

Score: CV_t (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

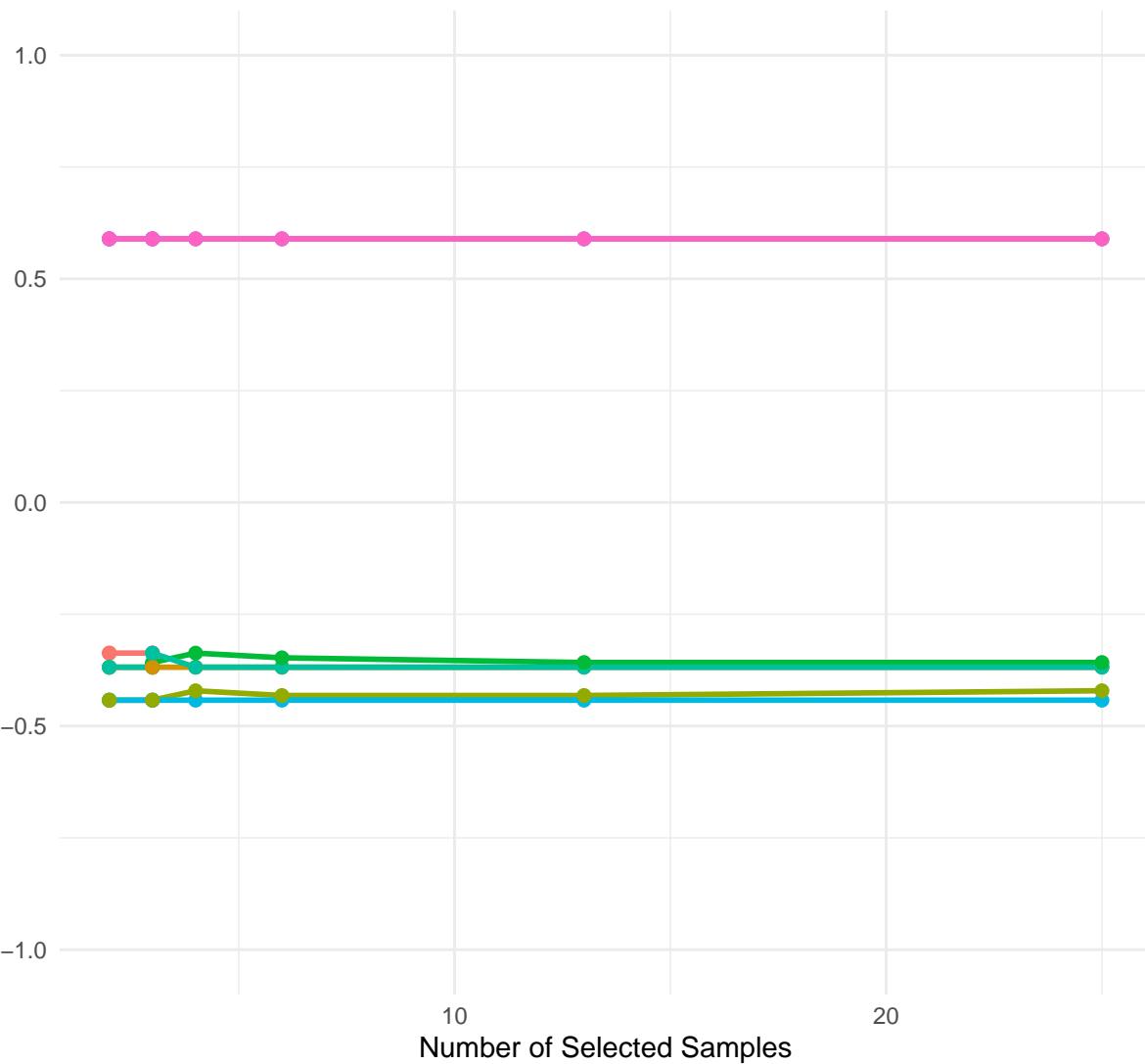
range

variance



Score: CV_t (Local, partial1 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

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

Score: CV_t (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

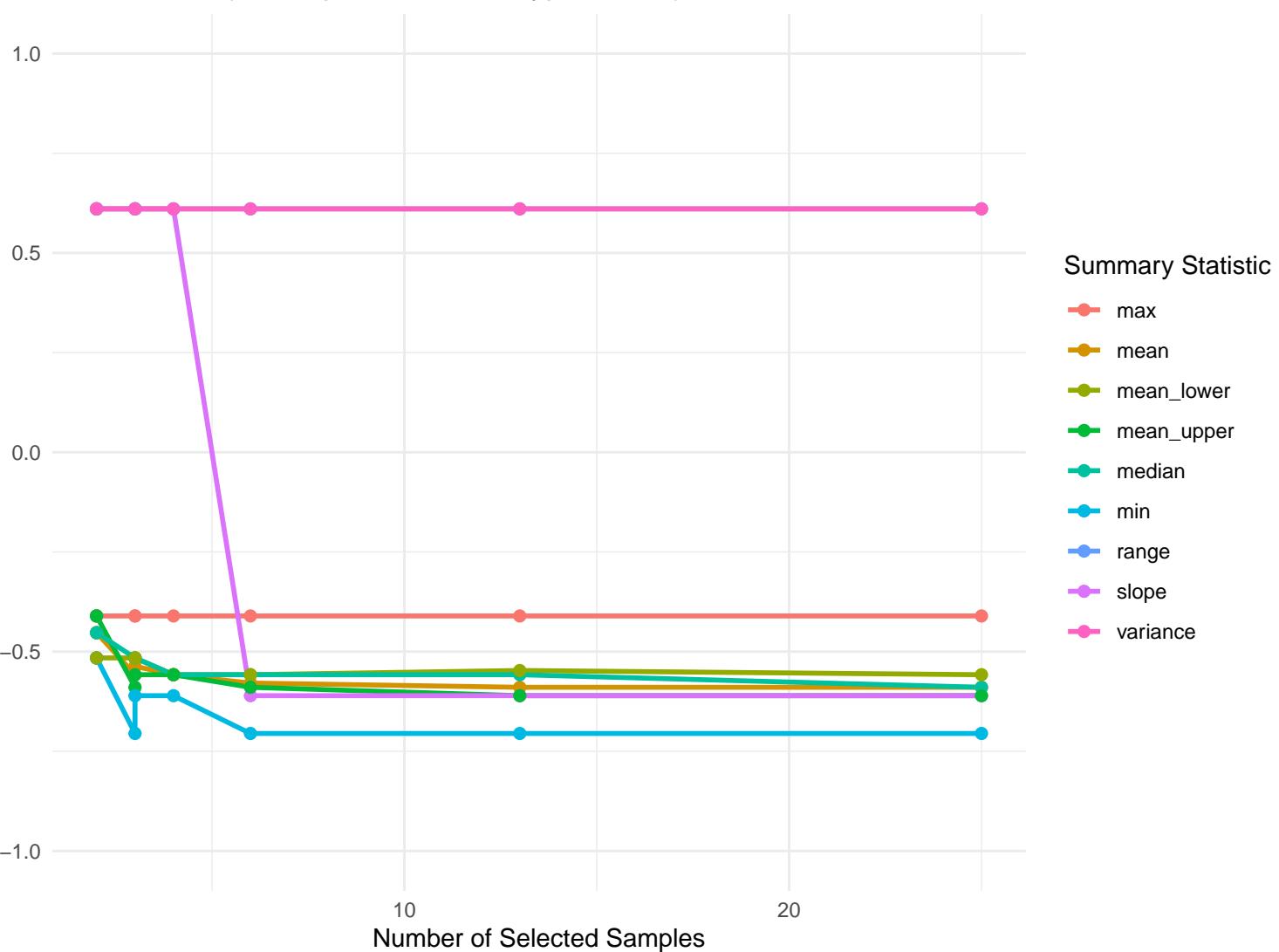
Summary Statistic

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



Score: CV_t (Local, partial1 , Genotype: wave)

Kendall Correlation



Score: RN (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

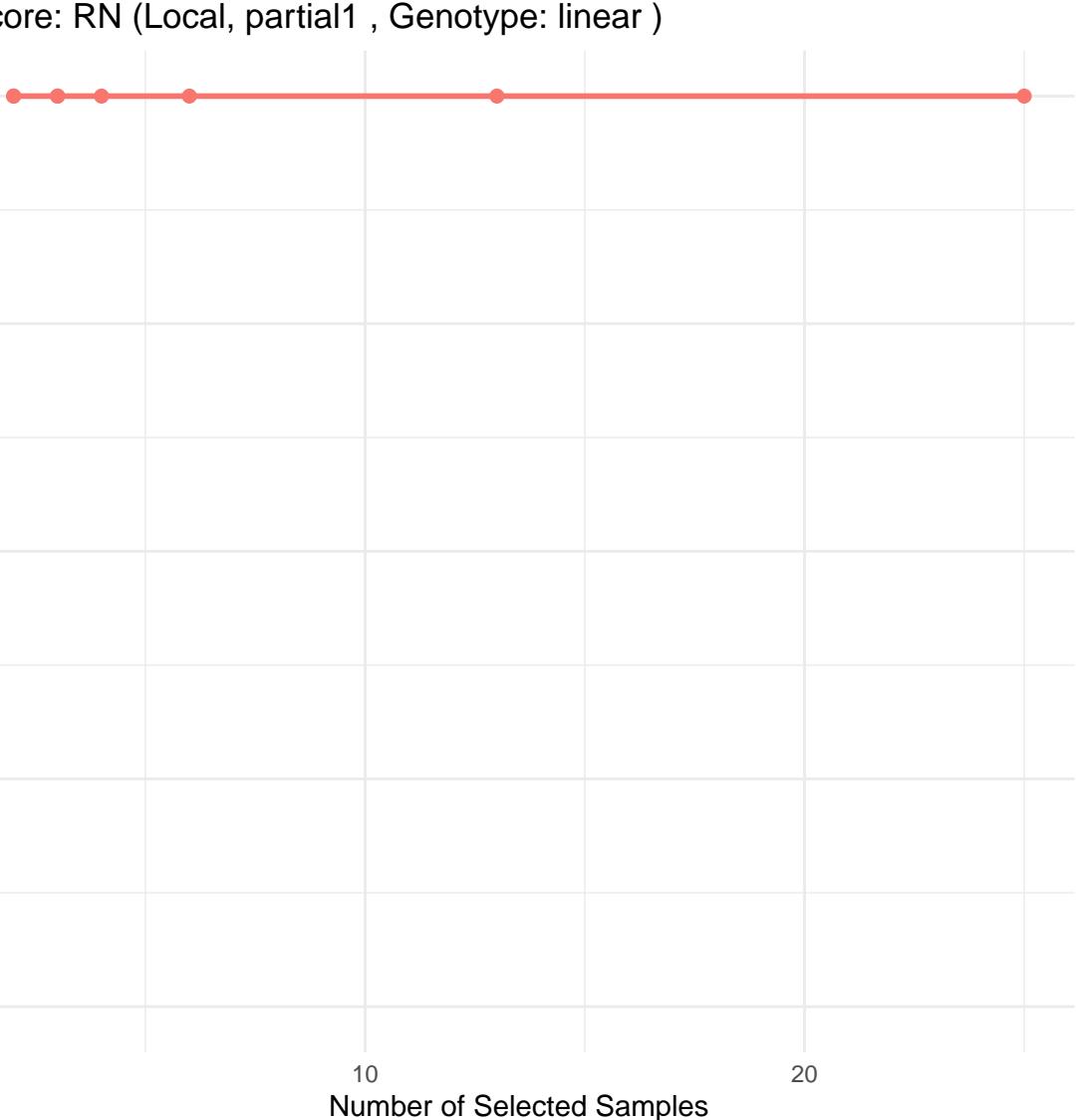
0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic
slope



Score: RN (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

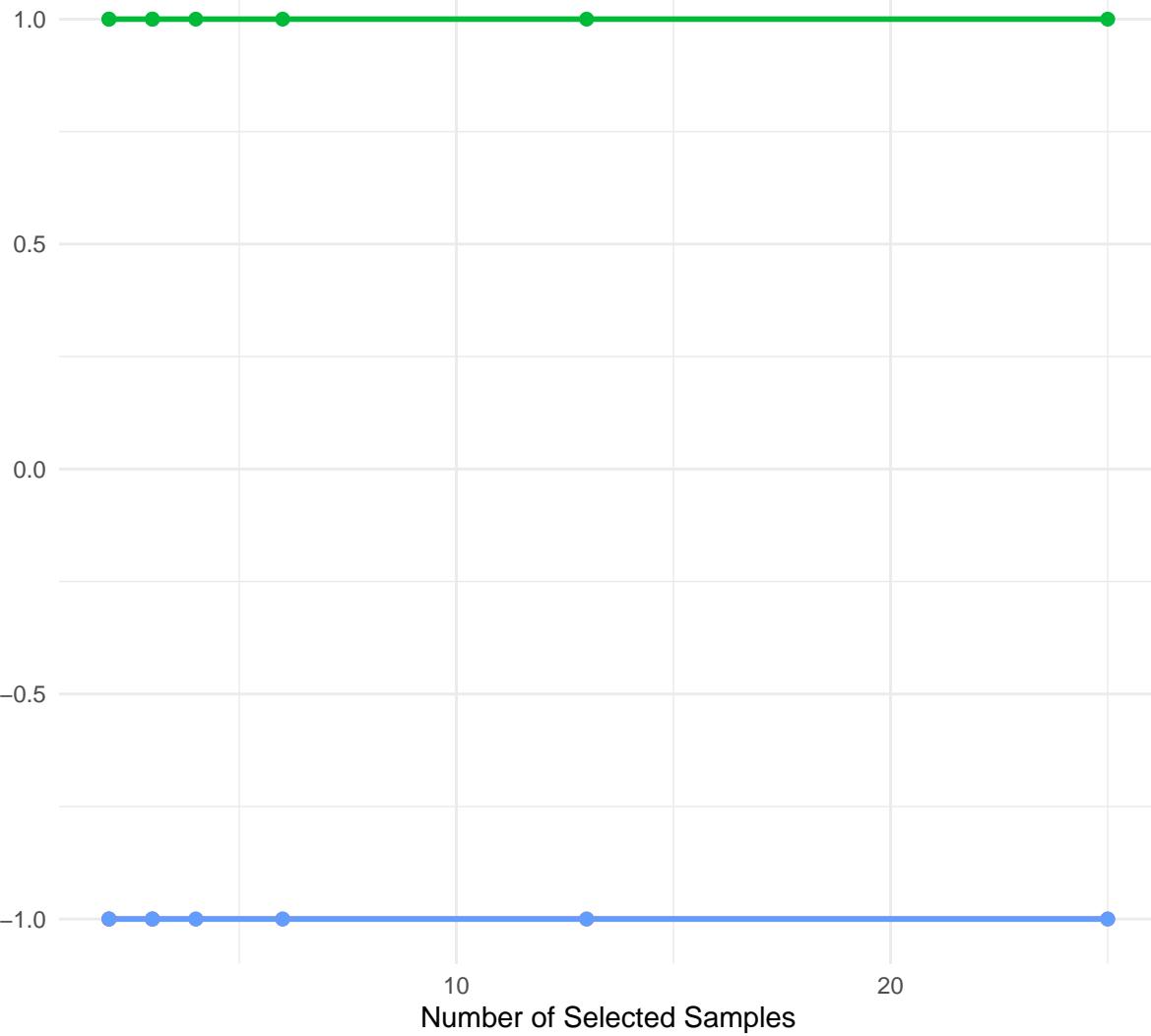
slope

variance



Score: RN (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

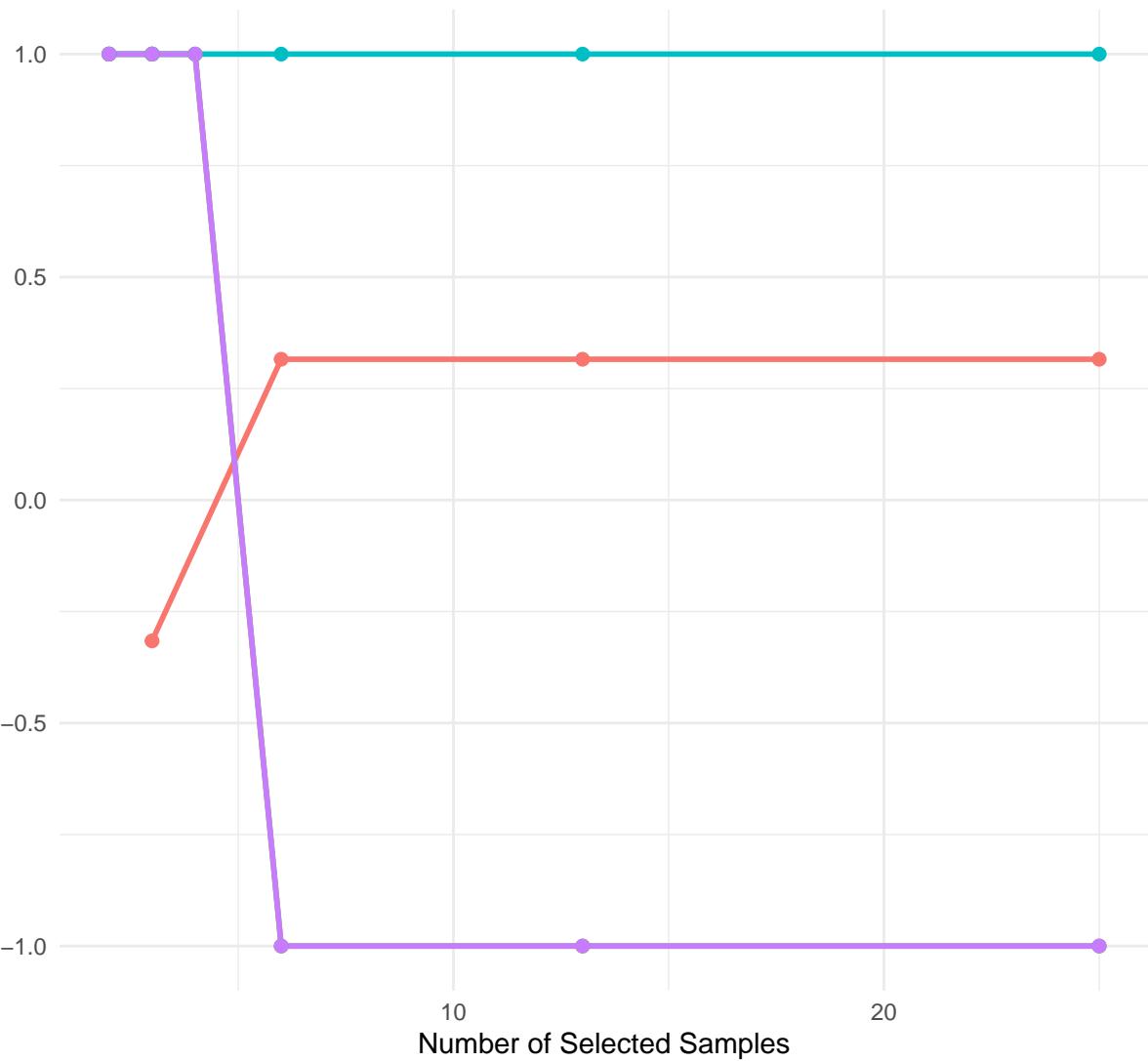
range

slope

variance

Score: RN (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

- min
- range
- slope
- variance

Score: RNN (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

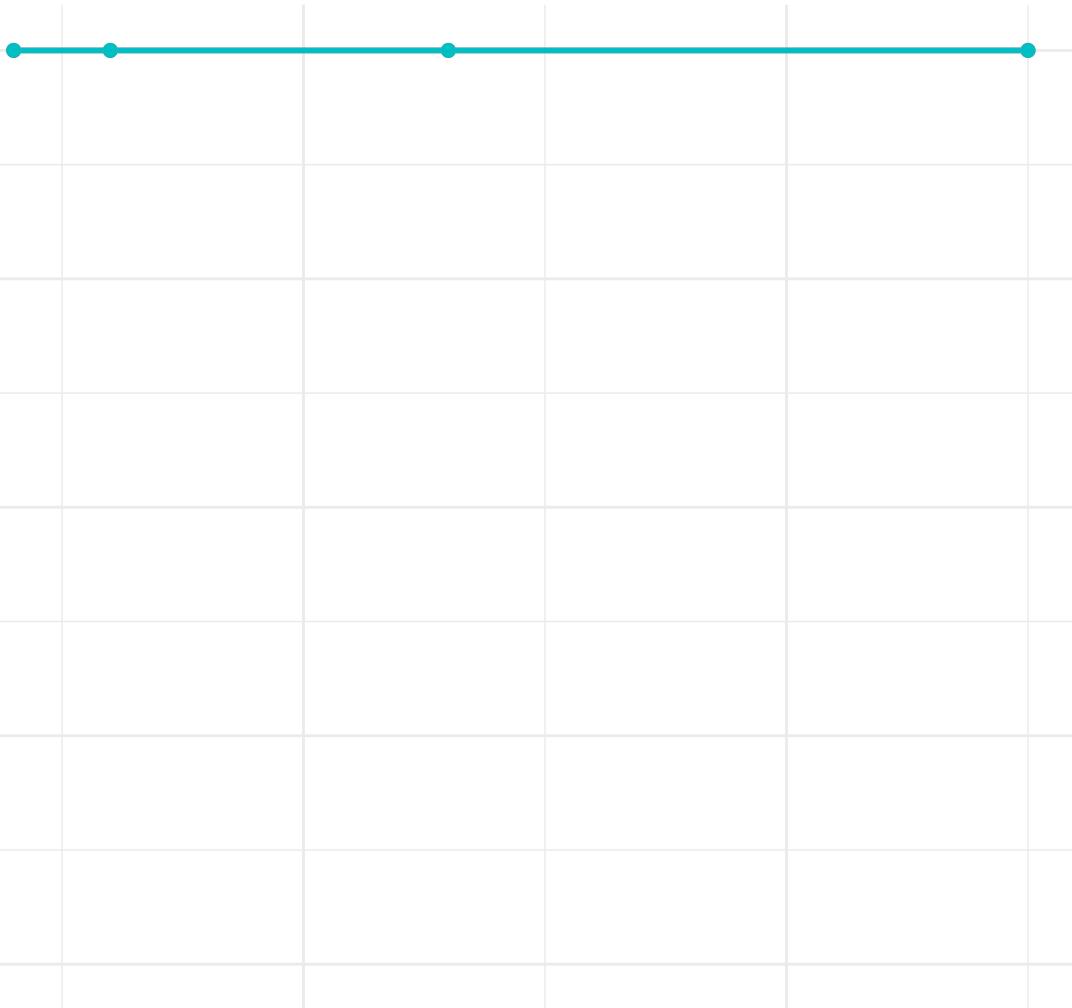
-1.0

Number of Selected Samples

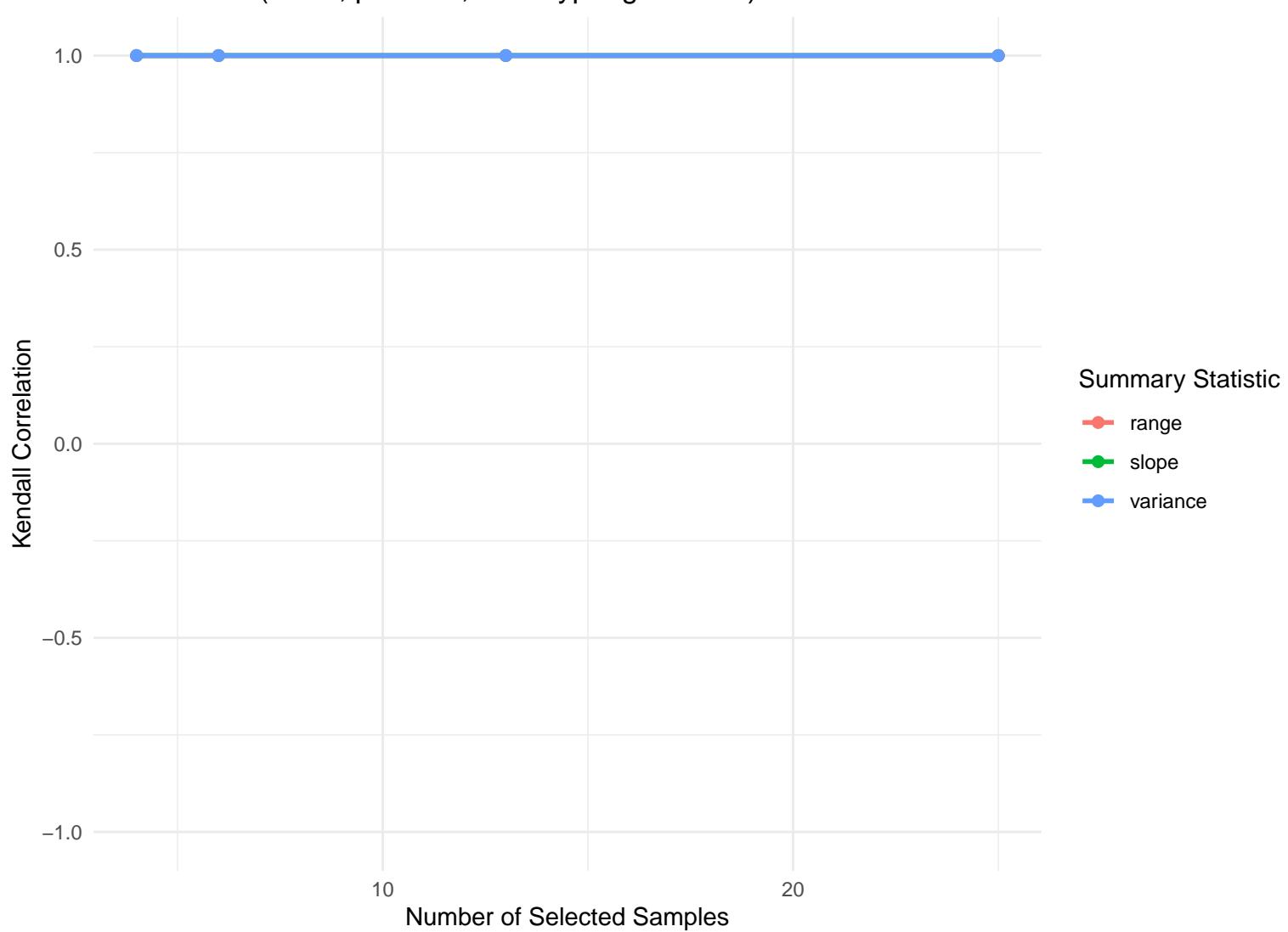
Summary Statistic

range

variance

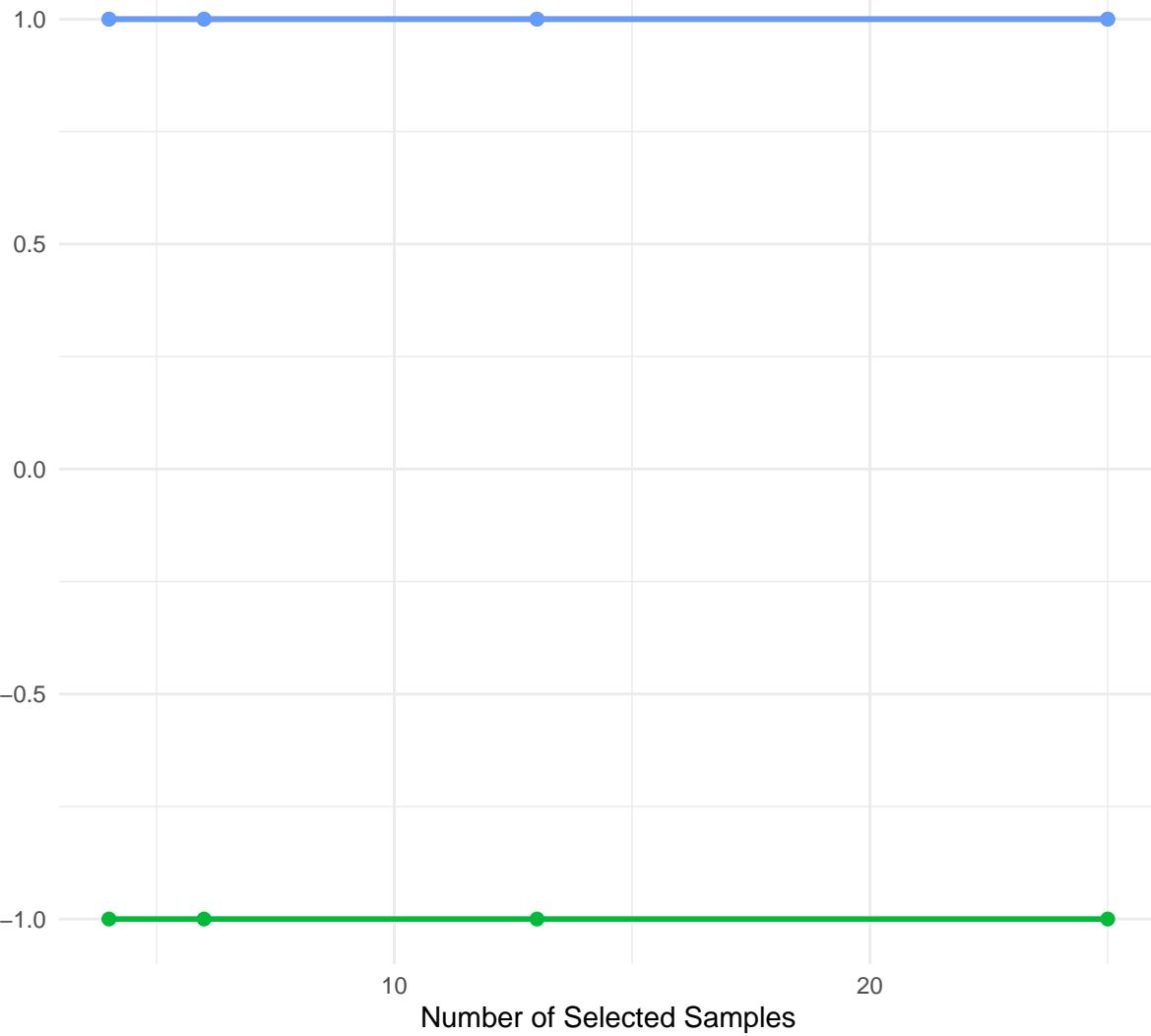


Score: RNN (Local, partial1 , Genotype: gaussian)



Score: RNN (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

range

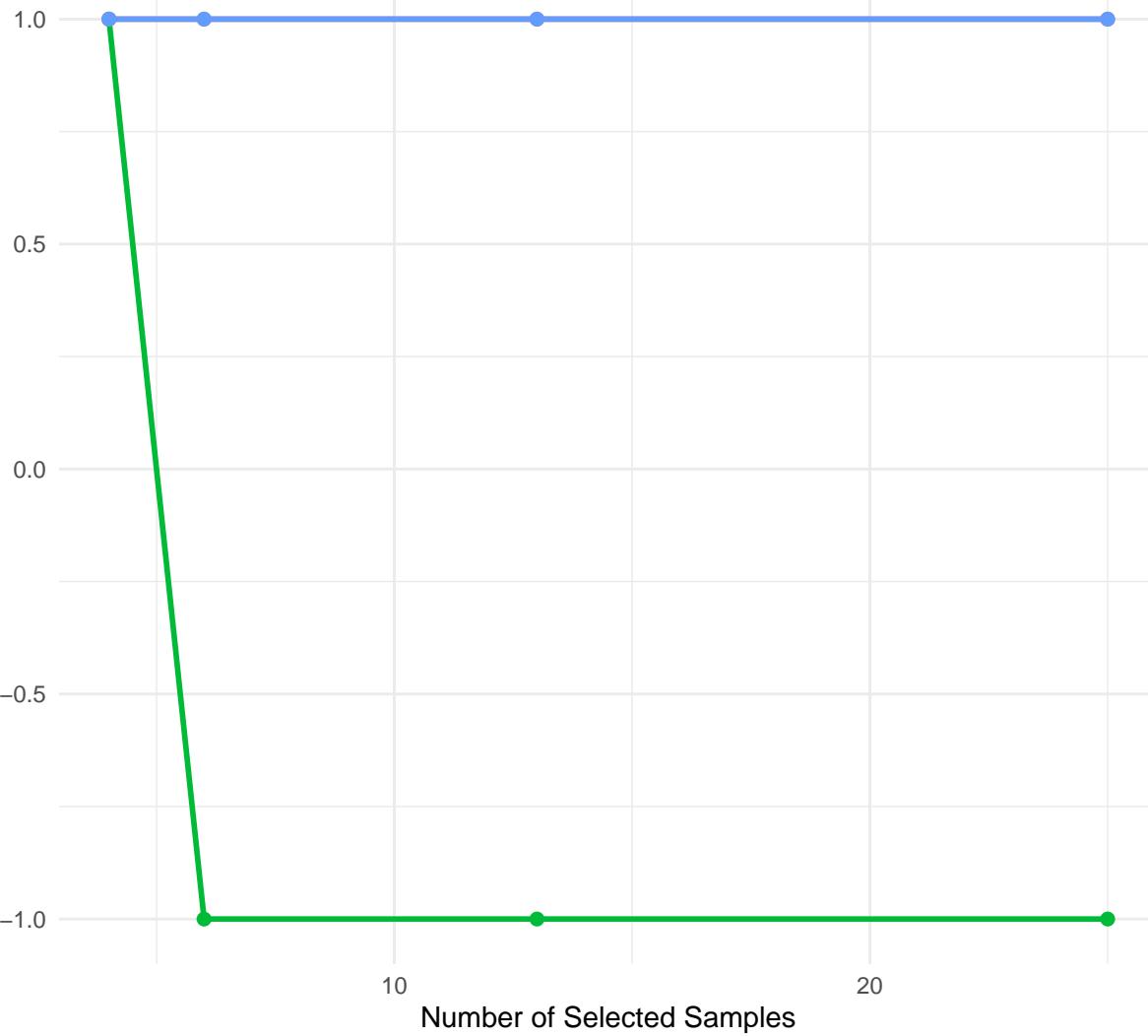
slope

variance

Number of Selected Samples

Score: RNN (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

range

slope

variance

Number of Selected Samples

Score: D_slope (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: D_slope (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

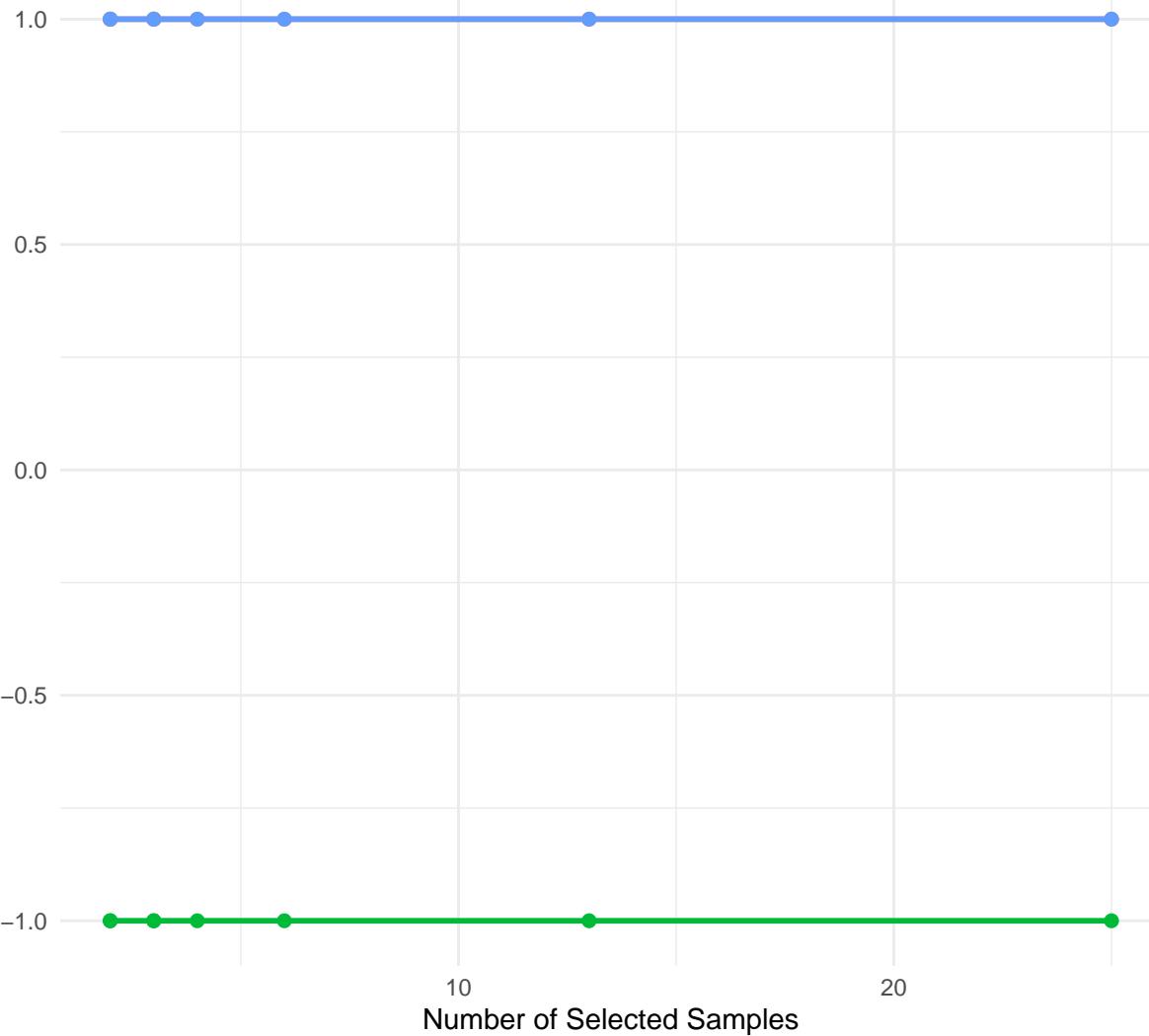
slope

variance



Score: D_slope (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

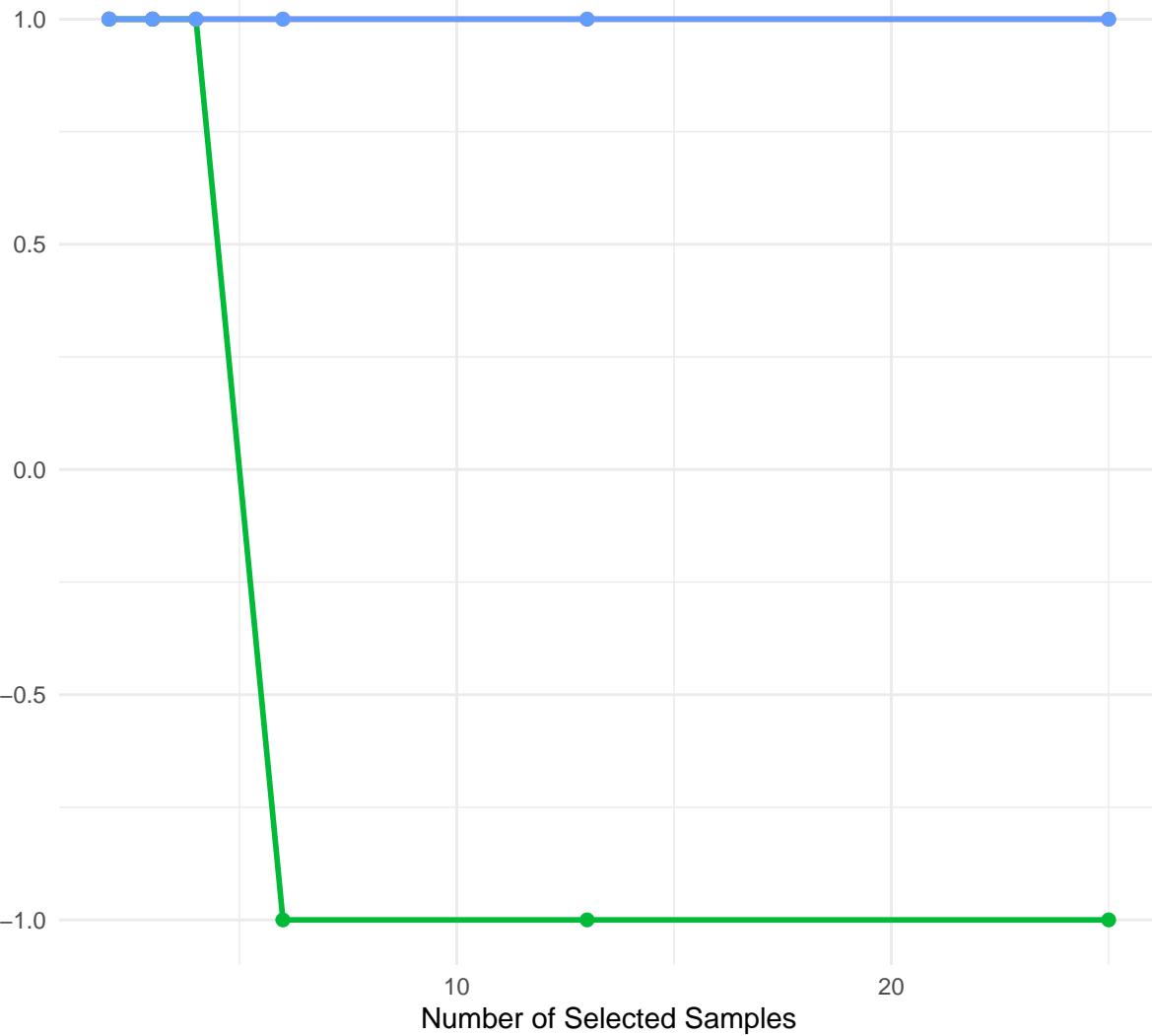
range

slope

variance

Score: D_slope (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

- range
- slope
- variance

Score: RC (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

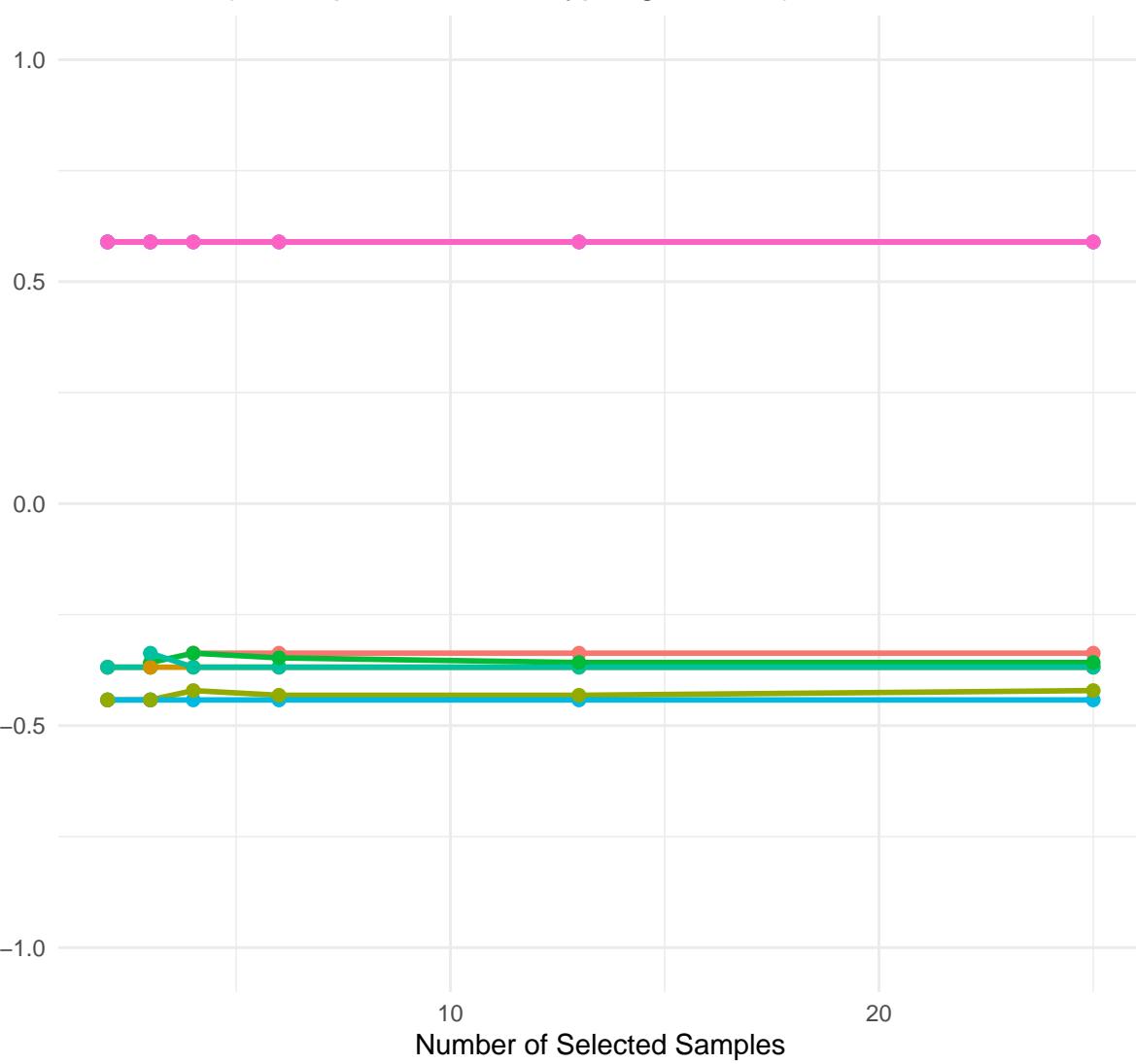
range

variance



Score: RC (Local, partial1 , Genotype: gaussian)

Kendall Correlation

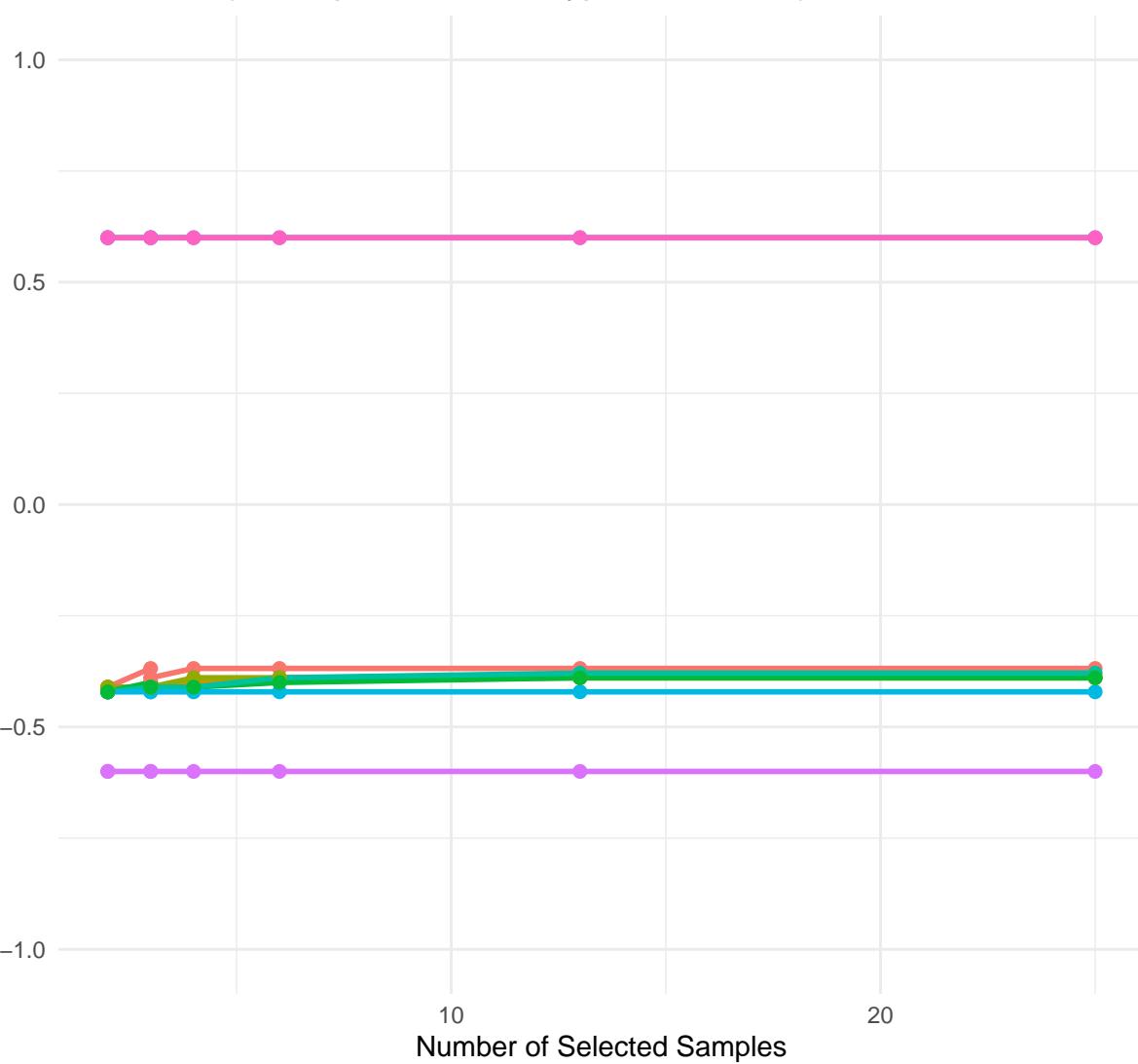


Summary Statistic

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

Score: RC (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

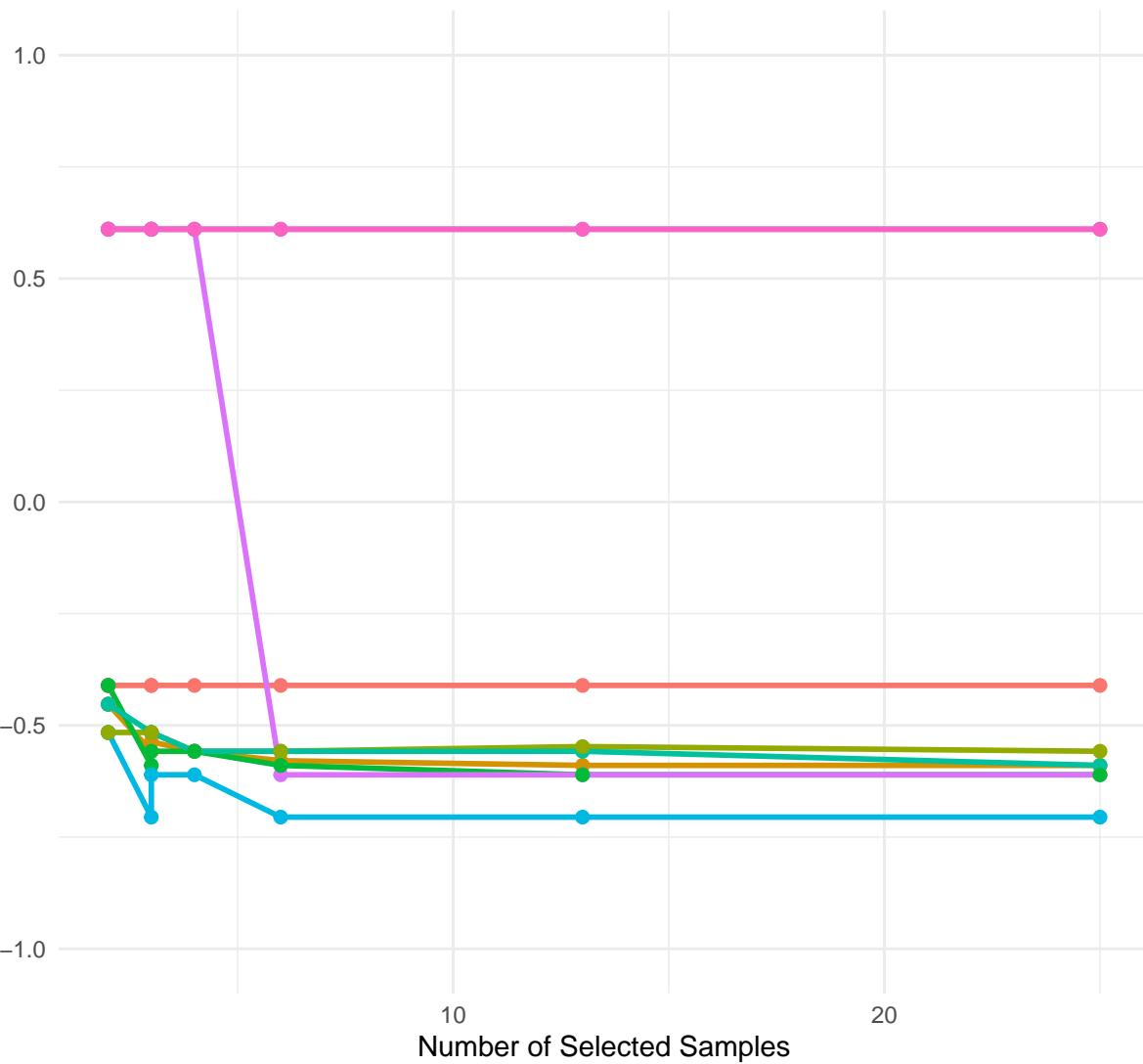


Summary Statistic

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

Score: RC (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: gPi (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

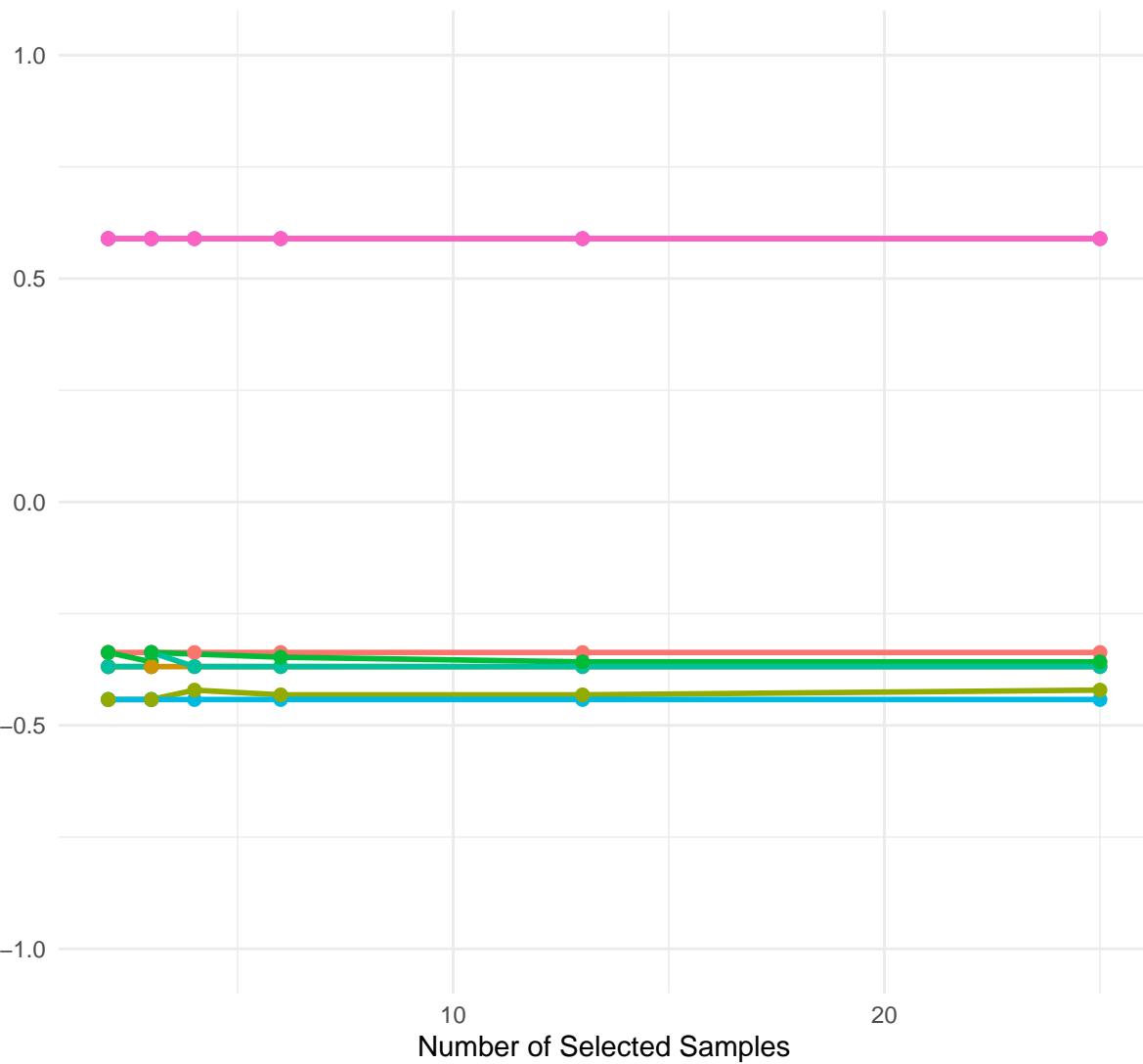
range

variance



Score: gPi (Local, partial1 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

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

Score: gPi (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

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

A horizontal pink line with circular markers at approximately 0.58, representing the variance statistic.

A horizontal green line with circular markers at approximately -0.45, representing the mean_upper statistic.

A horizontal blue line with circular markers at approximately -0.48, representing the median statistic.

A horizontal red line with circular markers at approximately -0.45, representing the max statistic.

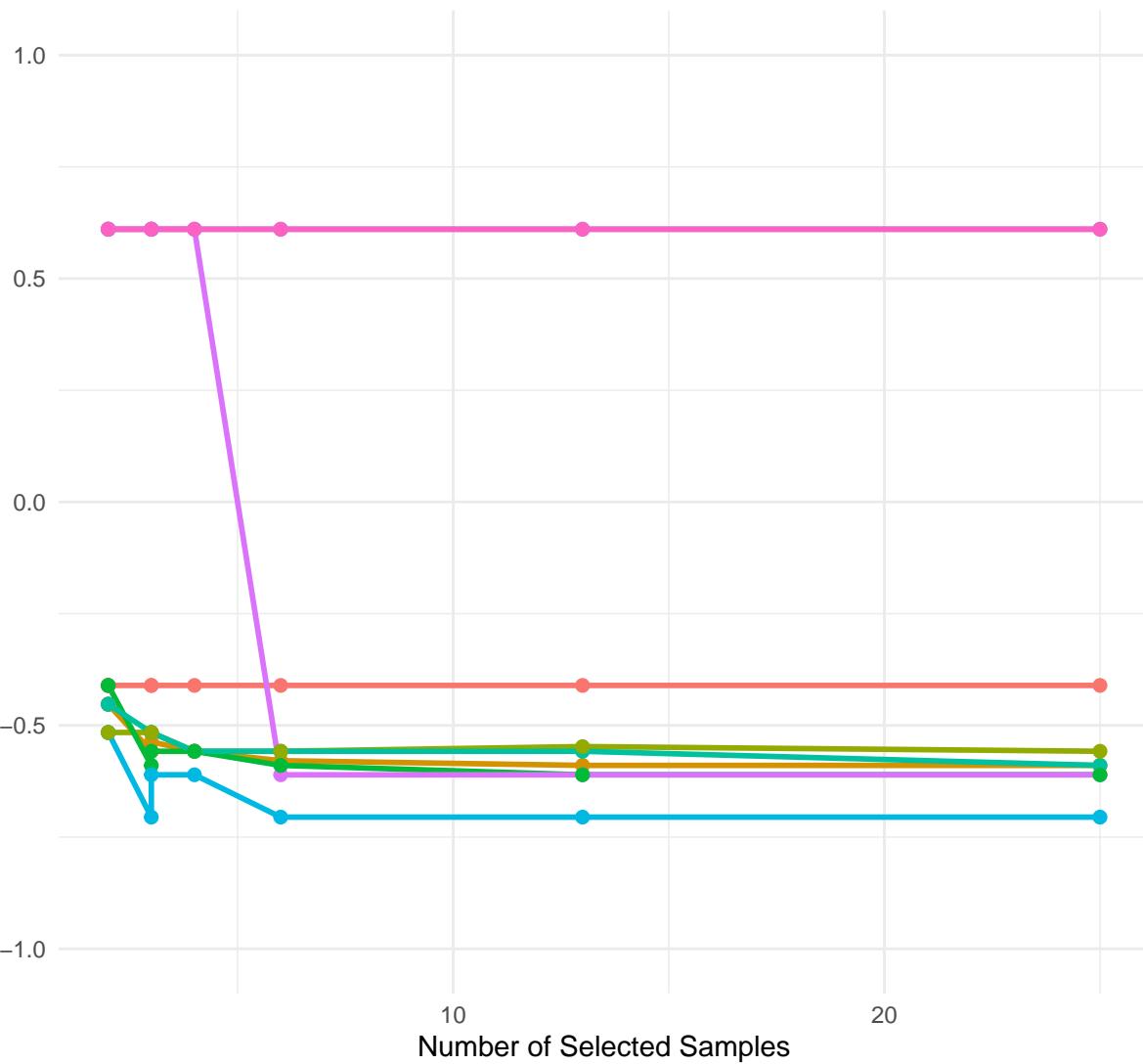
A horizontal orange line with circular markers at approximately -0.45, representing the mean statistic.

A horizontal yellow-green line with circular markers at approximately -0.45, representing the mean_lower statistic.

A horizontal purple line with circular markers at approximately -0.65, representing the slope statistic.

Score: gPi (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: PPF (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

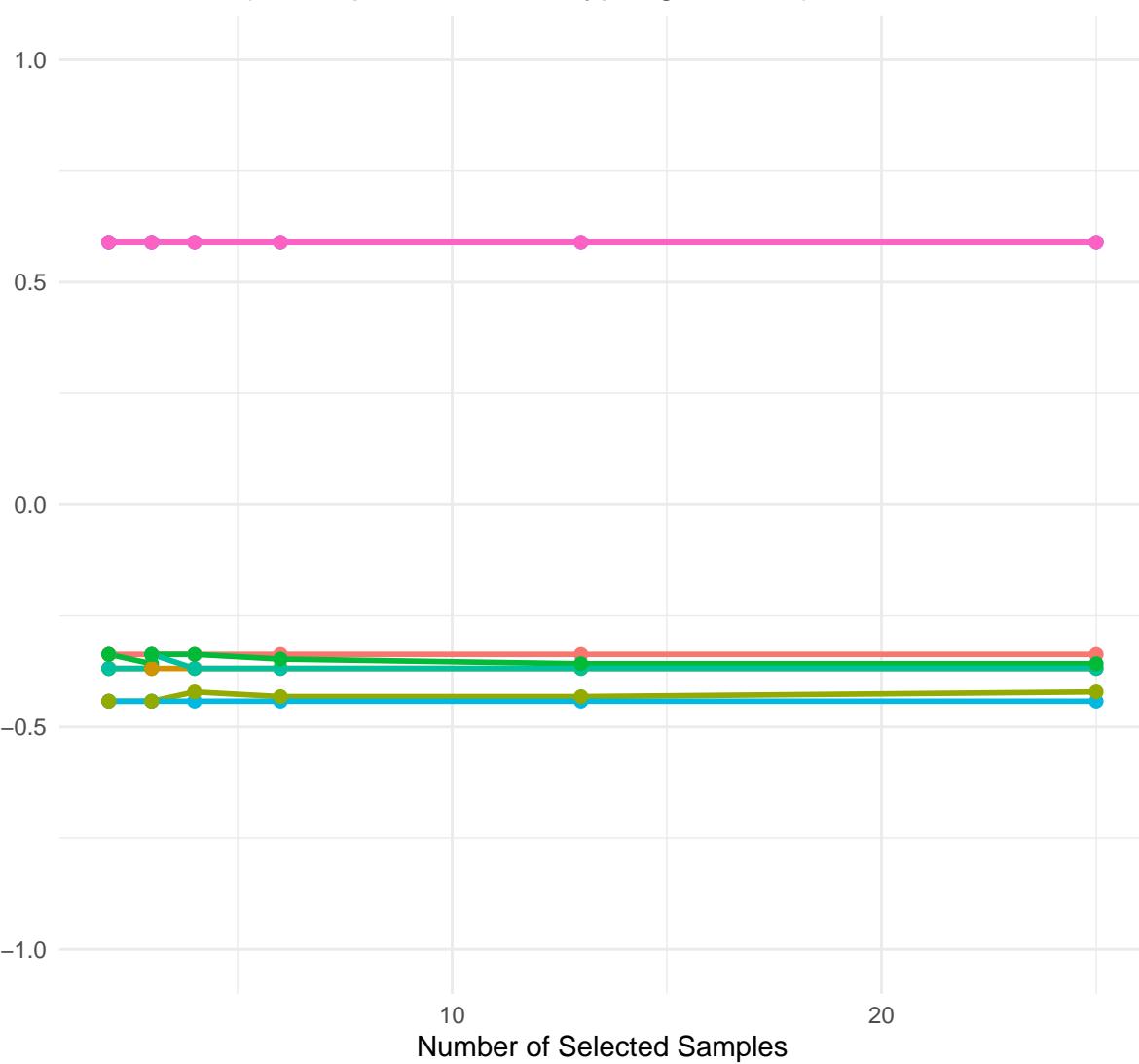
range

variance



Score: PPF (Local, partial1 , Genotype: gaussian)

Kendall Correlation

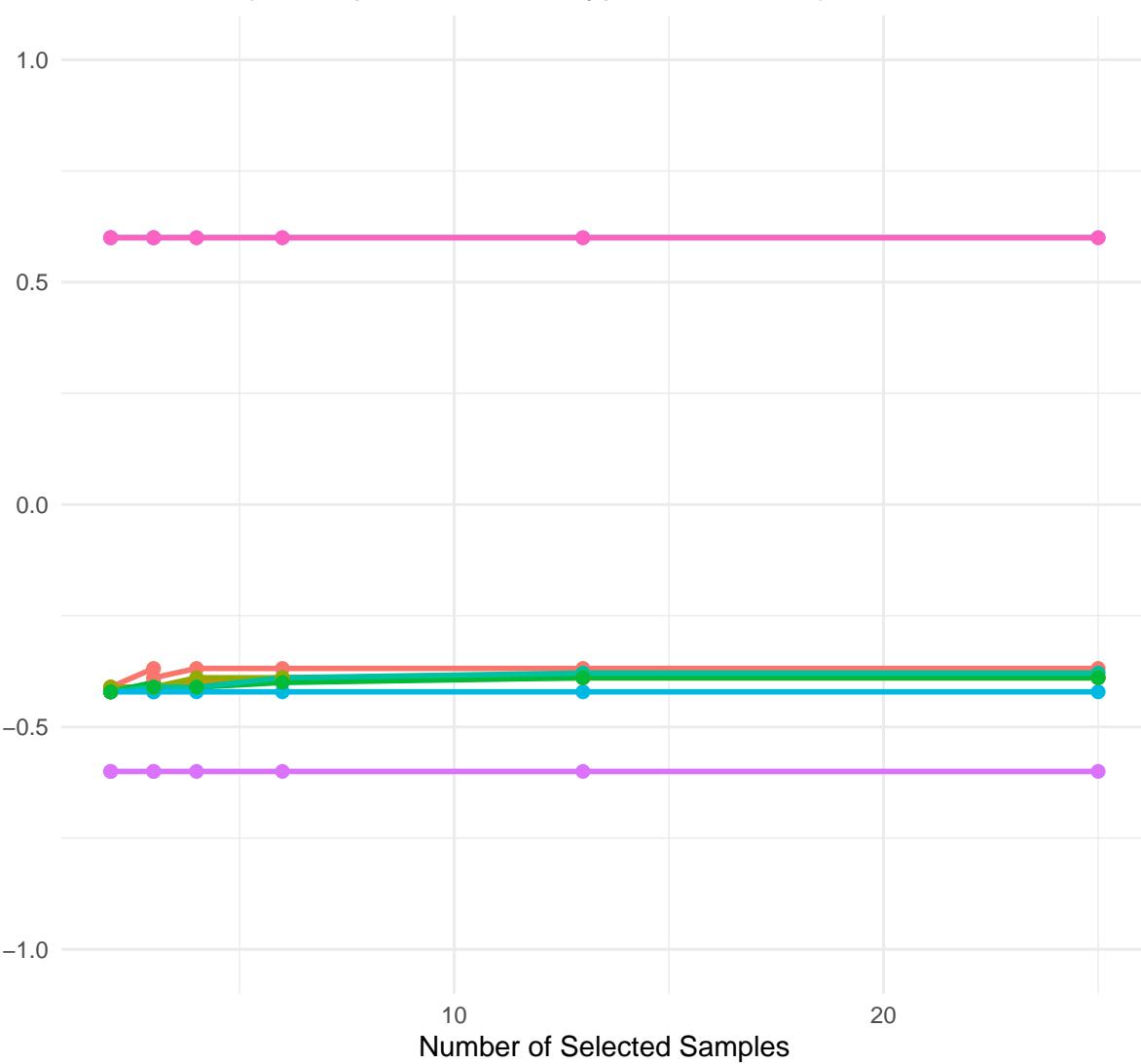


Summary Statistic

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

Score: PPF (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

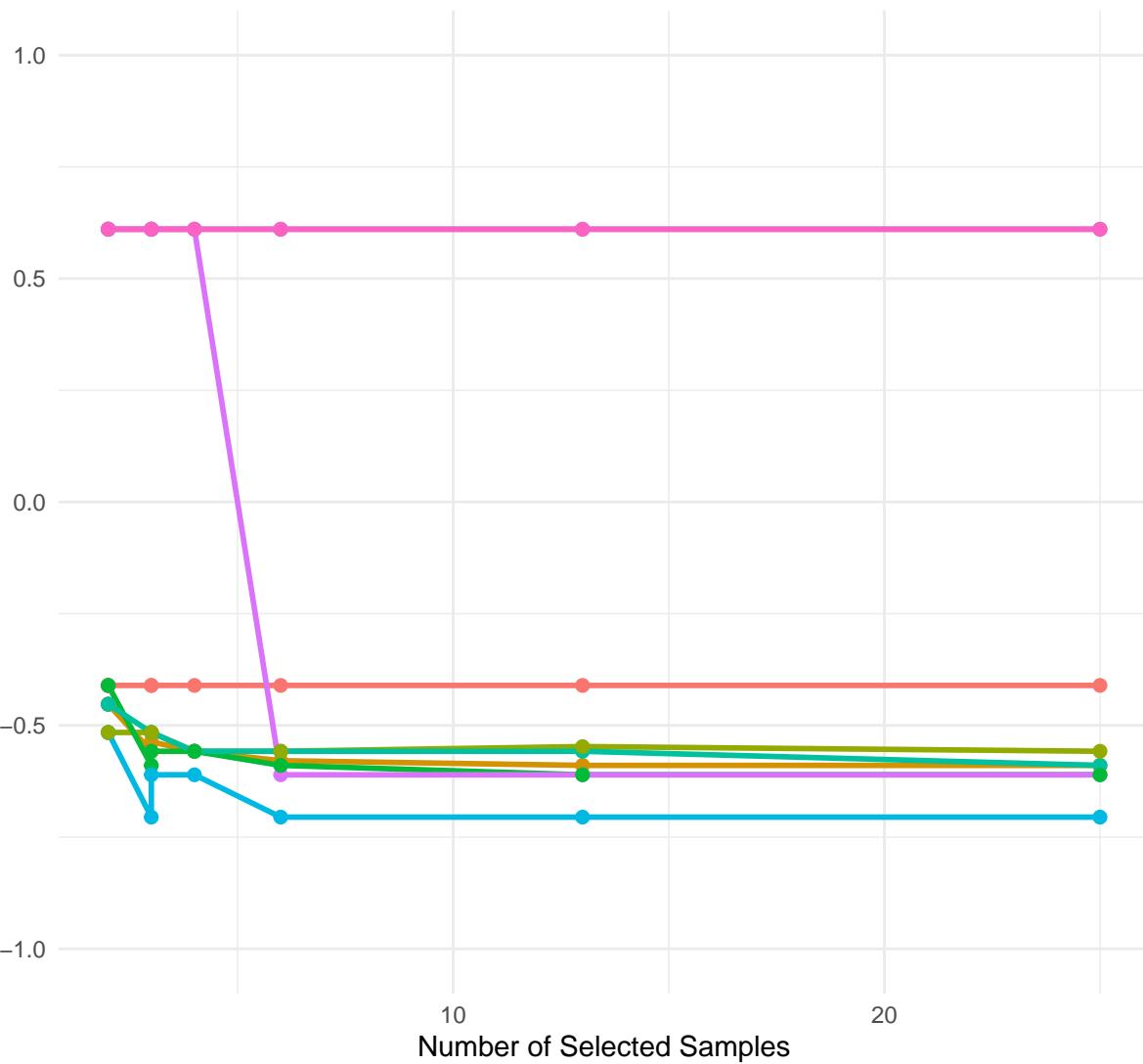


Summary Statistic

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

Score: PPF (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: PPi (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: PPi (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

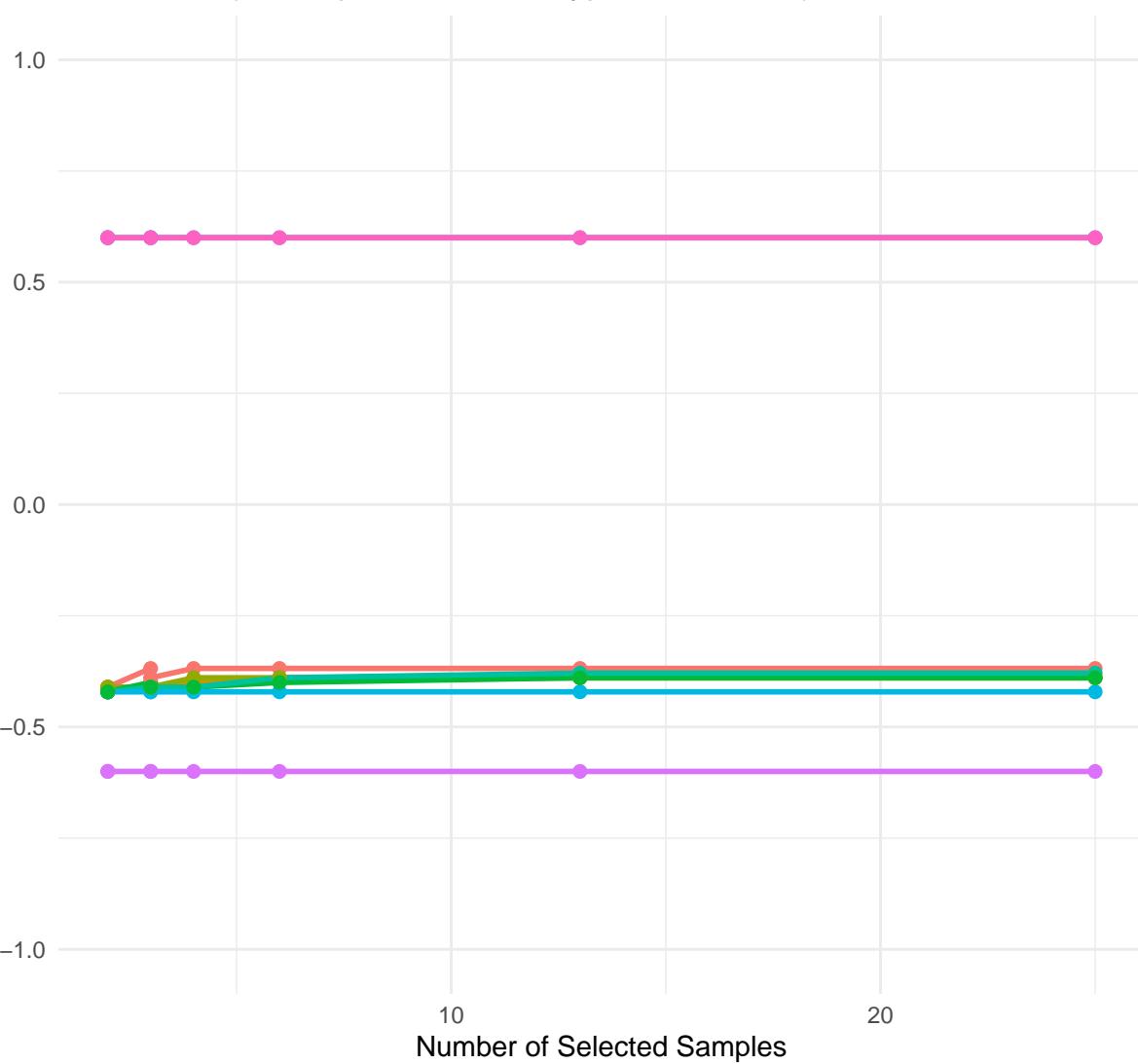
Summary Statistic

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



Score: PPi (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

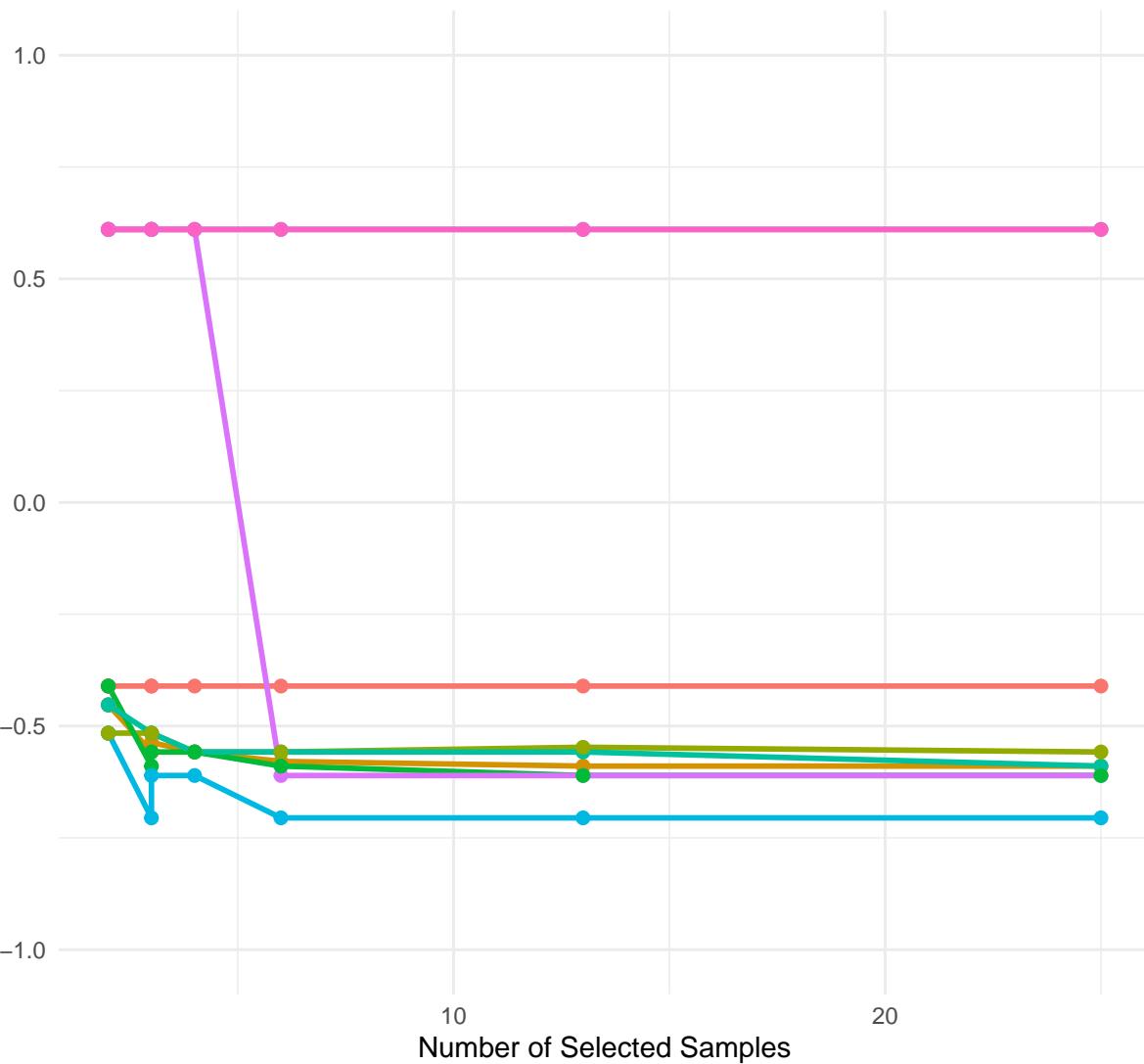


Summary Statistic

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

Score: PPi (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: Plmd (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

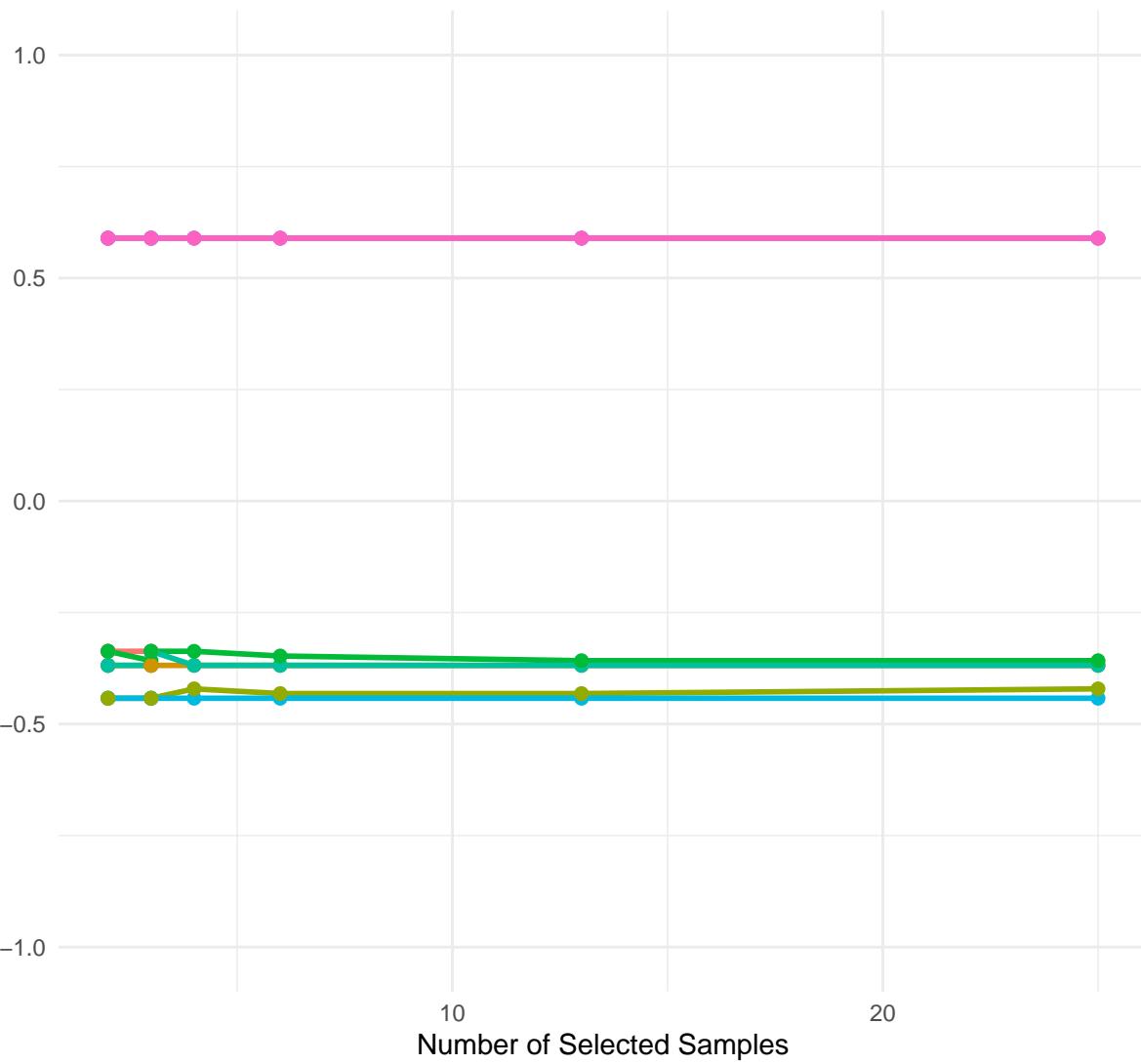
range

variance



Score: Plmd (Local, partial1 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

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

Score: Plmd (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

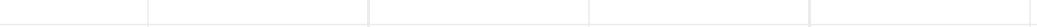
-0.5

-1.0

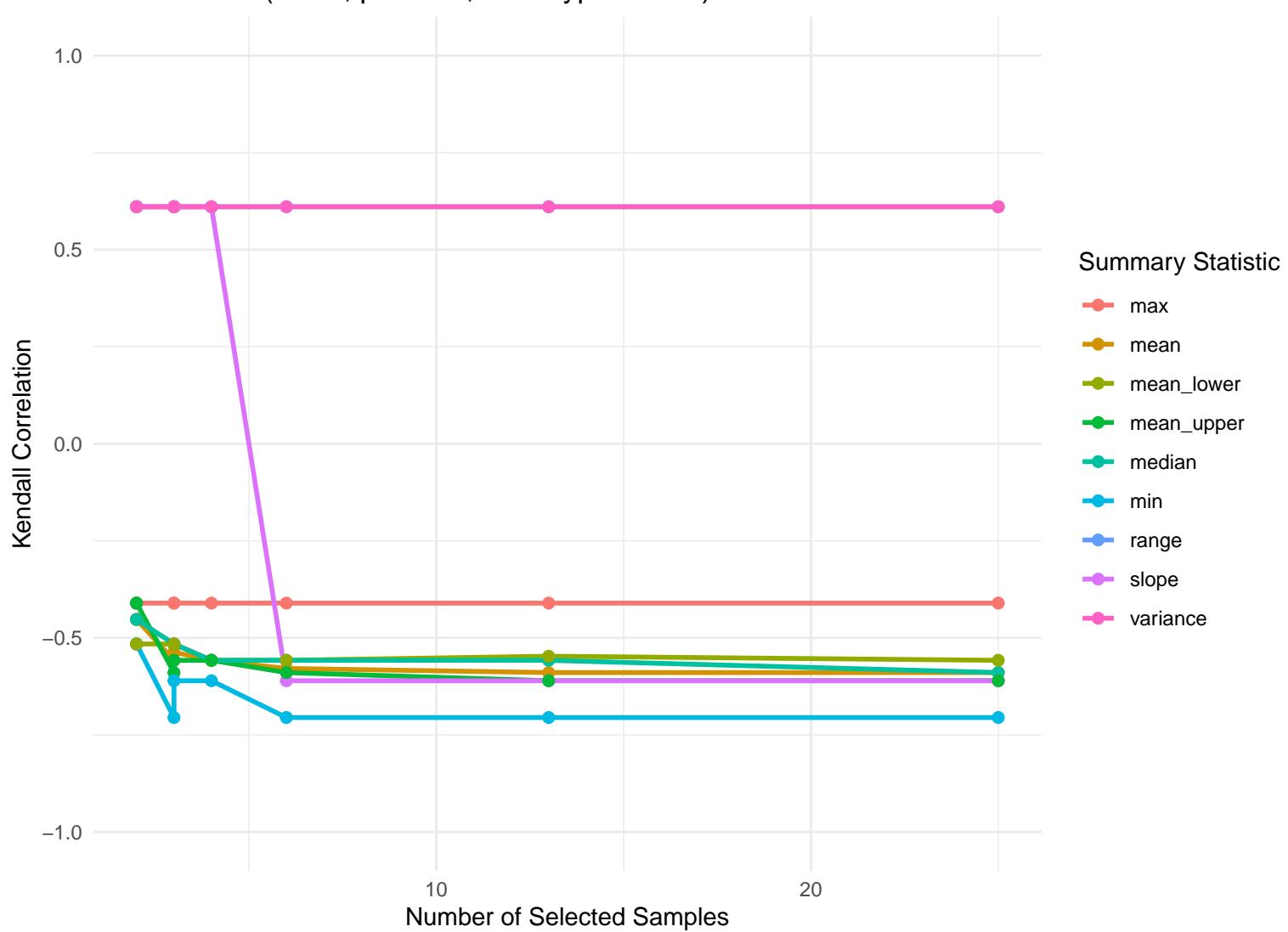
Number of Selected Samples

Summary Statistic

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



Score: Plmd (Local, partial1 , Genotype: wave)



Score: PILSM (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

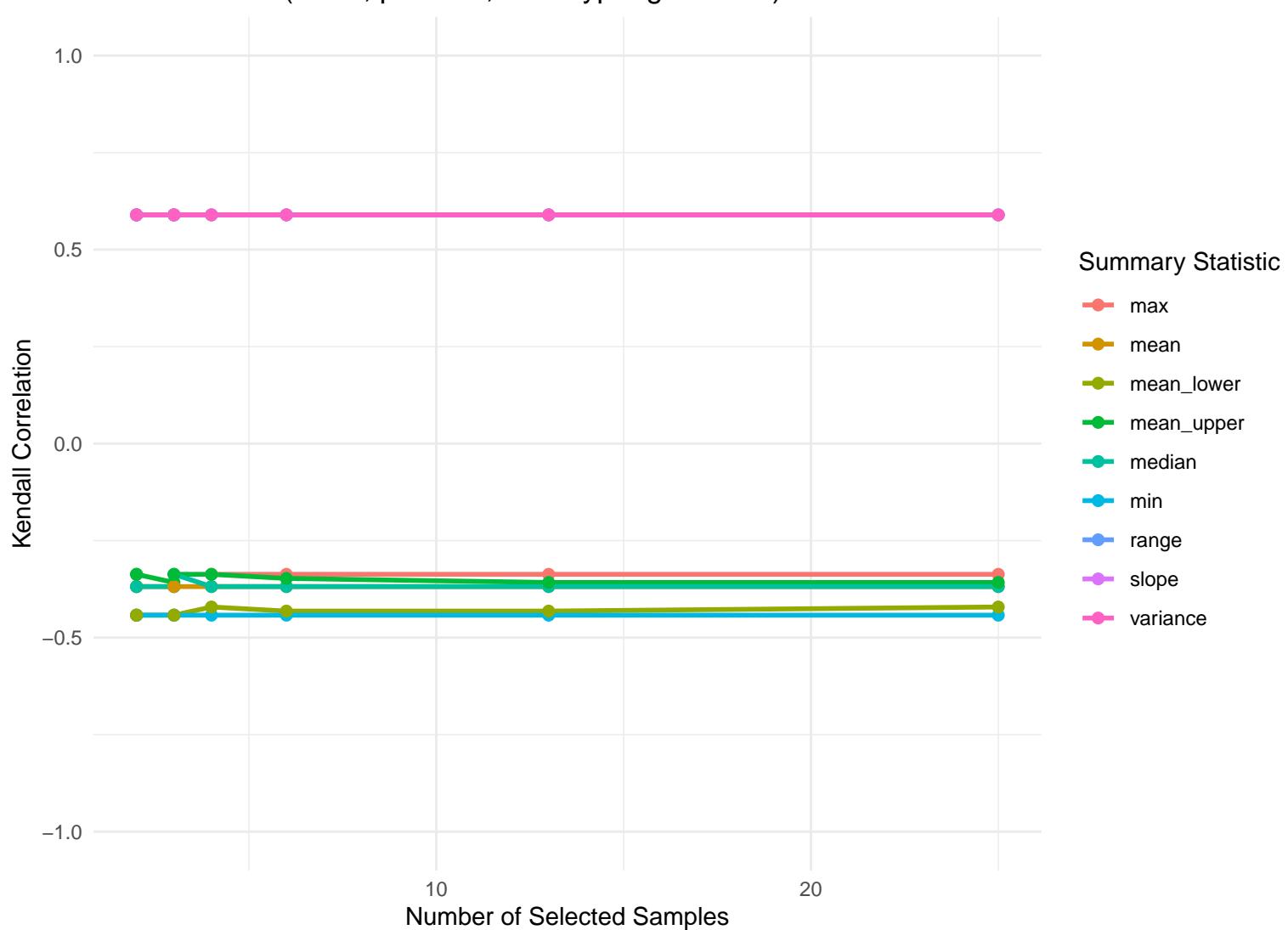
Summary Statistic

range

variance

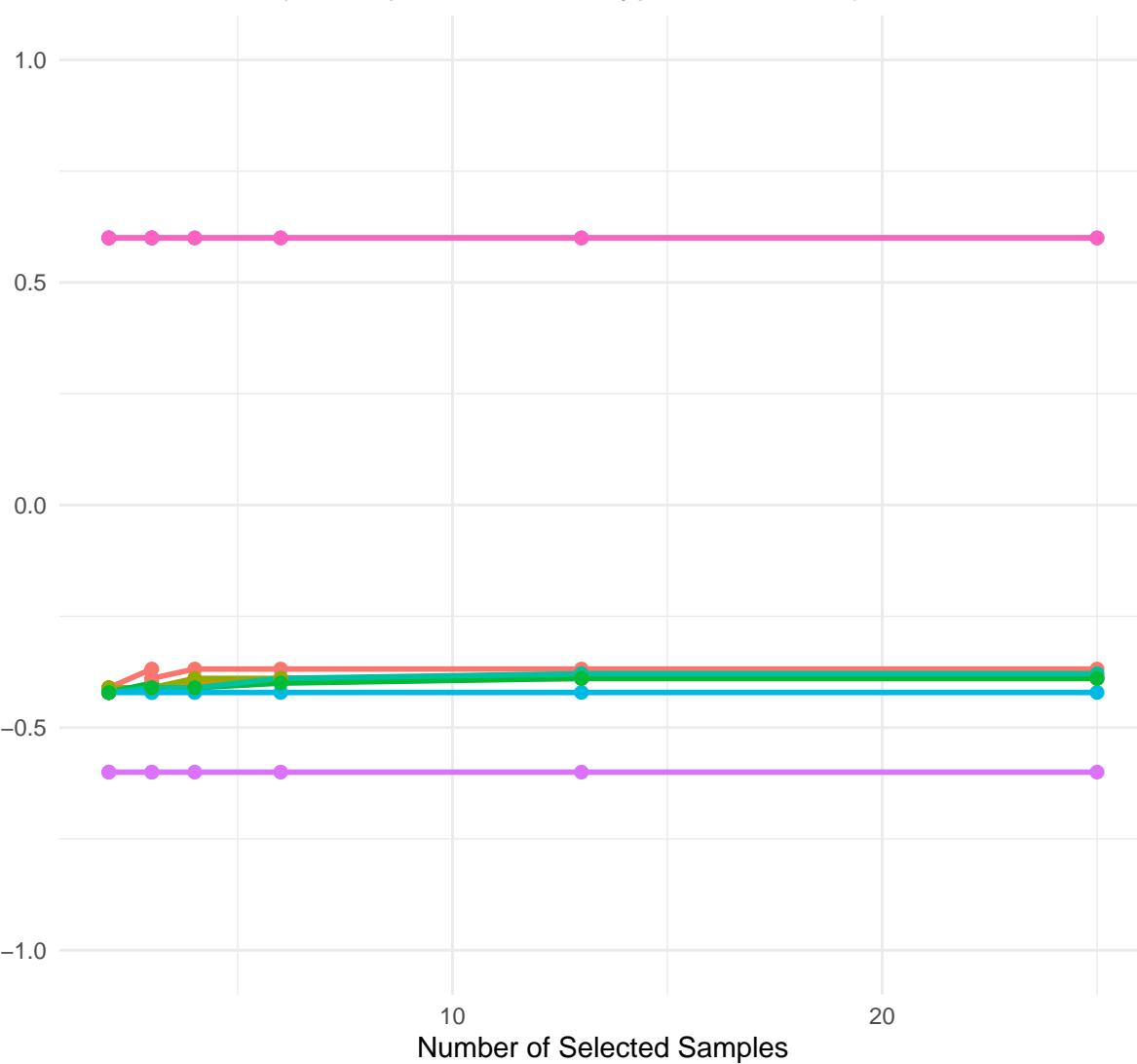


Score: PILSM (Local, partial1 , Genotype: gaussian)



Score: PILSM (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

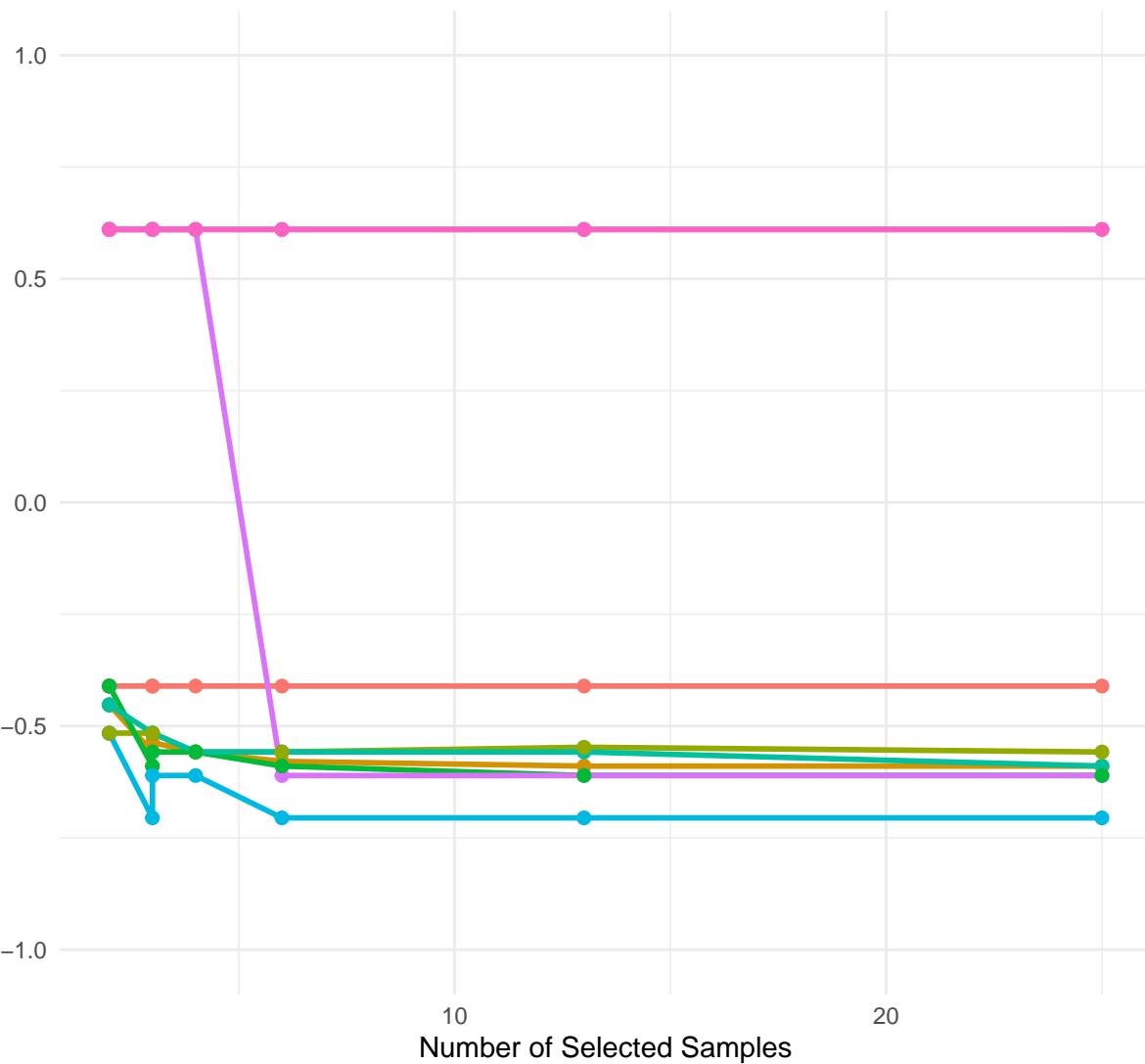


Summary Statistic

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

Score: PILSM (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: RTR (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

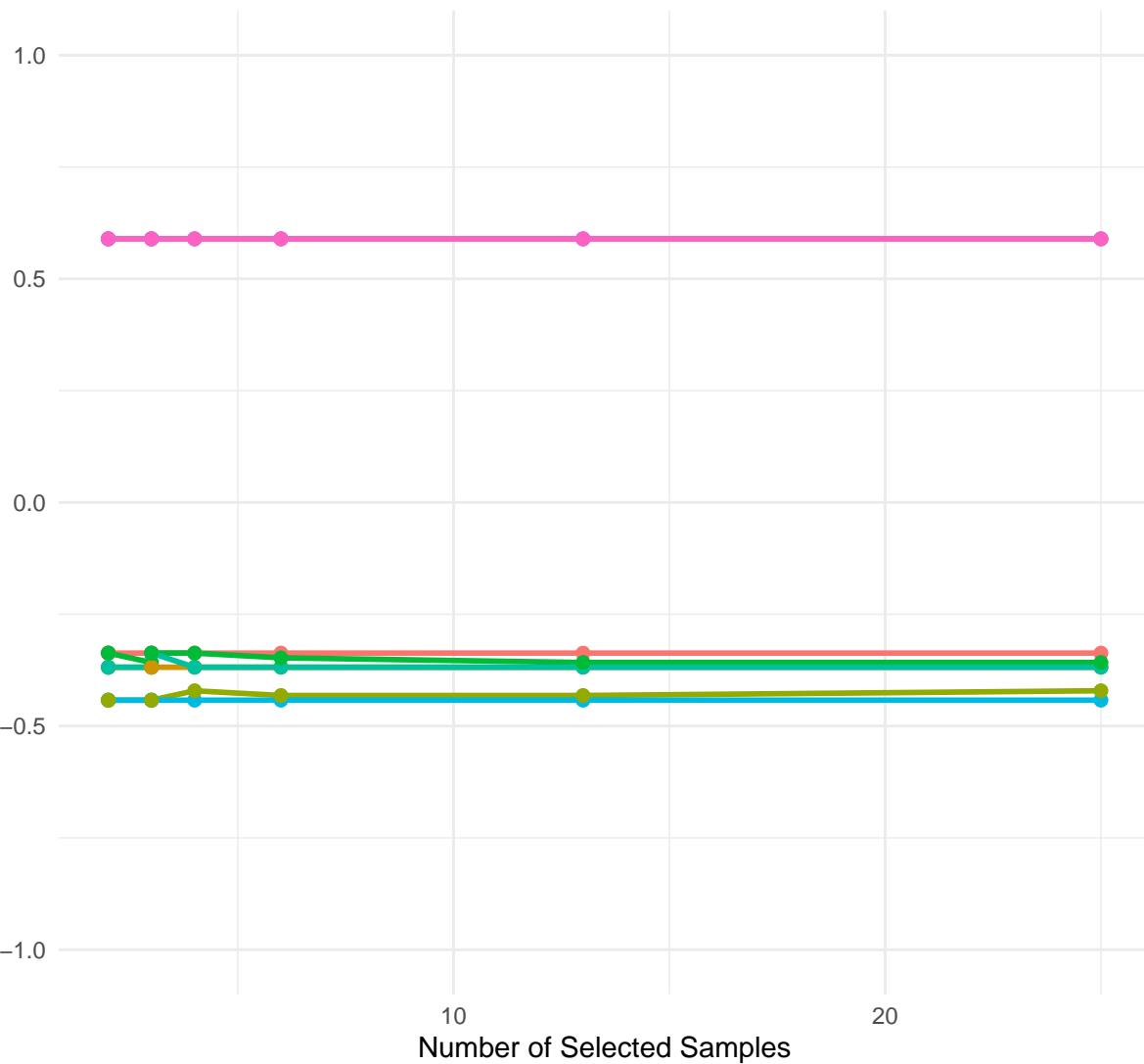
Summary Statistic

slope

Number of Selected Samples

Score: RTR (Local, partial1 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

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

Score: RTR (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

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

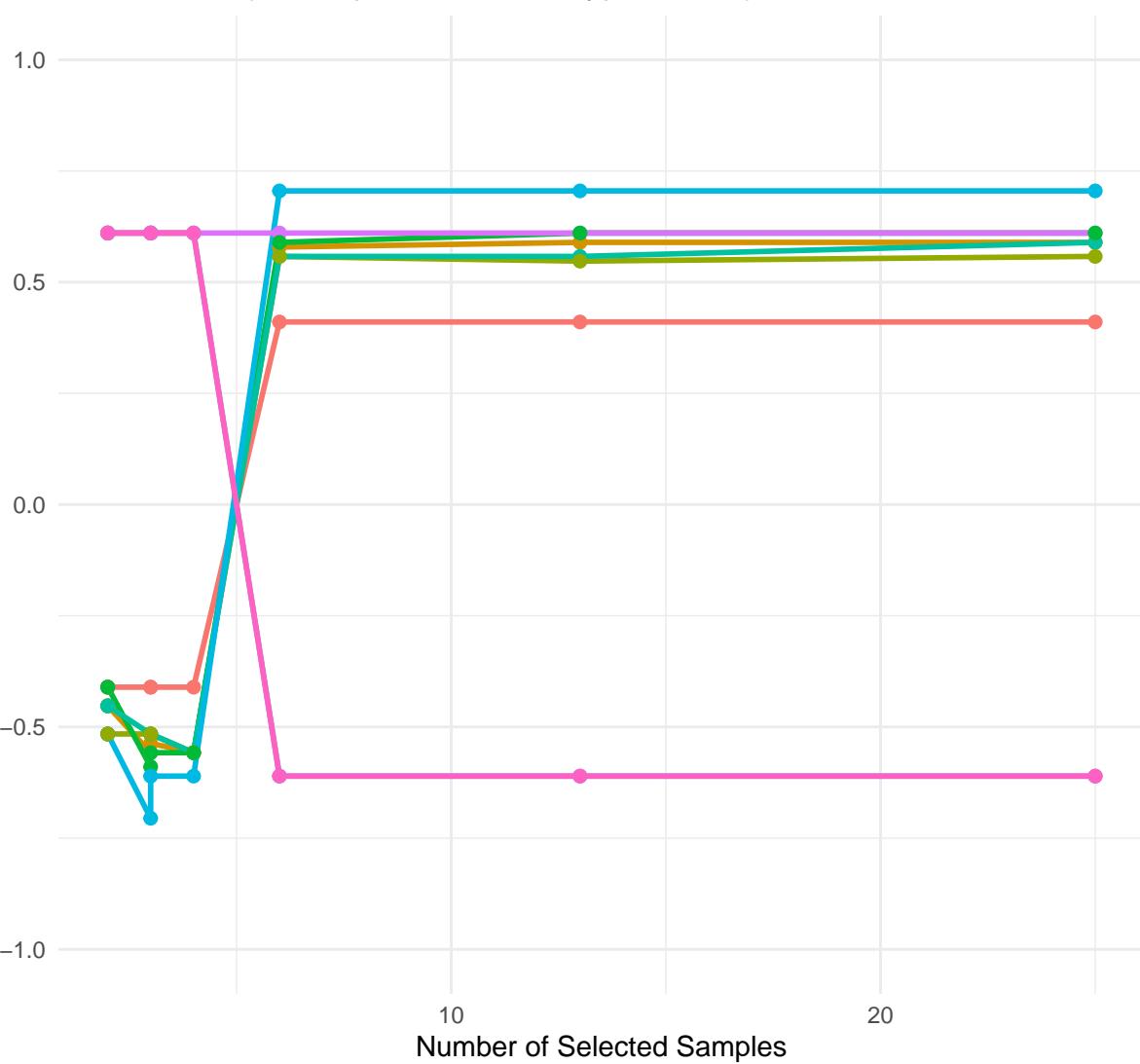
0.62 0.62 0.62 0.62 0.62 0.62

0.42 0.42 0.42 0.42 0.42 0.42 0.42 0.42 0.42

-0.62 -0.62 -0.62 -0.62 -0.62 -0.62

Score: RTR (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: PIR (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

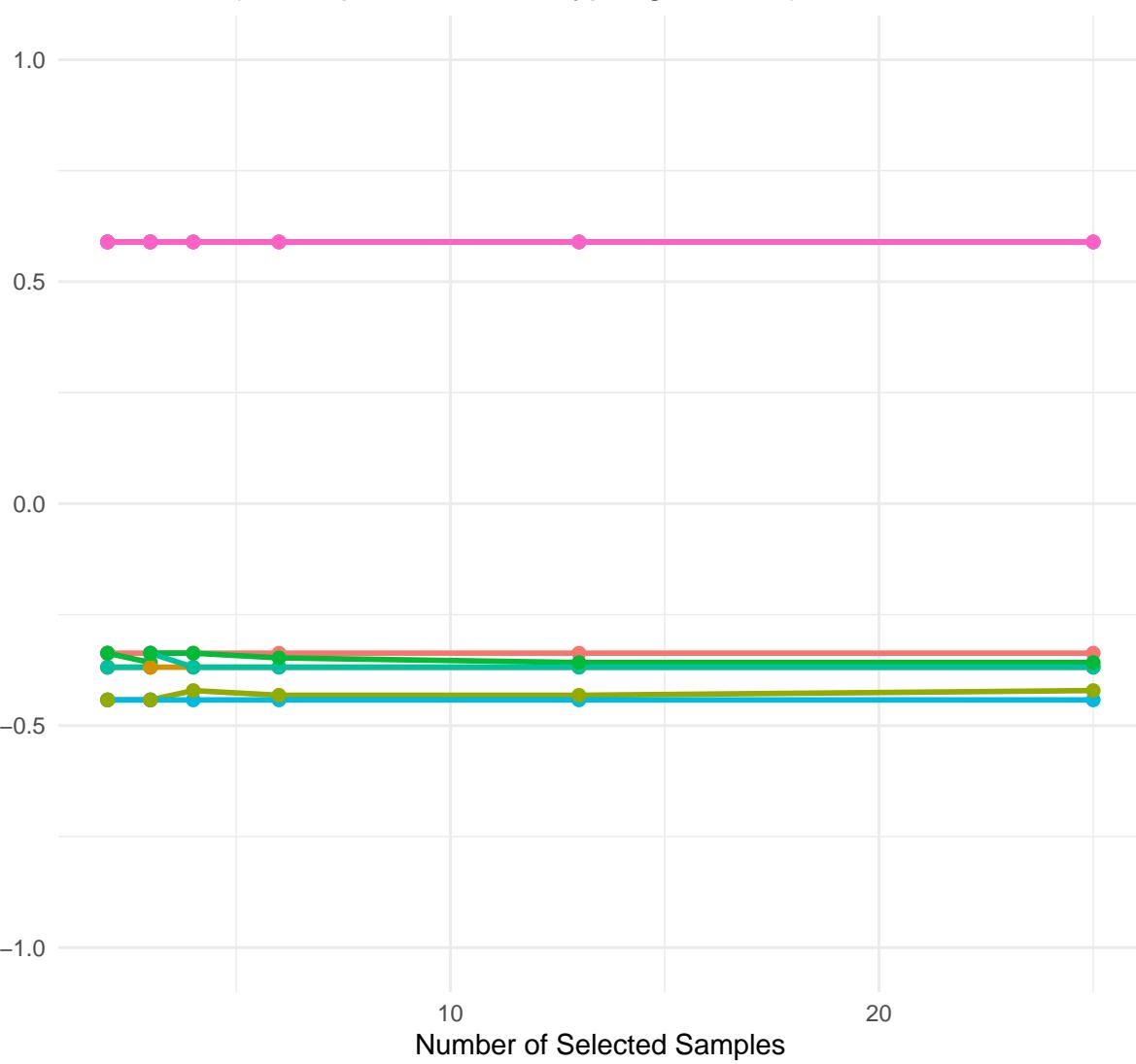
range

variance



Score: PIR (Local, partial1 , Genotype: gaussian)

Kendall Correlation

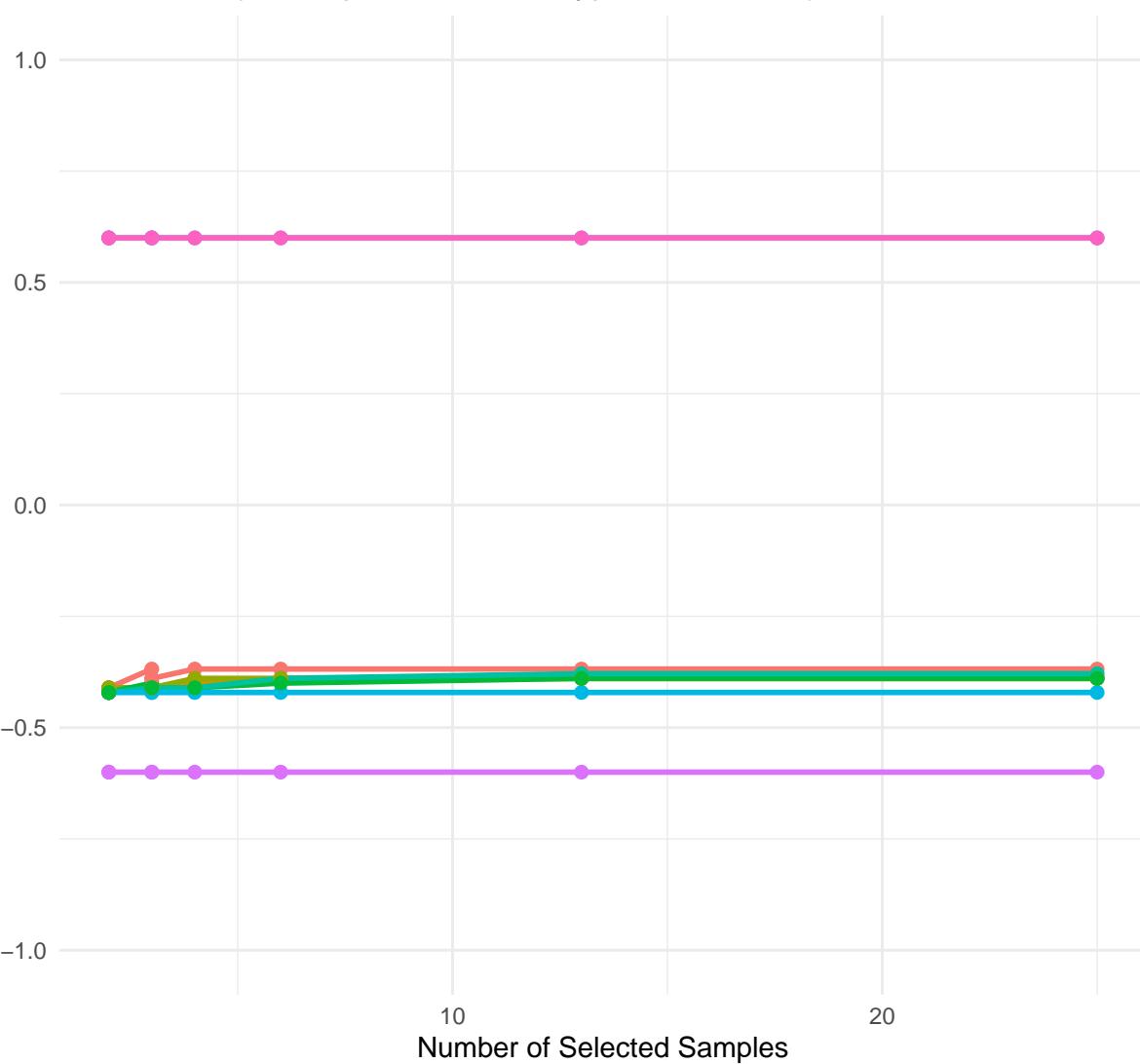


Summary Statistic

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

Score: PIR (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

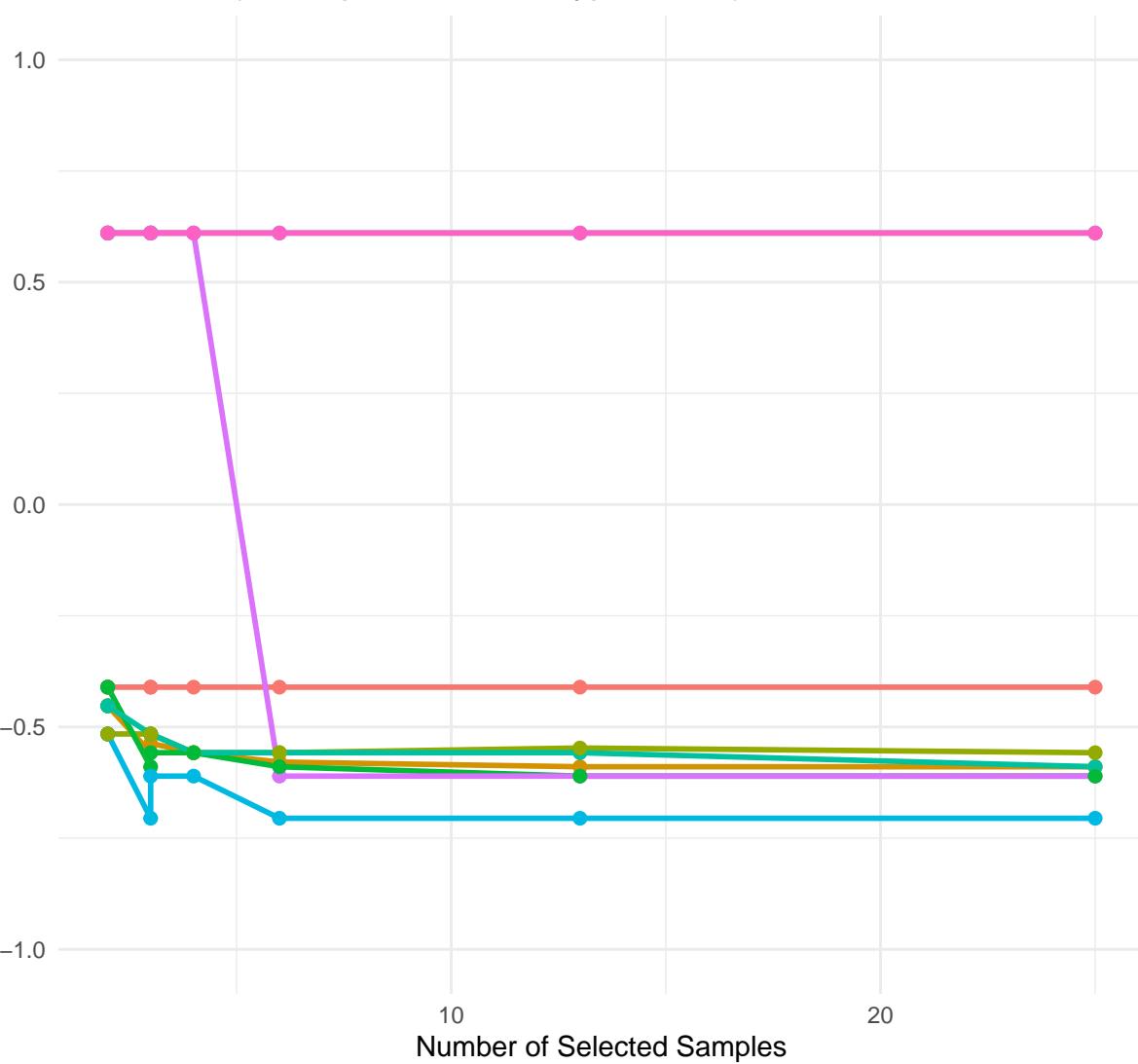


Summary Statistic

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

Score: PIR (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: RDPI (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

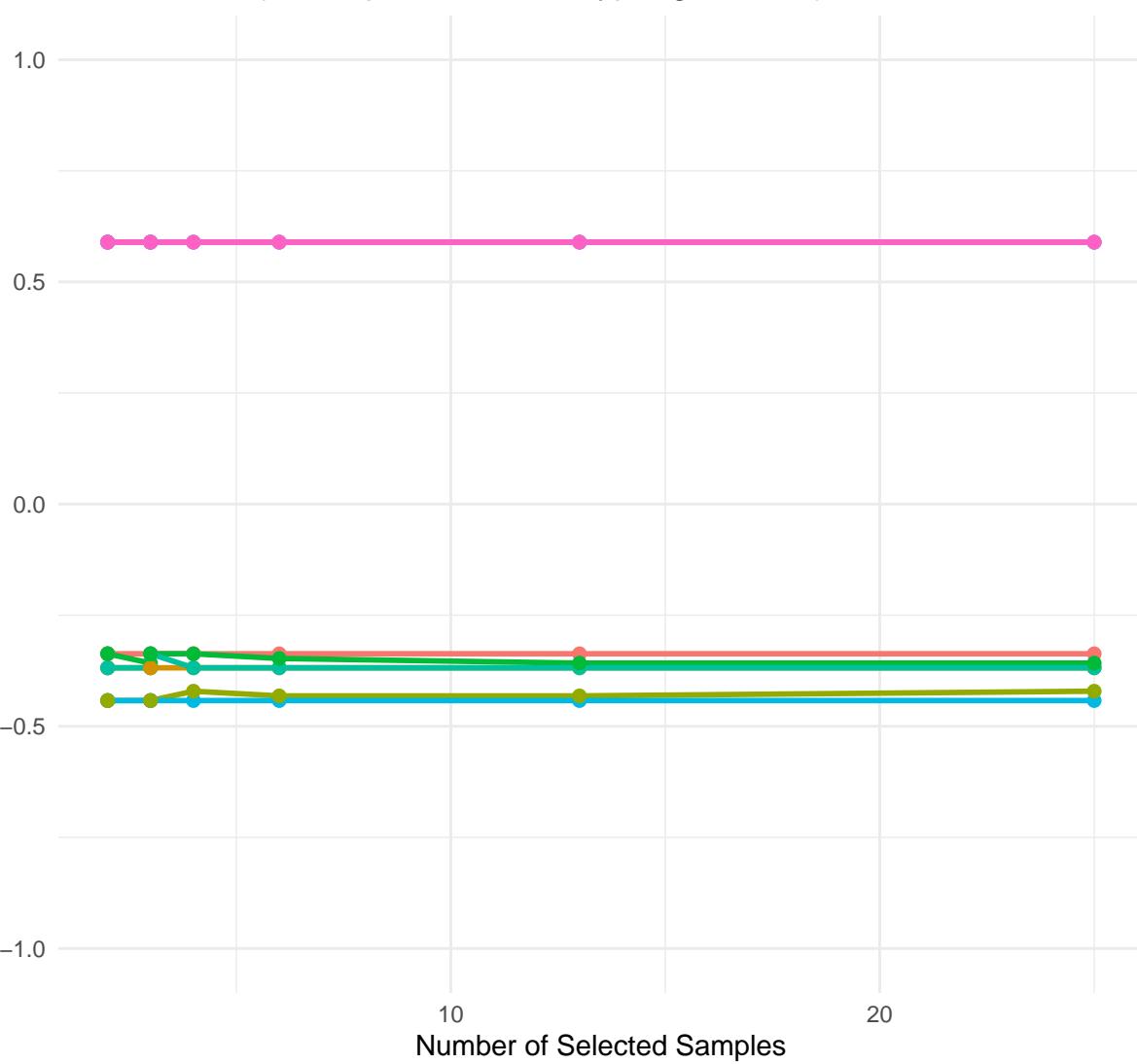
range

variance



Score: RDPI (Local, partial1 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

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

Score: RDPI (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

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

0.58

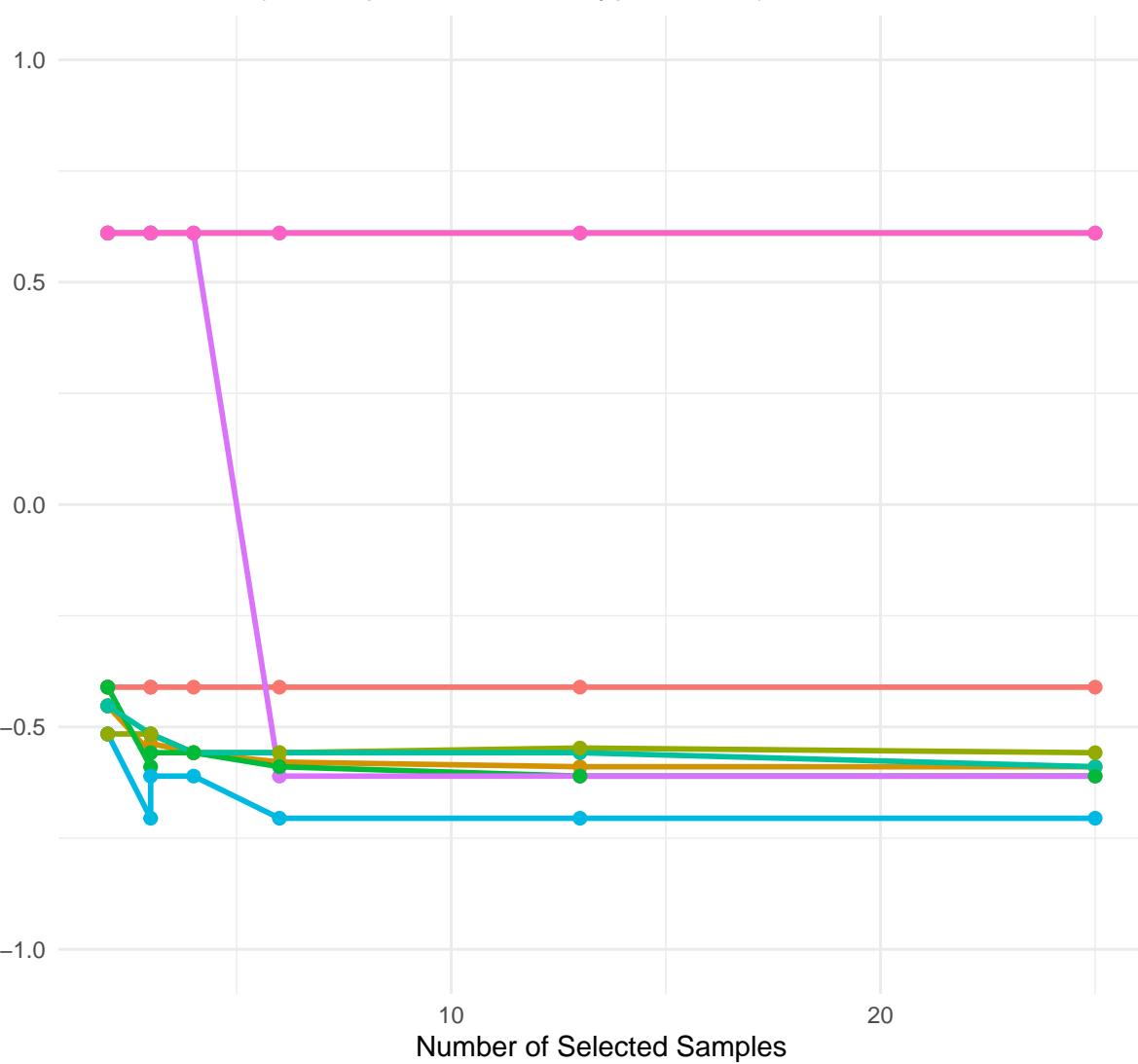
-0.48

-0.62

0.58

Score: RDPI (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: ESPI (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

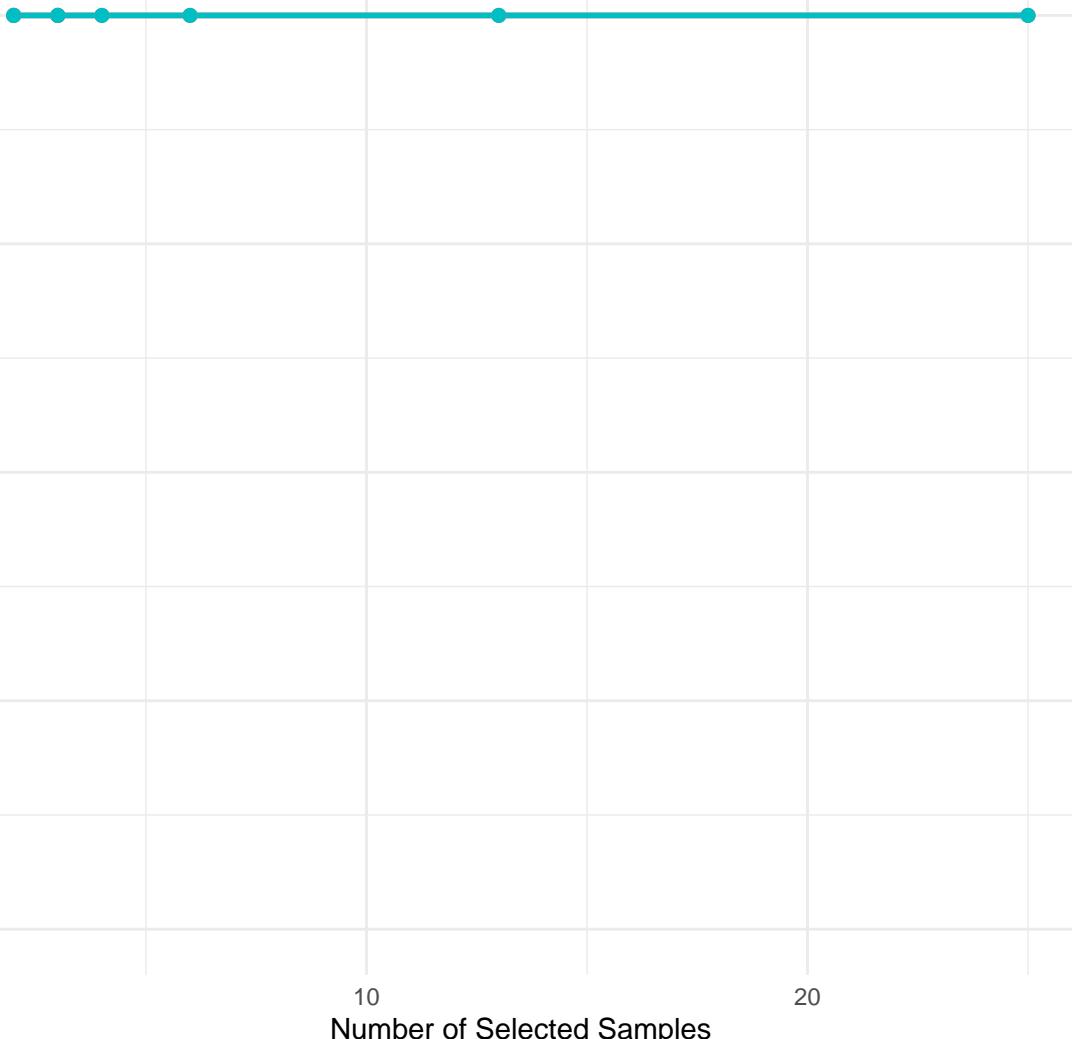
-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: ESPI (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

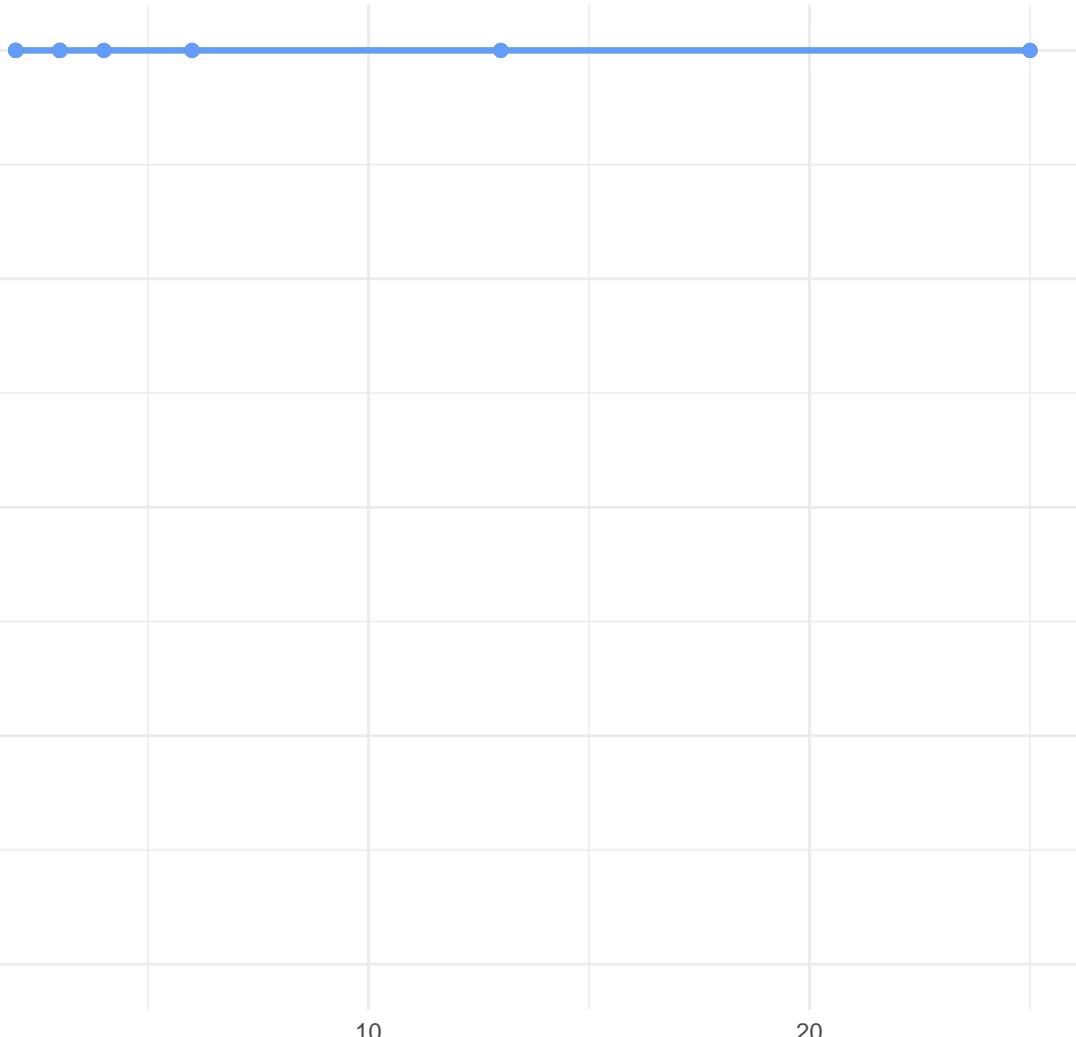
Number of Selected Samples

Summary Statistic

range

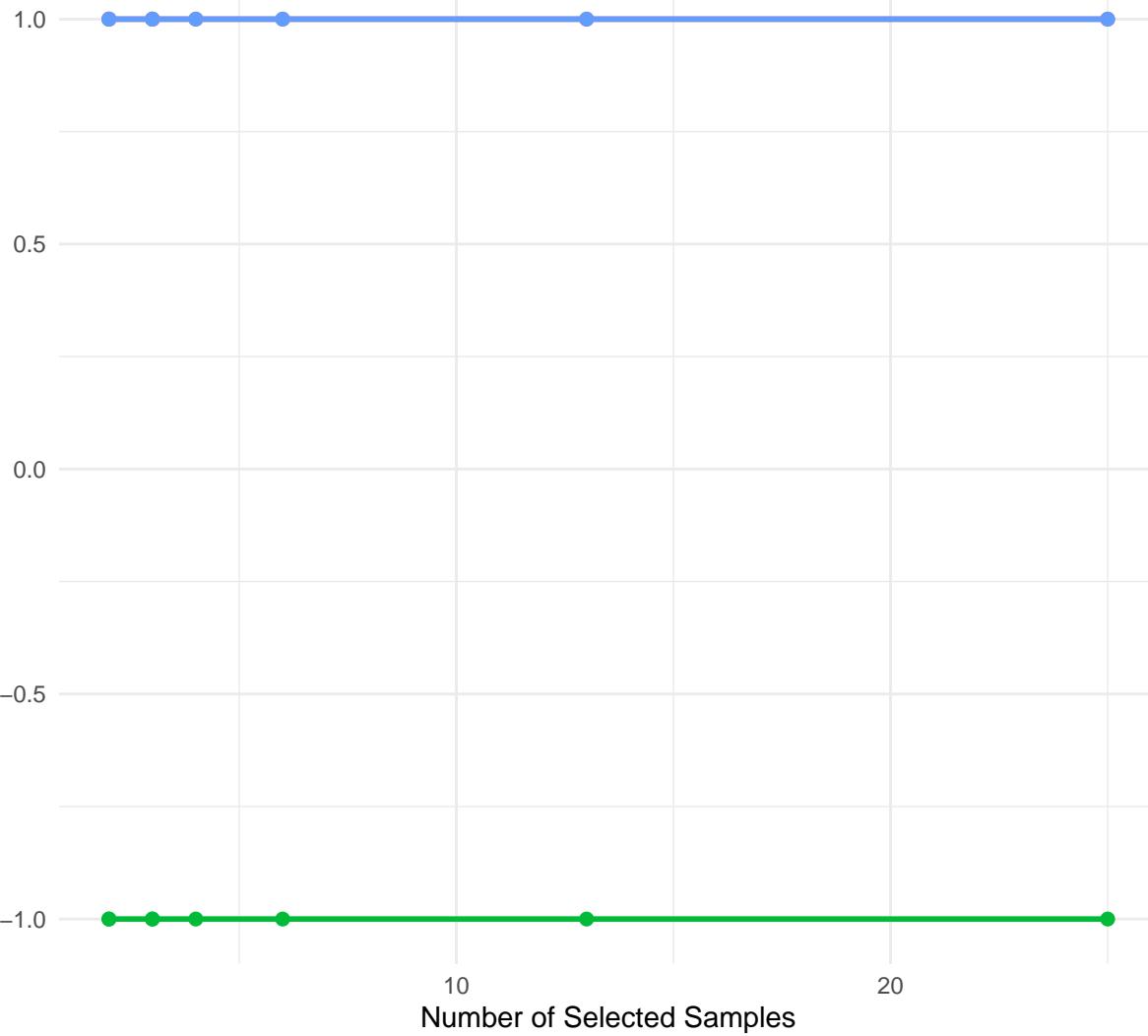
slope

variance



Score: ESPI (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

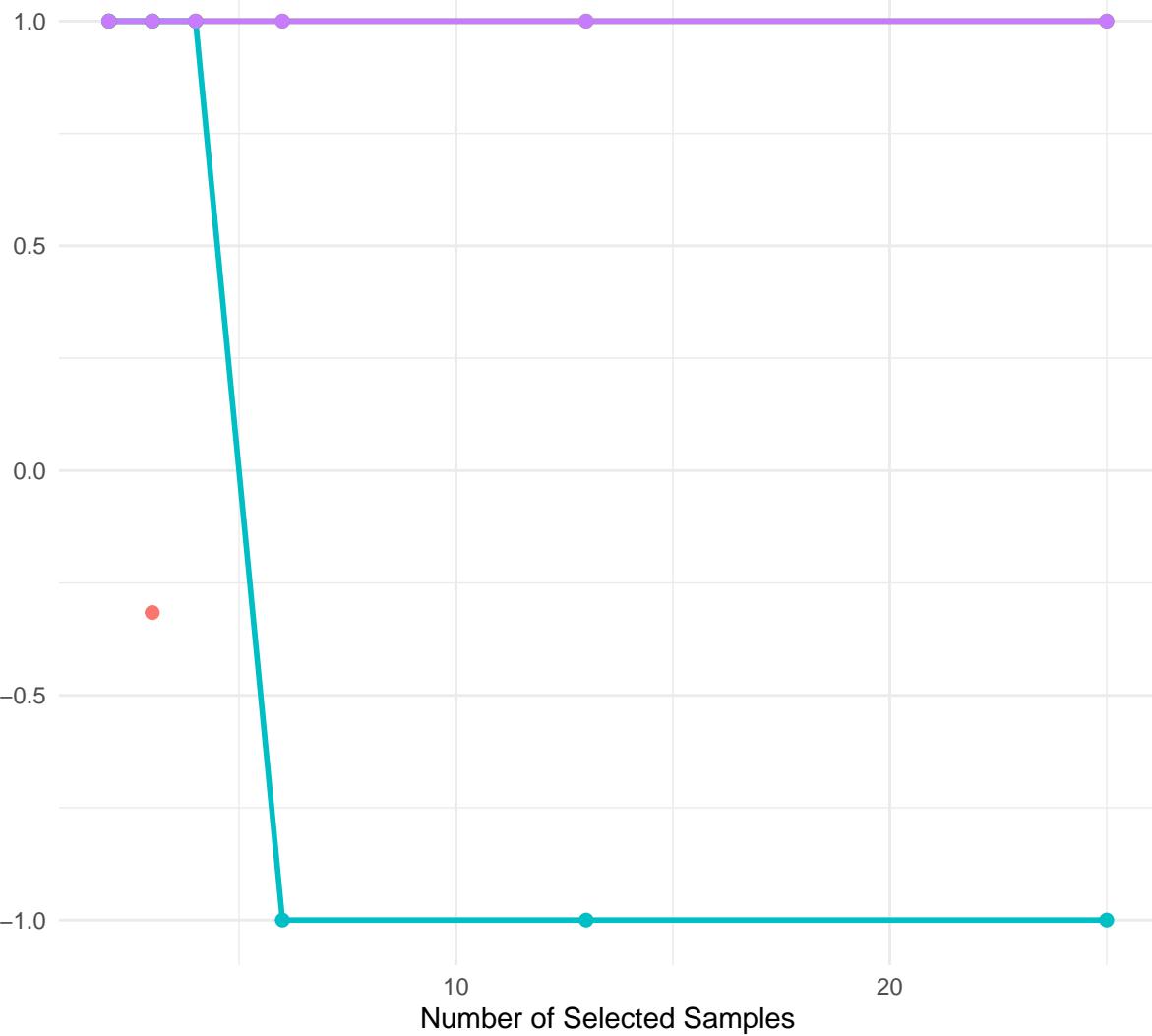
range

slope

variance

Score: ESPI (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

- min
- range
- slope
- variance

Score: ESPIID (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

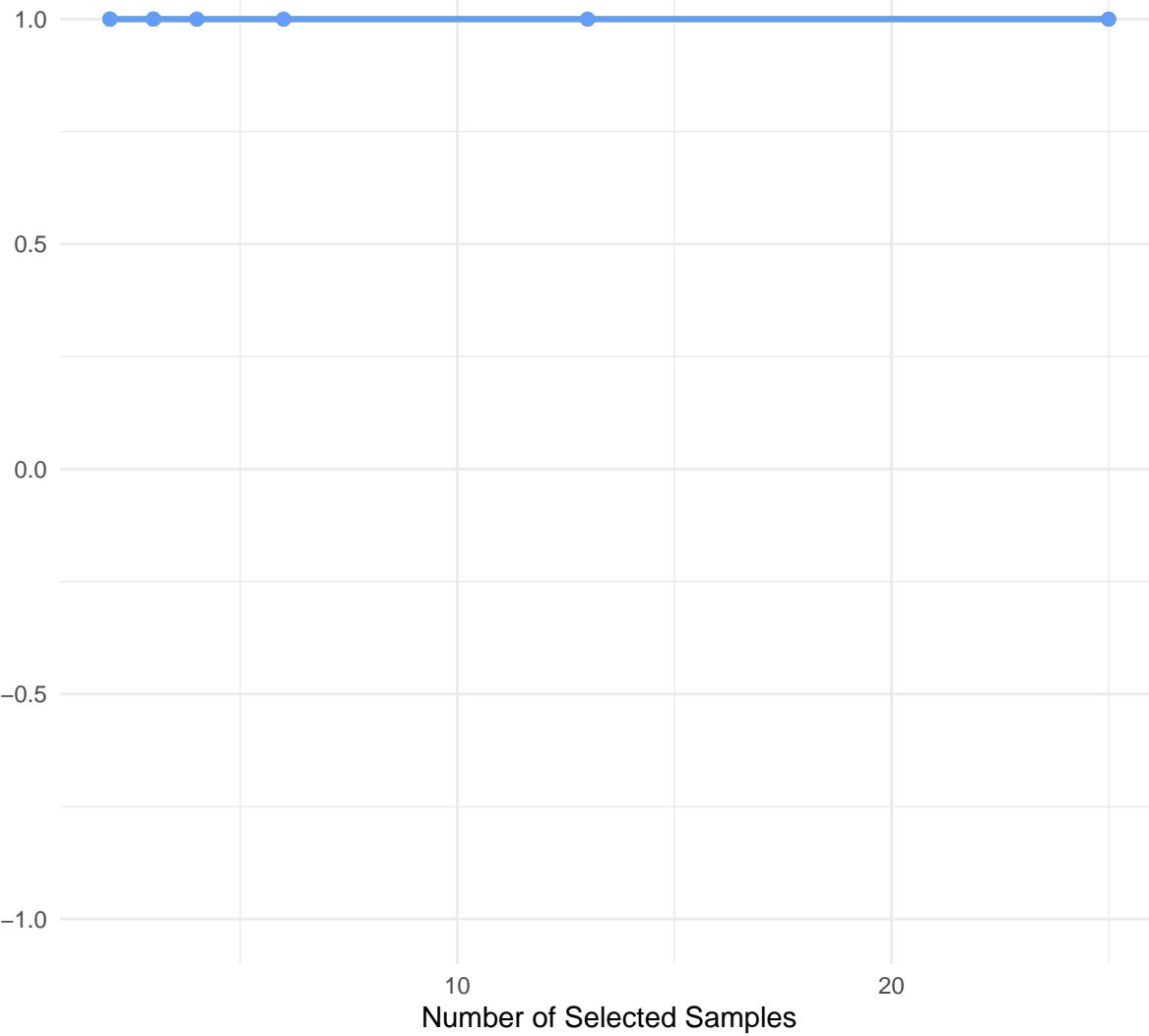
range

variance



Score: ESPIID (Local, partial1 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

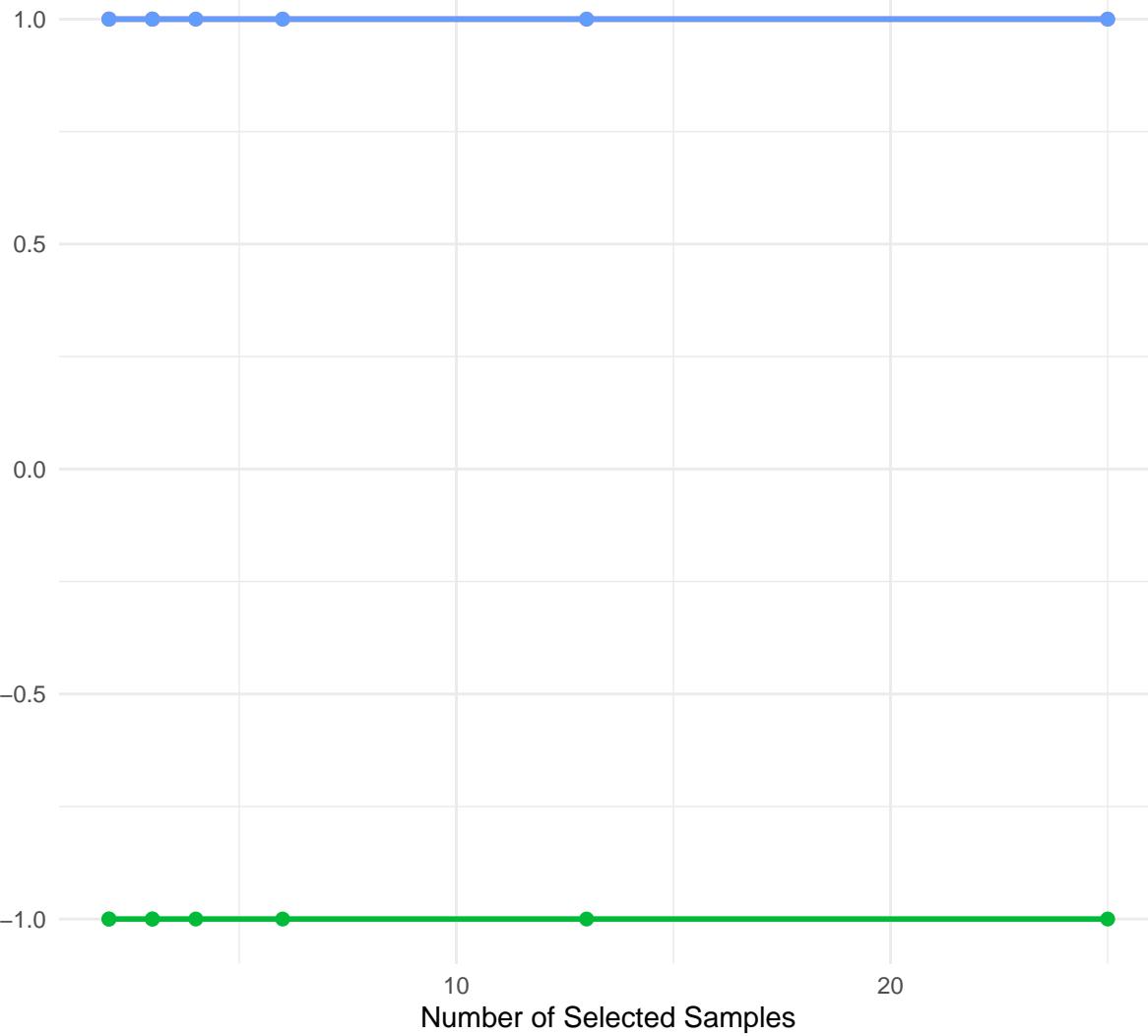
range

slope

variance

Score: ESPIID (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

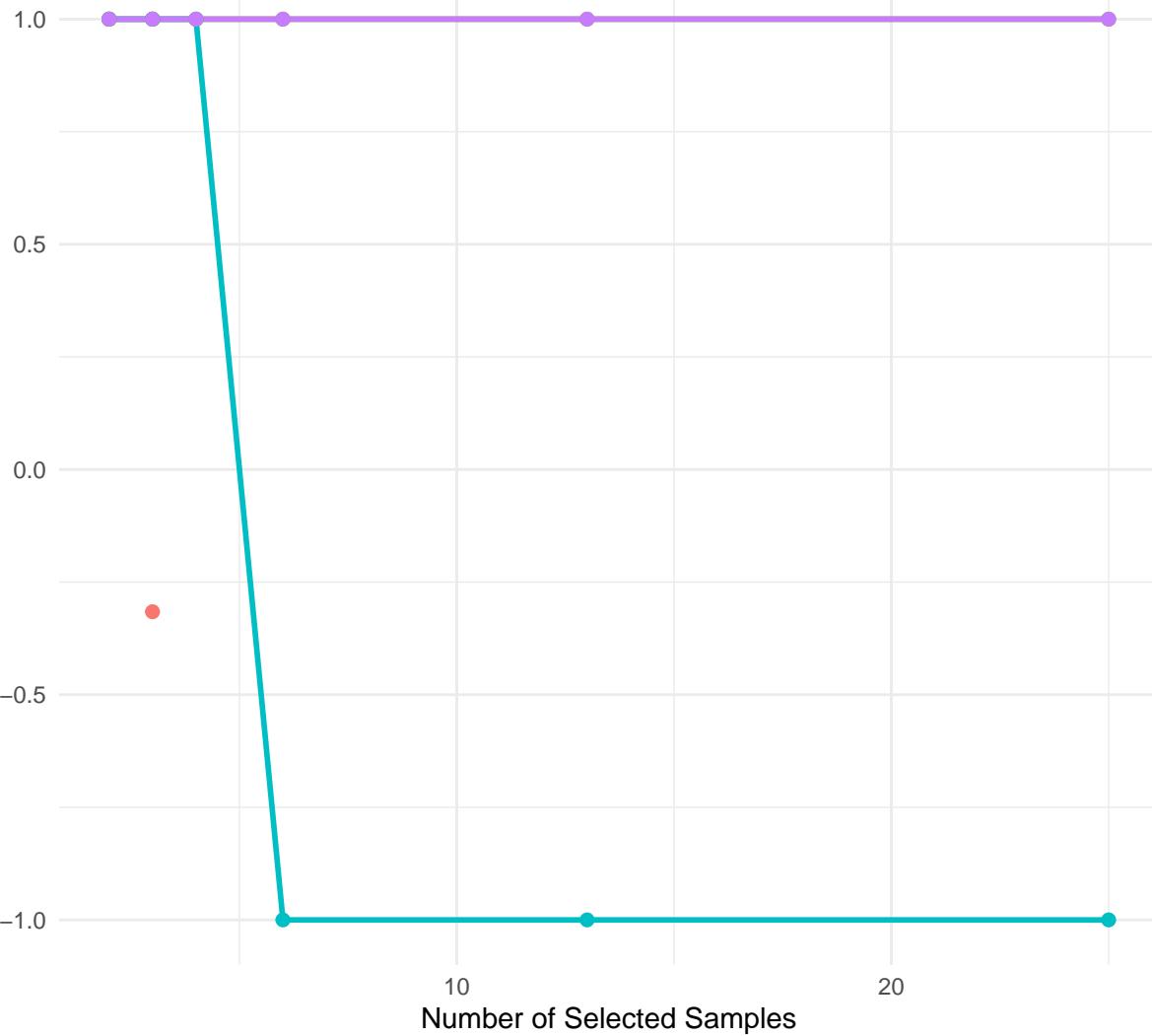
range

slope

variance

Score: ESPIID (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

- min
- range
- slope
- variance

Score: PSI (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Summary Statistic

range

variance

Number of Selected Samples

10

20

Score: PSI (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Summary Statistic

range

slope

variance

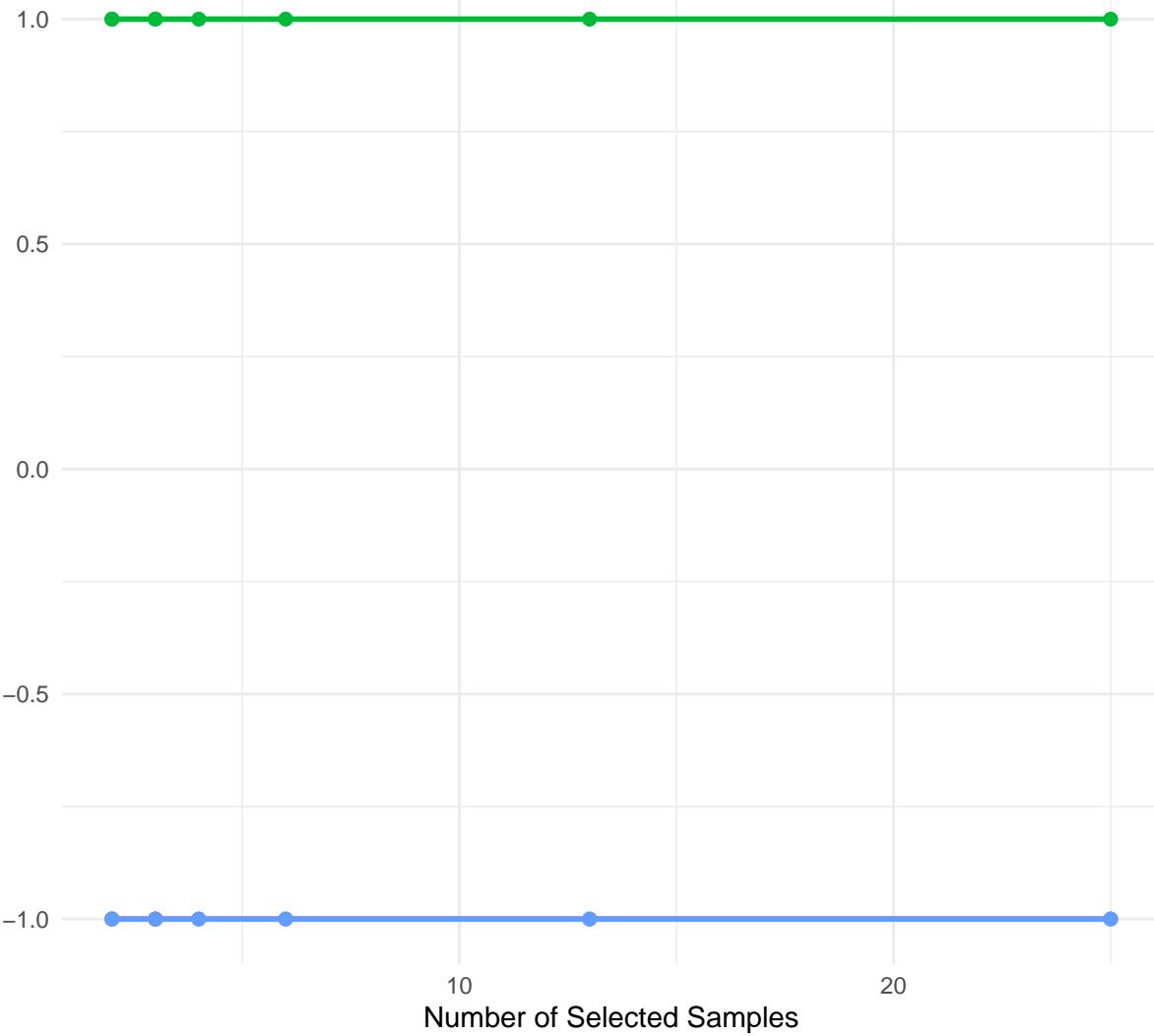
Number of Selected Samples

10 20



Score: PSI (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

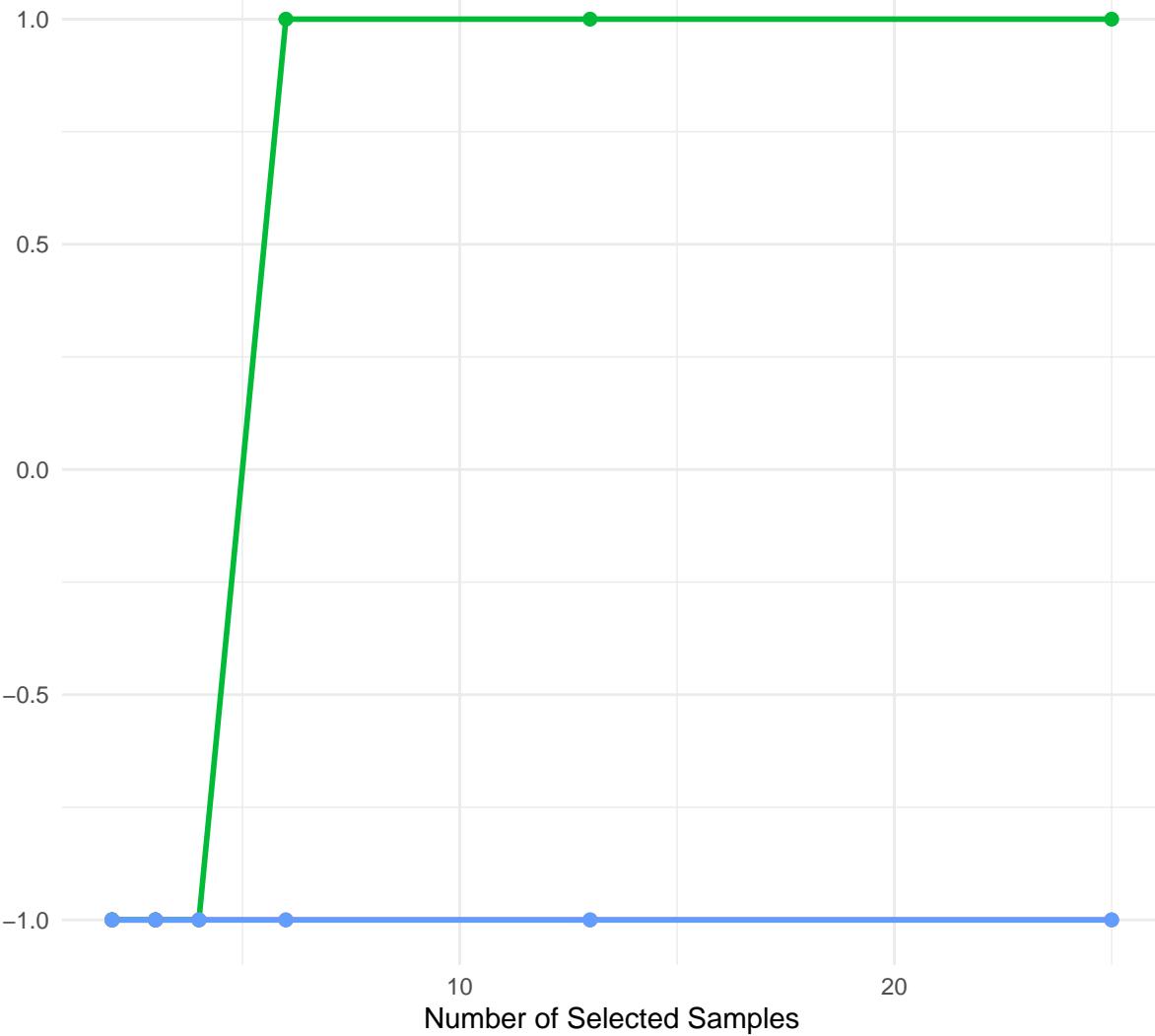
range

slope

variance

Score: PSI (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

- range
- slope
- variance

Score: RPI (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

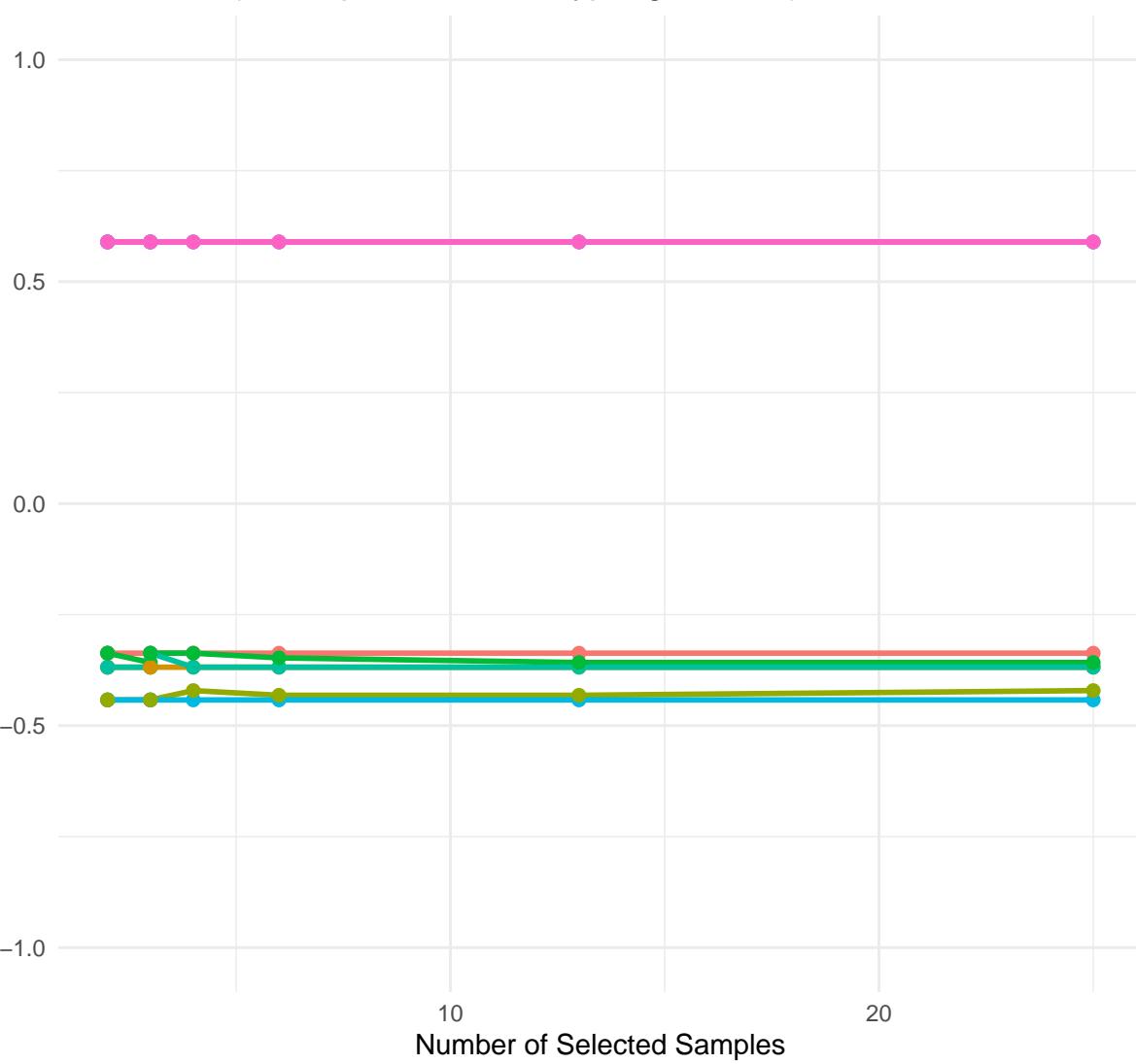
range

variance



Score: RPI (Local, partial1 , Genotype: gaussian)

Kendall Correlation

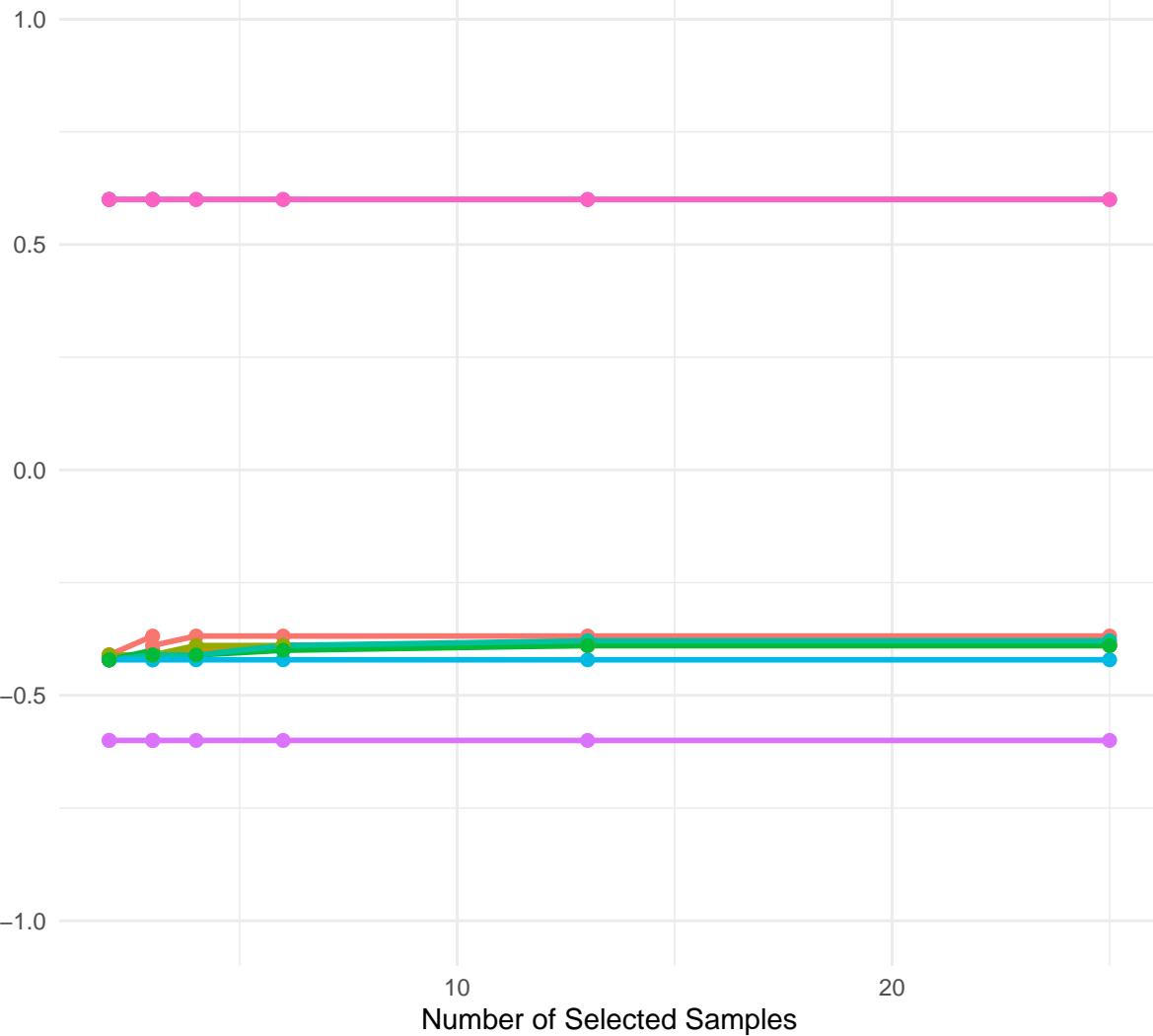


Summary Statistic

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

Score: RPI (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

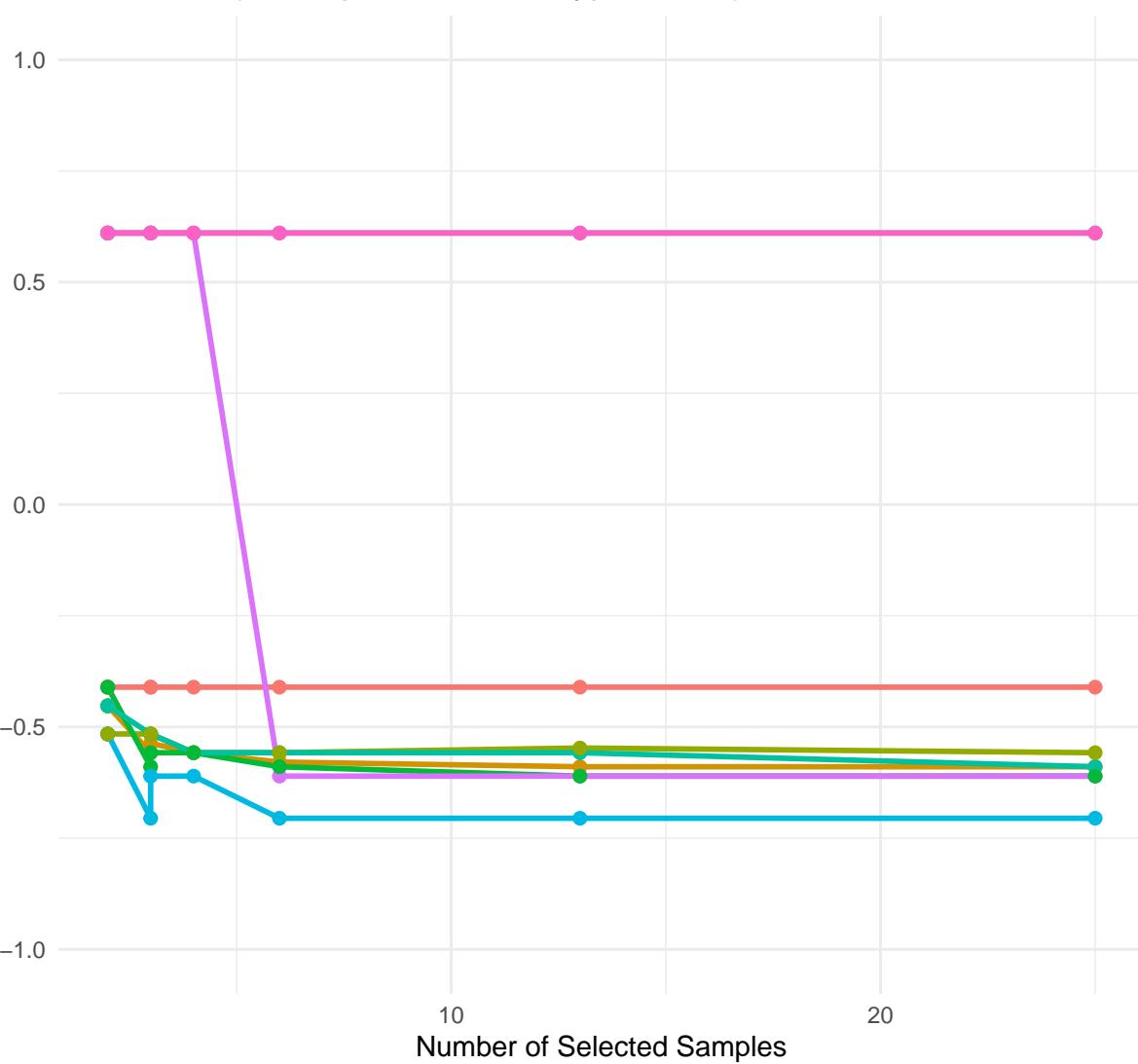


Summary Statistic

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

Score: RPI (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: PQ (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

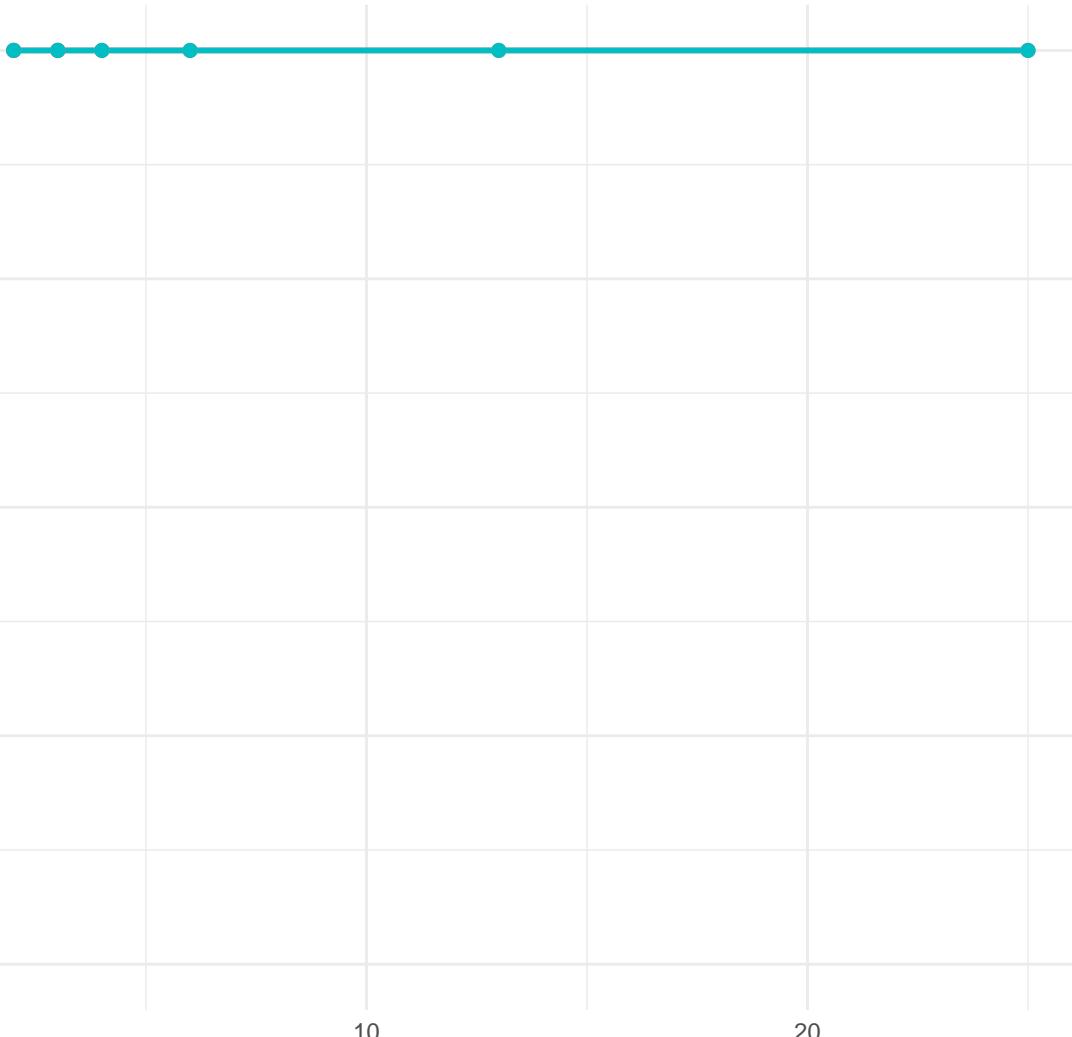
-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: PQ (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

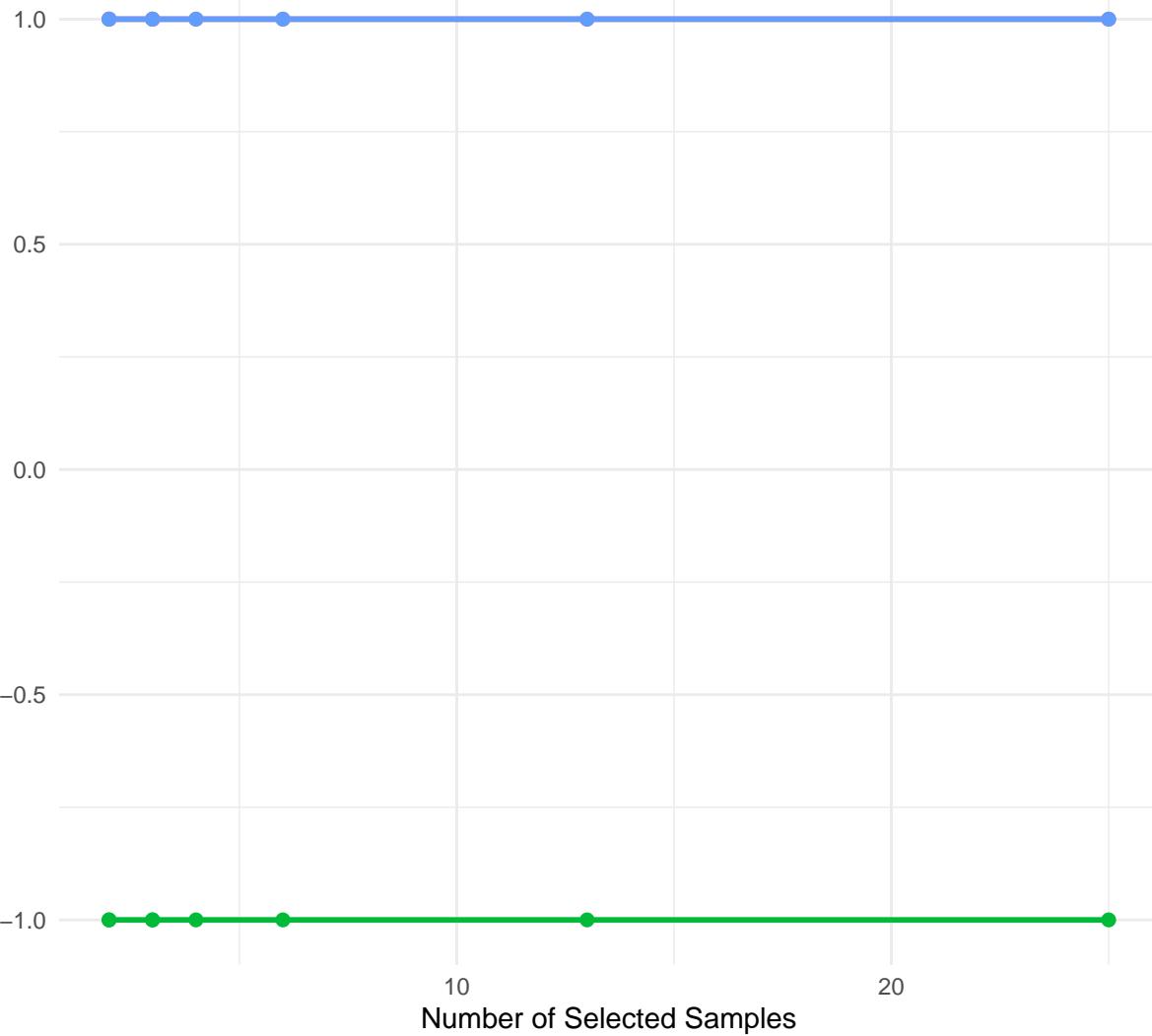
slope

variance



Score: PQ (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

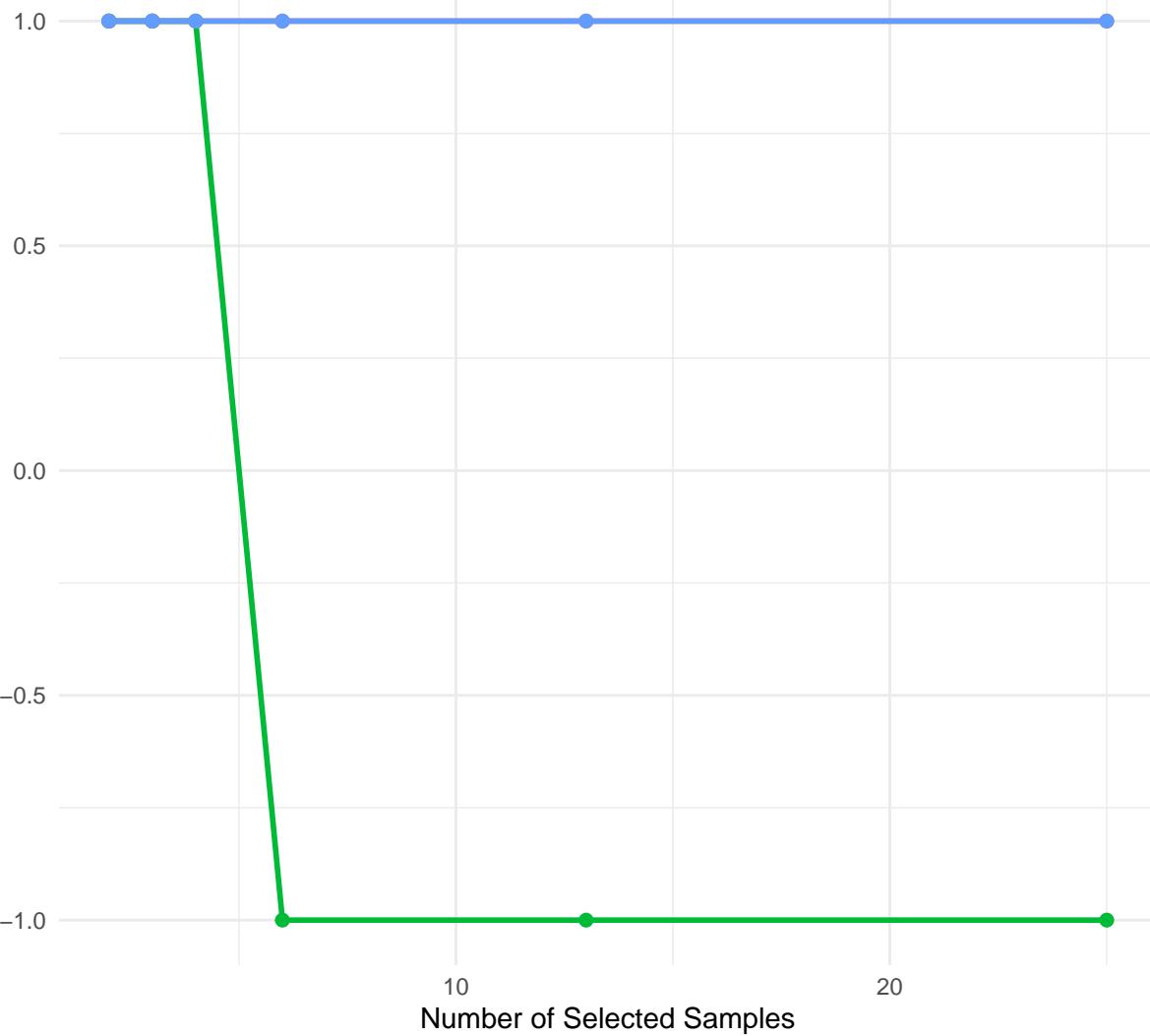
range

slope

variance

Score: PQ (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

- range
- slope
- variance

Score: PR (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

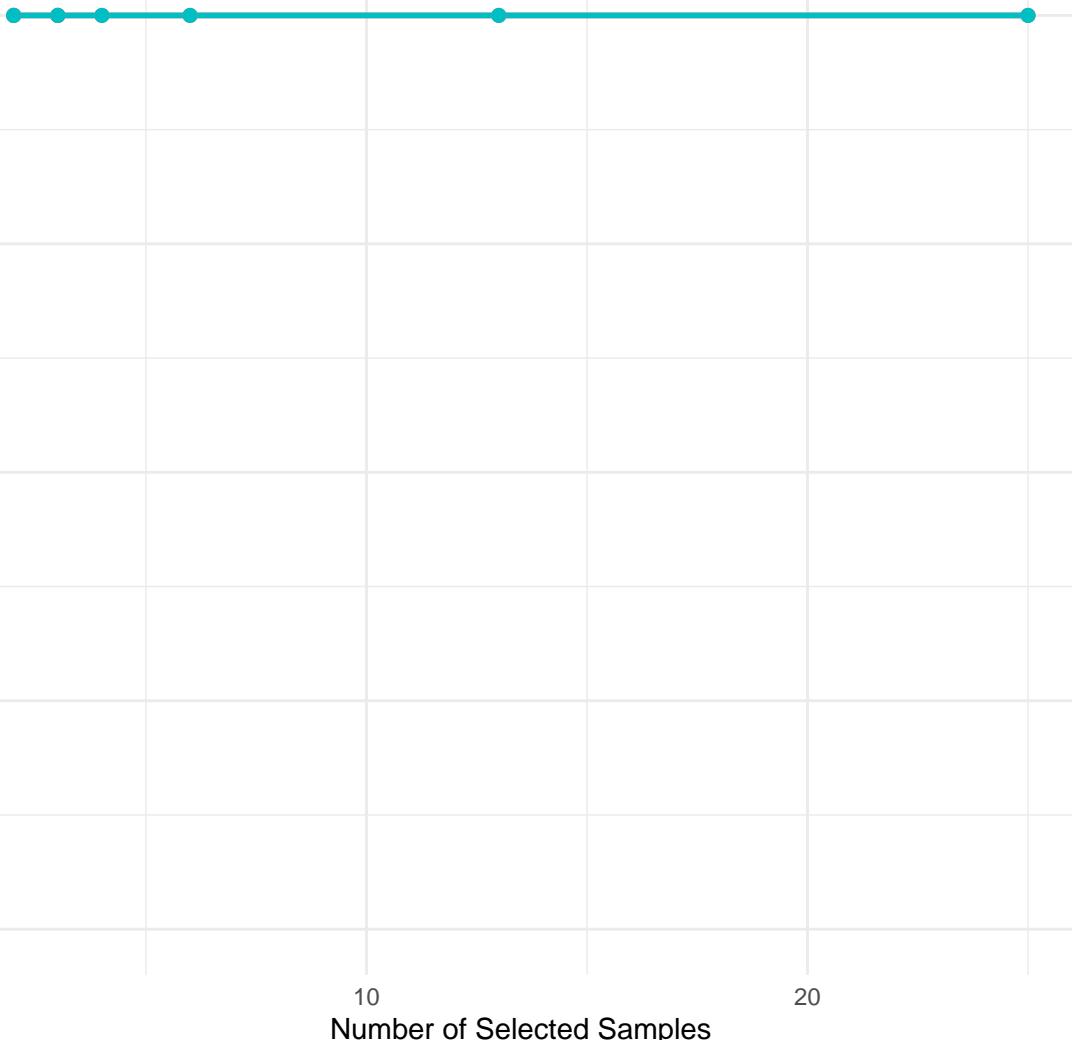
-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: PR (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

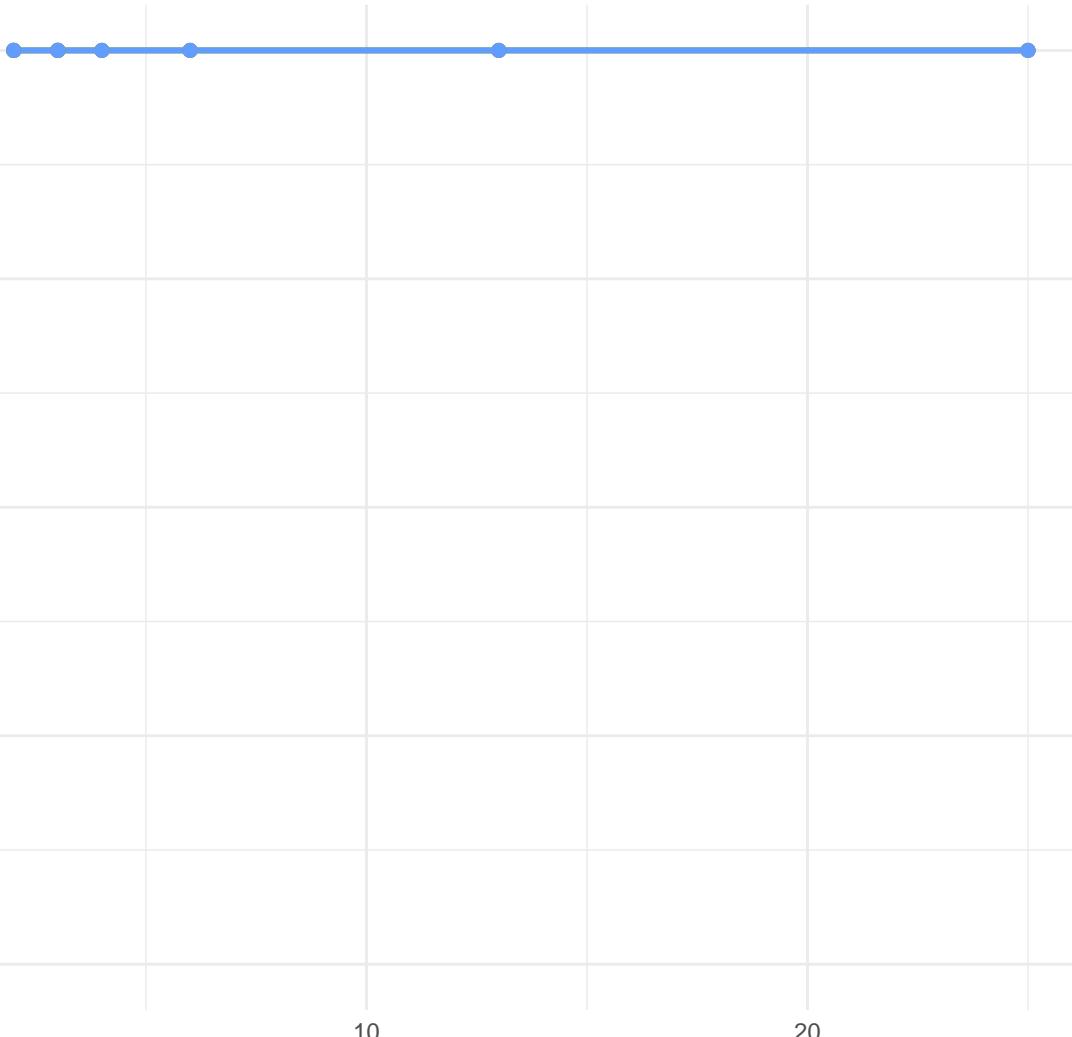
Number of Selected Samples

Summary Statistic

range

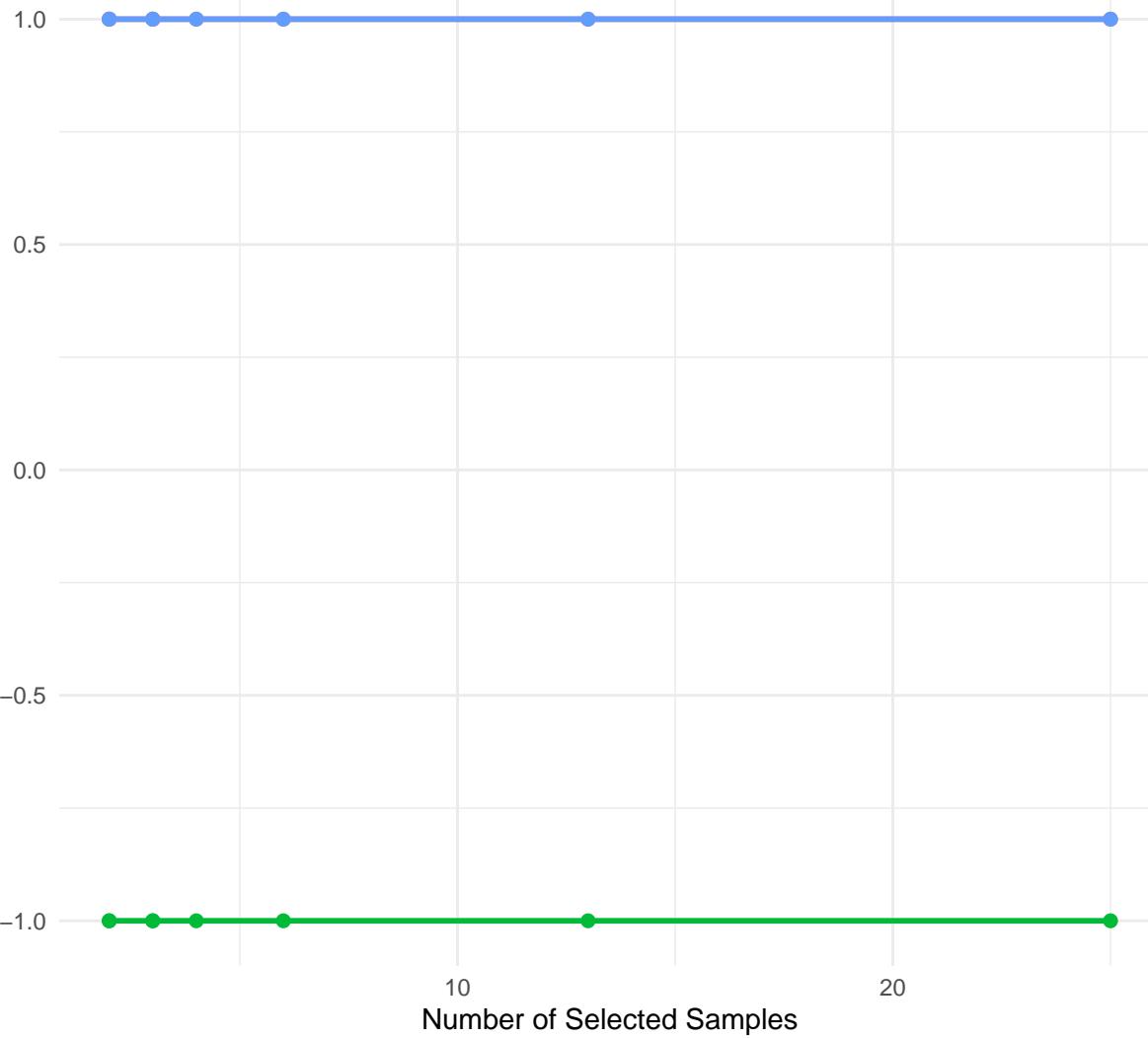
slope

variance



Score: PR (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

range

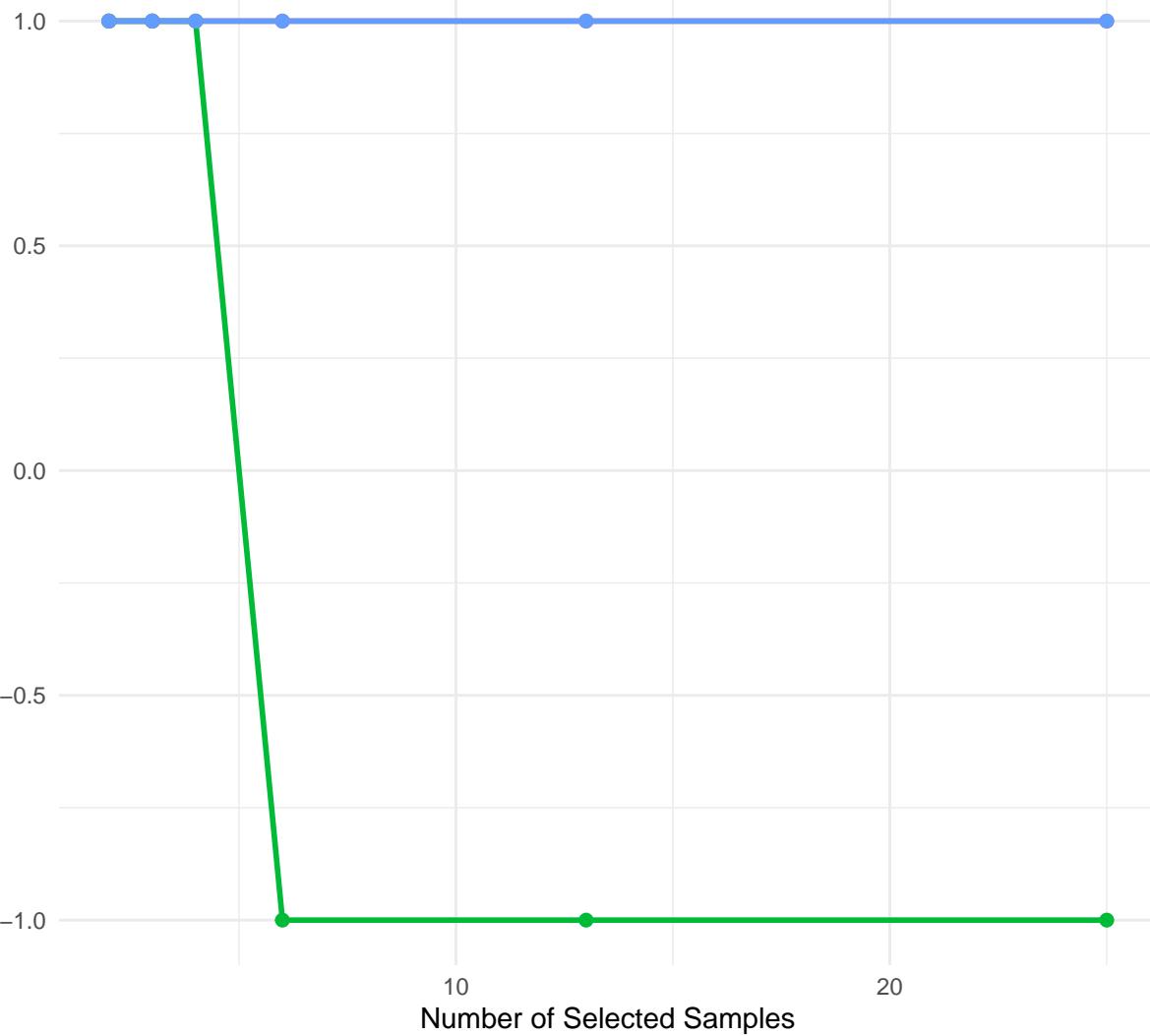
slope

variance

Number of Selected Samples

Score: PR (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

range

slope

variance

Score: NRW (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

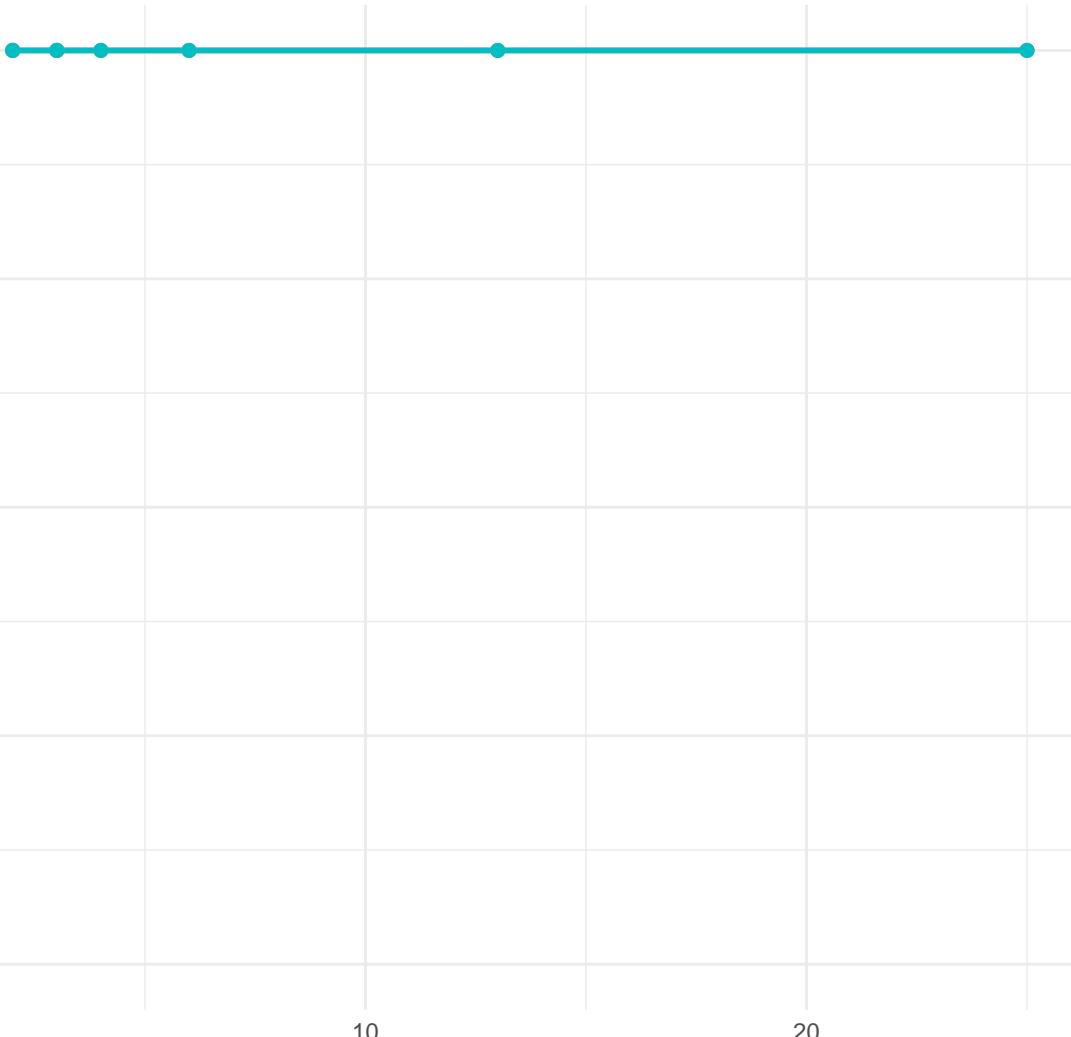
-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: NRW (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

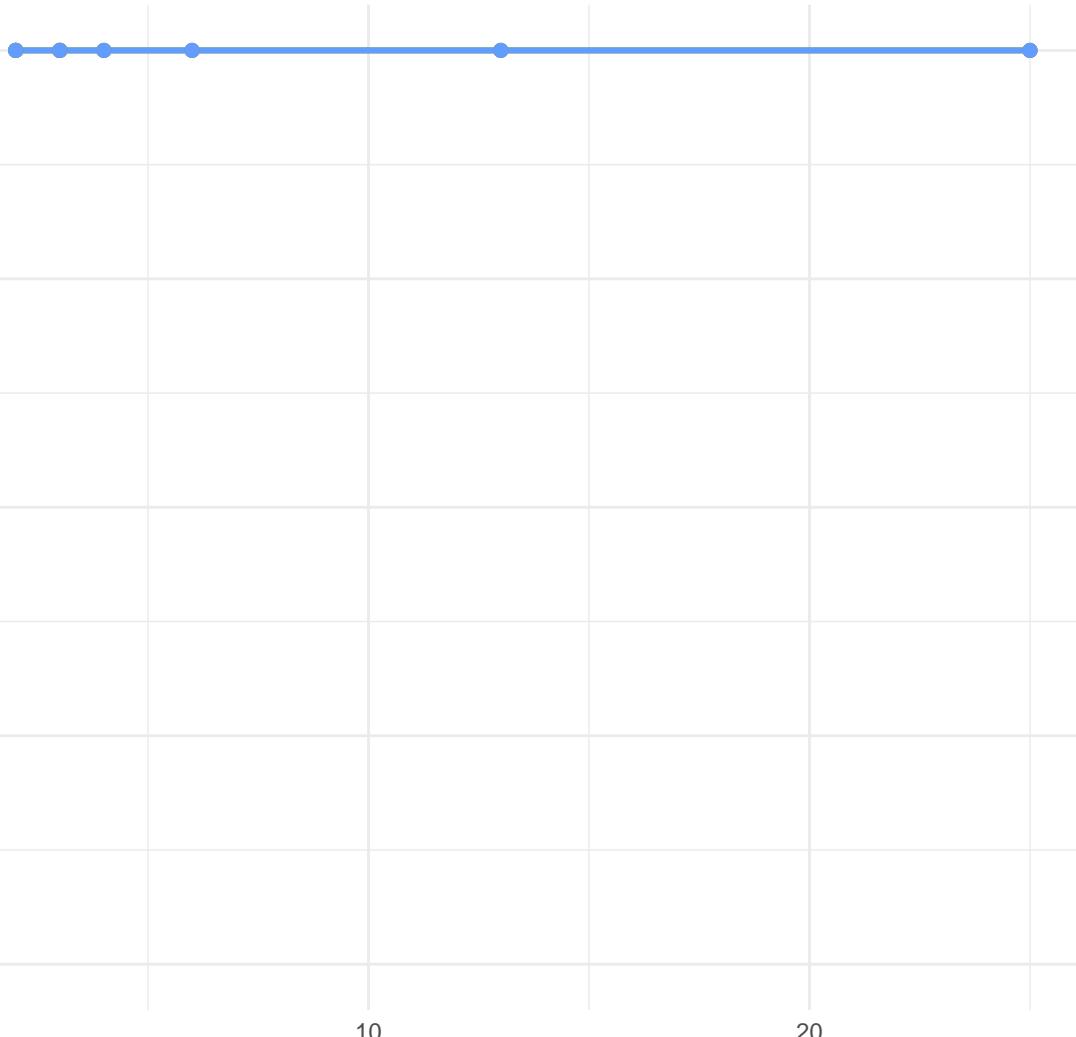
Number of Selected Samples

Summary Statistic

range

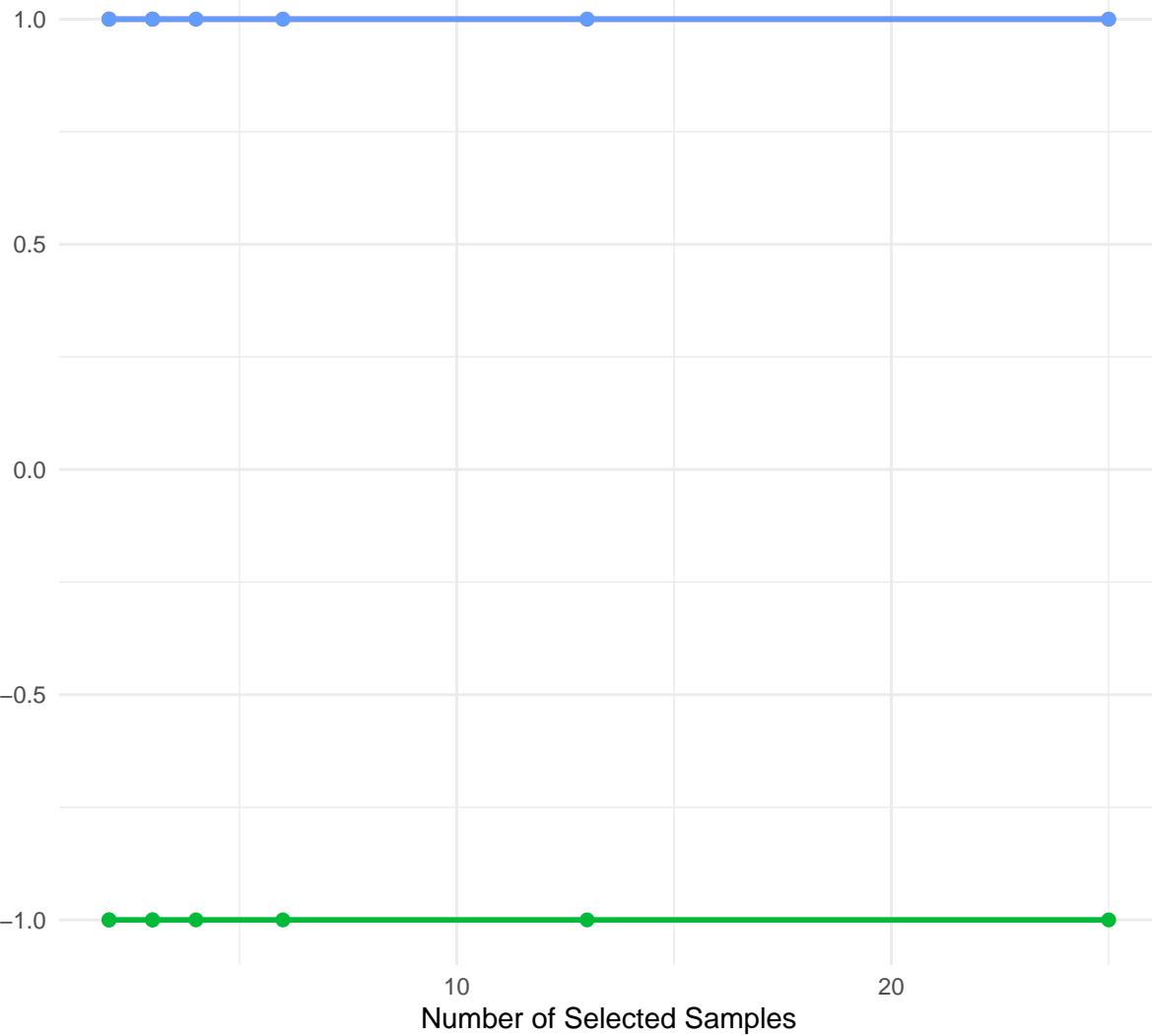
slope

variance



Score: NRW (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

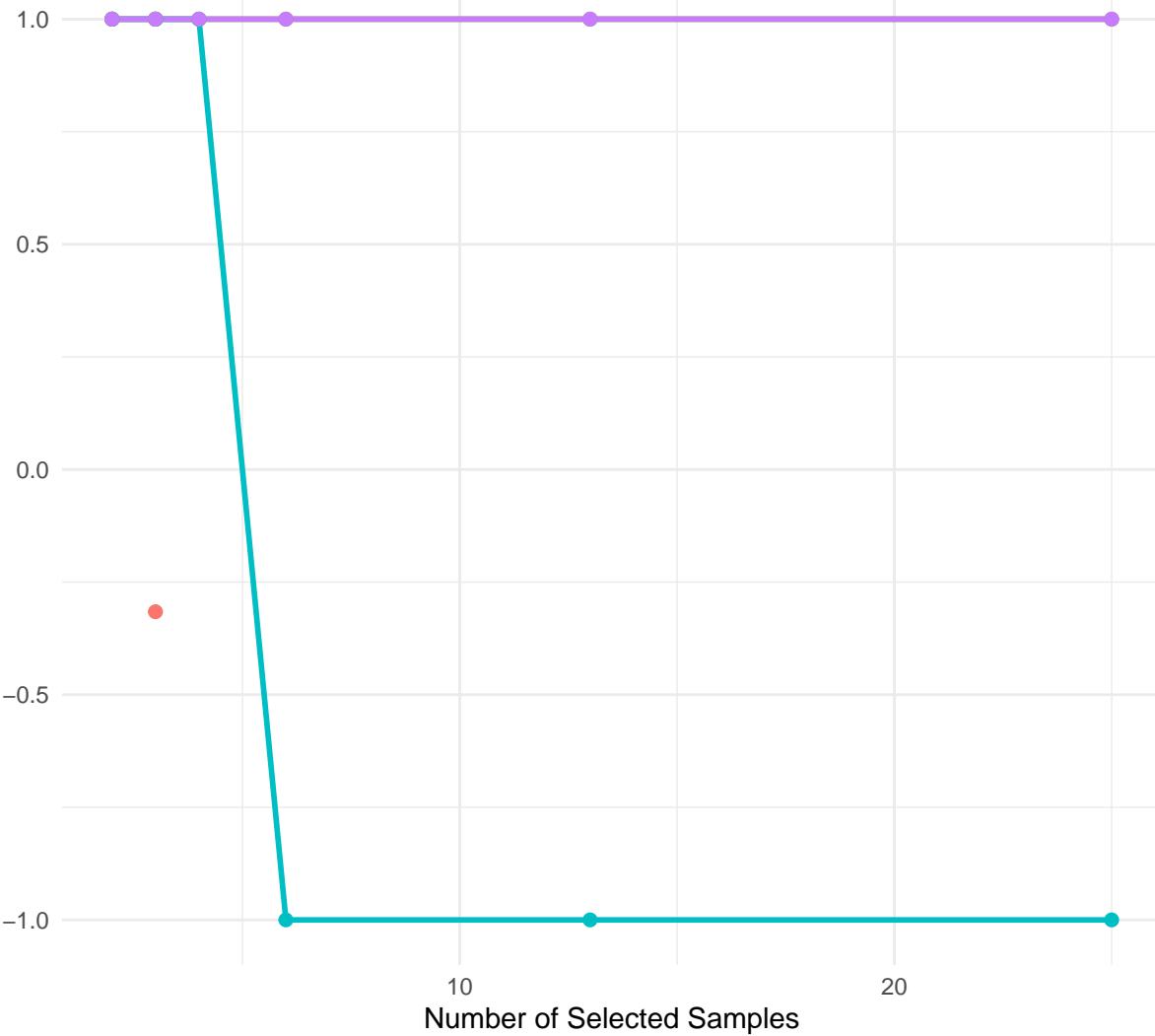
range

slope

variance

Score: NRW (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

- min
- range
- slope
- variance

Score: ESP (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

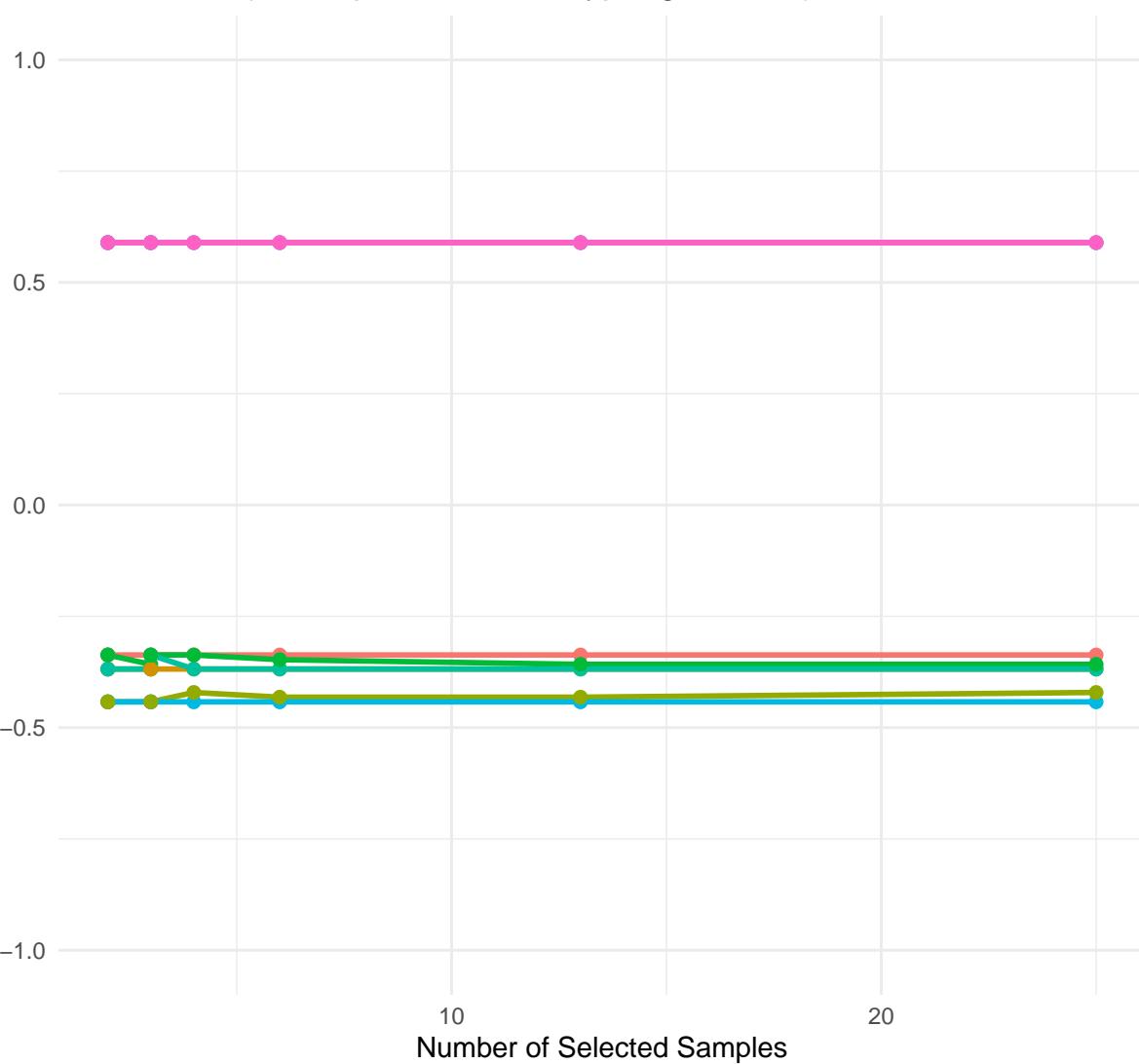
range

variance



Score: ESP (Local, partial1 , Genotype: gaussian)

Kendall Correlation

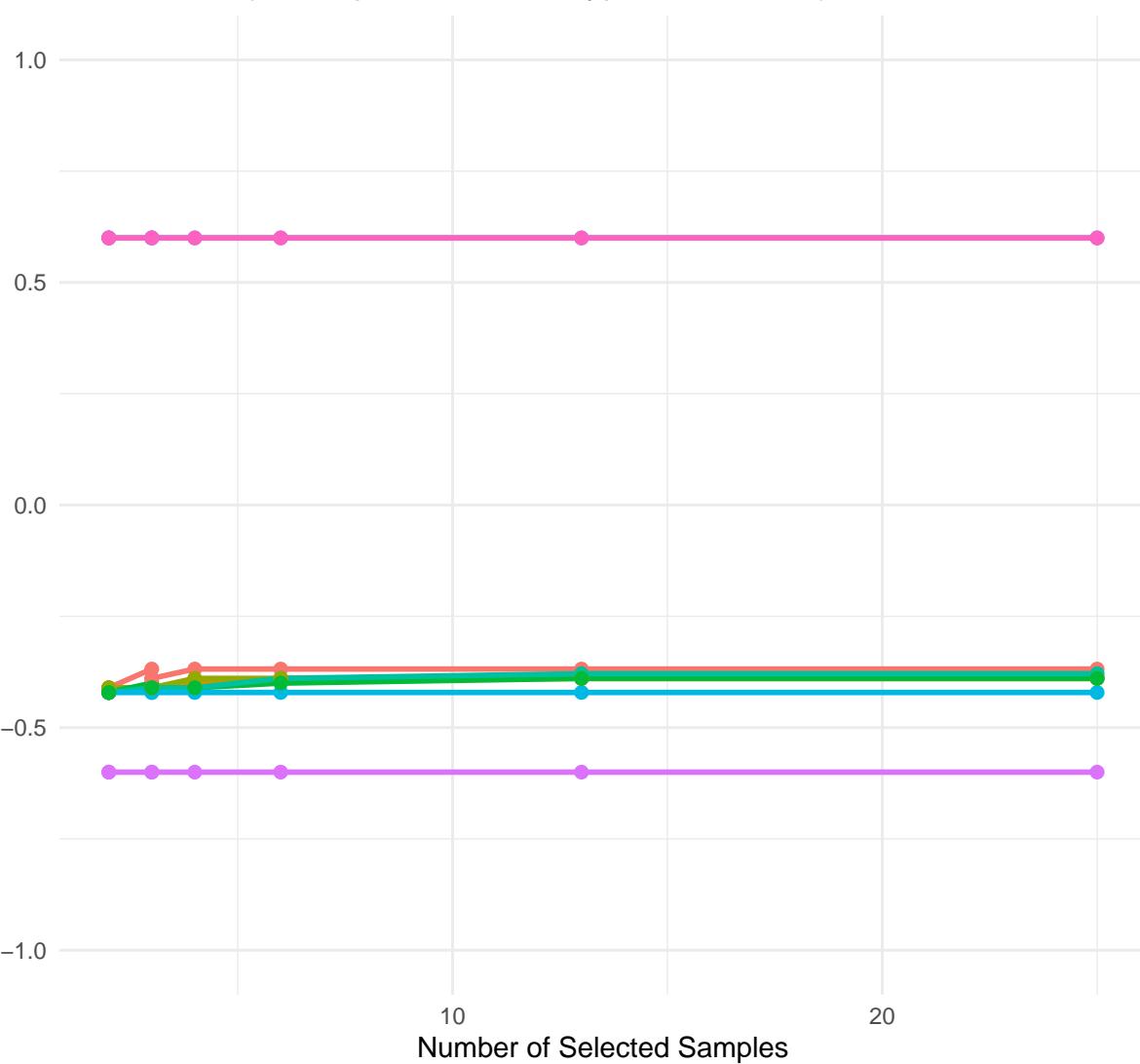


Summary Statistic

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

Score: ESP (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

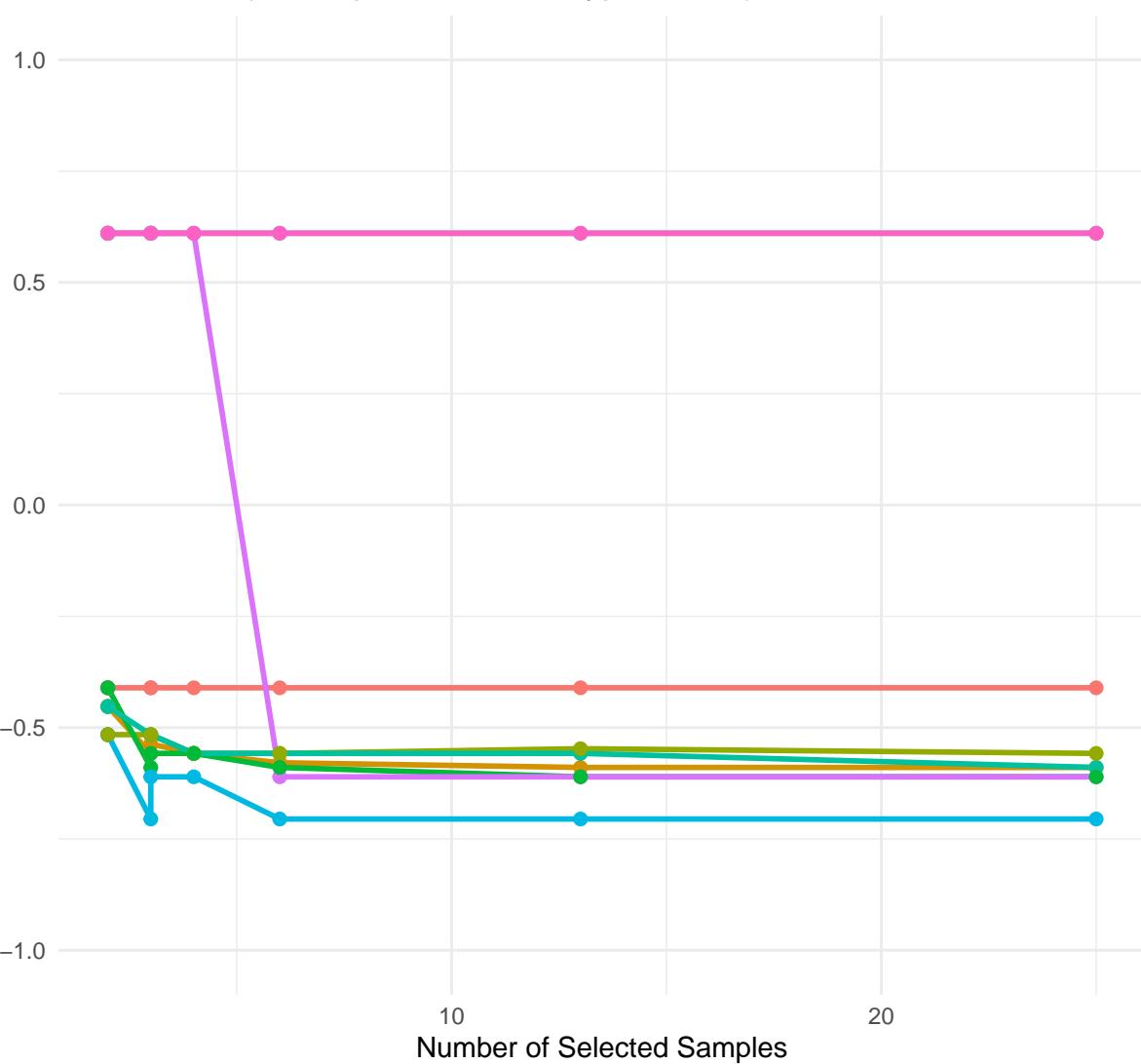


Summary Statistic

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

Score: ESP (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: CEV (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

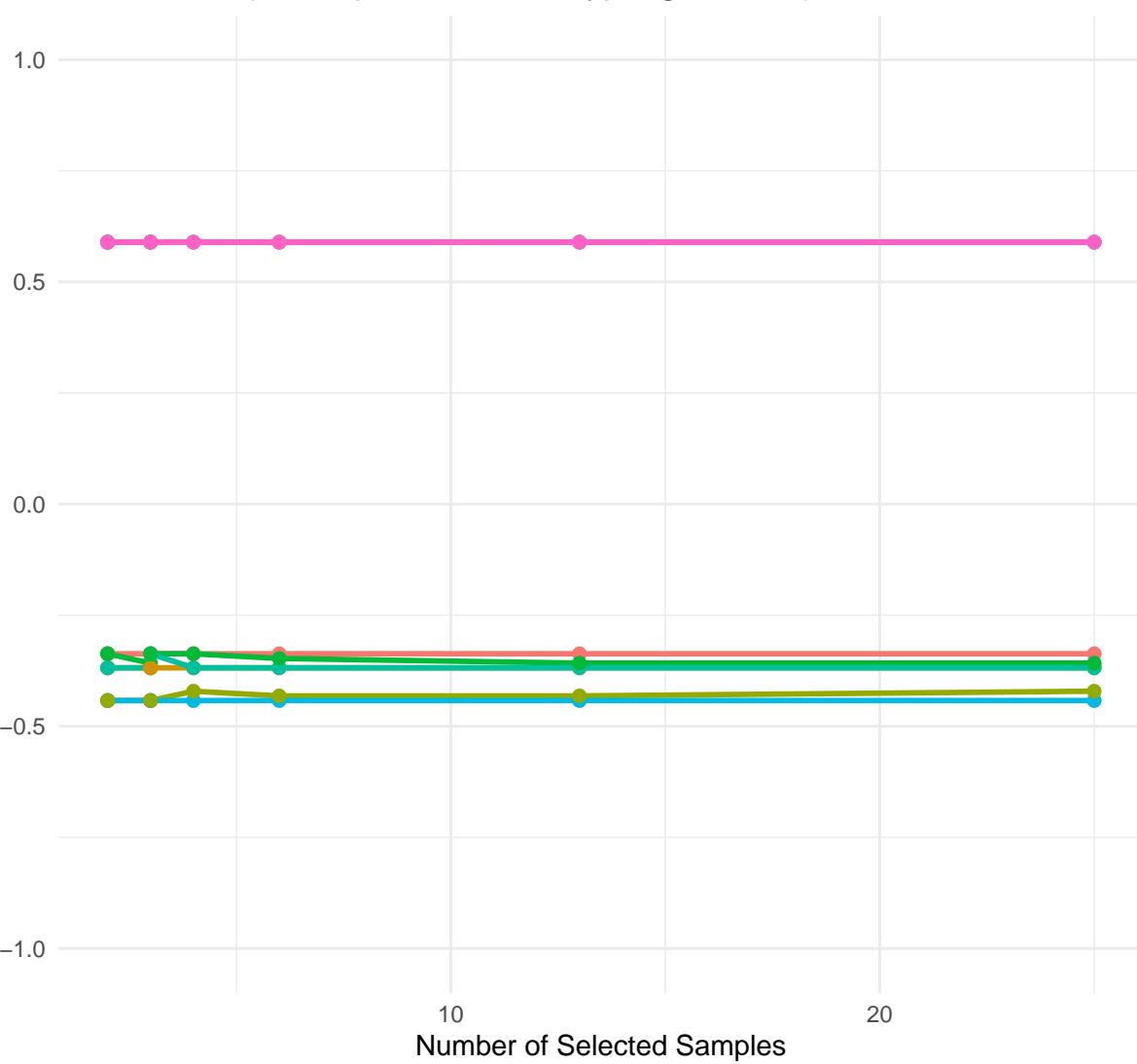
range

variance



Score: CEV (Local, partial1 , Genotype: gaussian)

Kendall Correlation

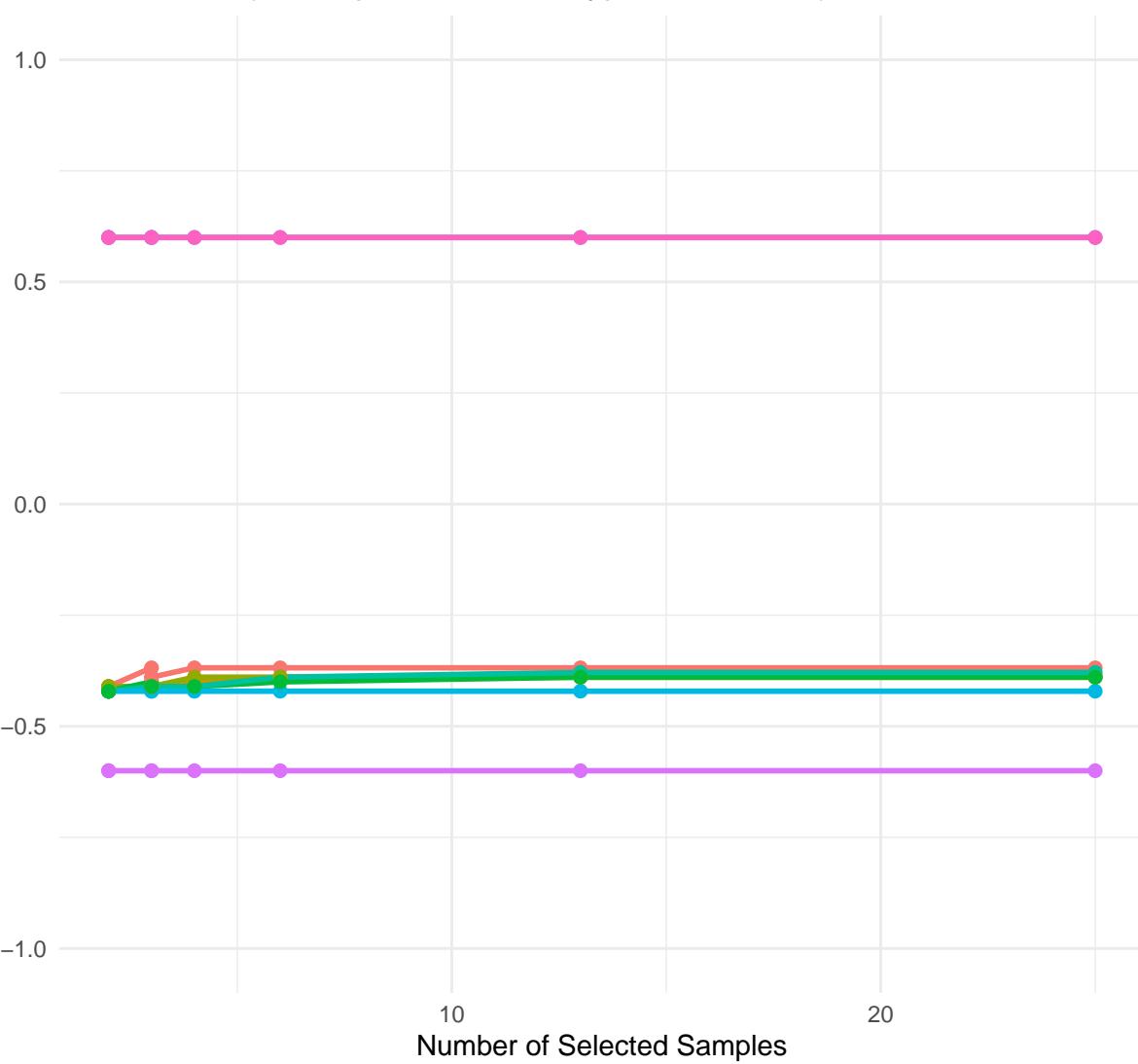


Summary Statistic

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

Score: CEV (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

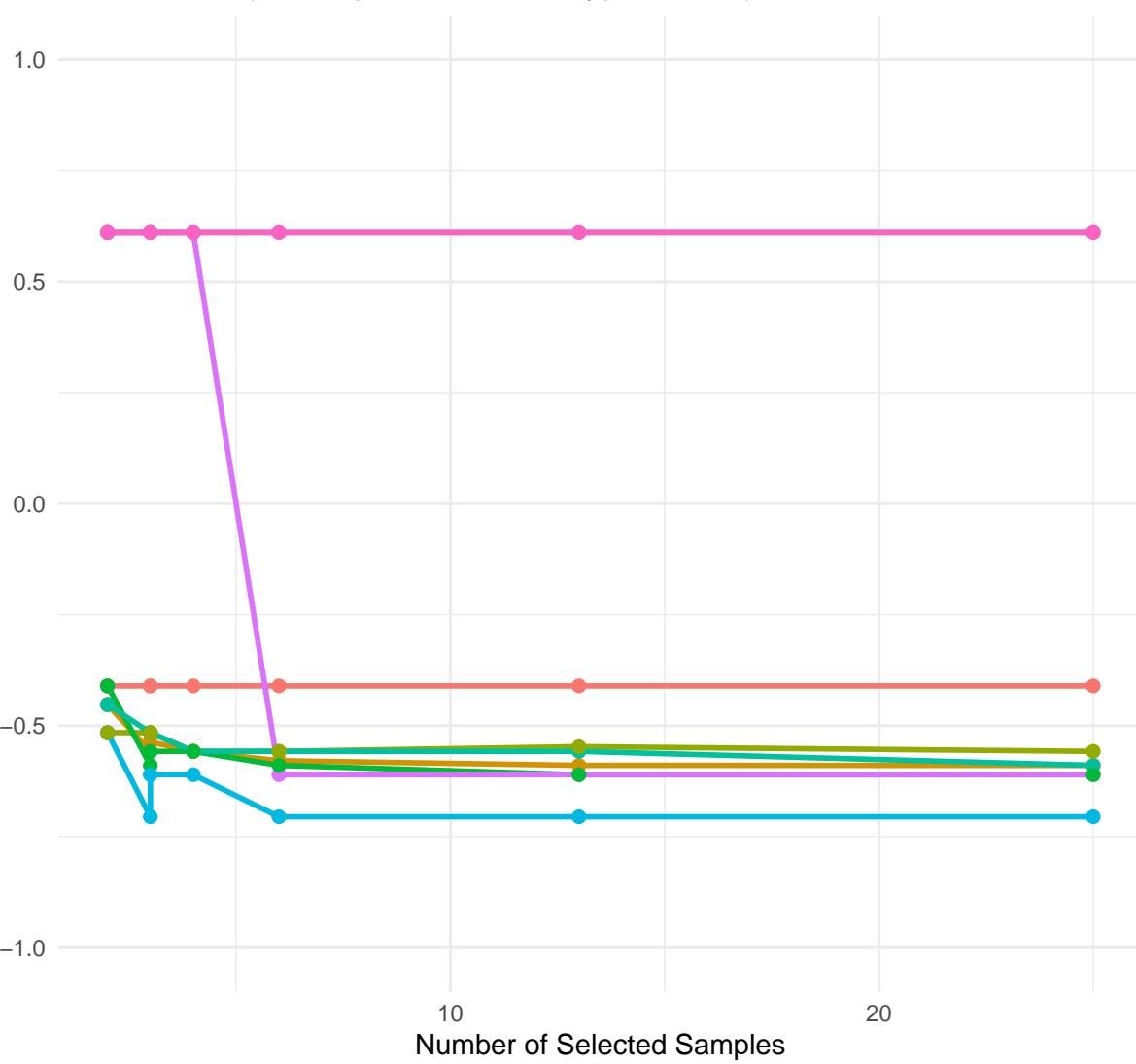


Summary Statistic

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

Score: CEV (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: PFI (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

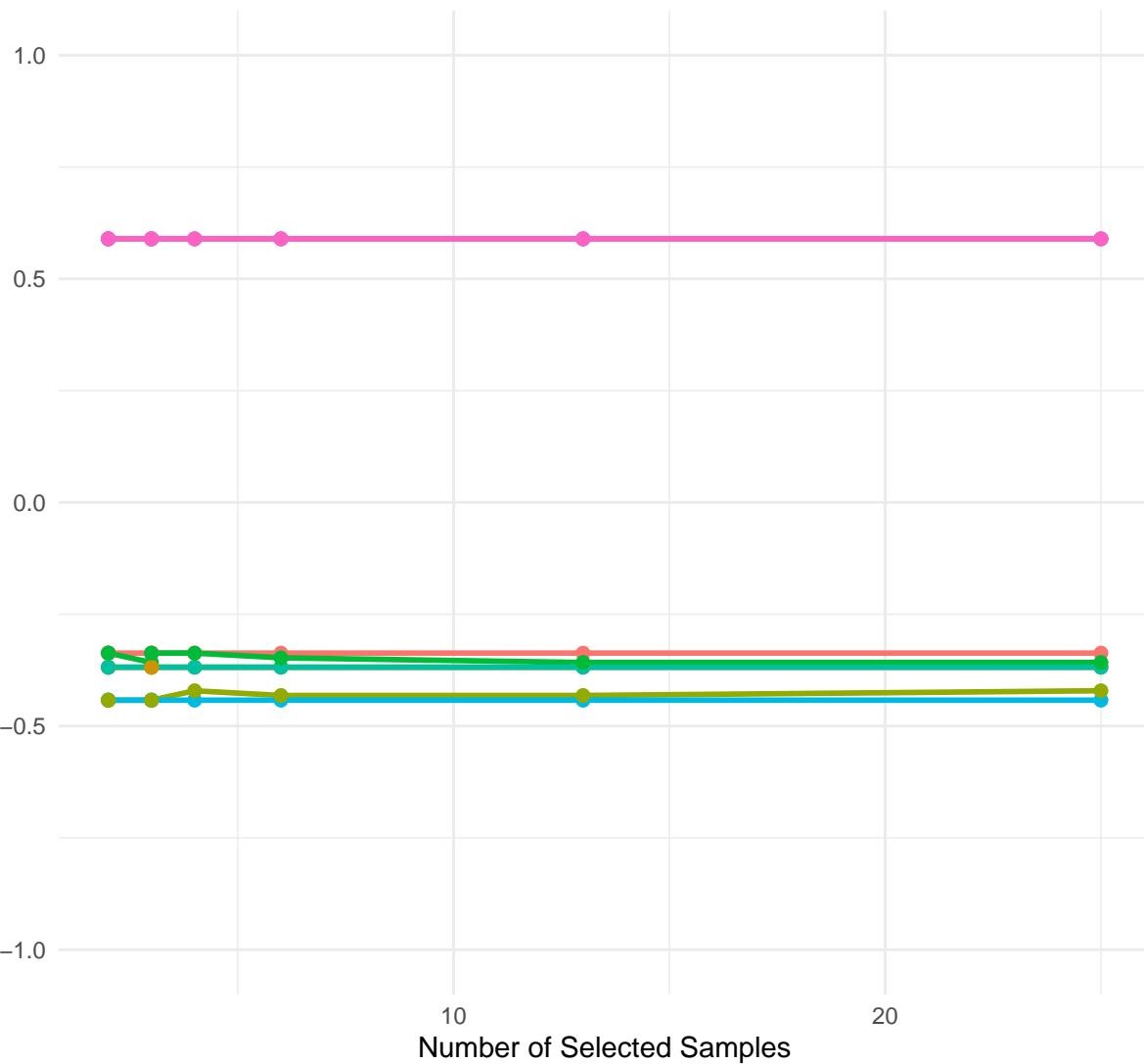
range

variance



Score: PFI (Local, partial1 , Genotype: gaussian)

Kendall Correlation

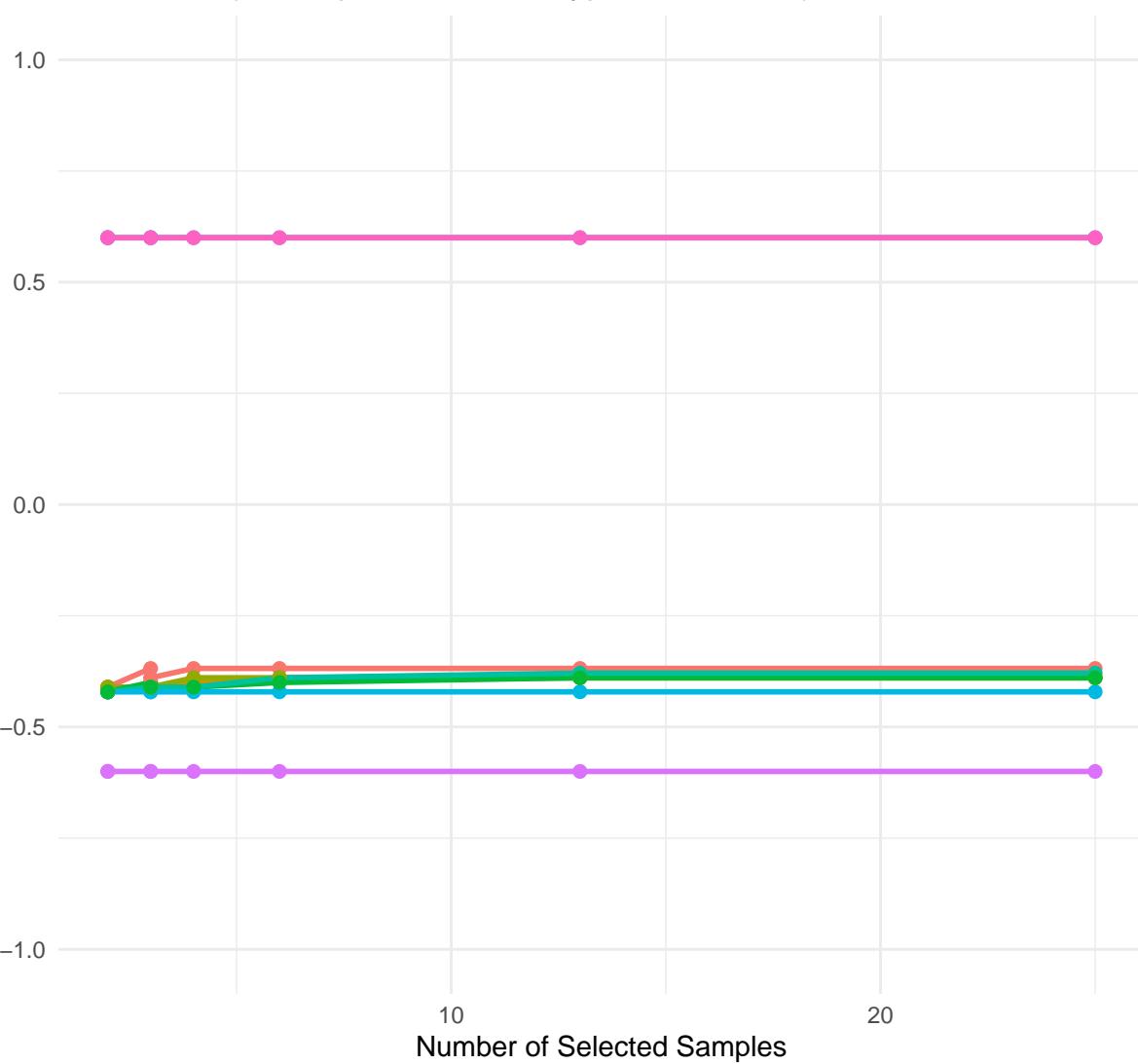


Summary Statistic

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

Score: PFI (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

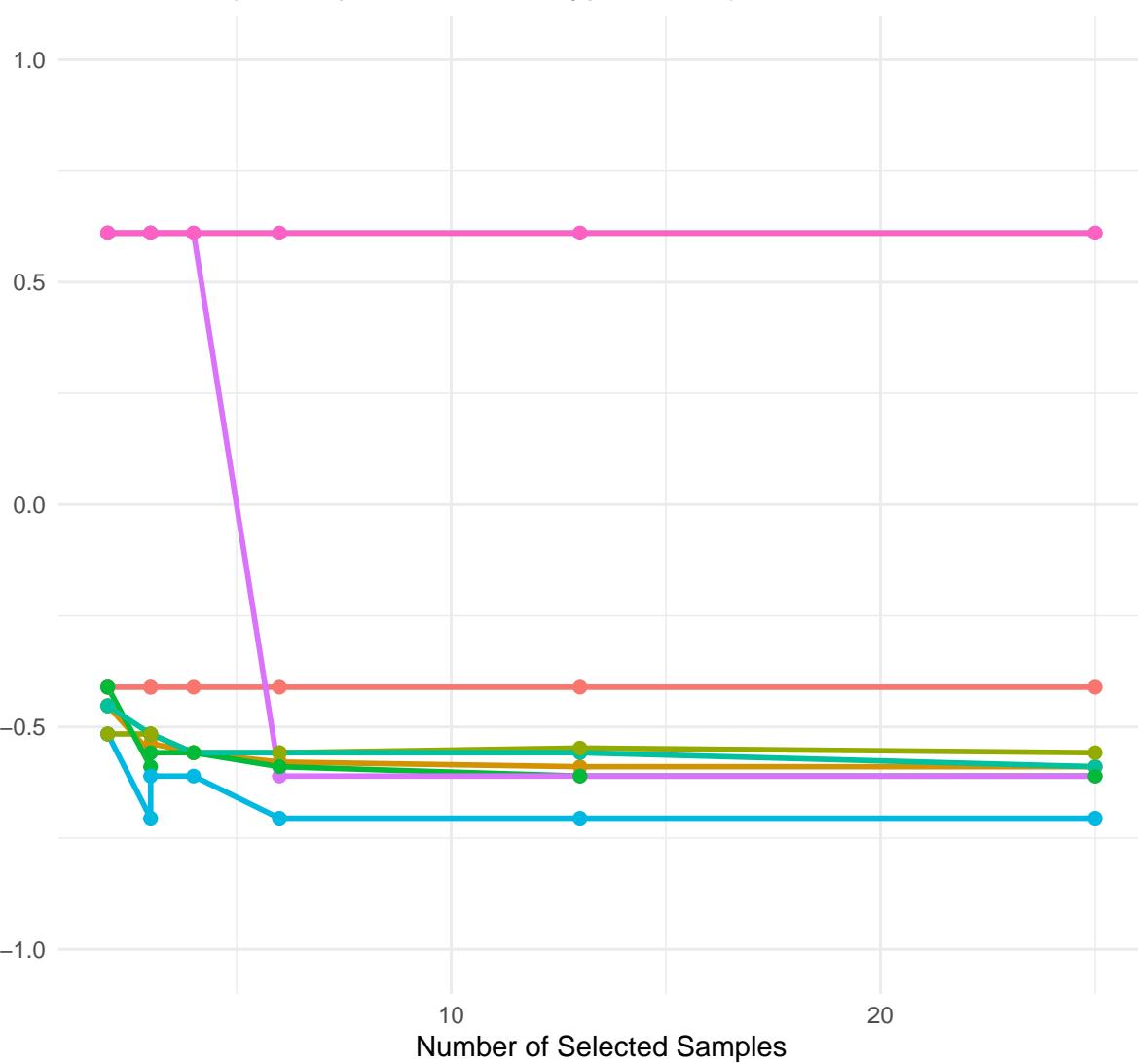


Summary Statistic

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

Score: PFI (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: APC (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: APC (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

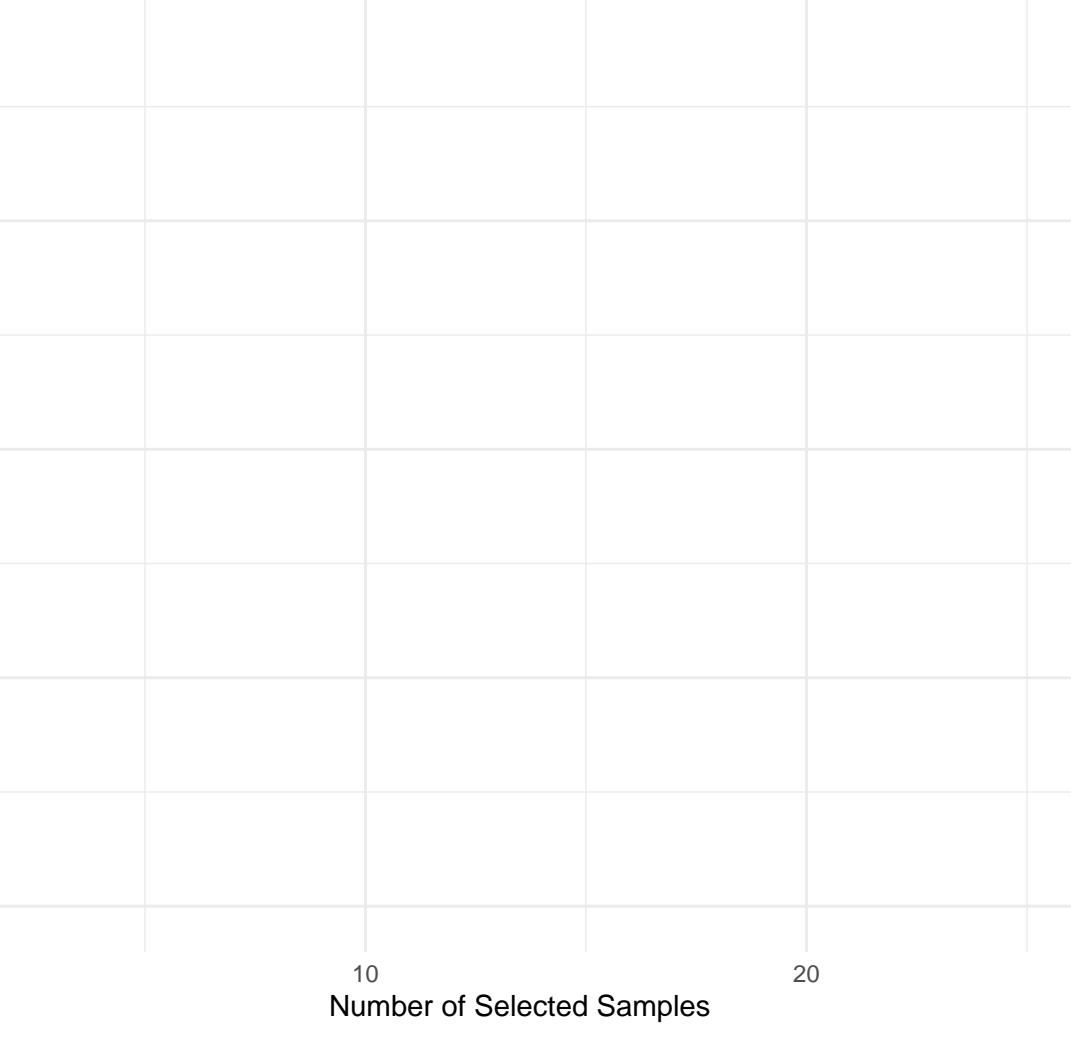
Number of Selected Samples

Summary Statistic

range

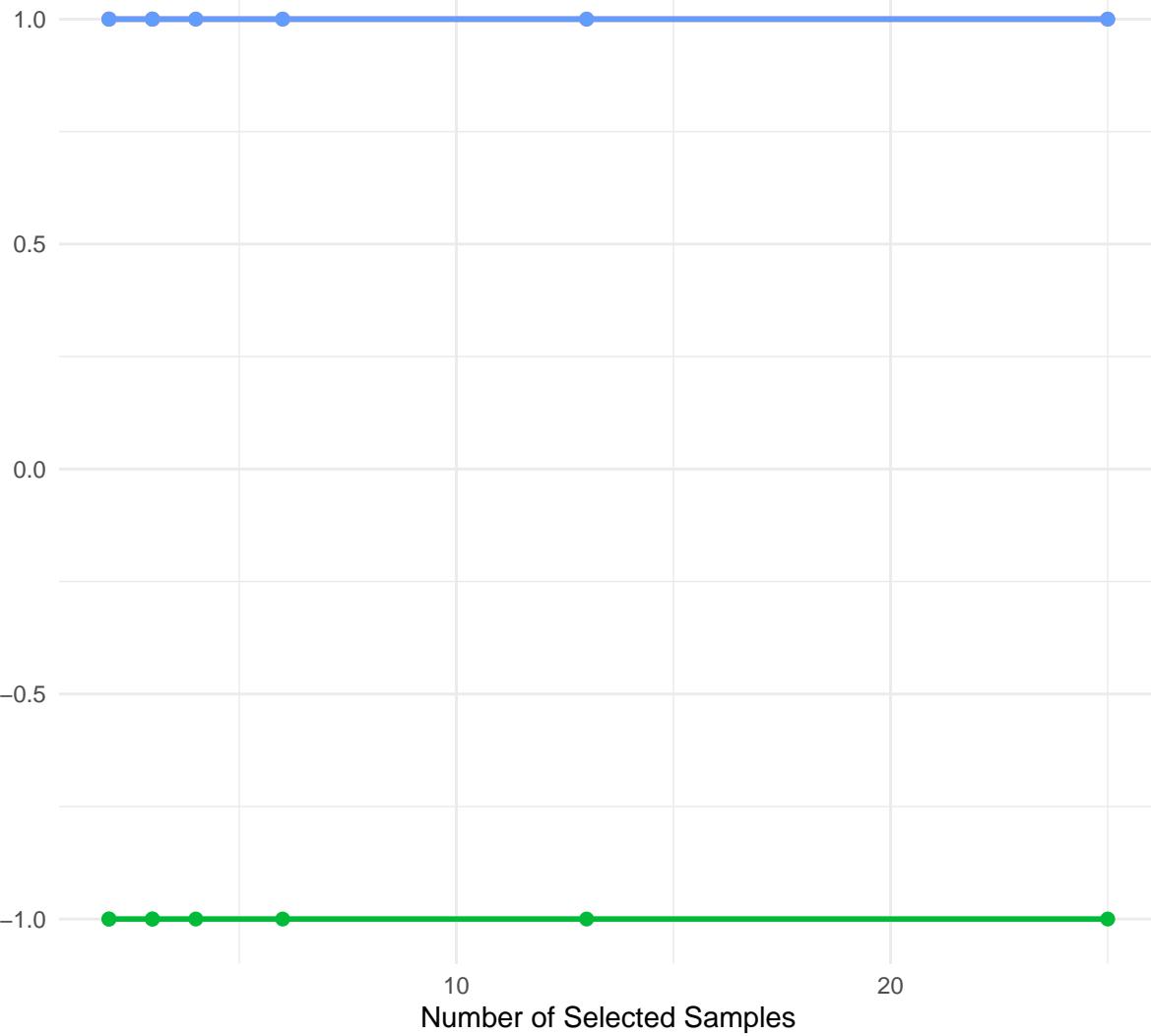
slope

variance



Score: APC (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

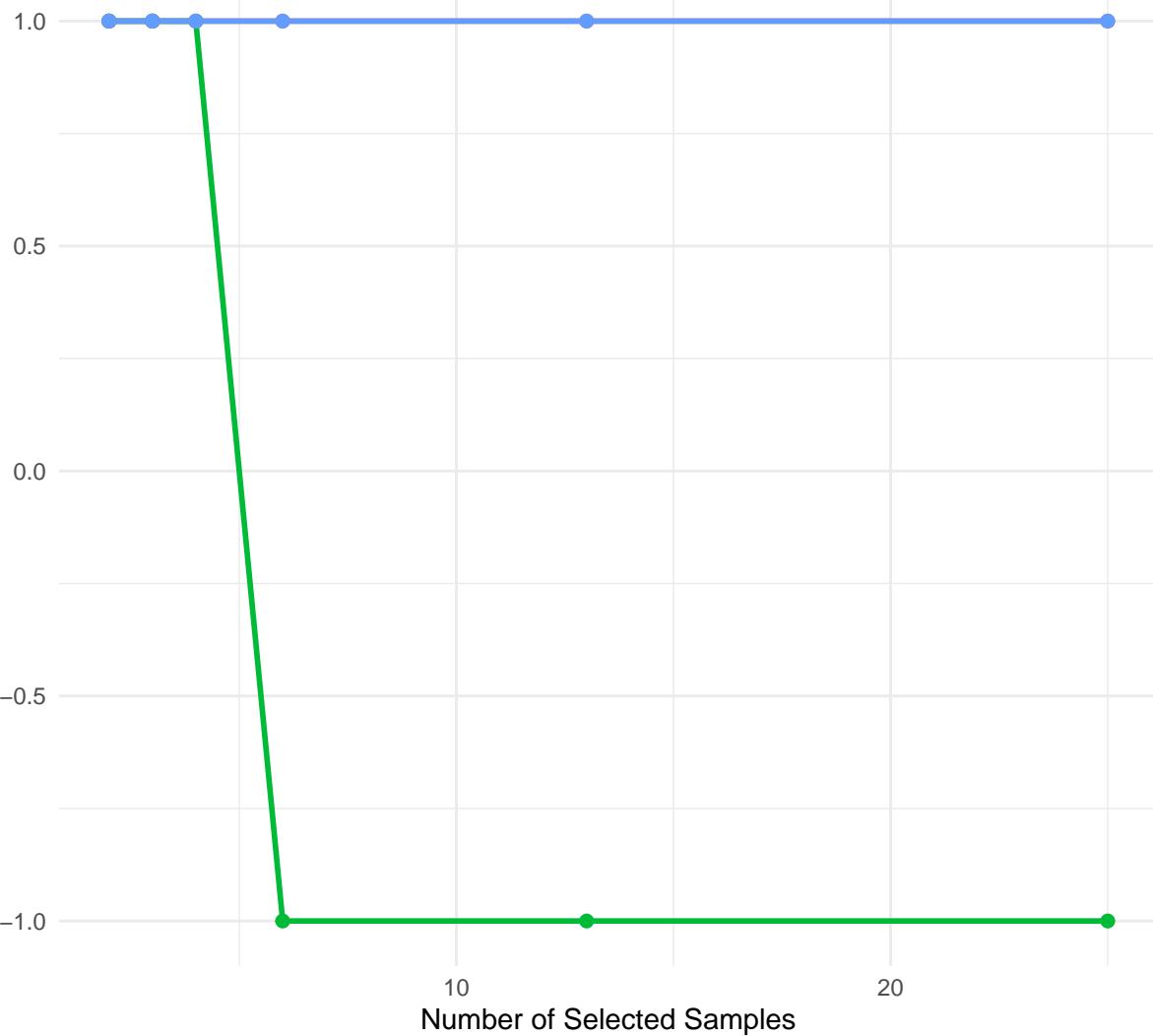
range

slope

variance

Score: APC (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

- range
- slope
- variance

Score: SI (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

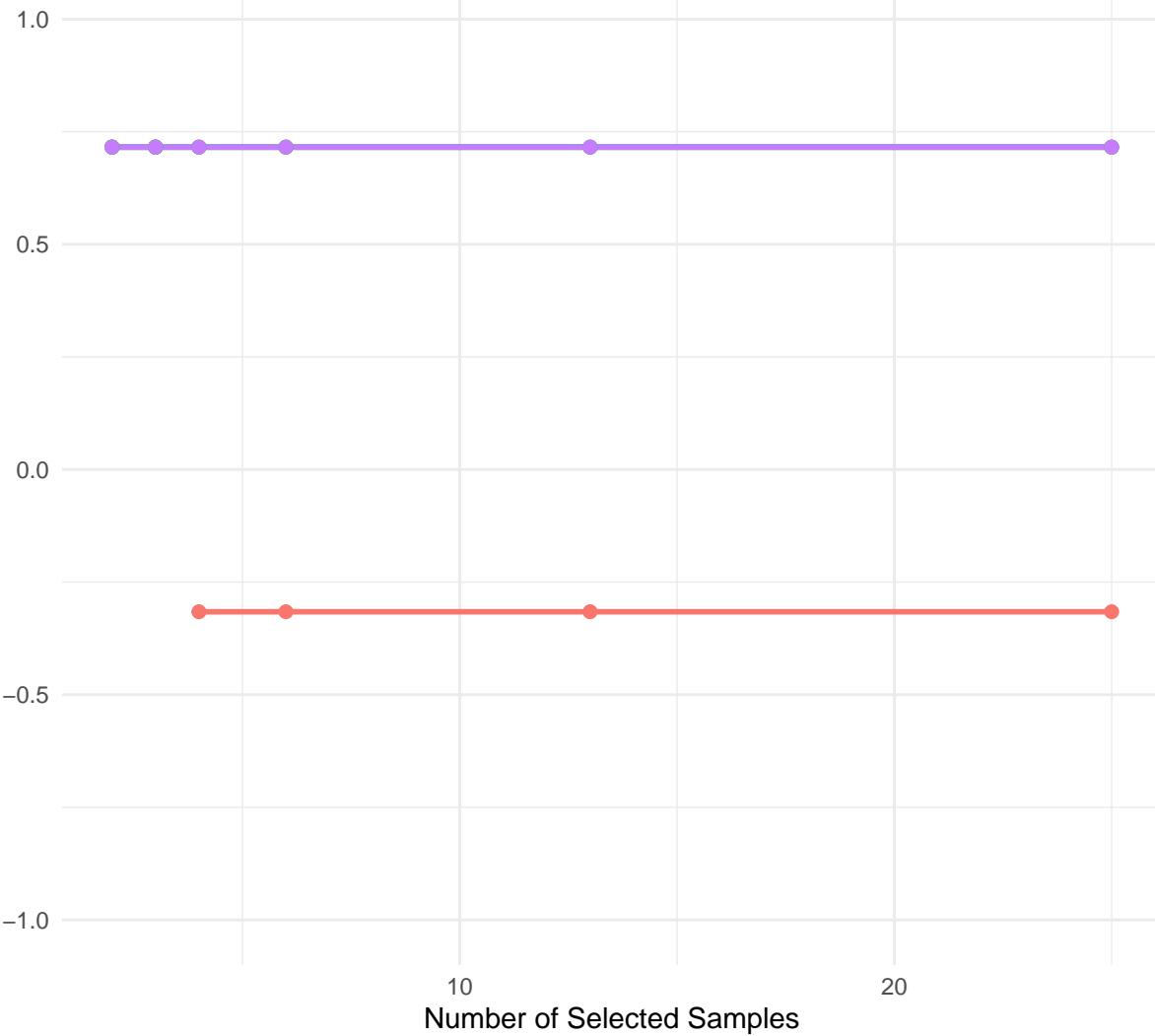
range

variance



Score: SI (Local, partial1 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

- min
- range
- slope
- variance

Score: SI (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

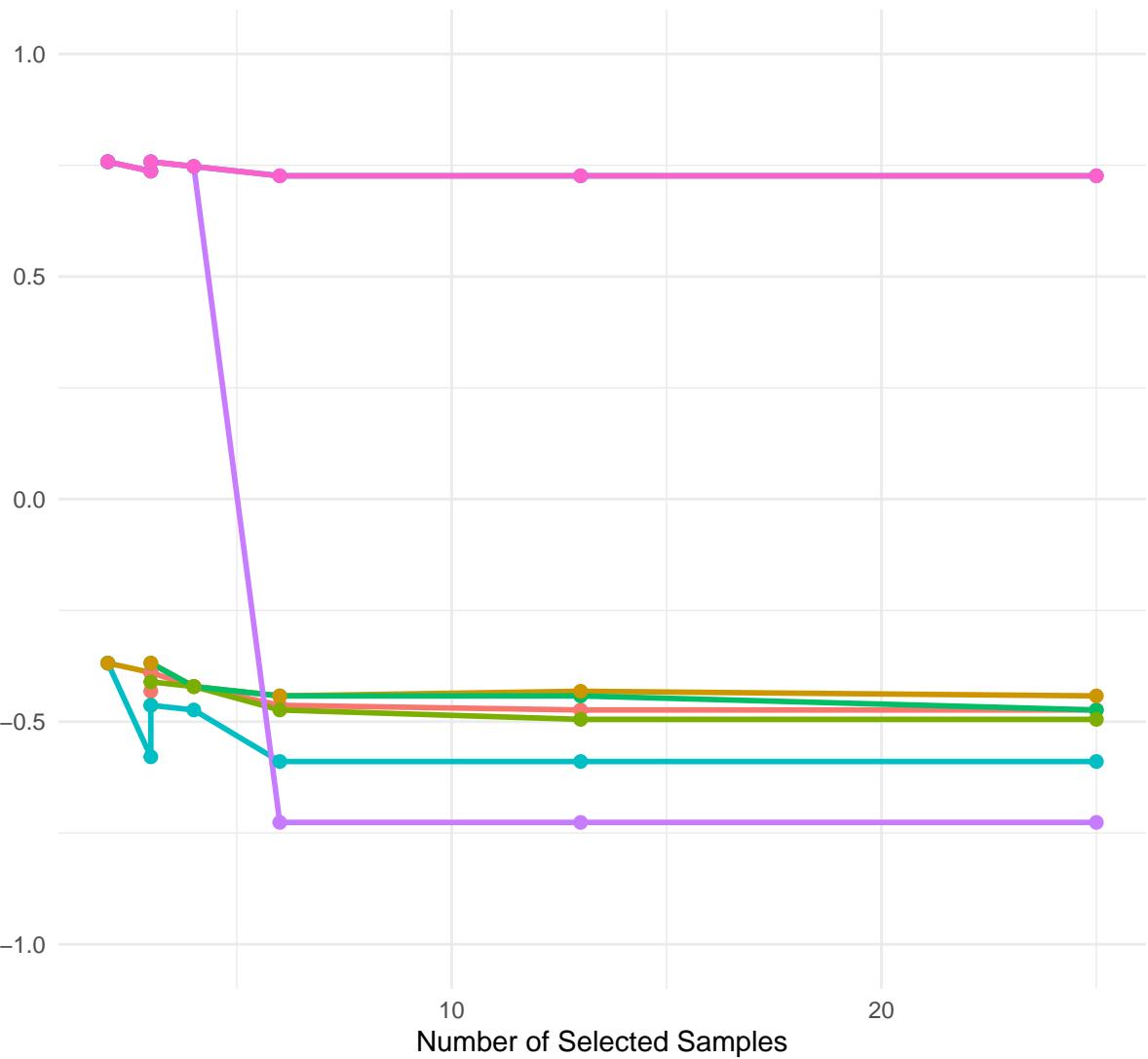
slope

variance



Score: SI (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Number of Selected Samples

Score: RSI (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: RSI (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

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

4.8 4.8 4.8 4.8 4.8 4.8 4.8 4.8 4.8

4.2 4.2 4.2 4.2 4.2 4.2 4.2 4.2 4.2

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3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

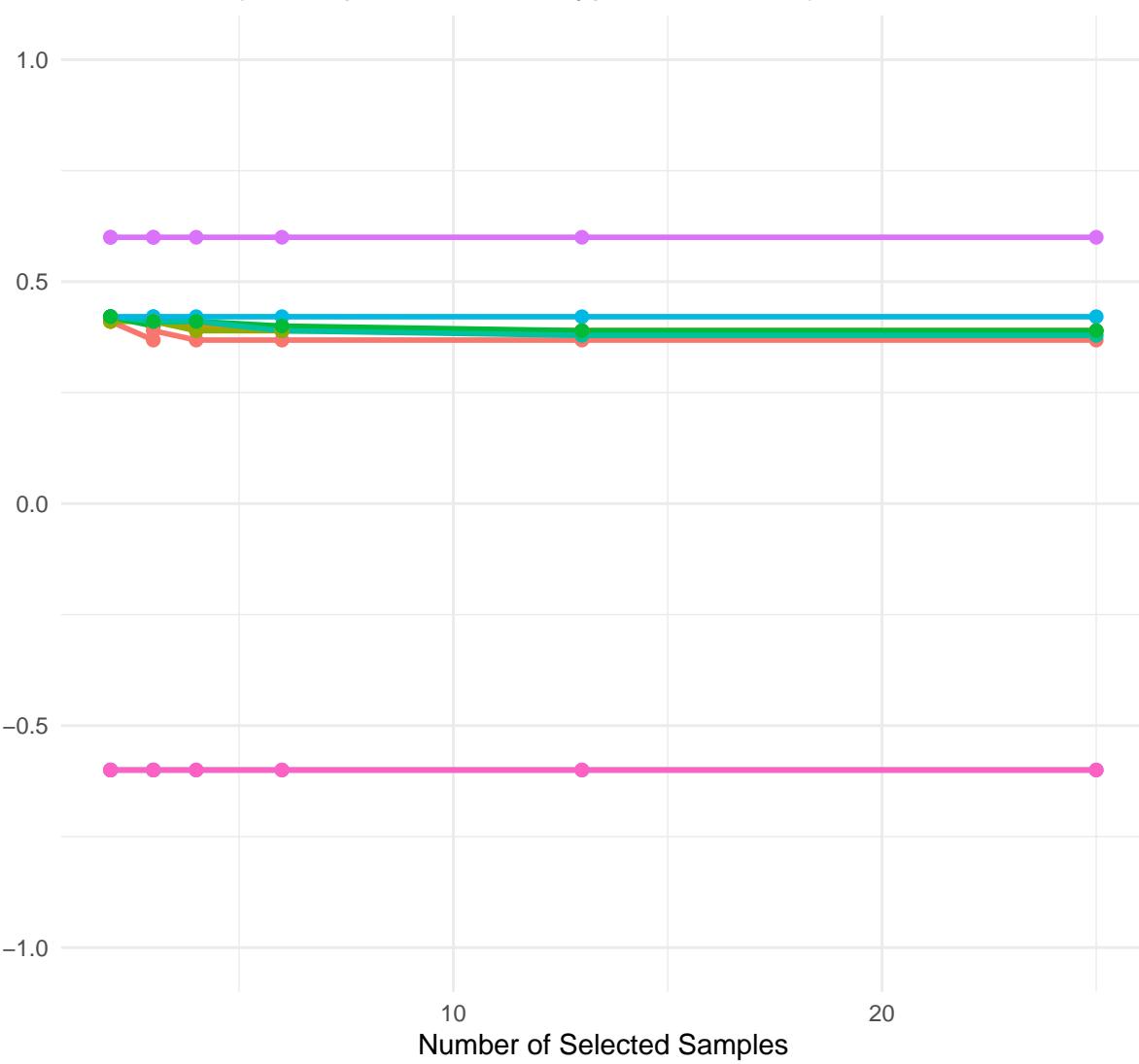
3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6 3.6

Score: RSI (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation

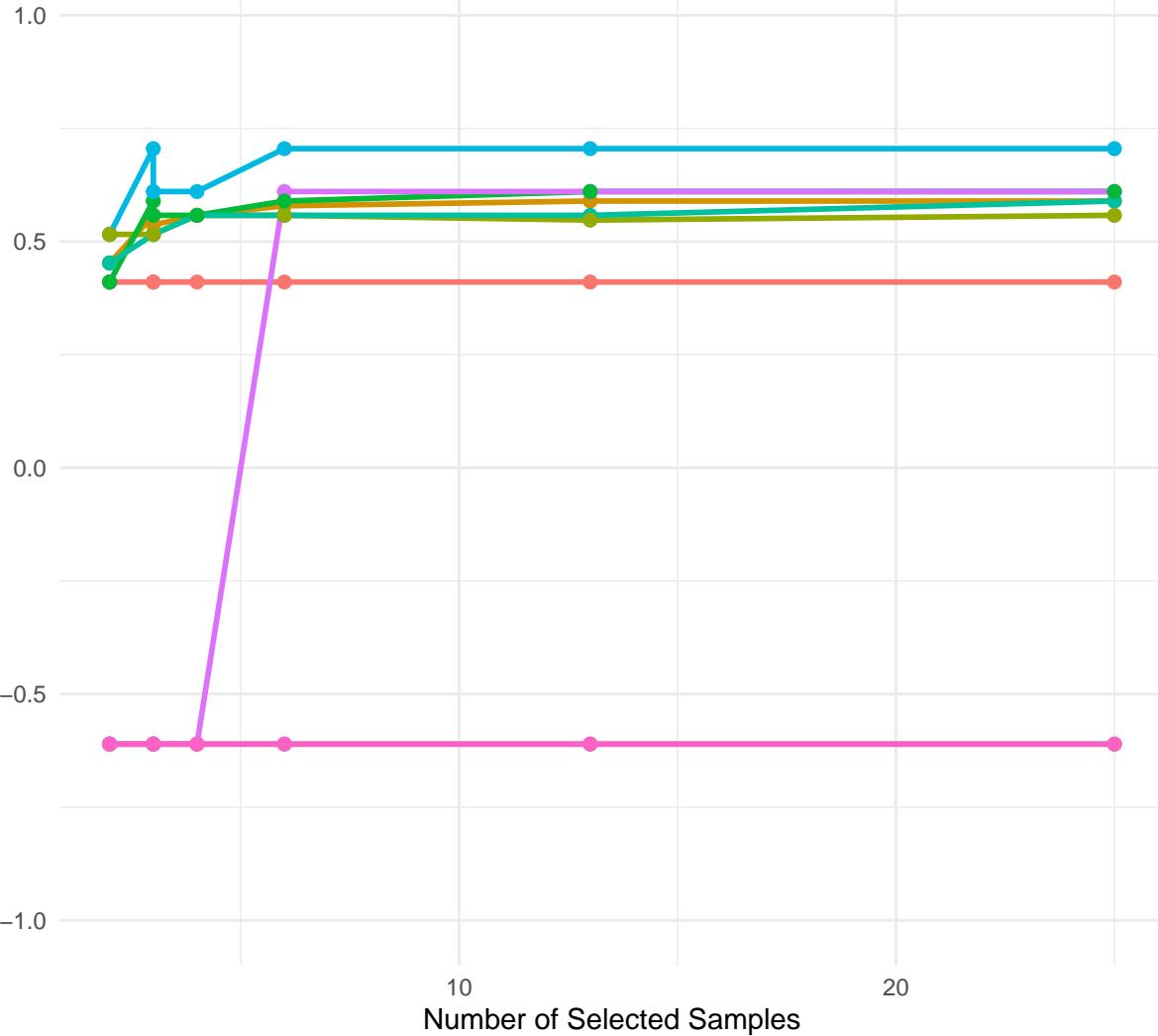


Summary Statistic

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

Score: RSI (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: EVS (Local, partial1 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

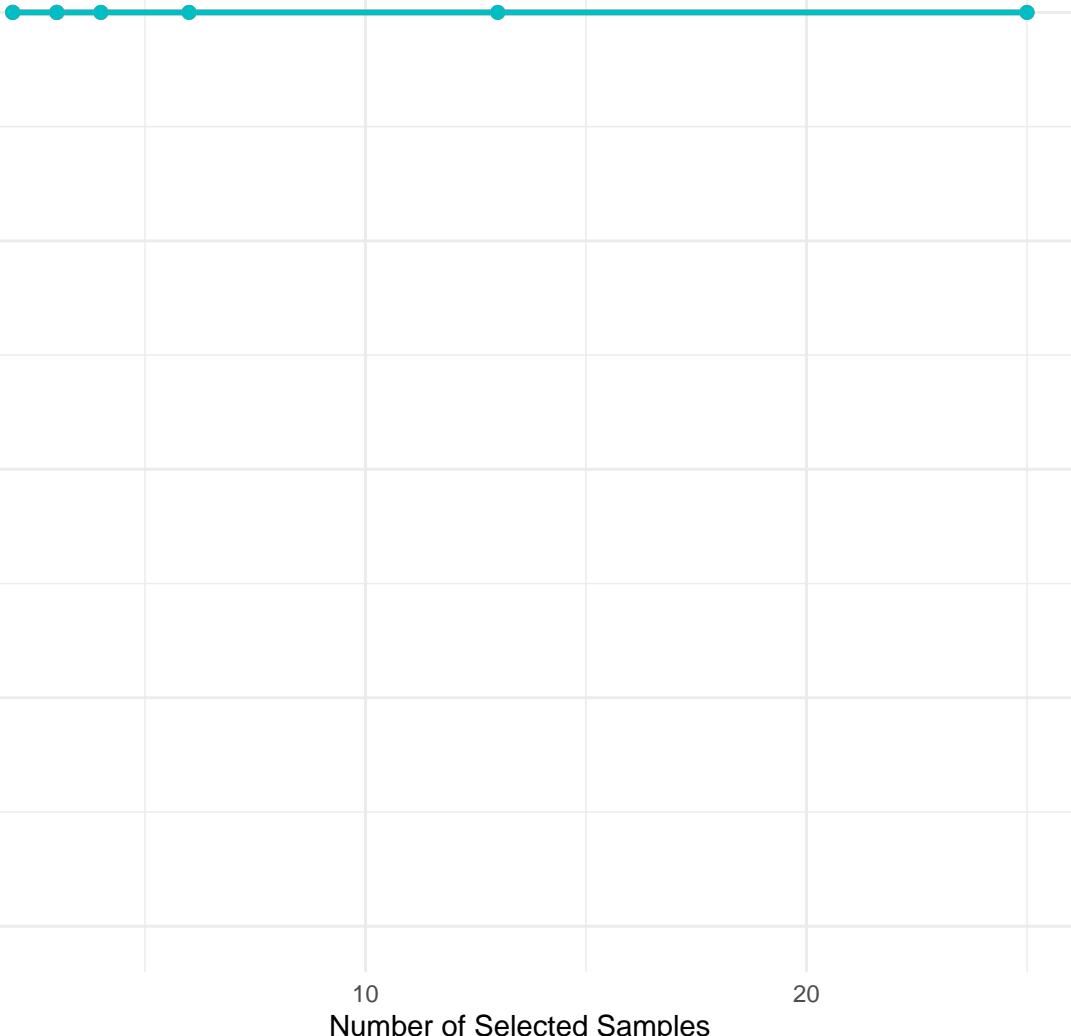
-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: EVS (Local, partial1 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

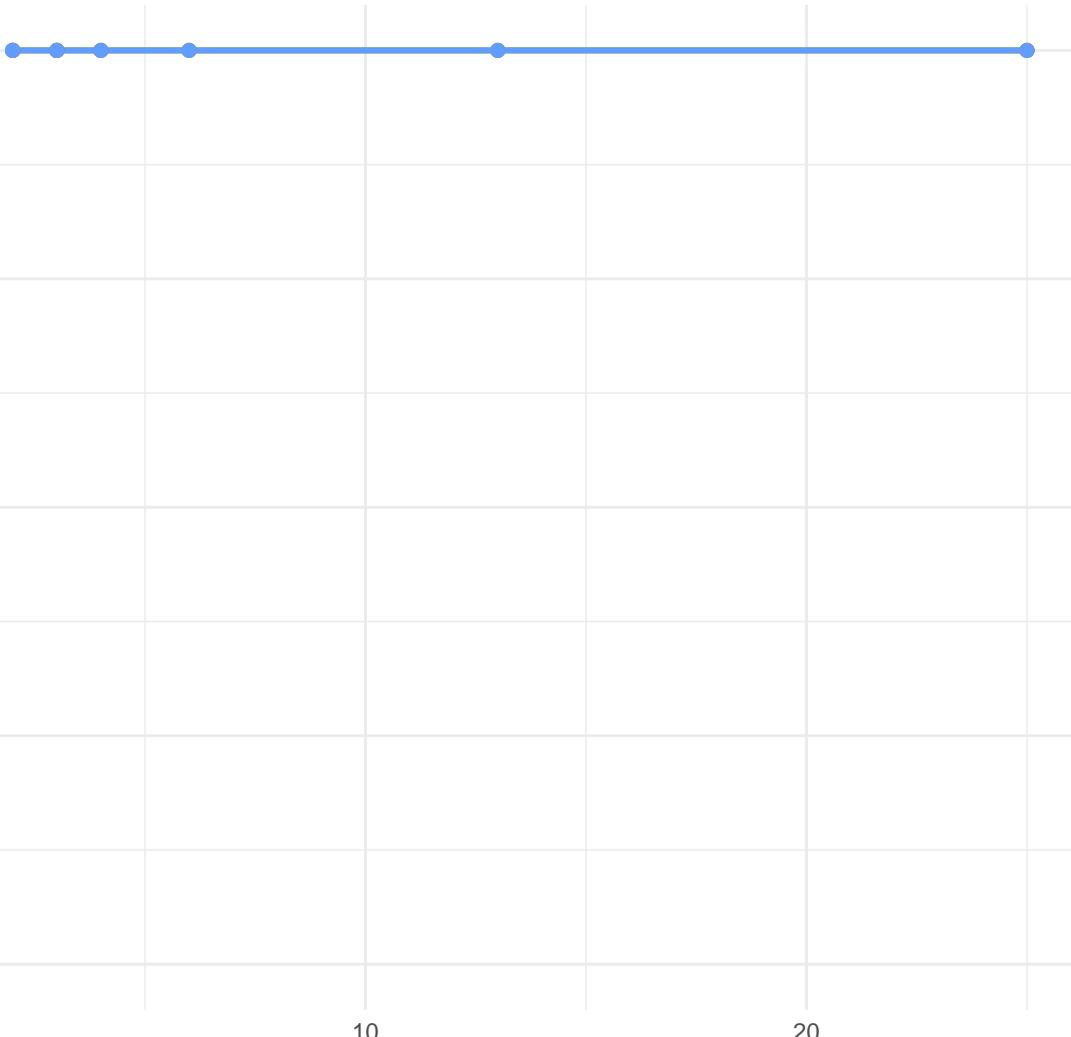
Number of Selected Samples

Summary Statistic

range

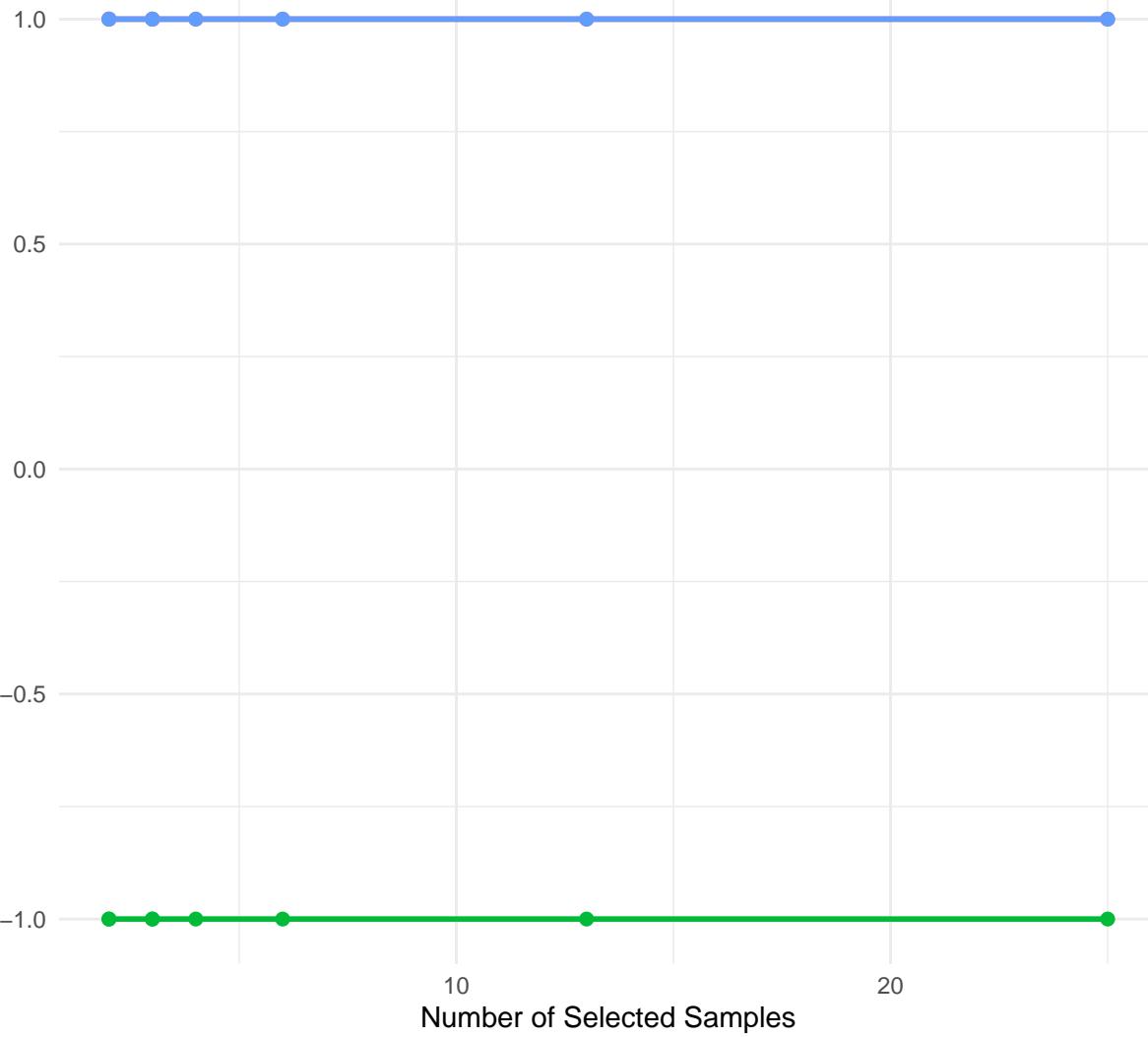
slope

variance



Score: EVS (Local, partial1 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

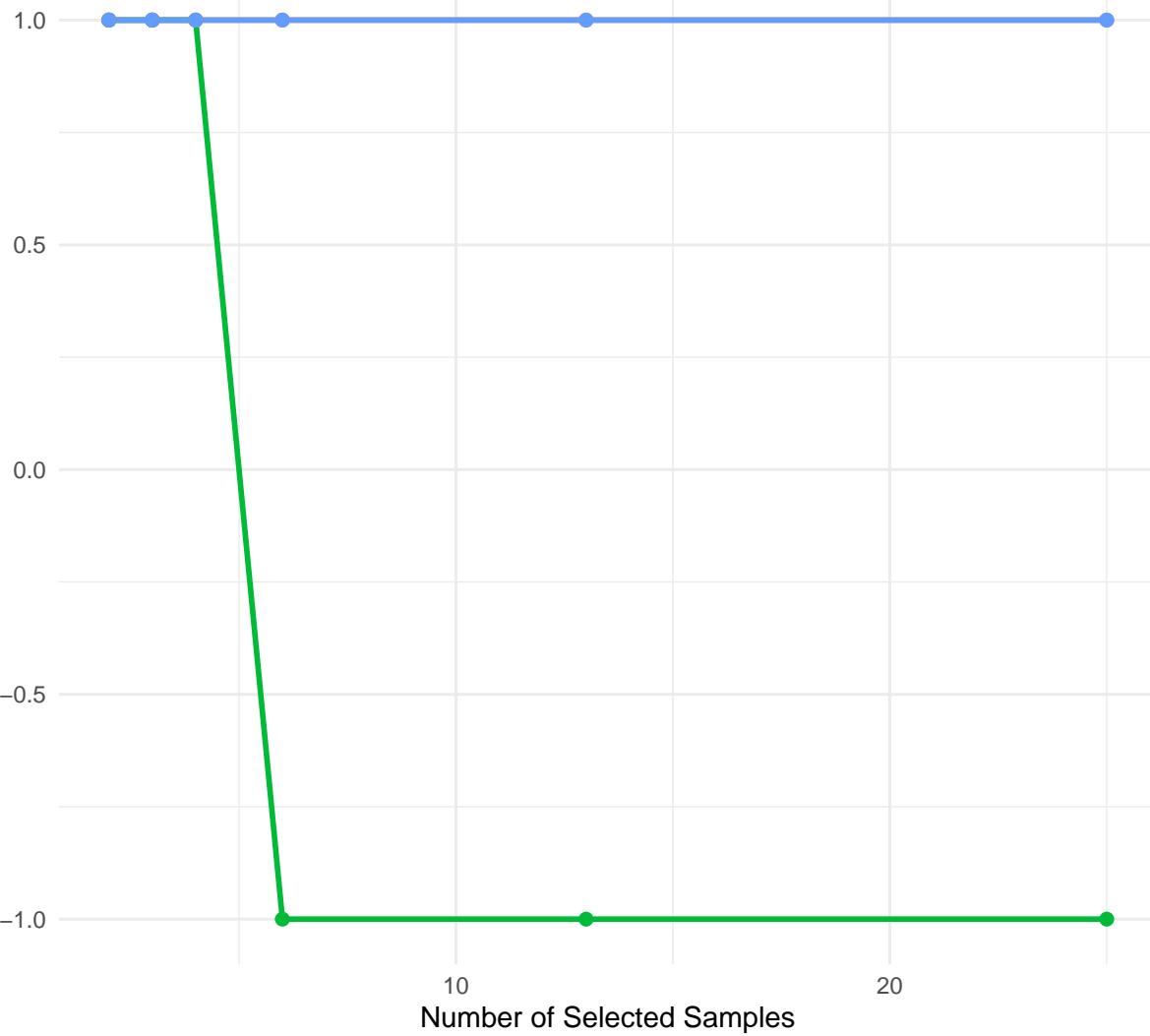
range

slope

variance

Score: EVS (Local, partial1 , Genotype: wave)

Kendall Correlation



Summary Statistic

range

slope

variance