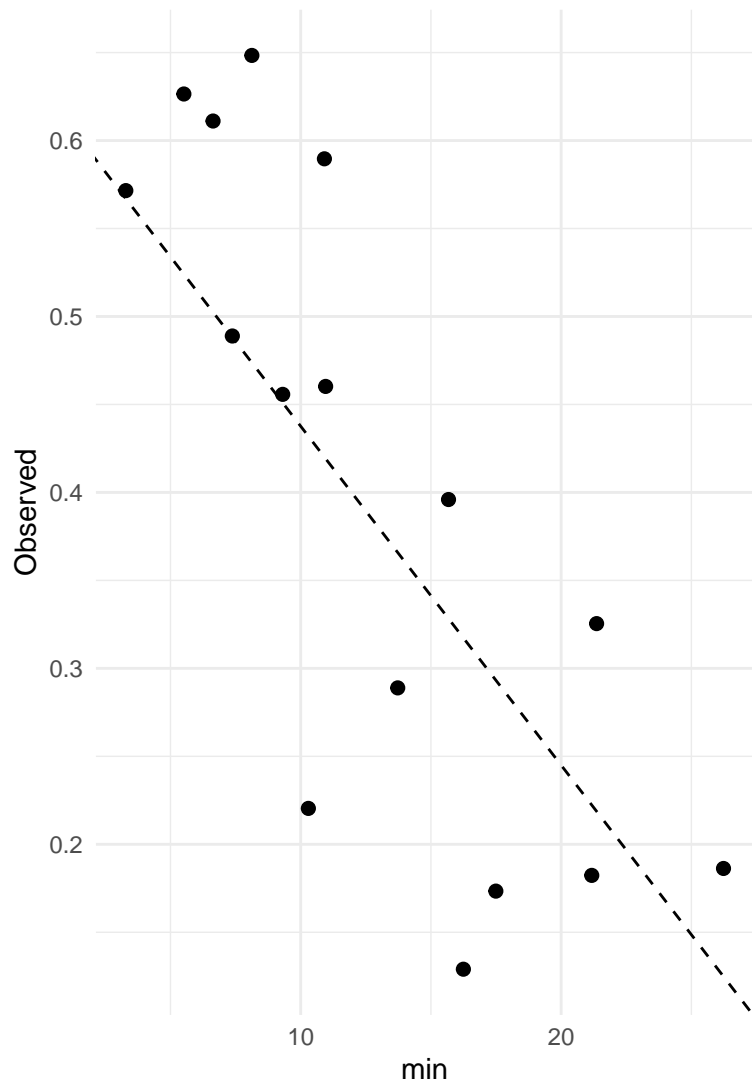
Mean_Upper Coefficients – Plmd



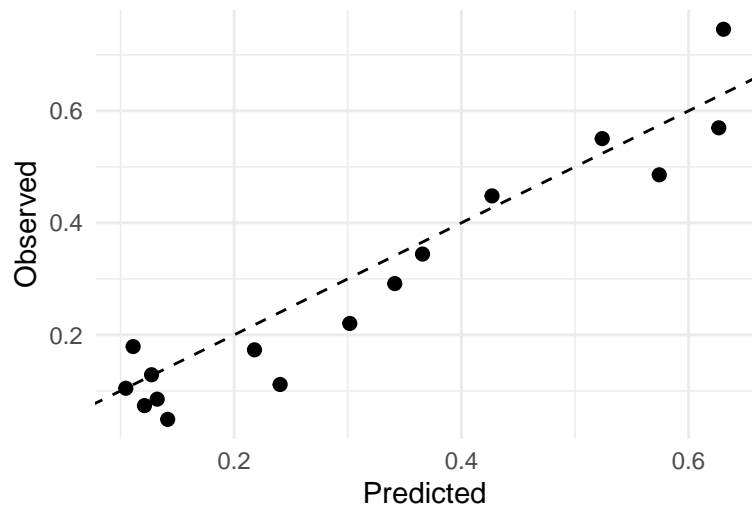
Ridge Coefficients – Plmd



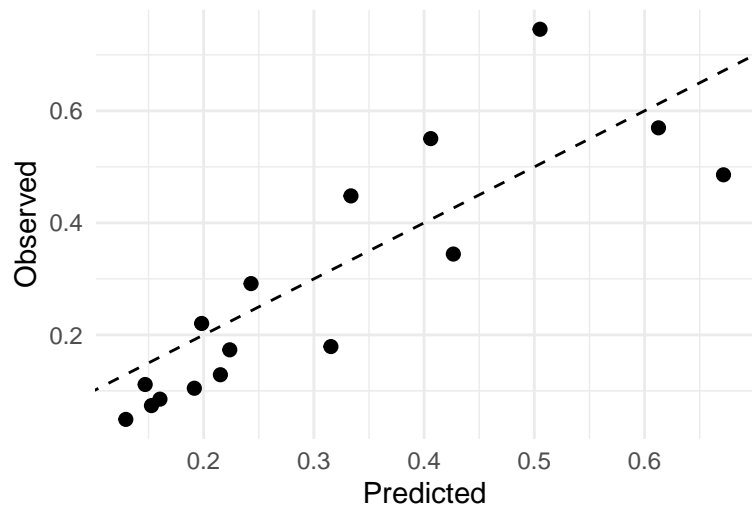


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{min}$ 

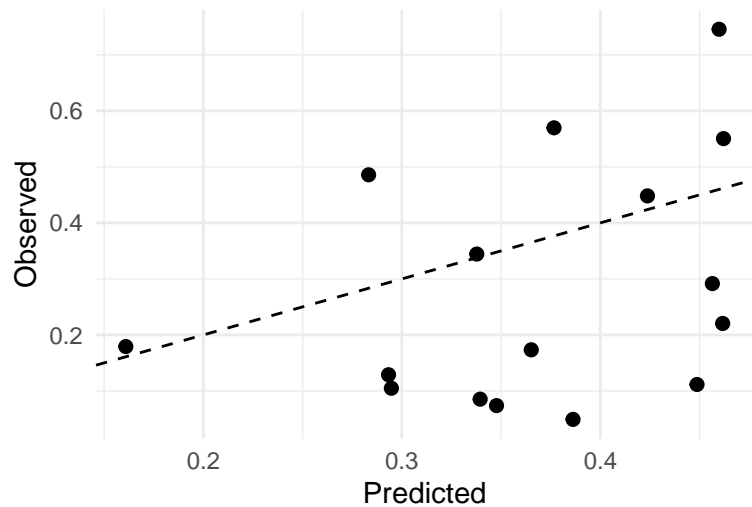
Full – PILSM



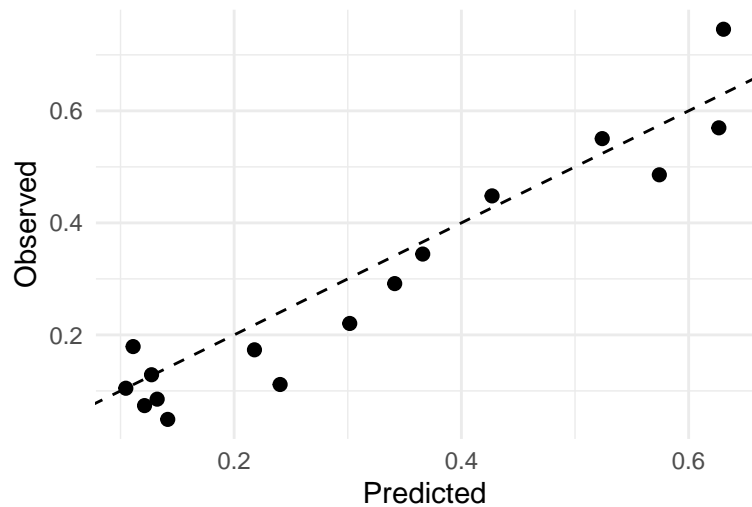
Range – PILSM



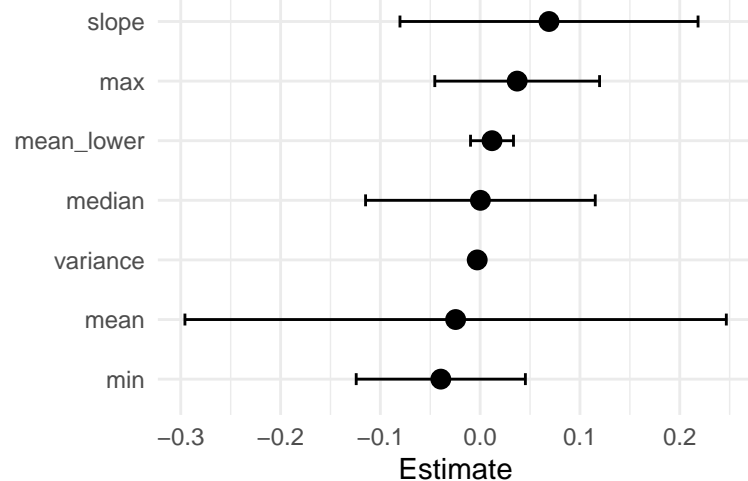
Mean_Upper – PILSM



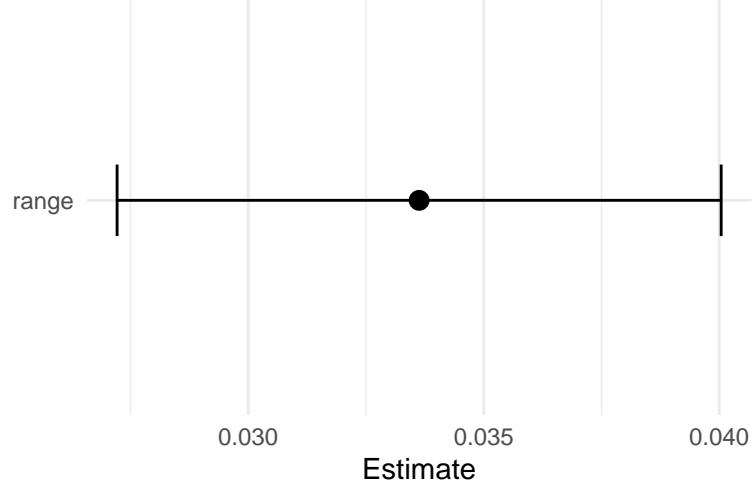
Ridge – PILSM



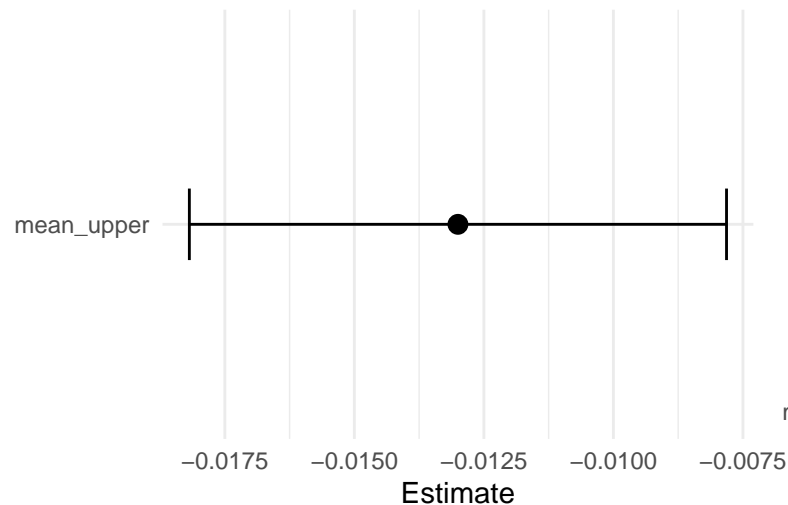
Full Coefficients – PILSM



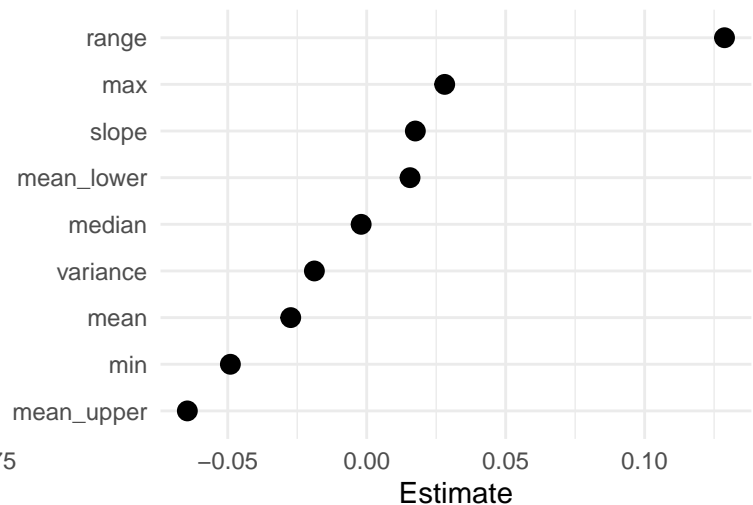
Range Coefficients – PILSM

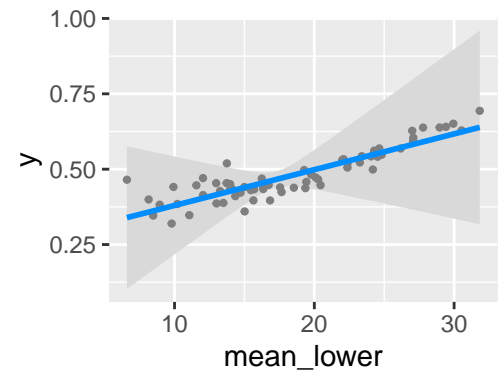
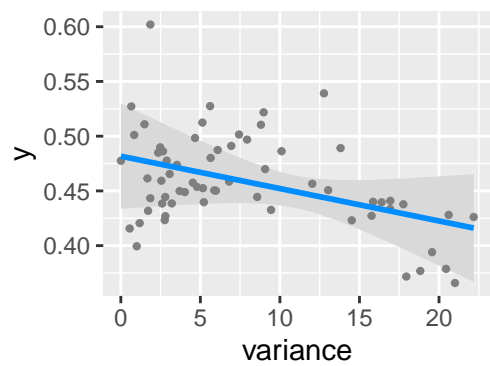
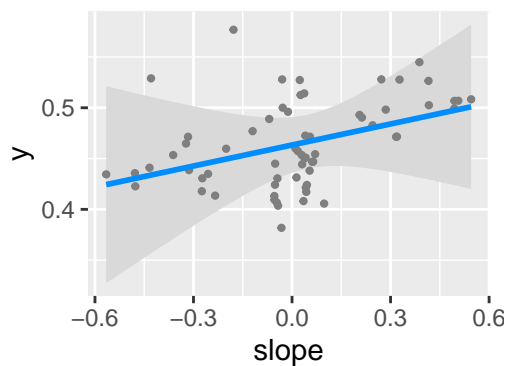
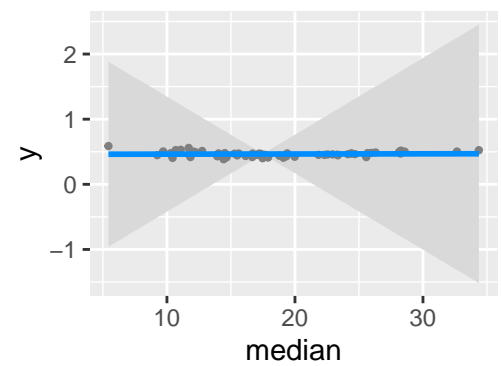
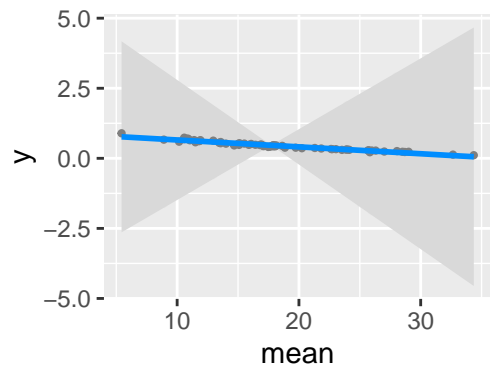
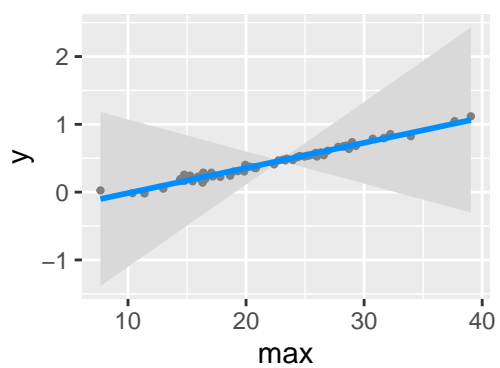
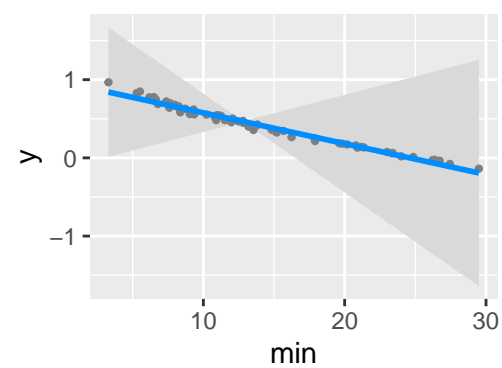


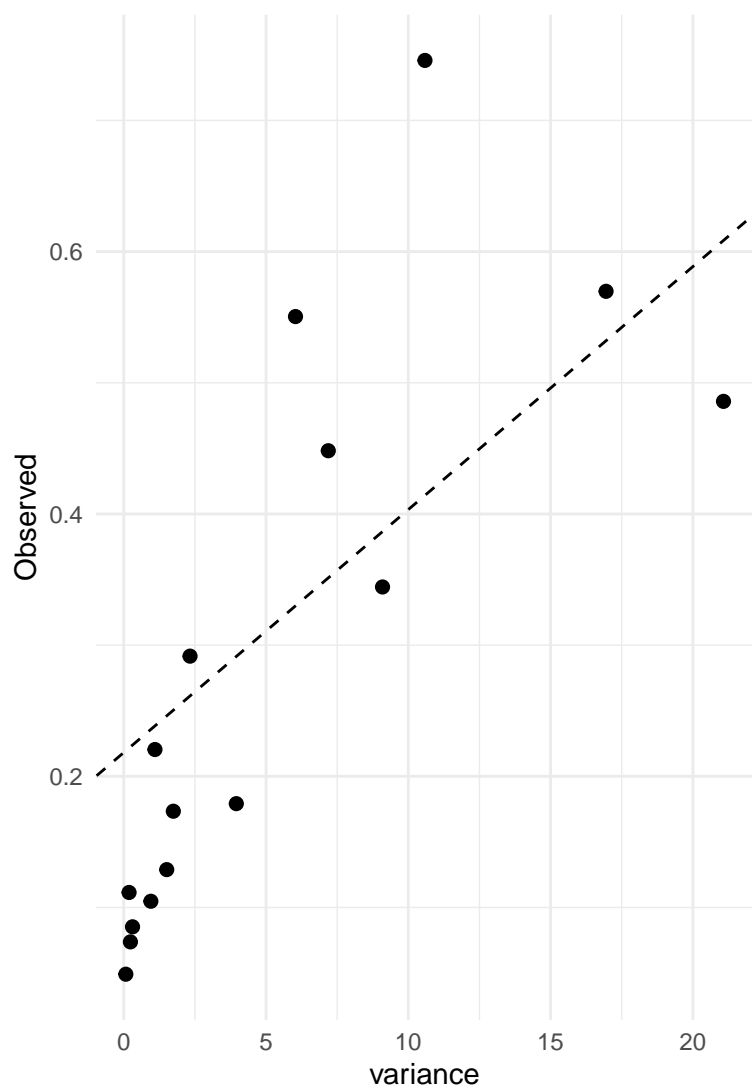
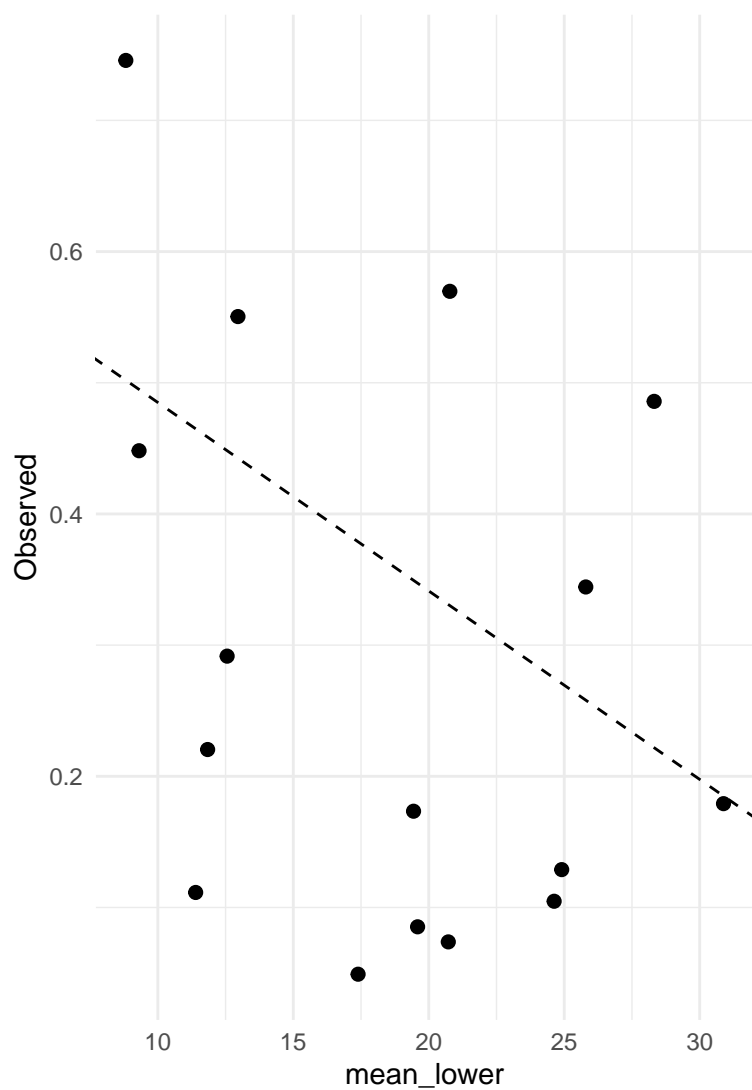
Mean_Upper Coefficients – PILSM



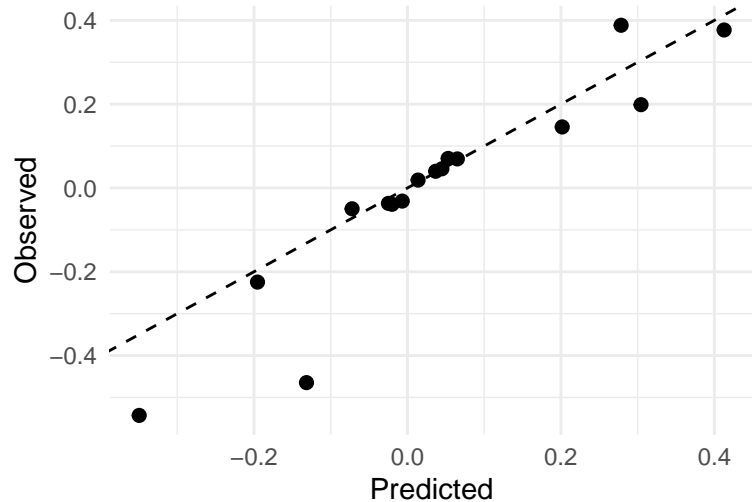
Ridge Coefficients – PILSM



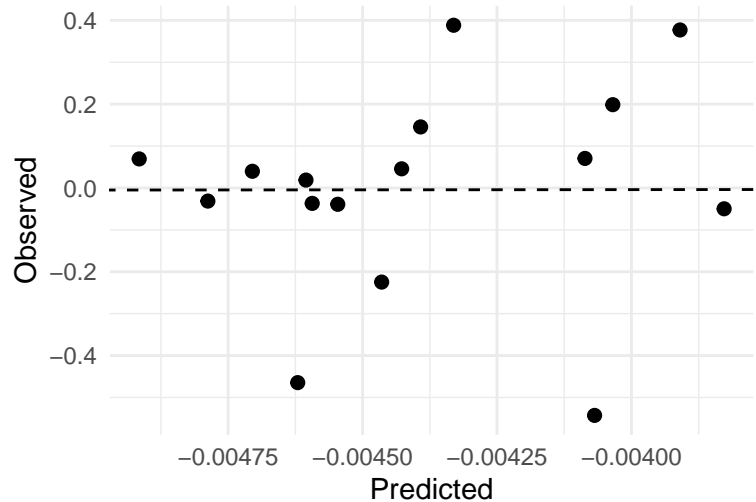


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{mean_lower}$ 

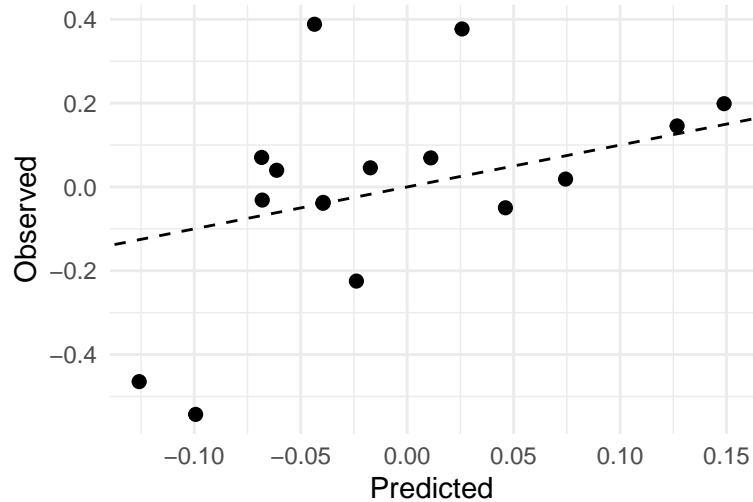
Full – RTR



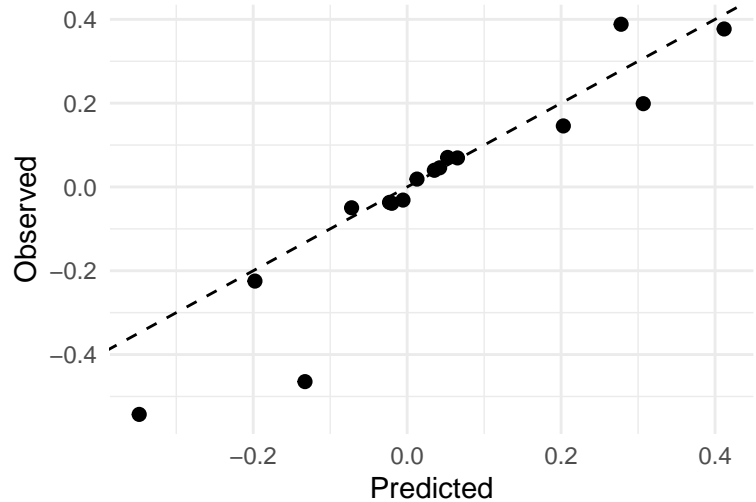
Range – RTR



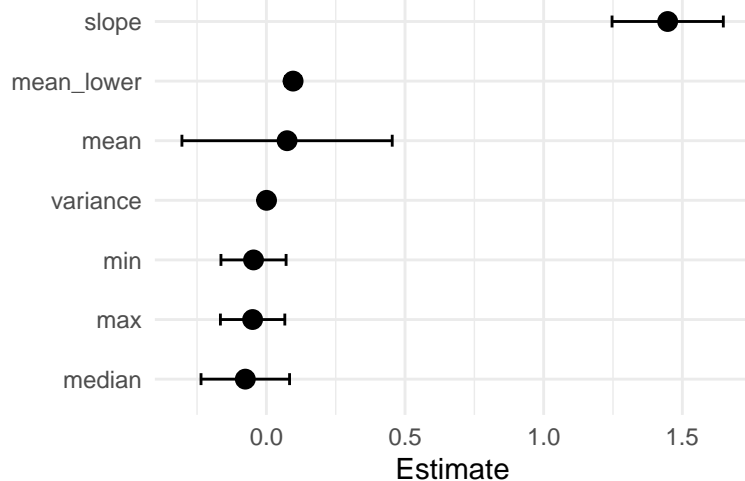
Mean_Upper – RTR



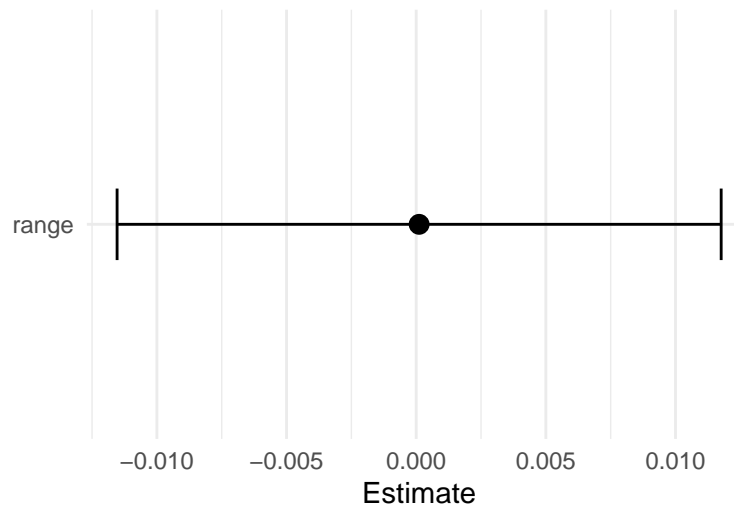
Ridge – RTR



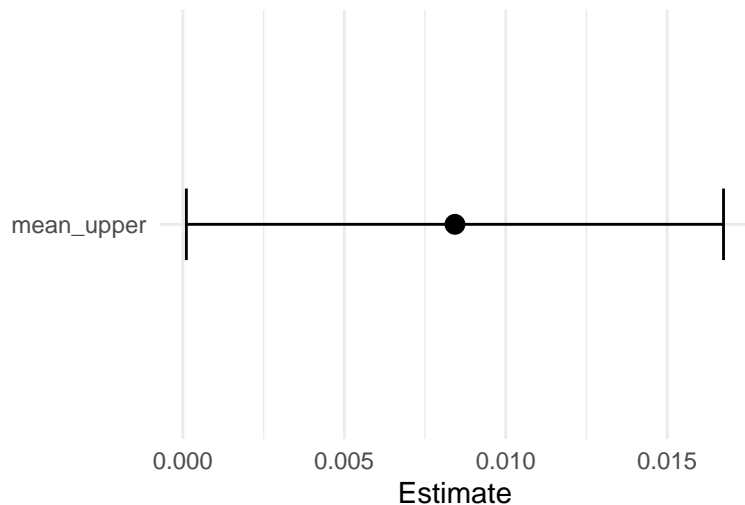
Full Coefficients – RTR



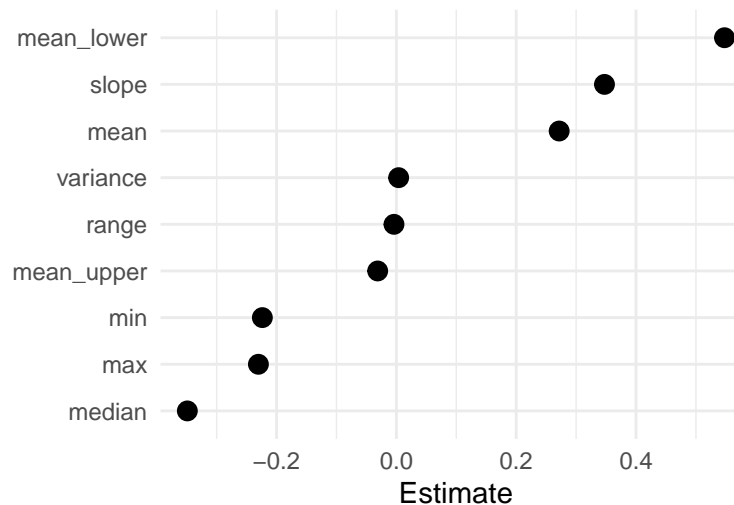
Range Coefficients – RTR

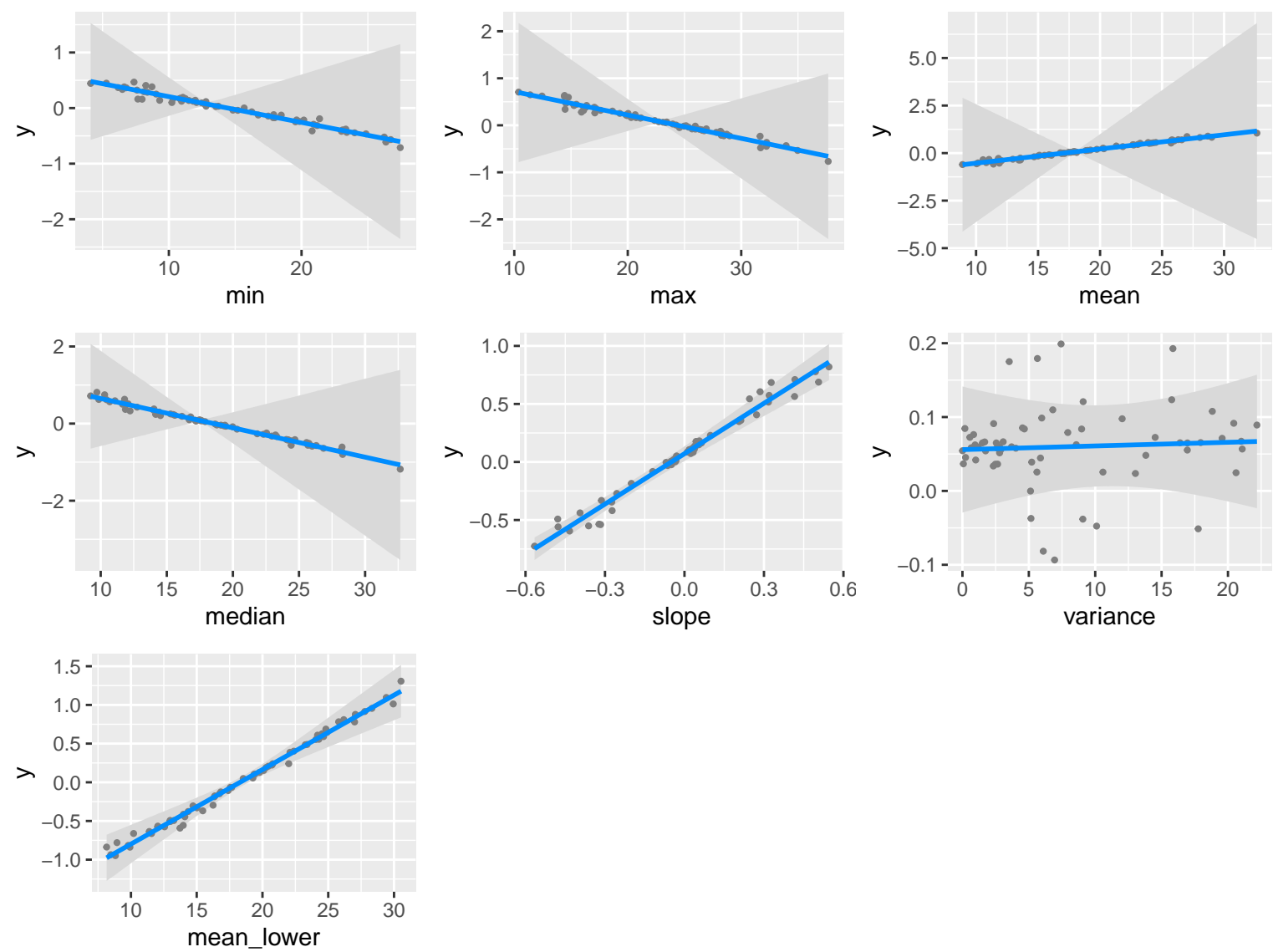


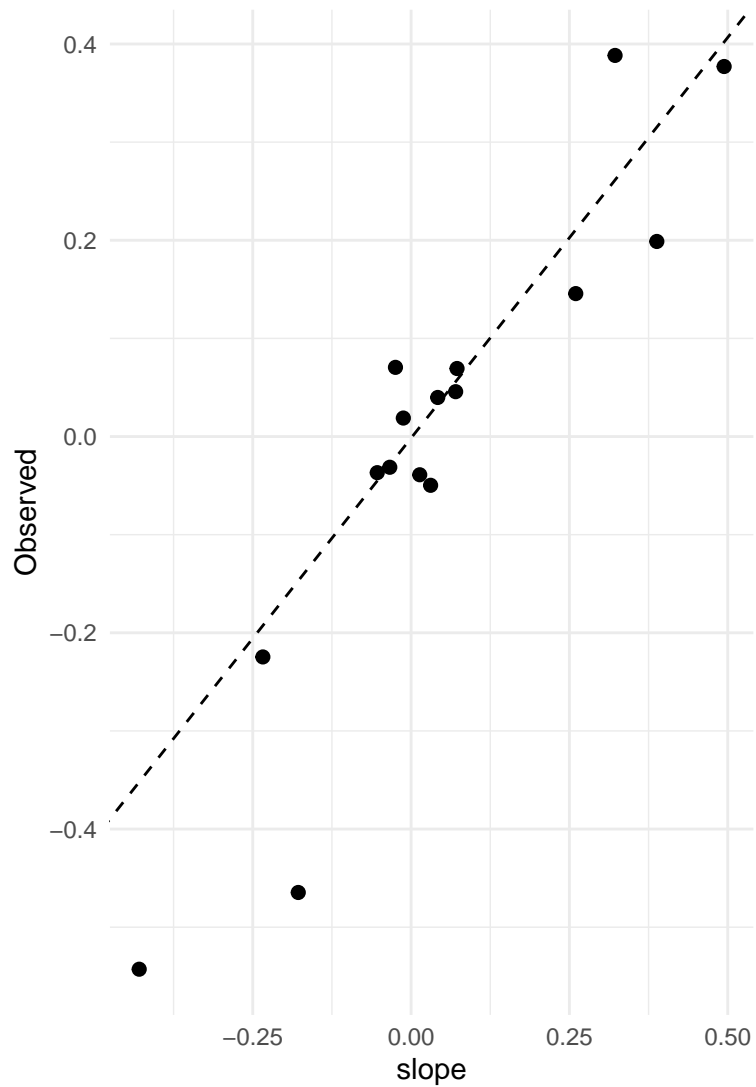
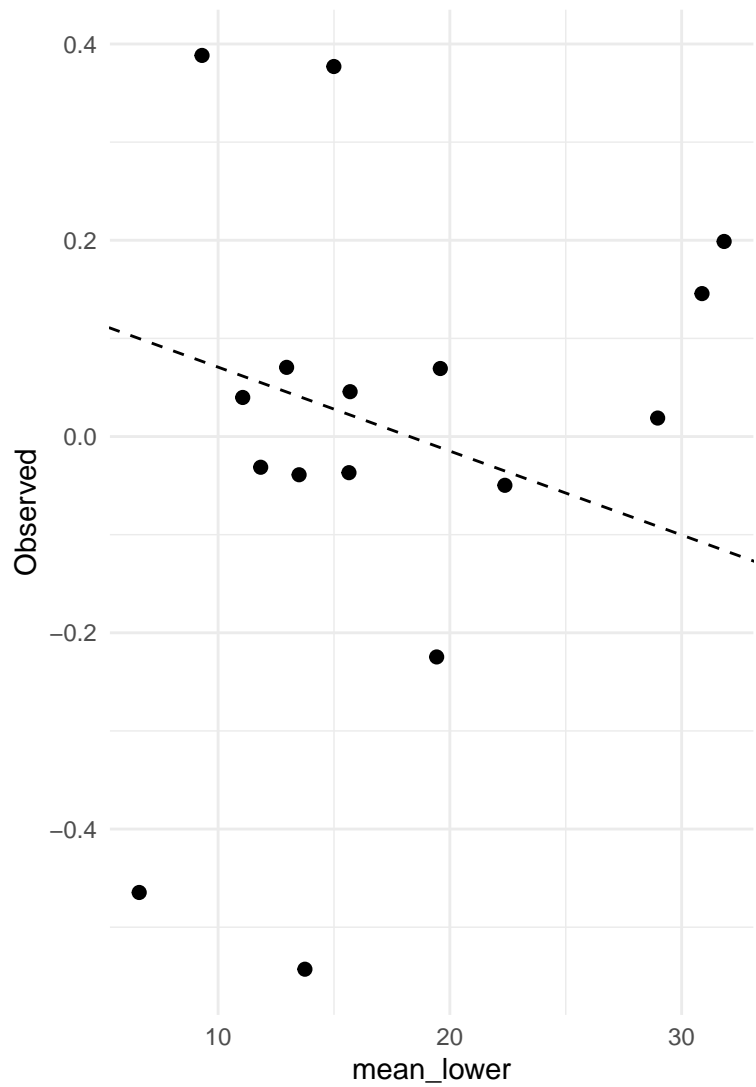
Mean_Upper Coefficients – RTR



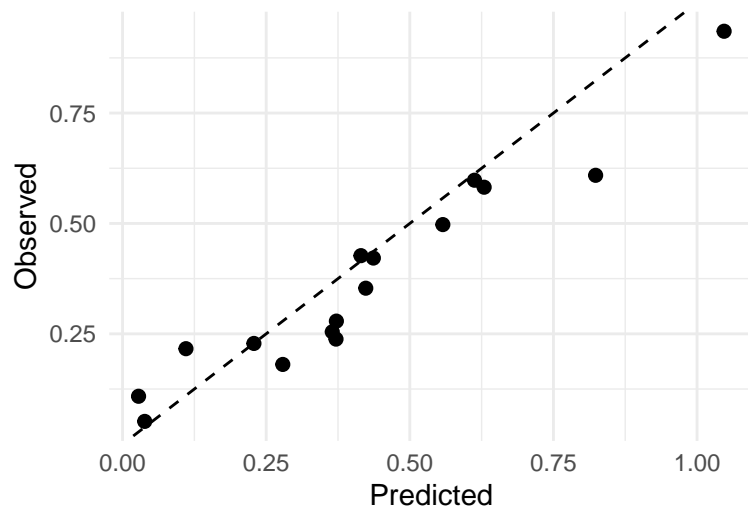
Ridge Coefficients – RTR



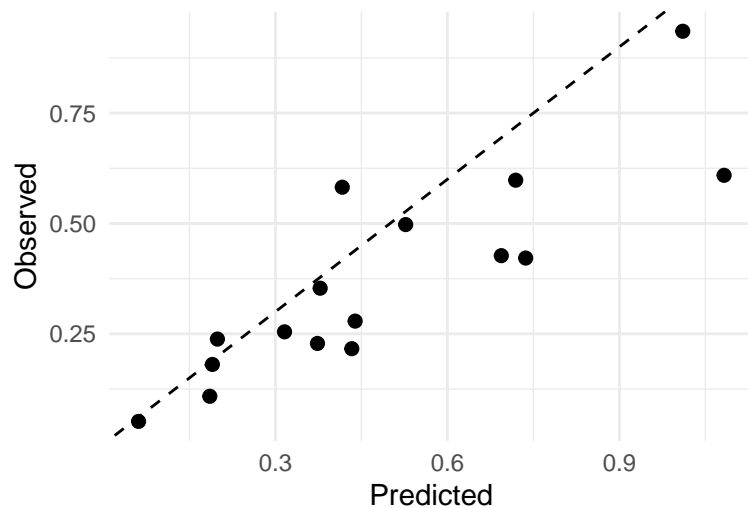


Simple Model: $y \sim \text{slope}$ Simple Model: $y \sim \text{mean_lower}$ 

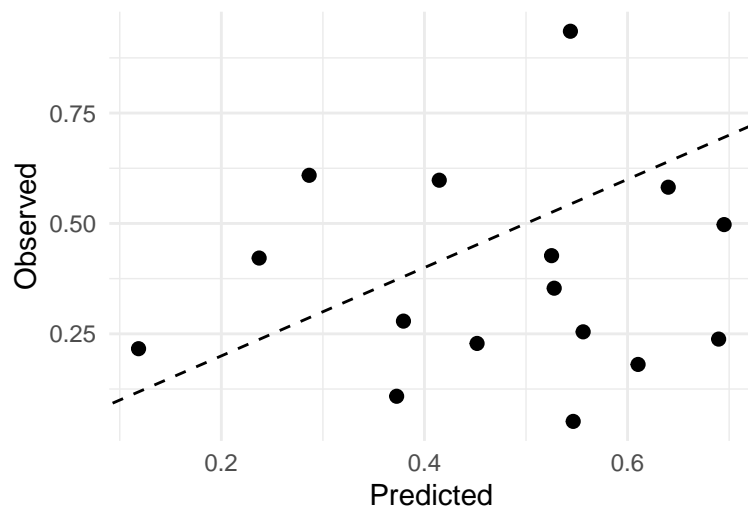
Full – PIR



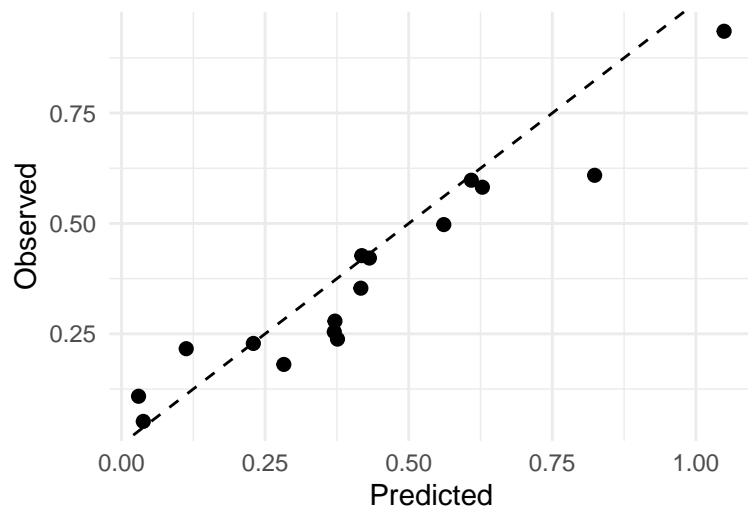
Range – PIR



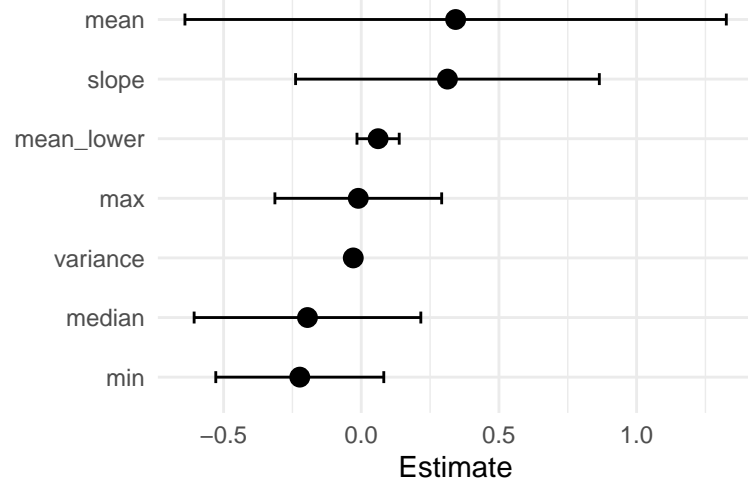
Mean_Upper – PIR



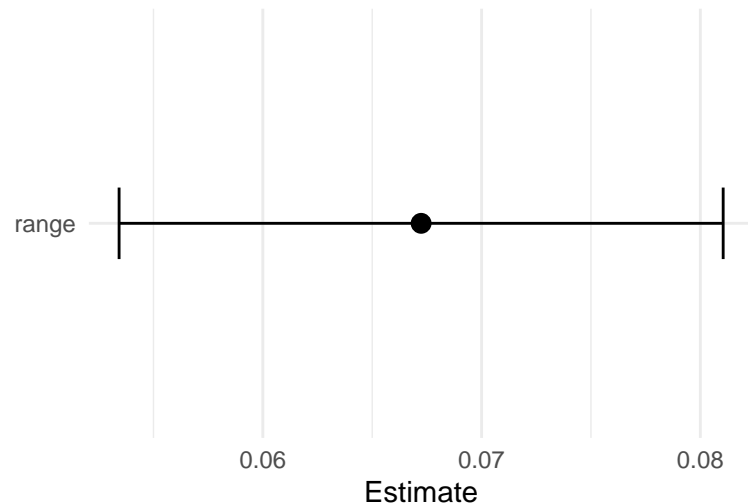
Ridge – PIR



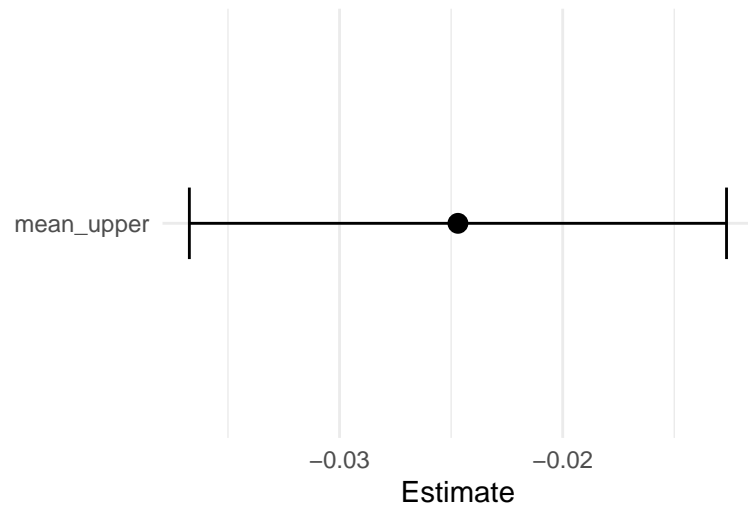
Full Coefficients – PIR



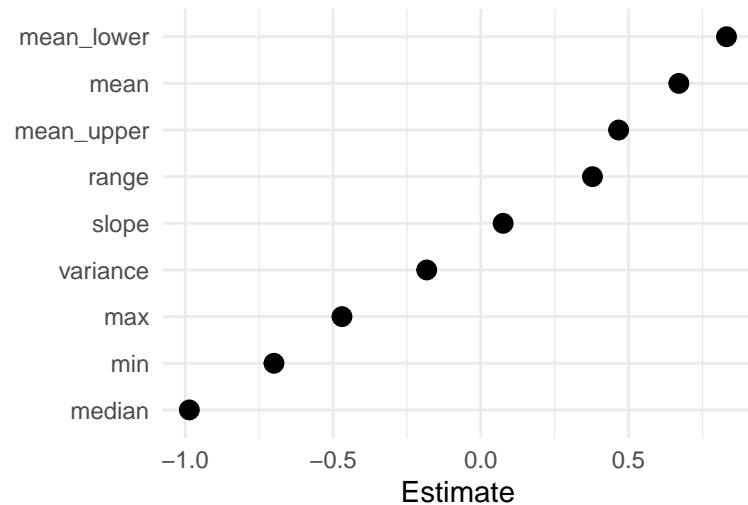
Range Coefficients – PIR

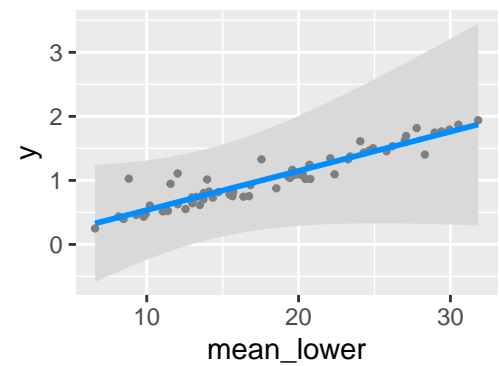
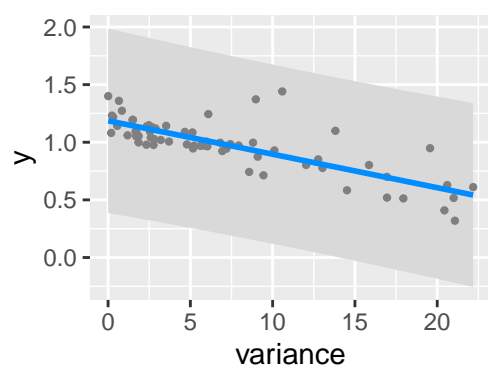
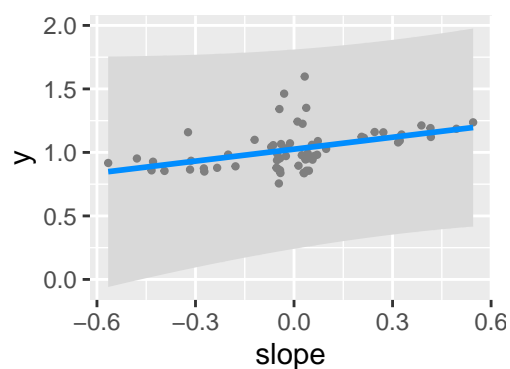
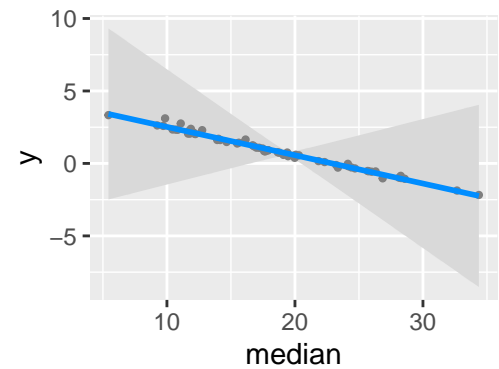
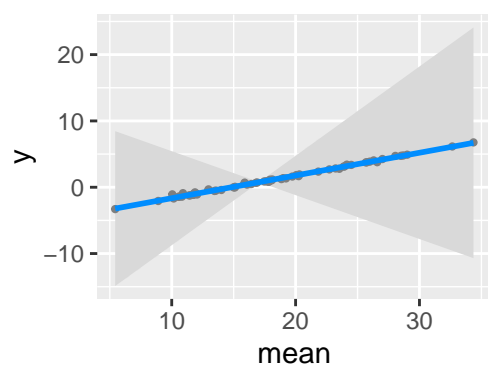
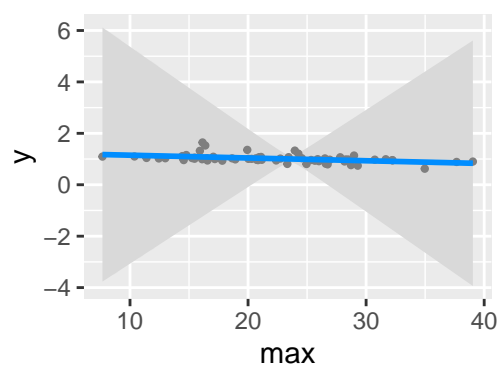
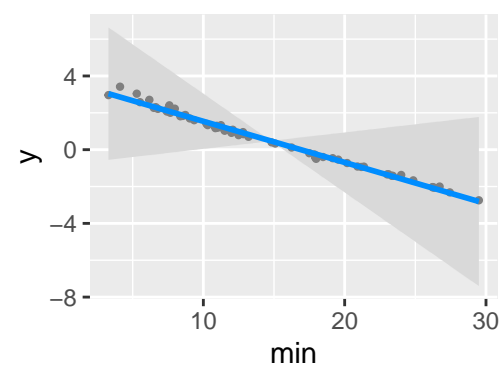


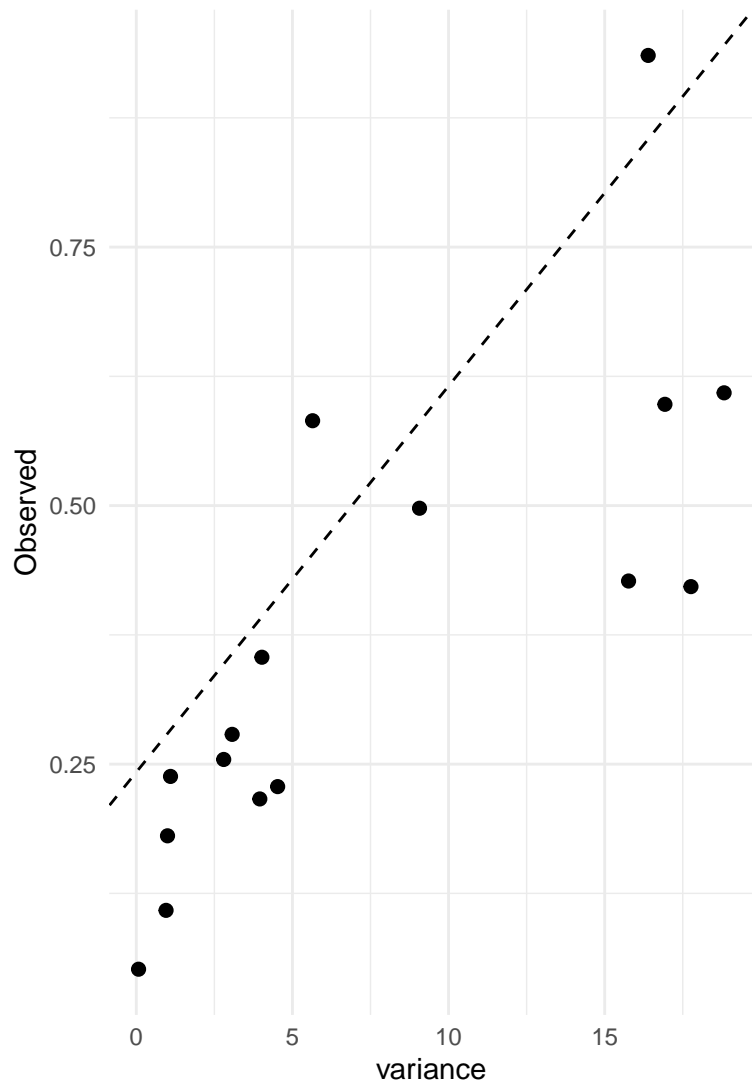
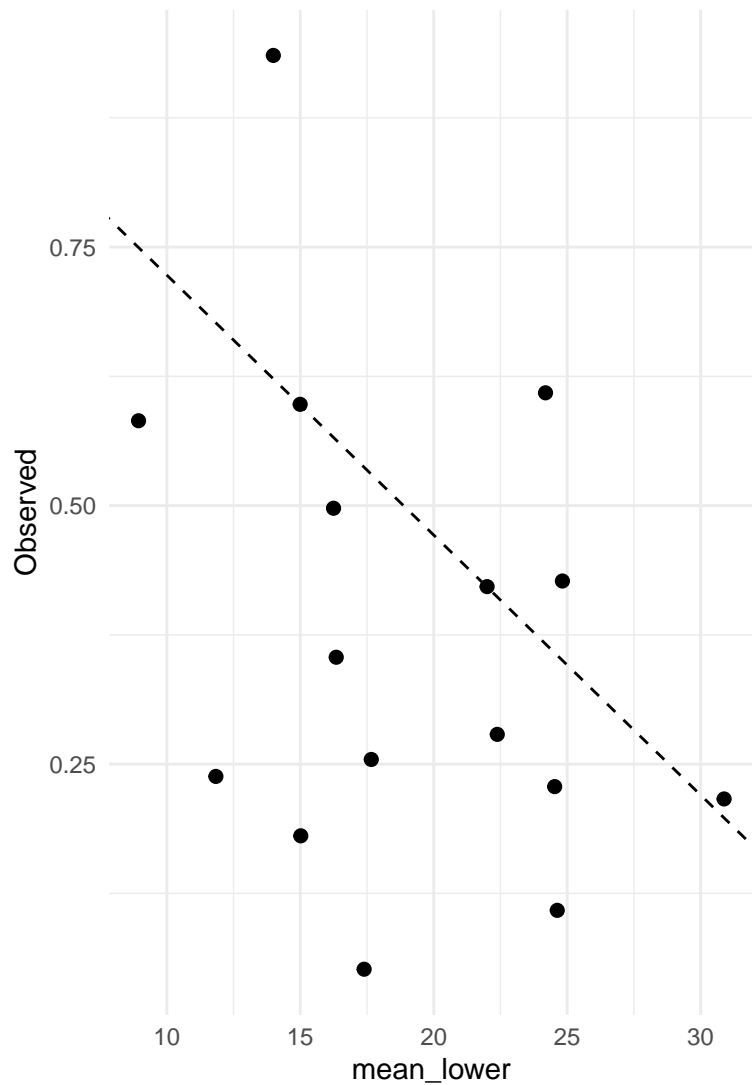
Mean_Upper Coefficients – PIR



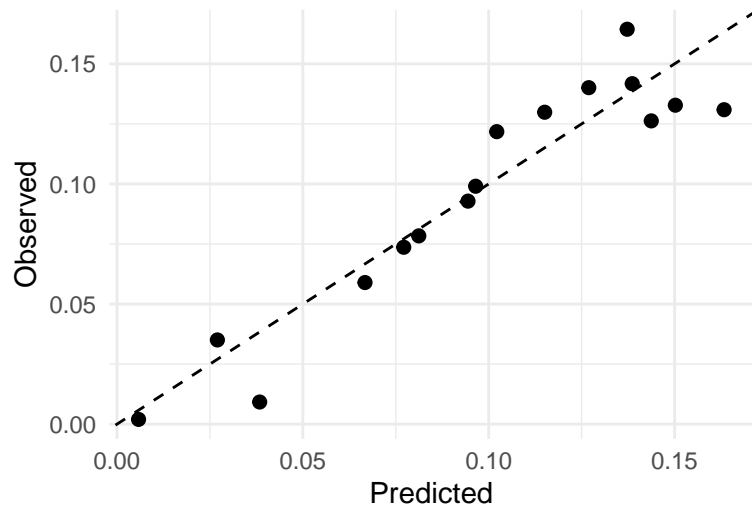
Ridge Coefficients – PIR



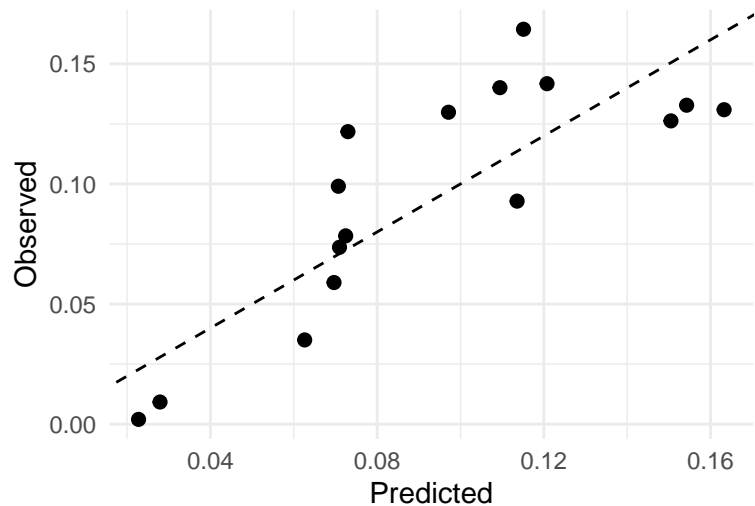


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{mean_lower}$ 

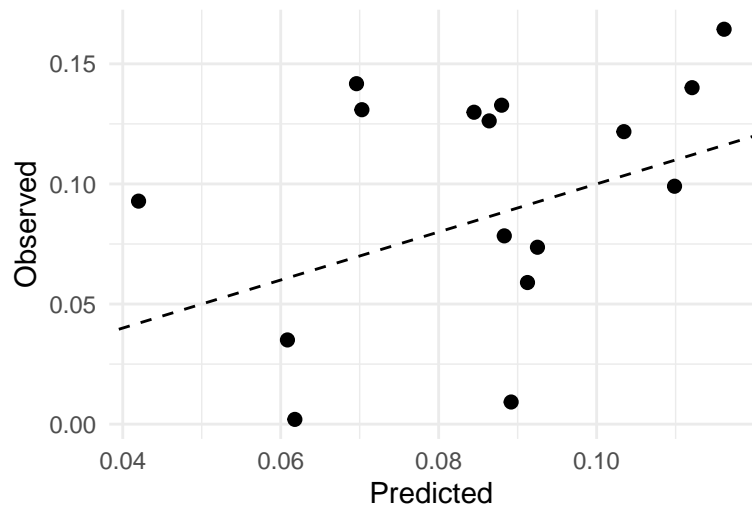
Full – RDPI



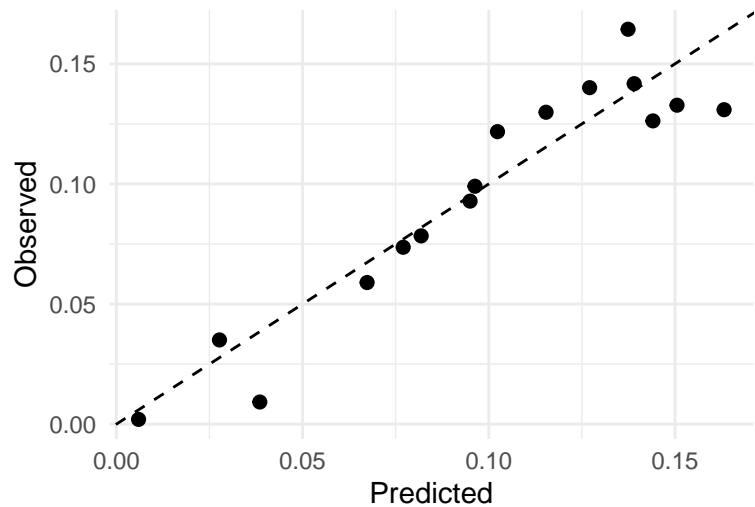
Range – RDPI



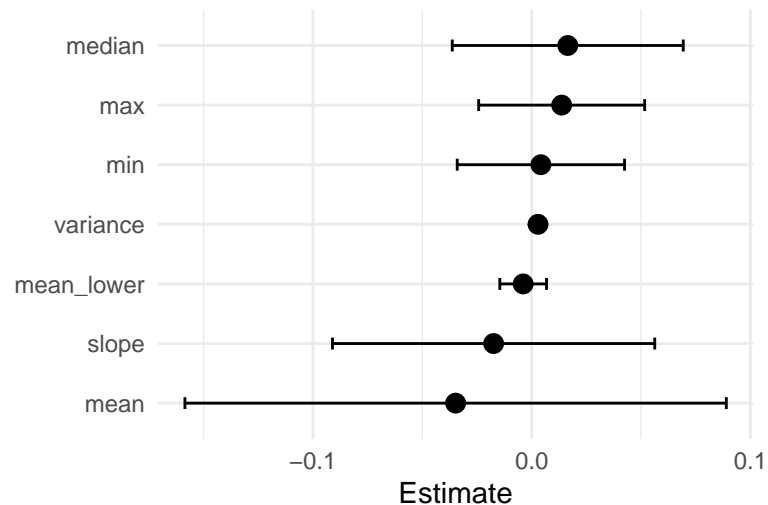
Mean_Upper – RDPI



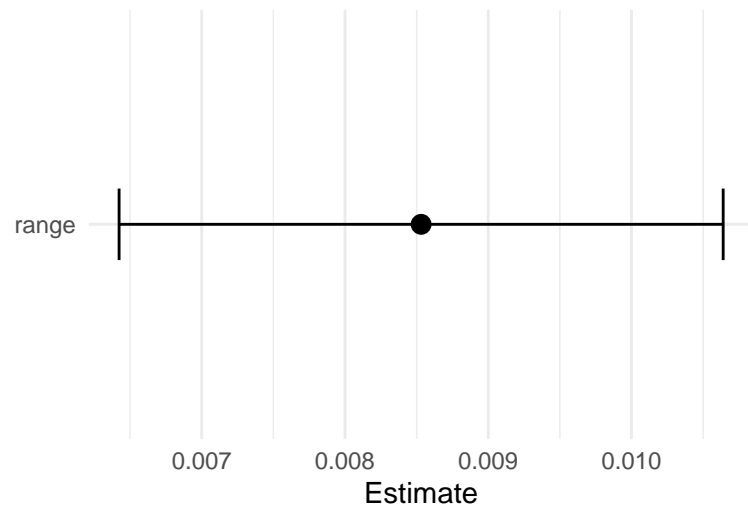
Ridge – RDPI



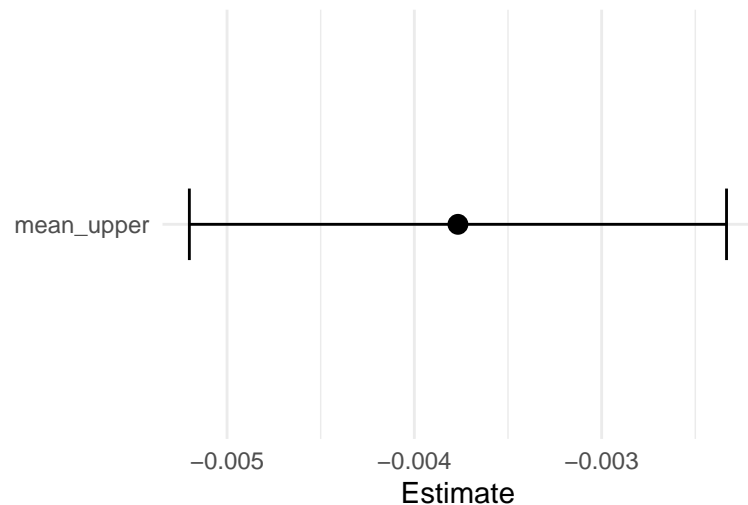
Full Coefficients – RDPI



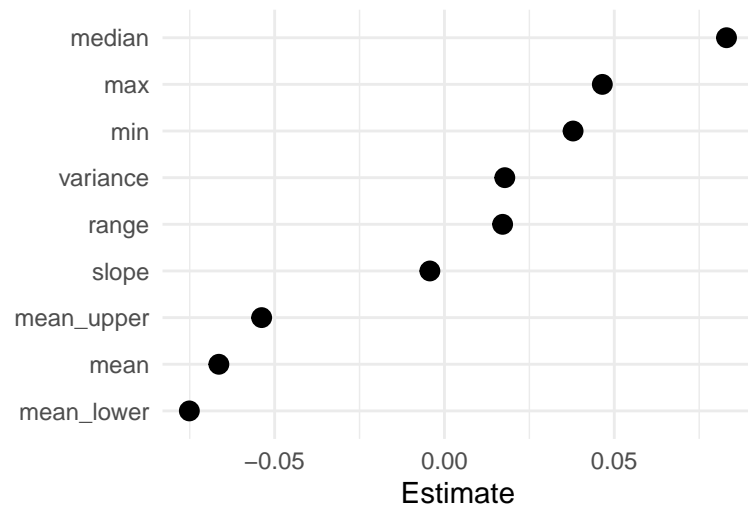
Range Coefficients – RDPI

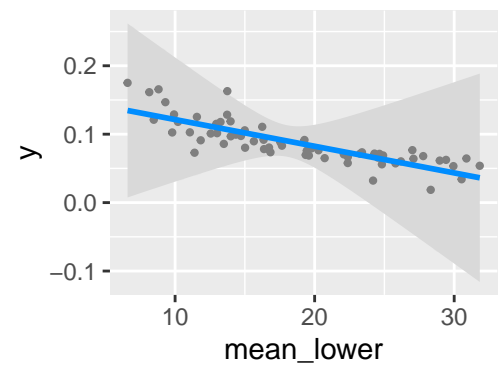
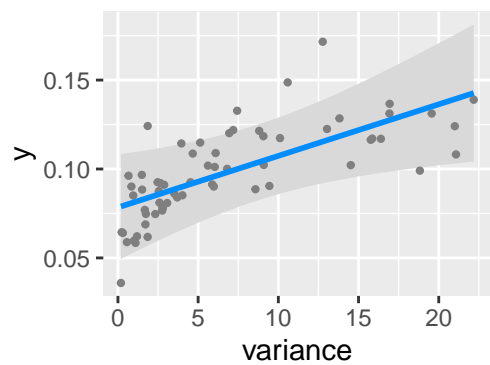
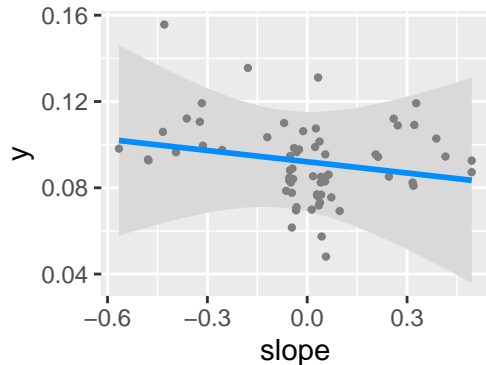
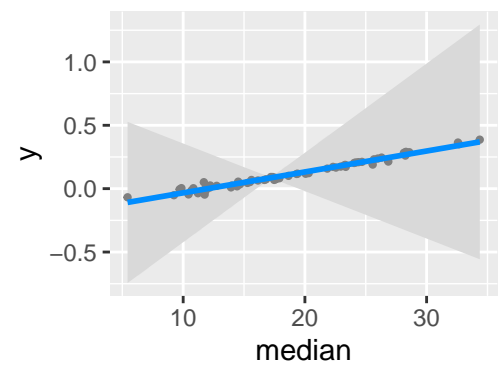
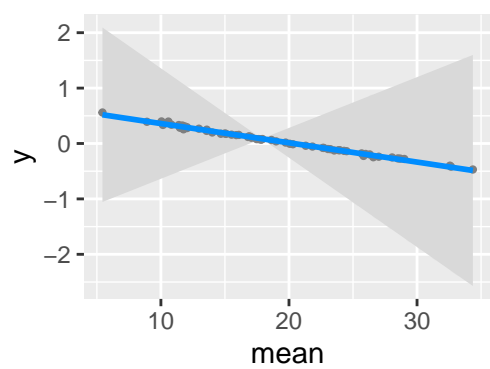
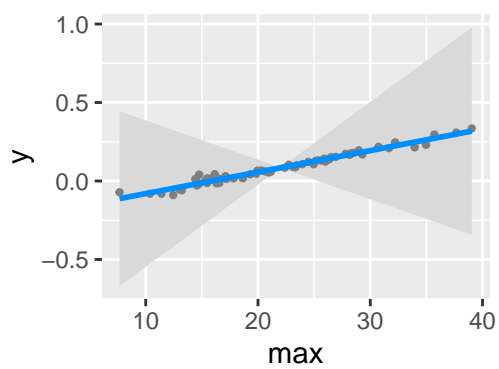
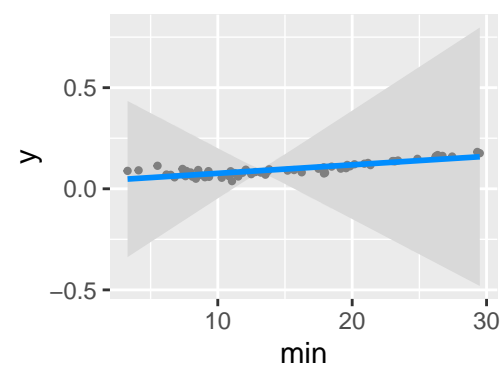


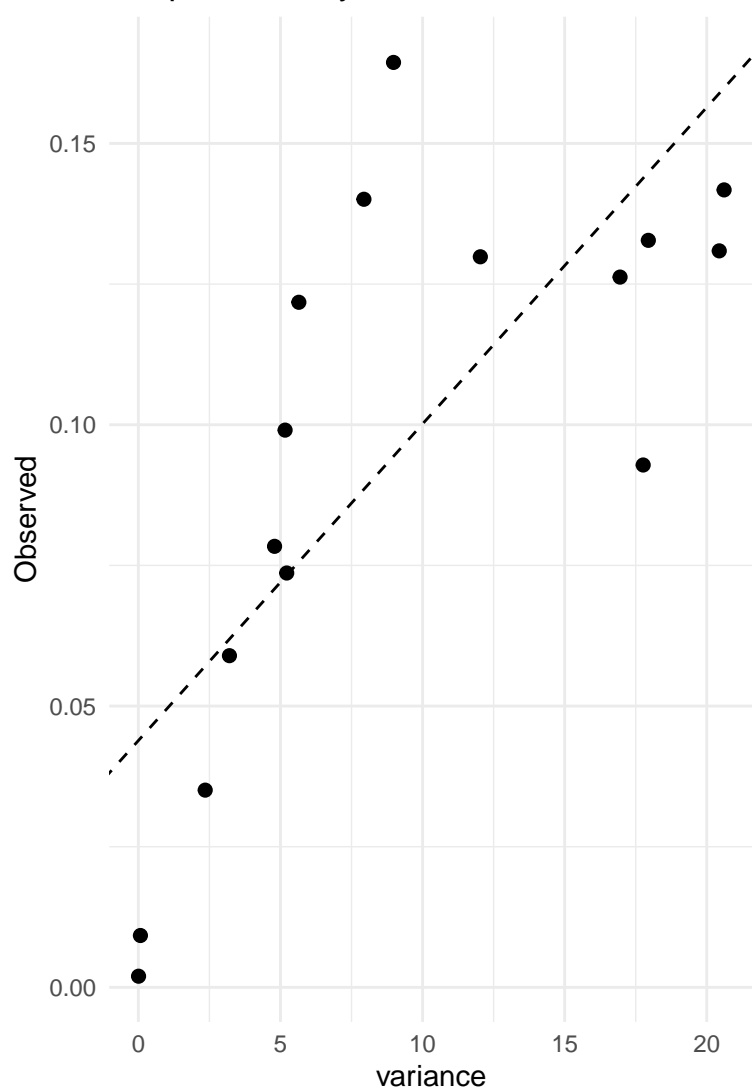
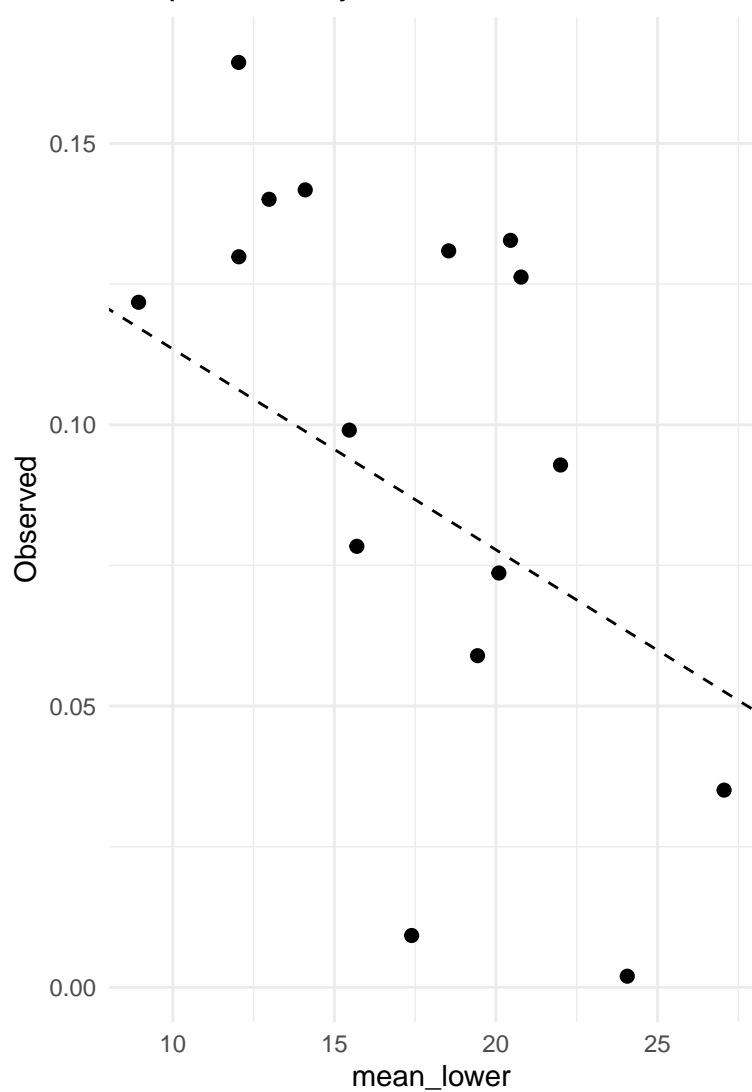
Mean_Upper Coefficients – RDPI



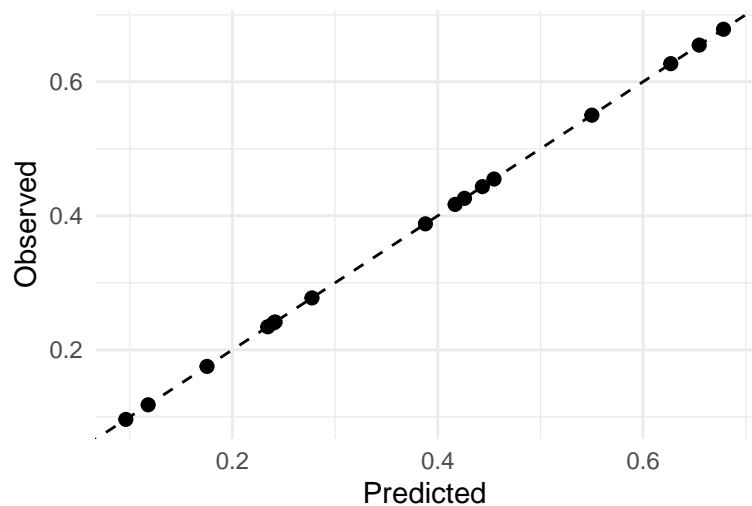
Ridge Coefficients – RDPI



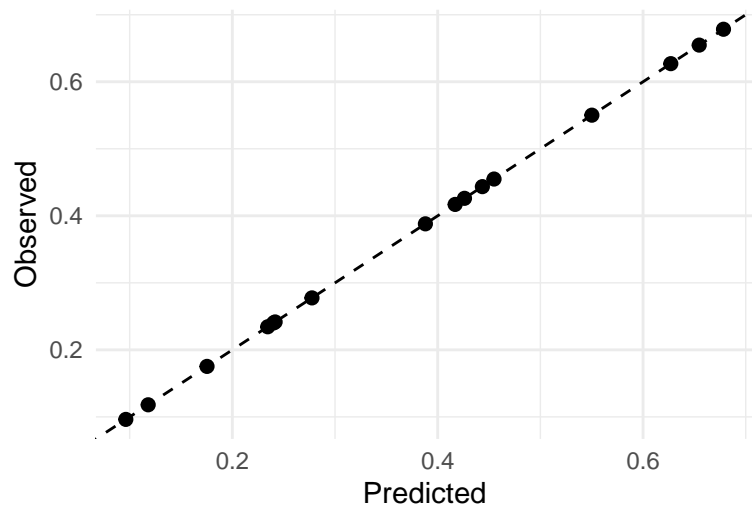


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{mean_lower}$ 

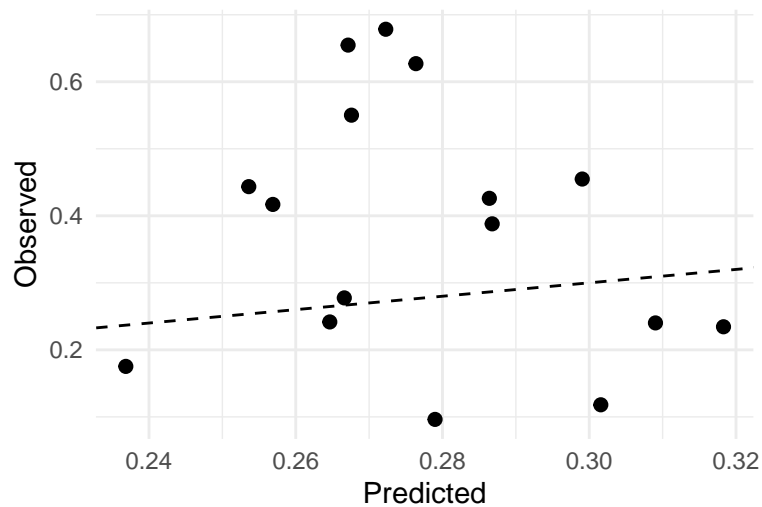
Full – ESPI



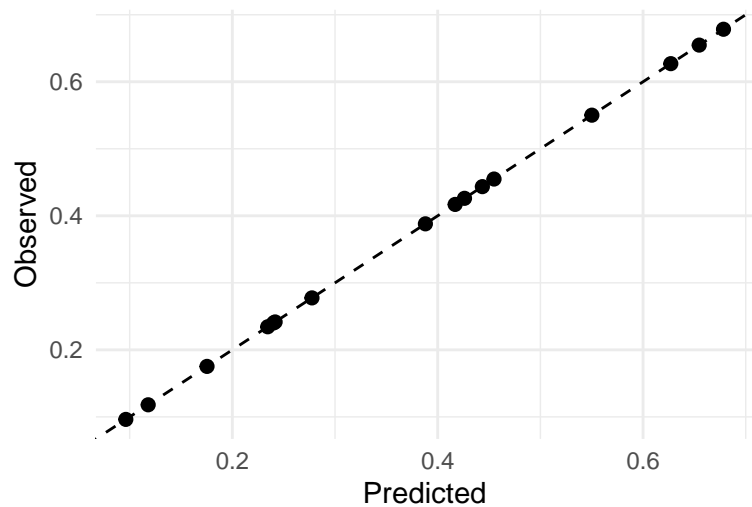
Range – ESPI



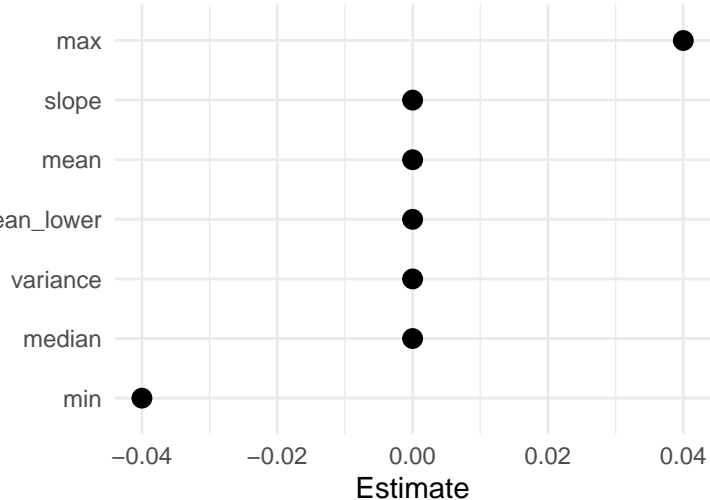
Mean_Upper – ESPI



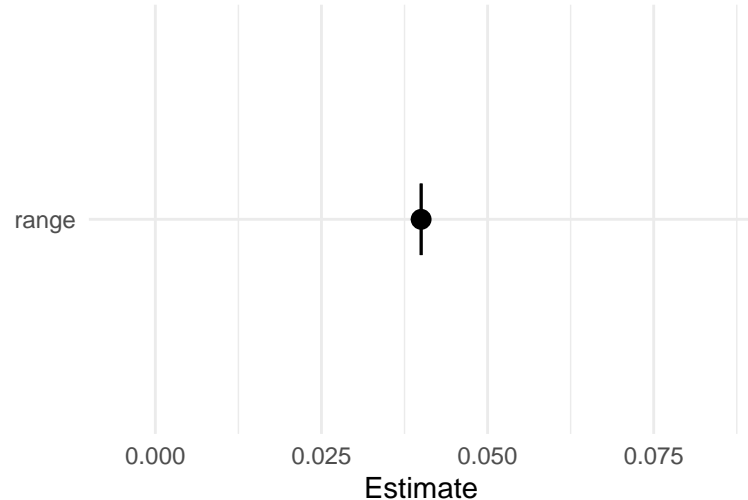
Ridge – ESPI



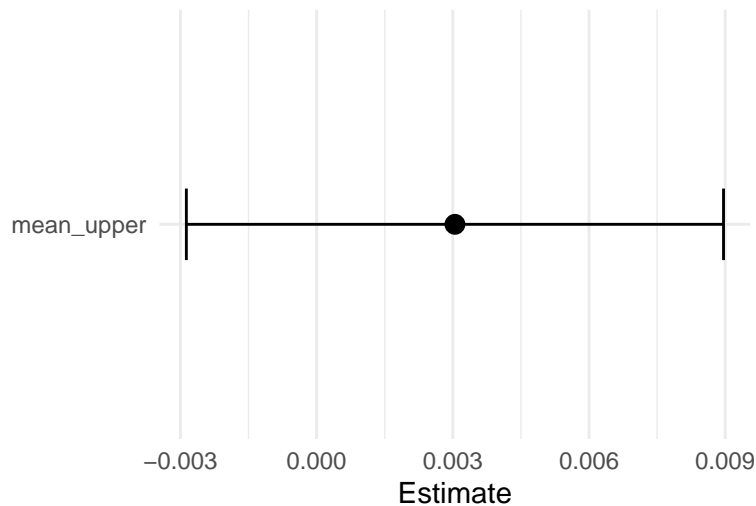
Full Coefficients – ESPI



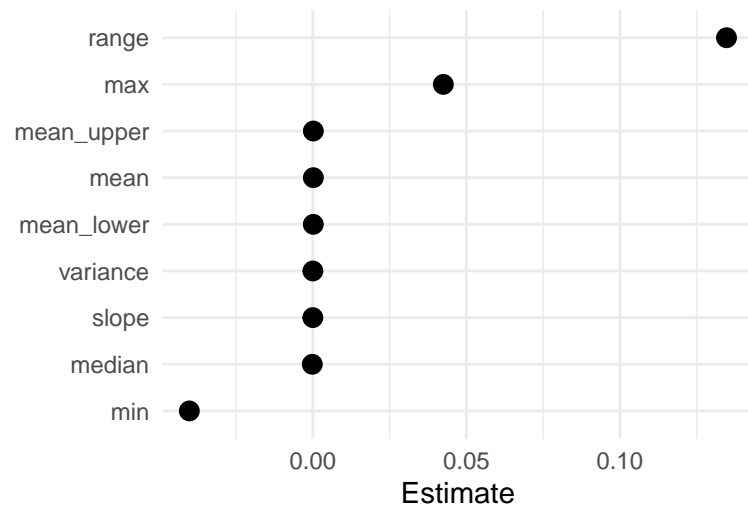
Range Coefficients – ESPI

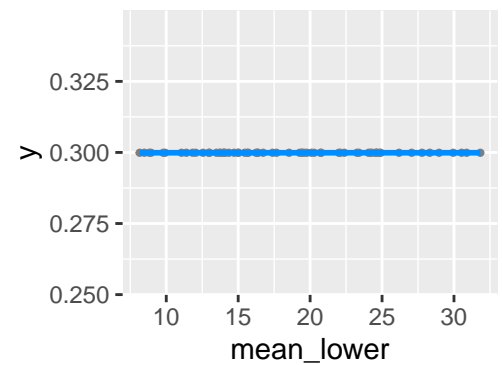
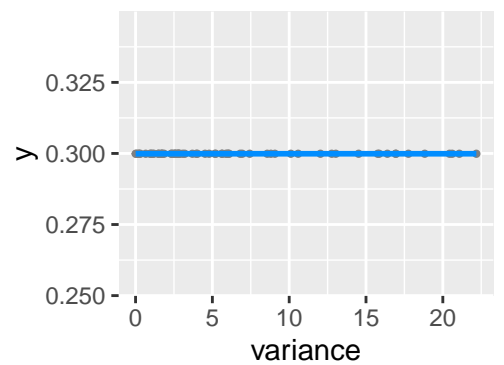
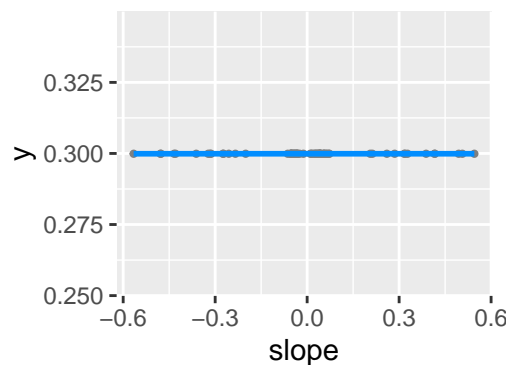
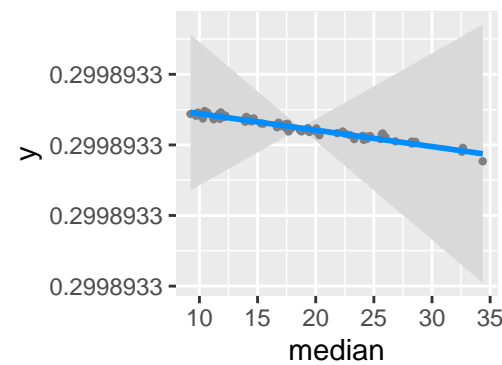
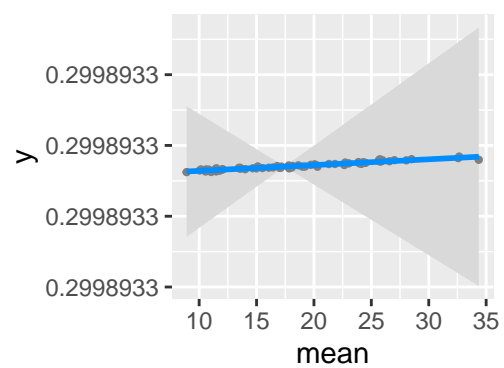
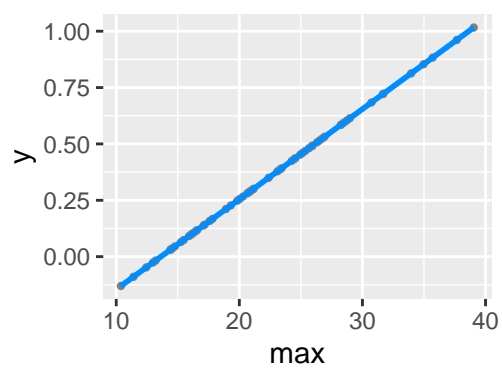
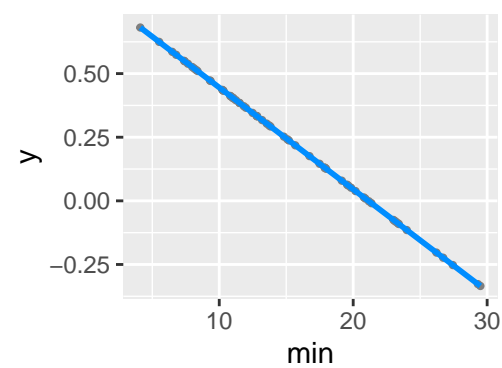


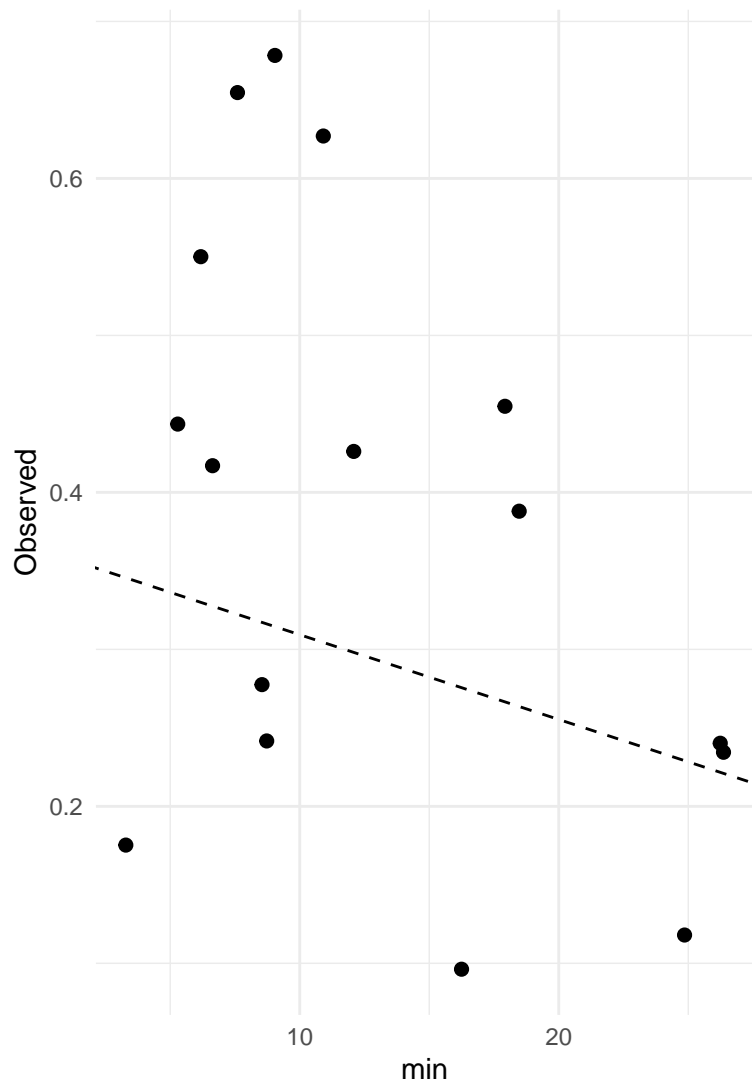
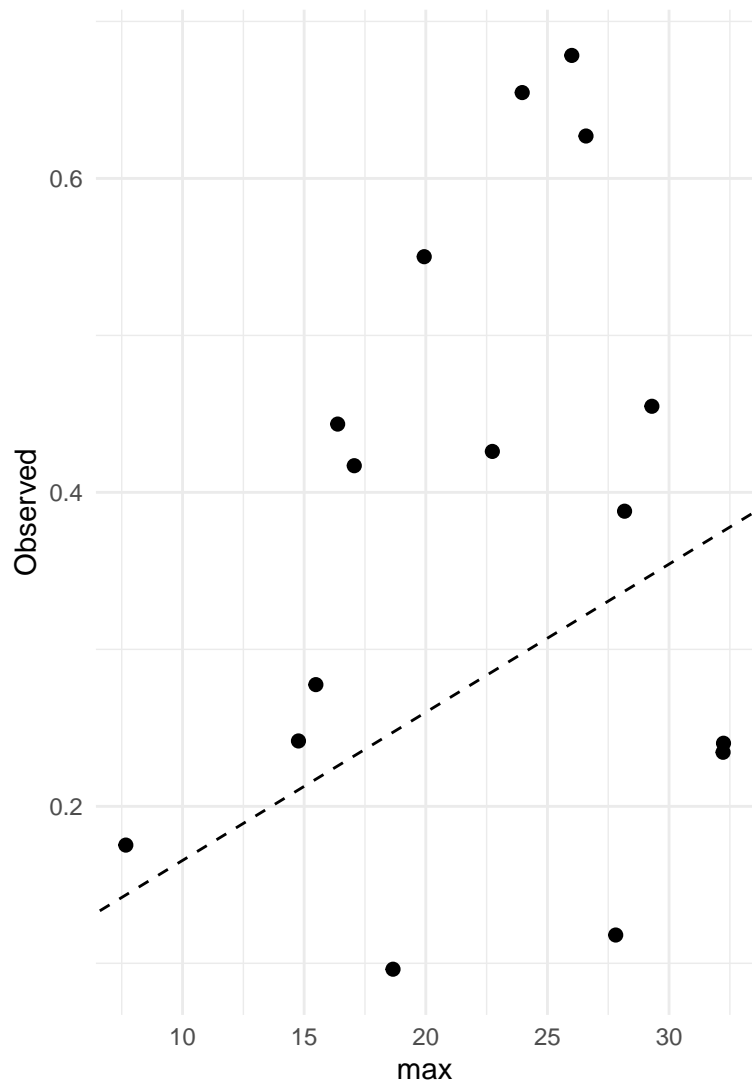
Mean_Upper Coefficients – ESPI



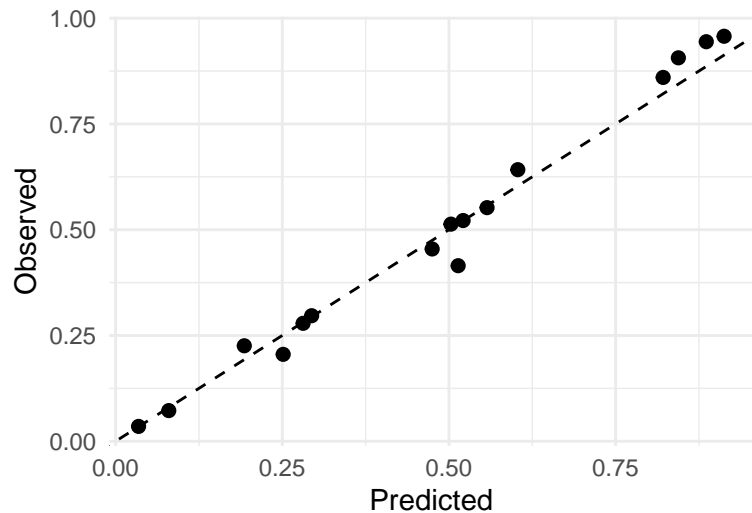
Ridge Coefficients – ESPI



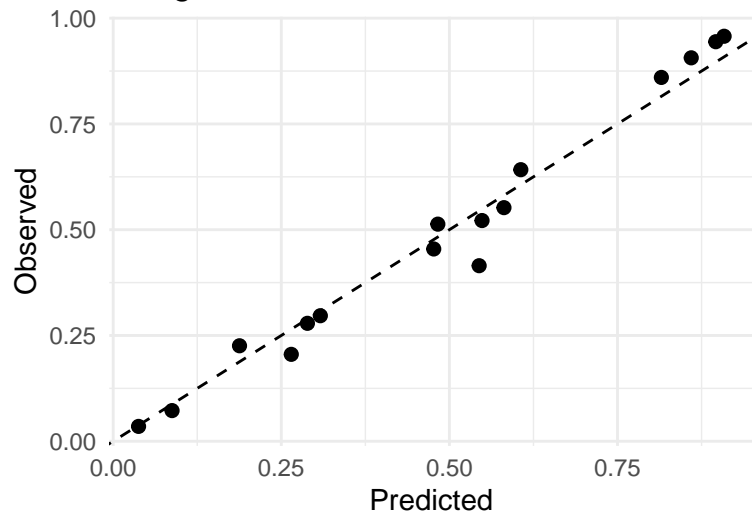


Simple Model: $y \sim \min$ Simple Model: $y \sim \max$ 

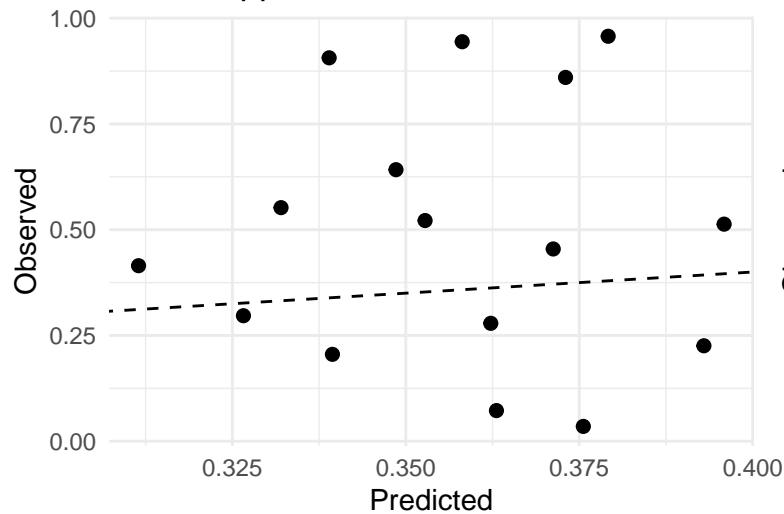
Full – ESPIID



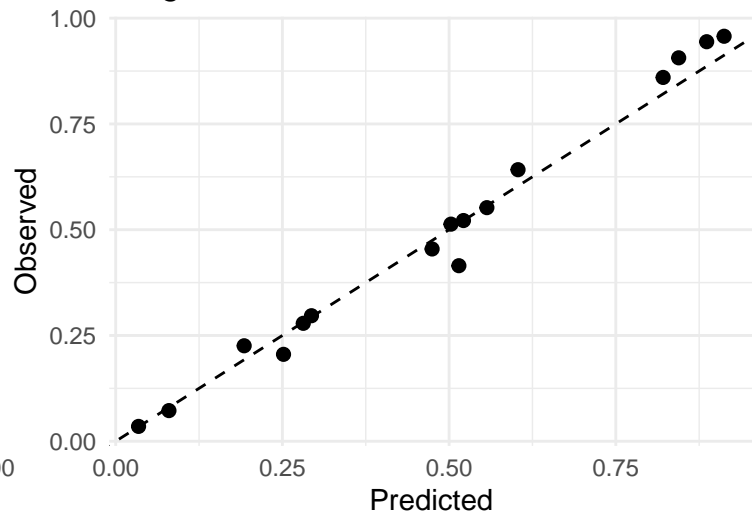
Range – ESPIID



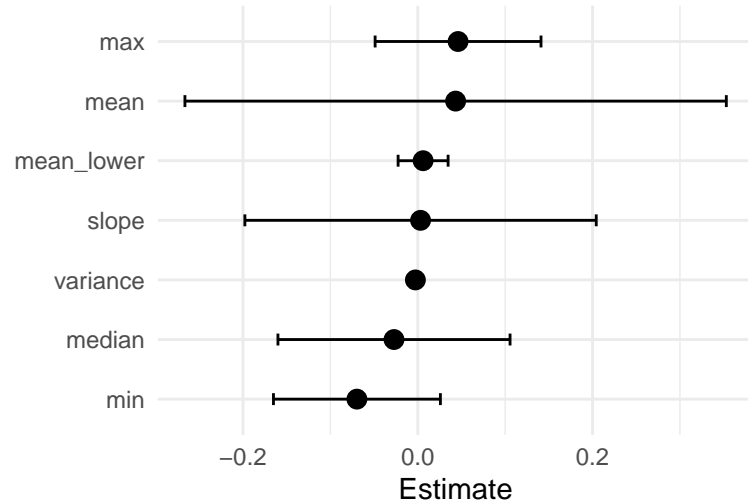
Mean_Upper – ESPIID



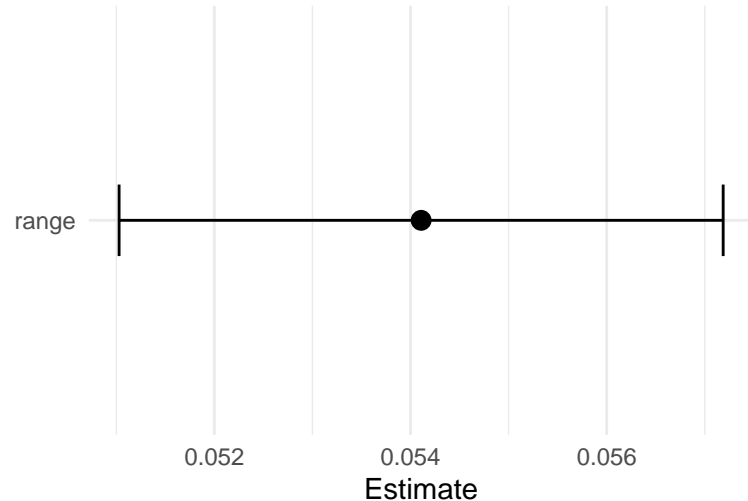
Ridge – ESPIID



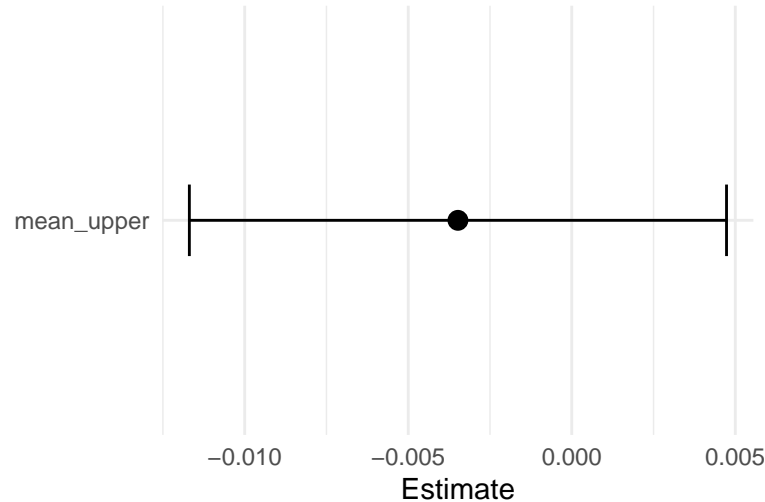
Full Coefficients – ESPIID



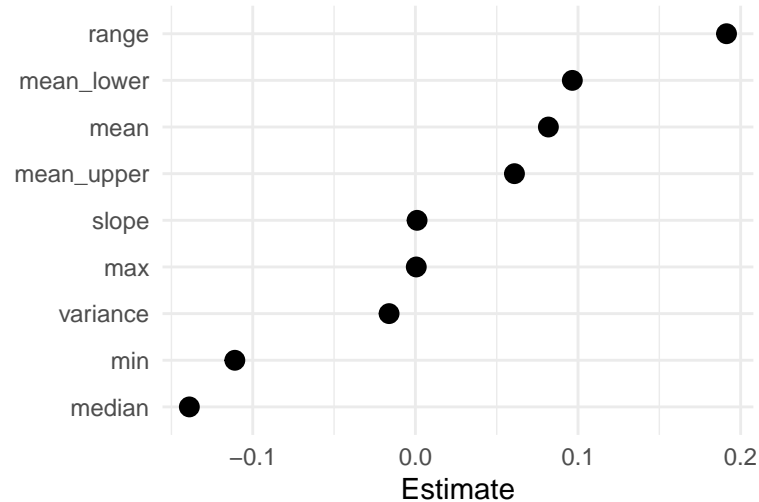
Range Coefficients – ESPIID

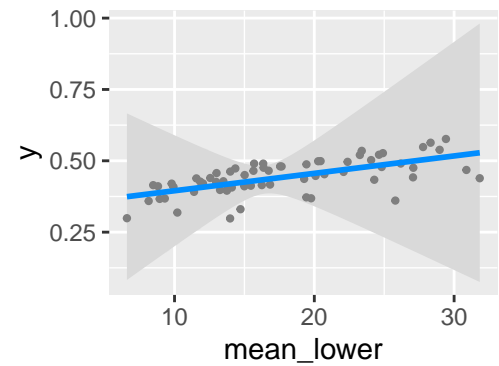
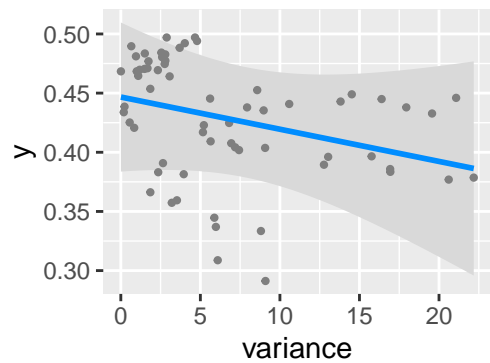
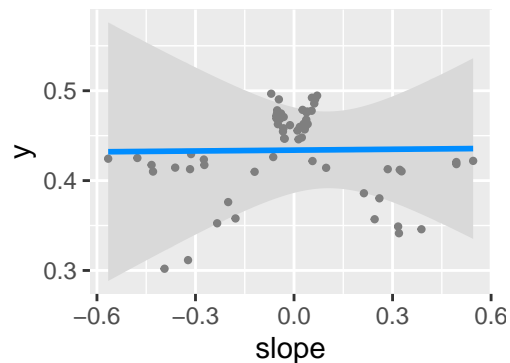
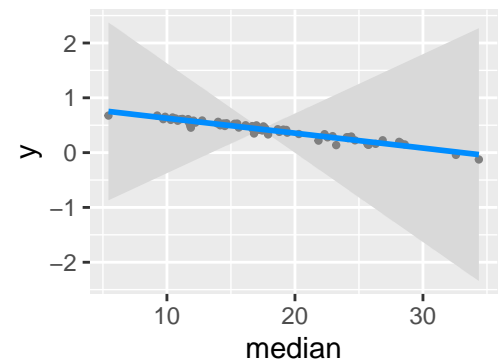
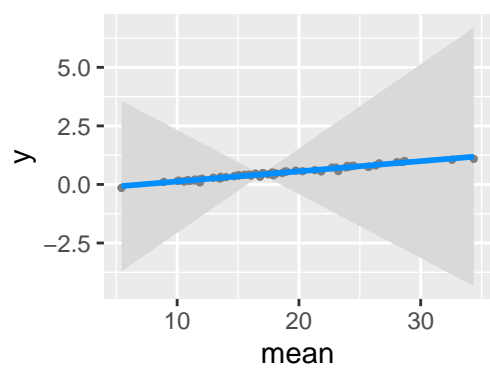
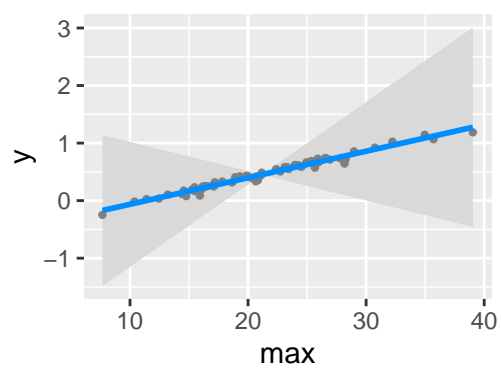
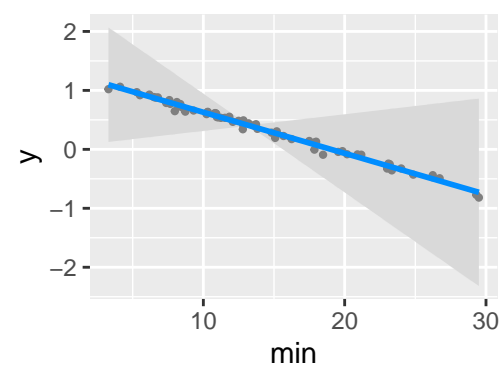


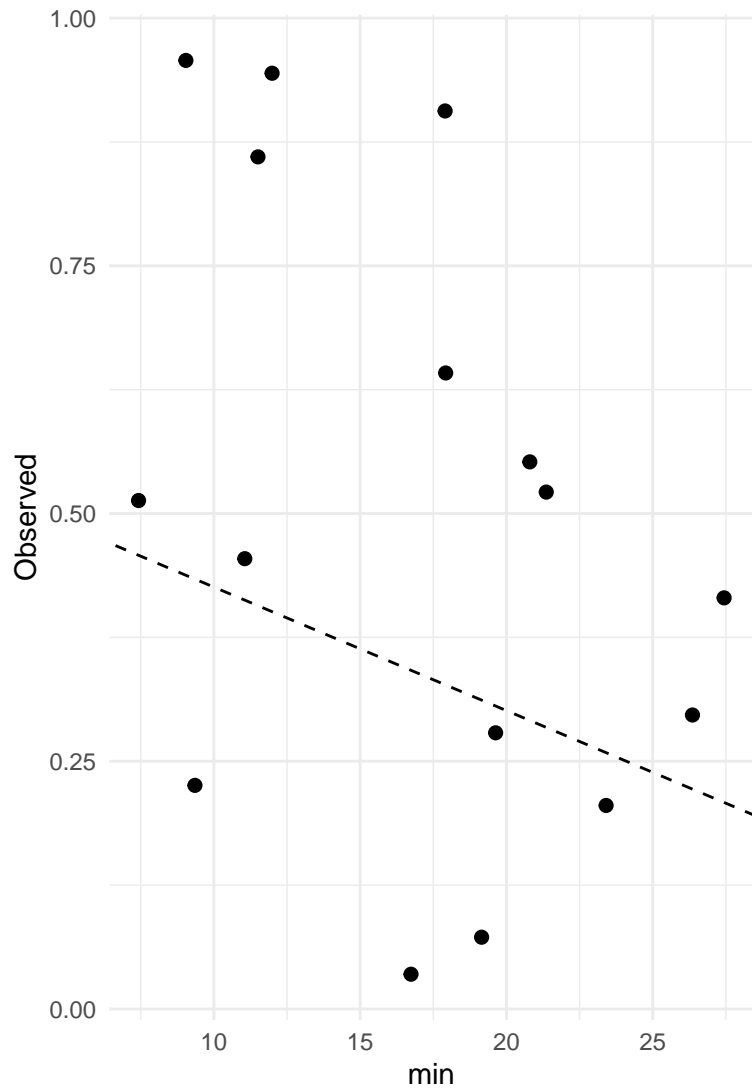
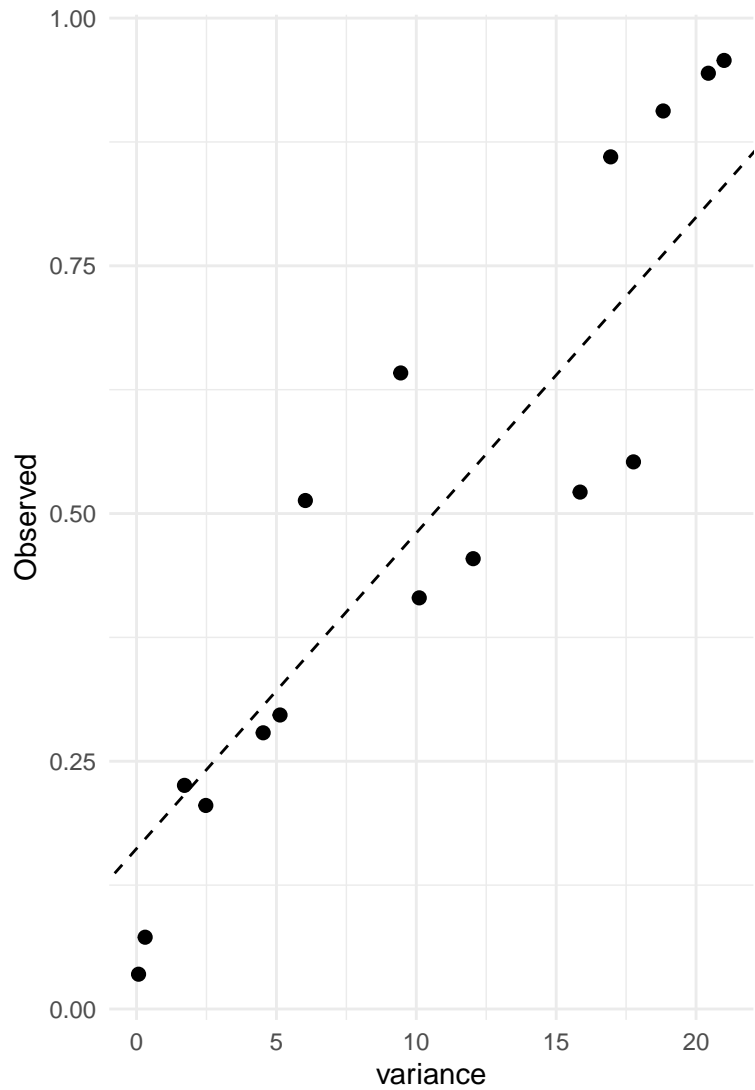
Mean_Upper Coefficients – ESPIID



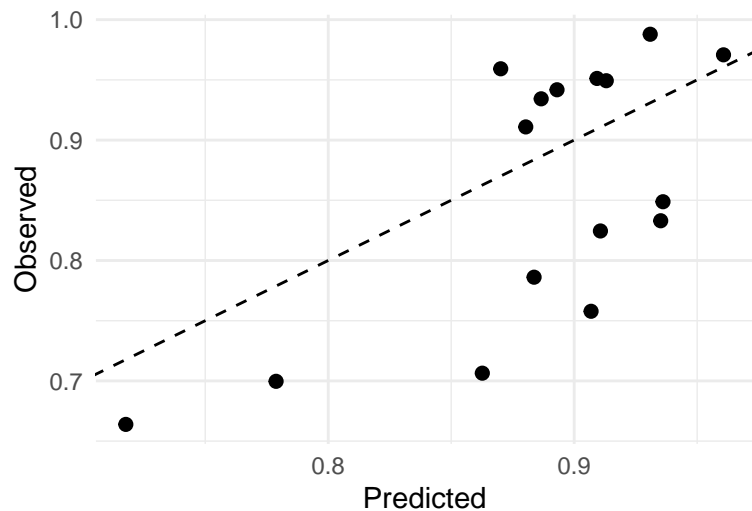
Ridge Coefficients – ESPIID



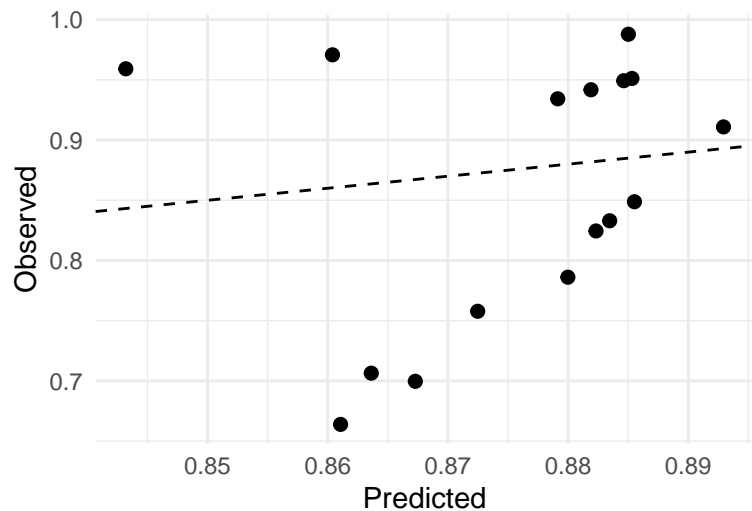


Simple Model: $y \sim \text{min}$ Simple Model: $y \sim \text{variance}$ 

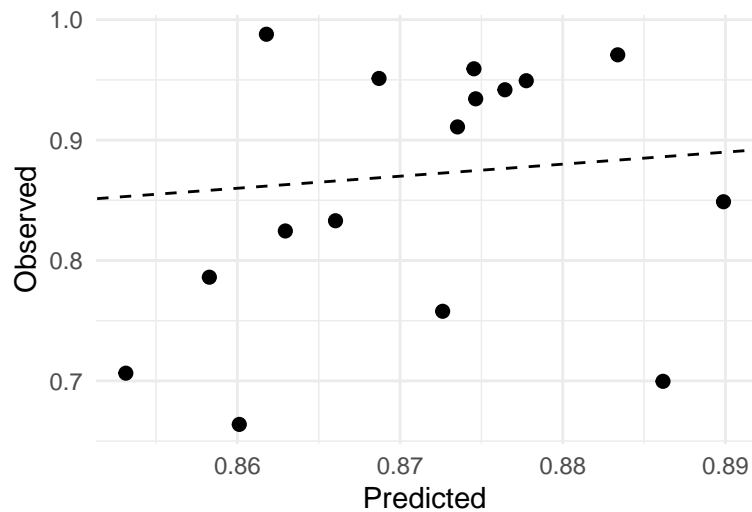
Full – PSI



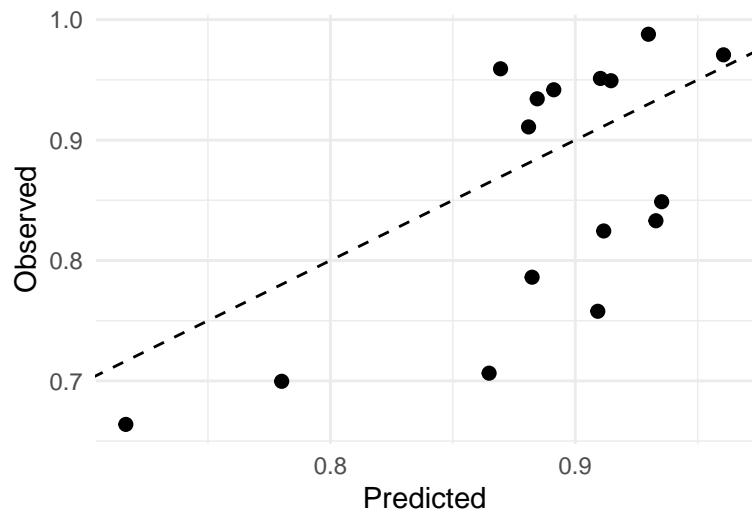
Range – PSI



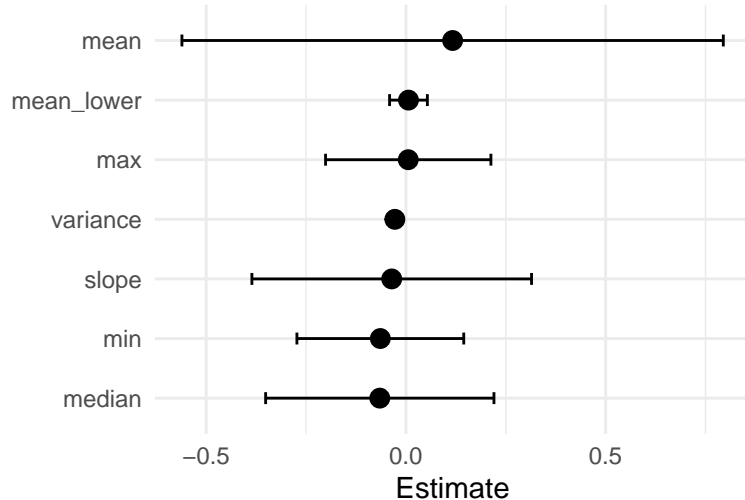
Mean_Upper – PSI



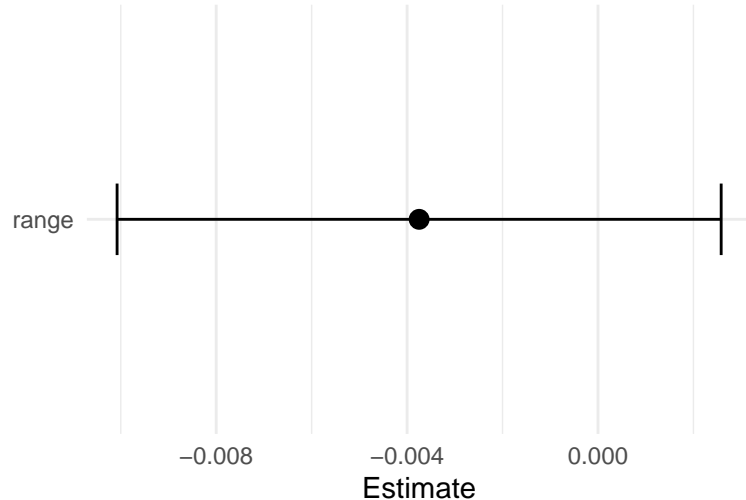
Ridge – PSI



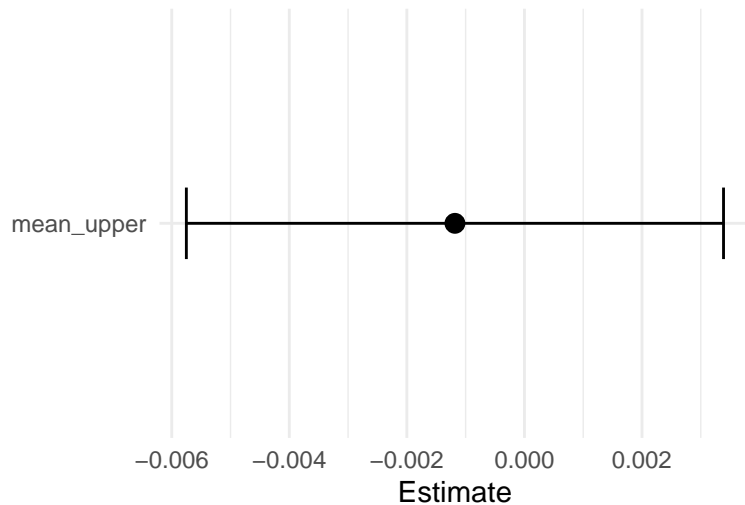
Full Coefficients – PSI



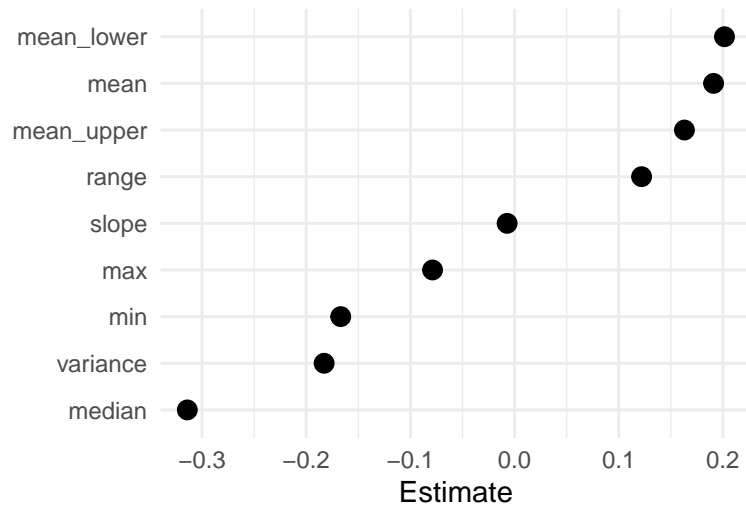
Range Coefficients – PSI

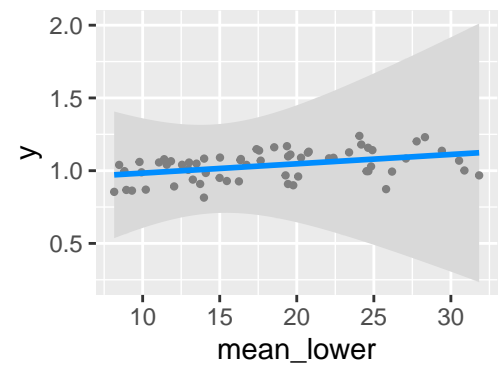
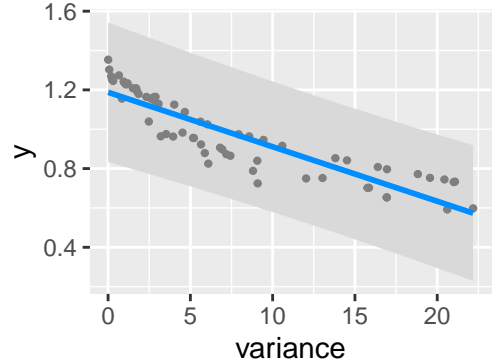
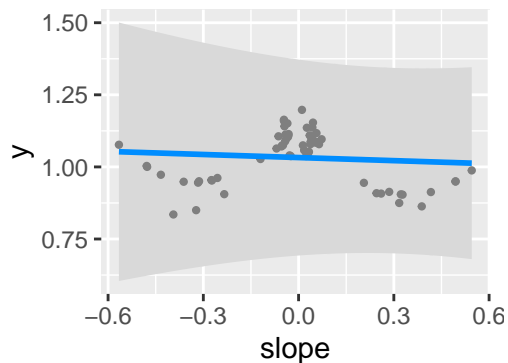
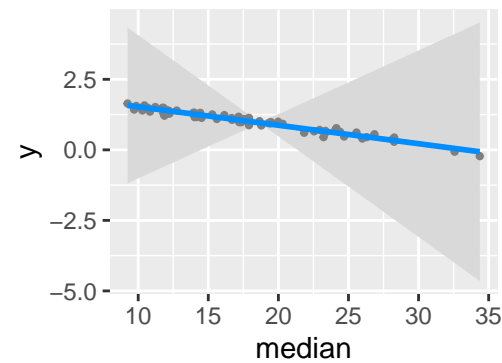
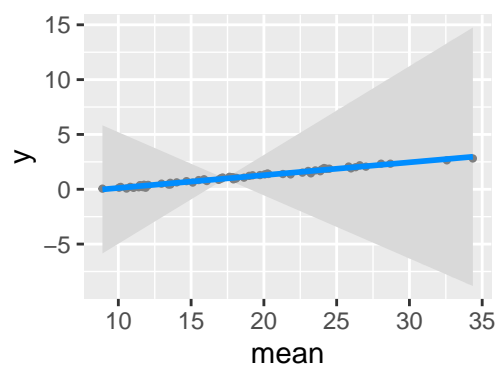
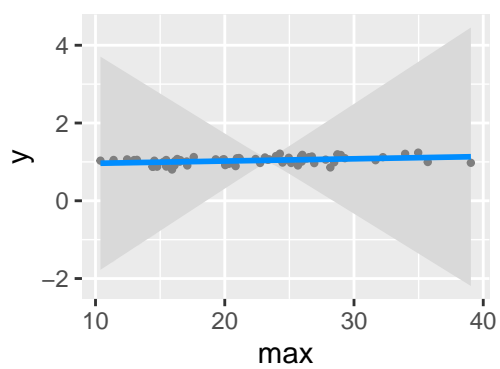
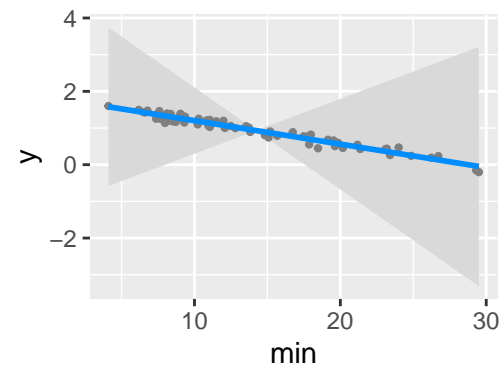


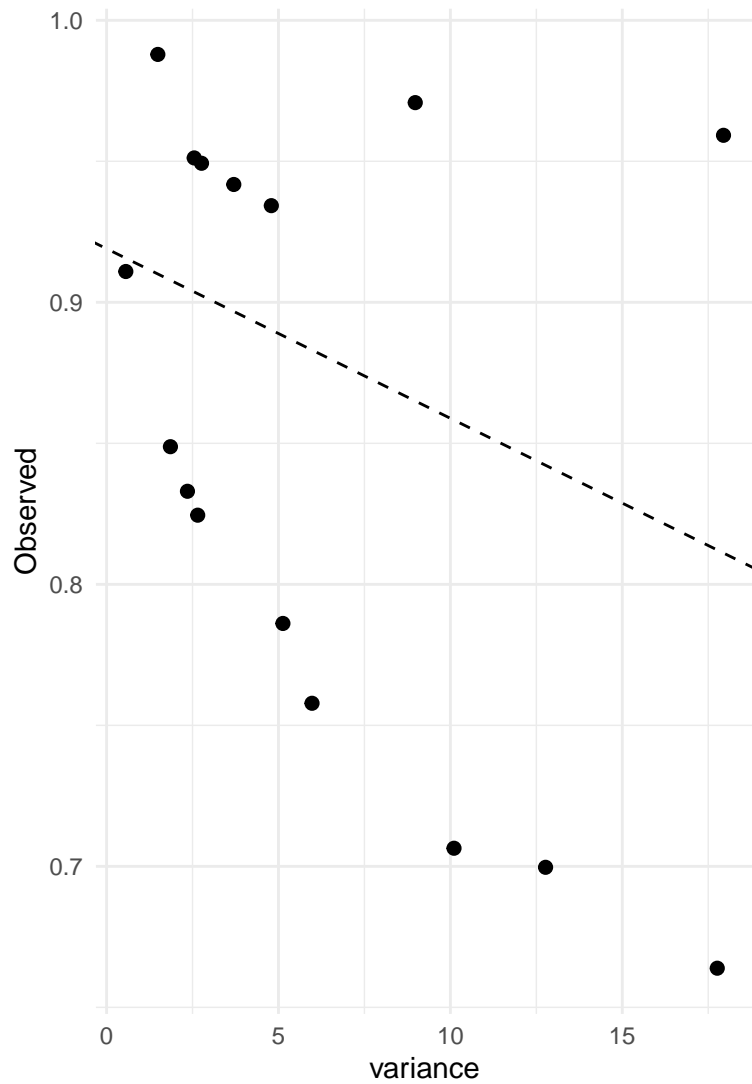
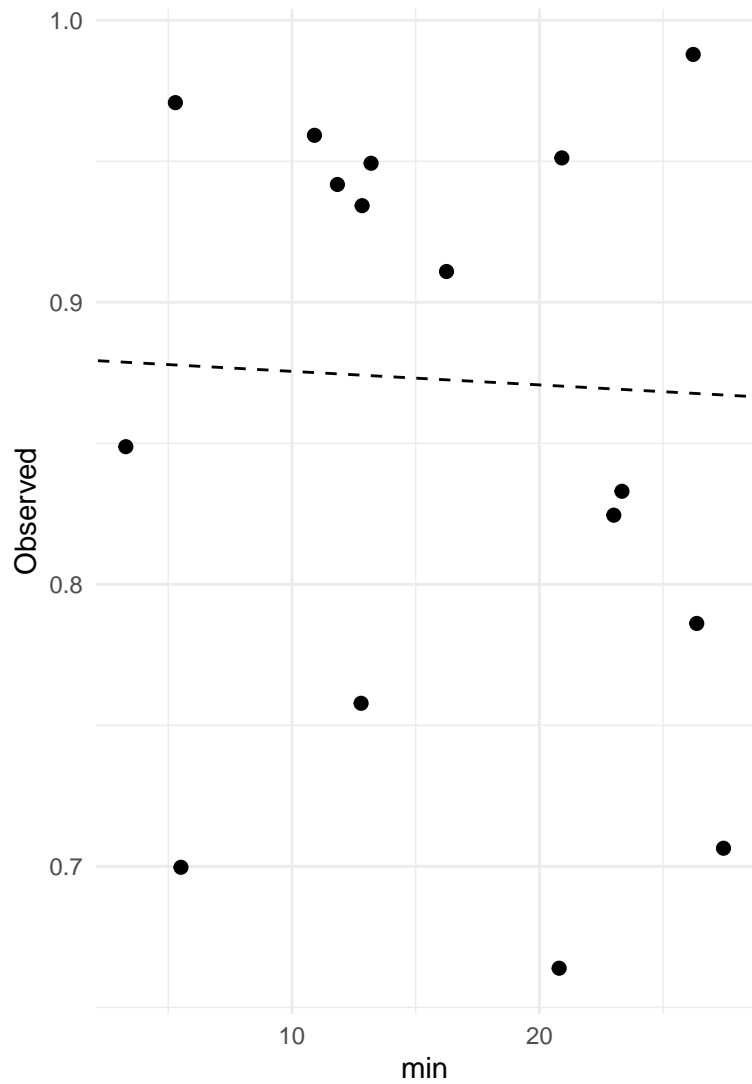
Mean_Upper Coefficients – PSI



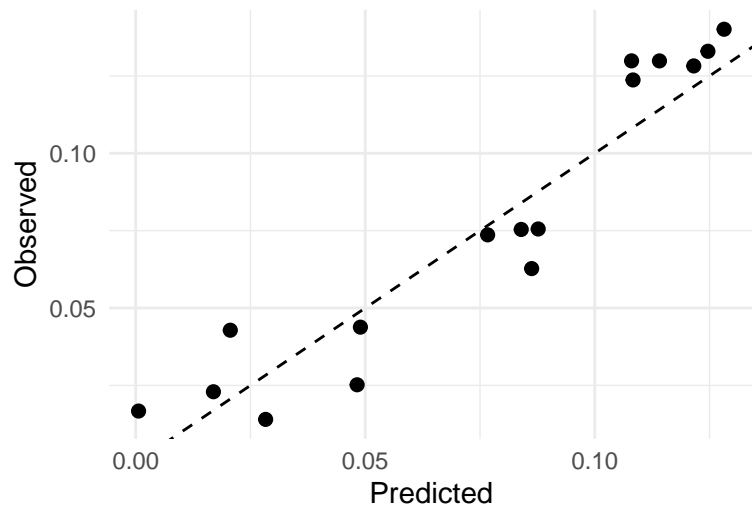
Ridge Coefficients – PSI



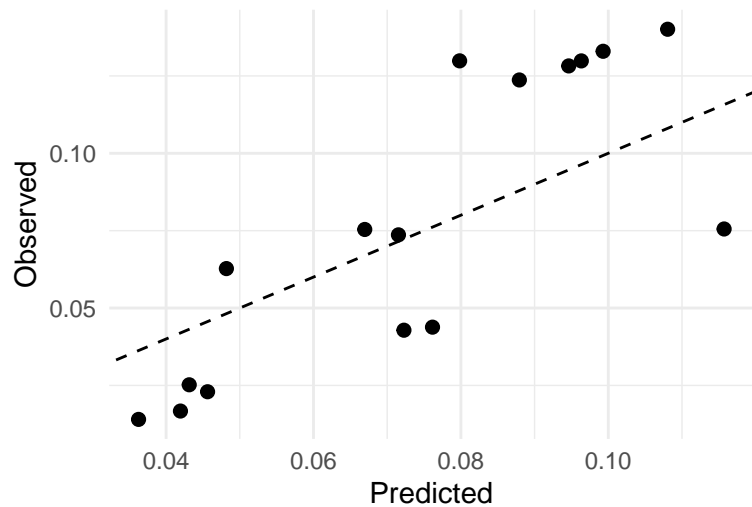


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{min}$ 

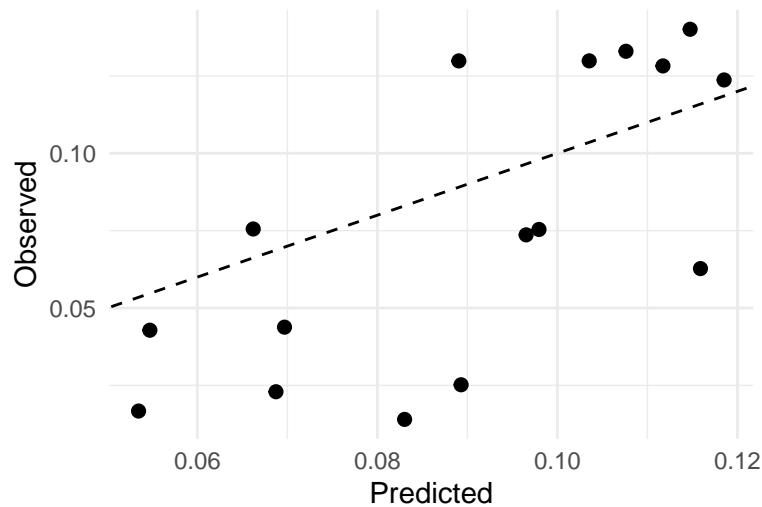
Full – RPI



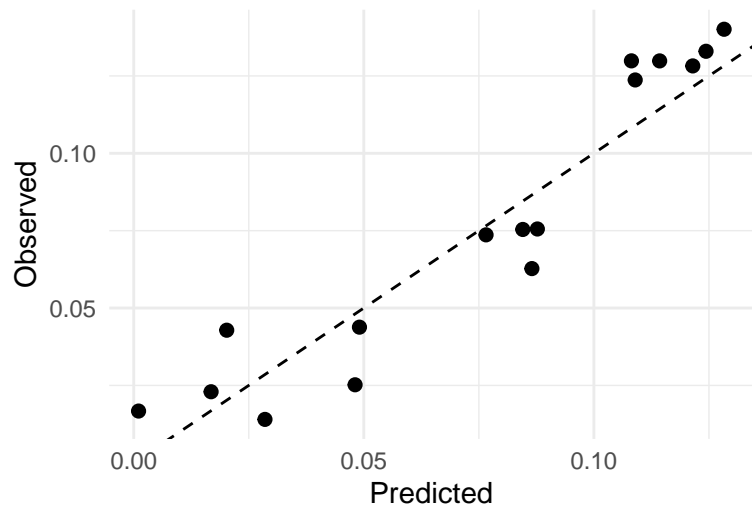
Range – RPI



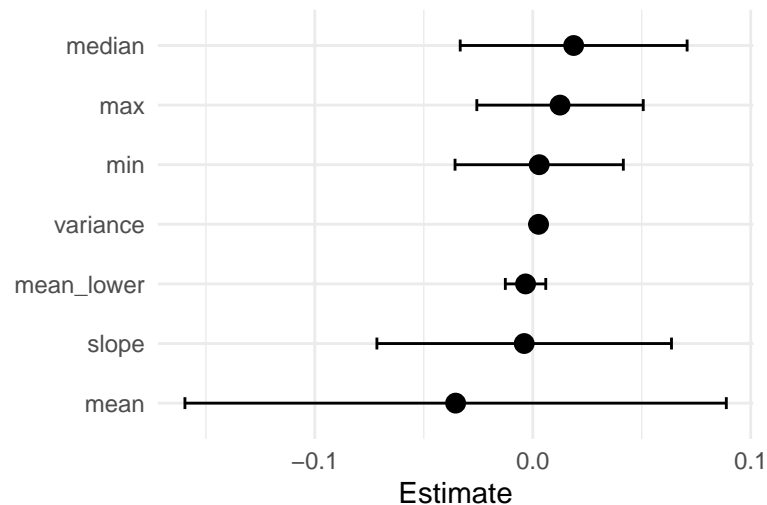
Mean_Upper – RPI



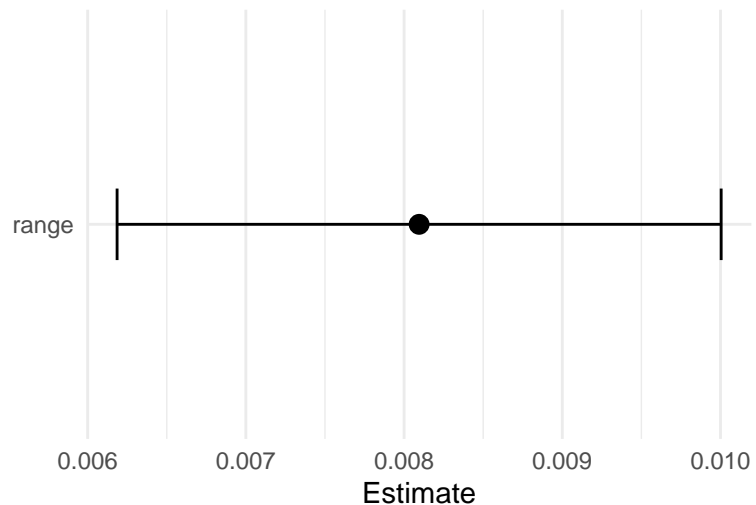
Ridge – RPI



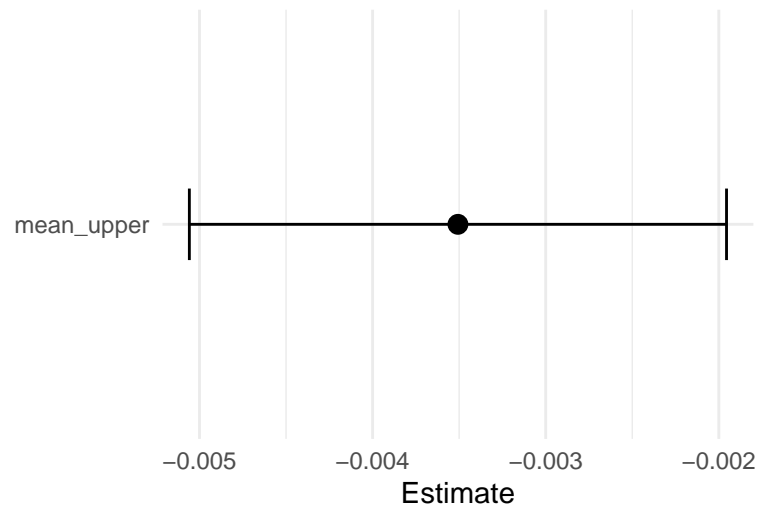
Full Coefficients – RPI



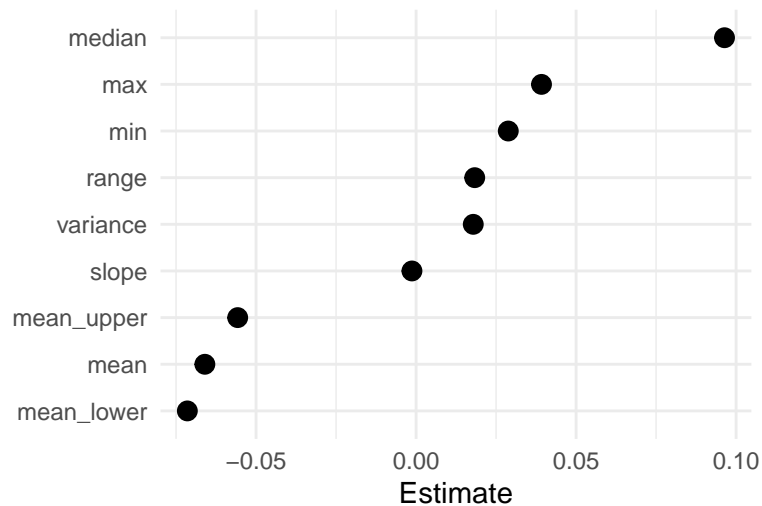
Range Coefficients – RPI

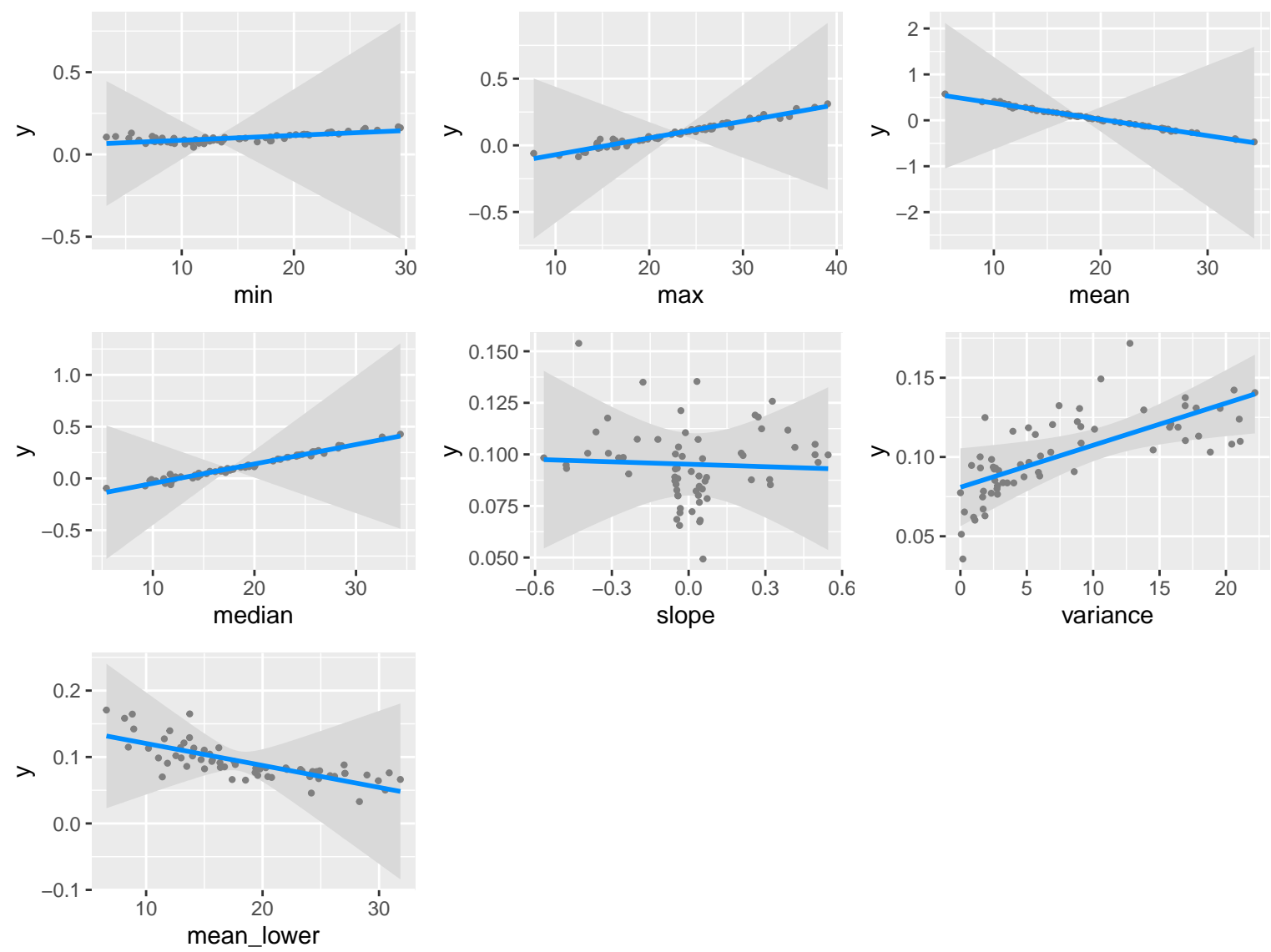


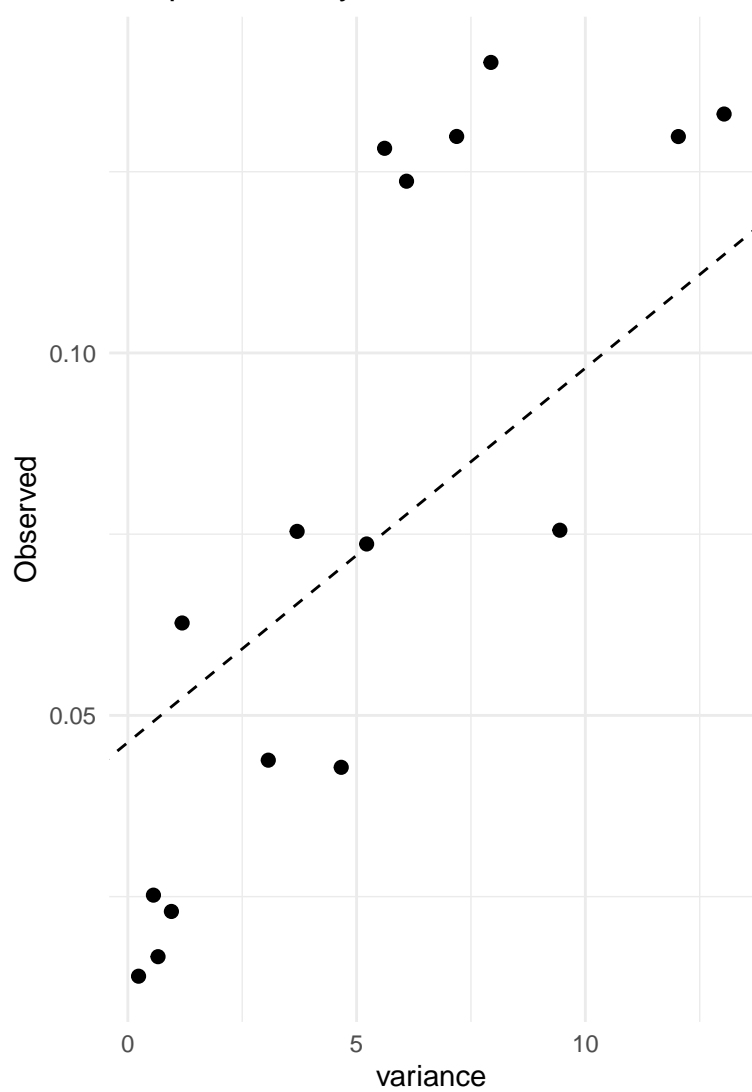
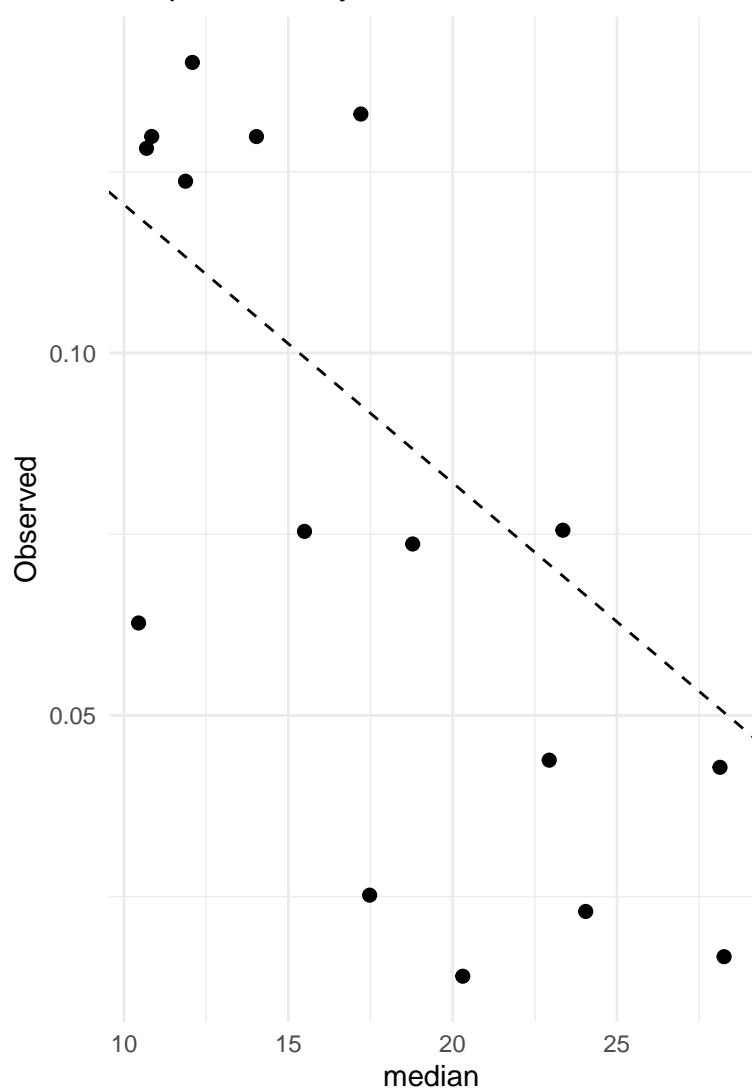
Mean_Upper Coefficients – RPI



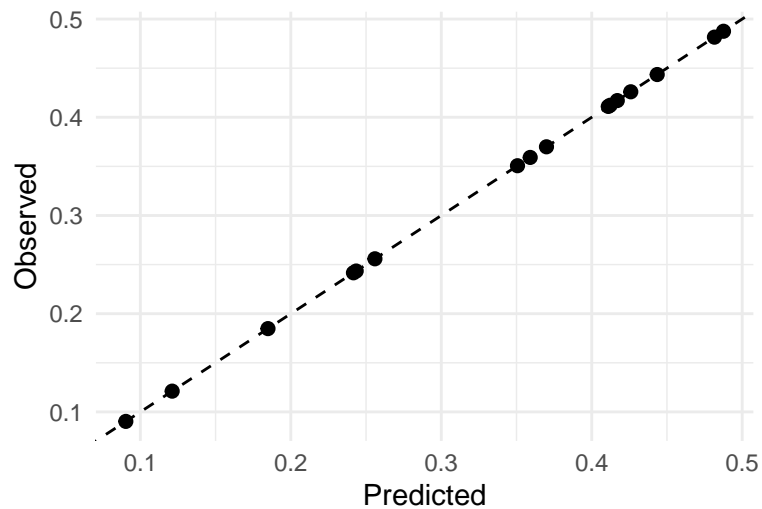
Ridge Coefficients – RPI



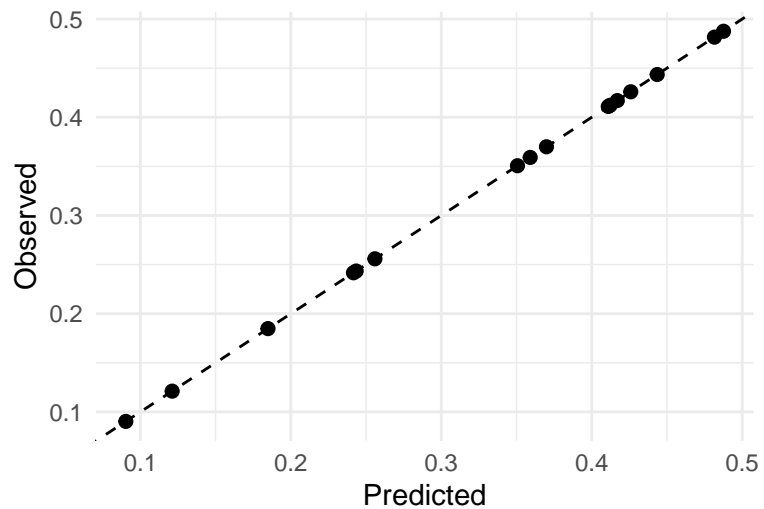


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{median}$ 

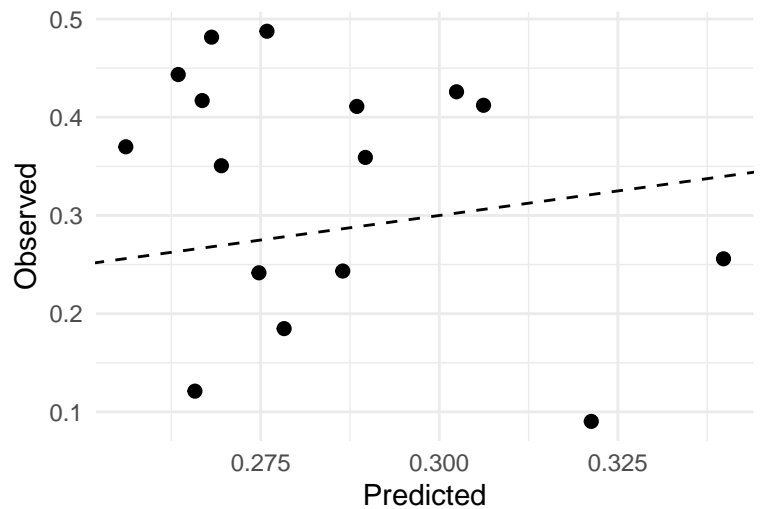
Full – PQ



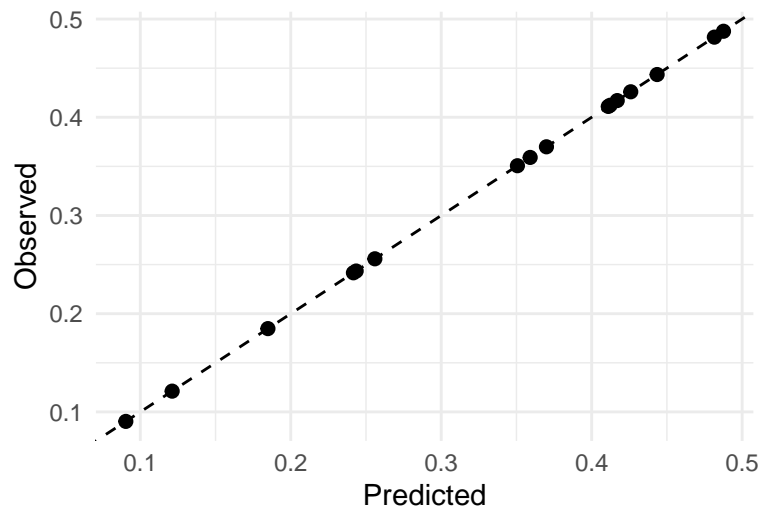
Range – PQ



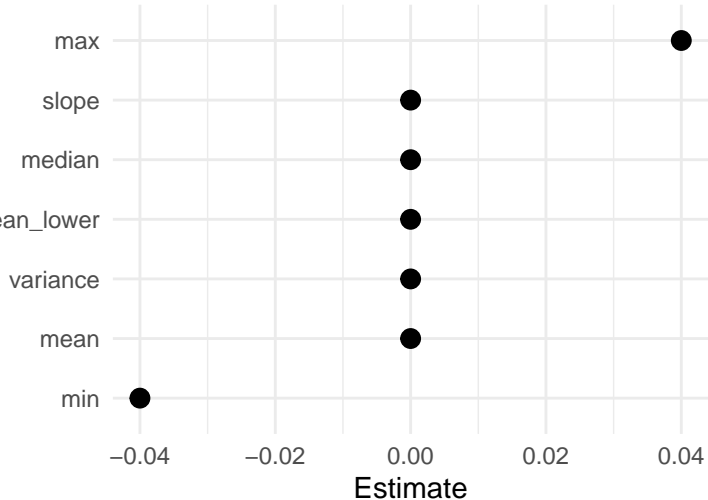
Mean_Upper – PQ



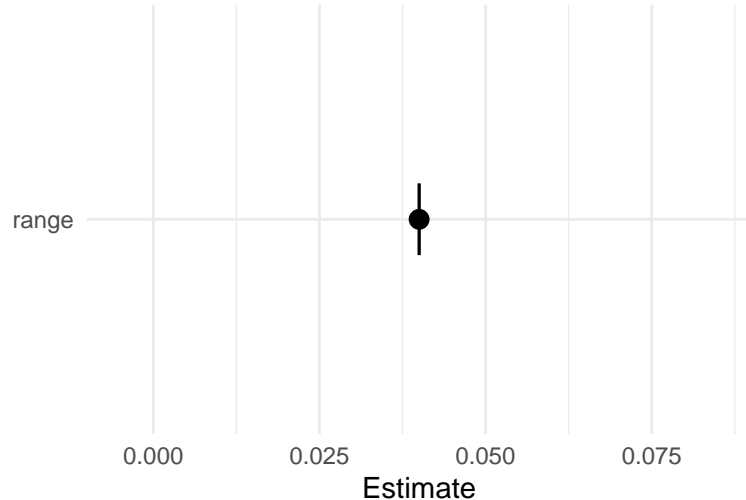
Ridge – PQ



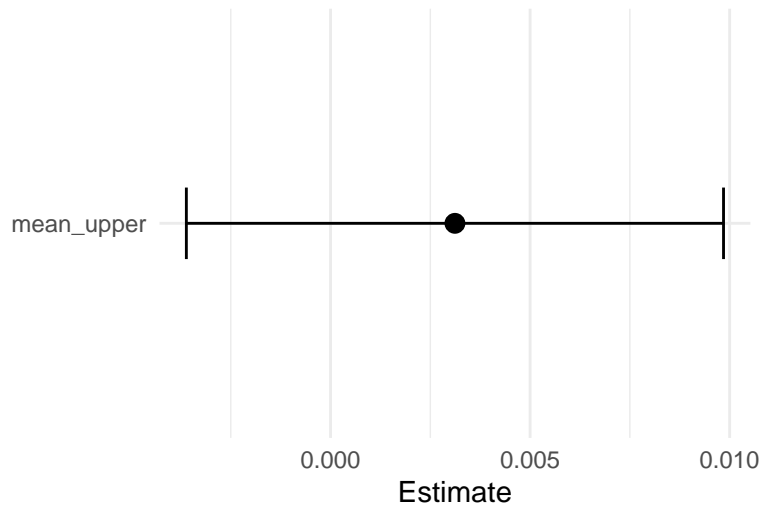
Full Coefficients – PQ



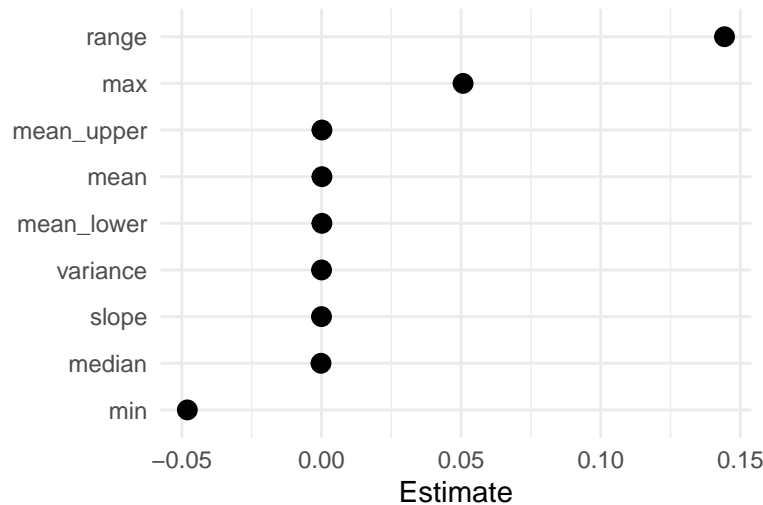
Range Coefficients – PQ

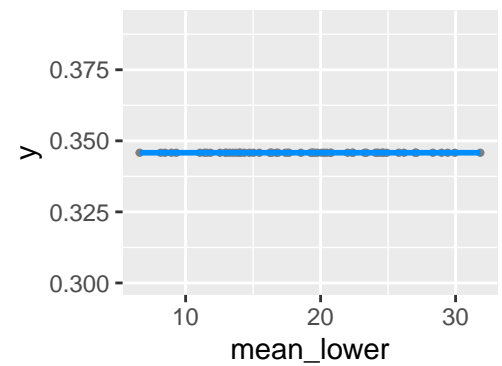
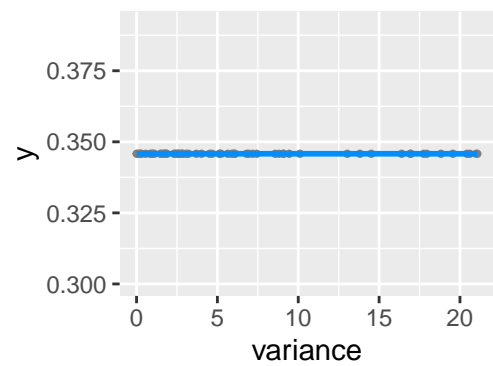
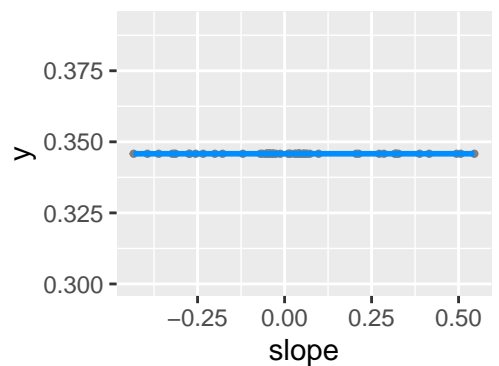
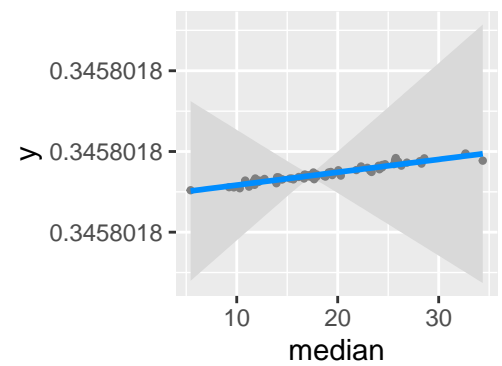
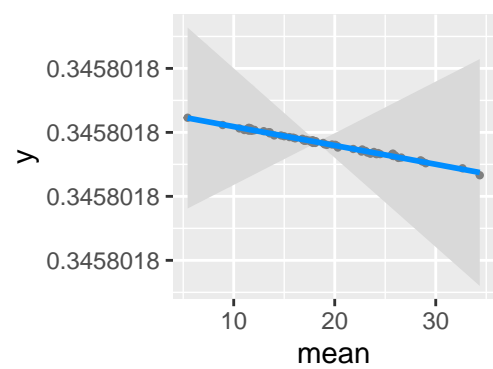
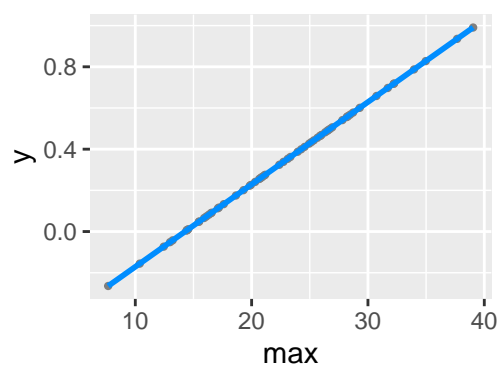
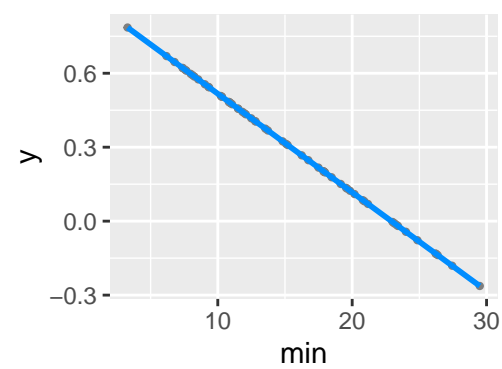


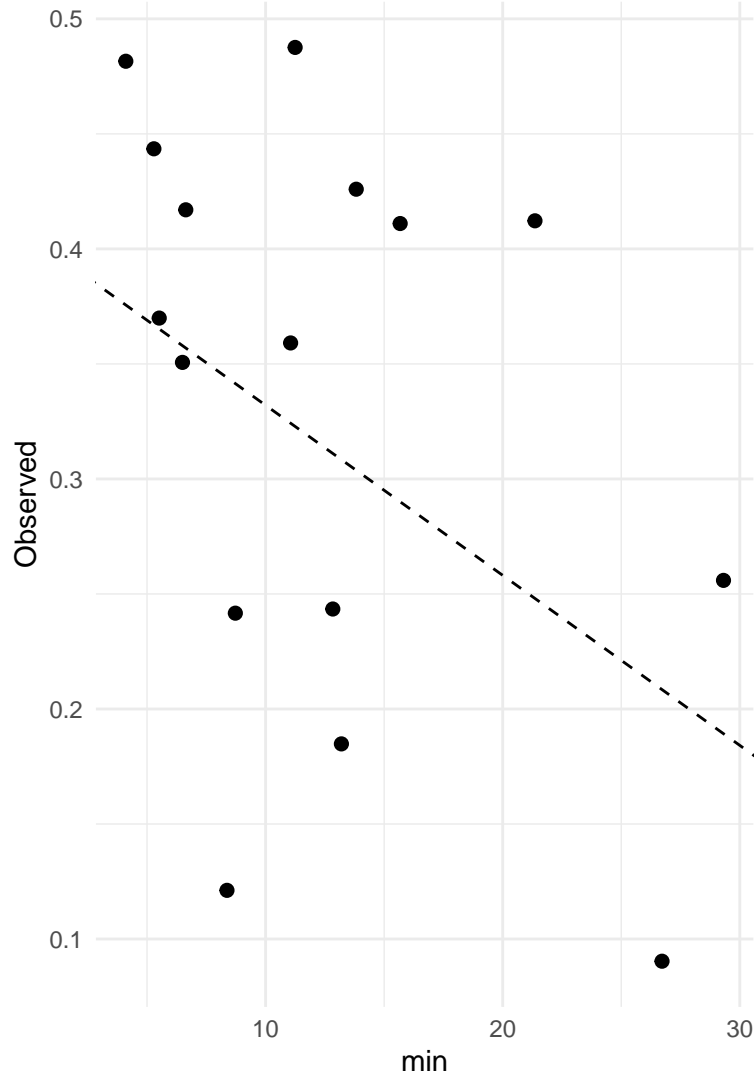
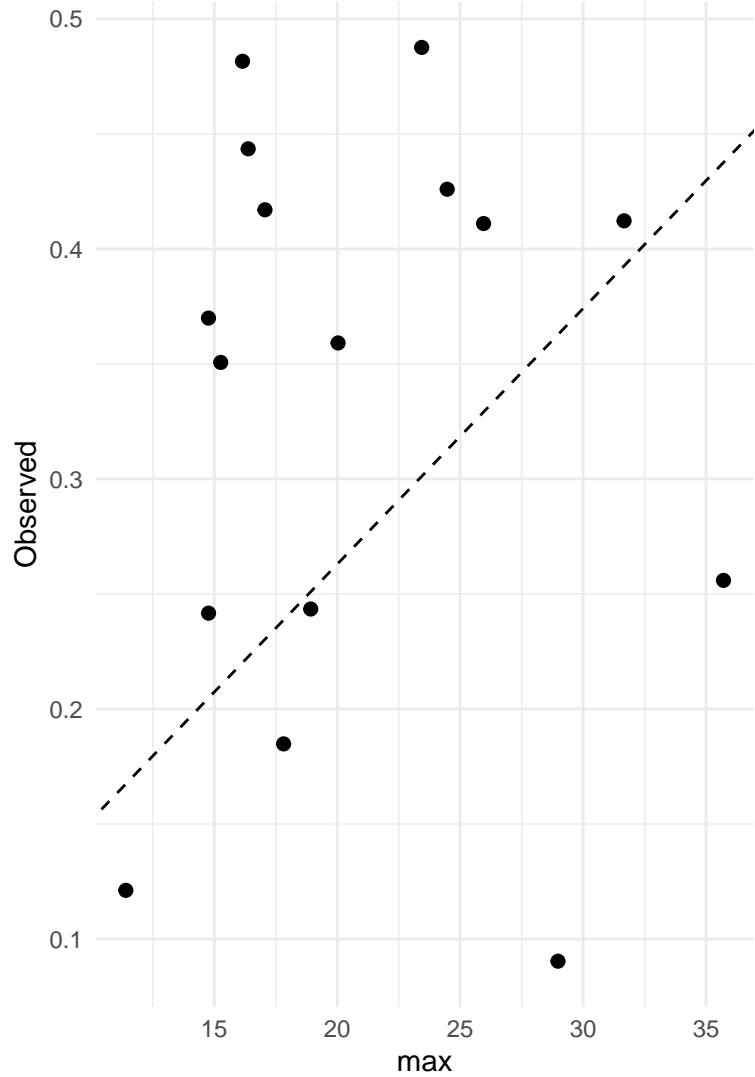
Mean_Upper Coefficients – PQ



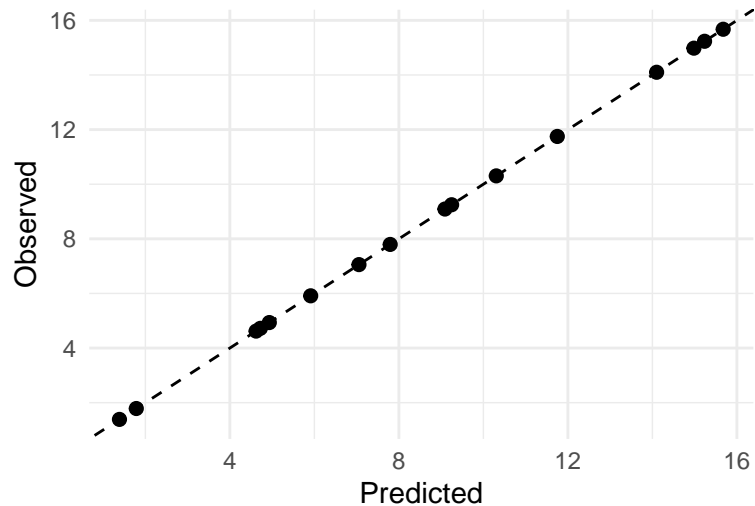
Ridge Coefficients – PQ



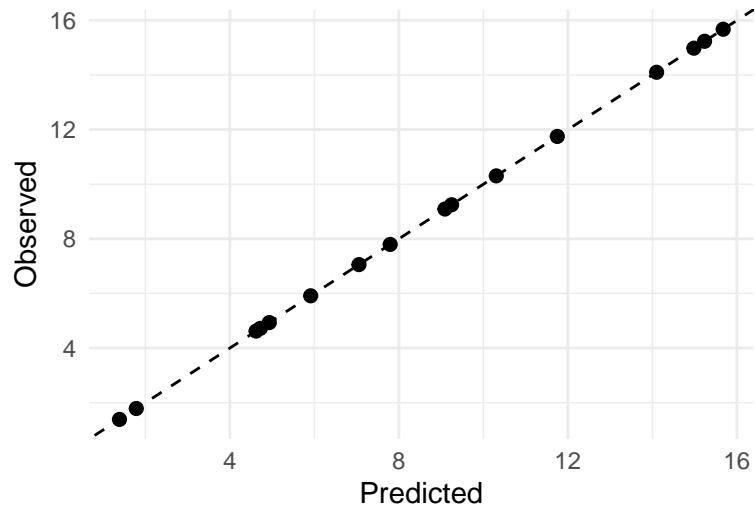


Simple Model: $y \sim \min$ Simple Model: $y \sim \max$ 

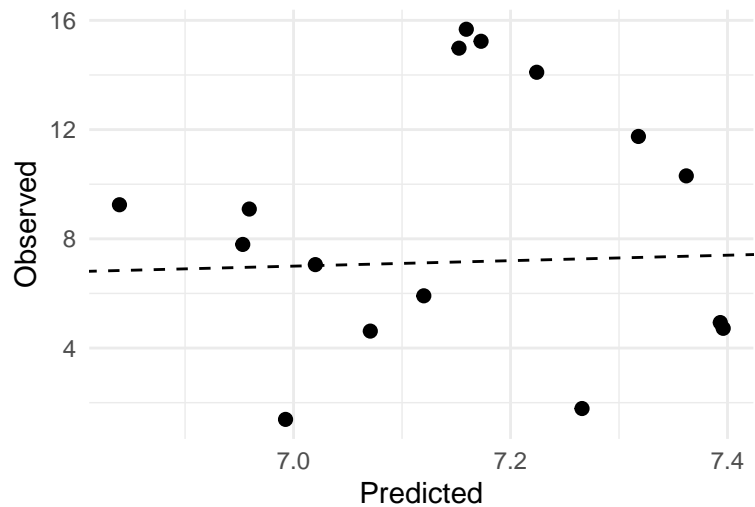
Full – PR



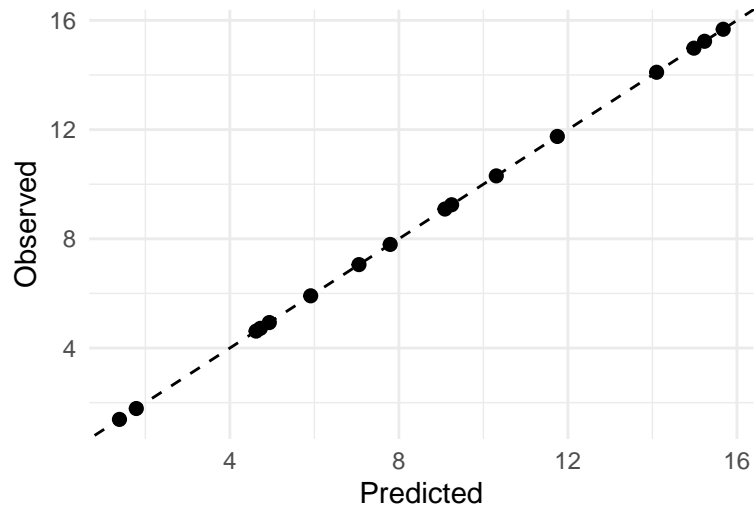
Range – PR



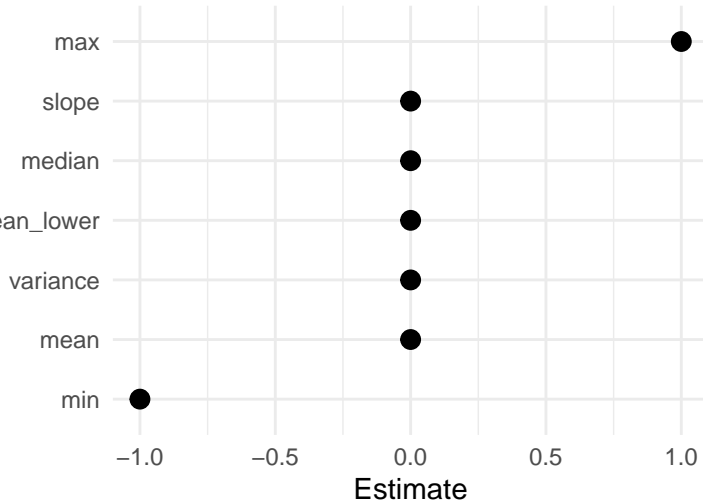
Mean_Upper – PR



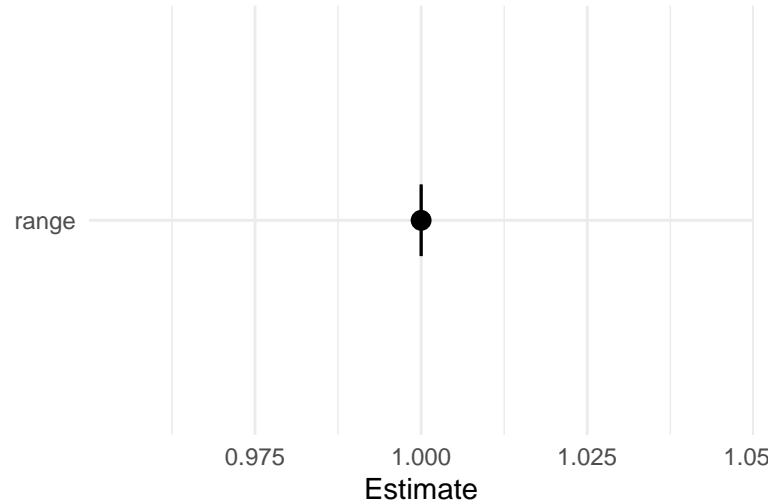
Ridge – PR



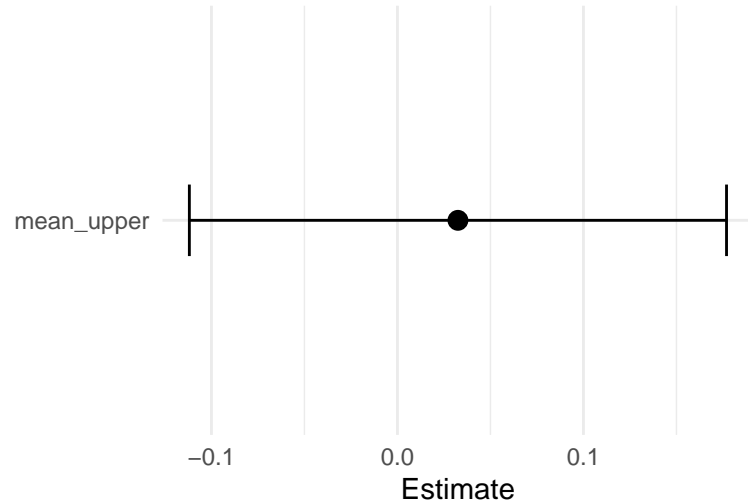
Full Coefficients – PR



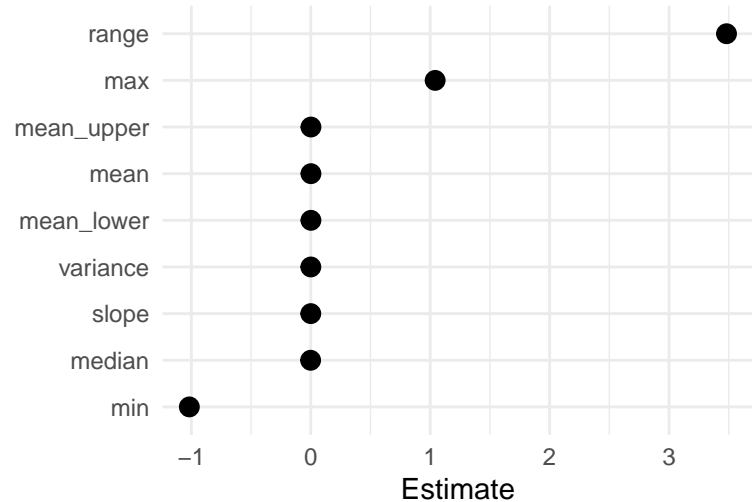
Range Coefficients – PR

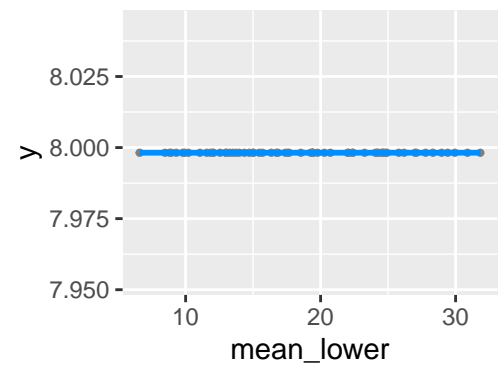
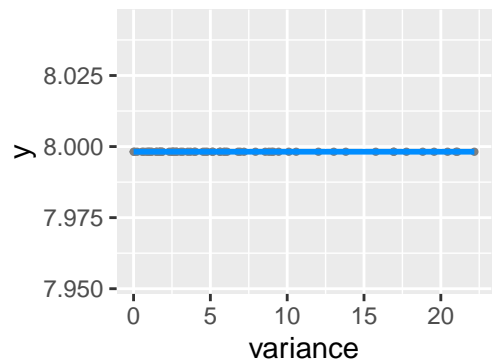
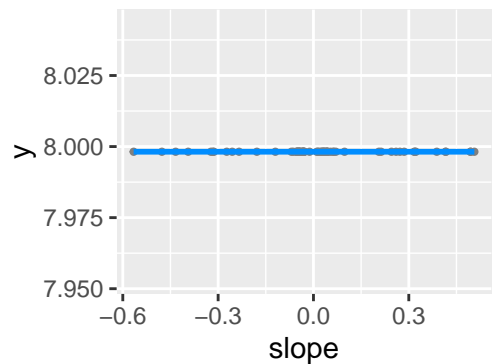
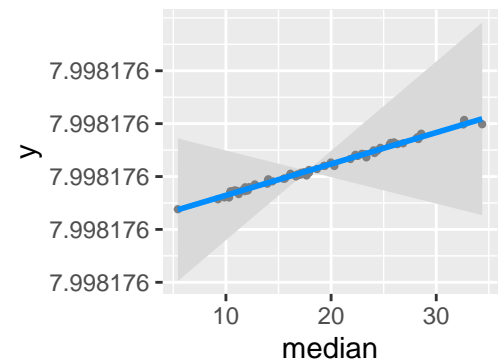
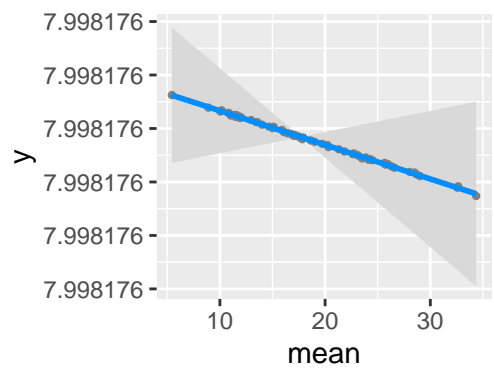
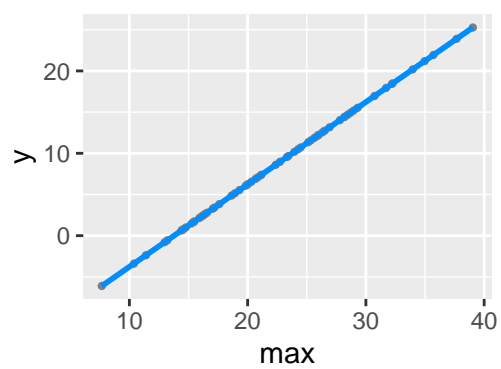
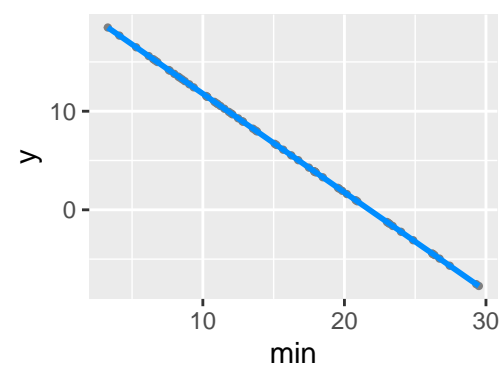


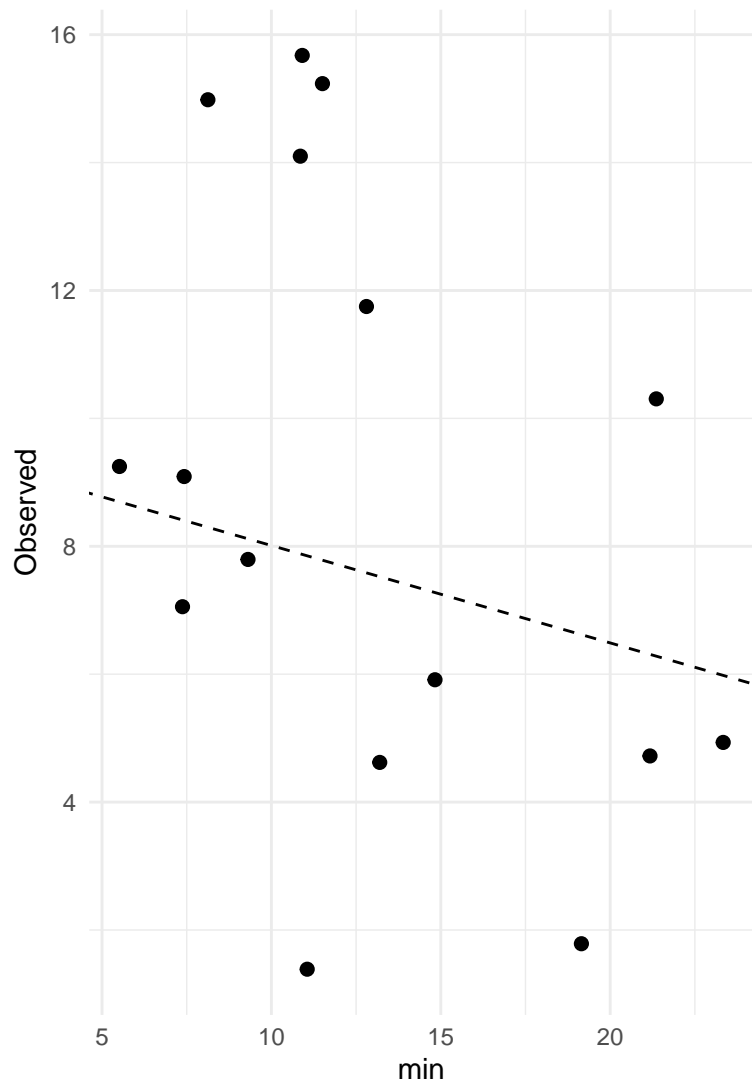
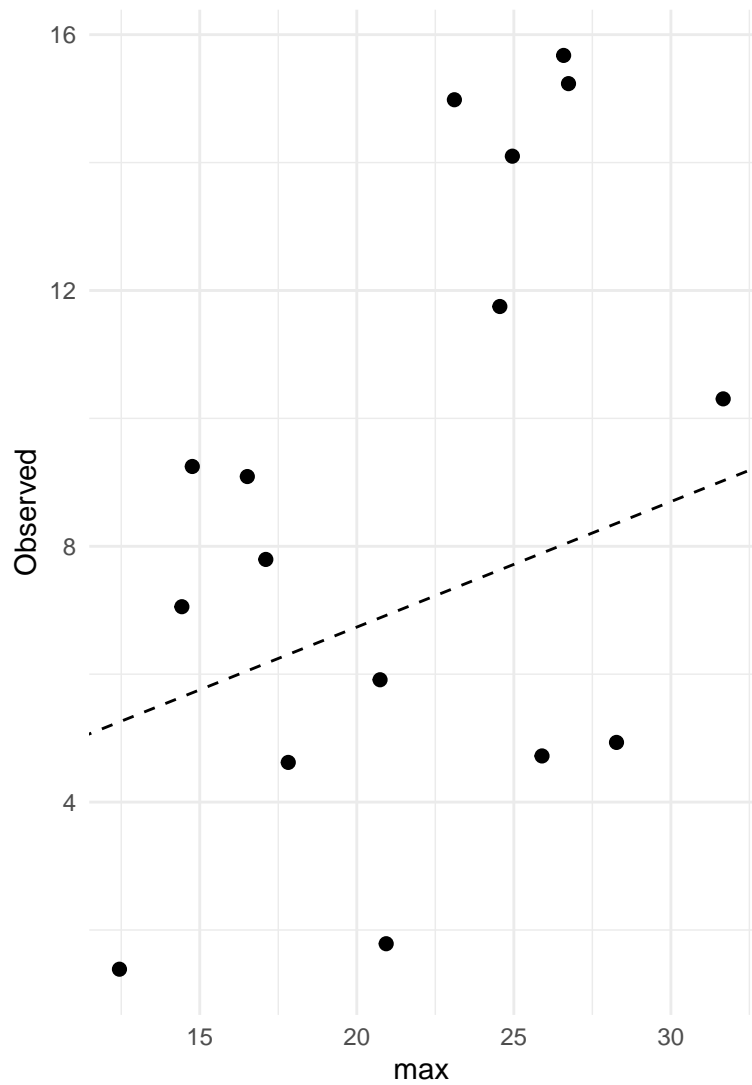
Mean_Upper Coefficients – PR



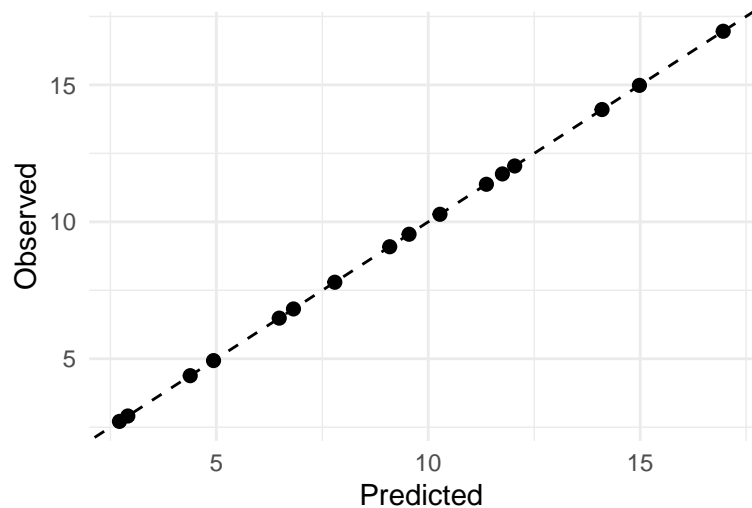
Ridge Coefficients – PR



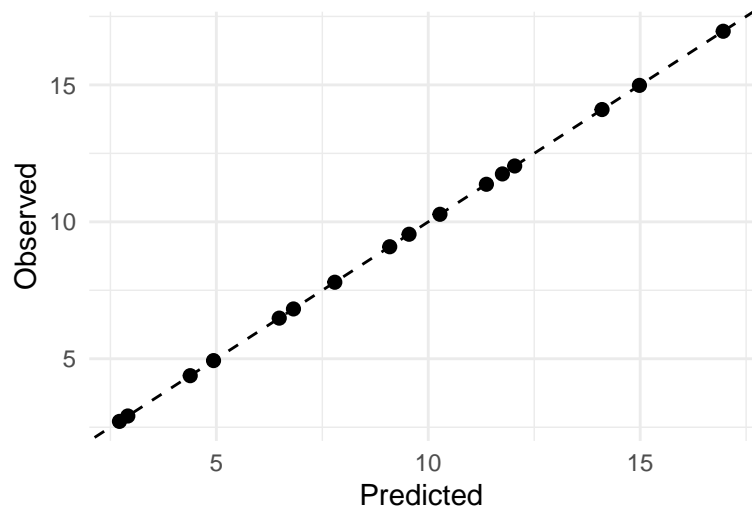


Simple Model: $y \sim \min$ Simple Model: $y \sim \max$ 

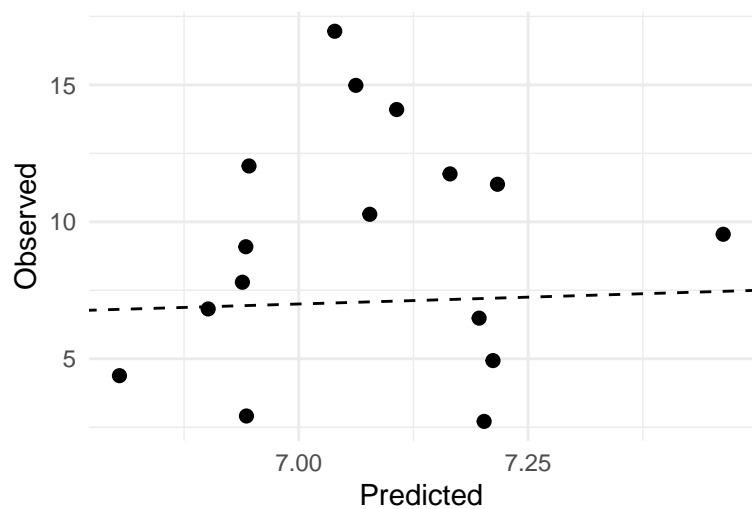
Full – NRW



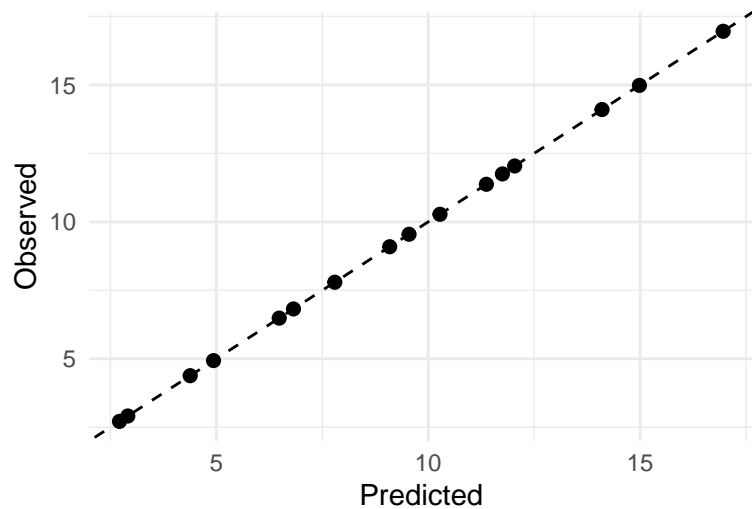
Range – NRW



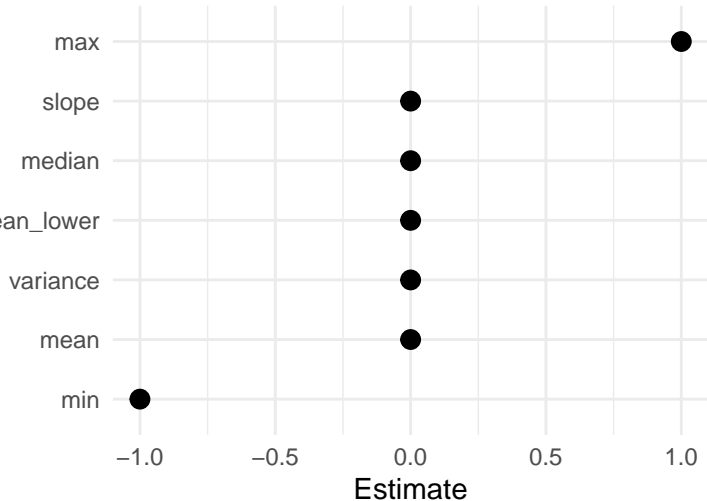
Mean_Upper – NRW



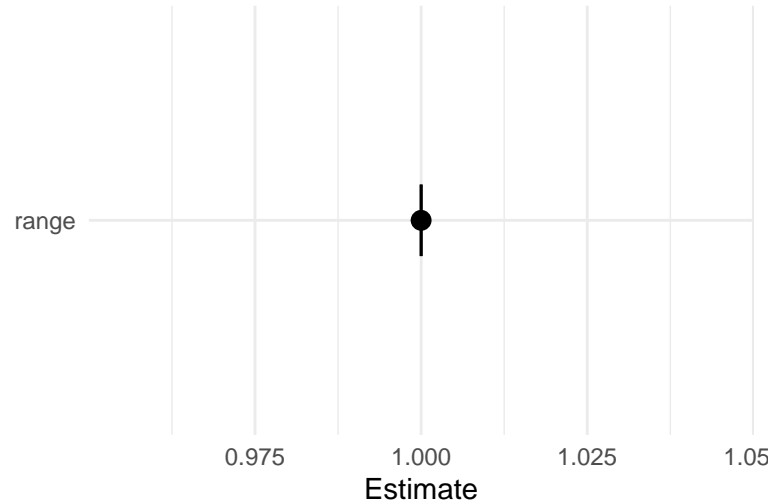
Ridge – NRW



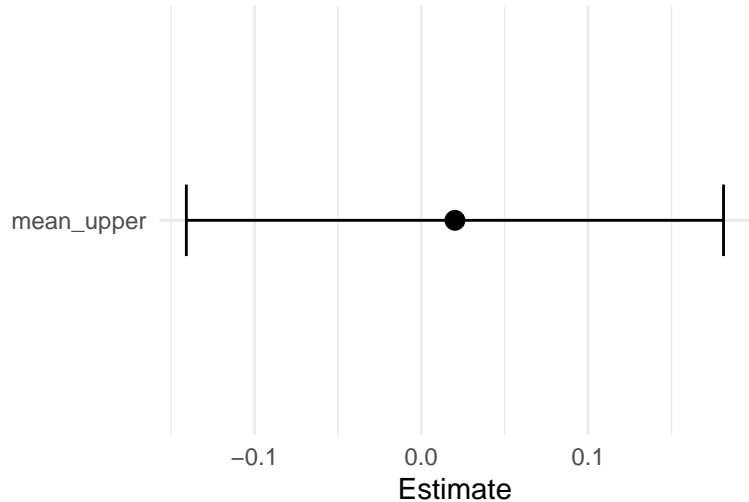
Full Coefficients – NRW



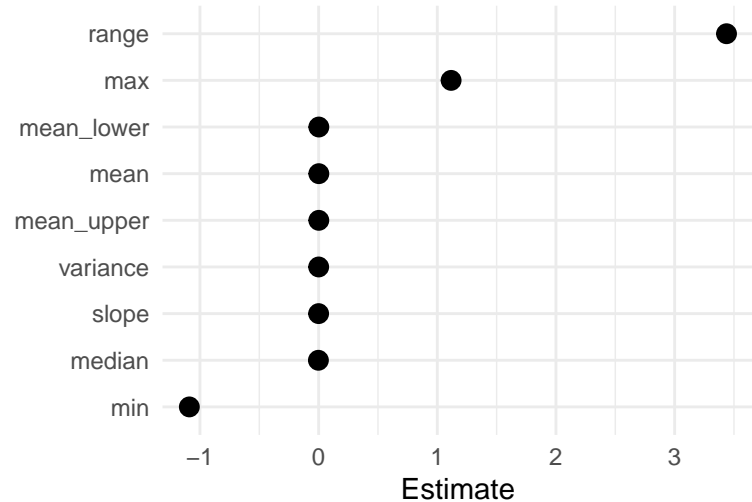
Range Coefficients – NRW

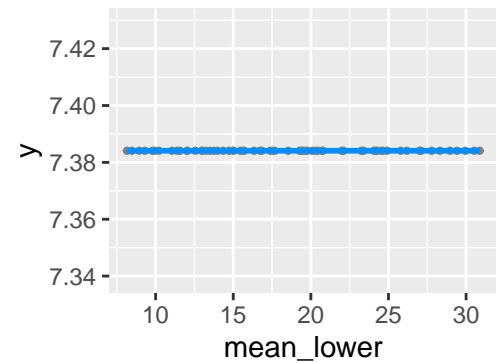
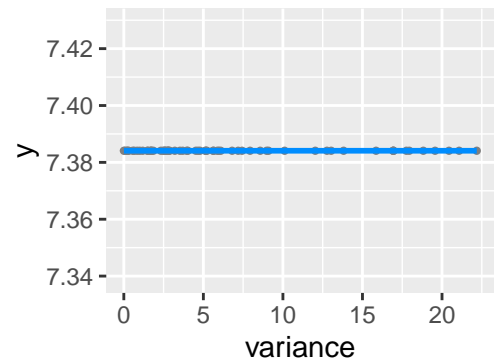
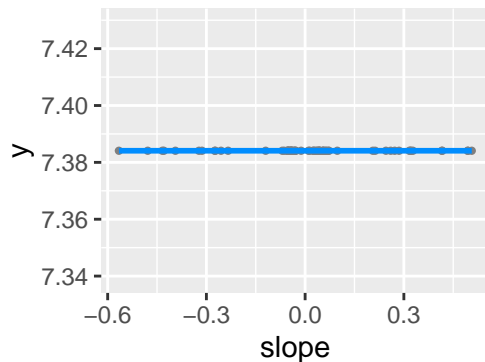
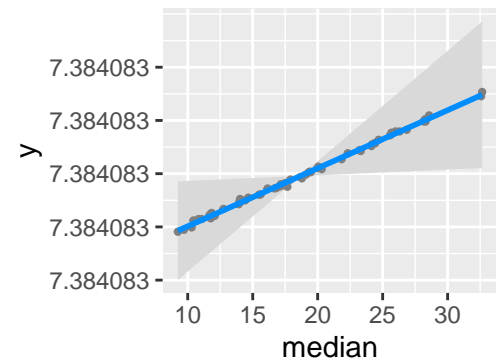
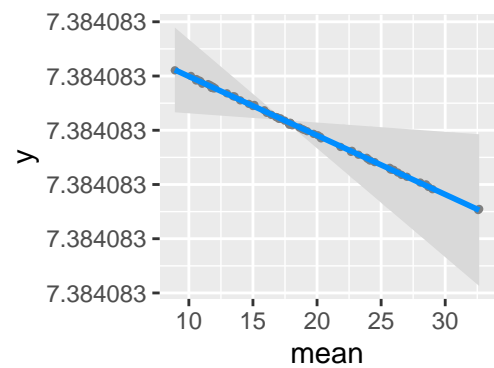
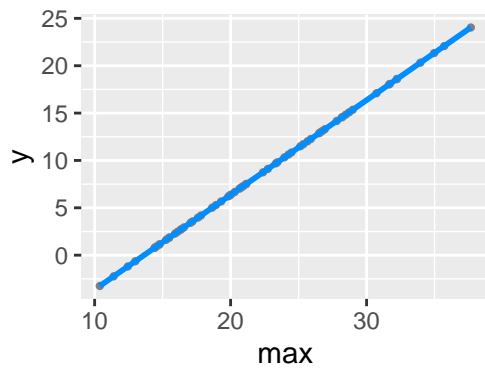
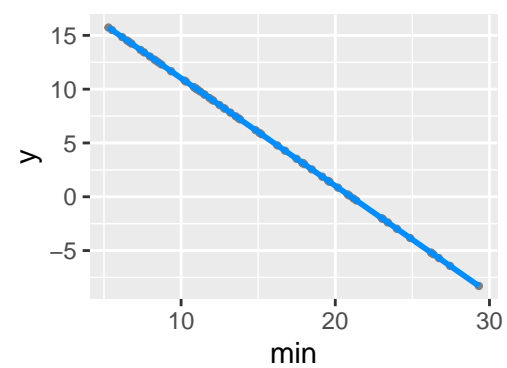


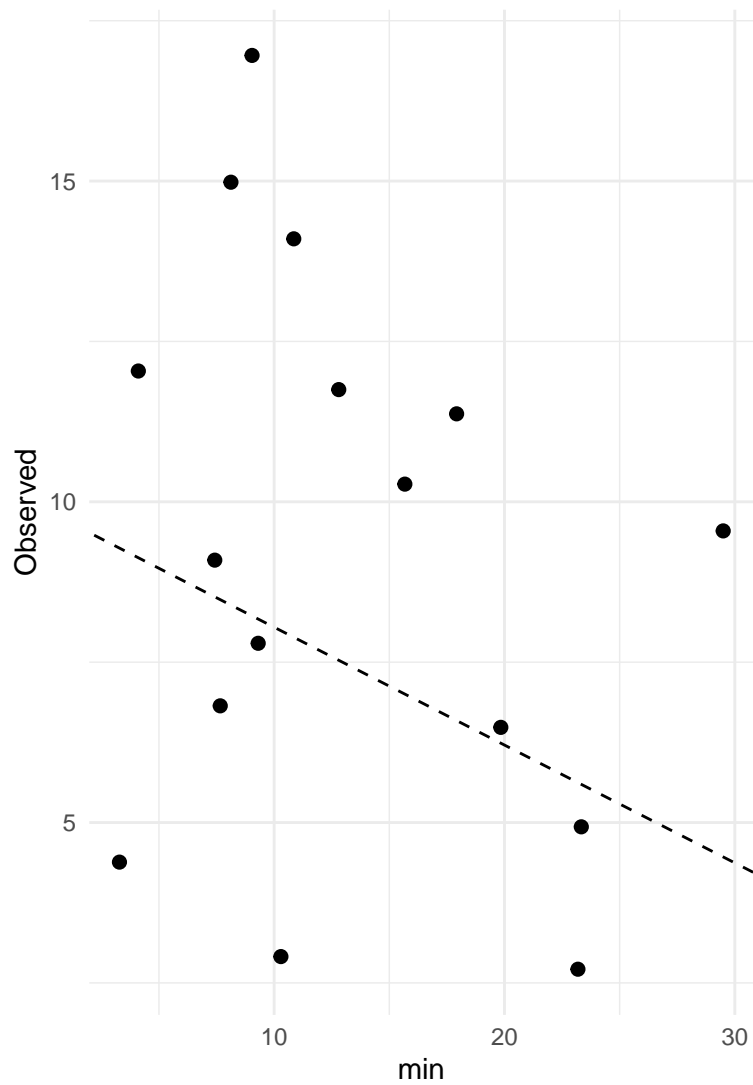
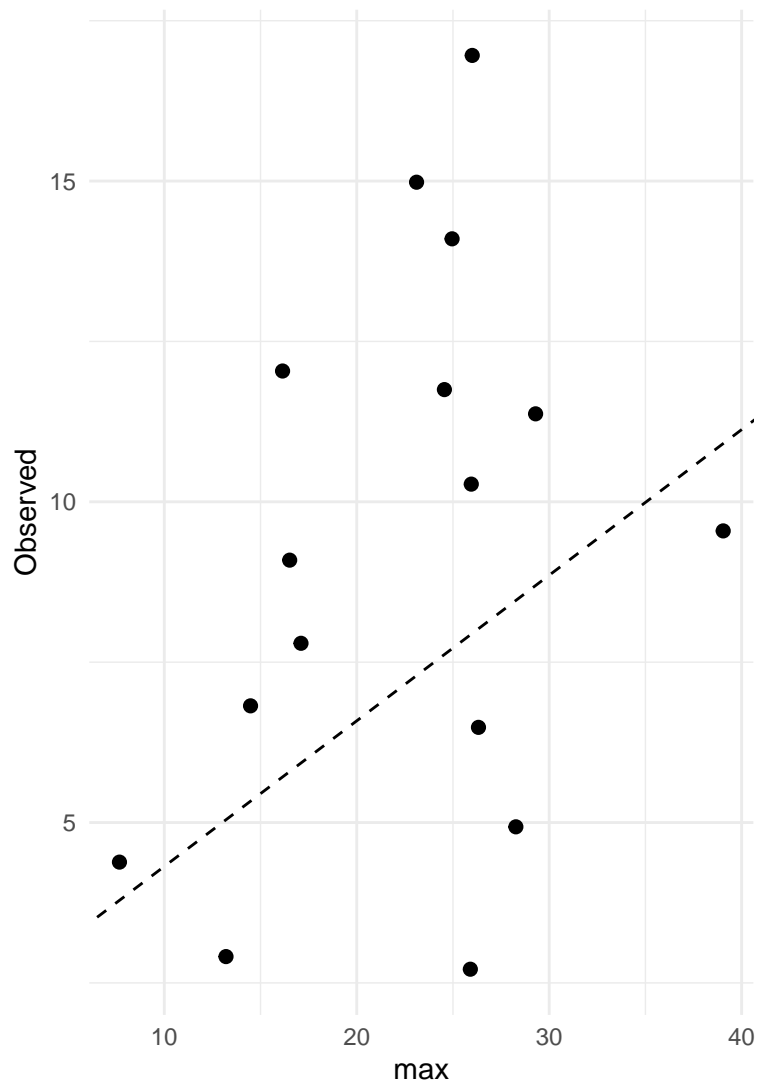
Mean_Upper Coefficients – NRW



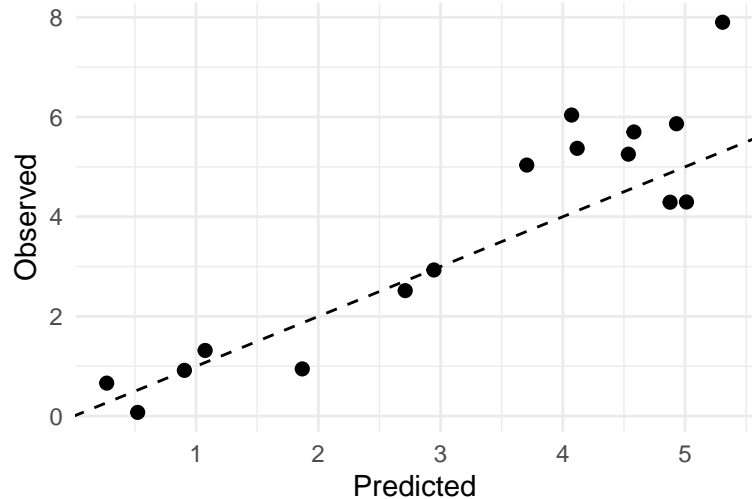
Ridge Coefficients – NRW



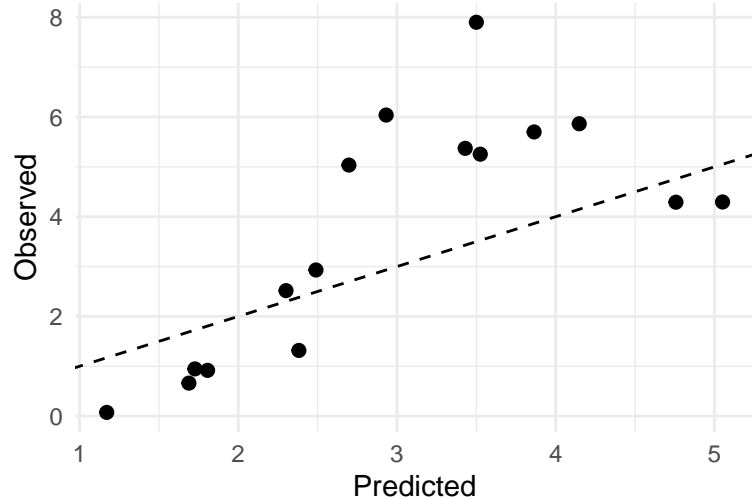


Simple Model: $y \sim \min$ Simple Model: $y \sim \max$ 

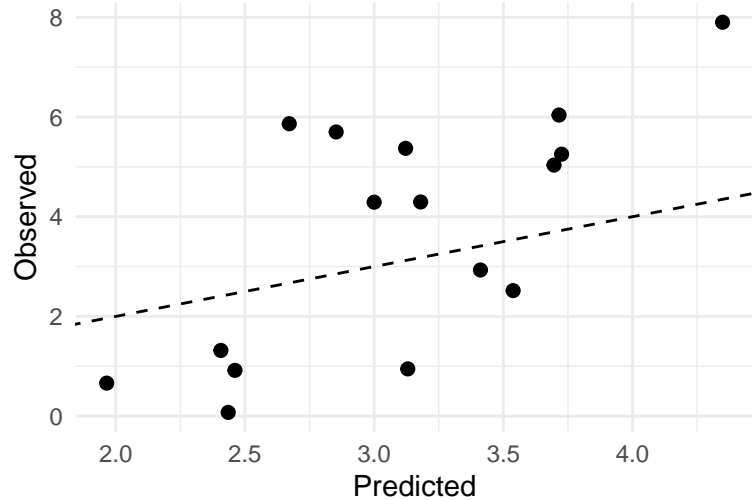
Full – ESP



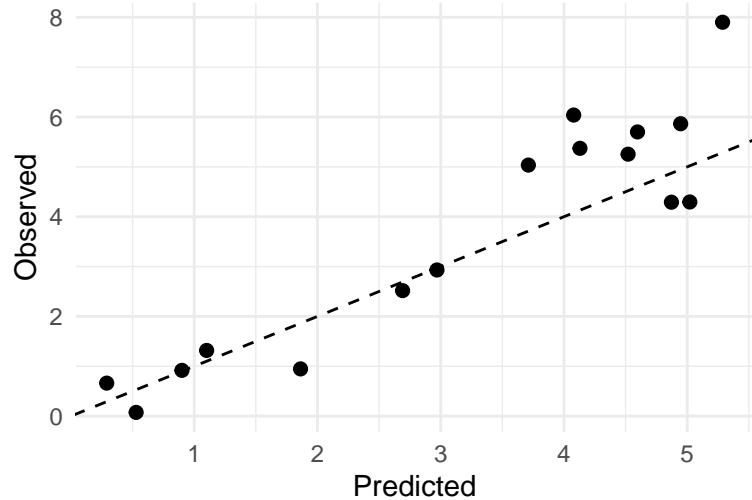
Range – ESP



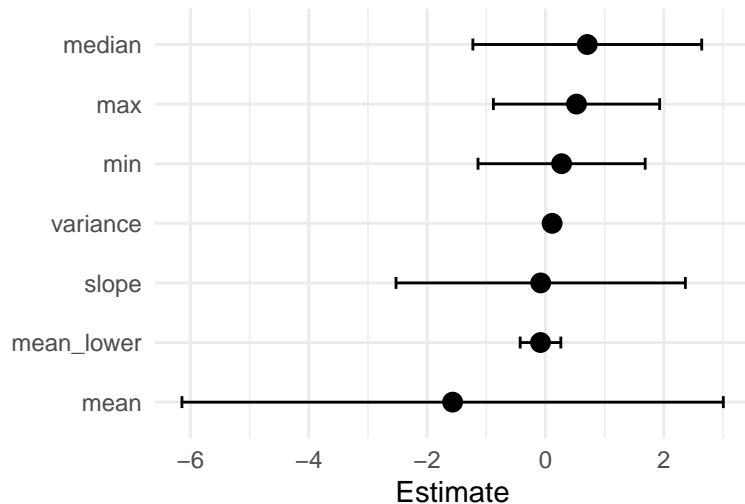
Mean_Upper – ESP



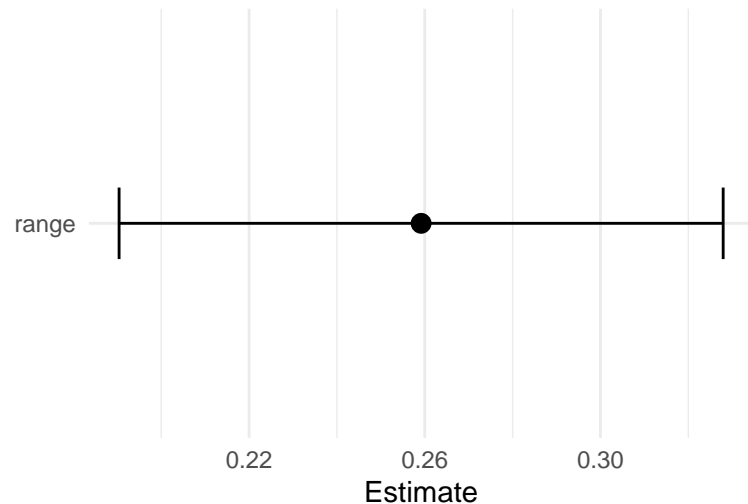
Ridge – ESP



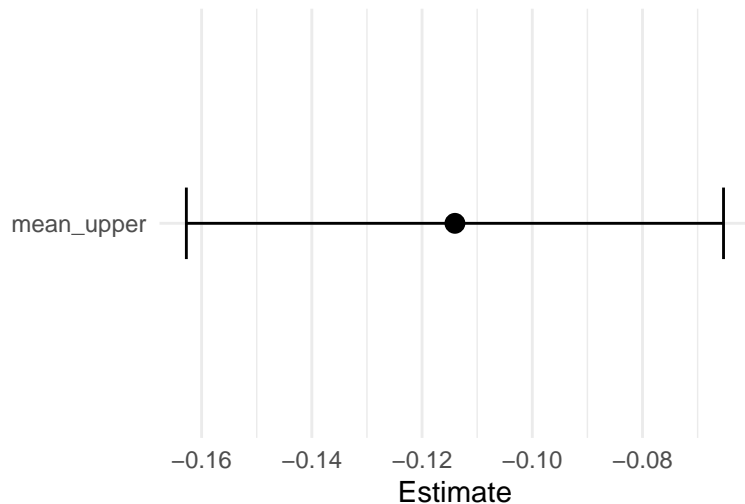
Full Coefficients – ESP



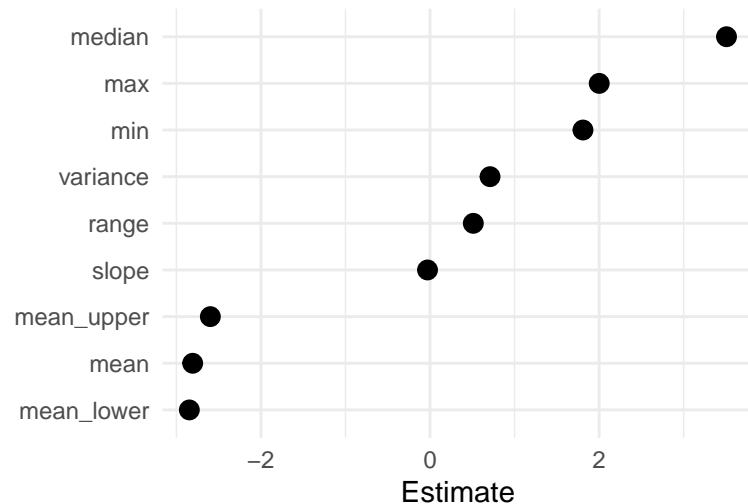
Range Coefficients – ESP

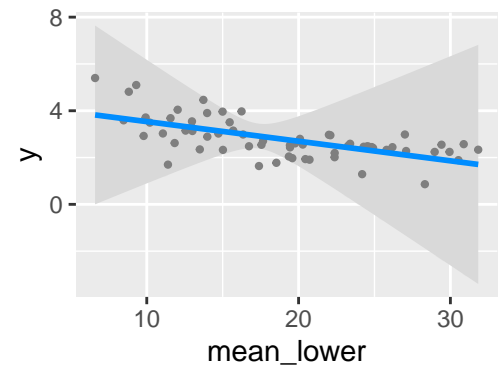
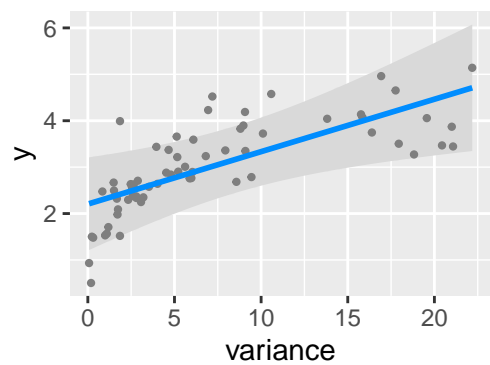
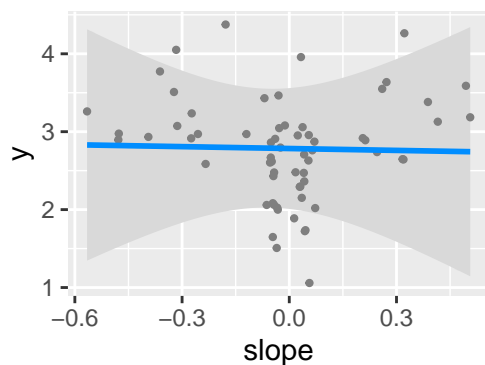
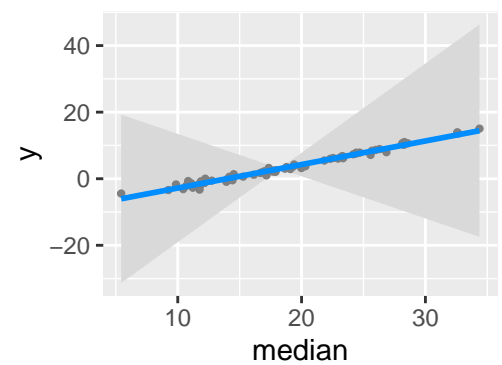
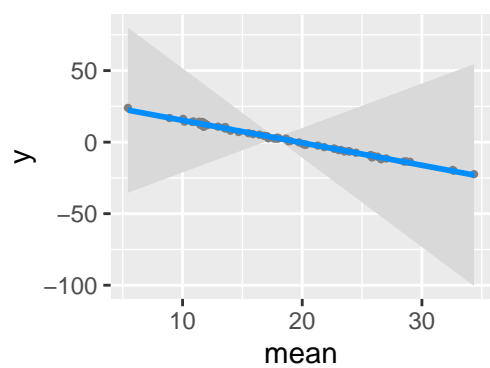
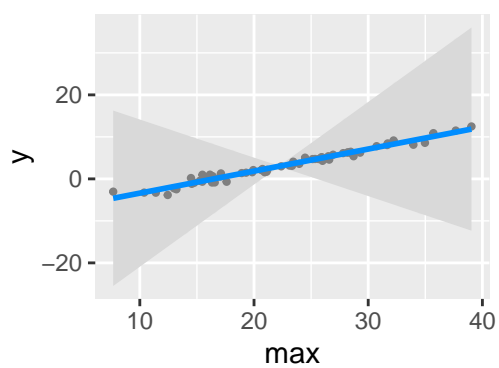
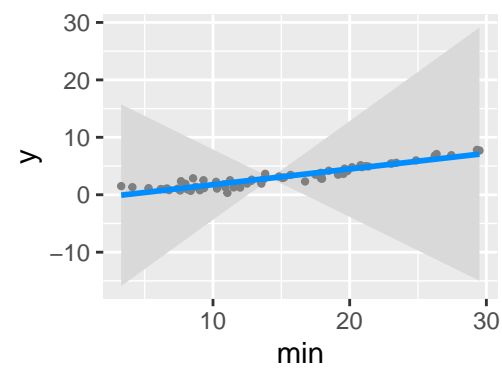


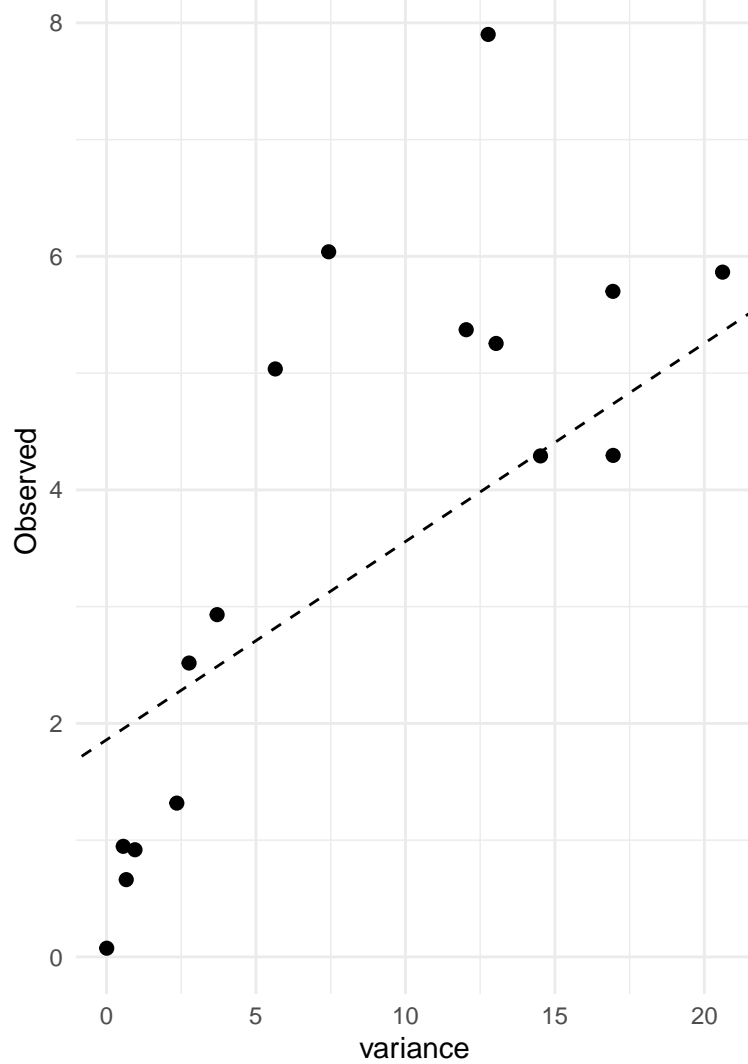
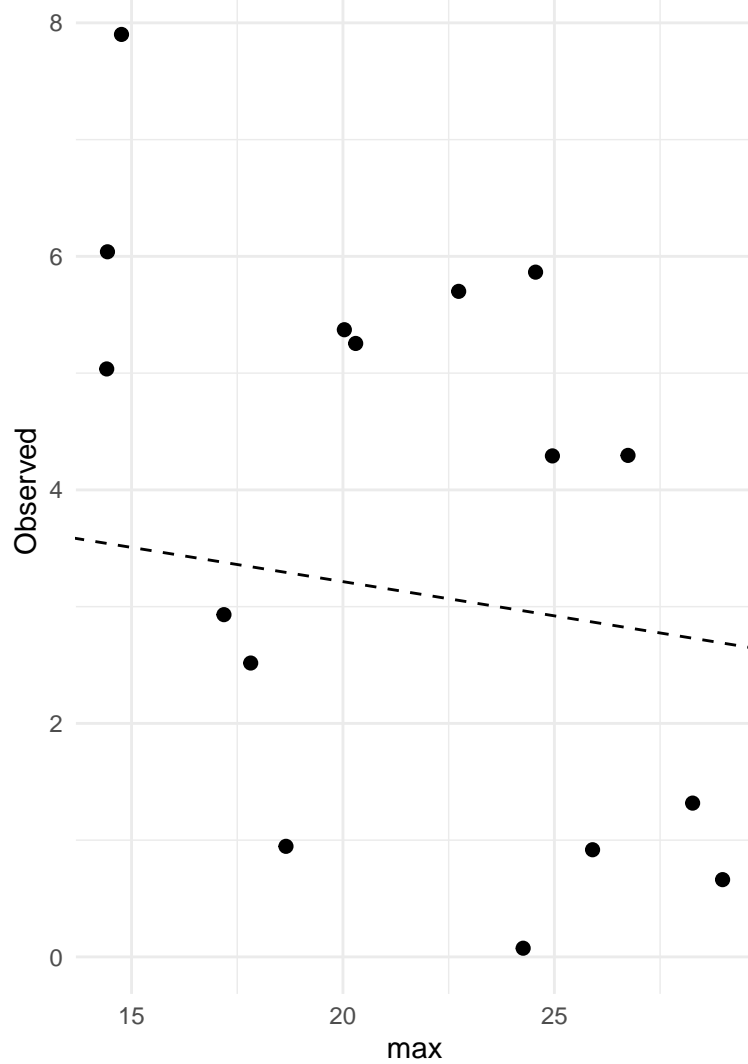
Mean_Upper Coefficients – ESP



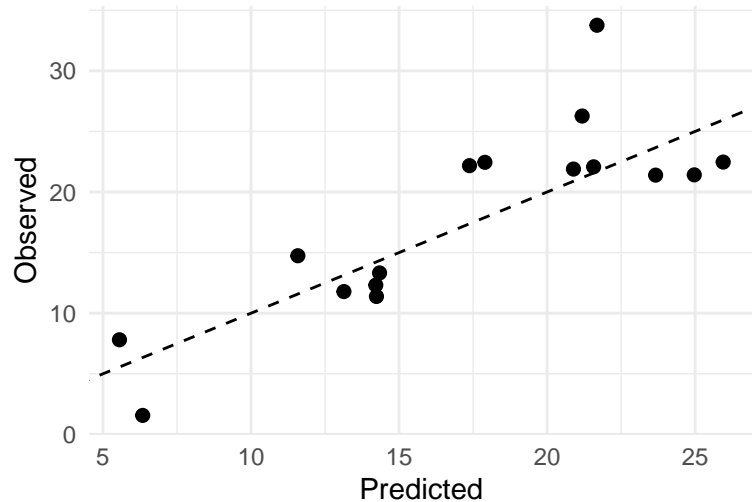
Ridge Coefficients – ESP



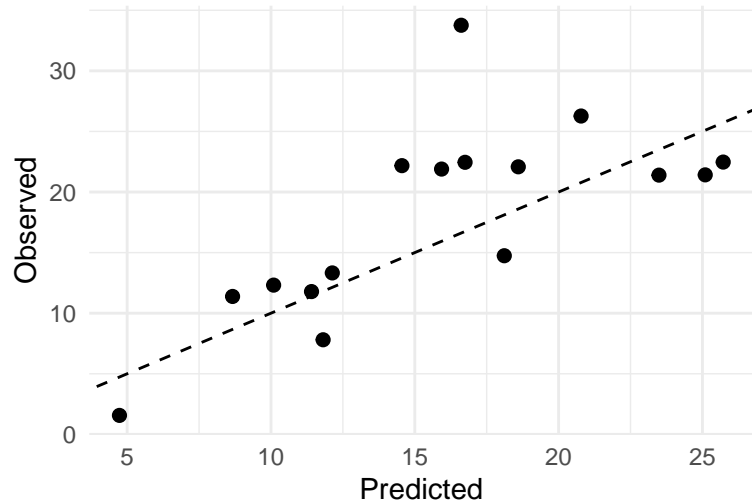


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{max}$ 

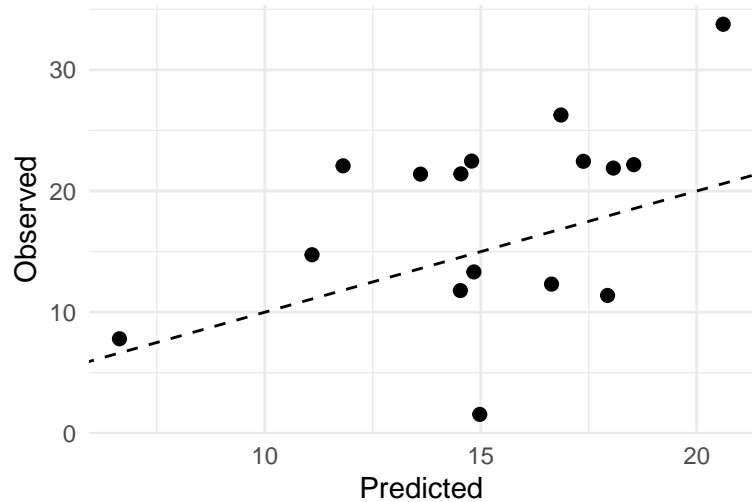
Full – CEV



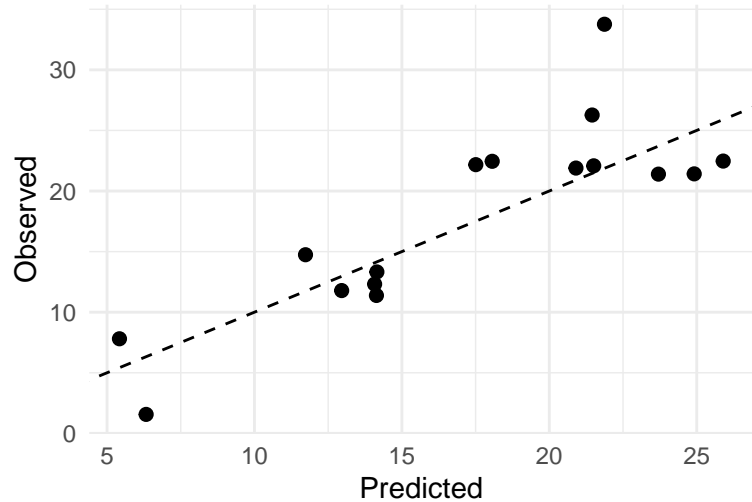
Range – CEV



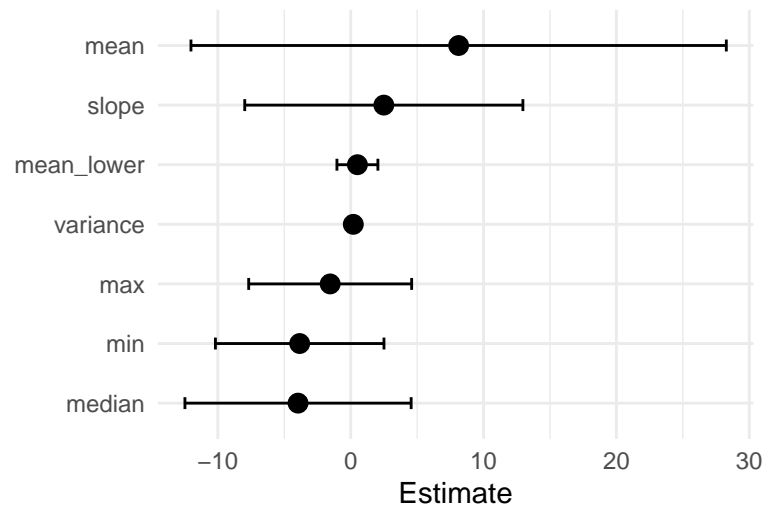
Mean_Upper – CEV



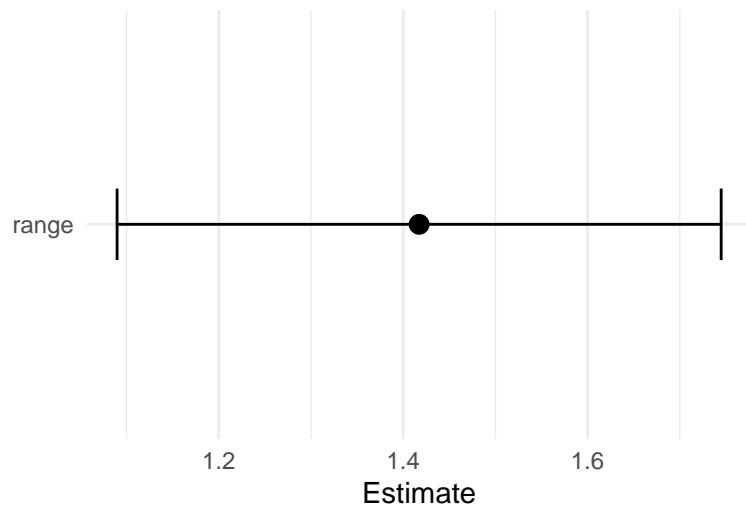
Ridge – CEV



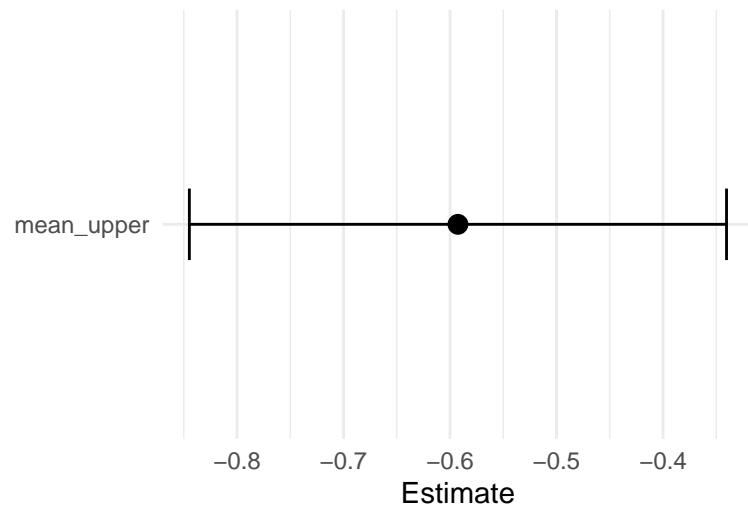
Full Coefficients – CEV



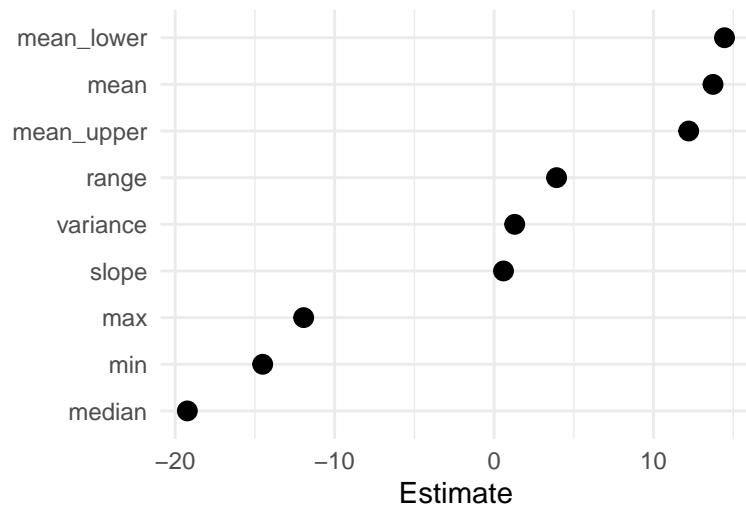
Range Coefficients – CEV

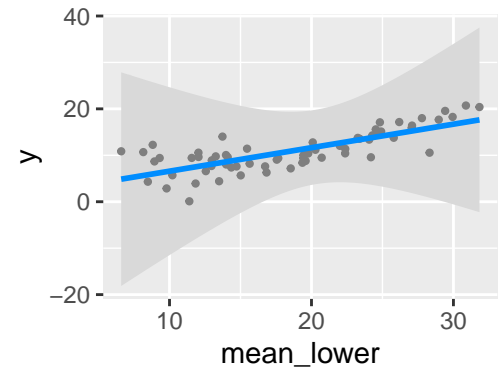
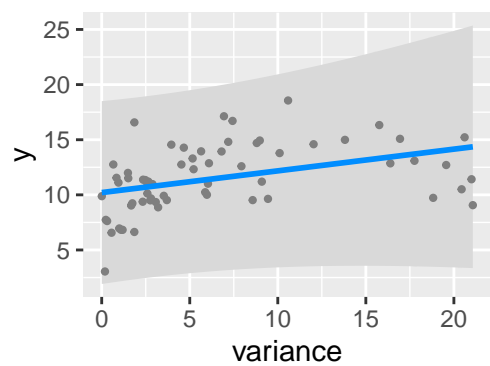
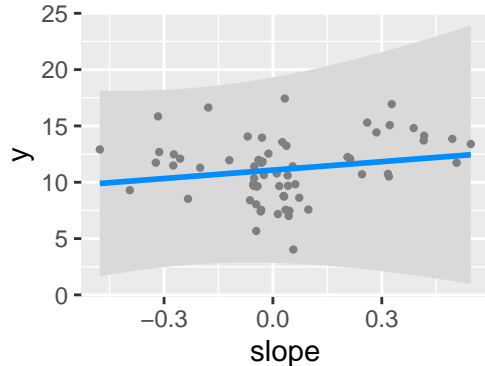
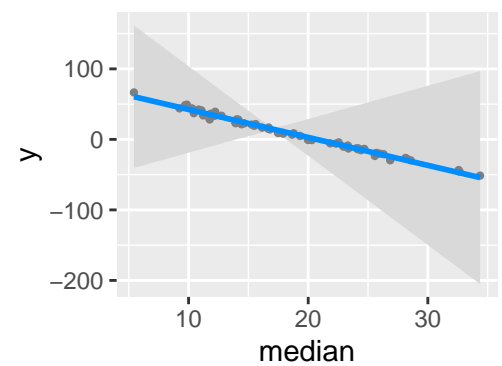
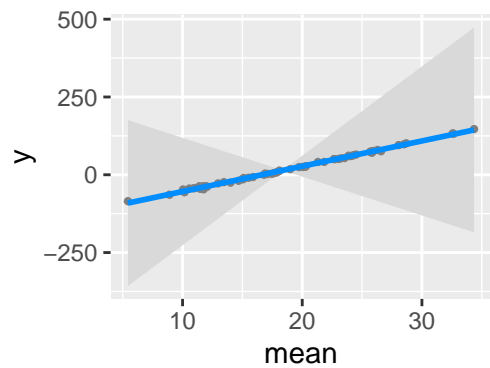
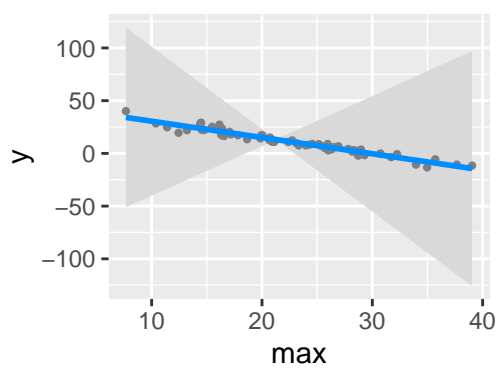
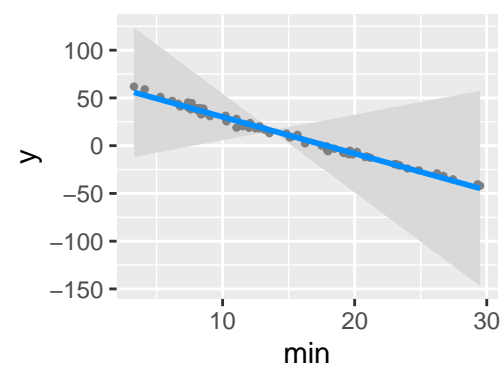


Mean_Upper Coefficients – CEV

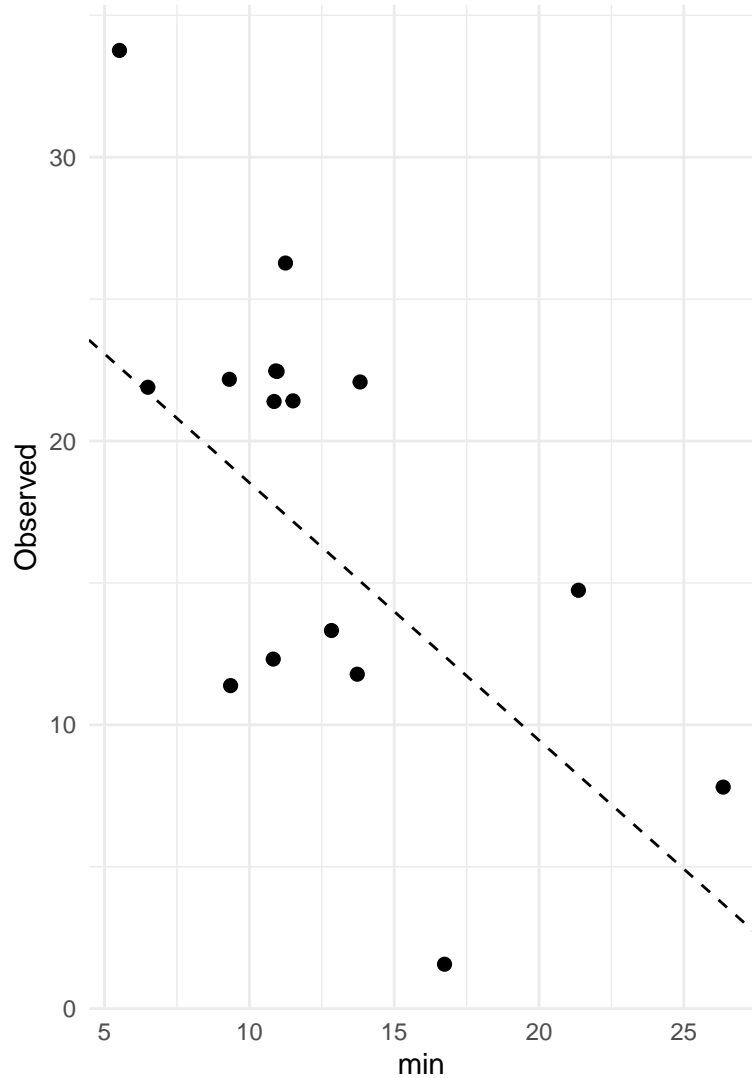


Ridge Coefficients – CEV

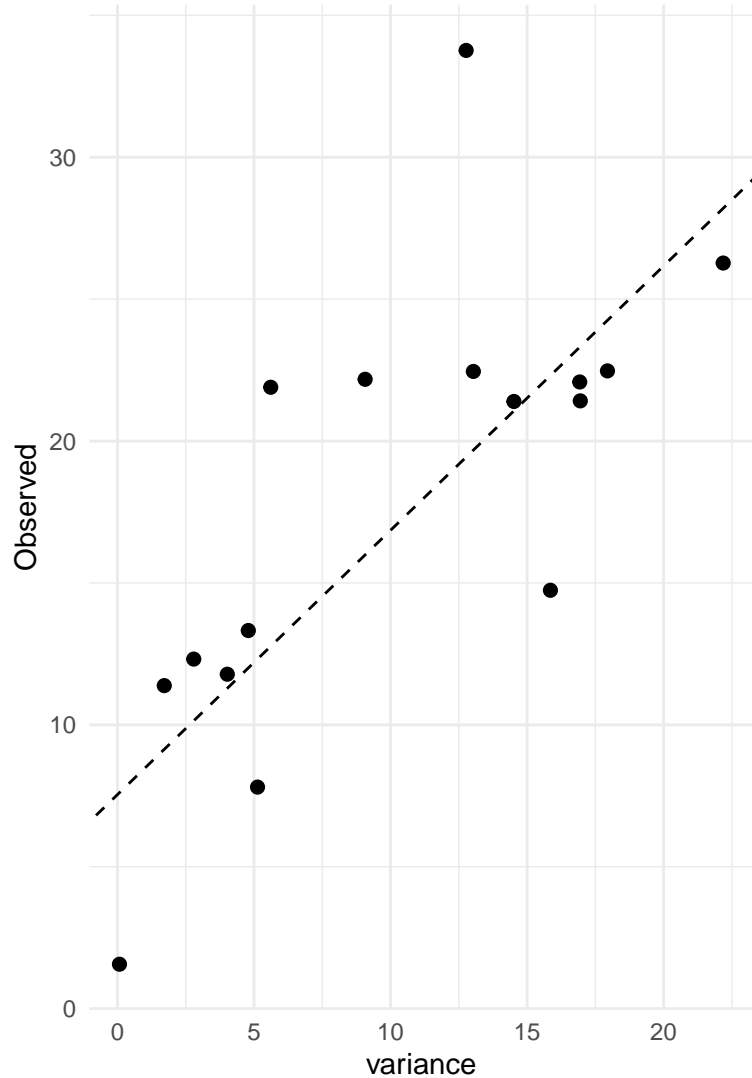




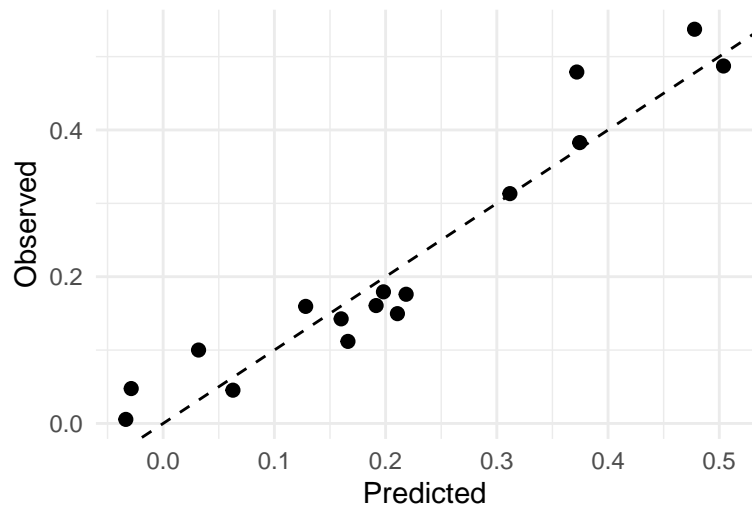
Simple Model: $y \sim \text{min}$



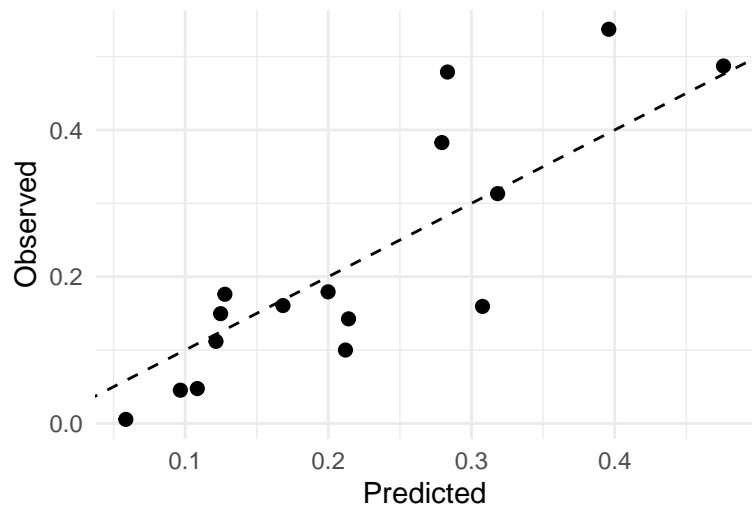
Simple Model: $y \sim \text{variance}$



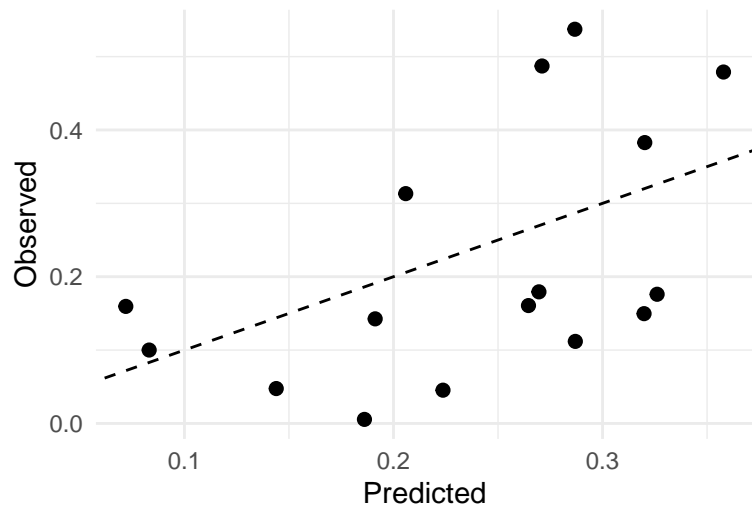
Full – PFI



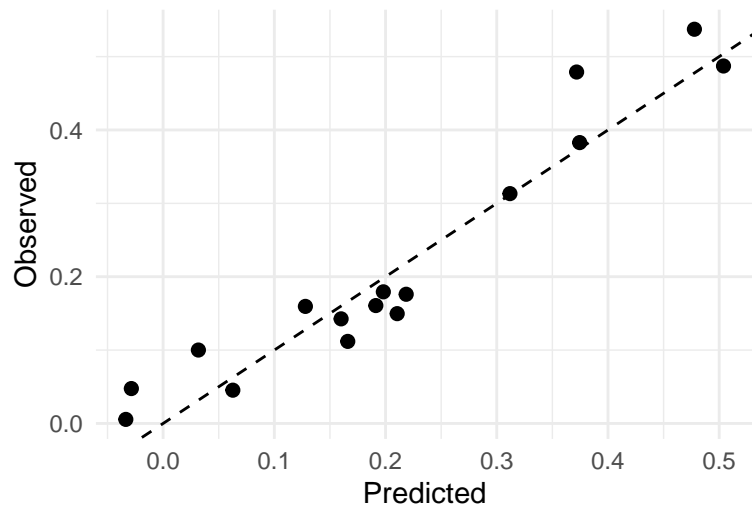
Range – PFI



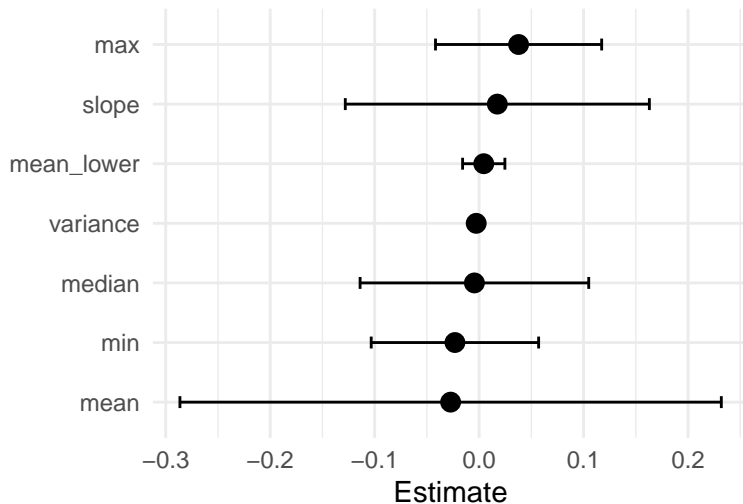
Mean_Upper – PFI



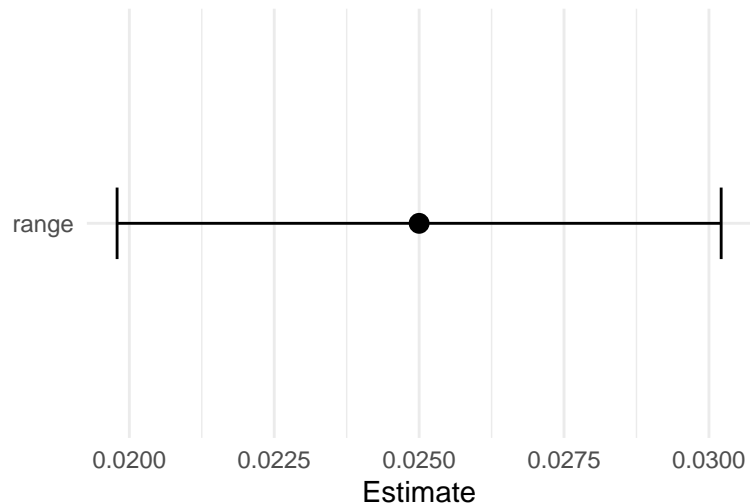
Ridge – PFI



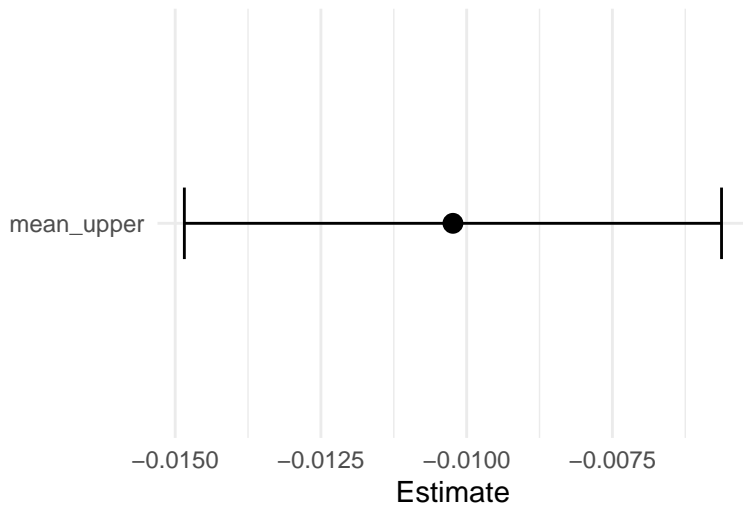
Full Coefficients – PFI



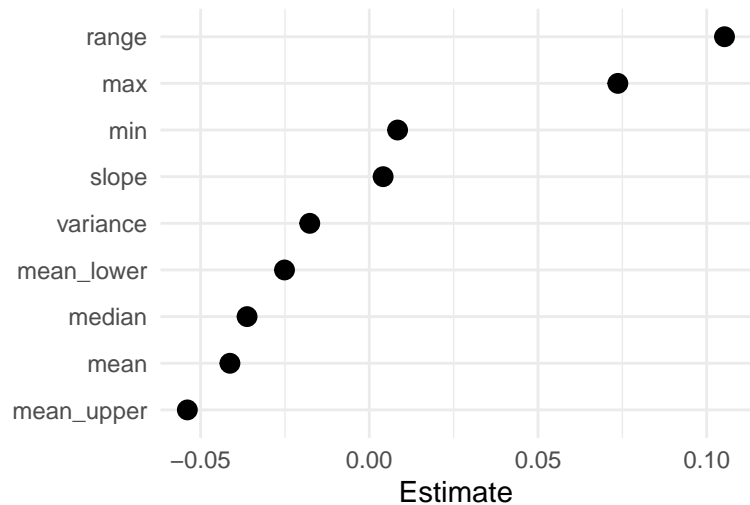
Range Coefficients – PFI

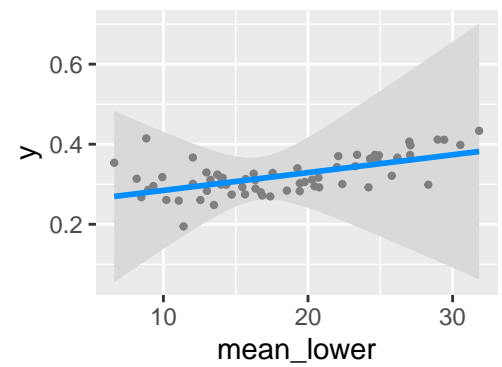
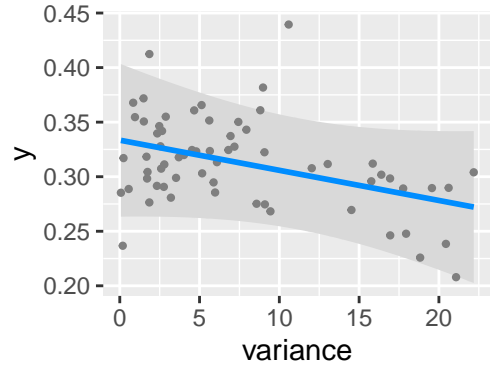
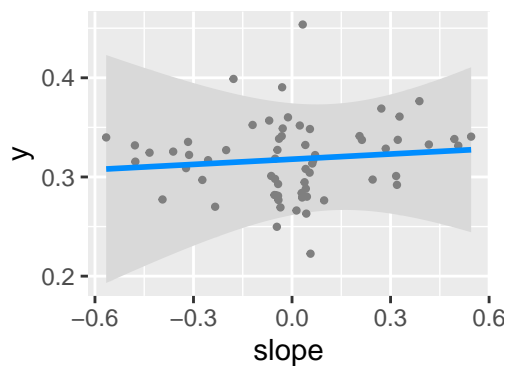
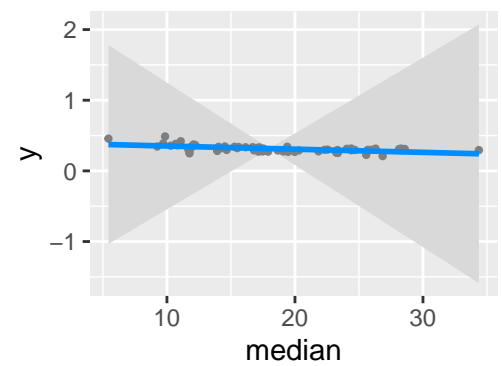
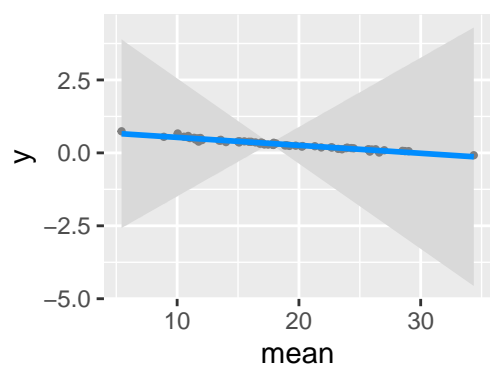
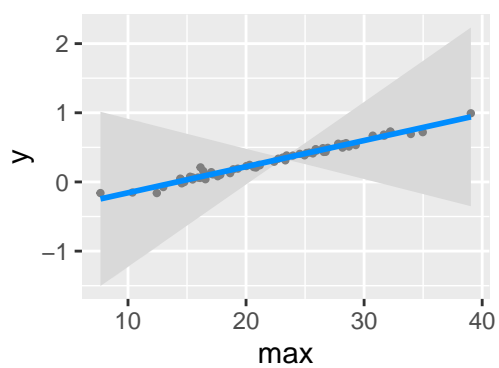
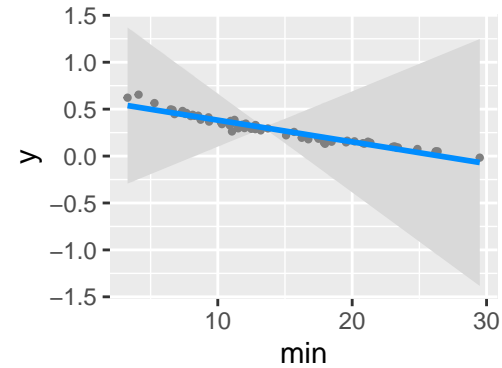


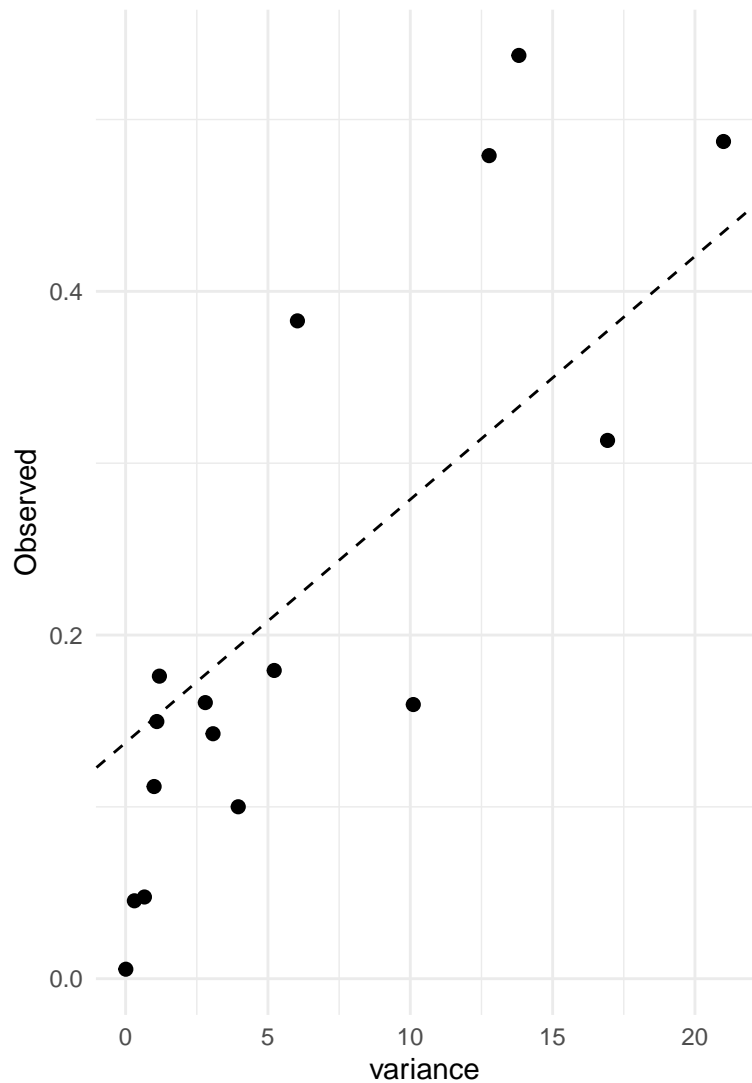
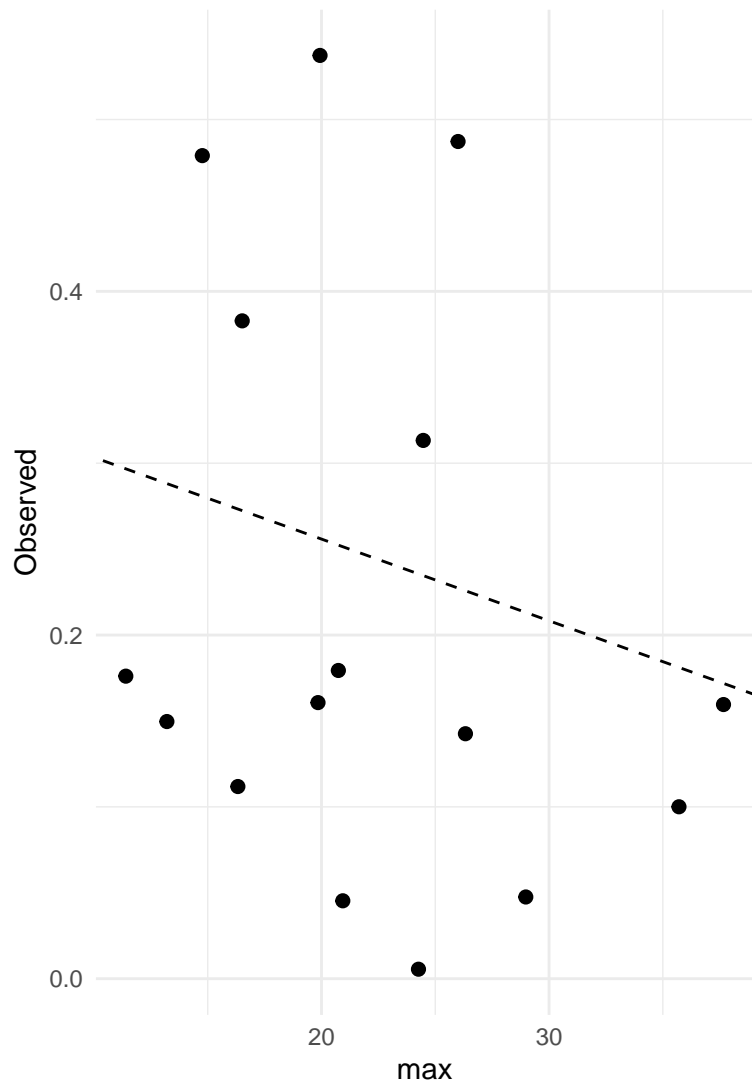
Mean_Upper Coefficients – PFI



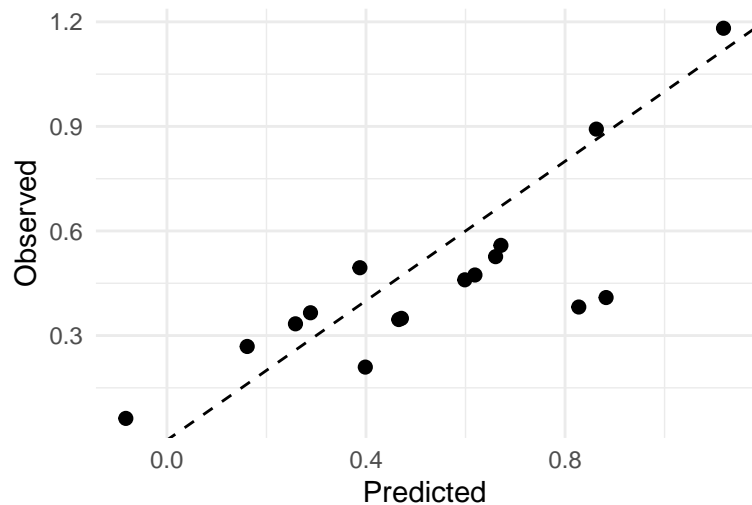
Ridge Coefficients – PFI



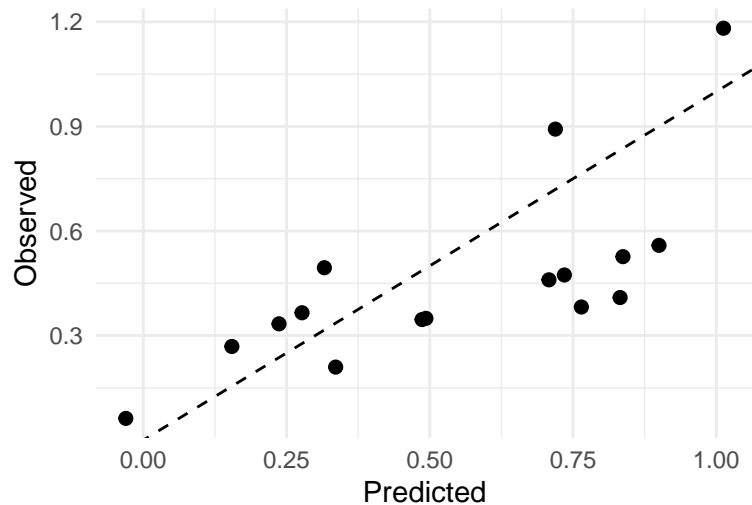


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{max}$ 

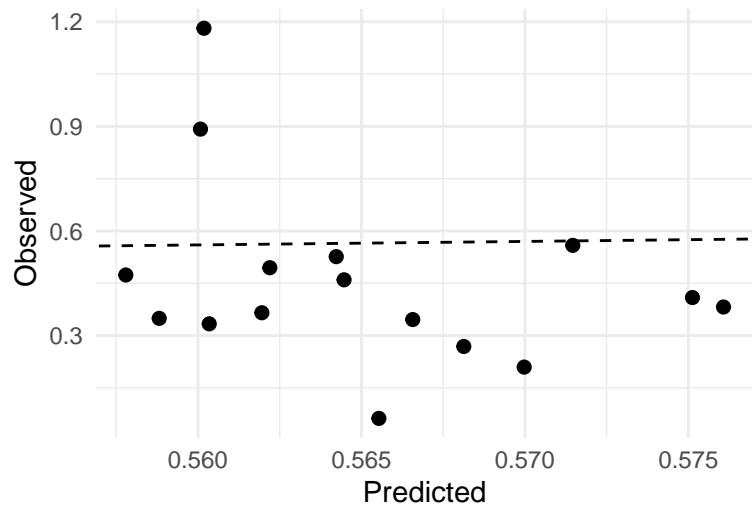
Full – APC



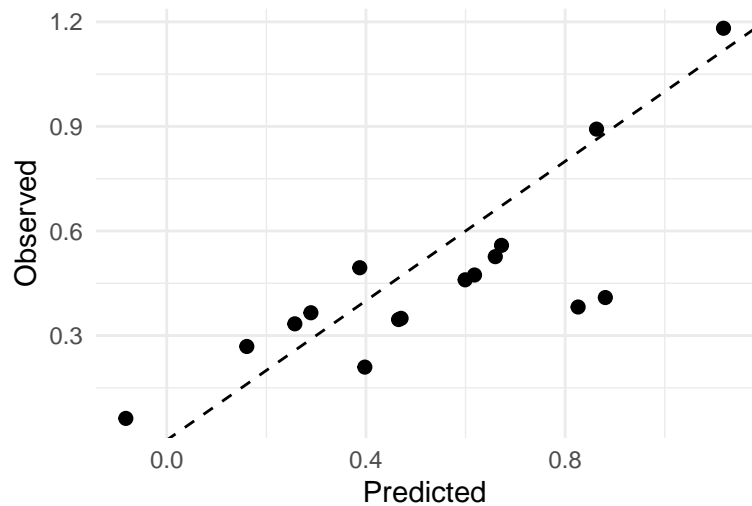
Range – APC



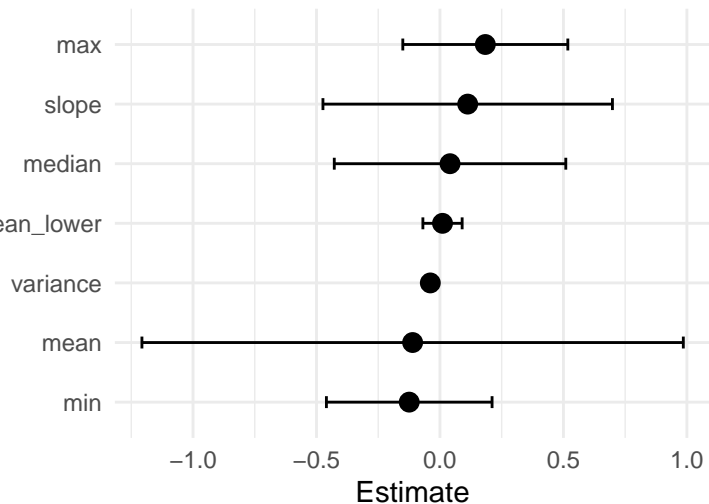
Mean_Upper – APC



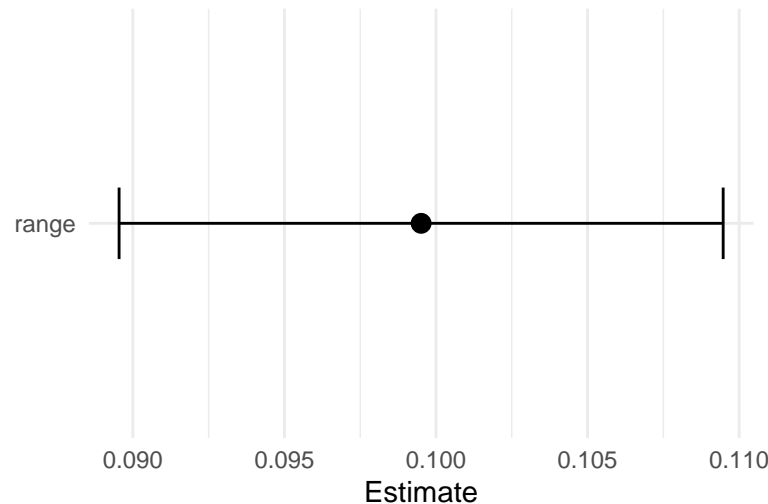
Ridge – APC



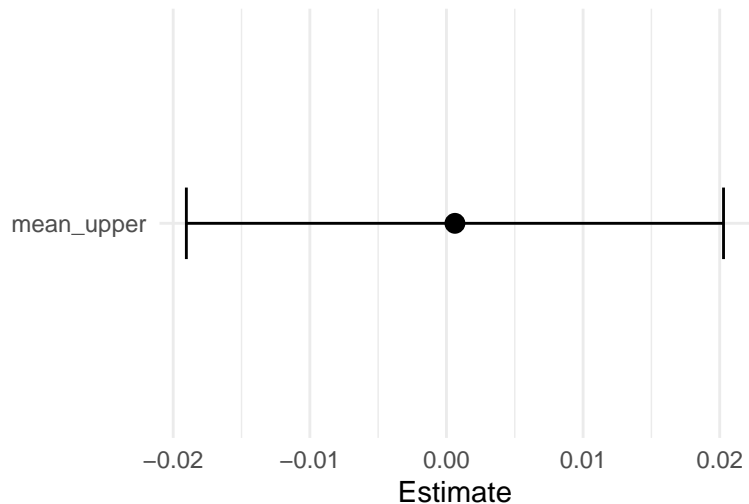
Full Coefficients – APC



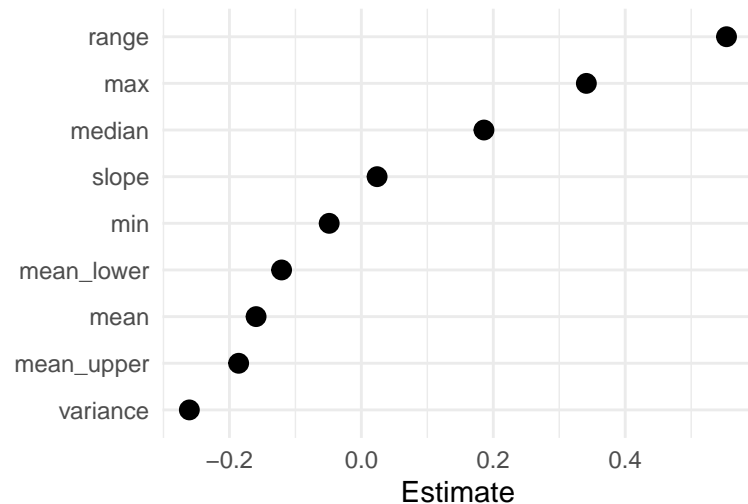
Range Coefficients – APC

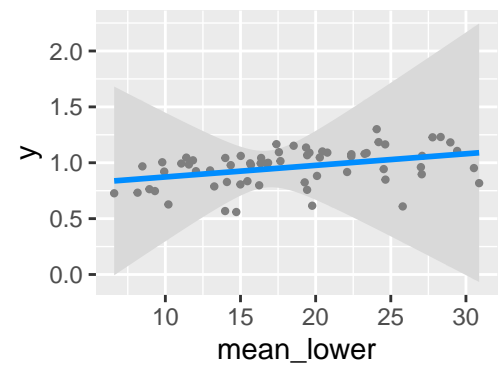
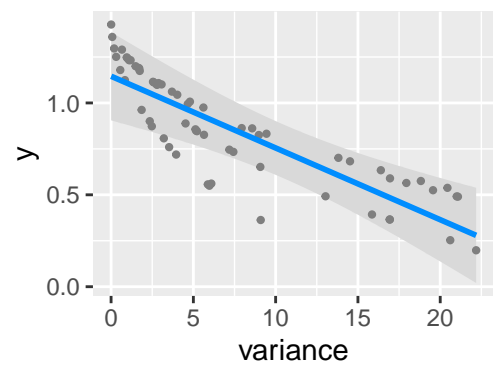
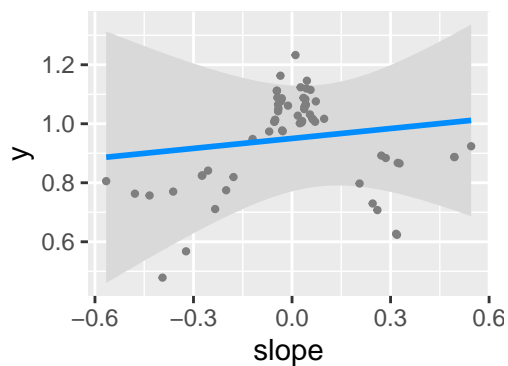
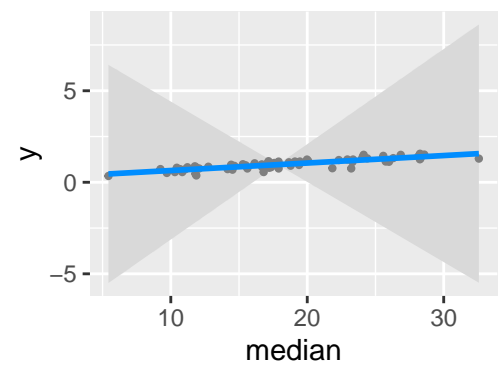
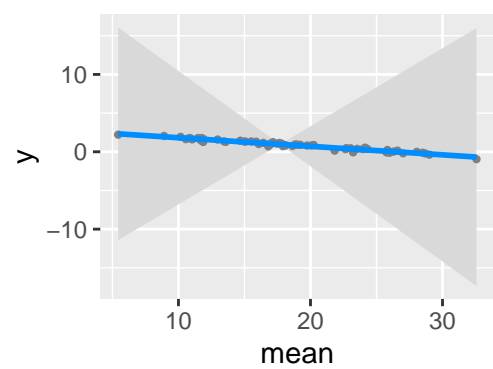
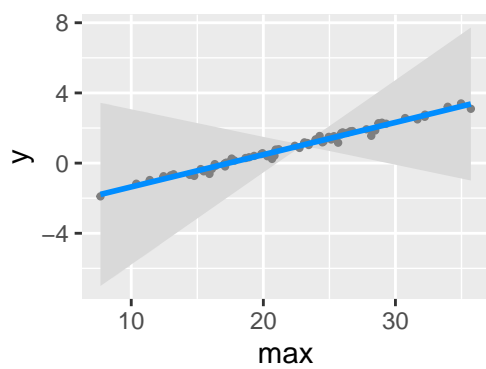
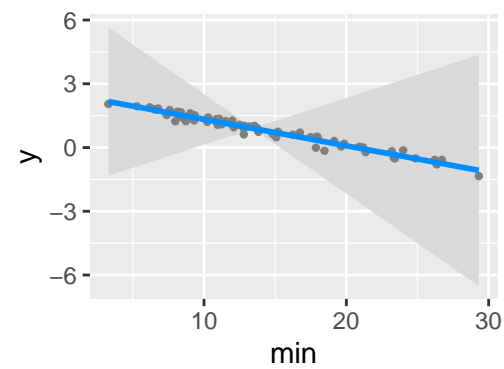


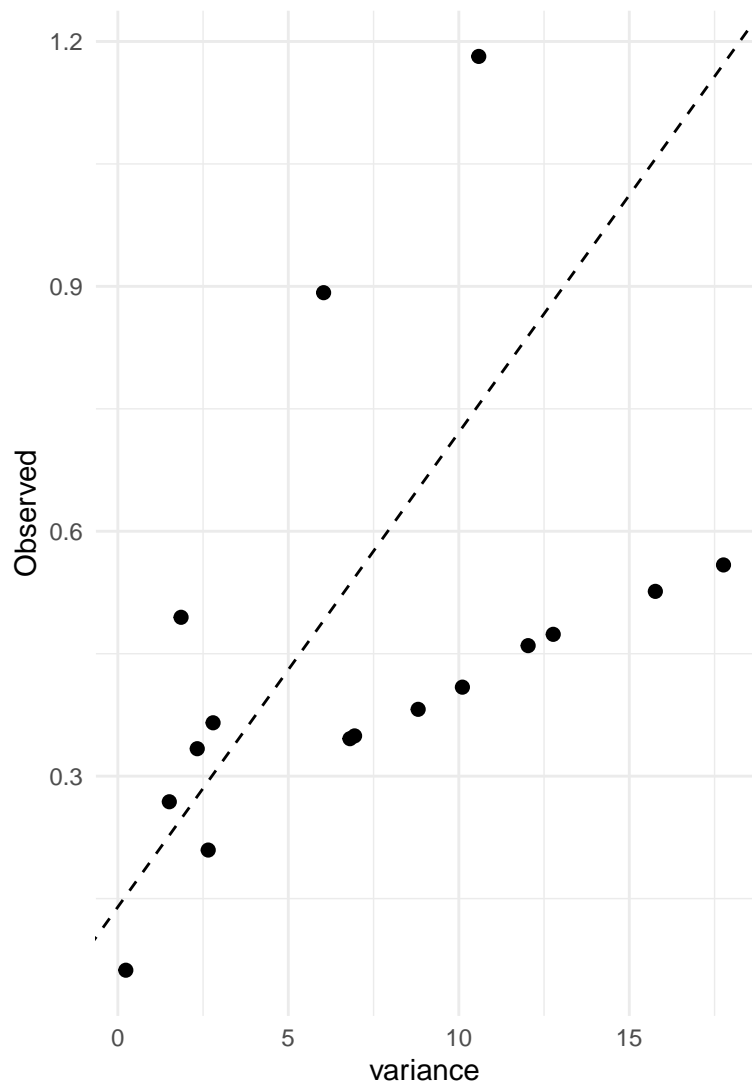
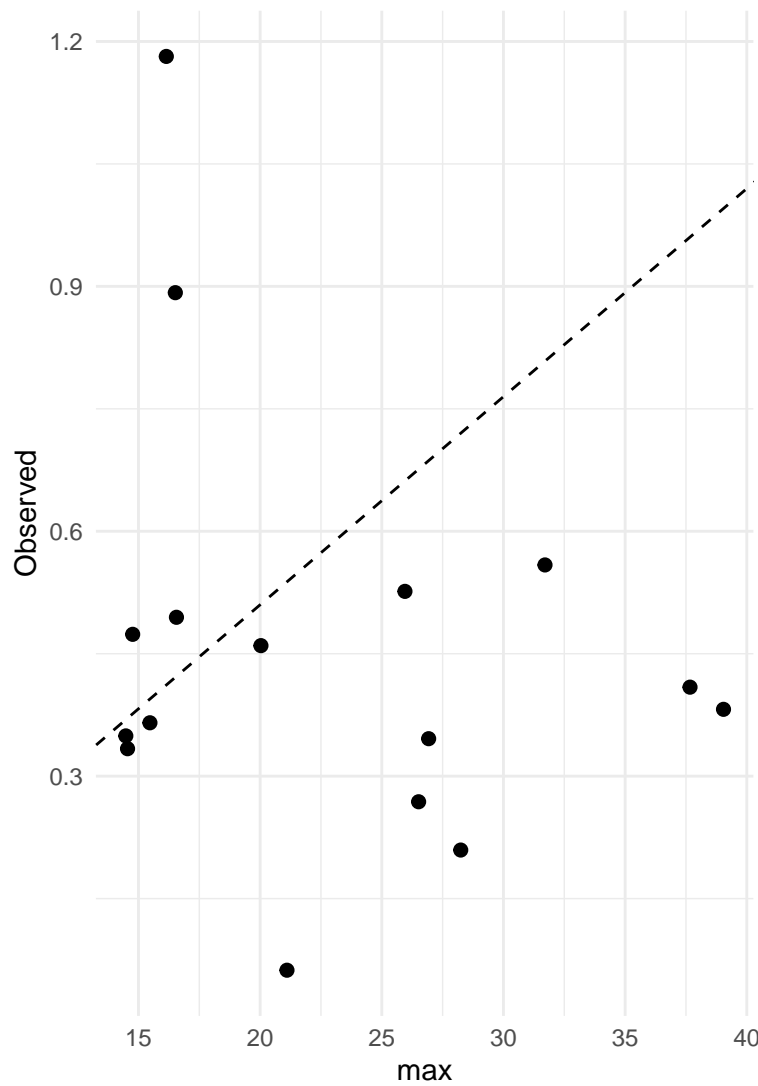
Mean_Upper Coefficients – APC



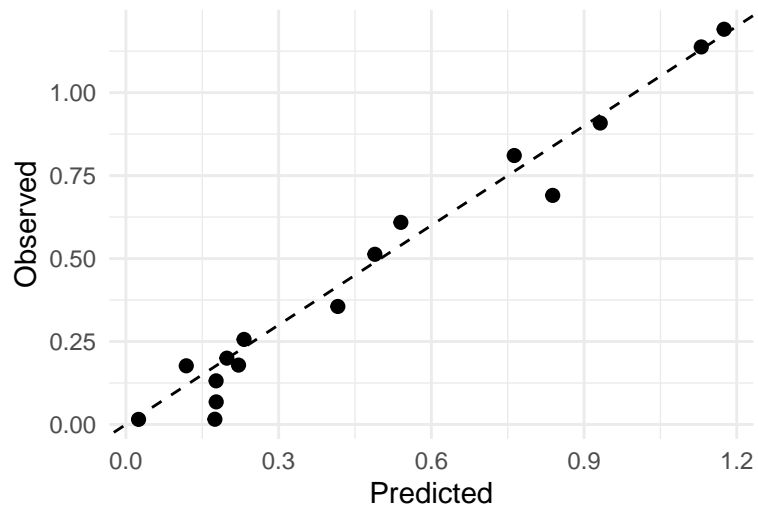
Ridge Coefficients – APC



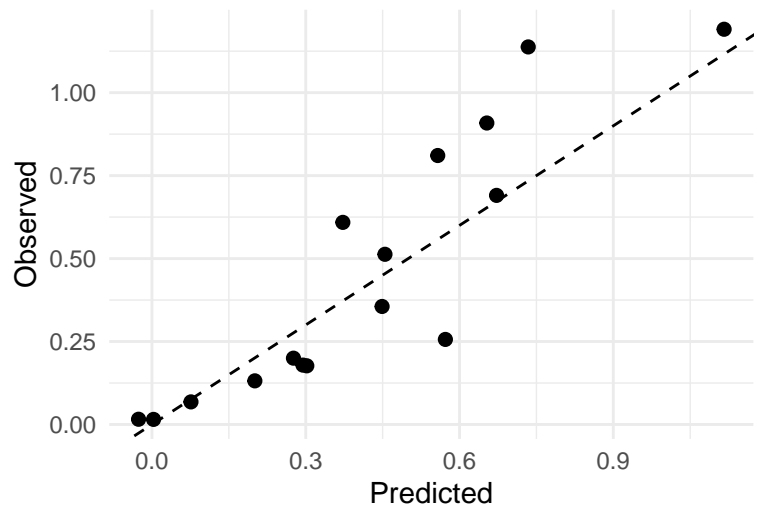


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{max}$ 

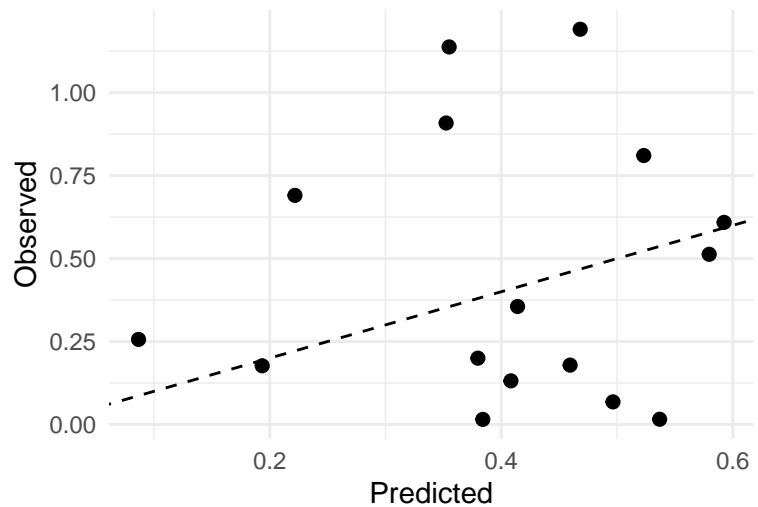
Full – SI



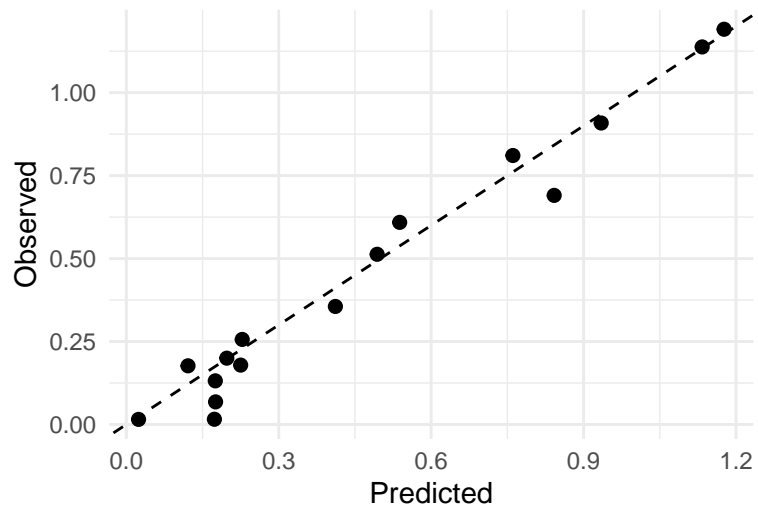
Range – SI



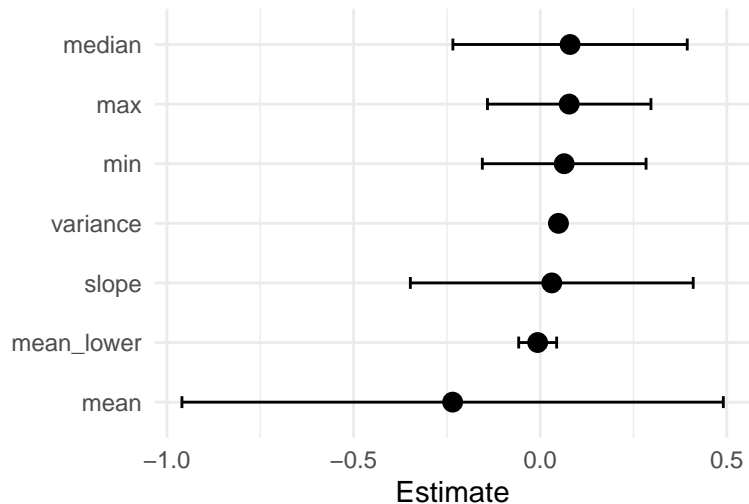
Mean_Upper – SI



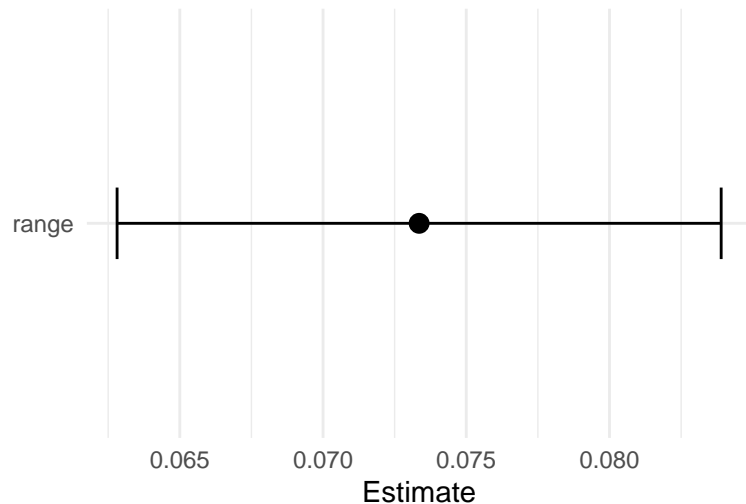
Ridge – SI



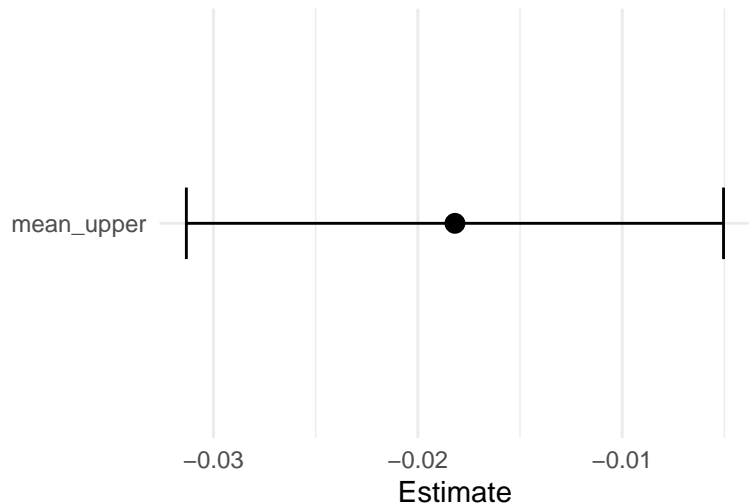
Full Coefficients – SI



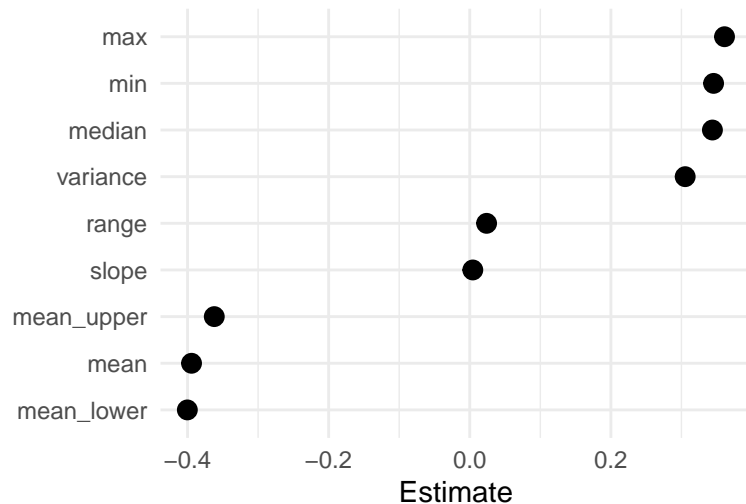
Range Coefficients – SI

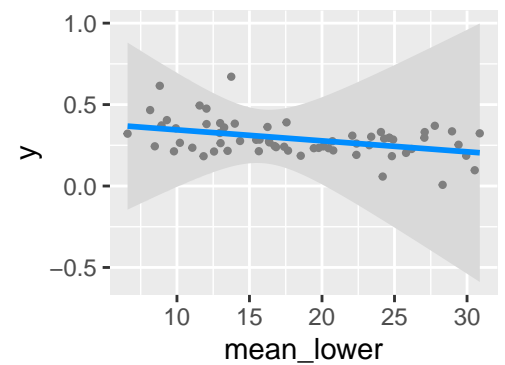
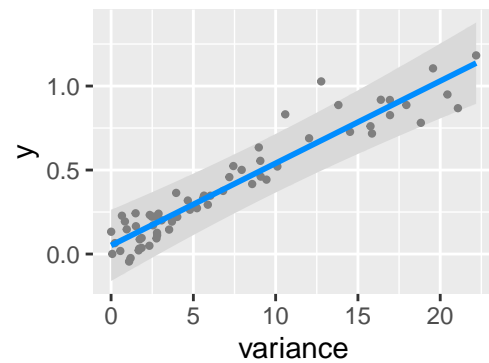
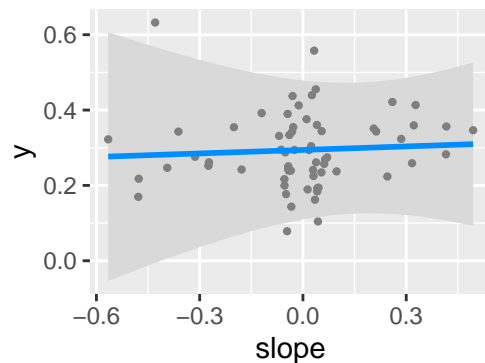
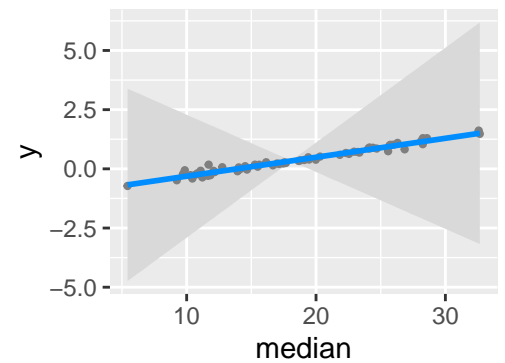
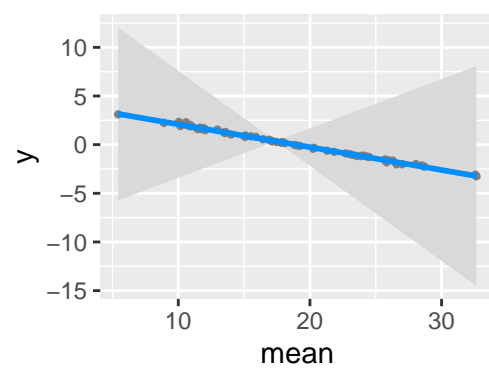
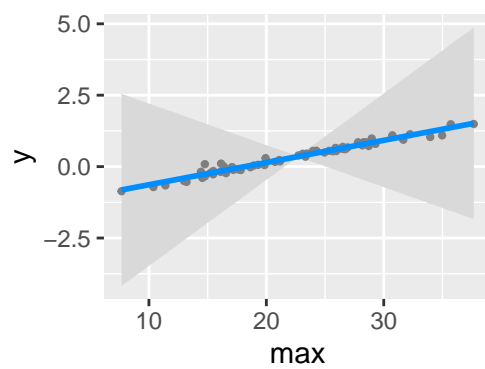
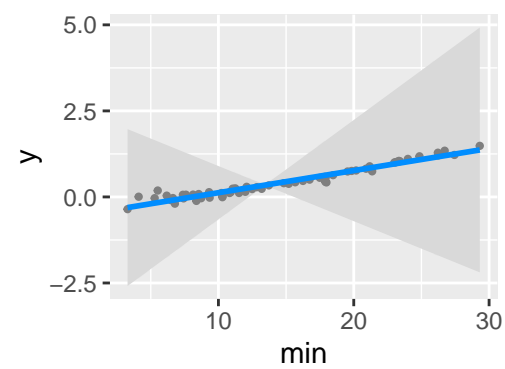


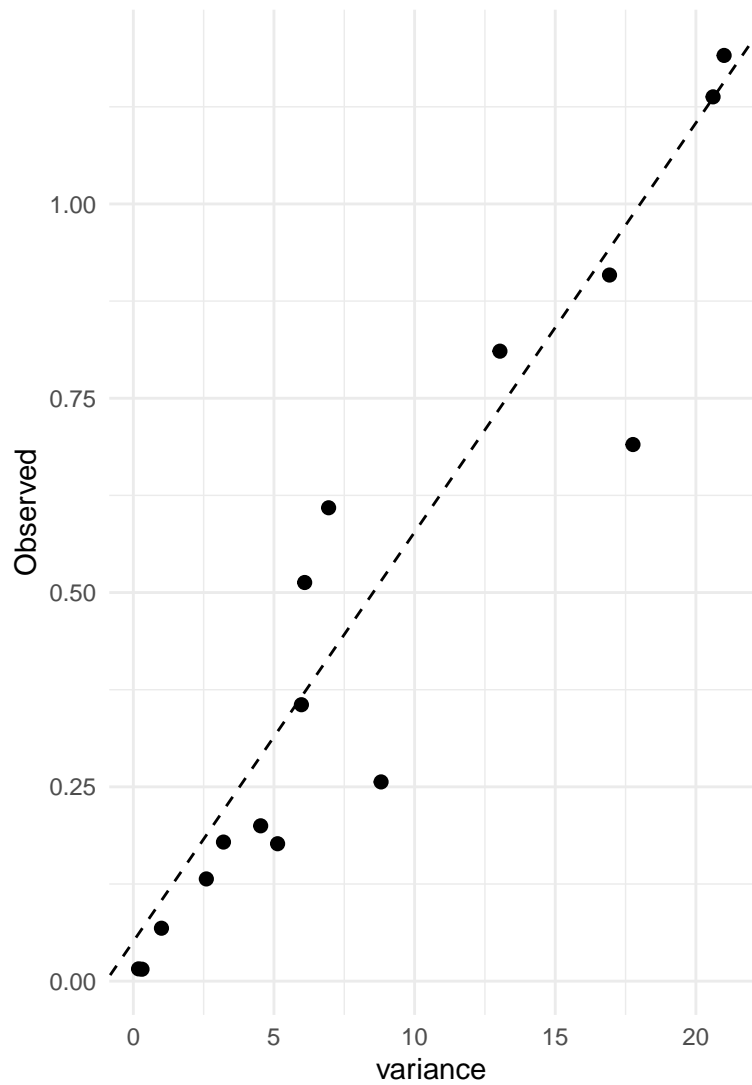
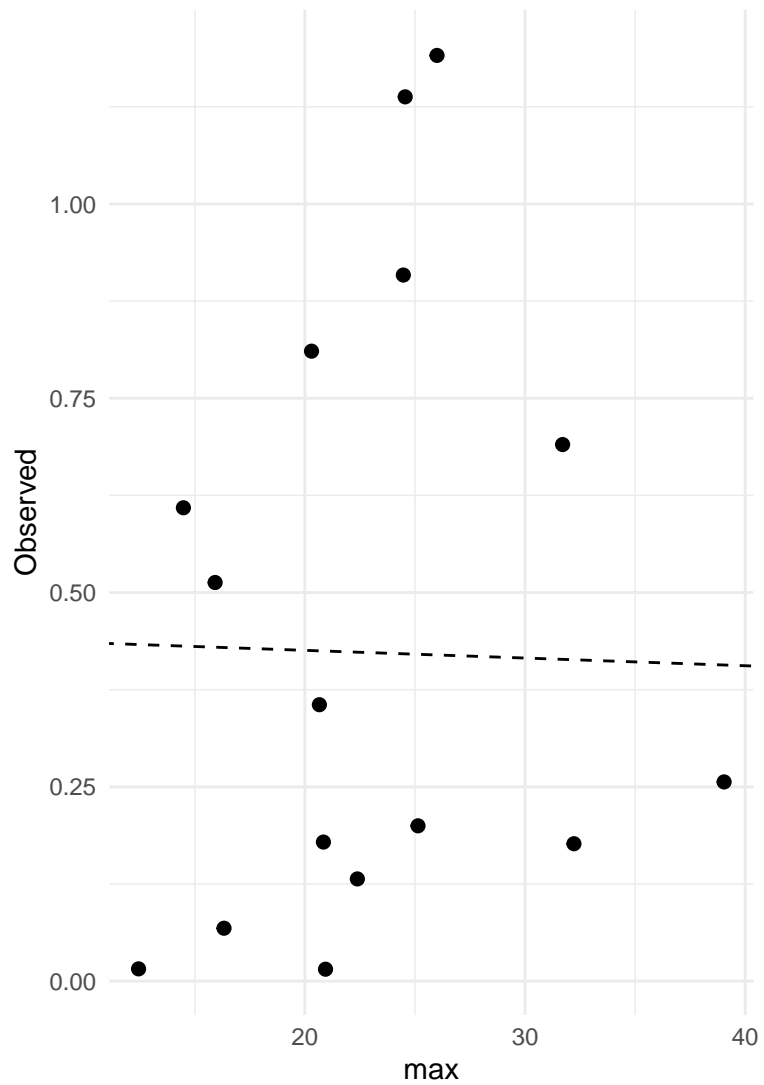
Mean_Upper Coefficients – SI



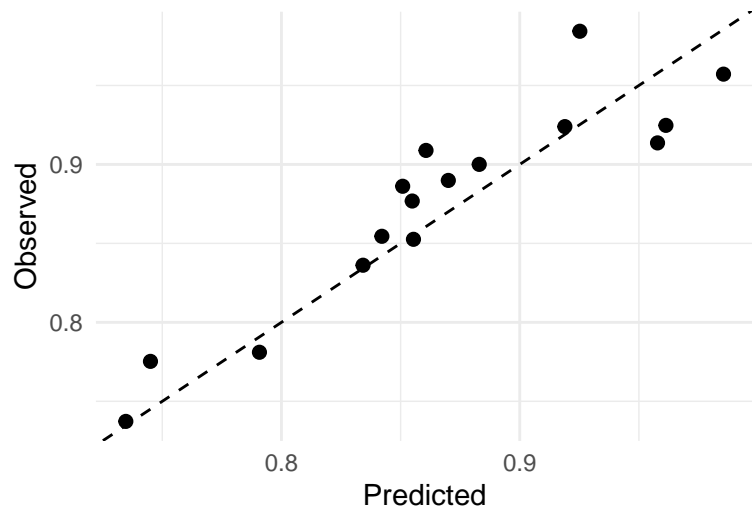
Ridge Coefficients – SI



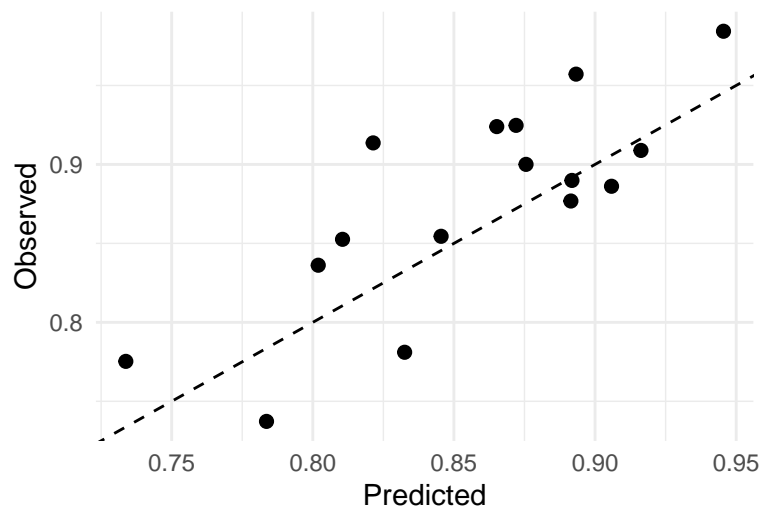


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{max}$ 

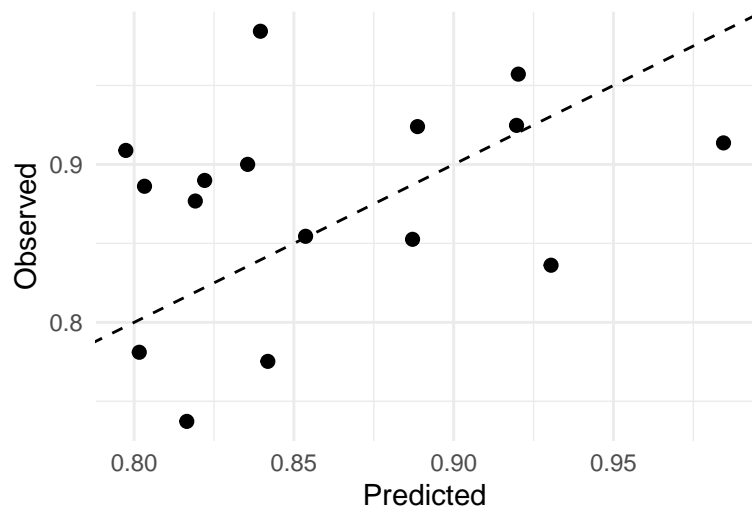
Full – RSI



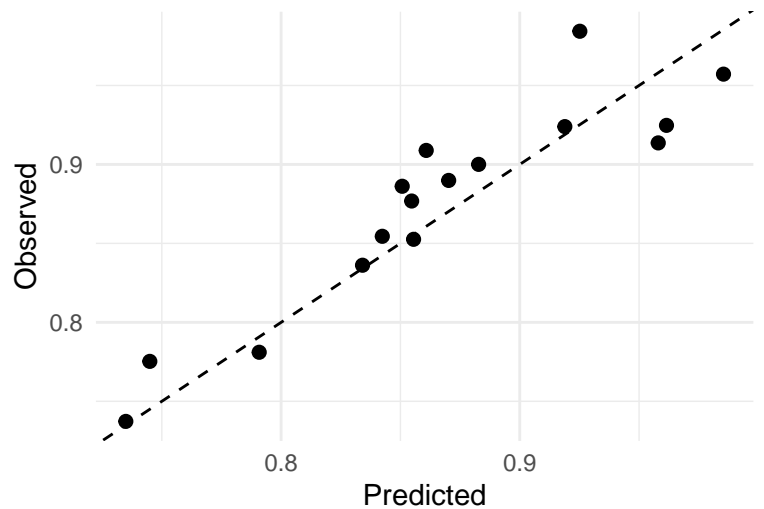
Range – RSI



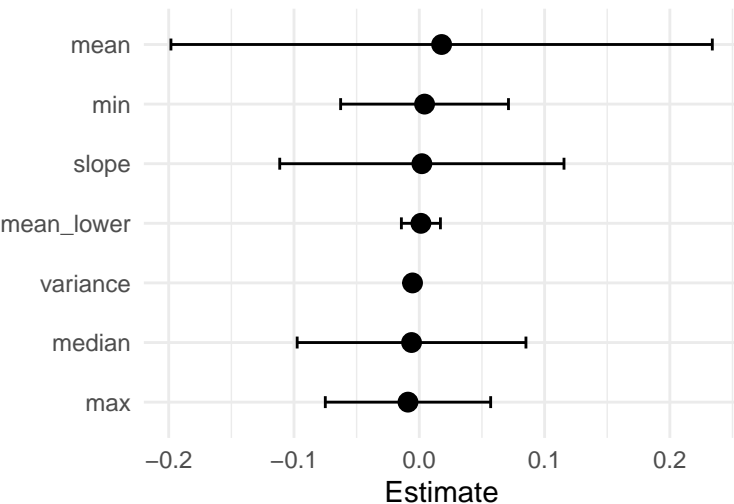
Mean_Upper – RSI



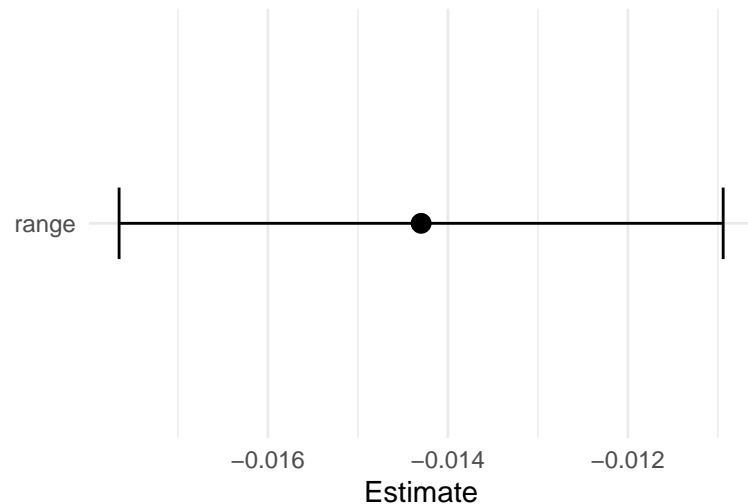
Ridge – RSI



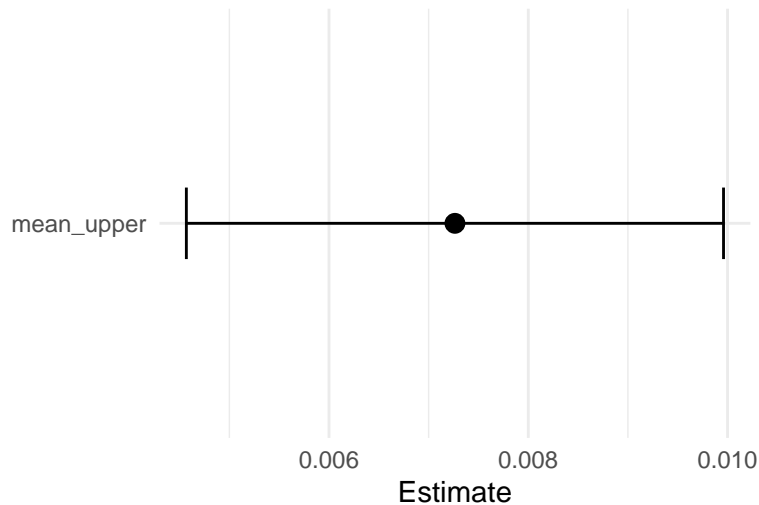
Full Coefficients – RSI



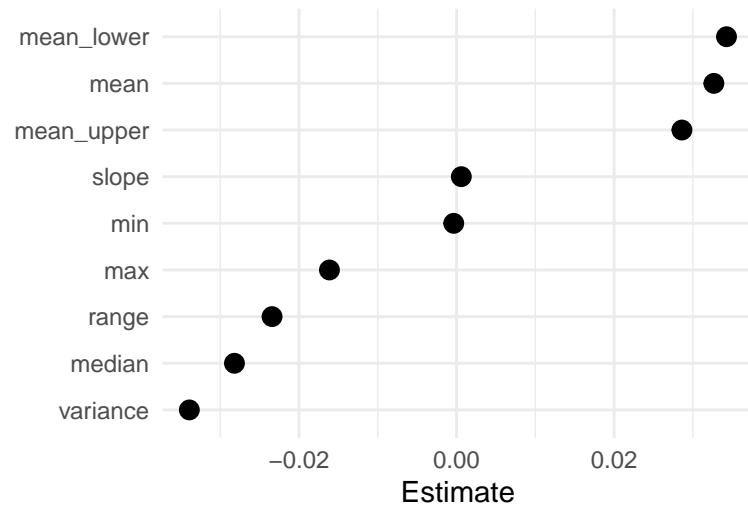
Range Coefficients – RSI

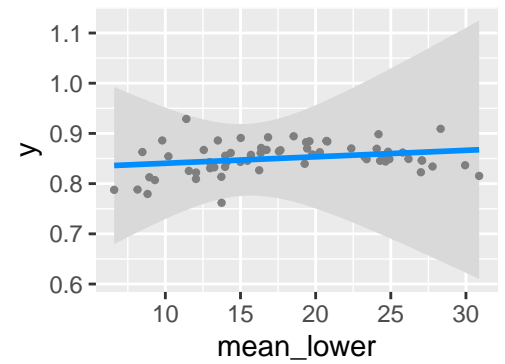
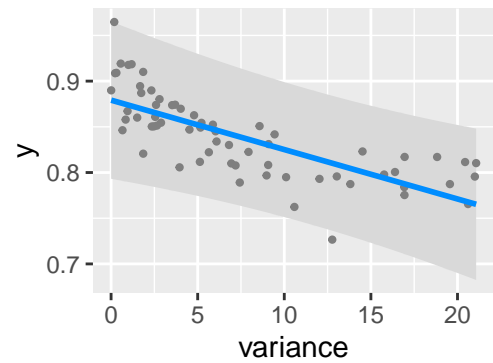
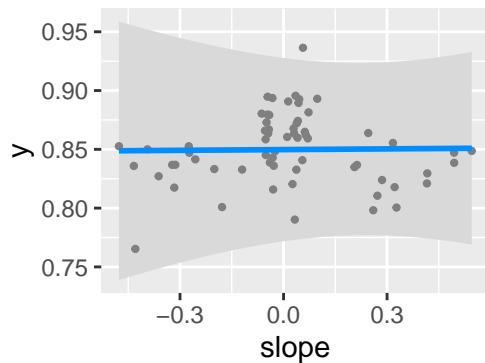
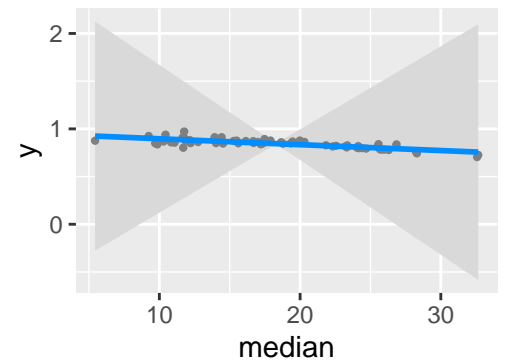
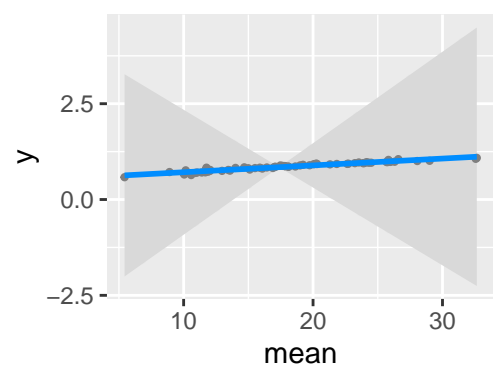
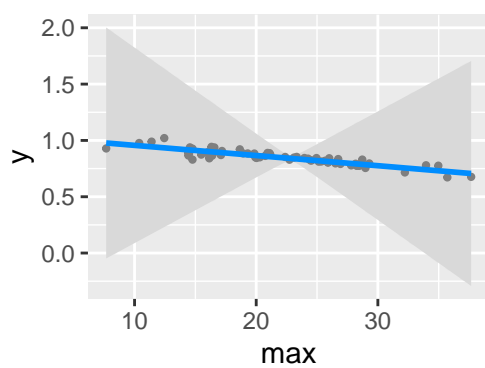
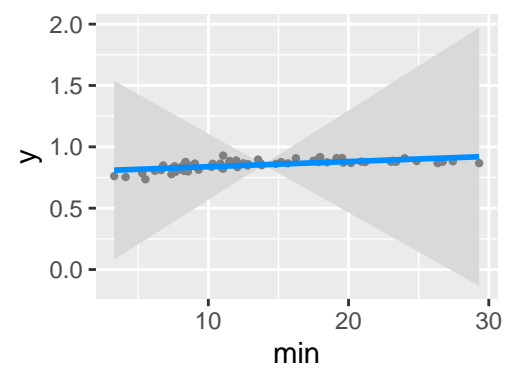


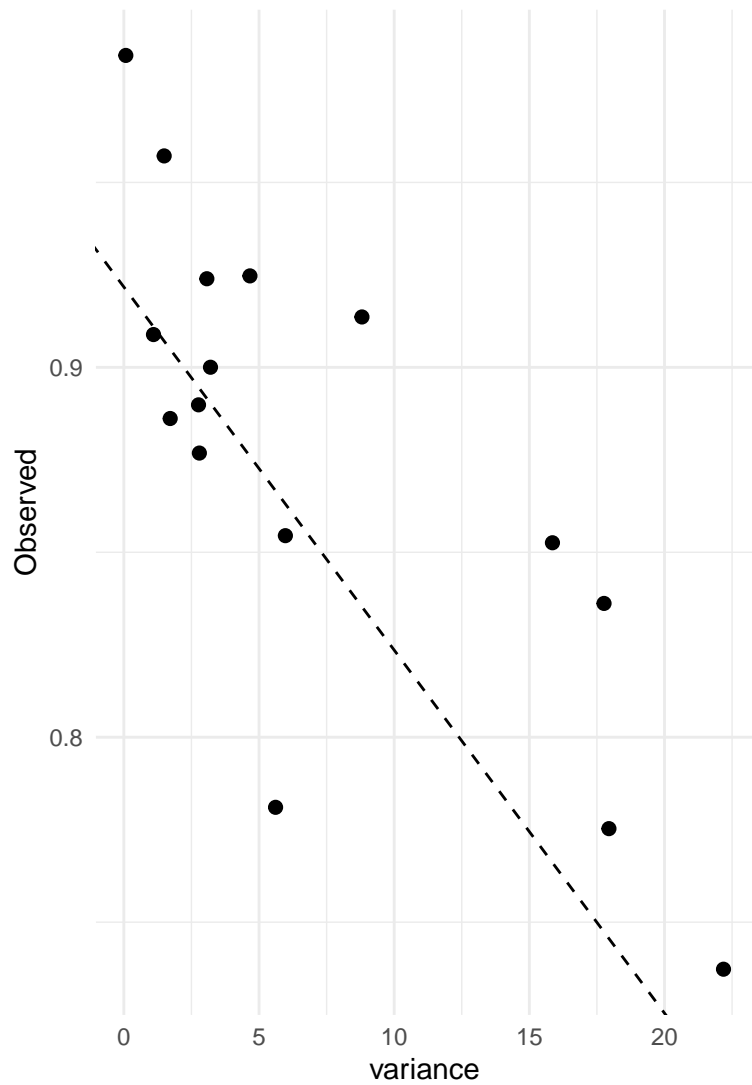
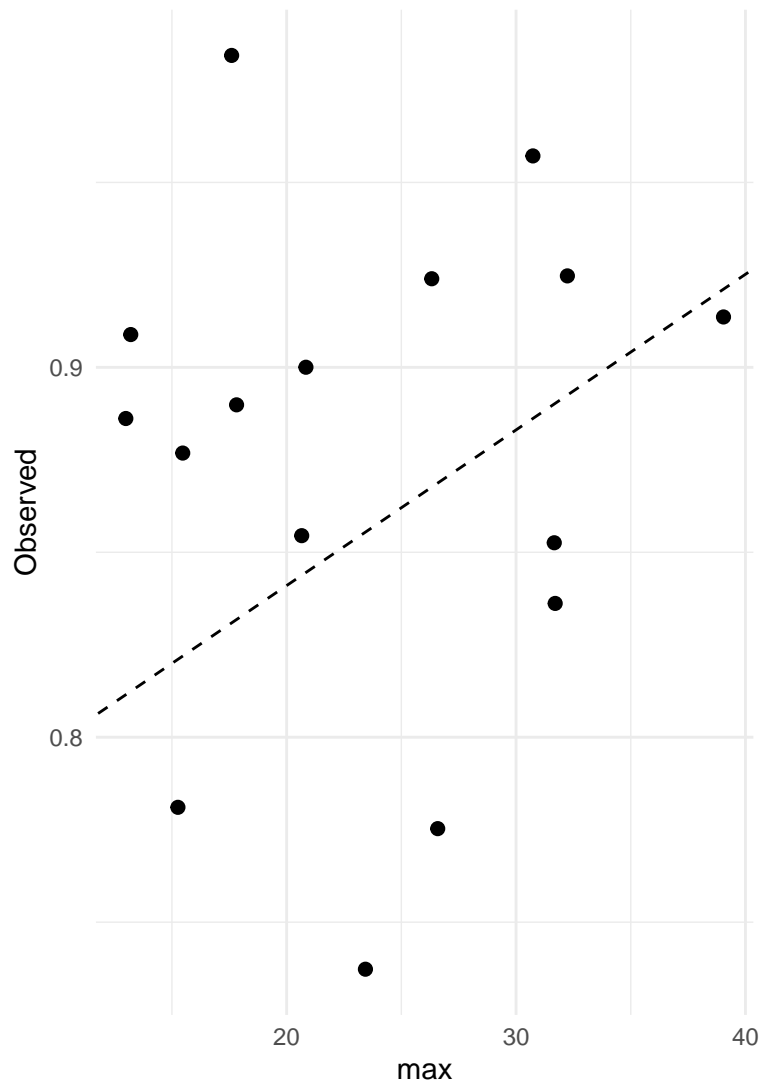
Mean_Upper Coefficients – RSI



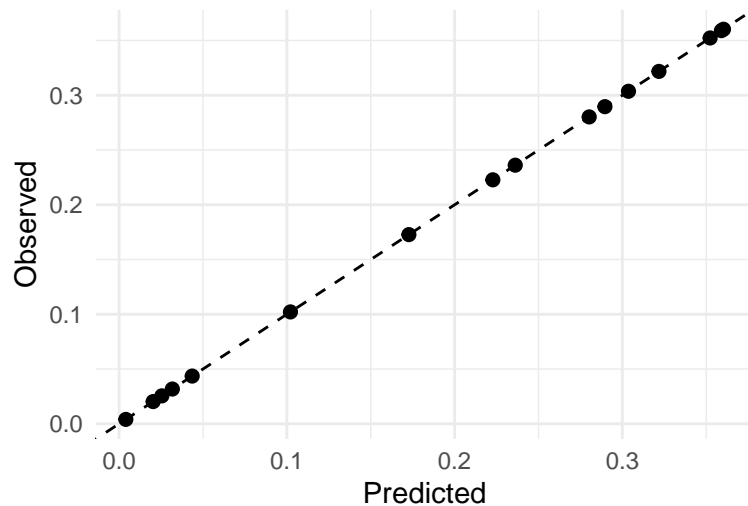
Ridge Coefficients – RSI



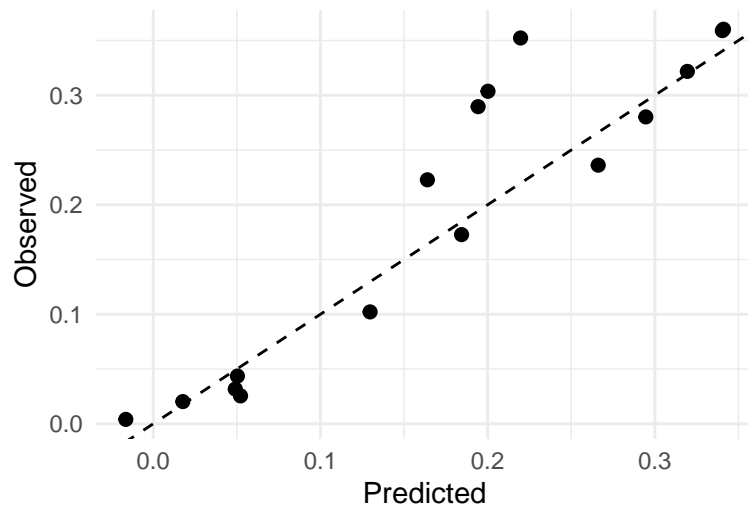


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{max}$ 

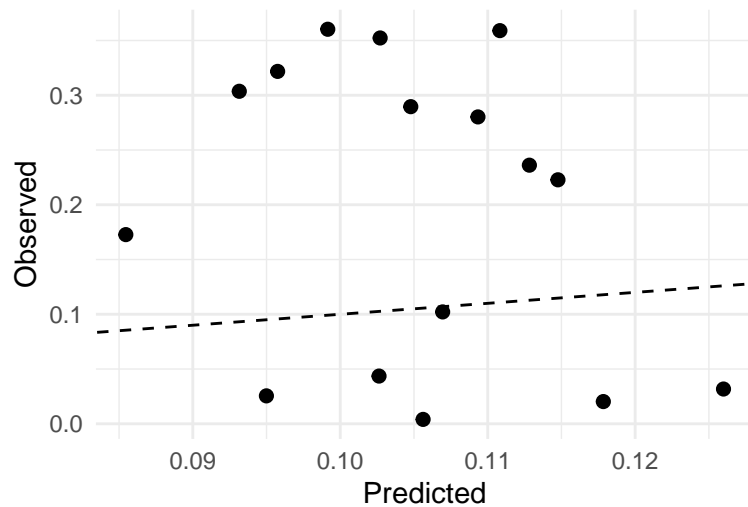
Full – EVS



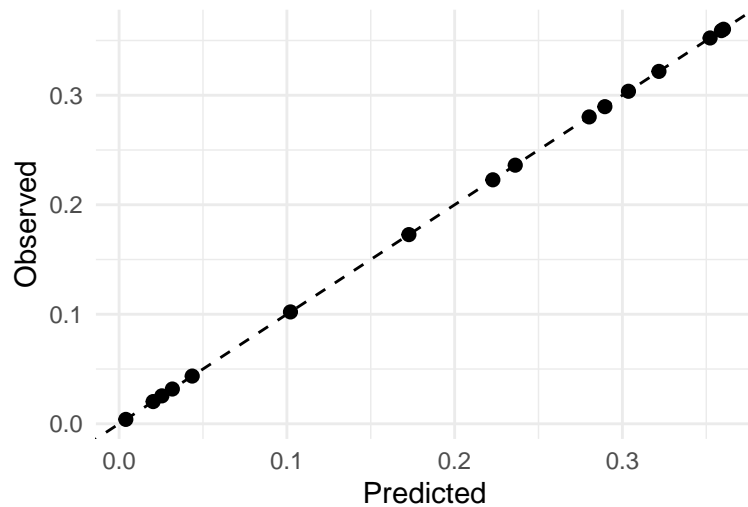
Range – EVS



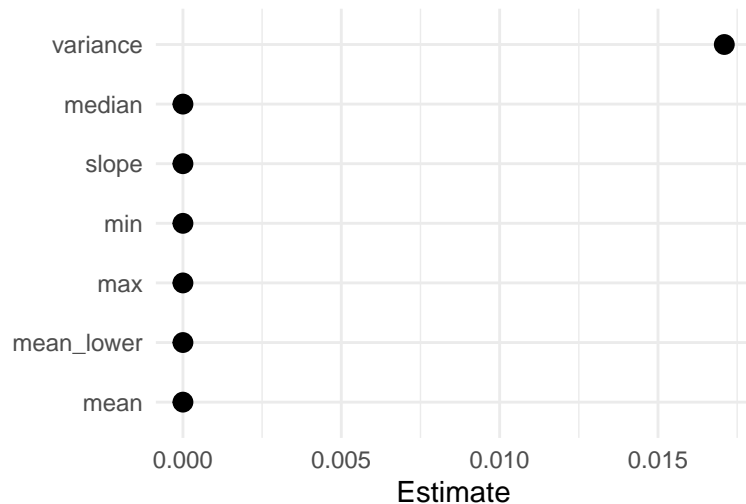
Mean_Upper – EVS



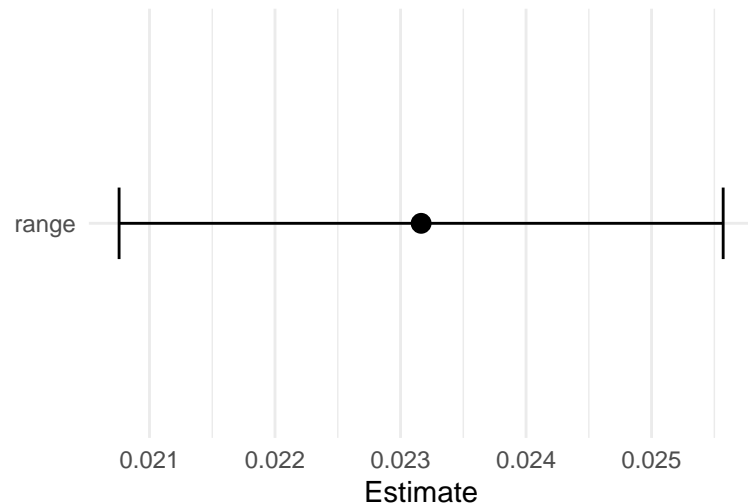
Ridge – EVS



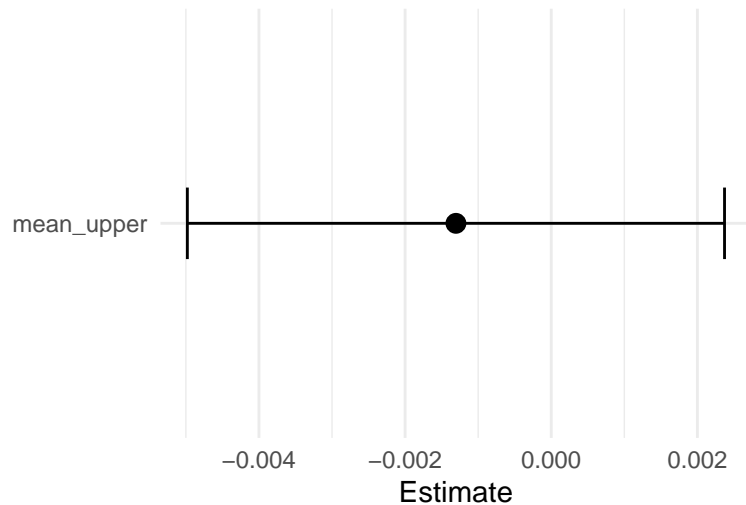
Full Coefficients – EVS



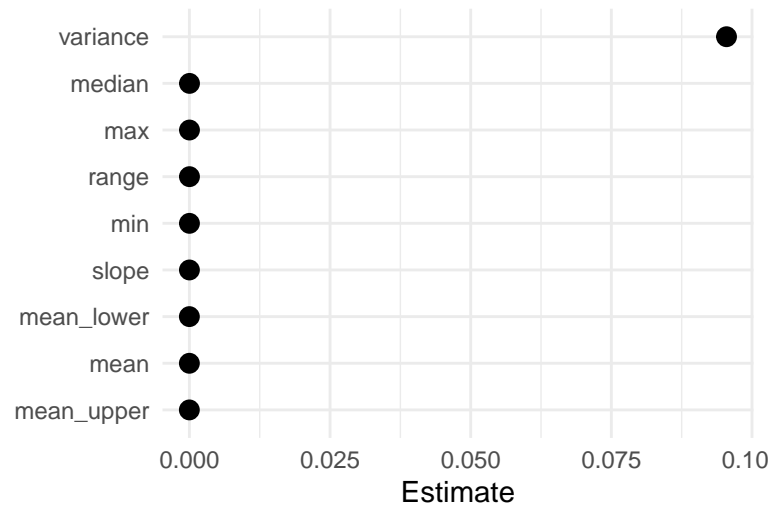
Range Coefficients – EVS

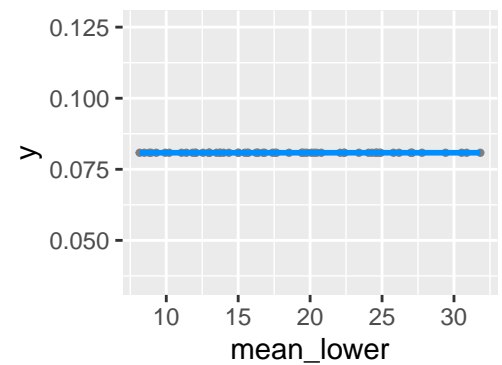
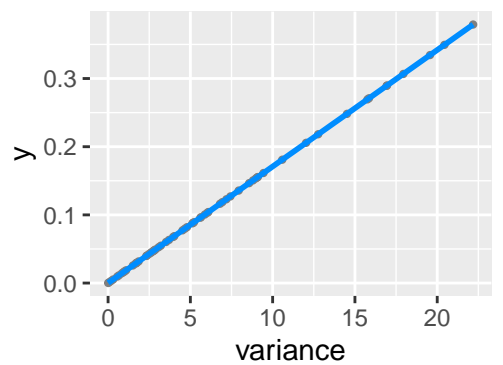
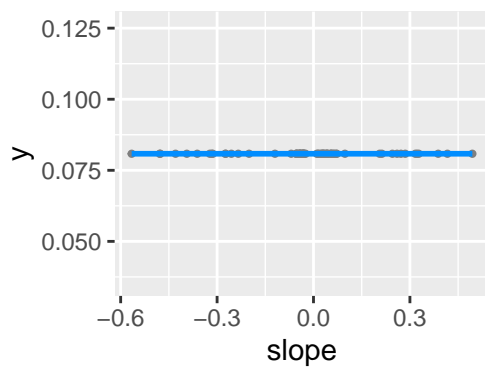
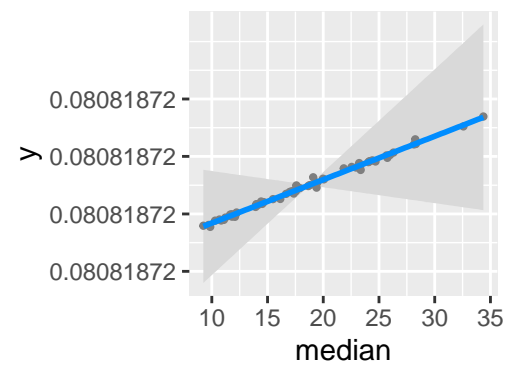
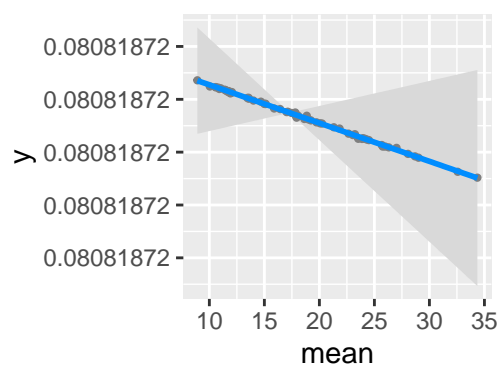
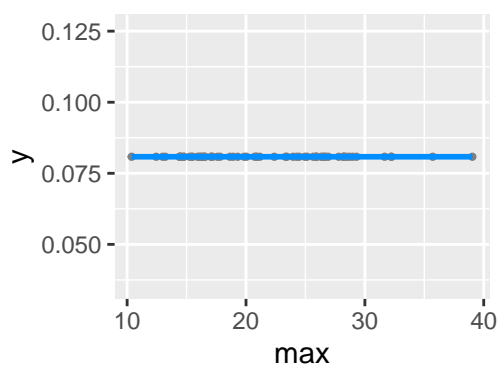
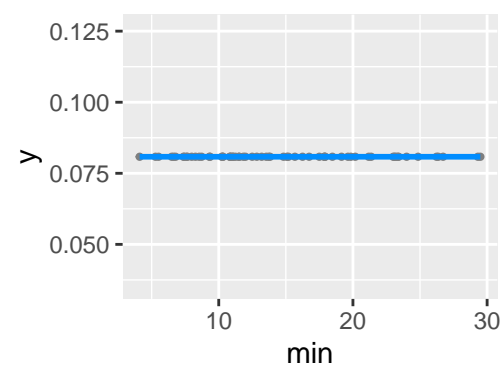


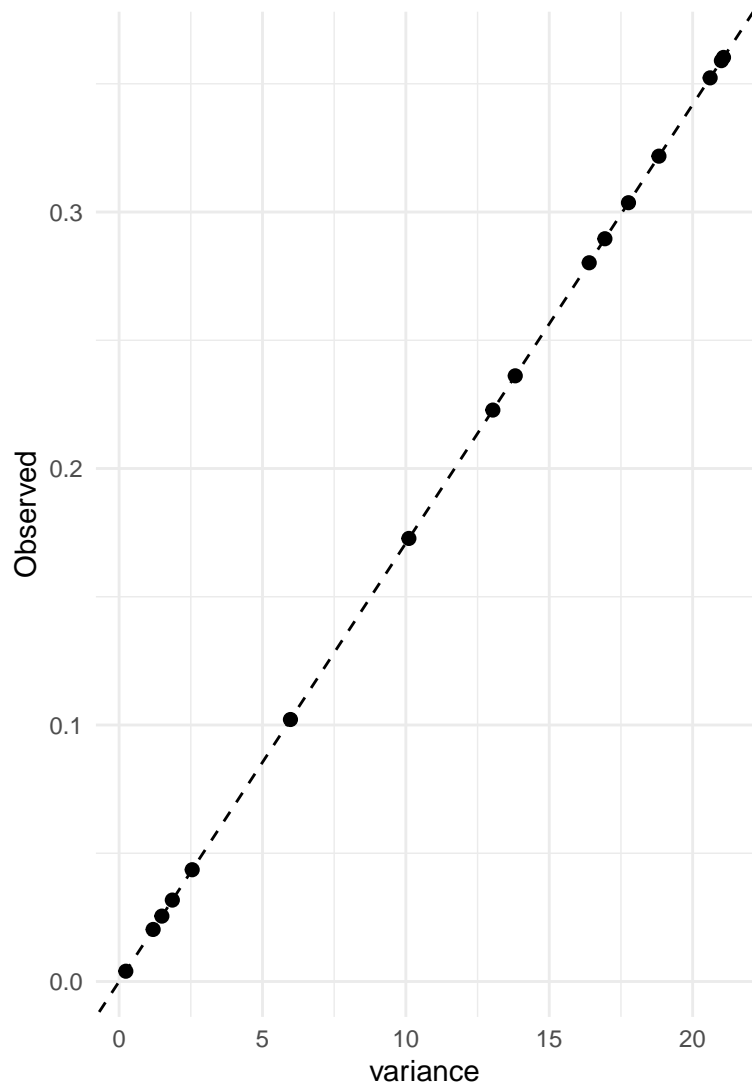
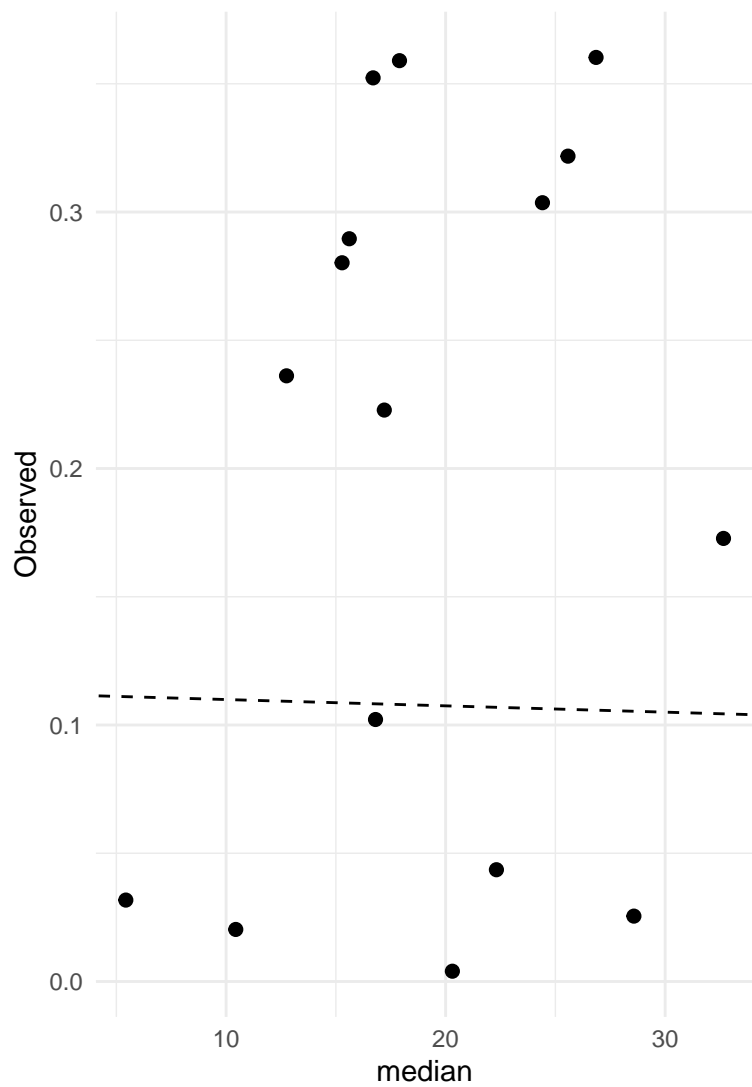
Mean_Upper Coefficients – EVS

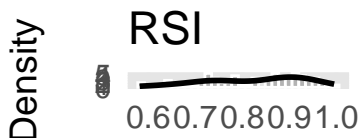
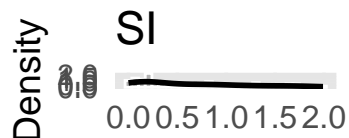
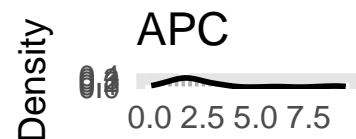
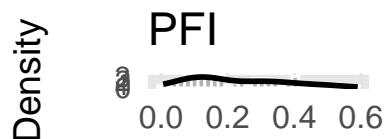
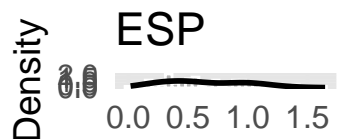
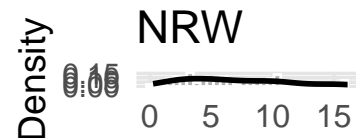
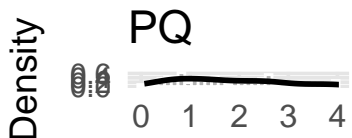
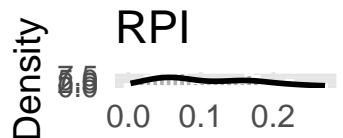
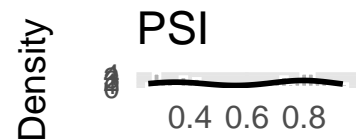
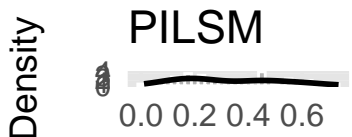
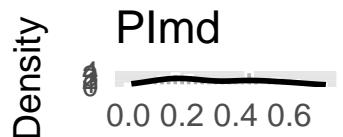
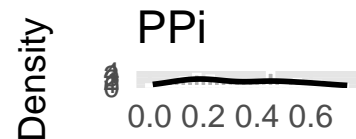
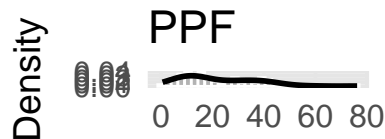
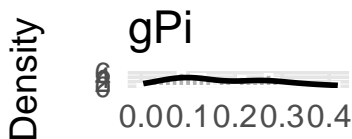
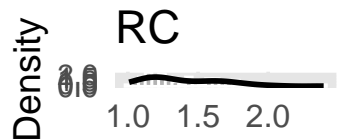
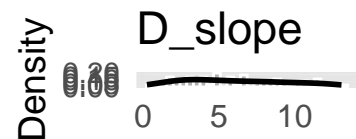
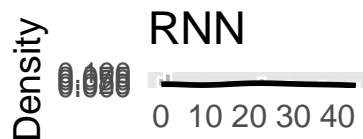
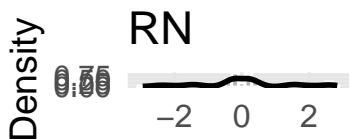
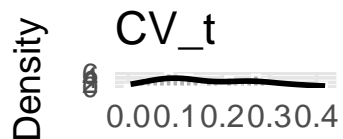


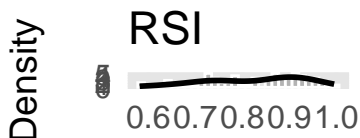
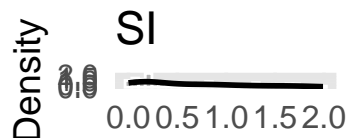
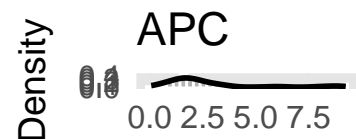
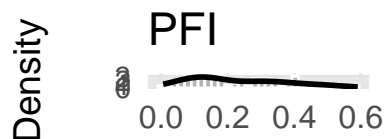
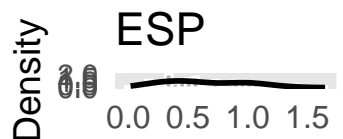
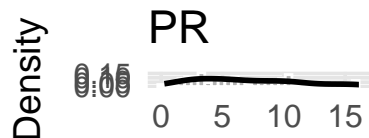
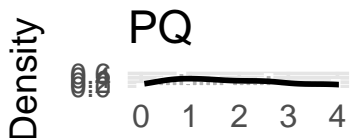
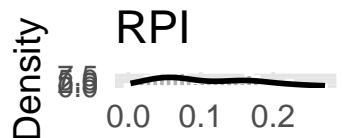
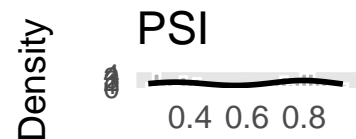
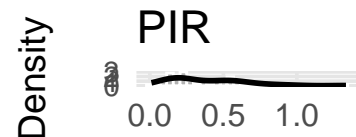
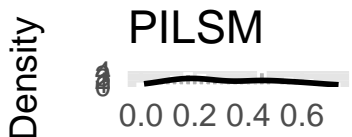
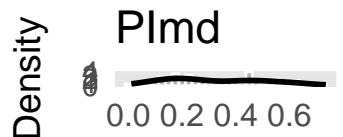
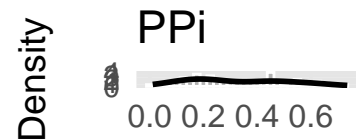
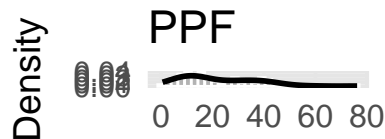
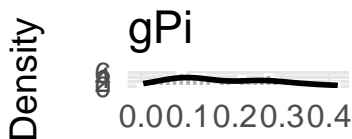
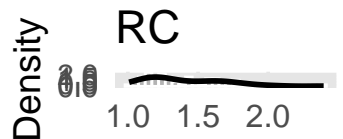
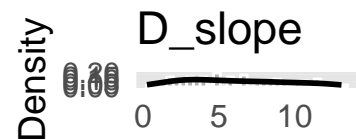
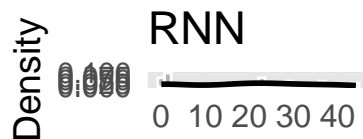
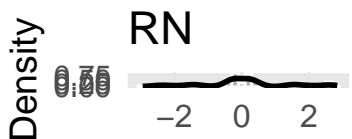
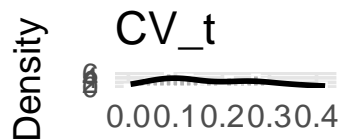
Ridge Coefficients – EVS



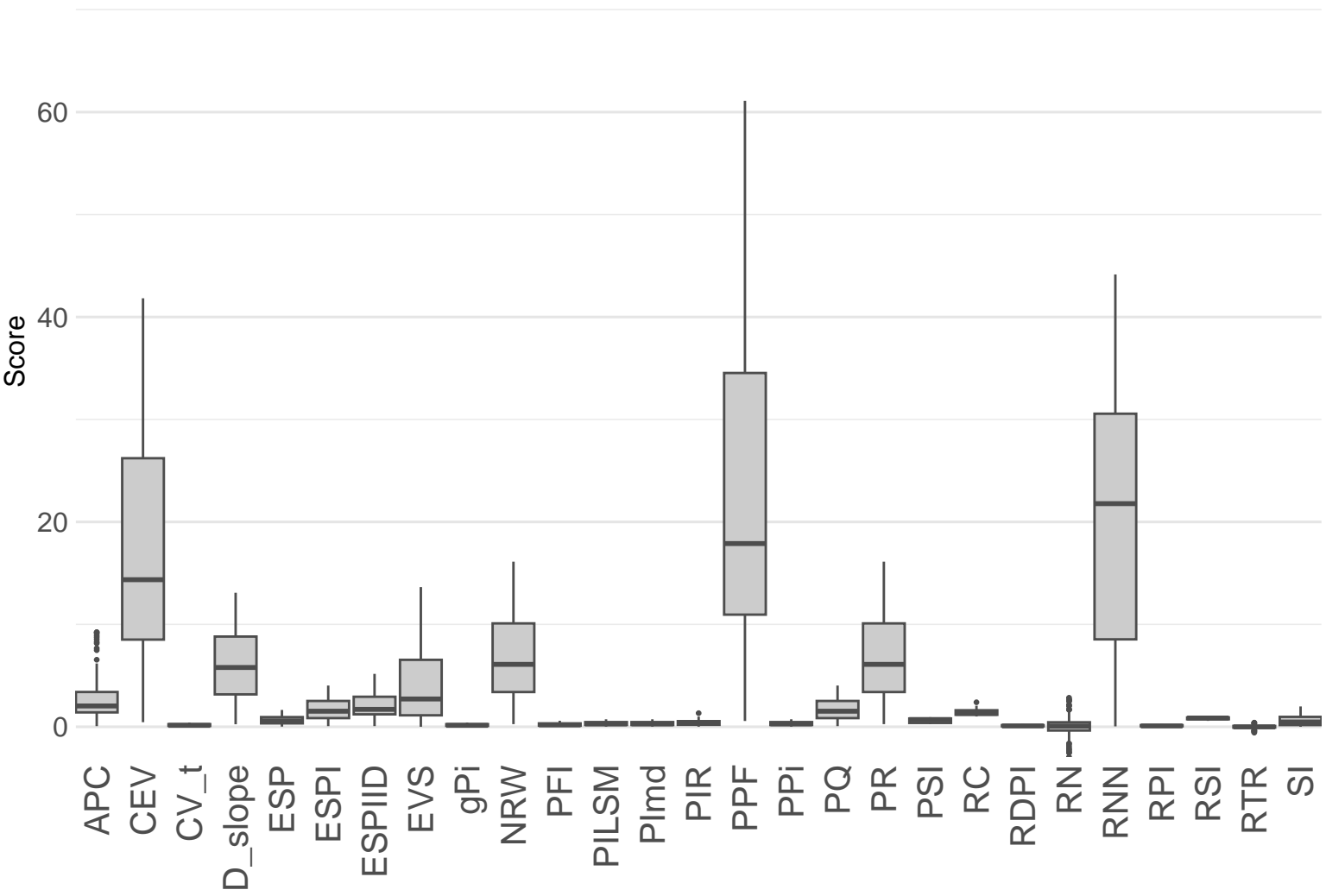


Simple Model: $y \sim \text{variance}$ Simple Model: $y \sim \text{median}$ 

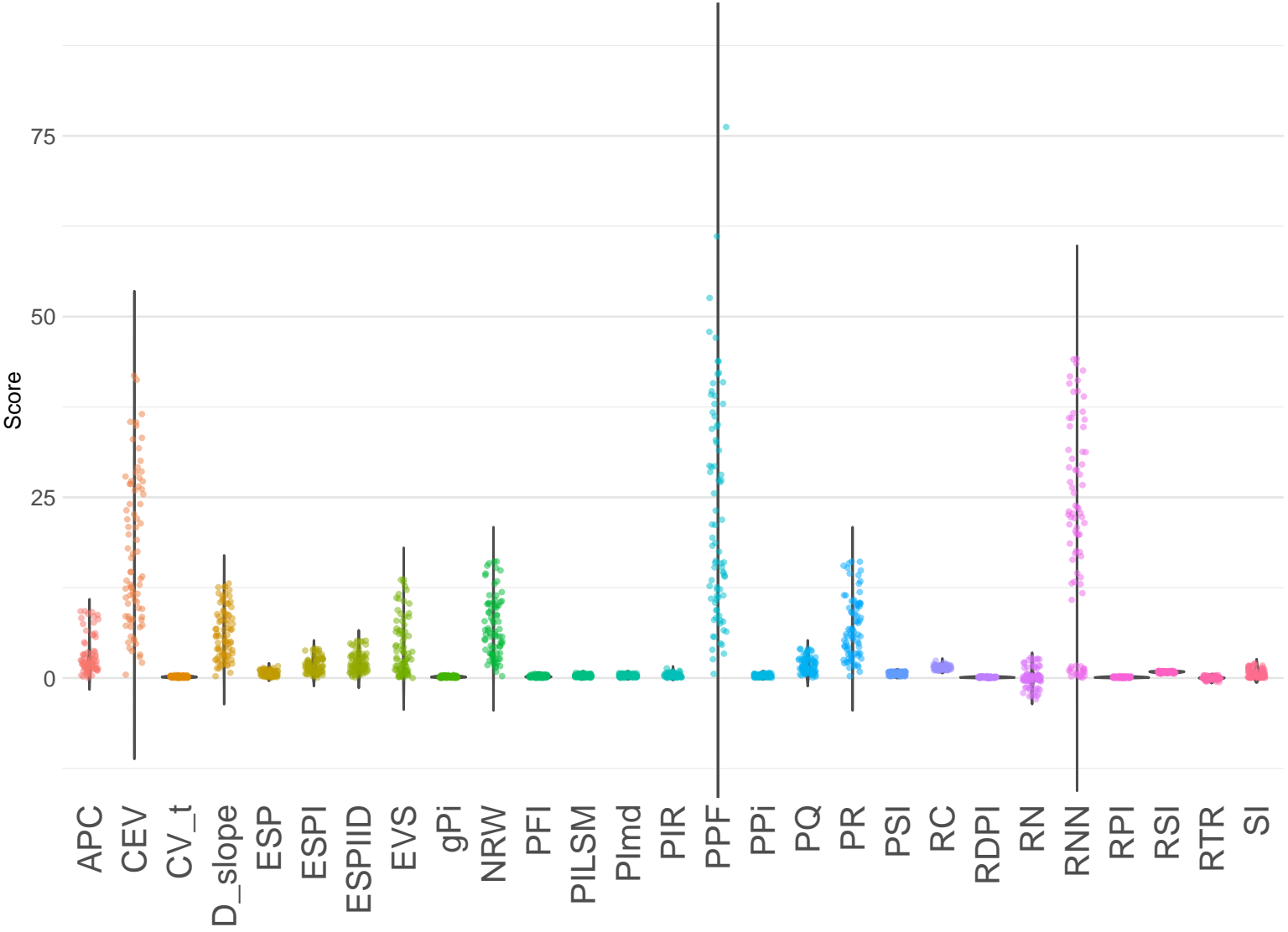




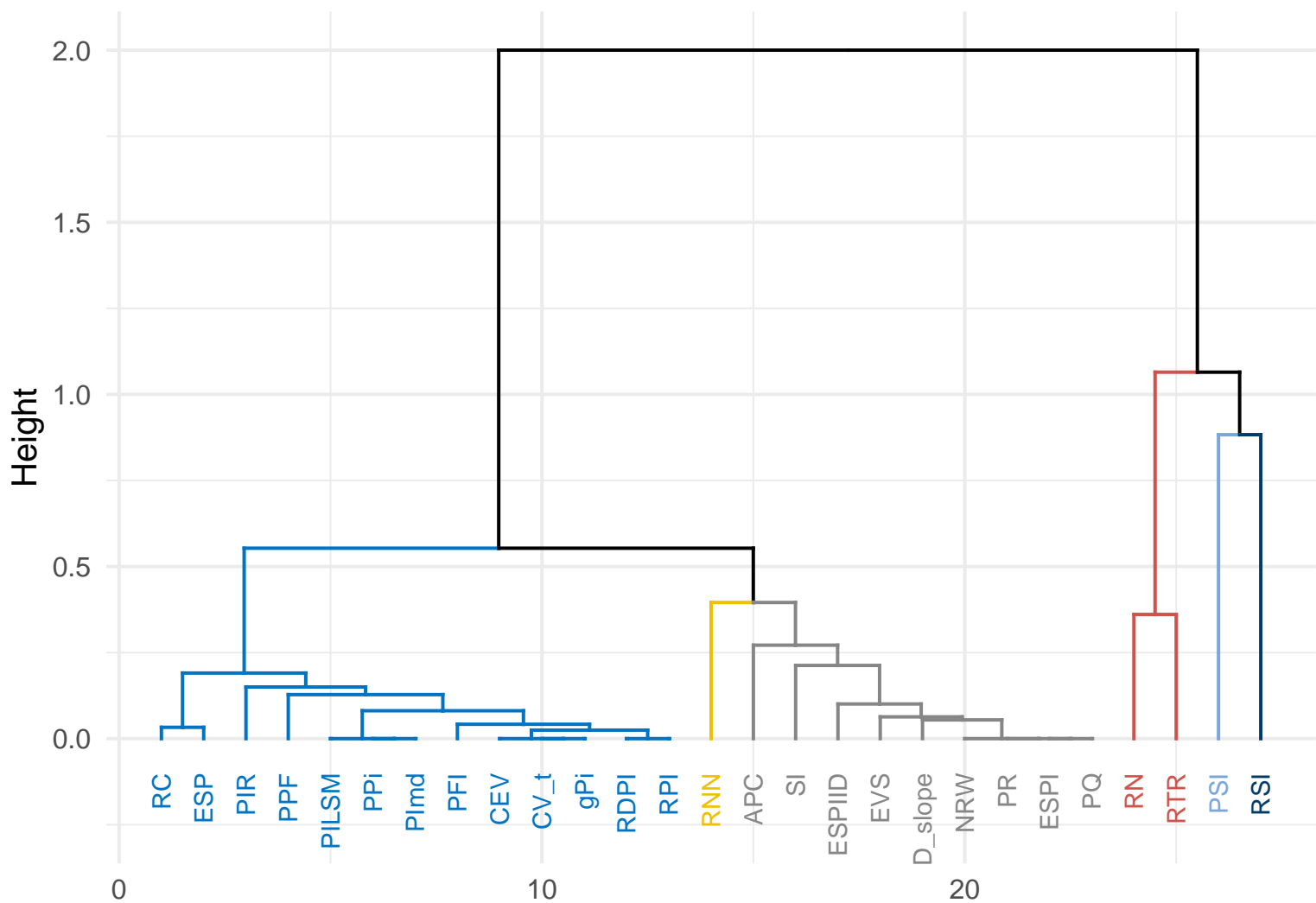
All Metrics Boxplot



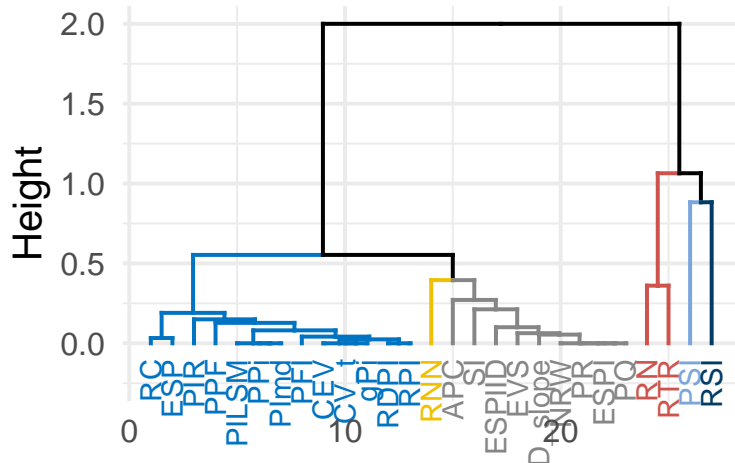
All Metrics Violin Plot



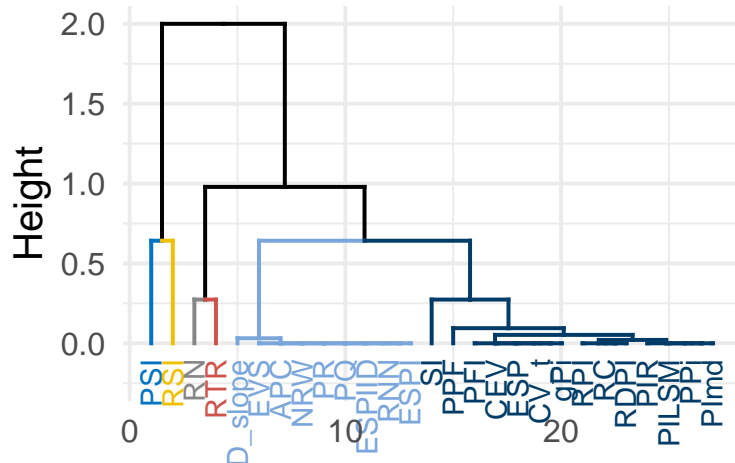
combined



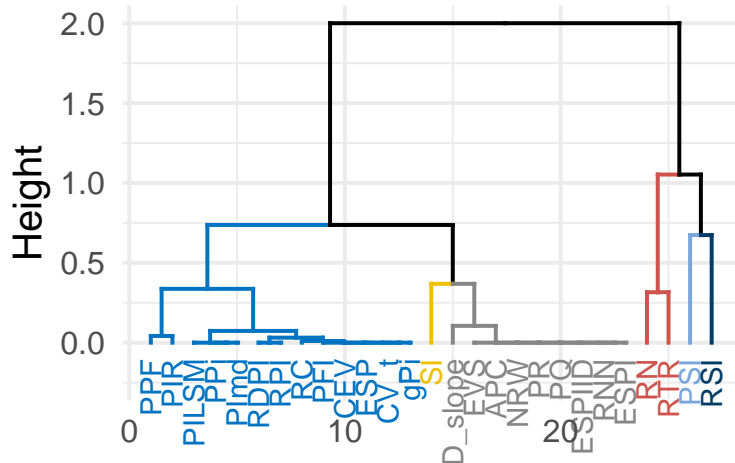
combined



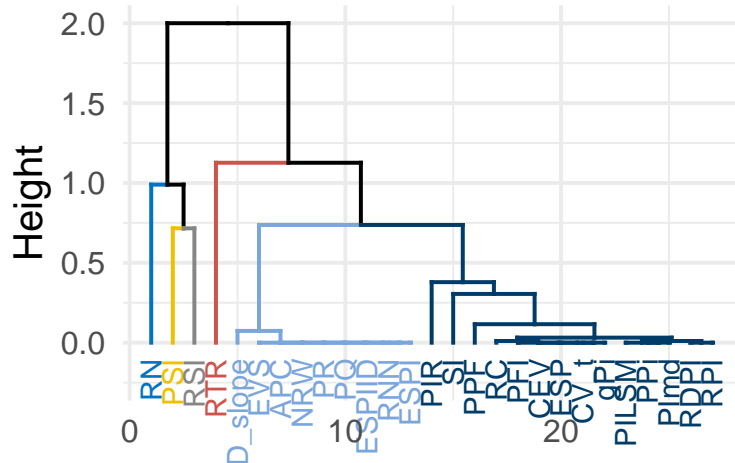
gaussian



sinusoidal



wave

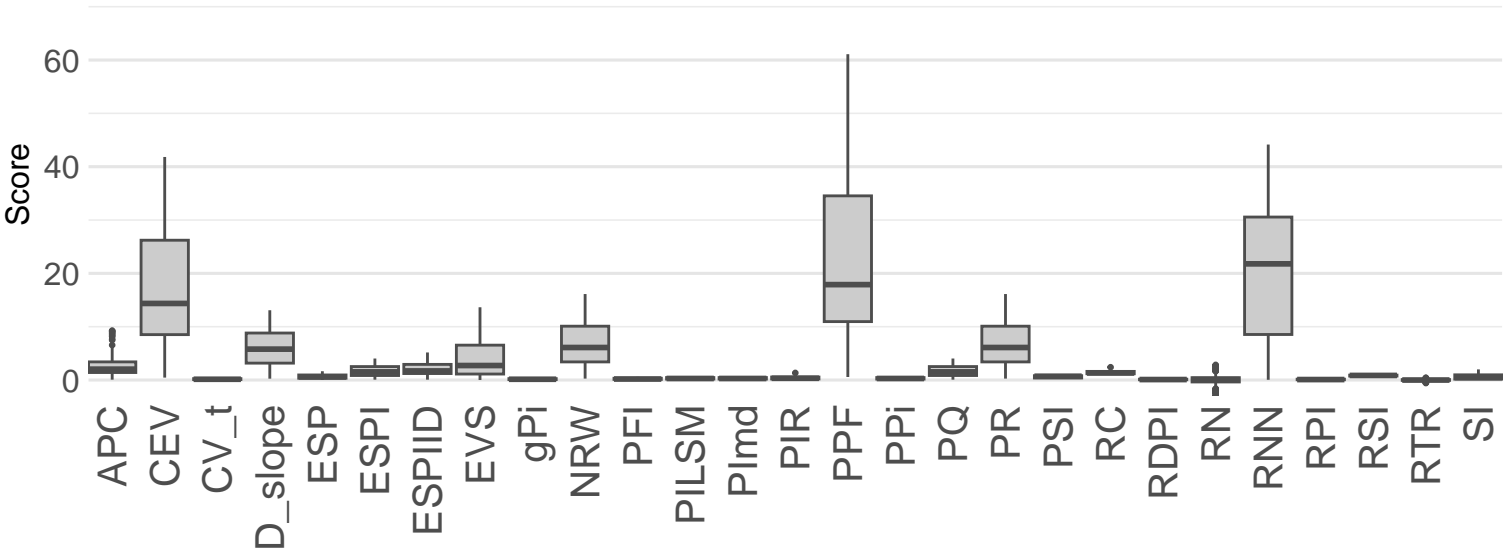


	combined	gaussian	linear	sinusoidal	wave
wave	0.811***	1.000***	1.000***	0.912***	NA
sinusoidal	0.897***	0.912***	0.912***	NA	0.912***
linear	0.811***	1.000***	NA	0.912***	1.000***
gaussian	0.811***	NA	1.000***	0.912***	1.000***
combined	NA	0.811***	0.811***	0.897***	0.811***

Plasticity Score distributions and hierarchical clustering of correlation distances

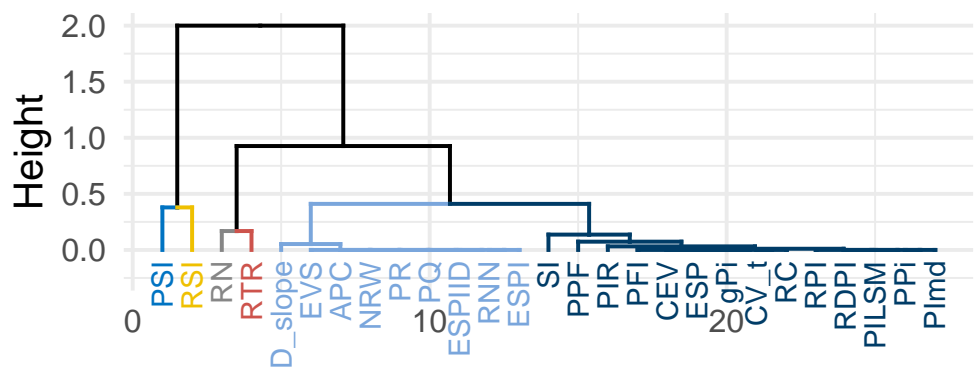
A

All Metrics Boxplot

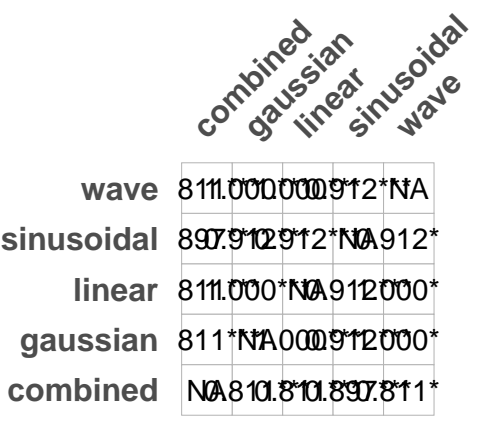


B

linear



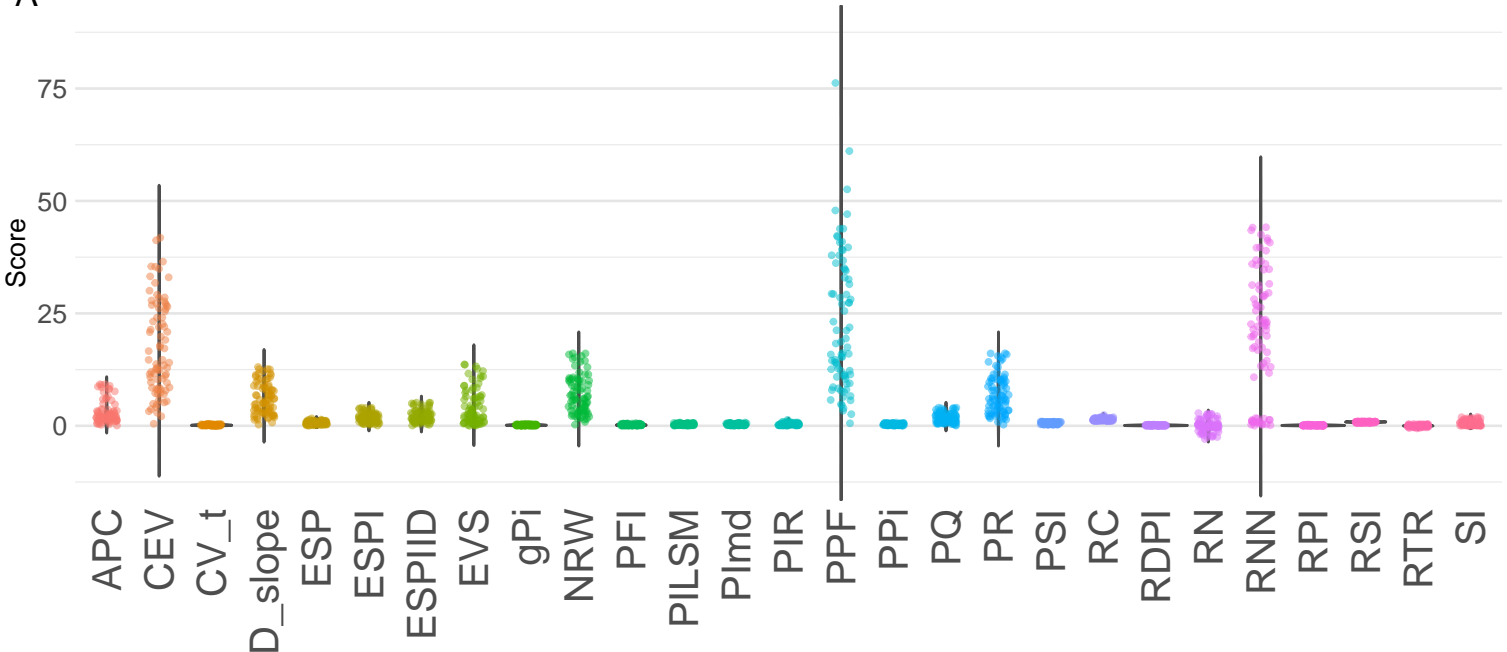
C



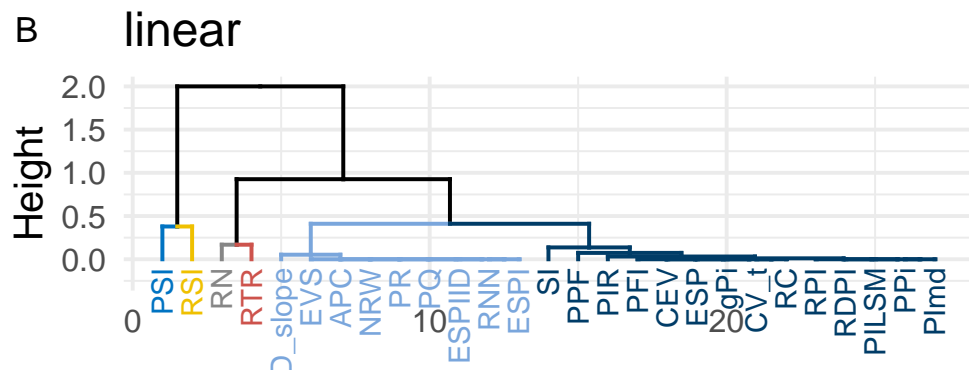
Plasticity Score distributions and hierarchical clustering of correlation distances

A

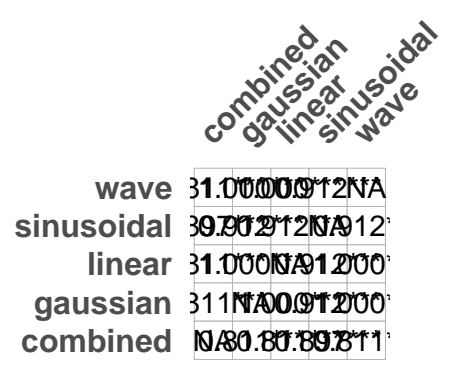
All Metrics Violin Plot



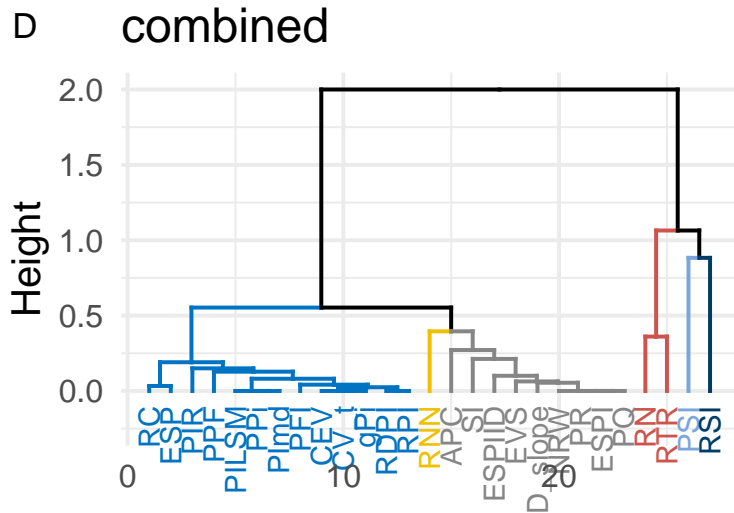
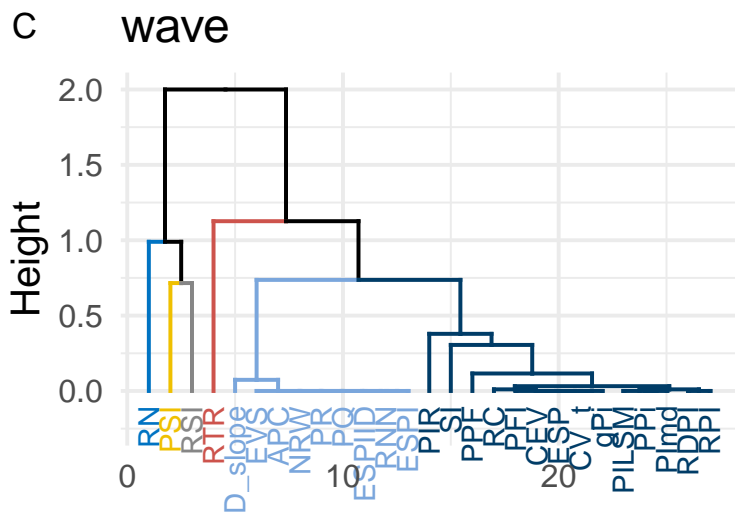
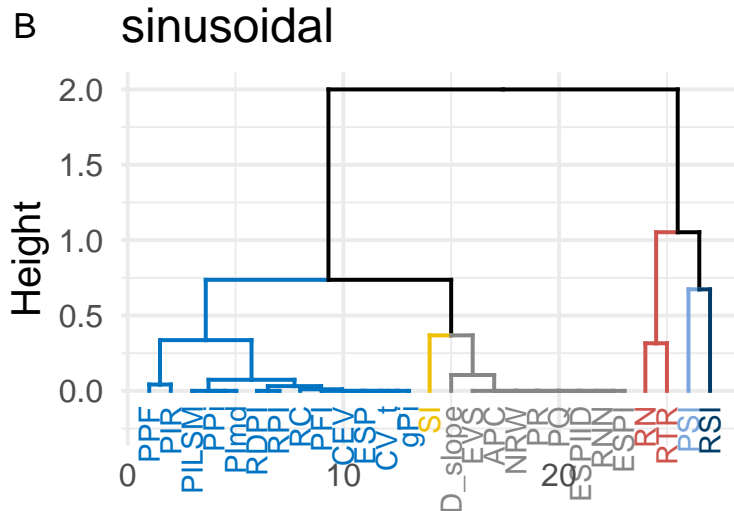
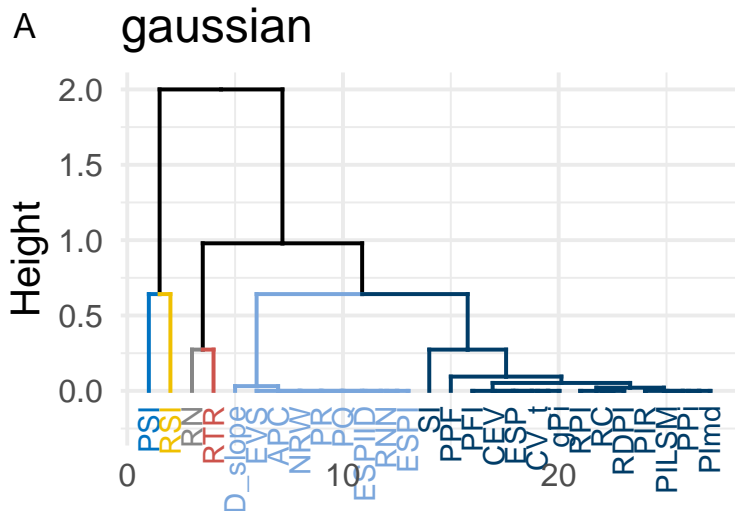
B



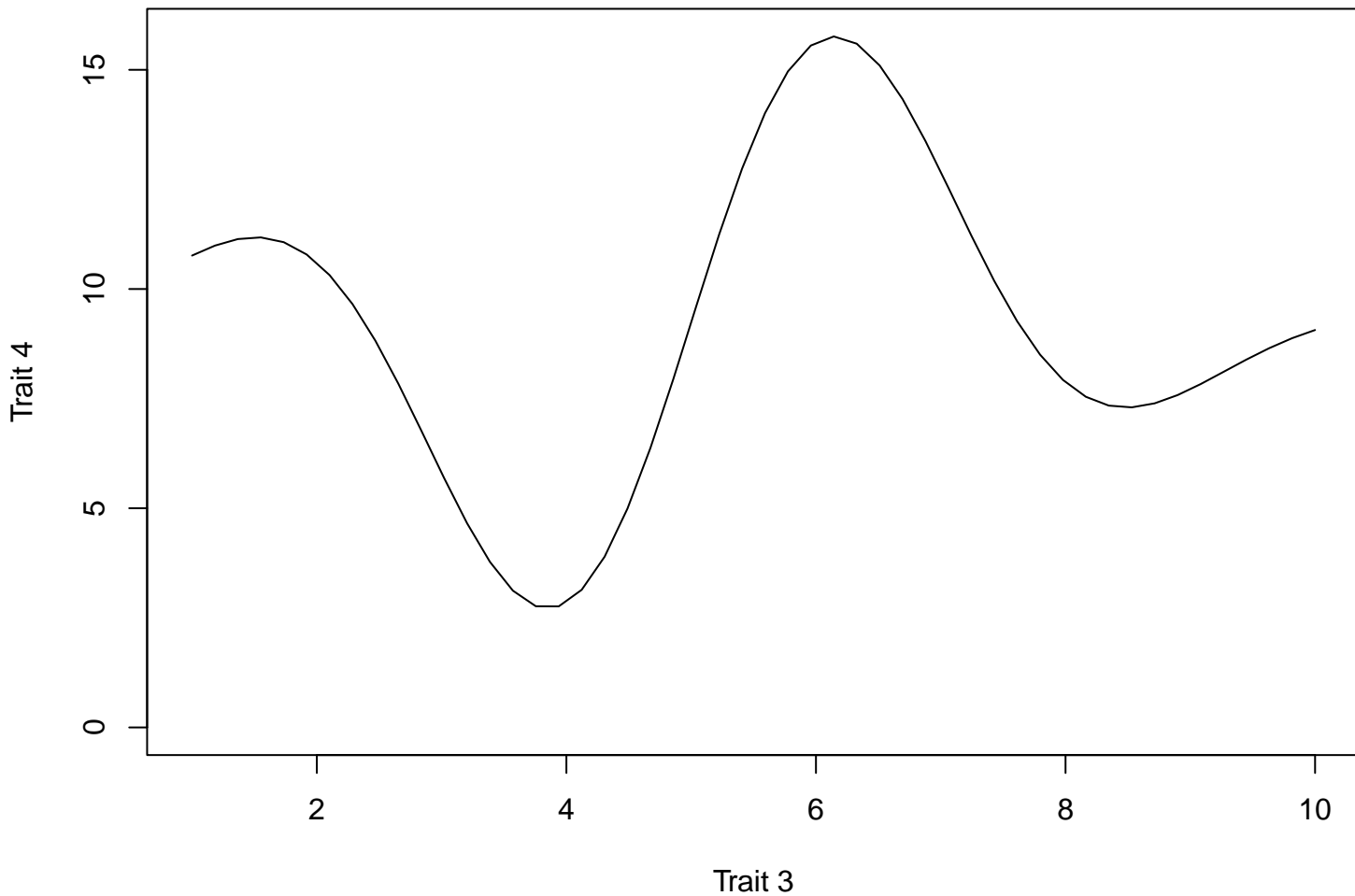
C



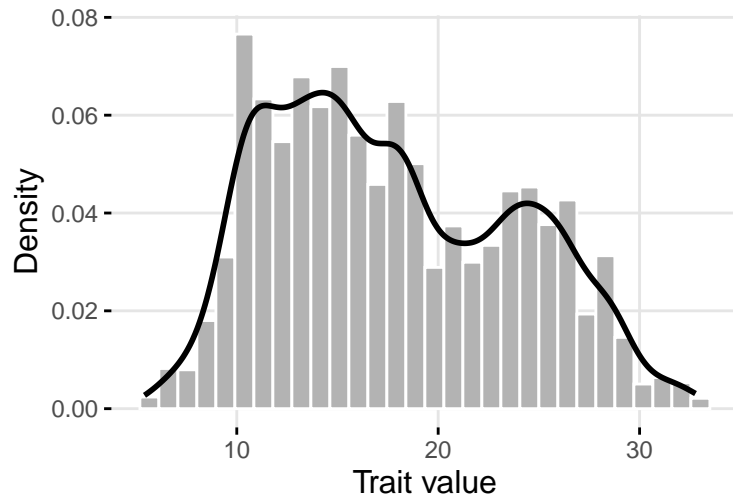
Hierarchical clustering of correlation distances



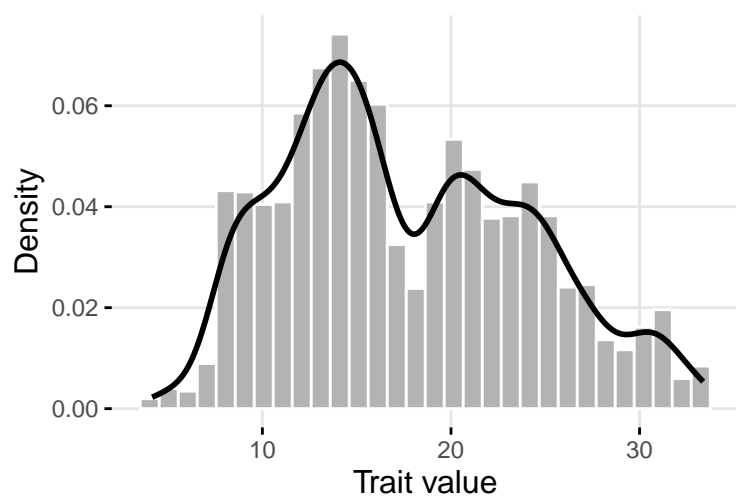
Rows 1001–1050: Trait 4 vs Trait 3



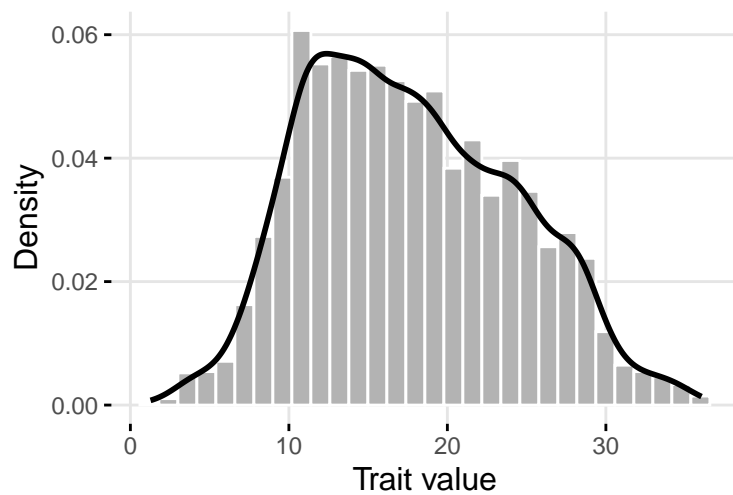
Gaussian



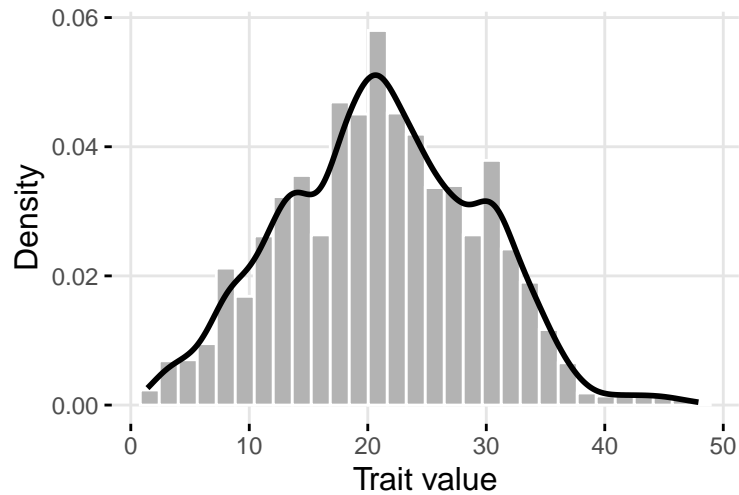
Sinusoidal



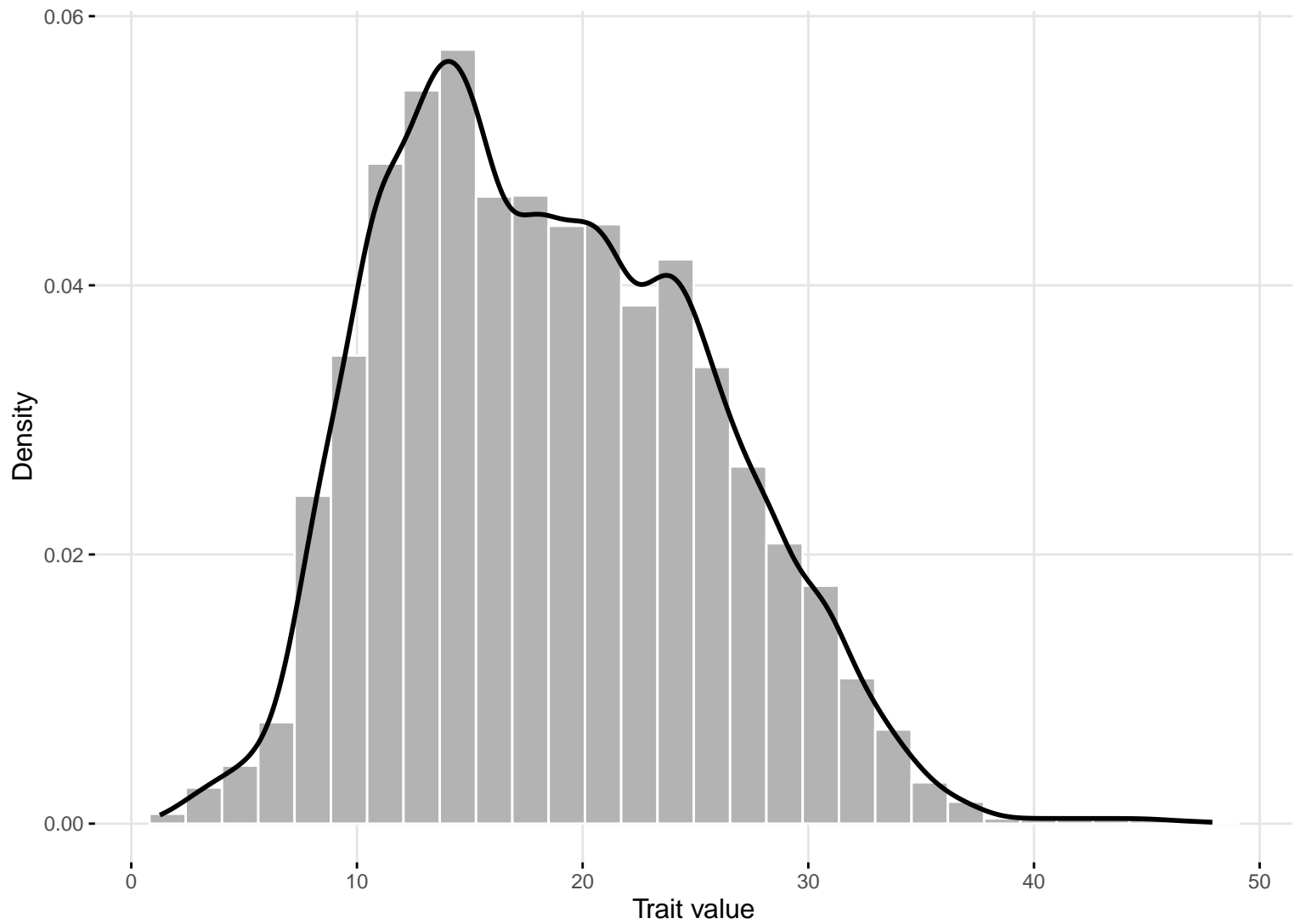
Wave

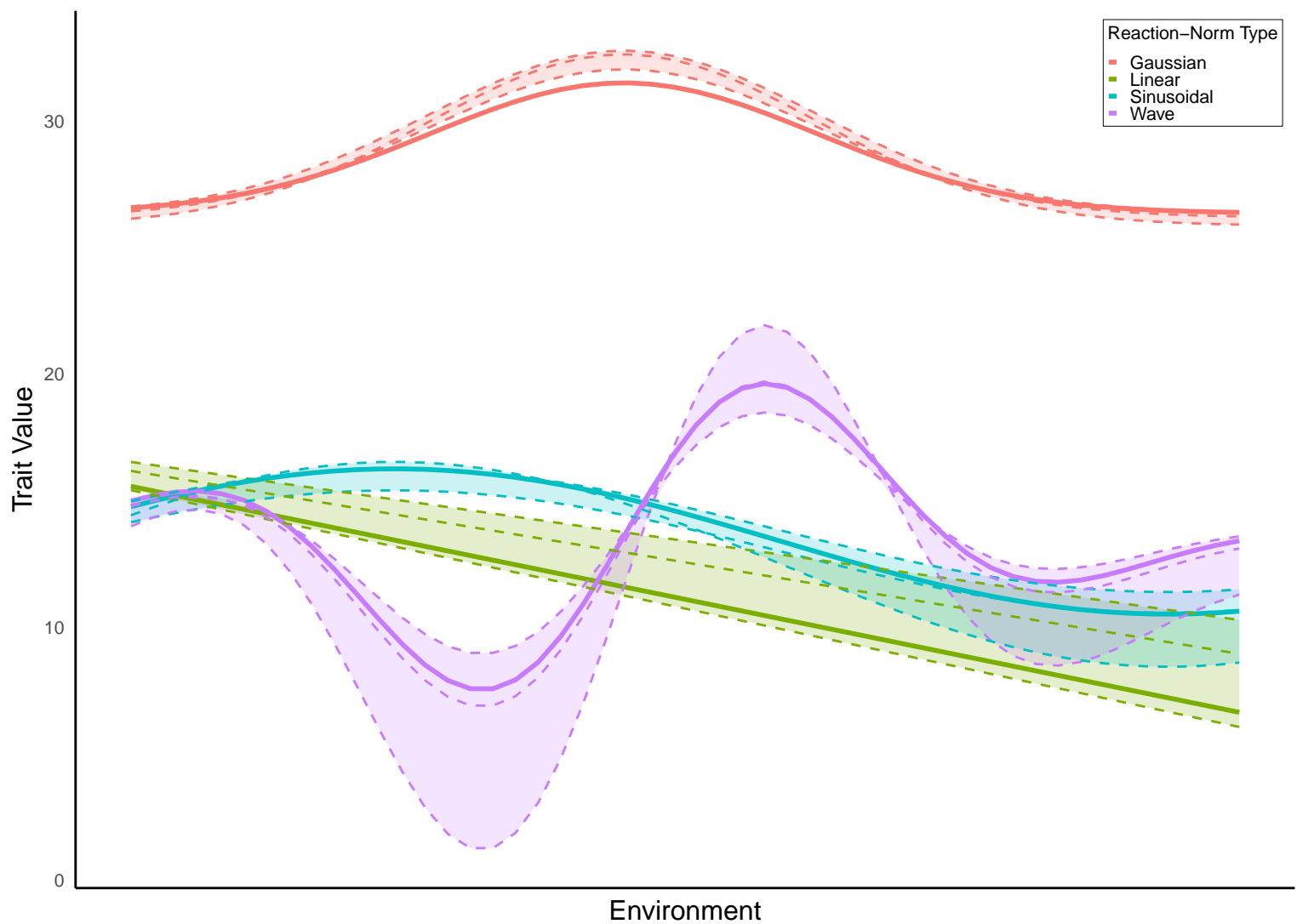


Linear

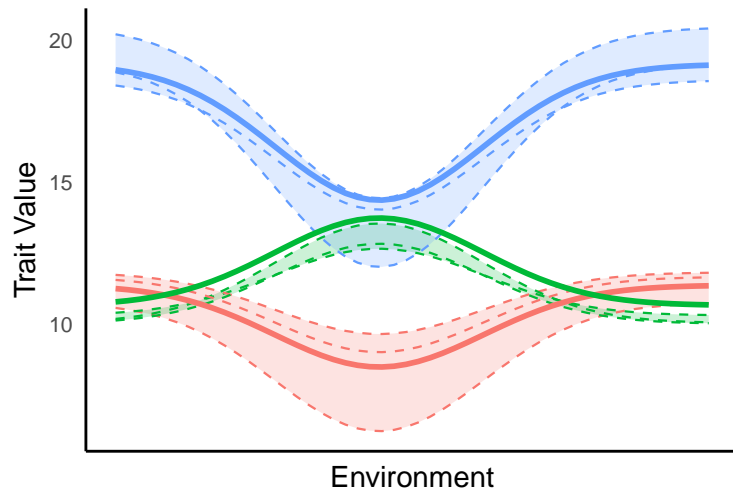


Combined Genotype Forms

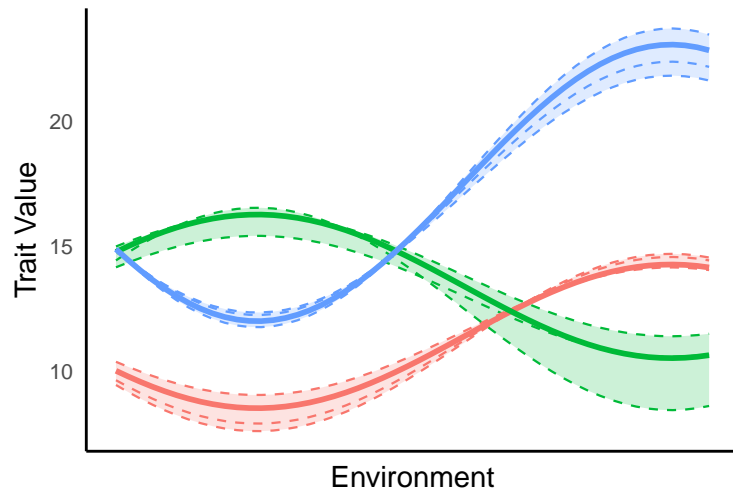




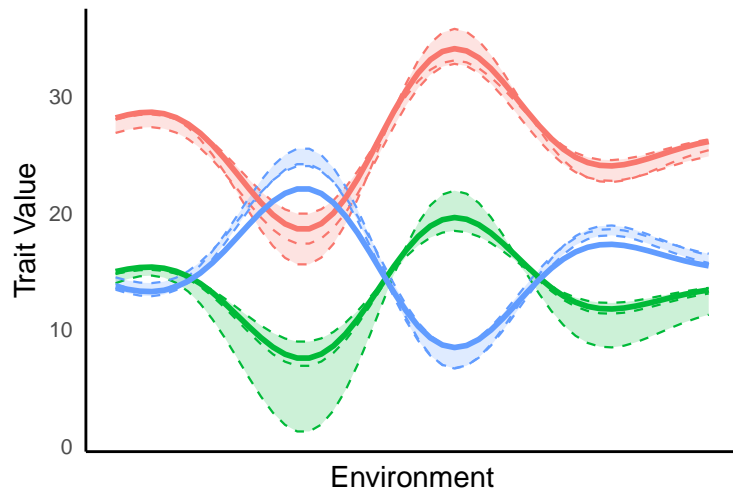
Gaussian



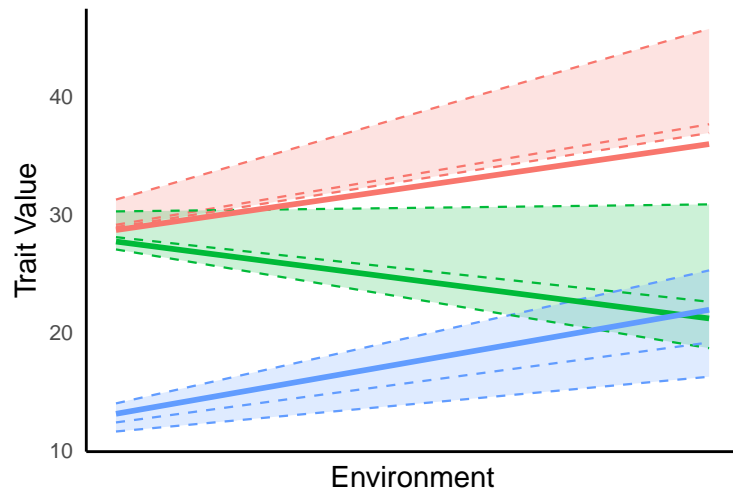
Sinusoidal



Wave

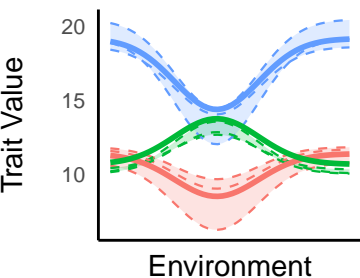


Linear

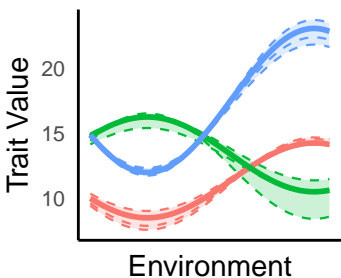


Reaction-Norm Profiles & Trait Distributions

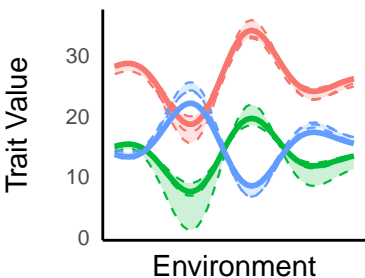
Gaussian



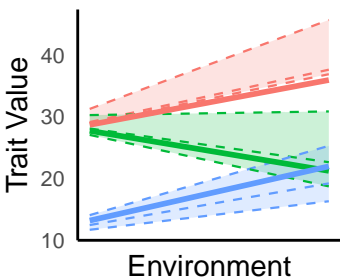
Sinusoidal



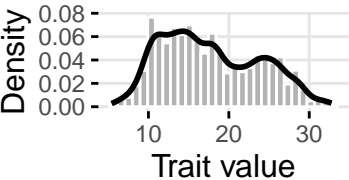
Wave



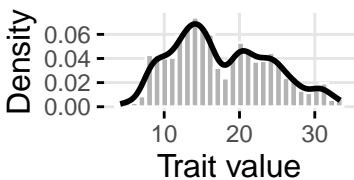
Linear



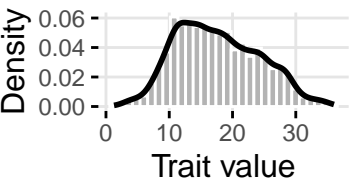
Gaussian



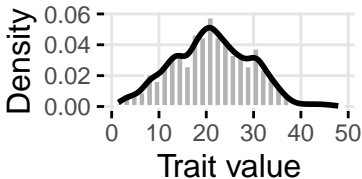
Sinusoidal



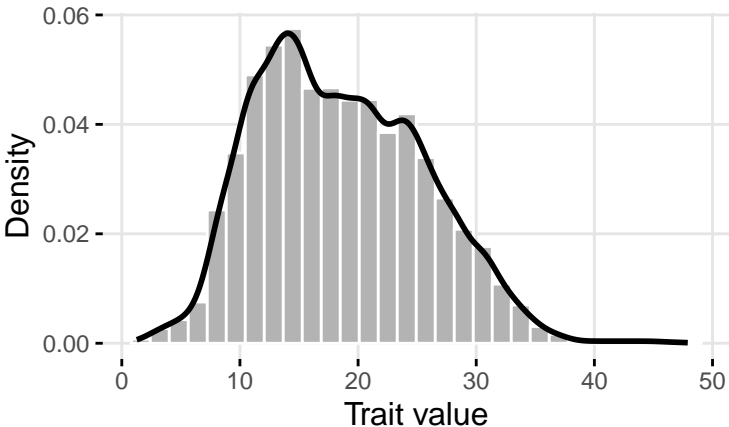
Wave



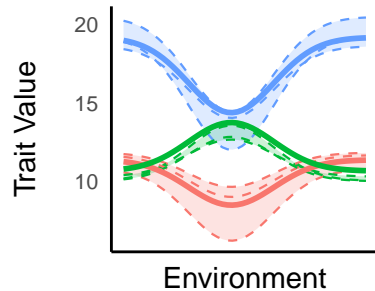
Linear



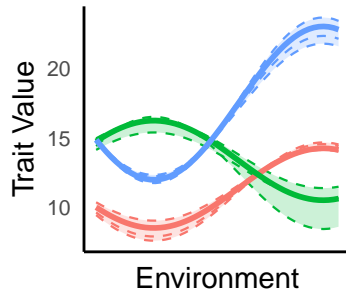
Combined Genotype Forms



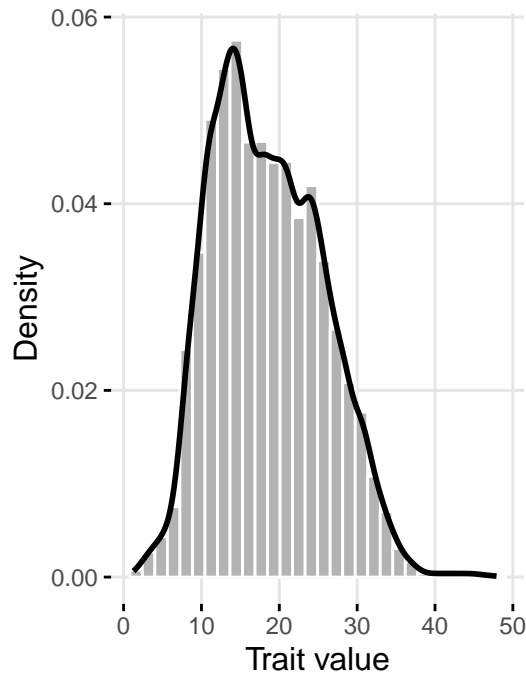
Gaussian



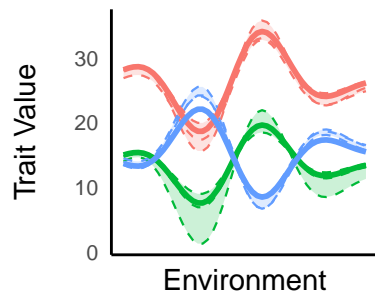
Sinusoidal



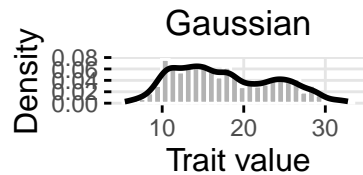
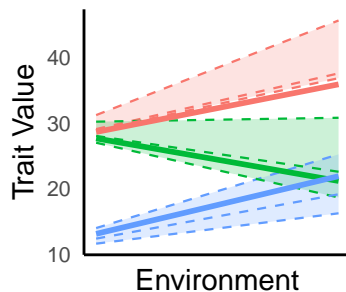
Combined Genotype Forms



Wave



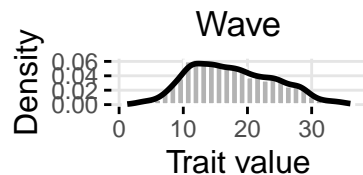
Linear



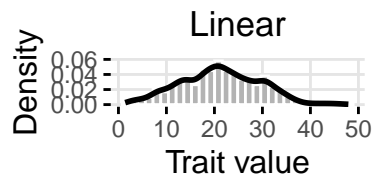
Gaussian



Sinusoidal



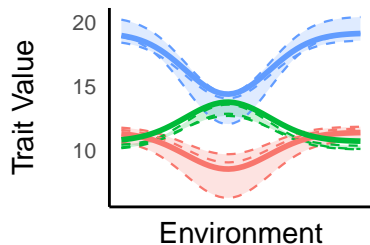
Wave



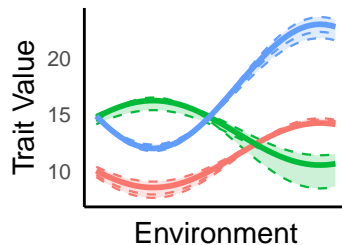
Linear

Alt B: Combined Above, Panels Below

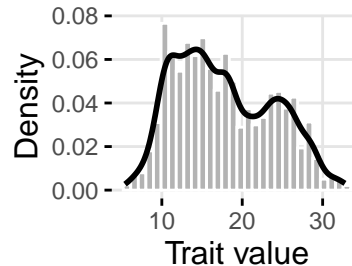
Gaussian



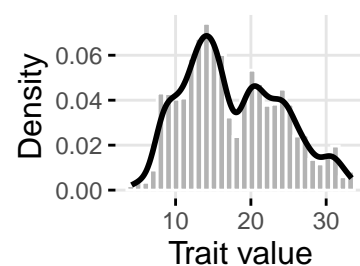
Sinusoidal



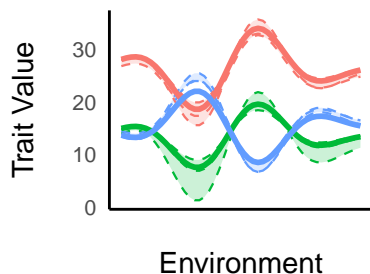
Gaussian



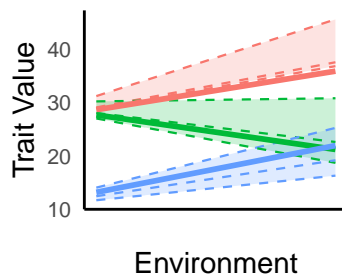
Sinusoidal



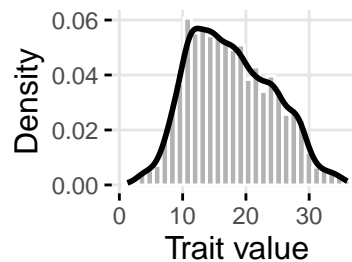
Wave



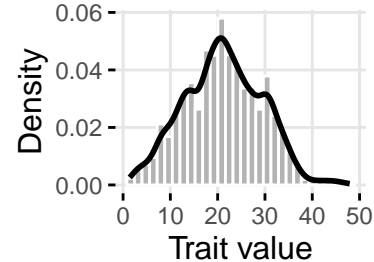
Linear



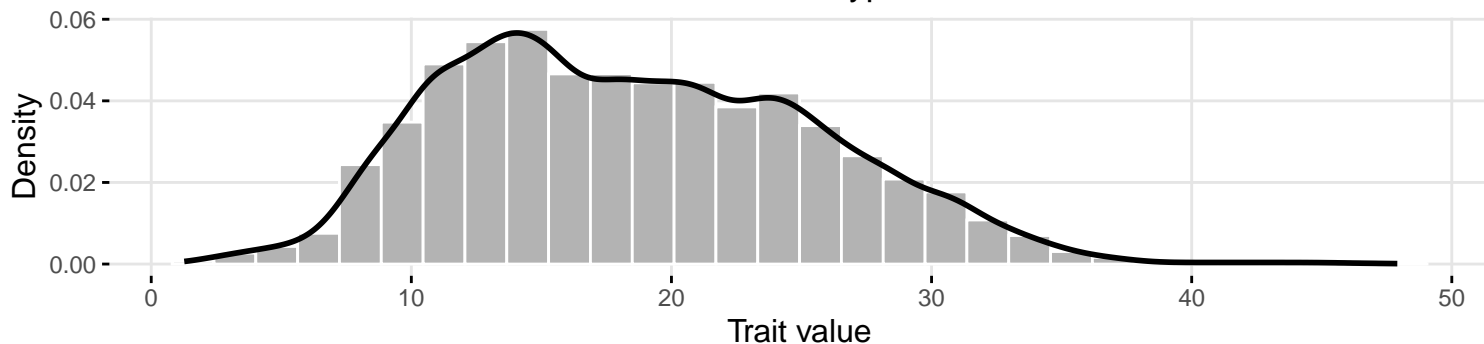
Wave



Linear



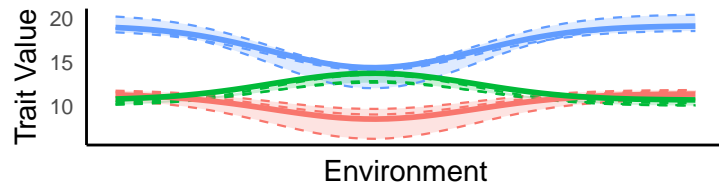
Combined Genotype Forms



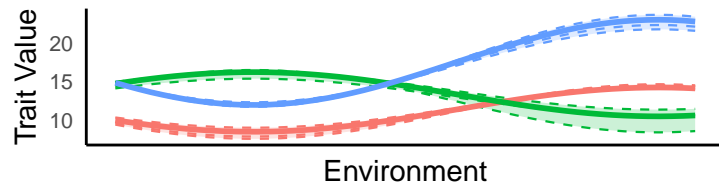
Reaction-Norm Profiles & Trait Distributions

A

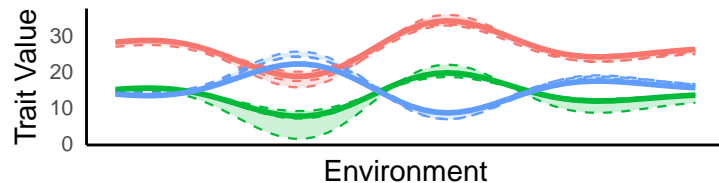
Gaussian



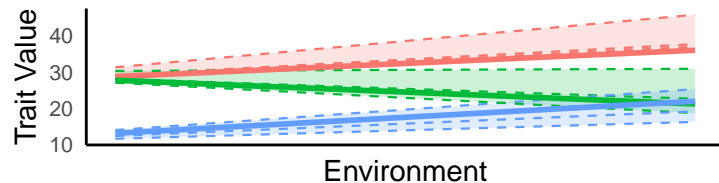
Sinusoidal



Wave

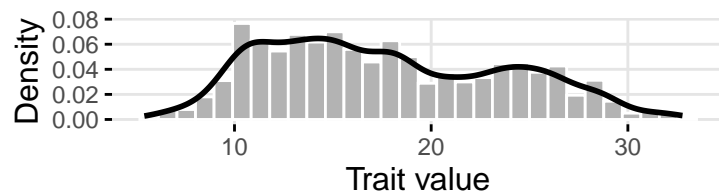


Linear

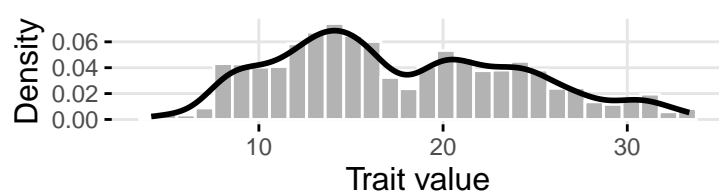


B

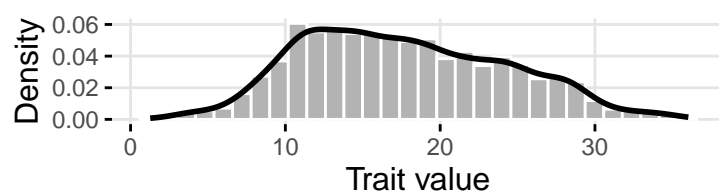
Gaussian



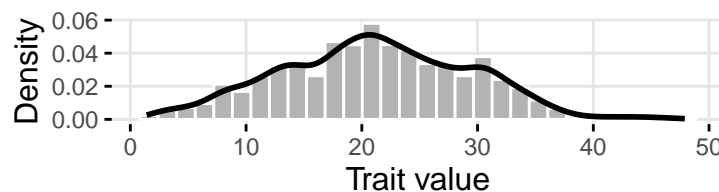
Sinusoidal



Wave

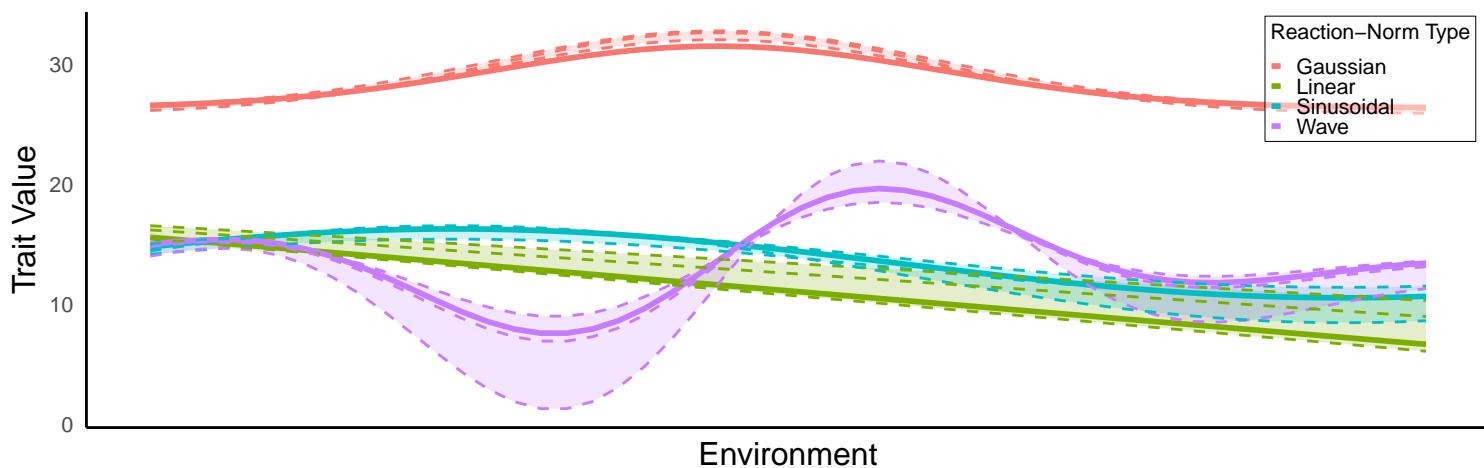


Linear

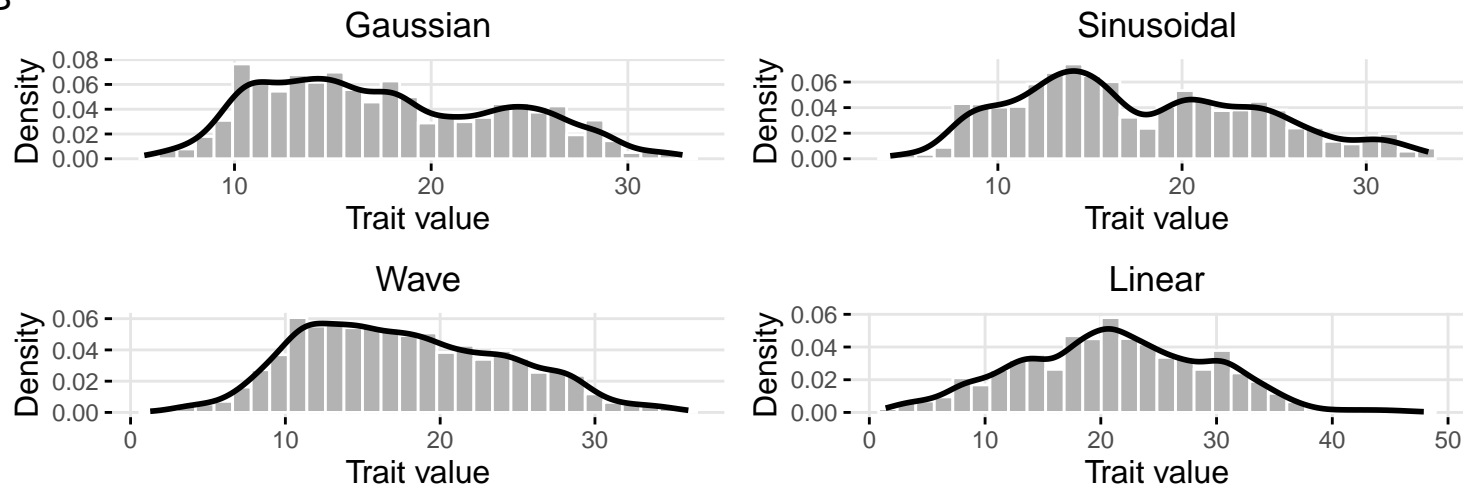


Reaction-Norm Profiles & Trait Distributions

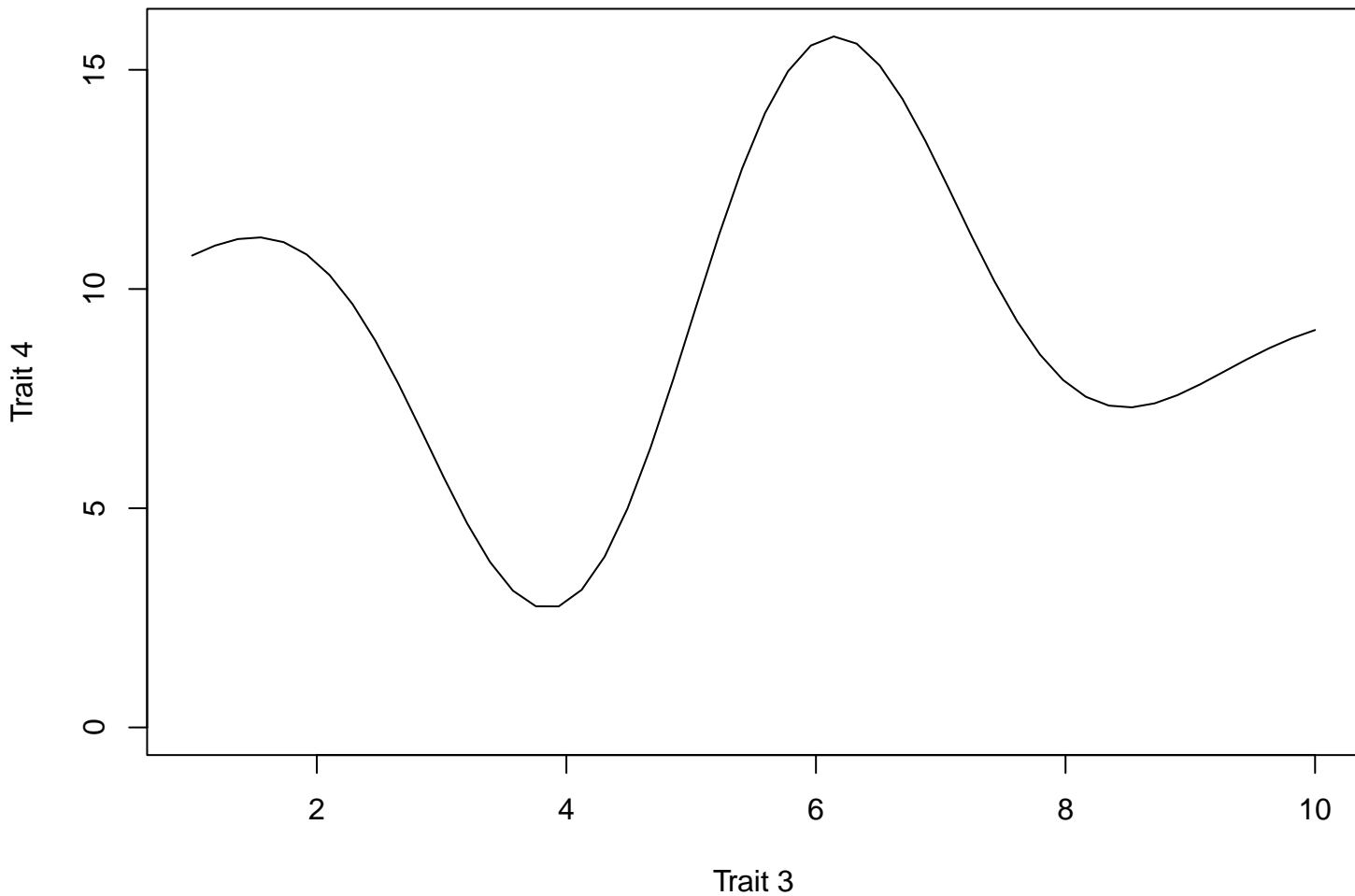
A



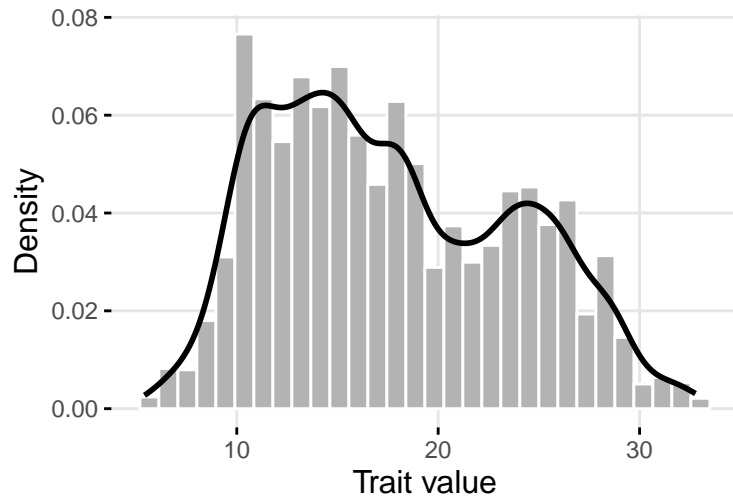
B



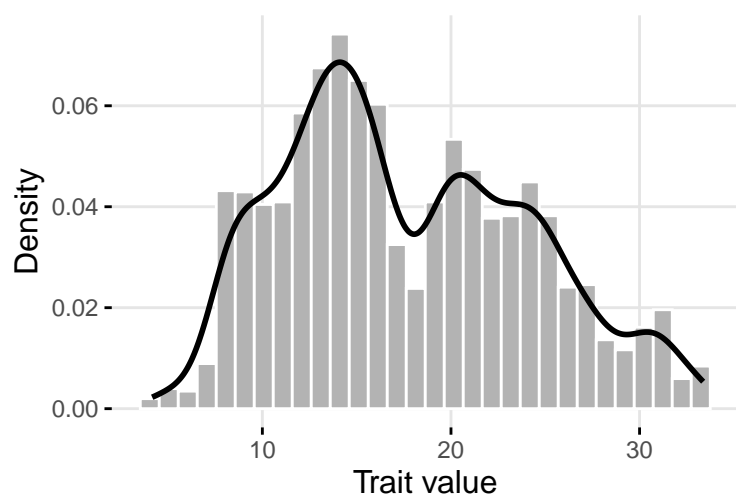
Rows 1001–1050: Trait 4 vs Trait 3



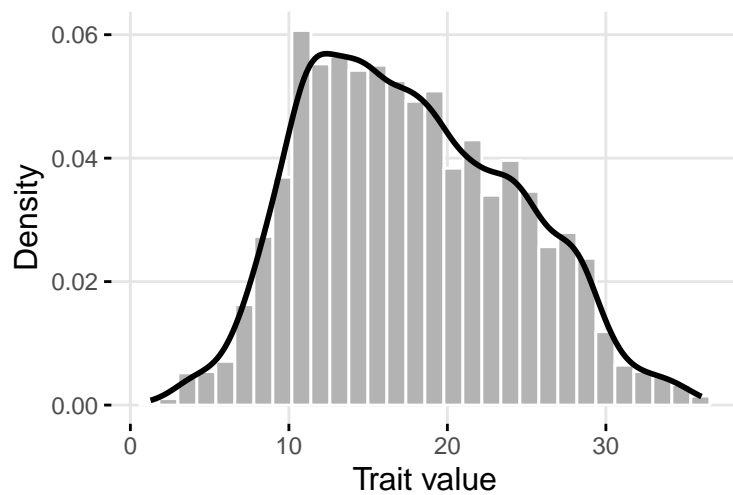
Gaussian



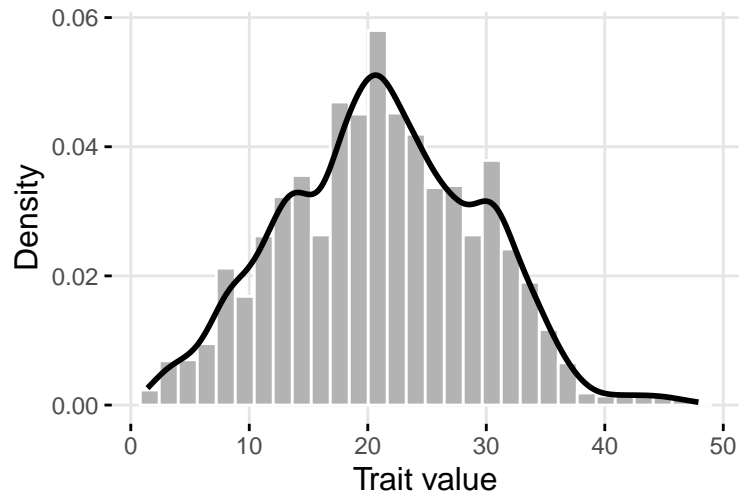
Sinusoidal



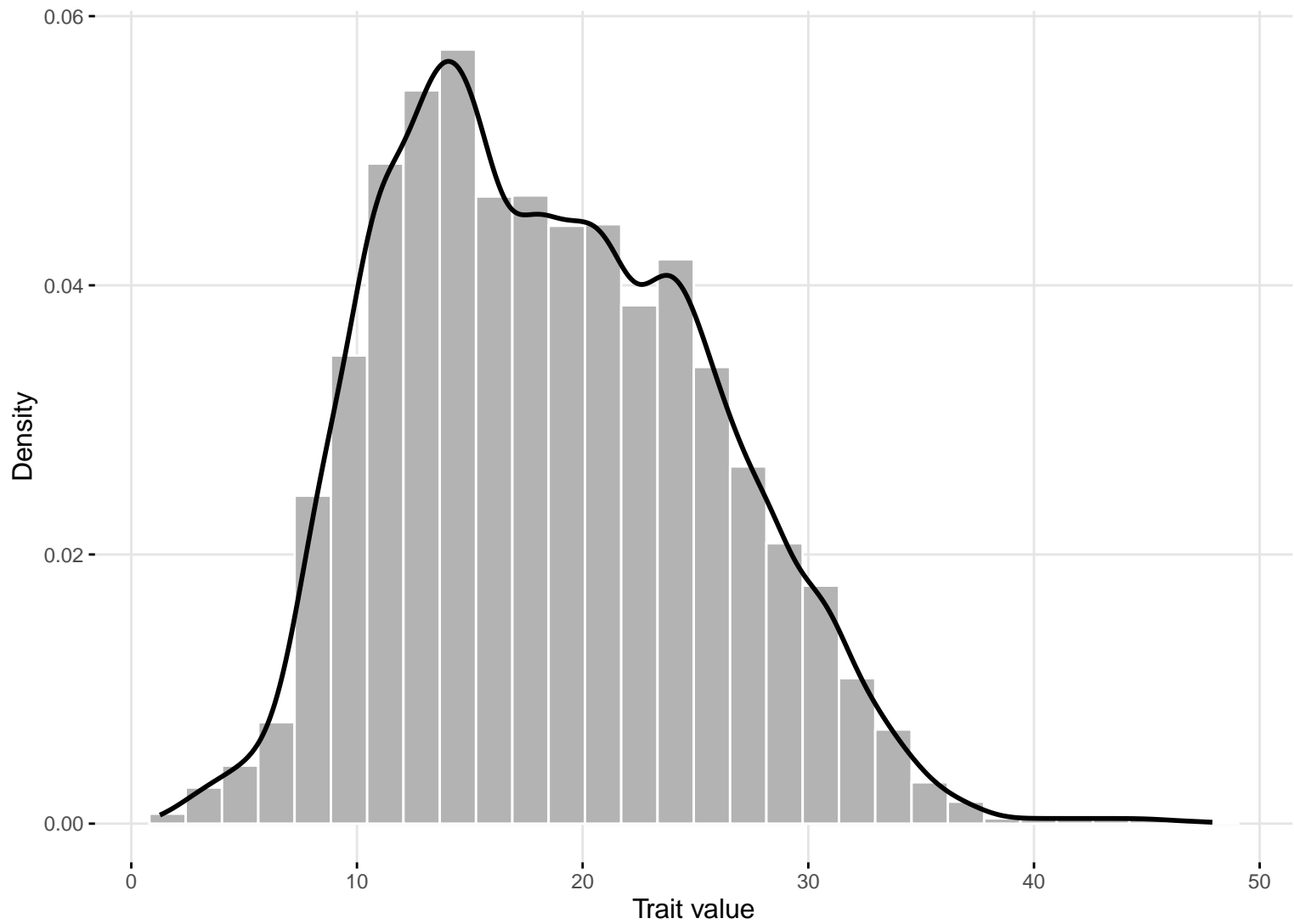
Wave

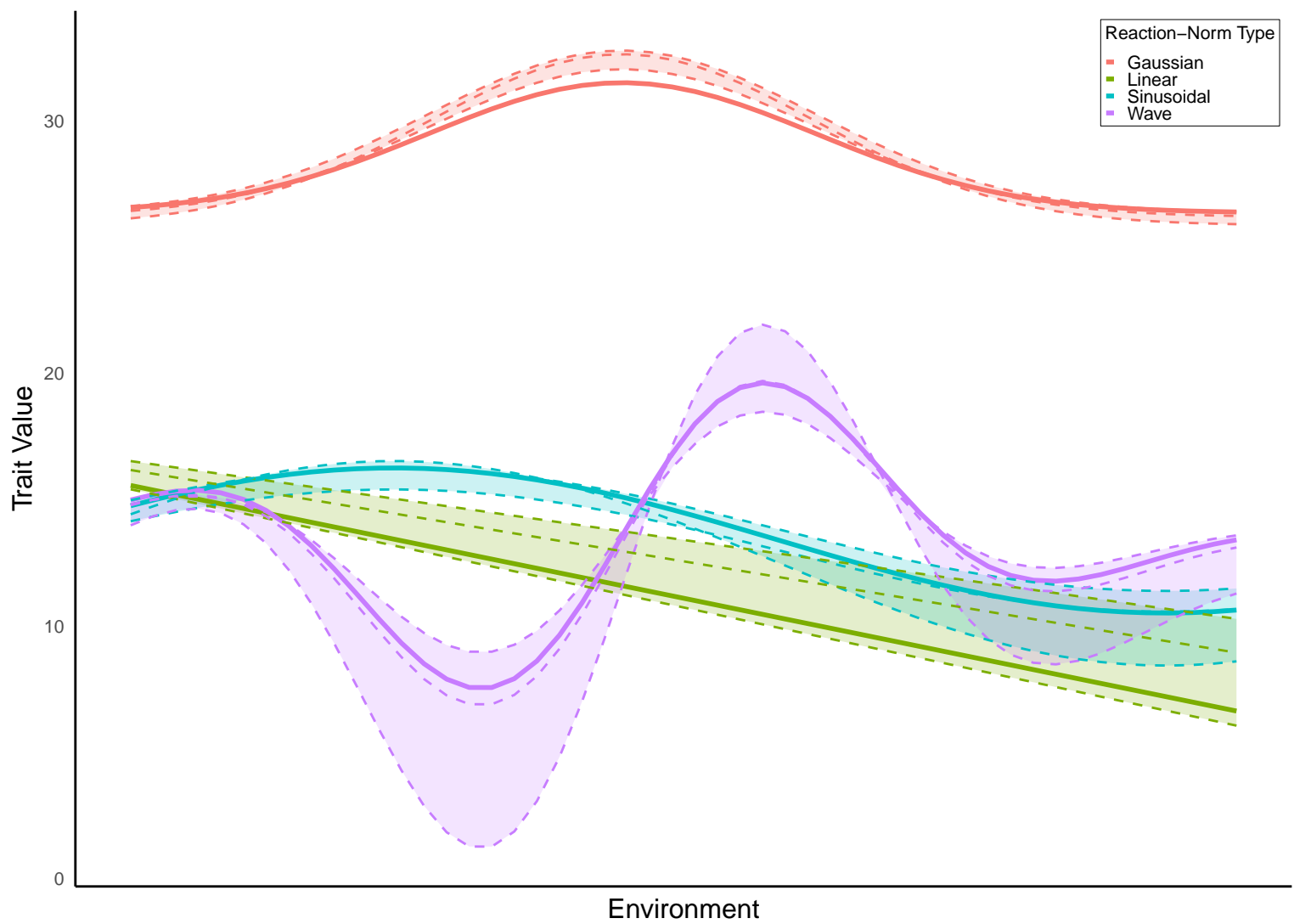


Linear

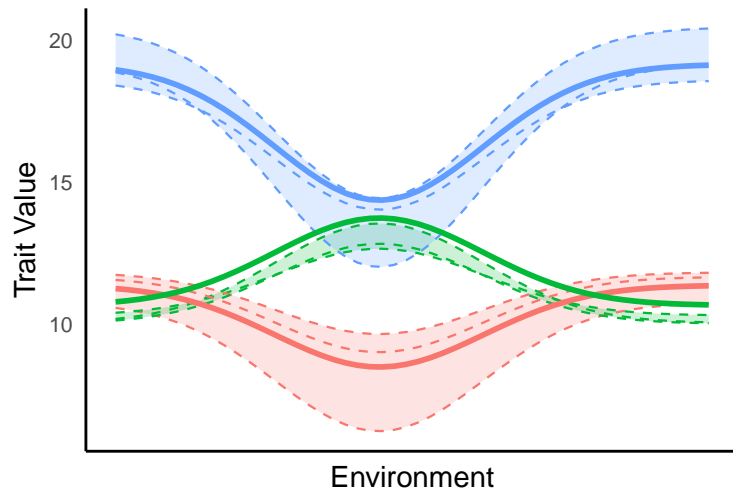


Combined Genotype Forms

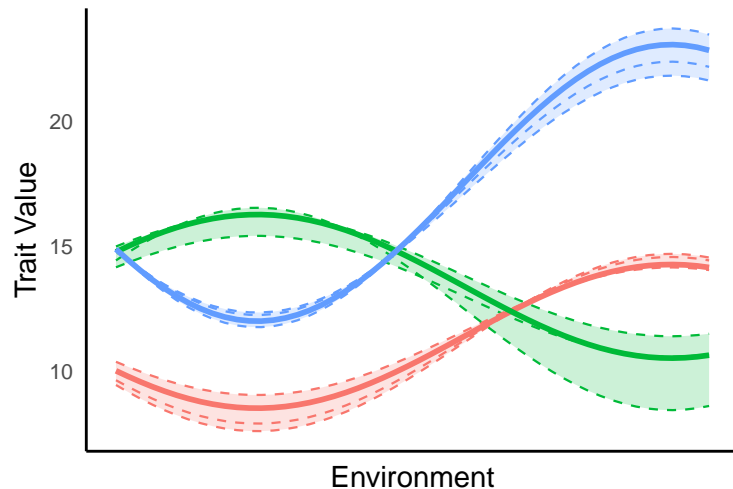




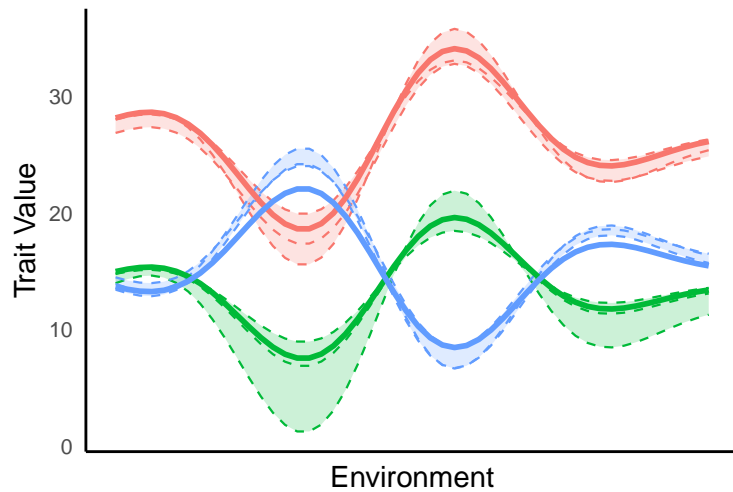
Gaussian



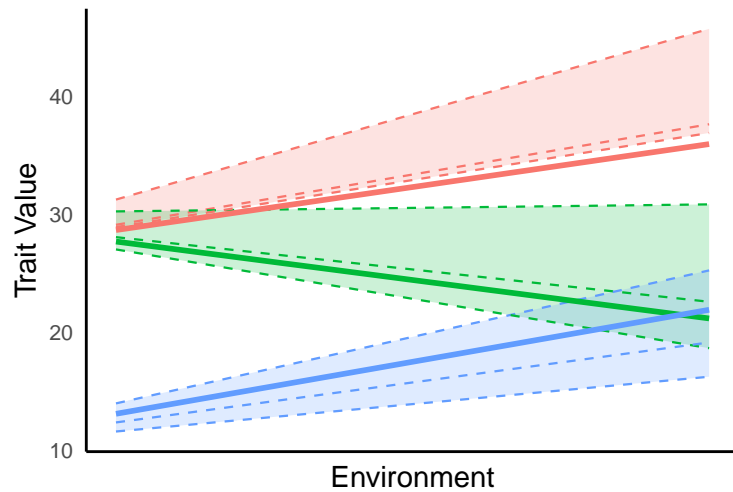
Sinusoidal



Wave

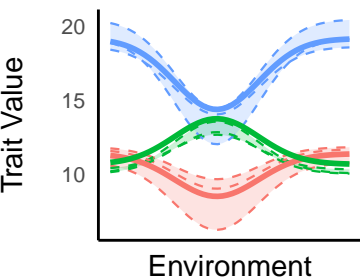


Linear

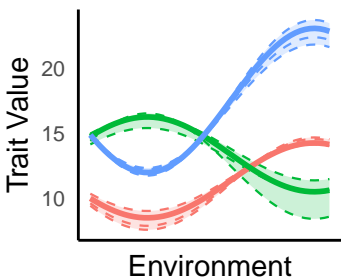


Reaction-Norm Profiles & Trait Distributions

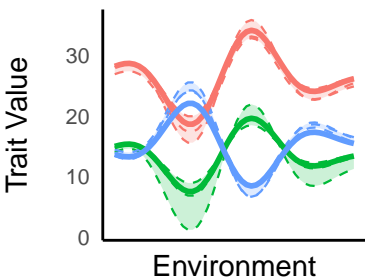
Gaussian



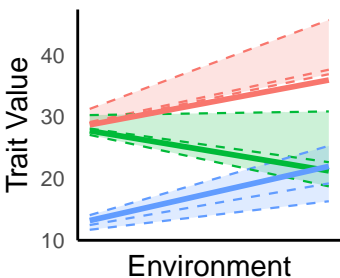
Sinusoidal



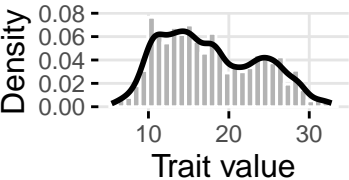
Wave



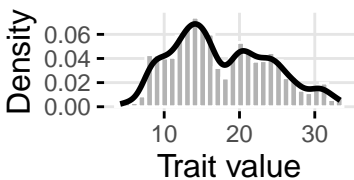
Linear



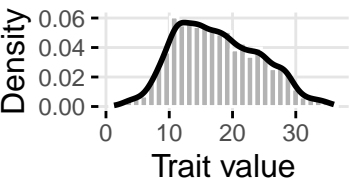
Gaussian



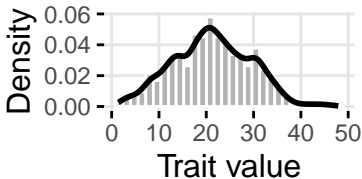
Sinusoidal



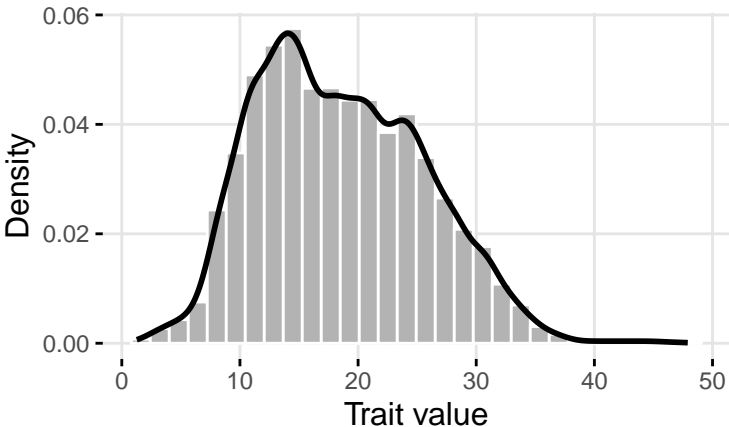
Wave



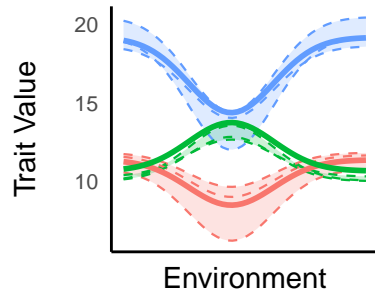
Linear



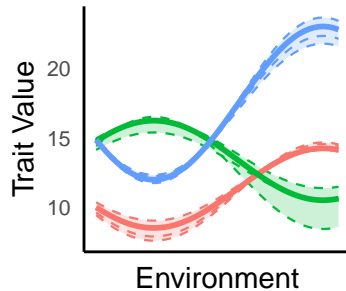
Combined Genotype Forms



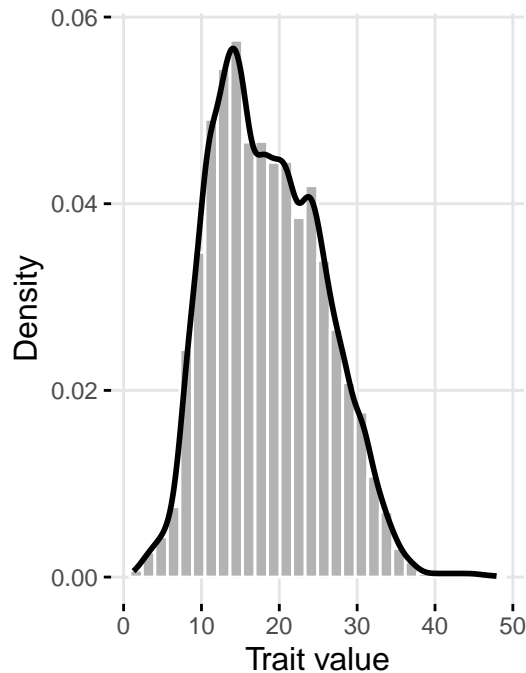
Gaussian



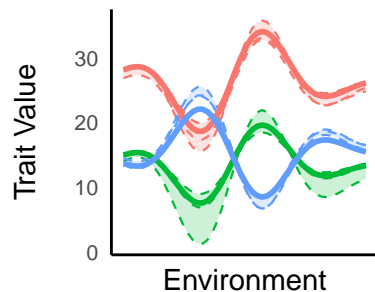
Sinusoidal



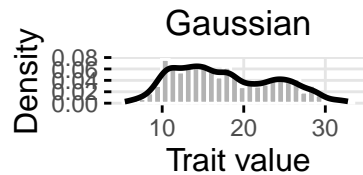
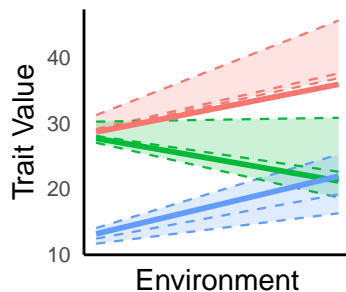
Combined Genotype Forms



Wave



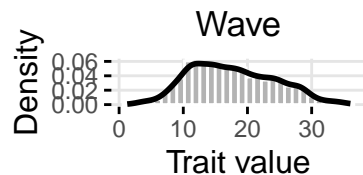
Linear



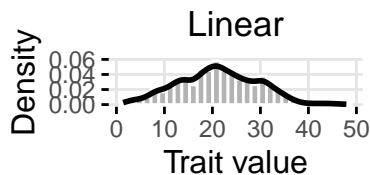
Gaussian



Sinusoidal



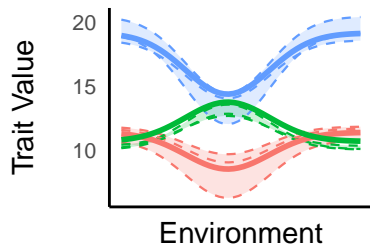
Wave



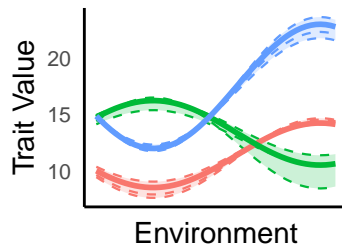
Linear

Alt B: Combined Above, Panels Below

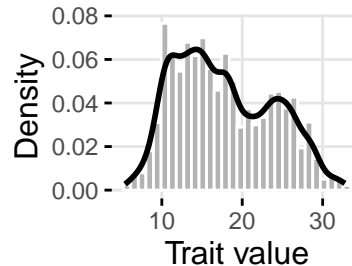
Gaussian



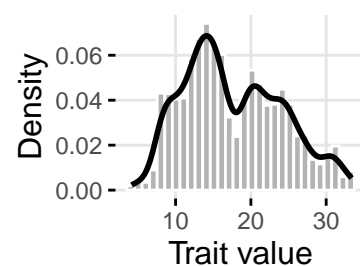
Sinusoidal



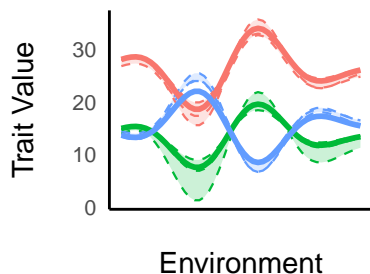
Gaussian



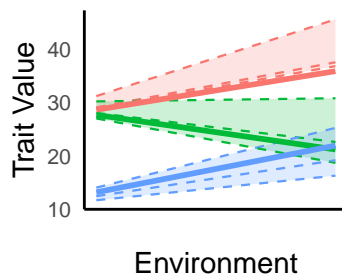
Sinusoidal



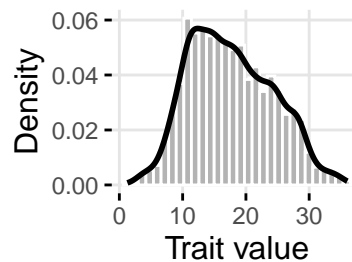
Wave



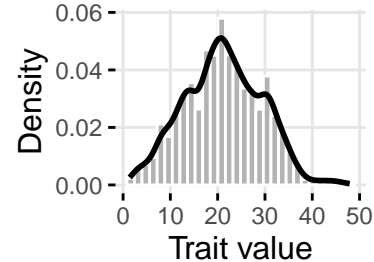
Linear



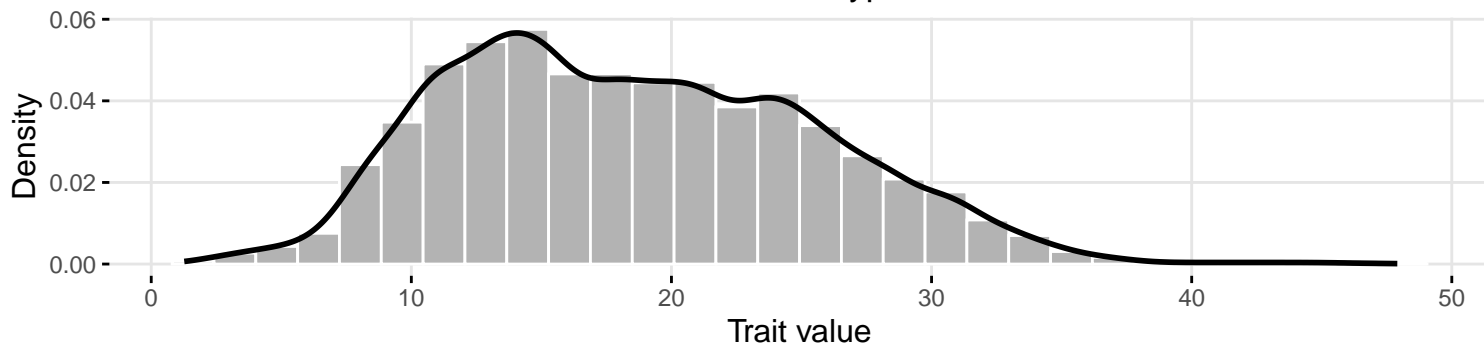
Wave



Linear



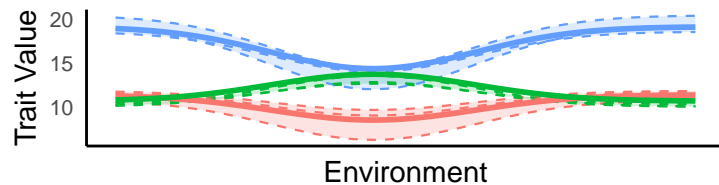
Combined Genotype Forms



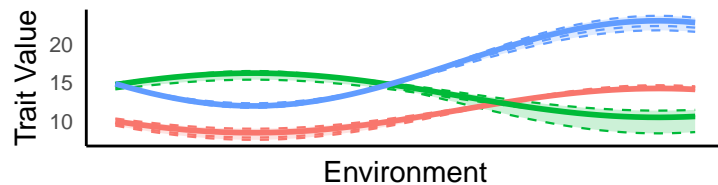
Reaction-Norm Profiles & Trait Distributions

A

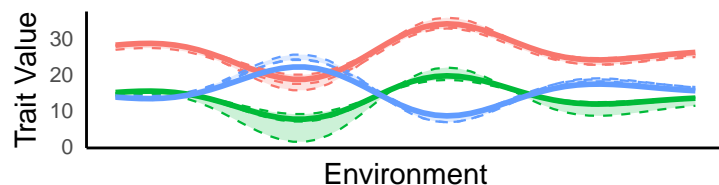
Gaussian



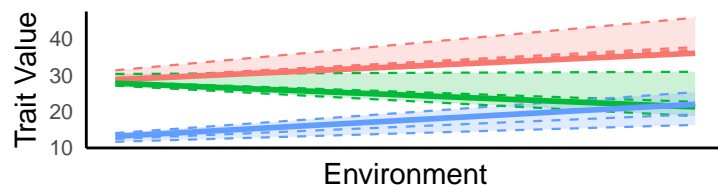
Sinusoidal



Wave

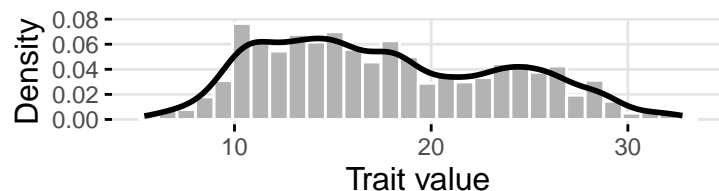


Linear

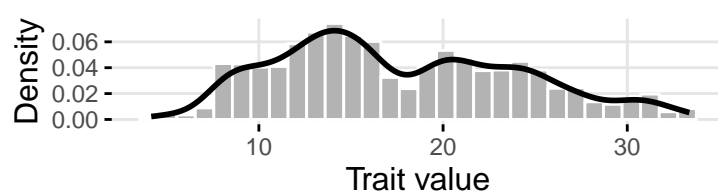


B

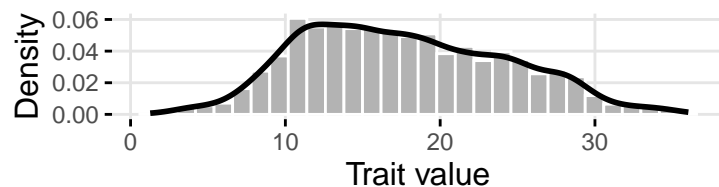
Gaussian



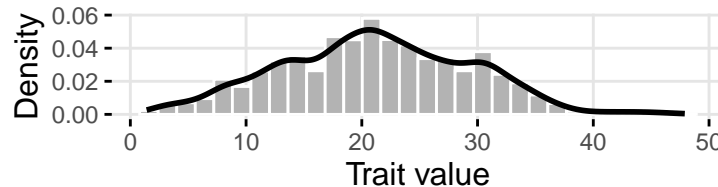
Sinusoidal



Wave

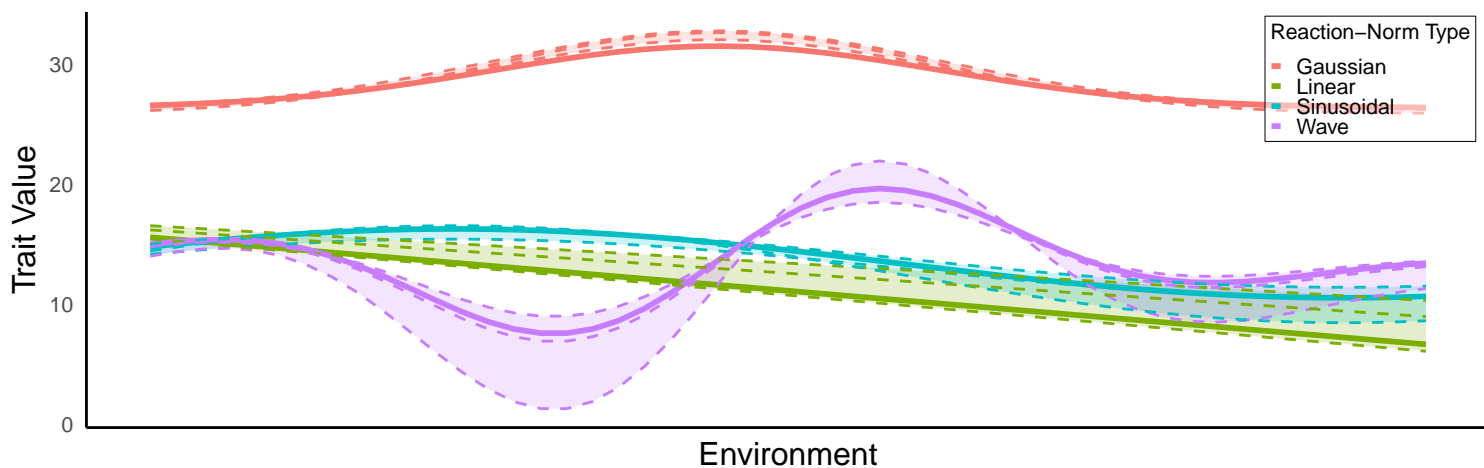


Linear

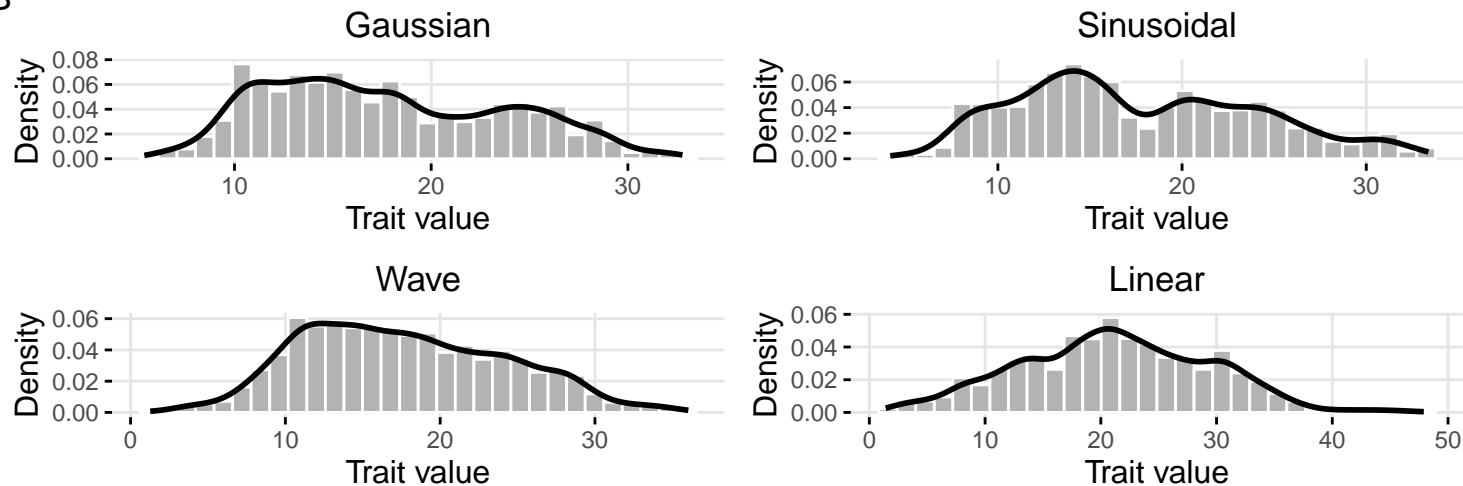


Reaction-Norm Profiles & Trait Distributions

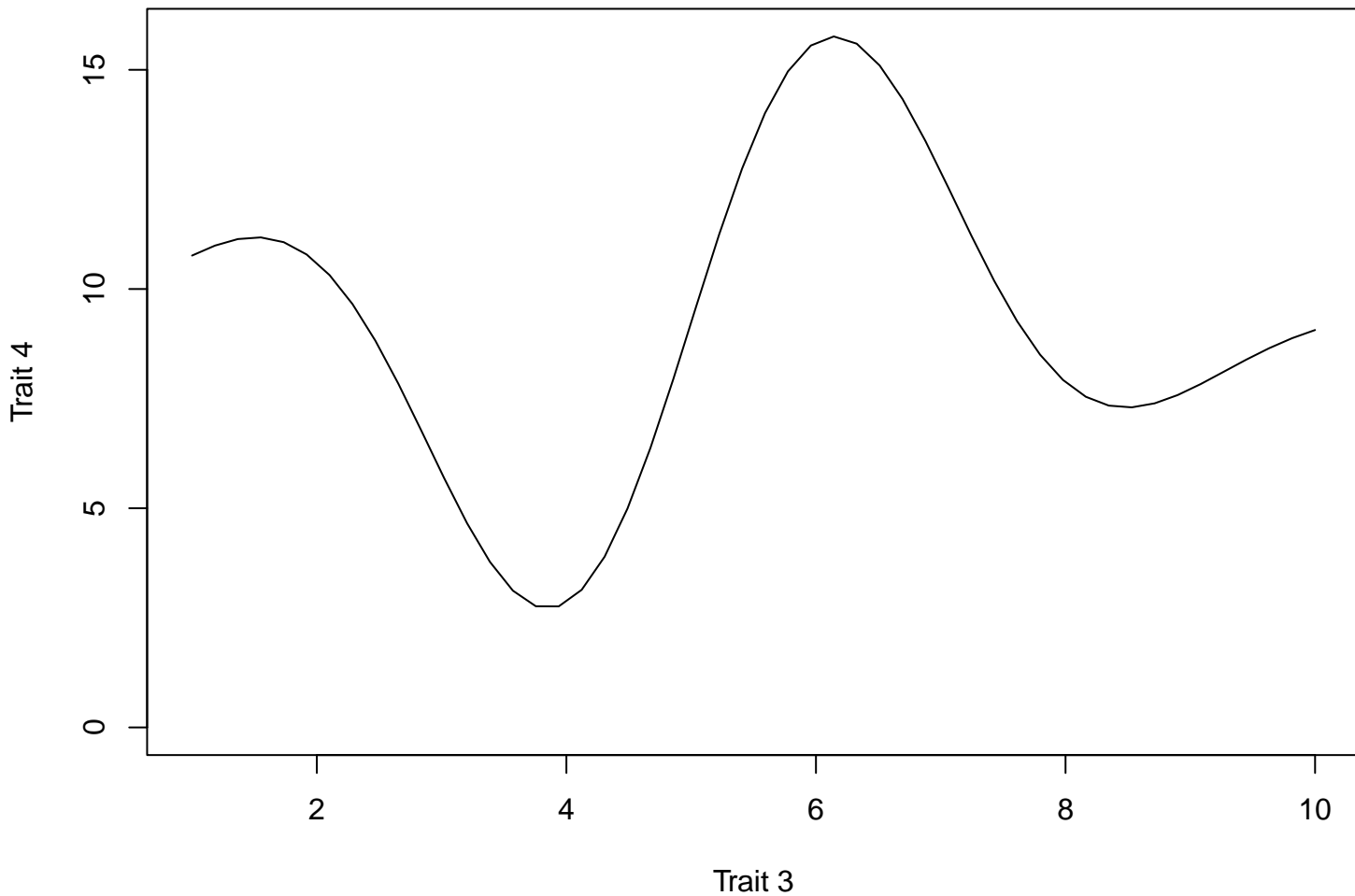
A



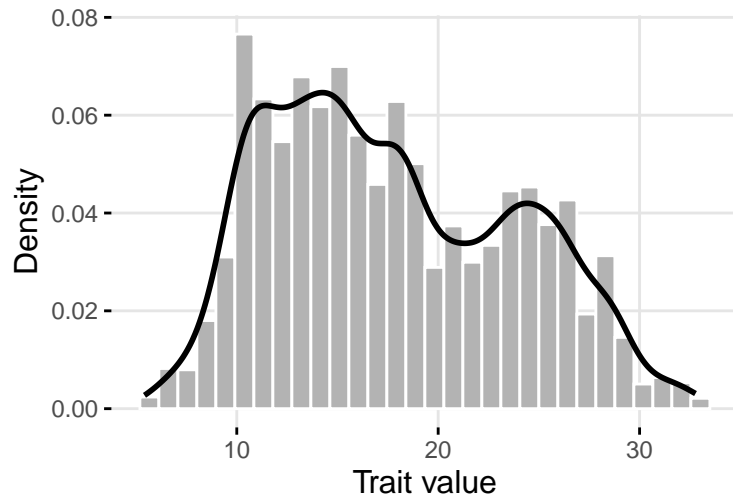
B



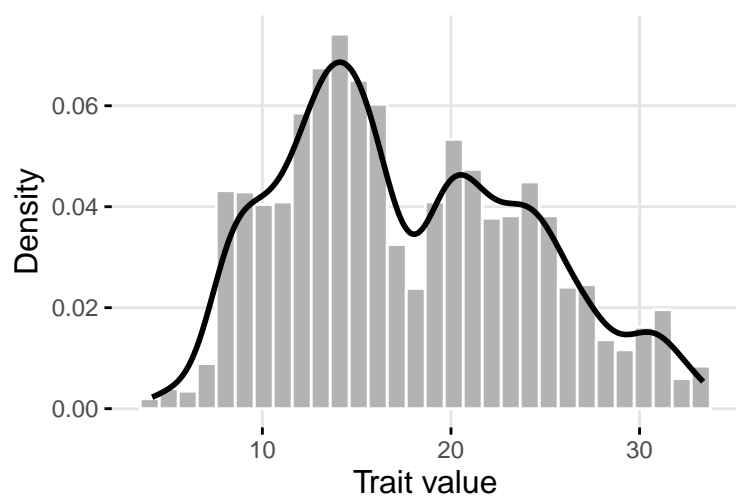
Rows 1001–1050: Trait 4 vs Trait 3



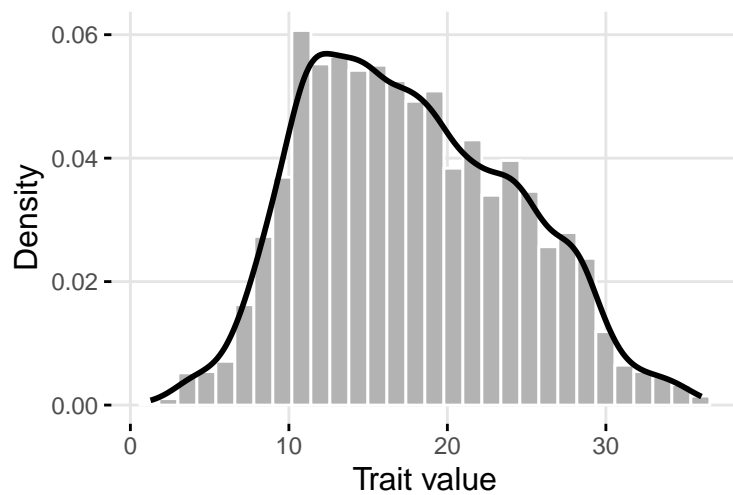
Gaussian



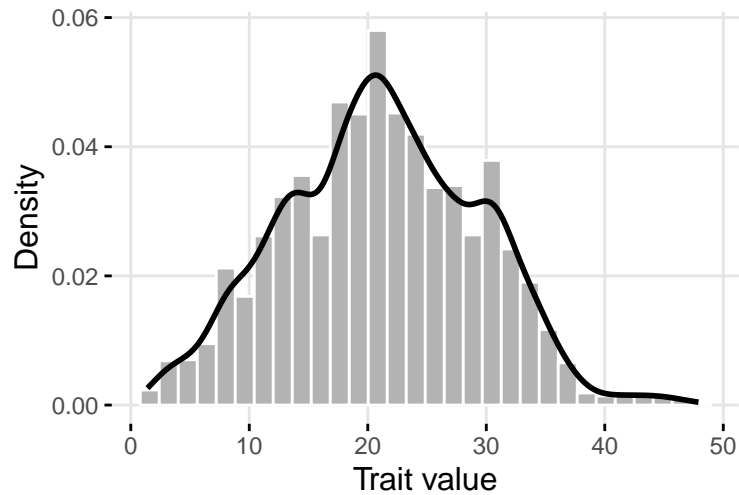
Sinusoidal



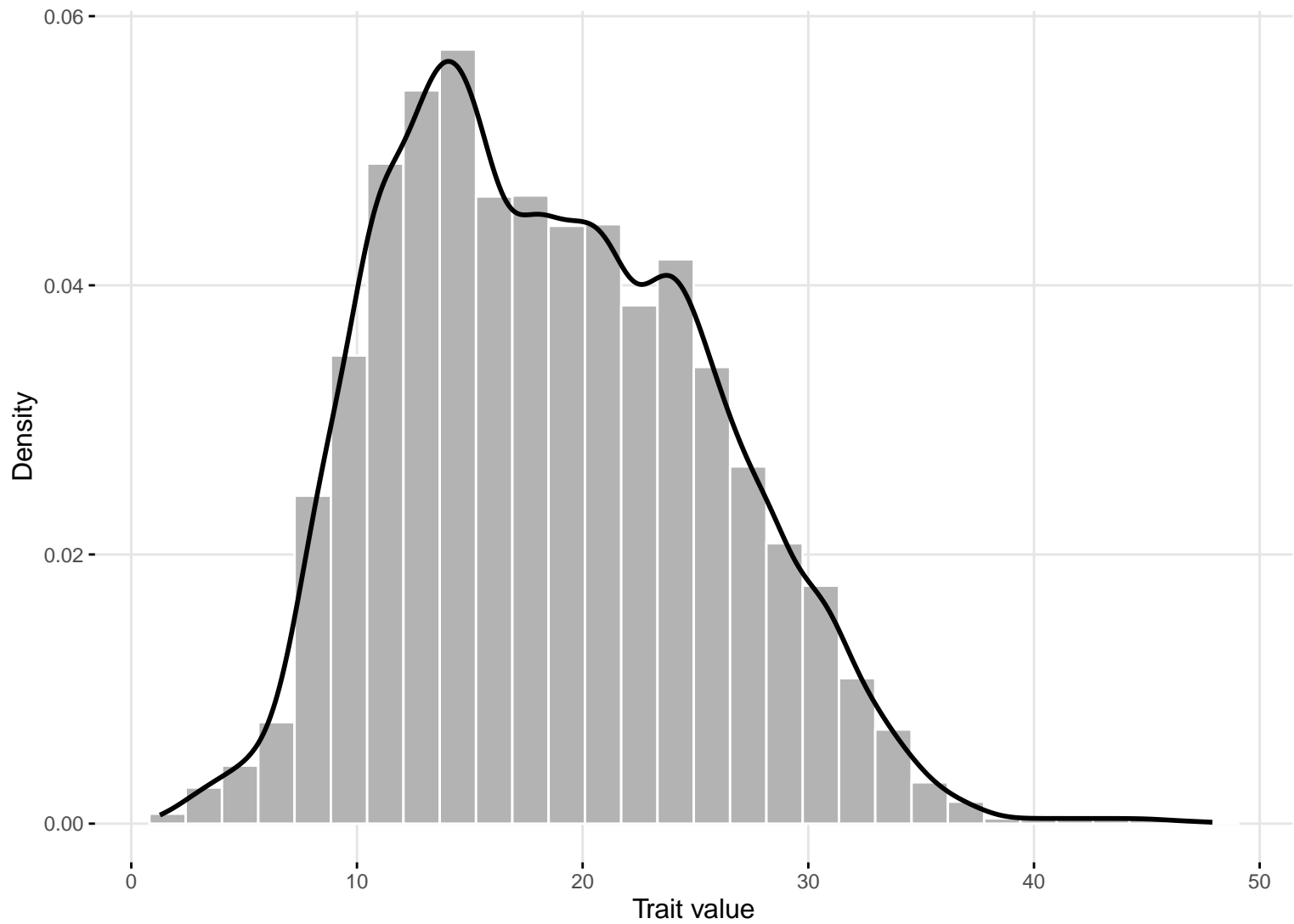
Wave

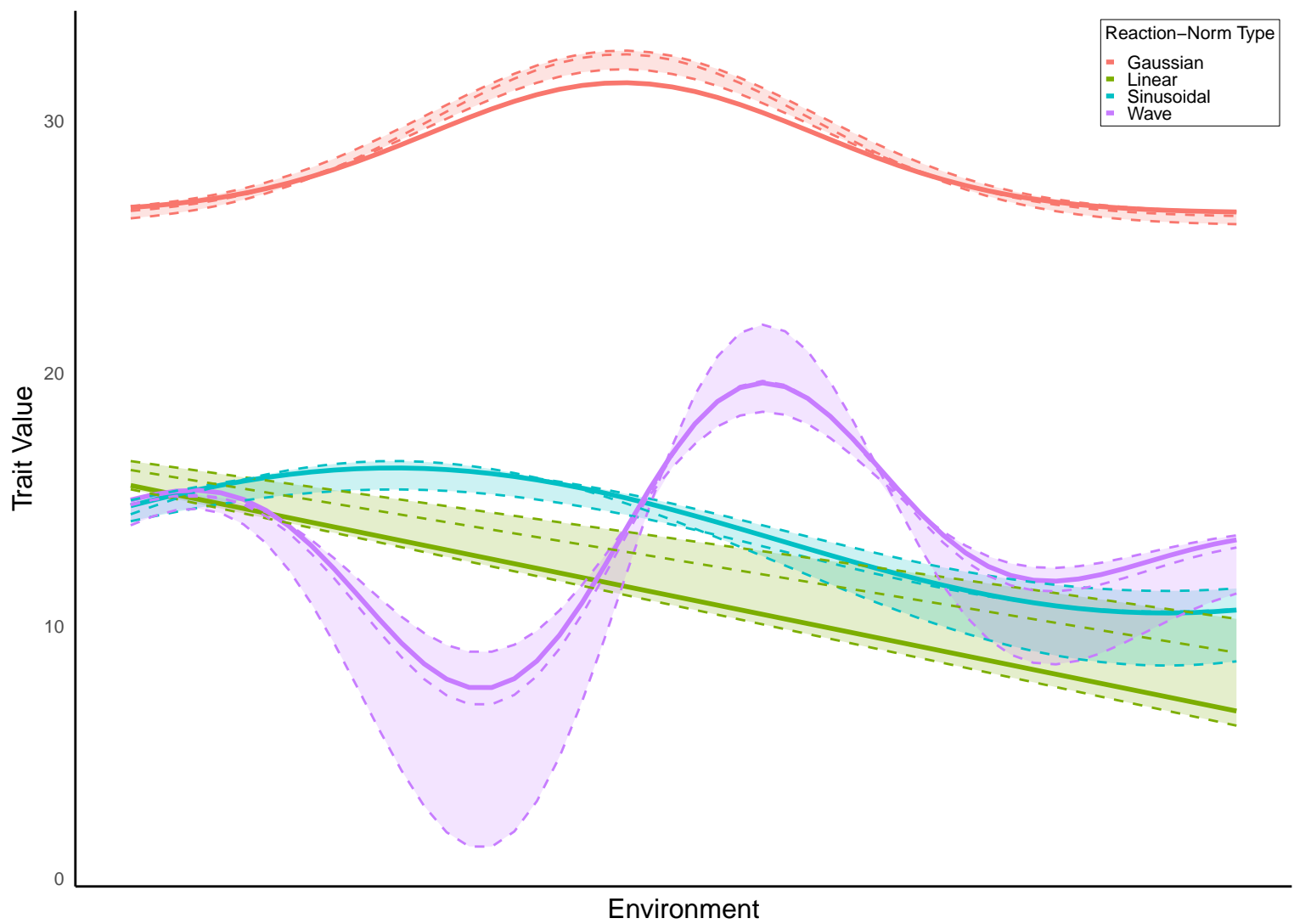


Linear

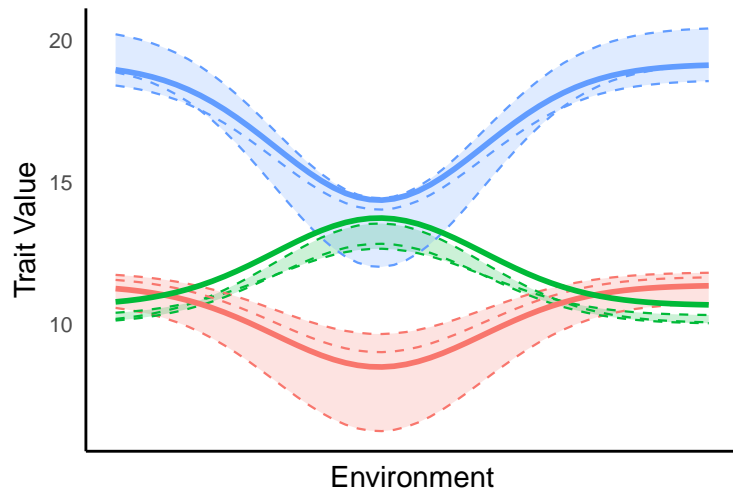


Combined Genotype Forms

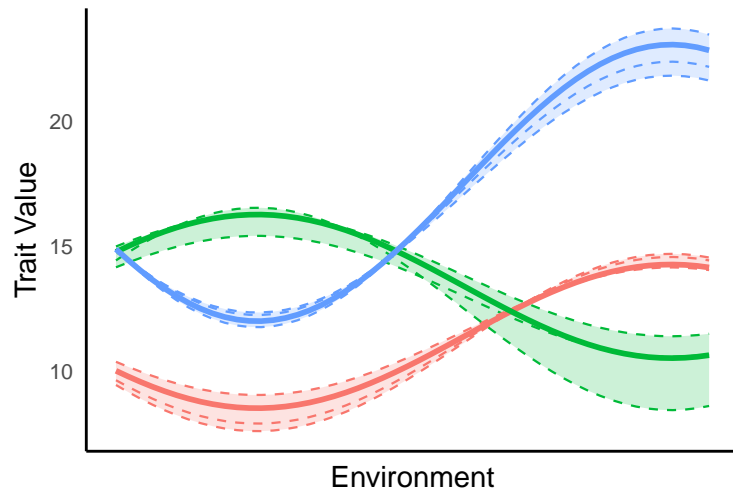




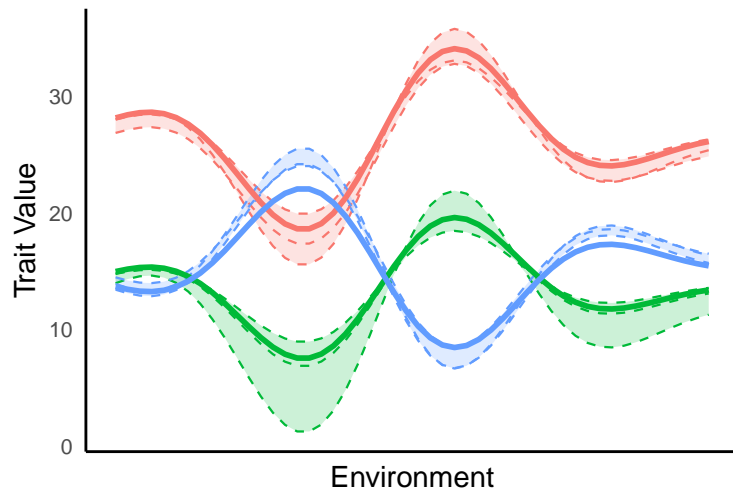
Gaussian



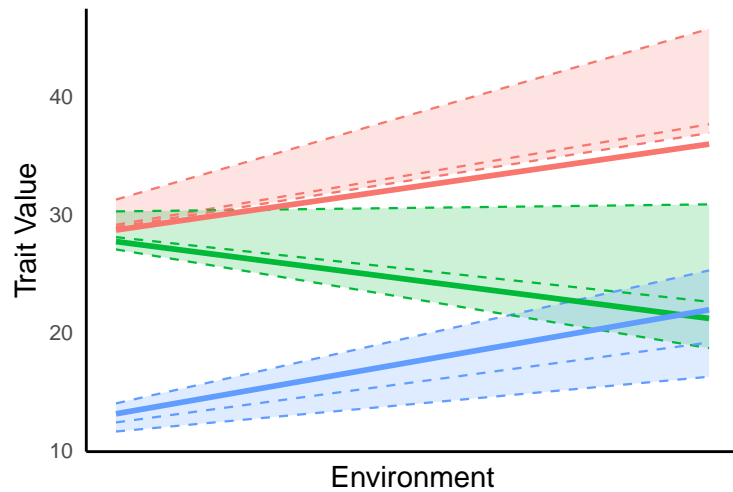
Sinusoidal



Wave

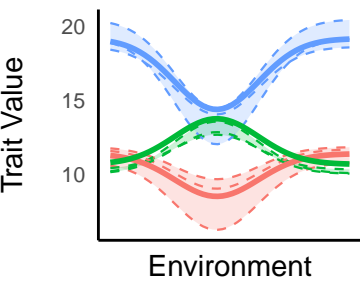


Linear

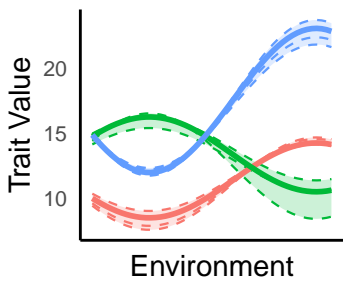


Reaction-Norm Profiles & Trait Distributions

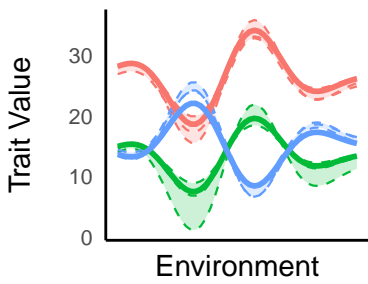
Gaussian



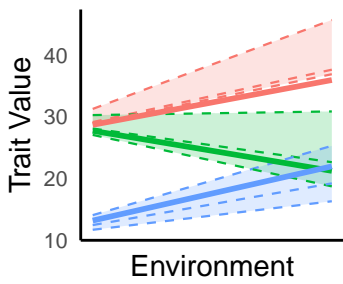
Sinusoidal



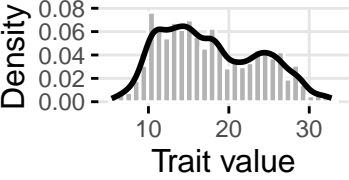
Wave



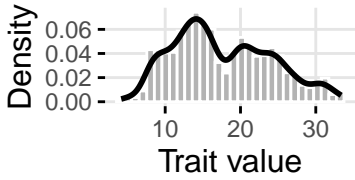
Linear



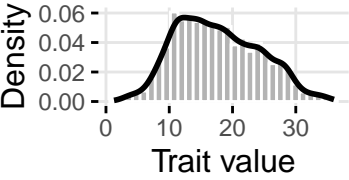
Gaussian



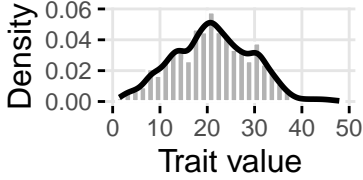
Sinusoidal



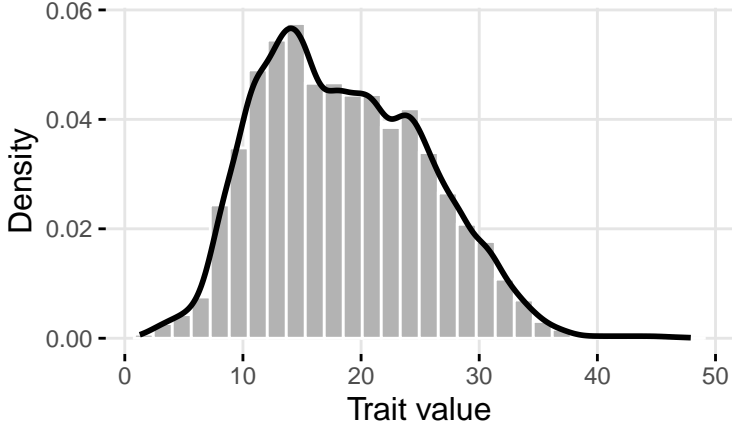
Wave



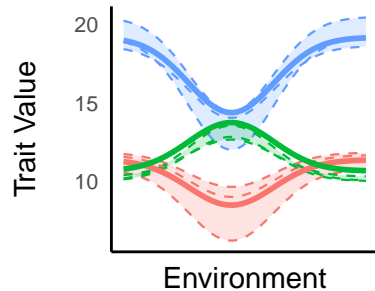
Linear



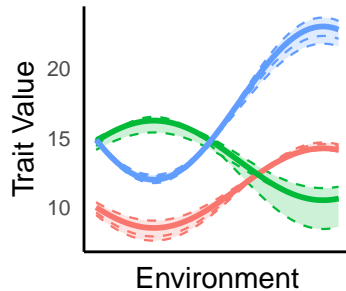
Combined Genotype Forms



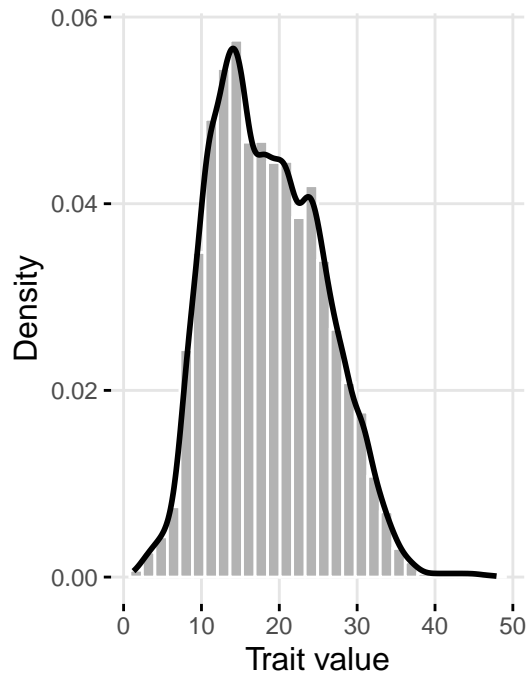
Gaussian



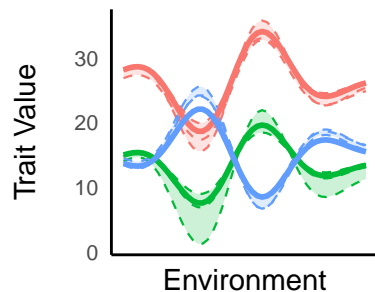
Sinusoidal



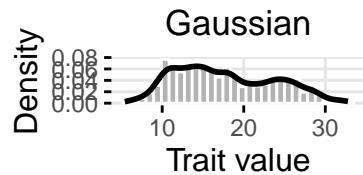
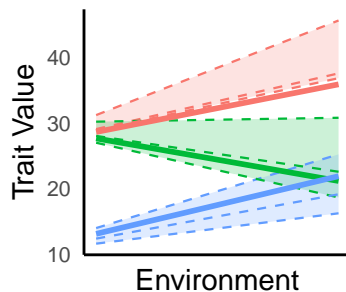
Combined Genotype Forms



Wave



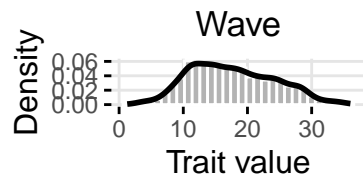
Linear



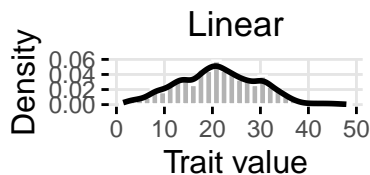
Gaussian



Sinusoidal



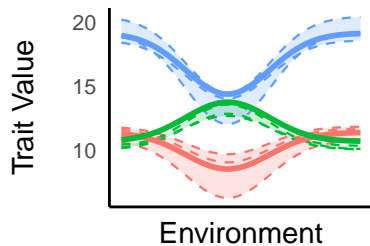
Wave



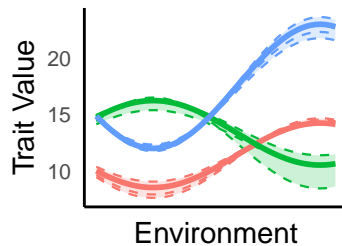
Linear

Alt B: Combined Above, Panels Below

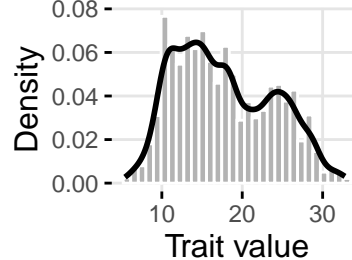
Gaussian



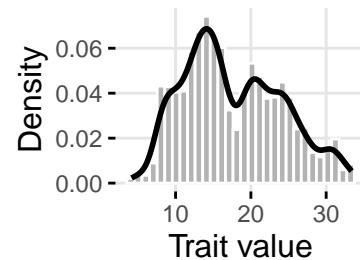
Sinusoidal



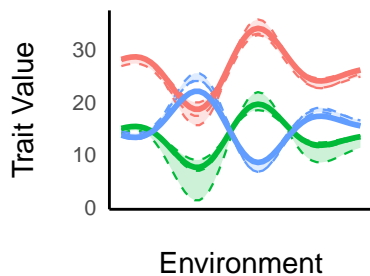
Gaussian



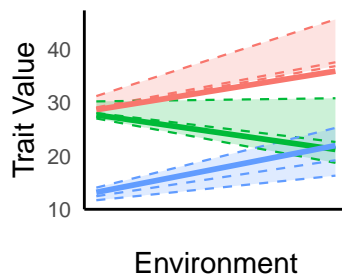
Sinusoidal



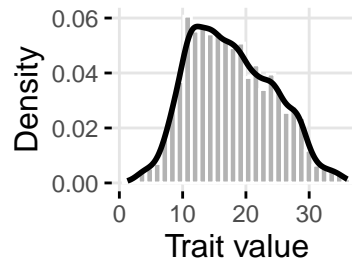
Wave



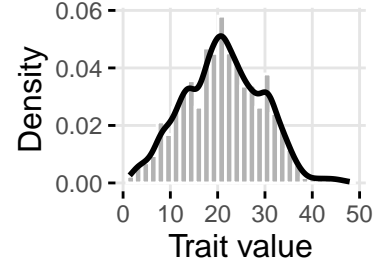
Linear



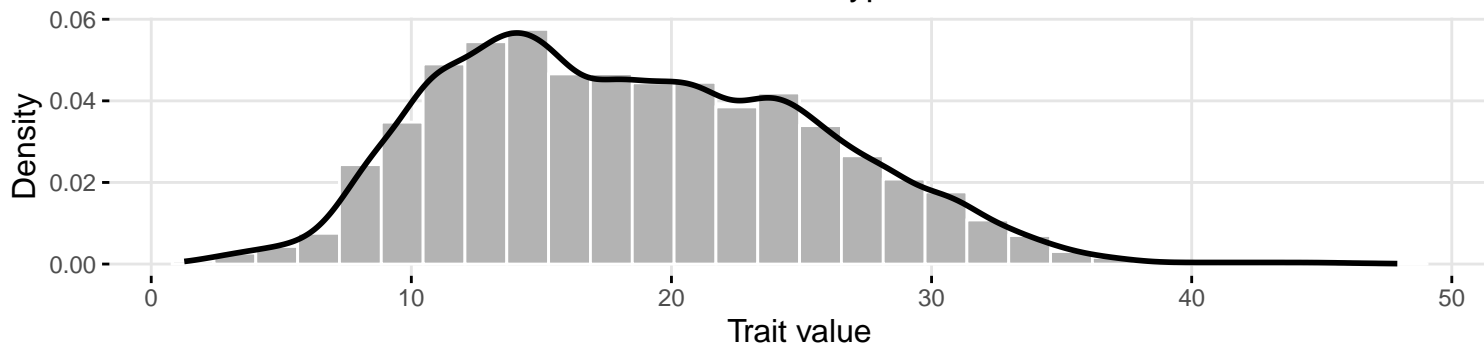
Wave



Linear



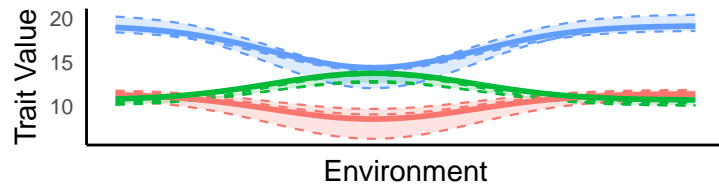
Combined Genotype Forms



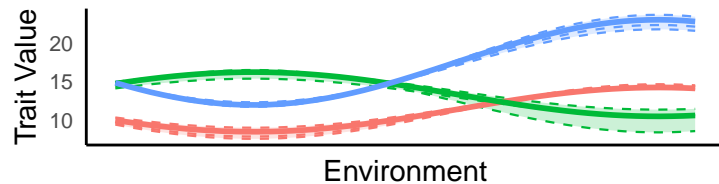
Reaction-Norm Profiles & Trait Distributions

A

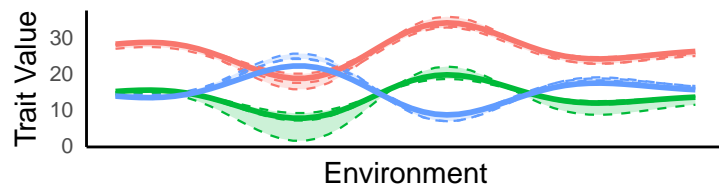
Gaussian



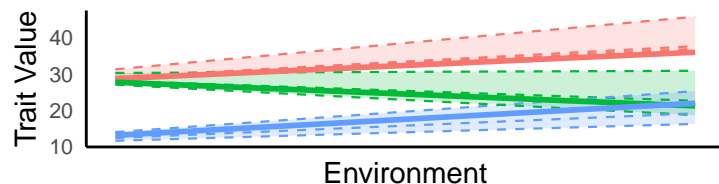
Sinusoidal



Wave

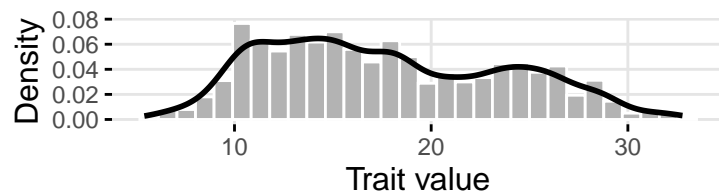


Linear

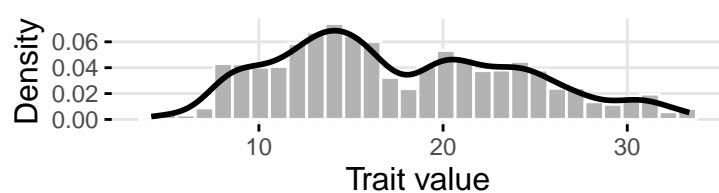


B

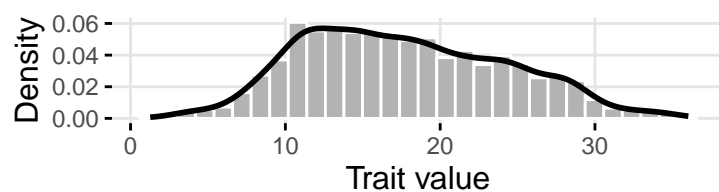
Gaussian



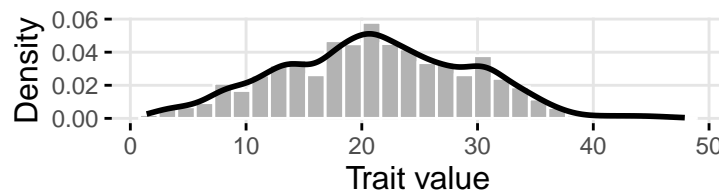
Sinusoidal



Wave

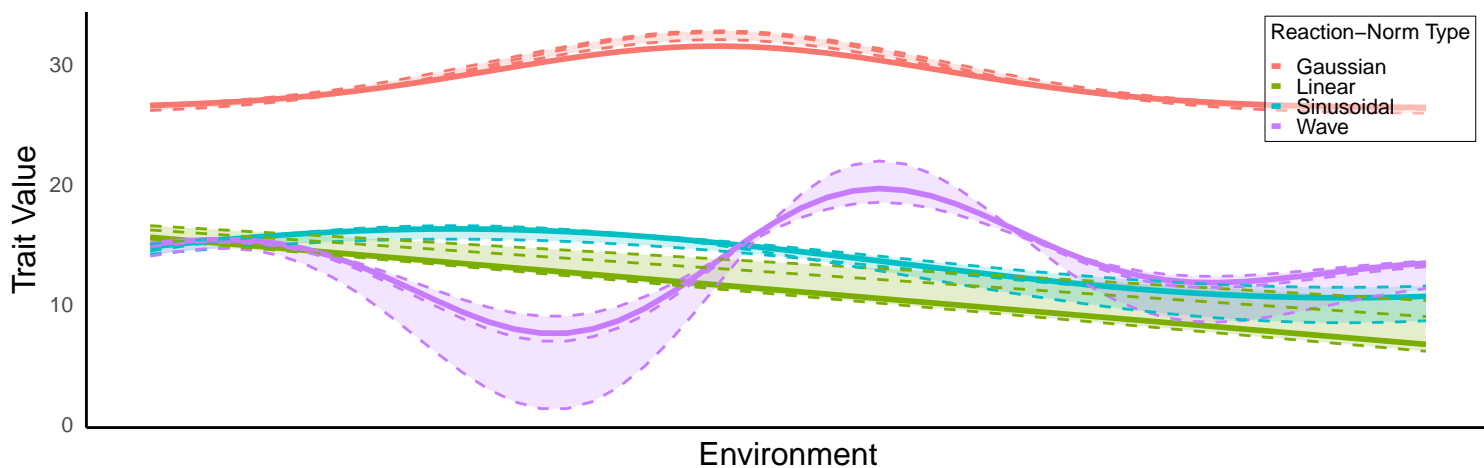


Linear



Reaction-Norm Profiles & Trait Distributions

A



B

