

Score: CV_t (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

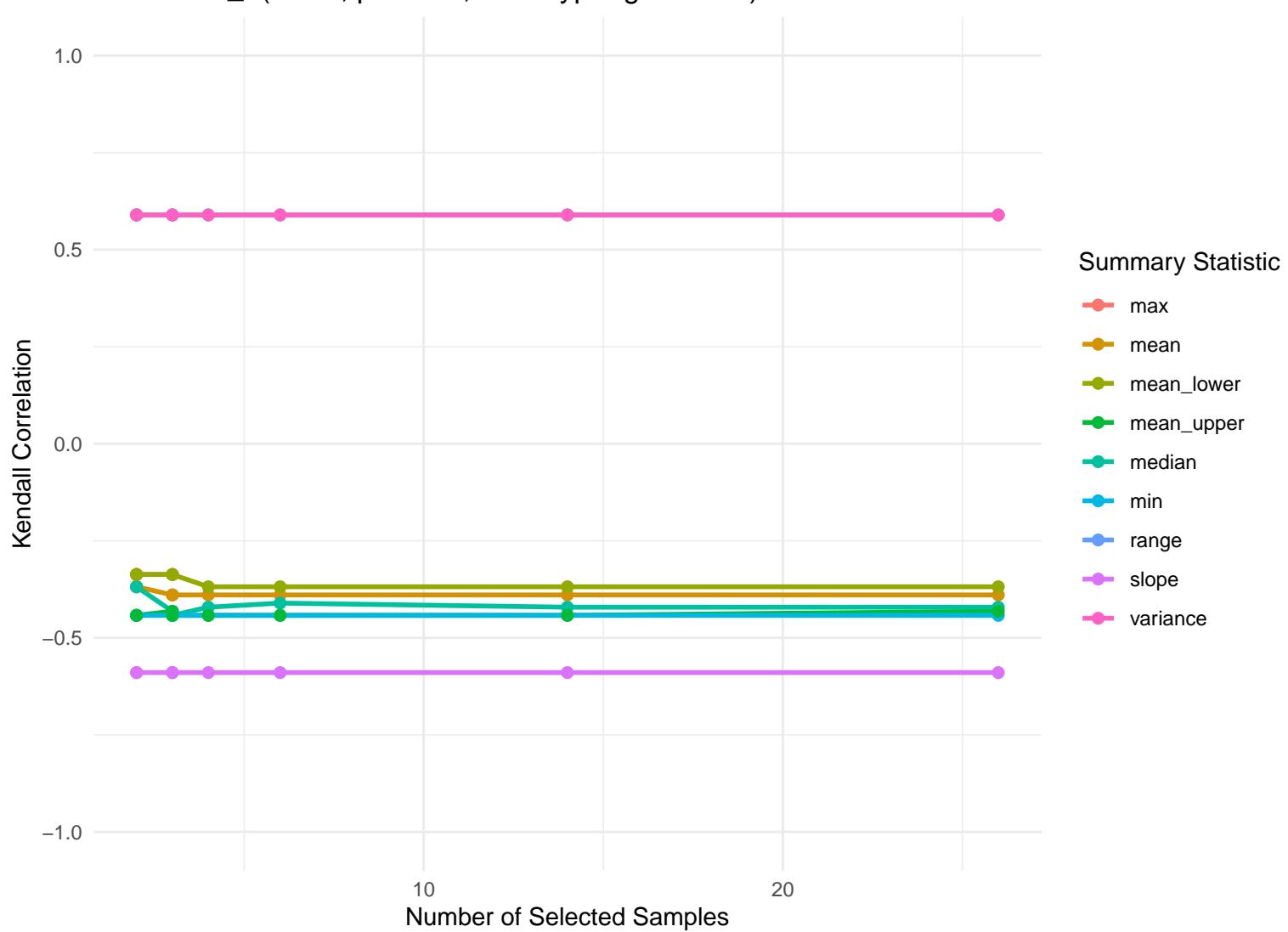
Summary Statistic

range

variance

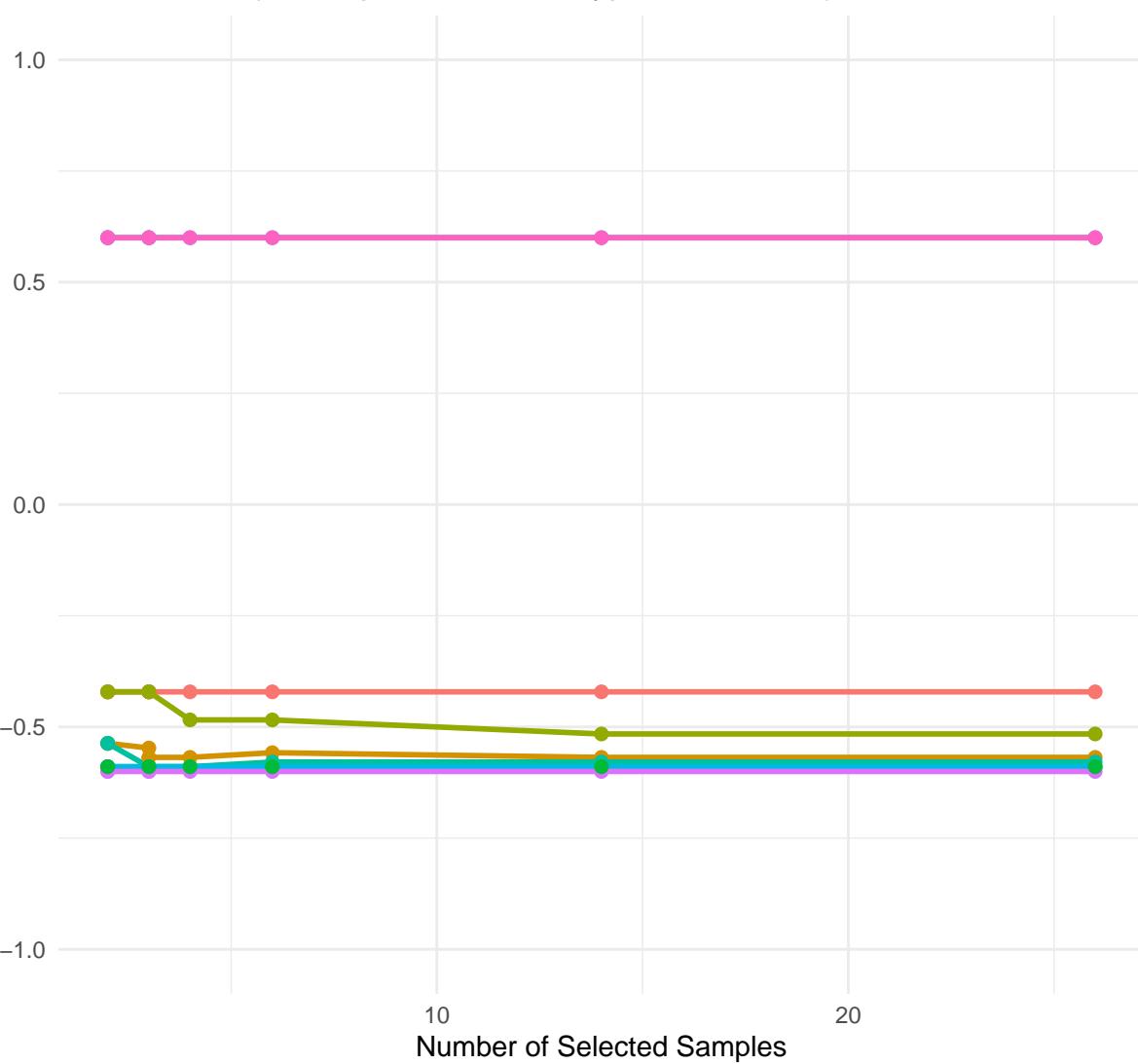


Score: CV_t (Local, partial2 , Genotype: gaussian)



Score: CV_t (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

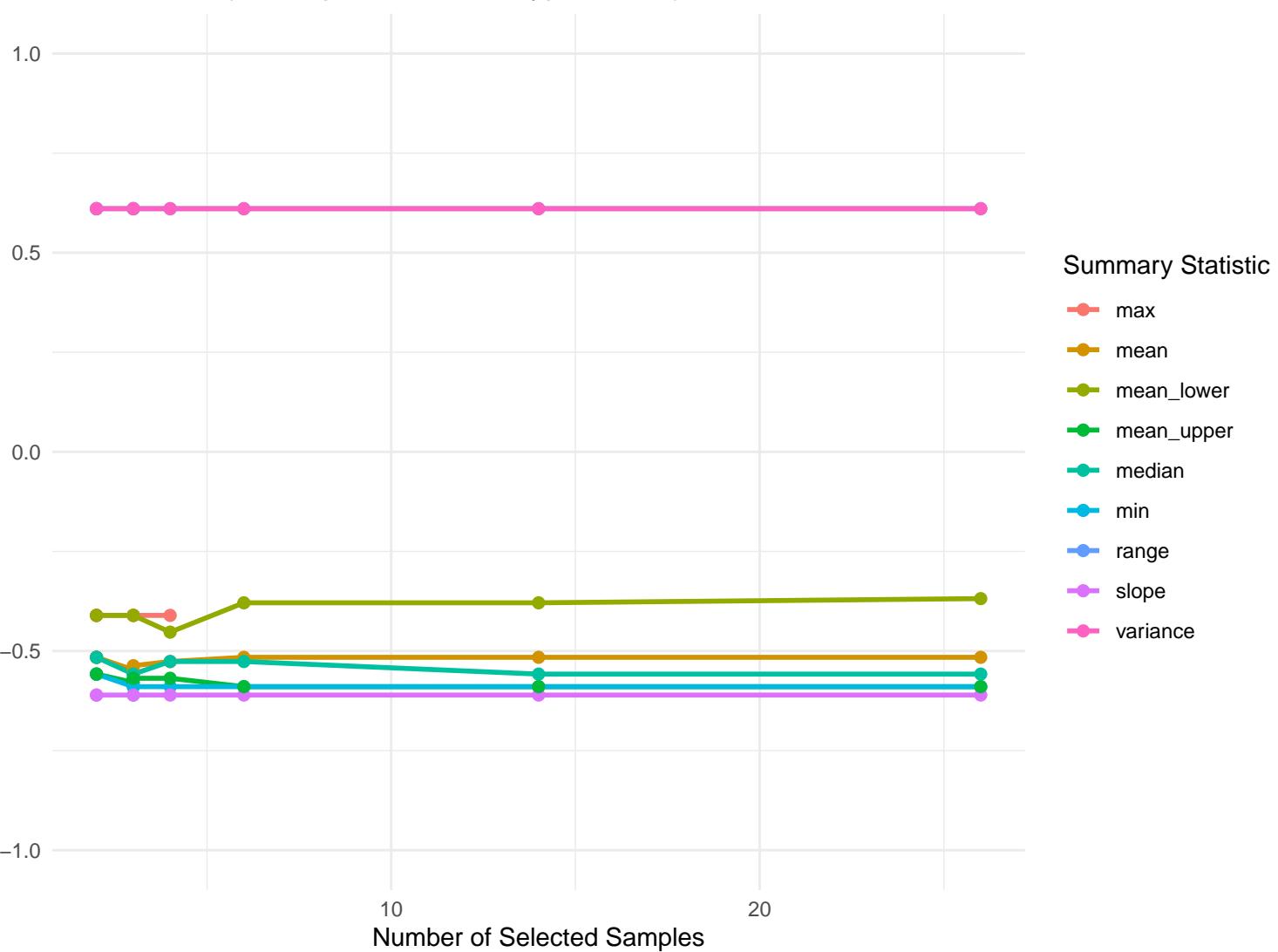


Summary Statistic

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

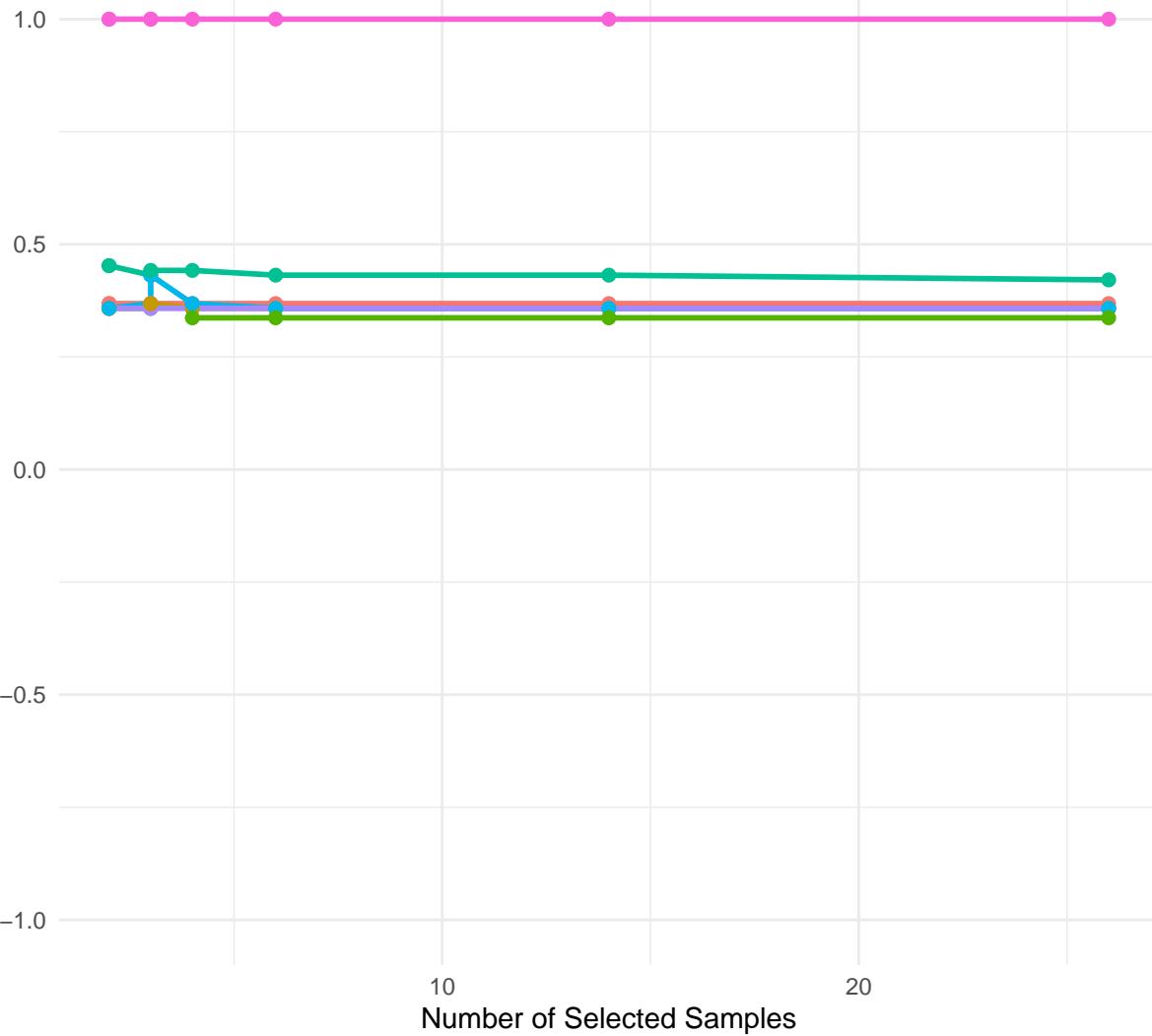
Score: CV_t (Local, partial2 , Genotype: wave)

Kendall Correlation



Score: RN (Local, partial2 , Genotype: linear)

Kendall Correlation

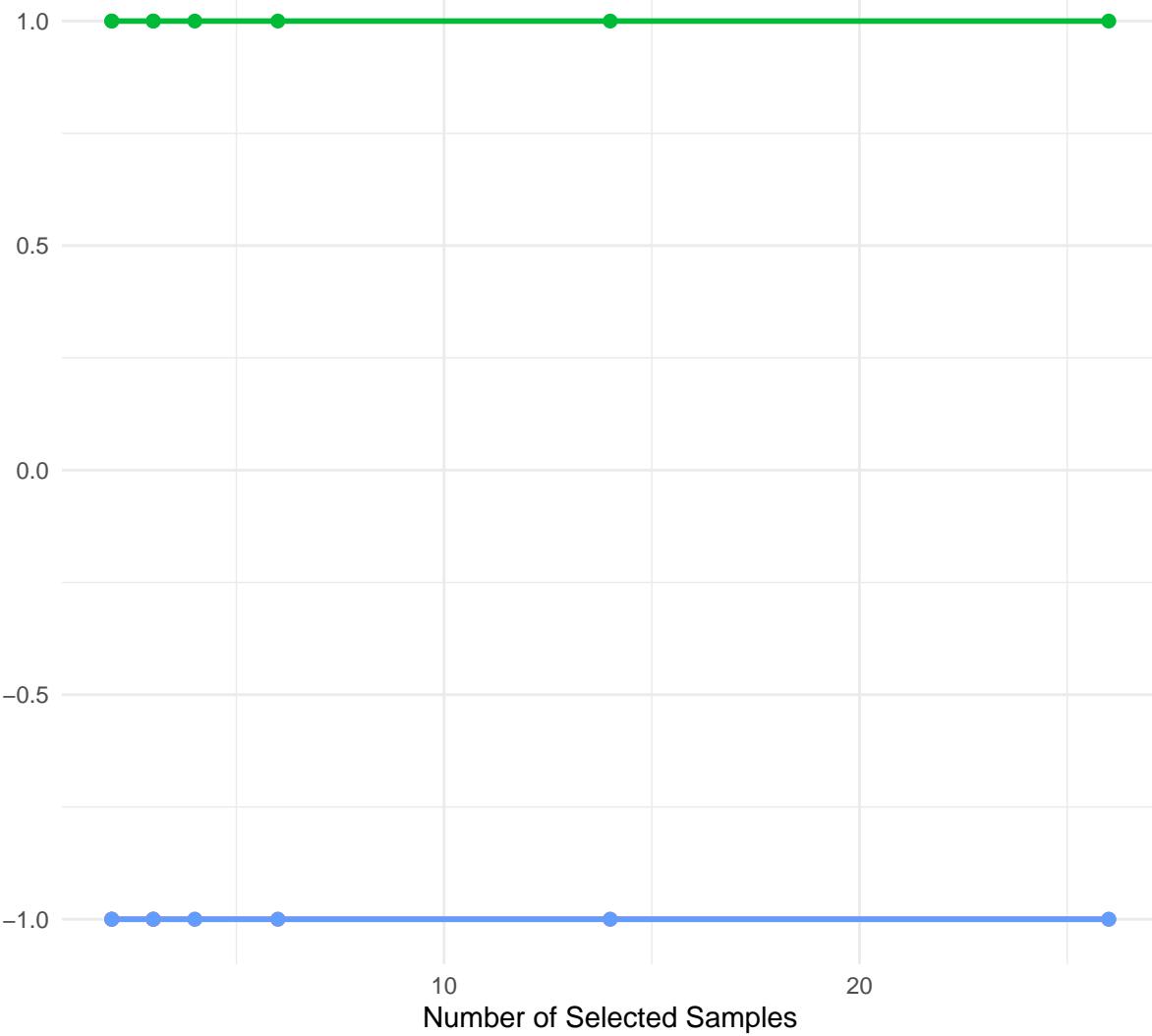


Summary Statistic

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

Score: RN (Local, partial2 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

range

slope

variance

Score: RN (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

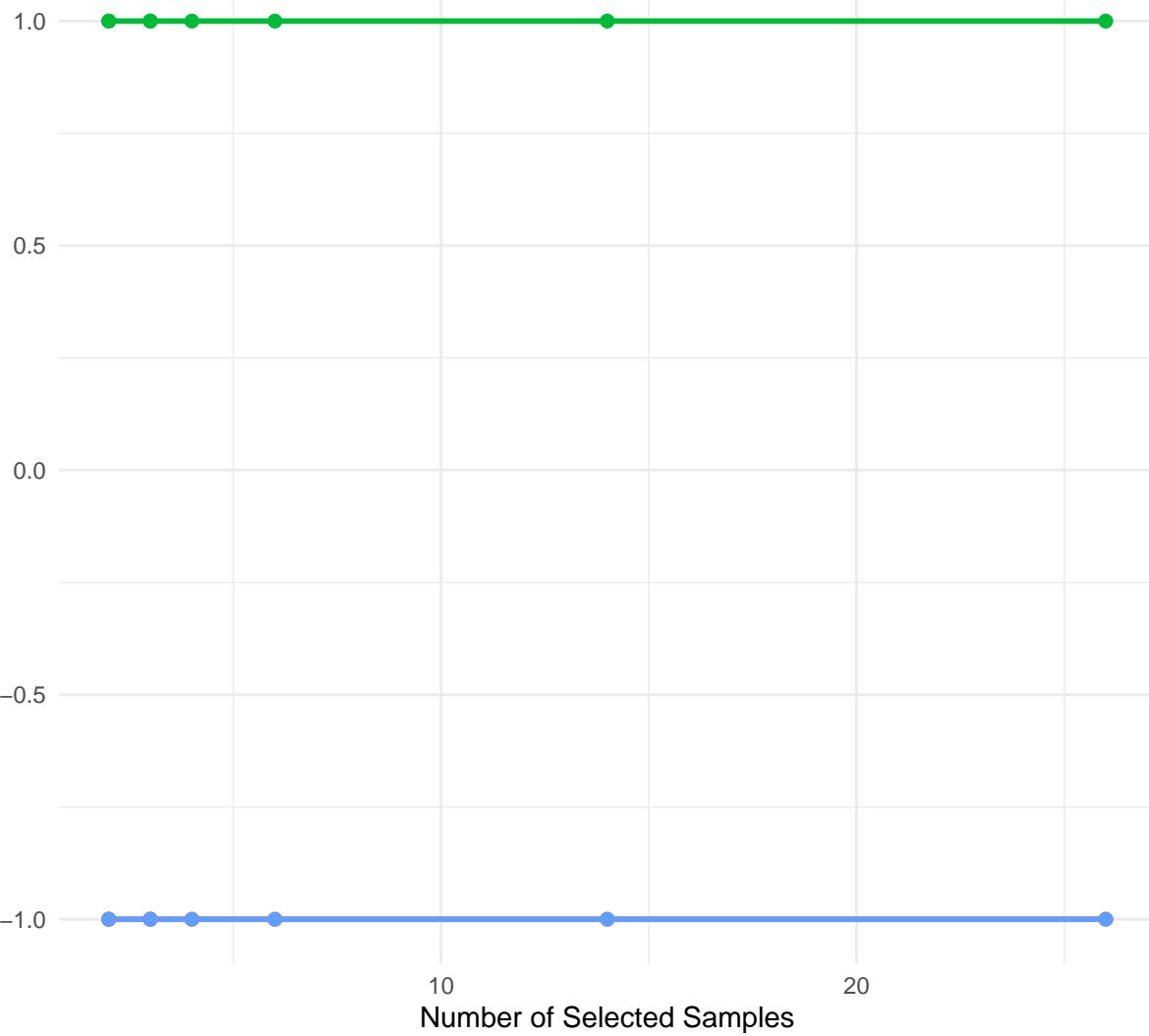
slope

variance



Score: RN (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

- range
- slope
- variance

Score: RNN (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

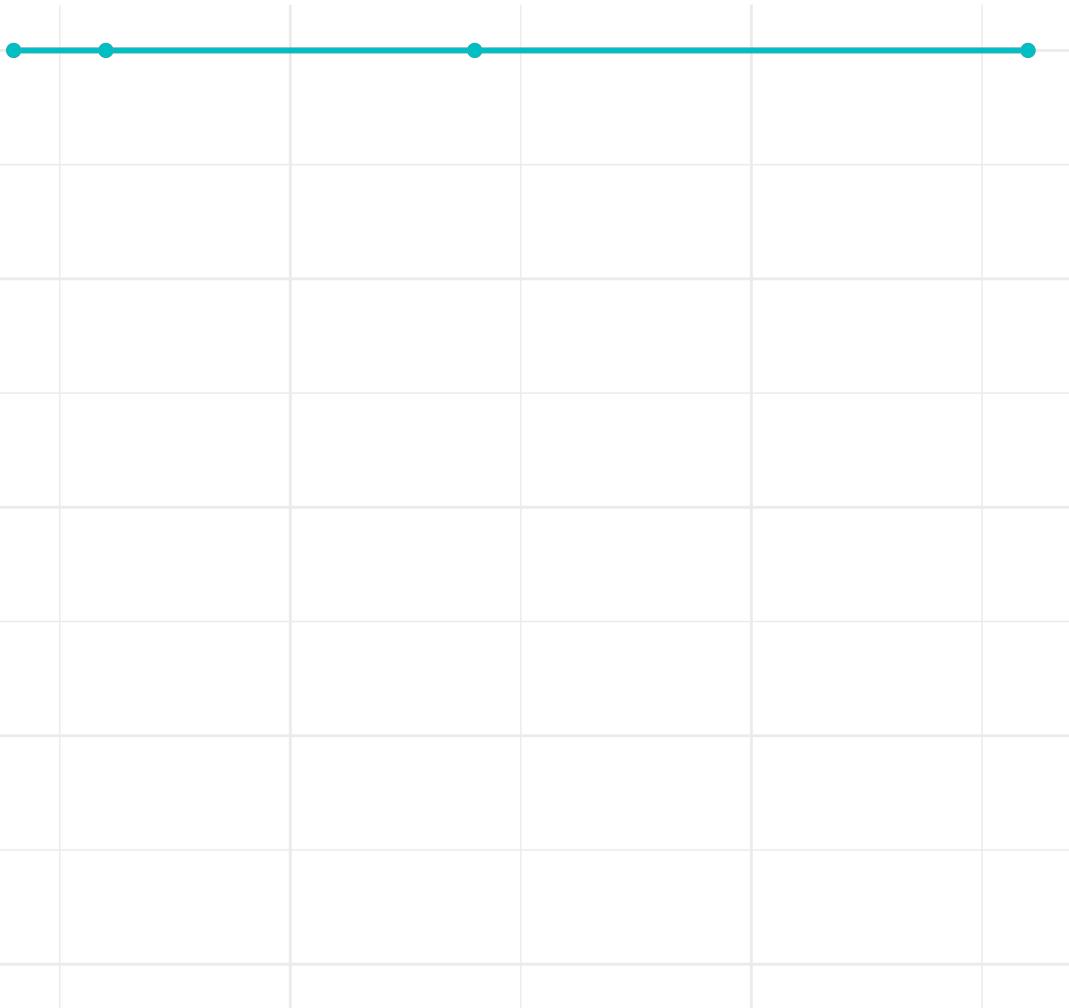
-1.0

Number of Selected Samples

Summary Statistic

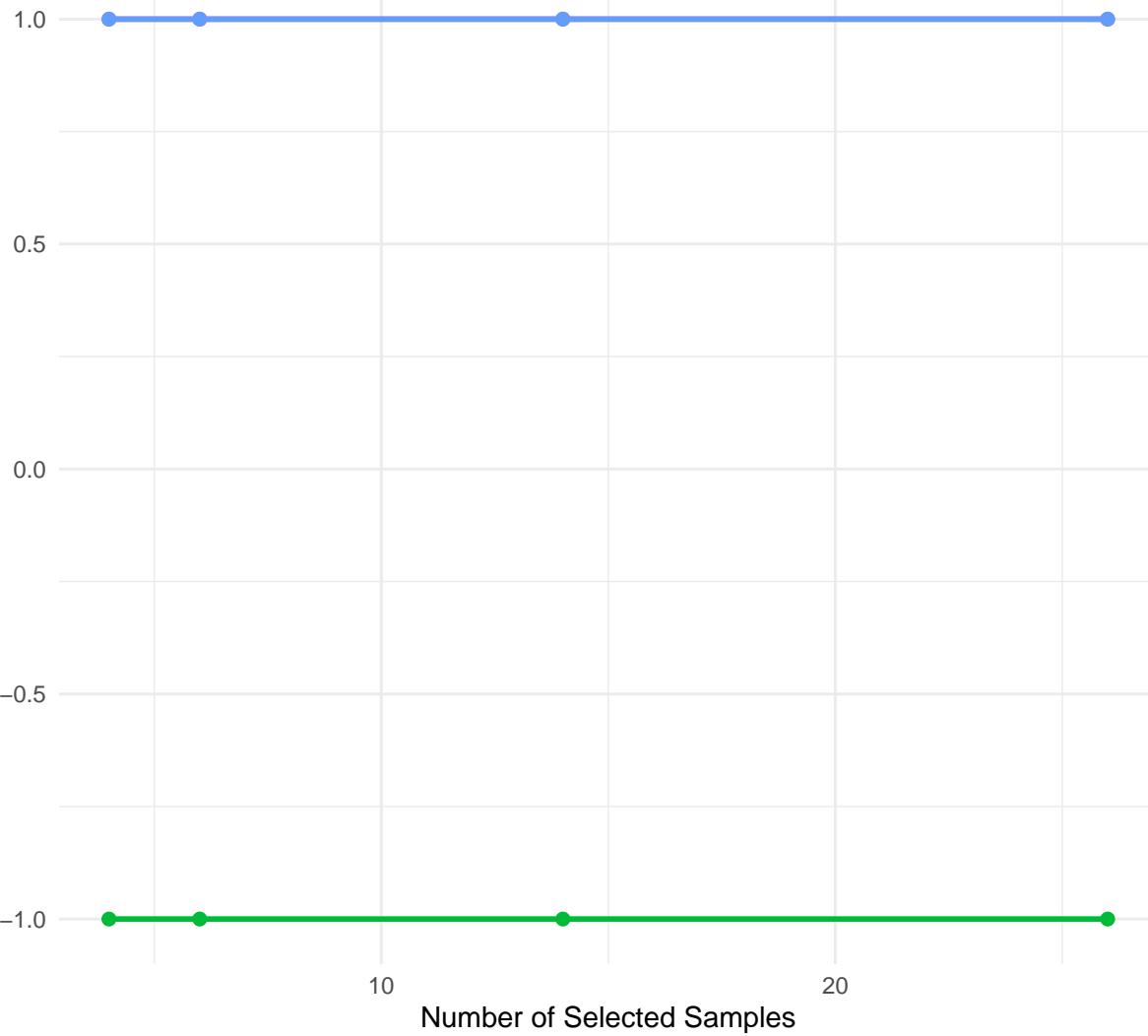
range

variance



Score: RNN (Local, partial2 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

range

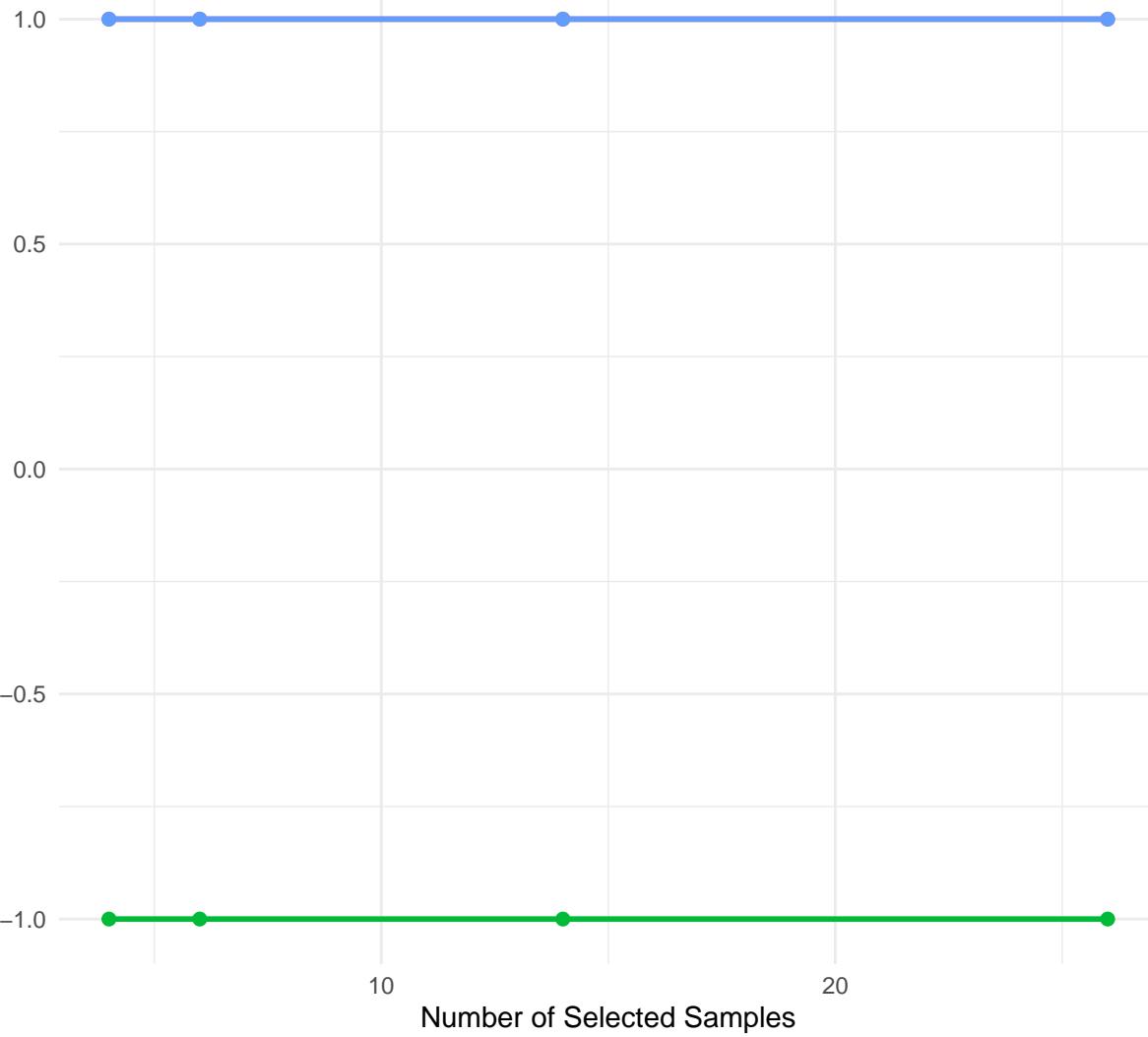
slope

variance

Number of Selected Samples

Score: RNN (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

range

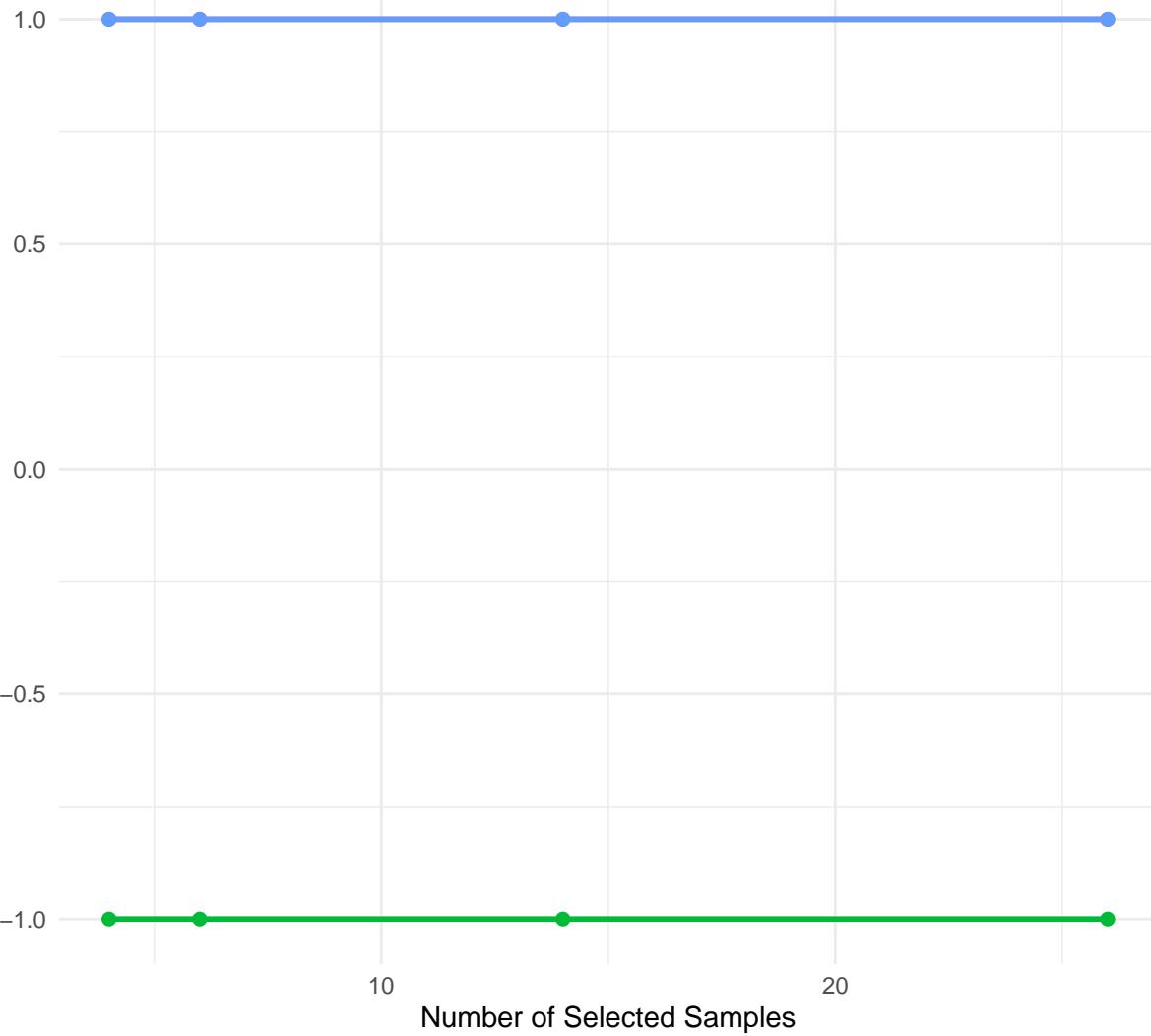
slope

variance

Number of Selected Samples

Score: RNN (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

range

slope

variance

Number of Selected Samples

Score: D_slope (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

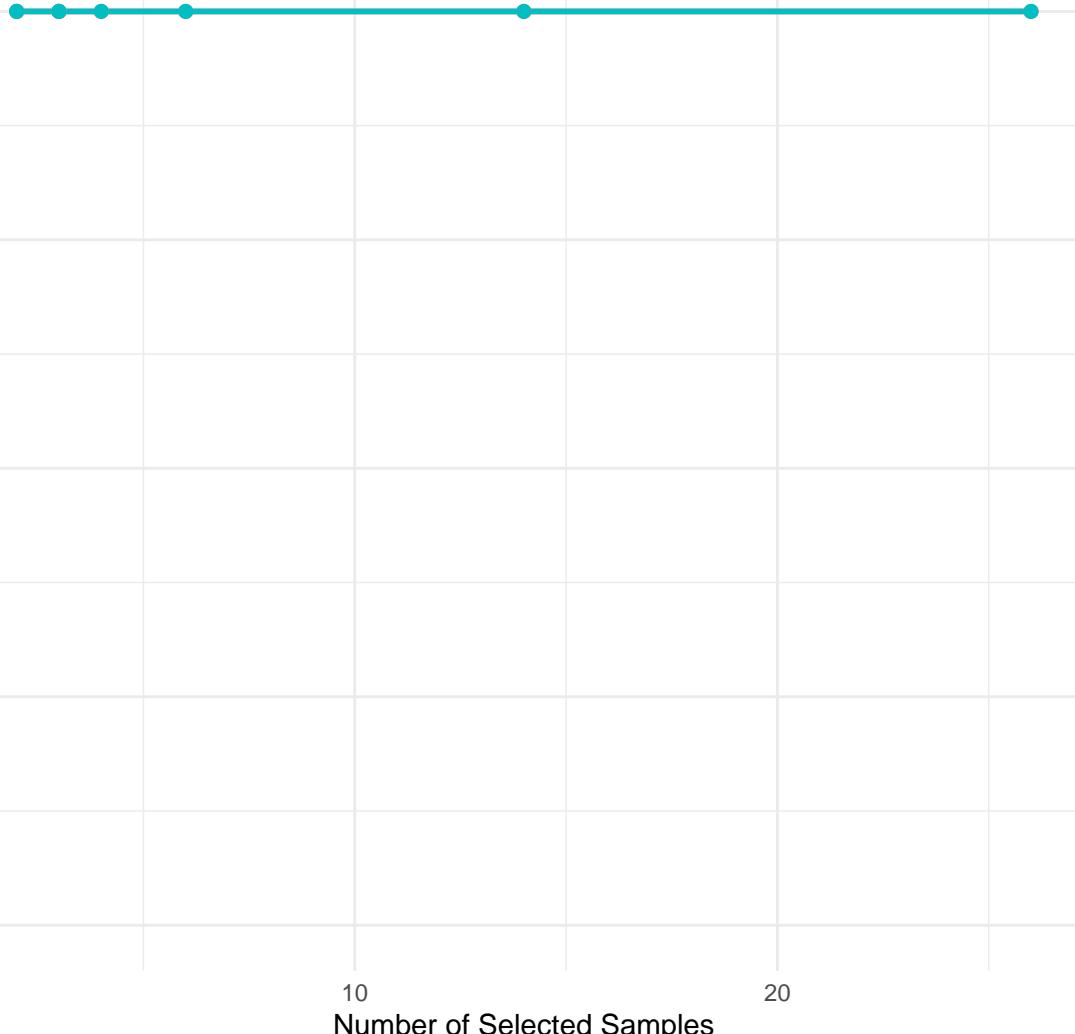
-1.0

Number of Selected Samples

Summary Statistic

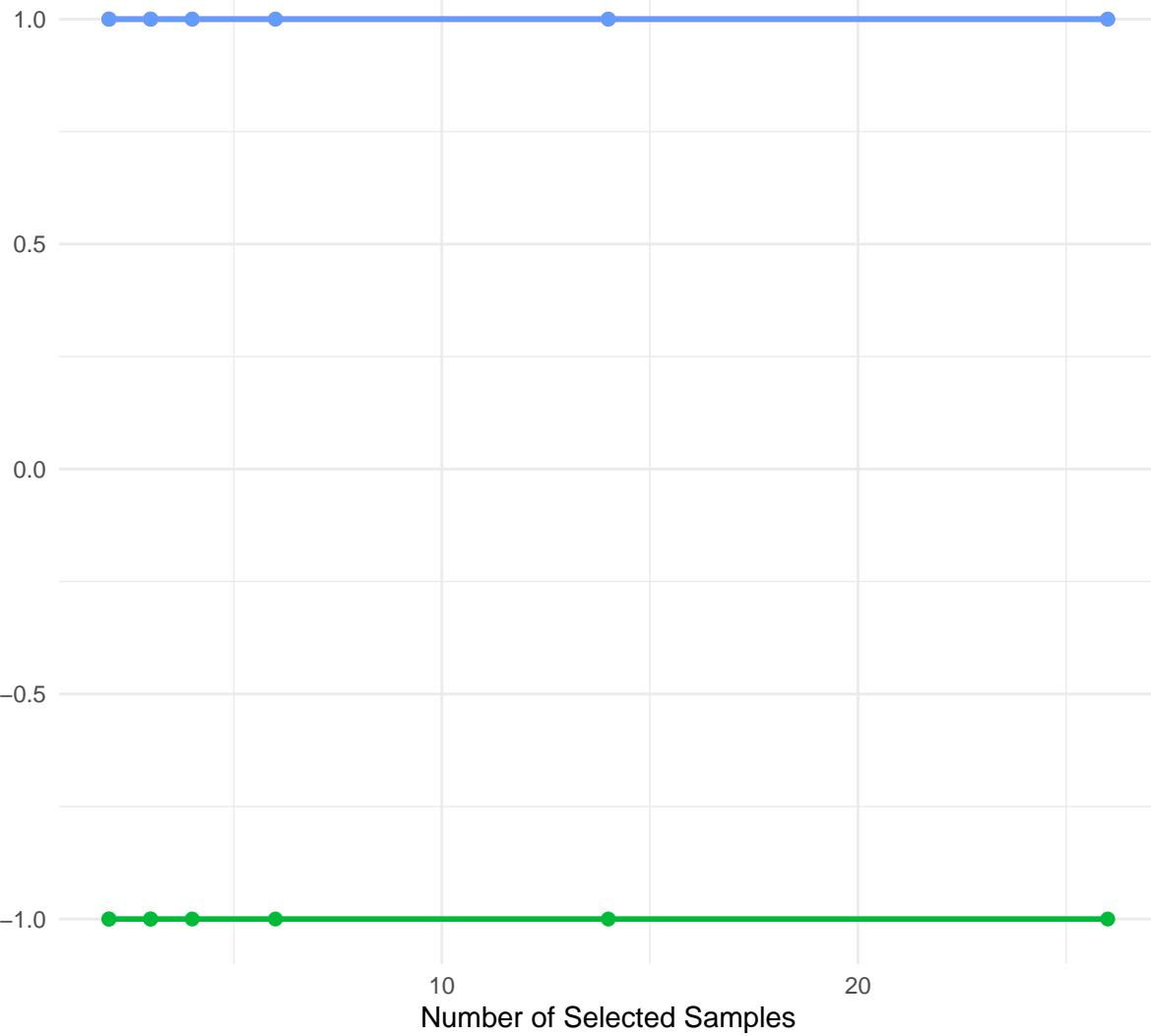
range

variance



Score: D_slope (Local, partial2 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

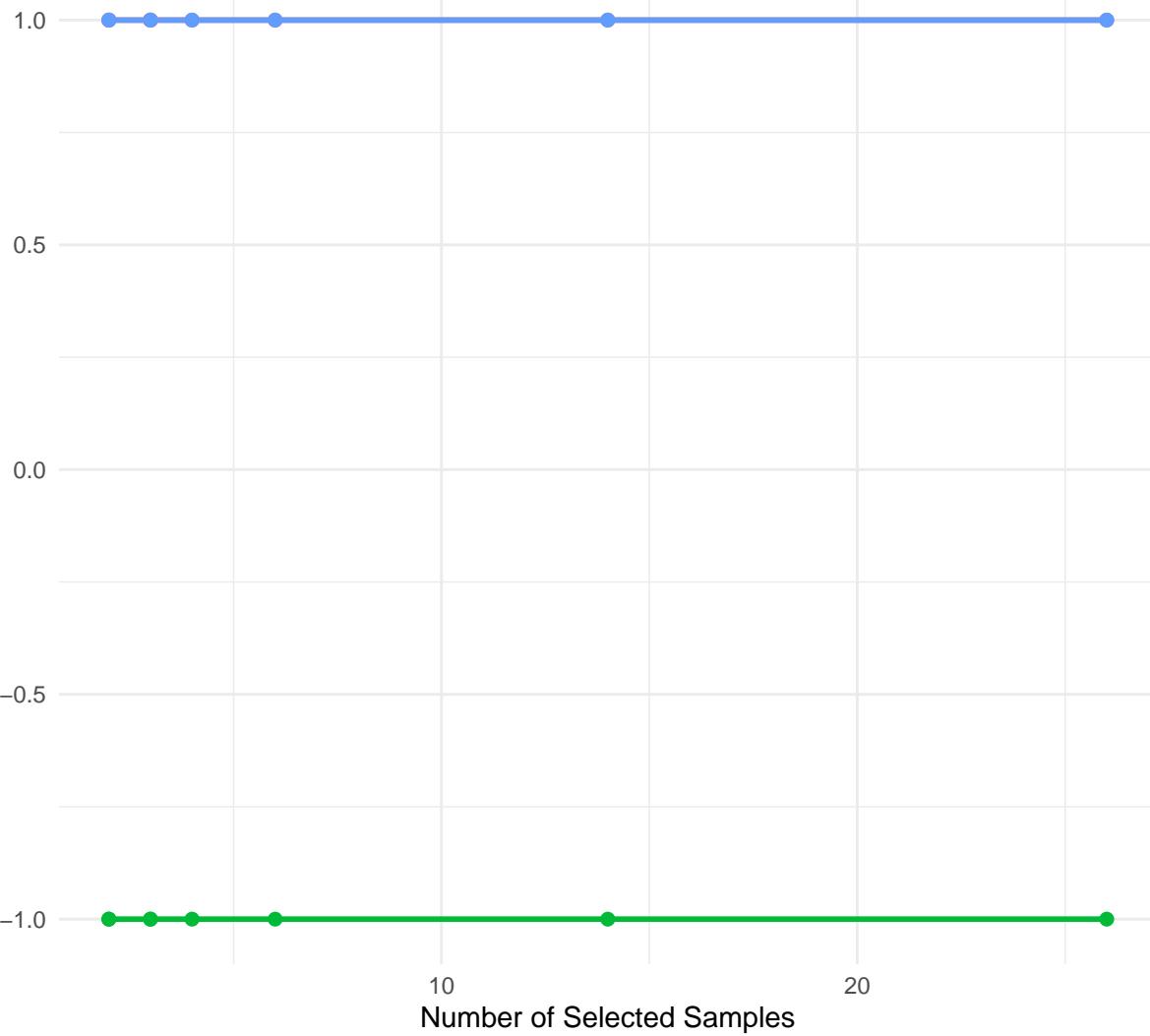
range

slope

variance

Score: D_slope (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

range

slope

variance

Score: D_slope (Local, partial2 , Genotype: wave)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Summary Statistic

range

slope

variance

Number of Selected Samples

Score: RC (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

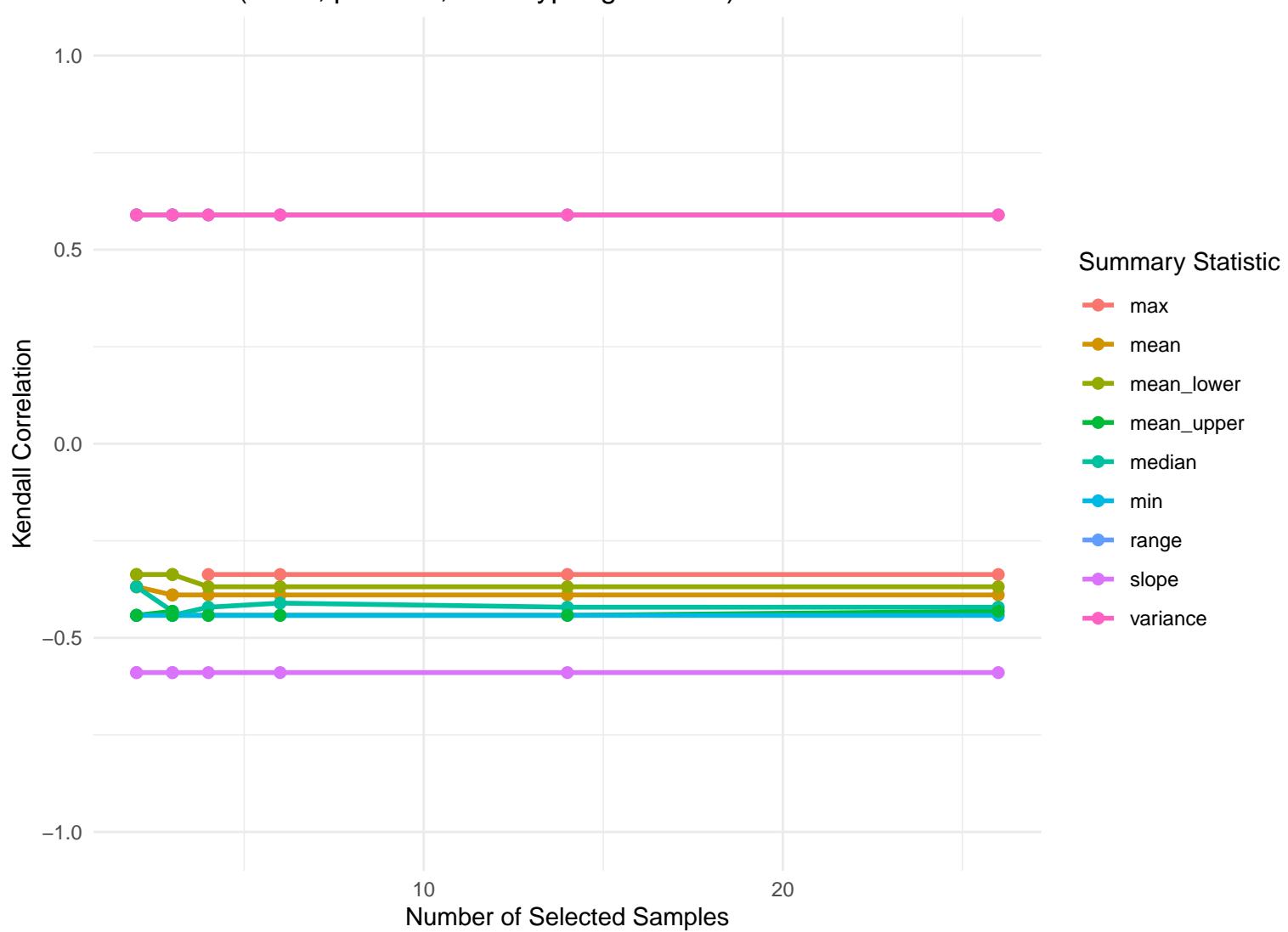
Summary Statistic

range

variance

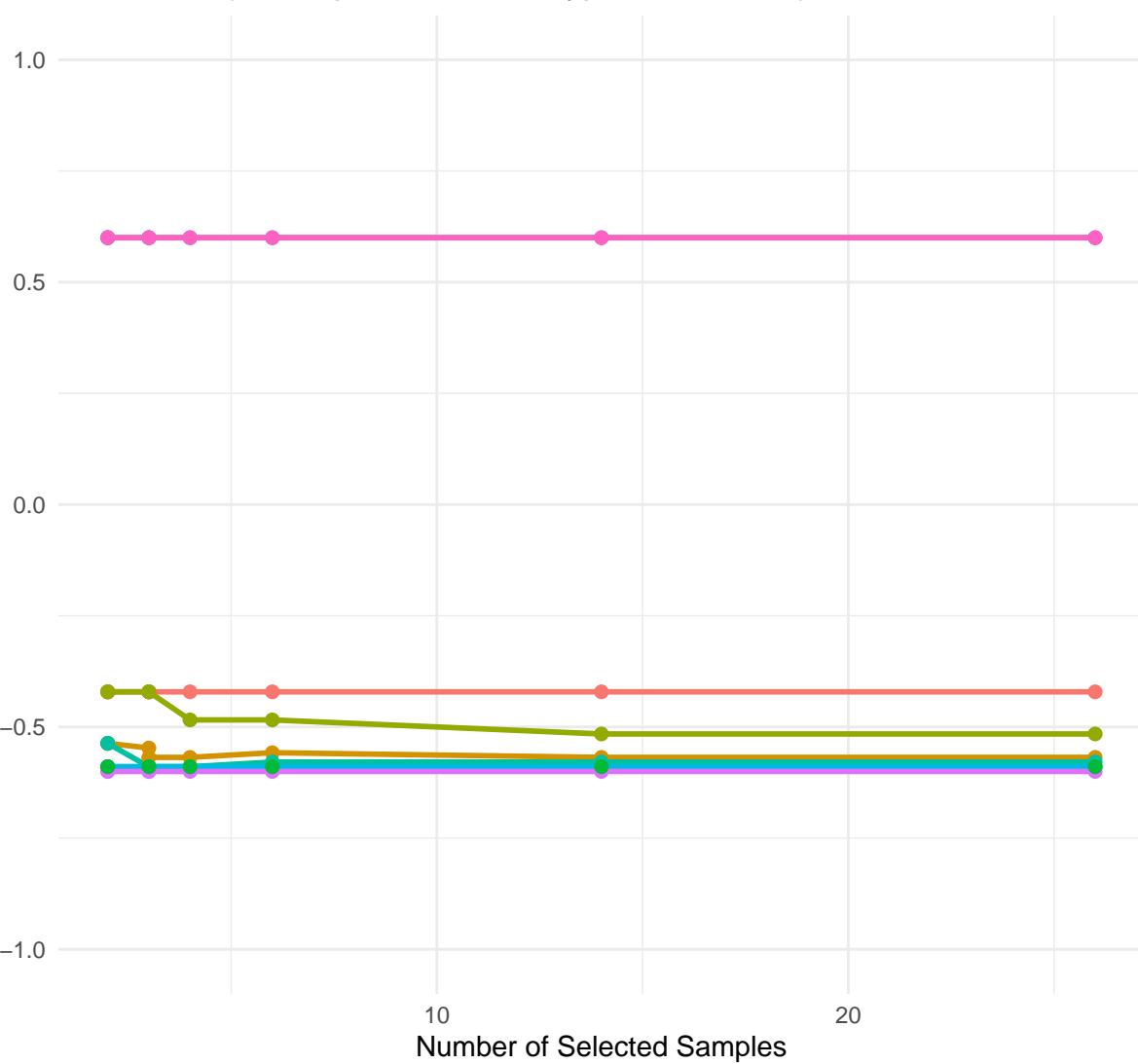


Score: RC (Local, partial2 , Genotype: gaussian)



Score: RC (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

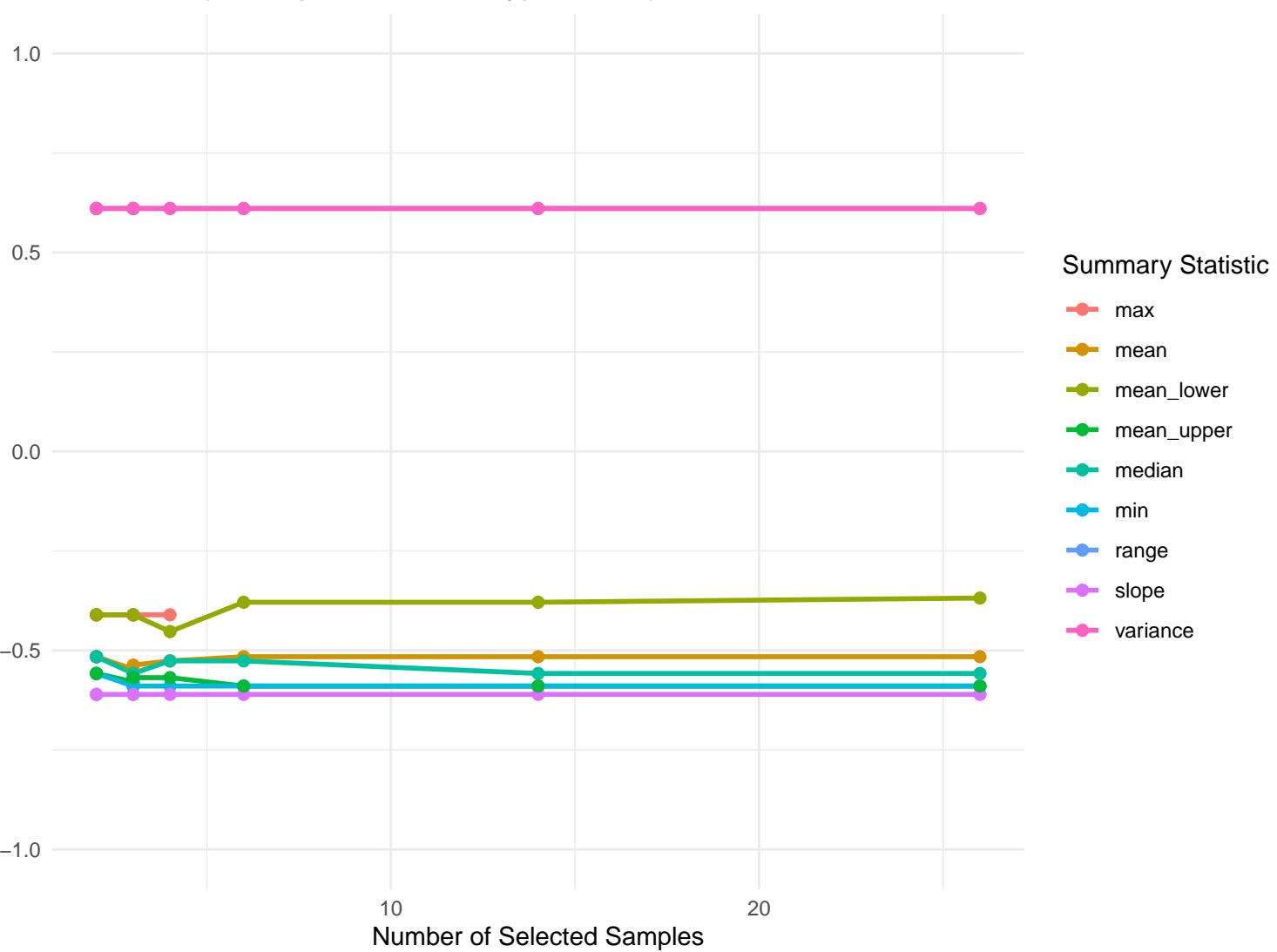


Summary Statistic

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

Score: RC (Local, partial2 , Genotype: wave)

Kendall Correlation



Score: gPi (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

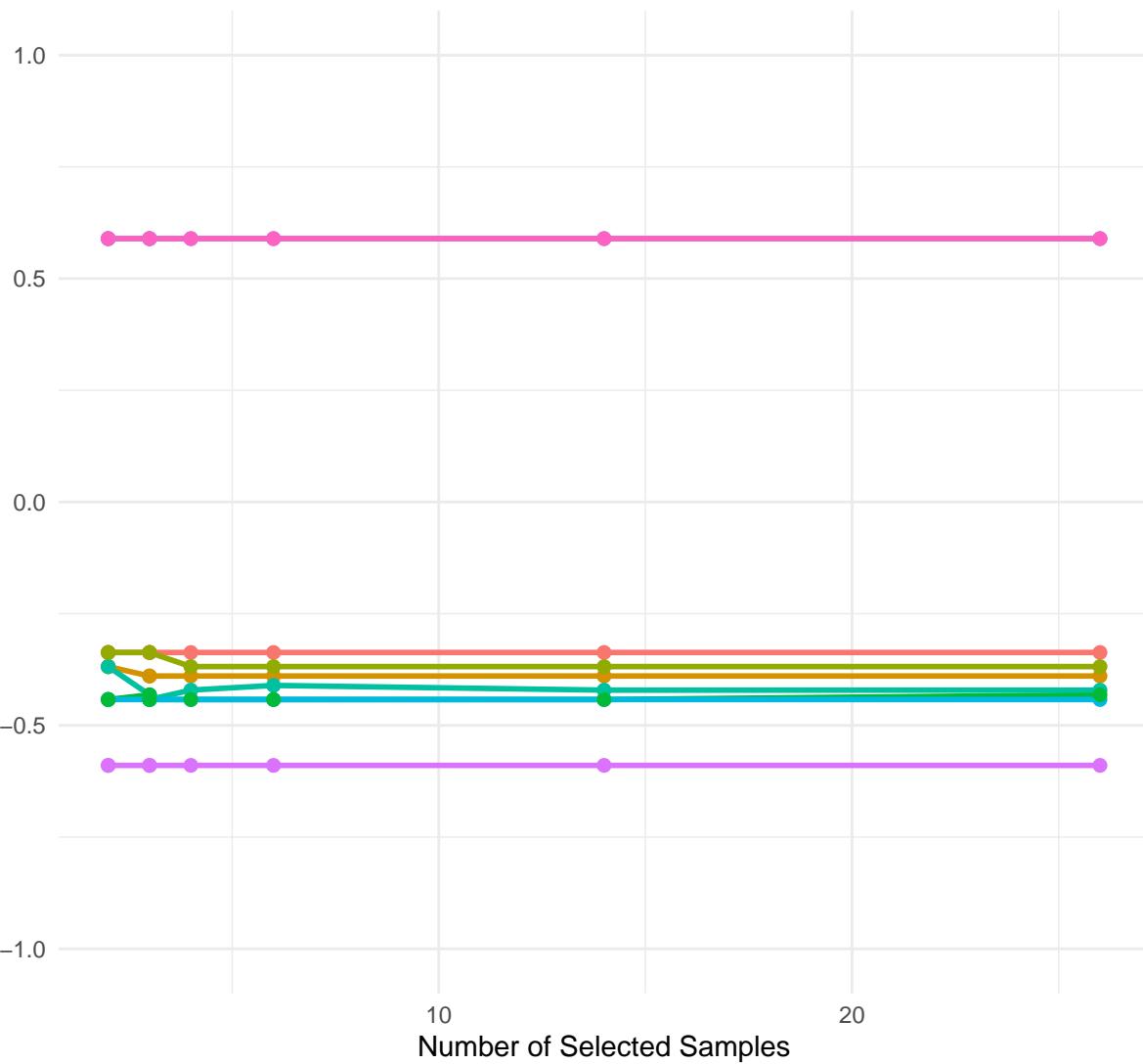
range

variance



Score: gPi (Local, partial2 , Genotype: gaussian)

Kendall Correlation

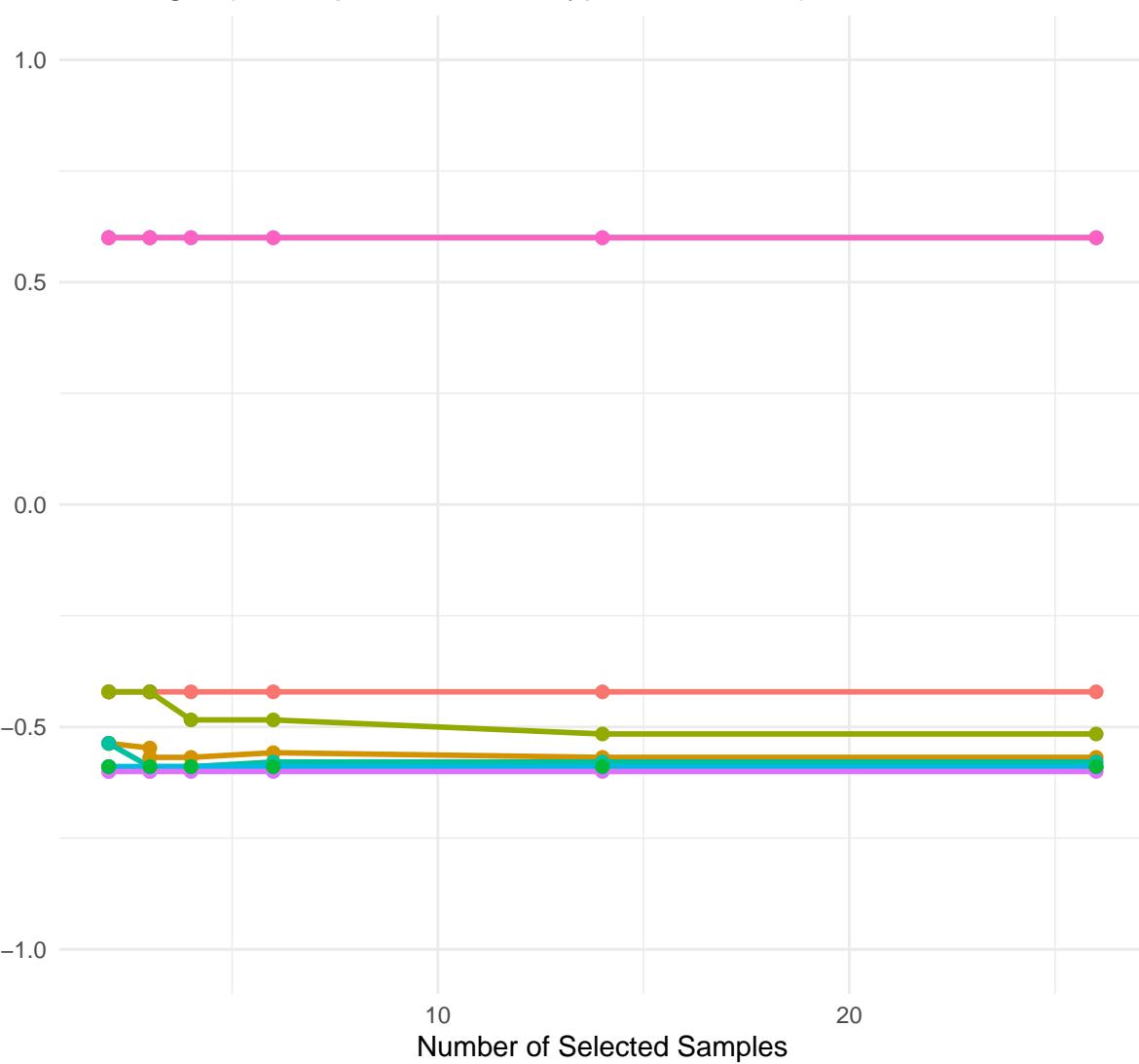


Summary Statistic

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

Score: gPi (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

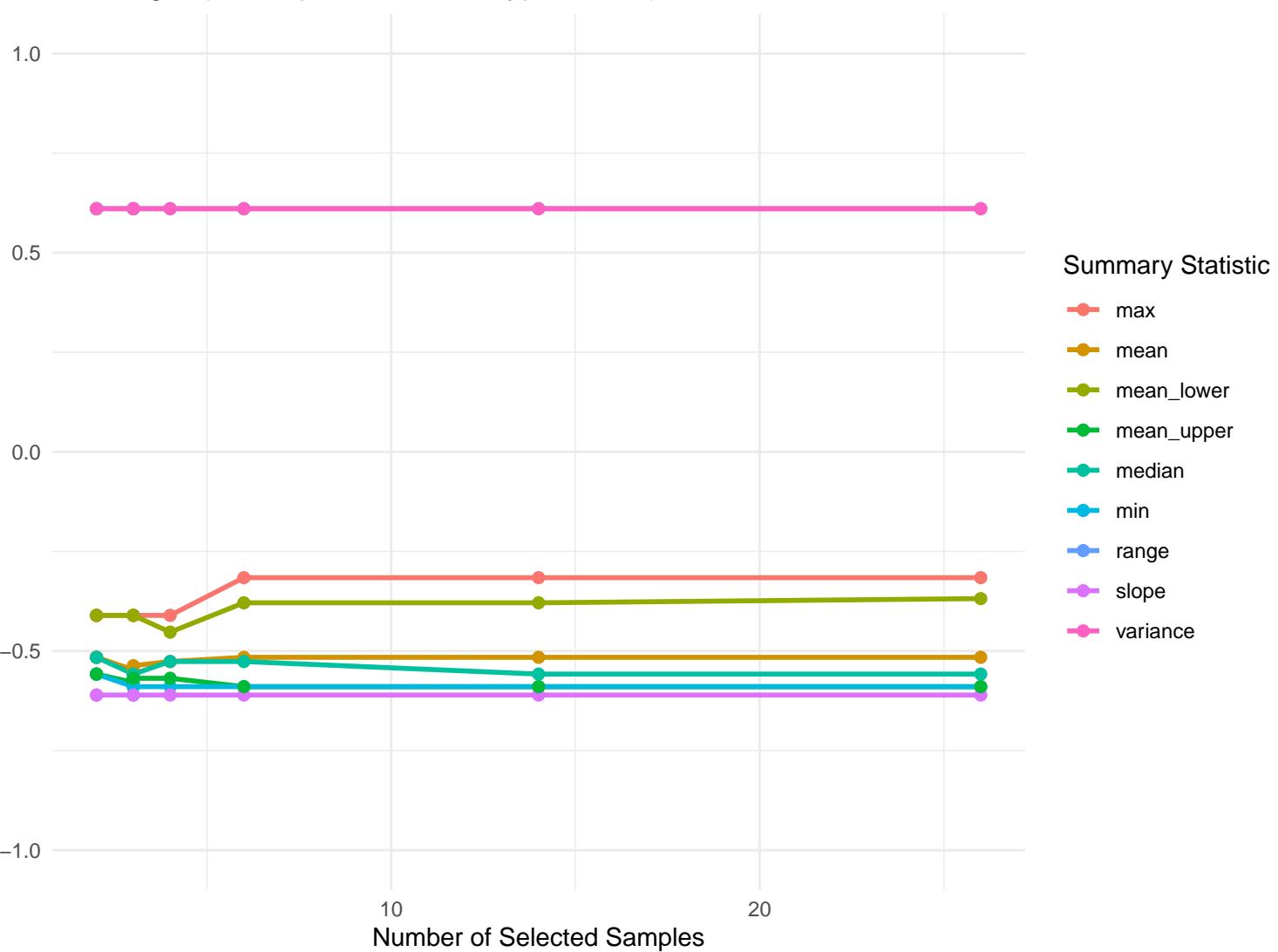


Summary Statistic

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

Score: gPi (Local, partial2 , Genotype: wave)

Kendall Correlation



Score: PPF (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

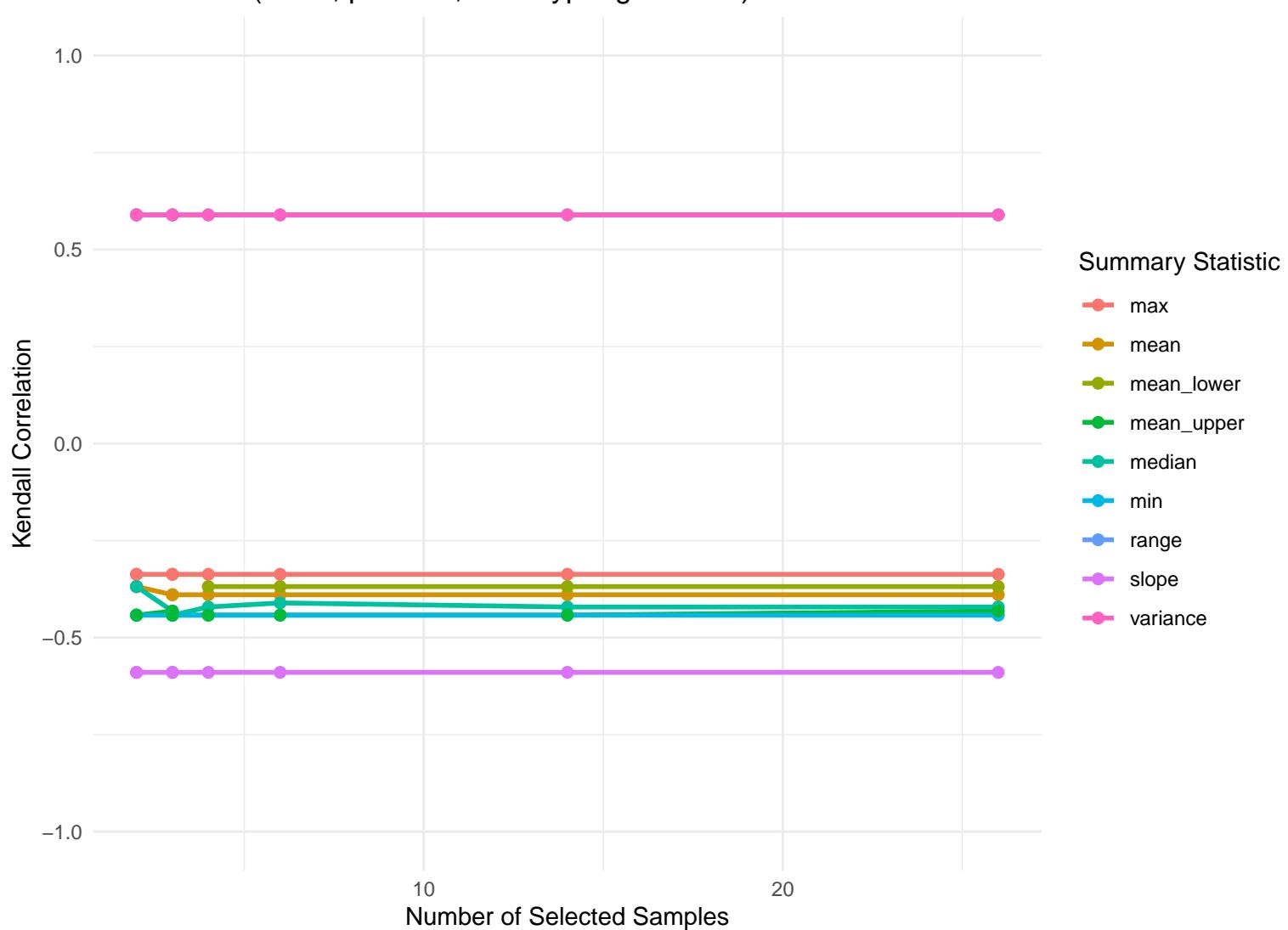
Summary Statistic

range

variance

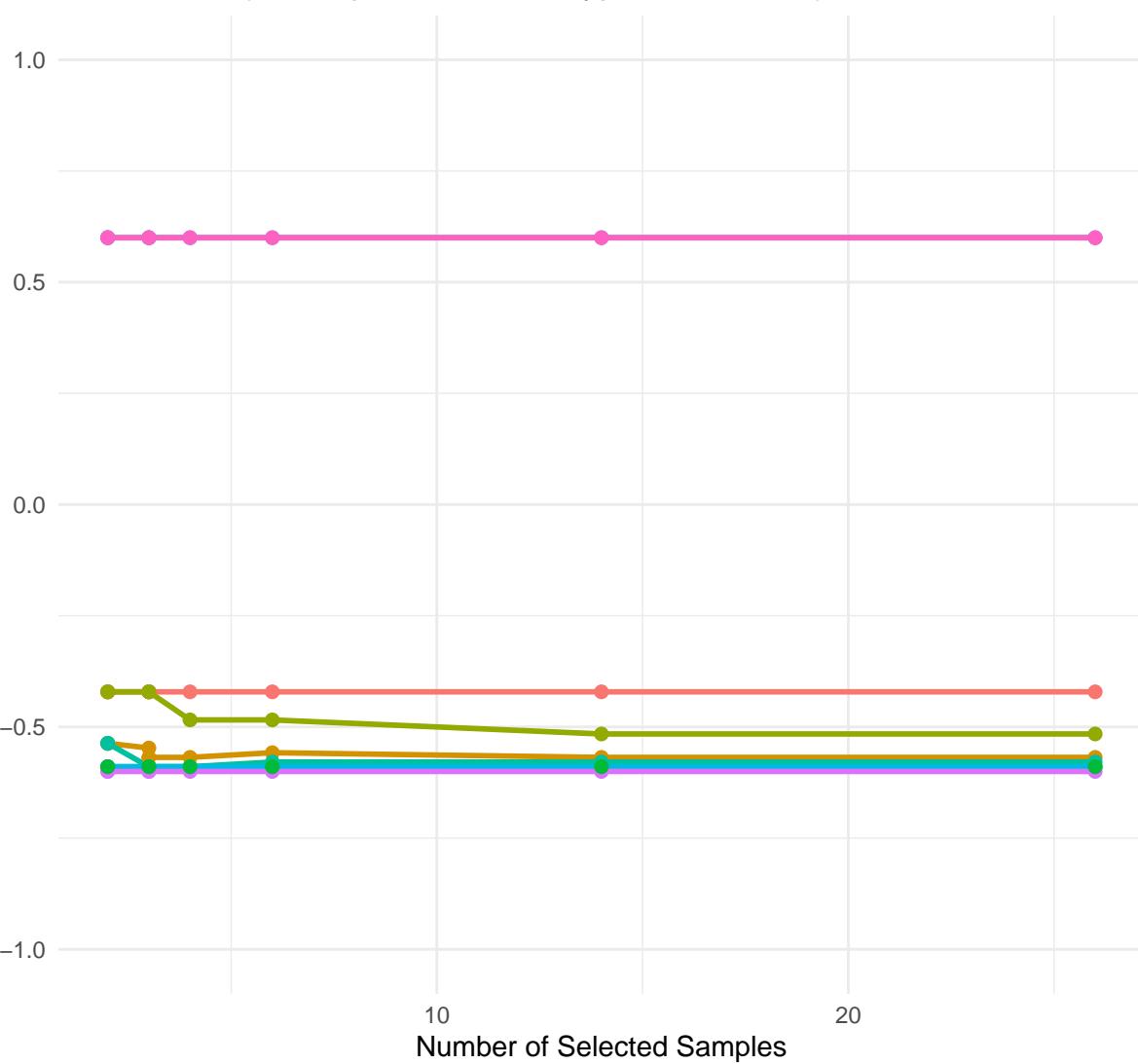


Score: PPF (Local, partial2 , Genotype: gaussian)



Score: PPF (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

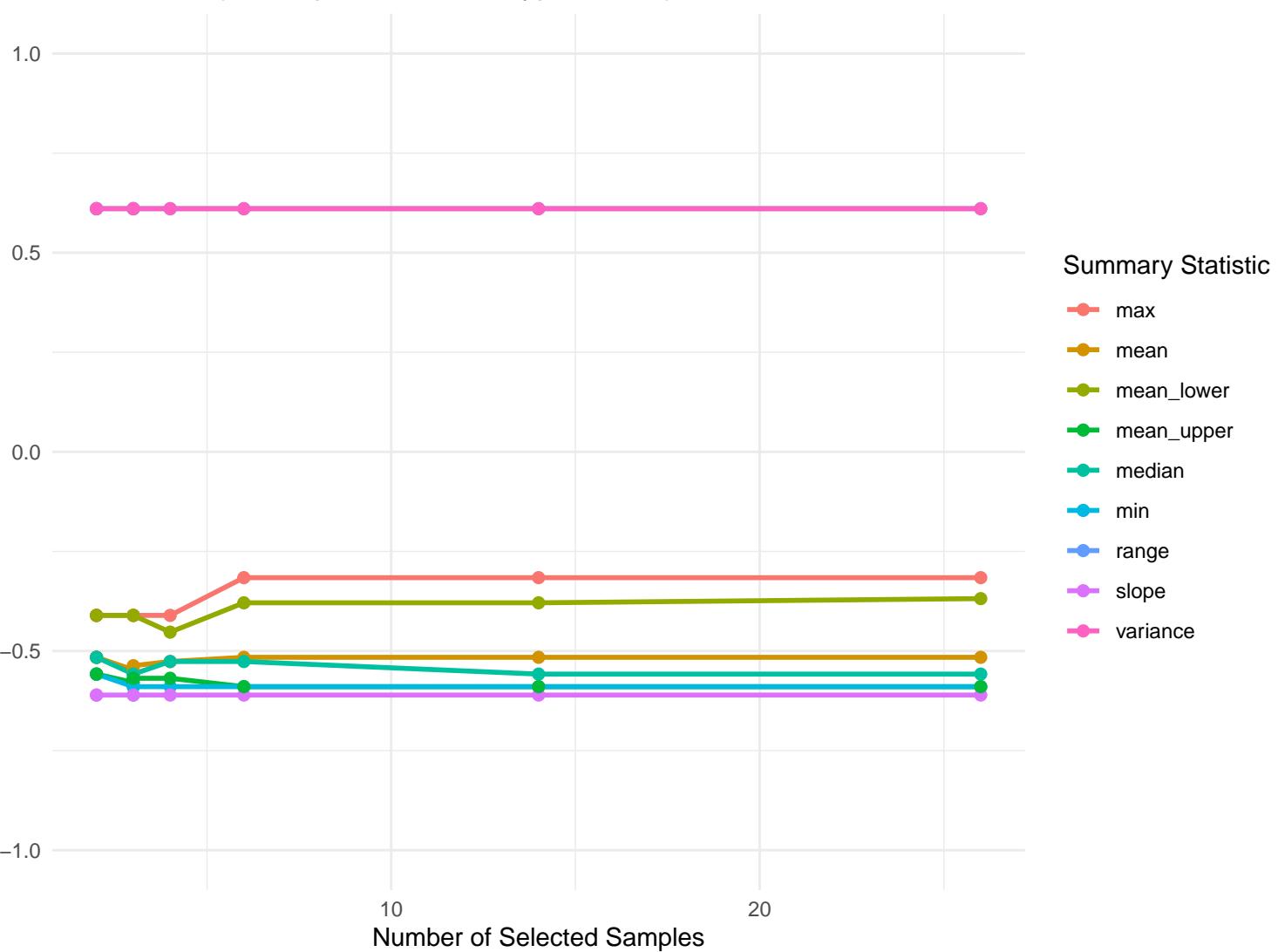


Summary Statistic

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

Score: PPF (Local, partial2 , Genotype: wave)

Kendall Correlation



Score: PPi (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

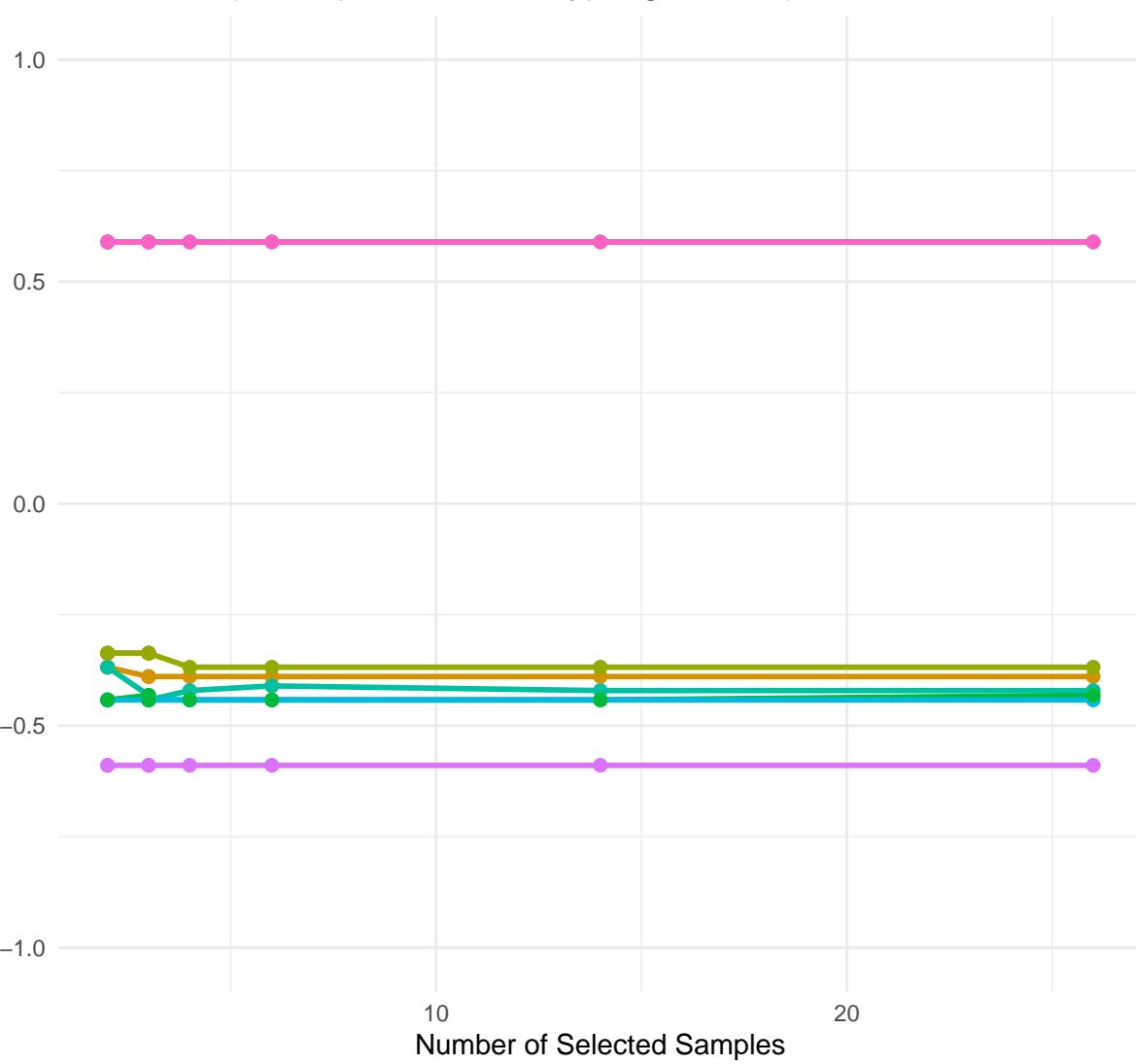
range

variance



Score: PPi (Local, partial2 , Genotype: gaussian)

Kendall Correlation

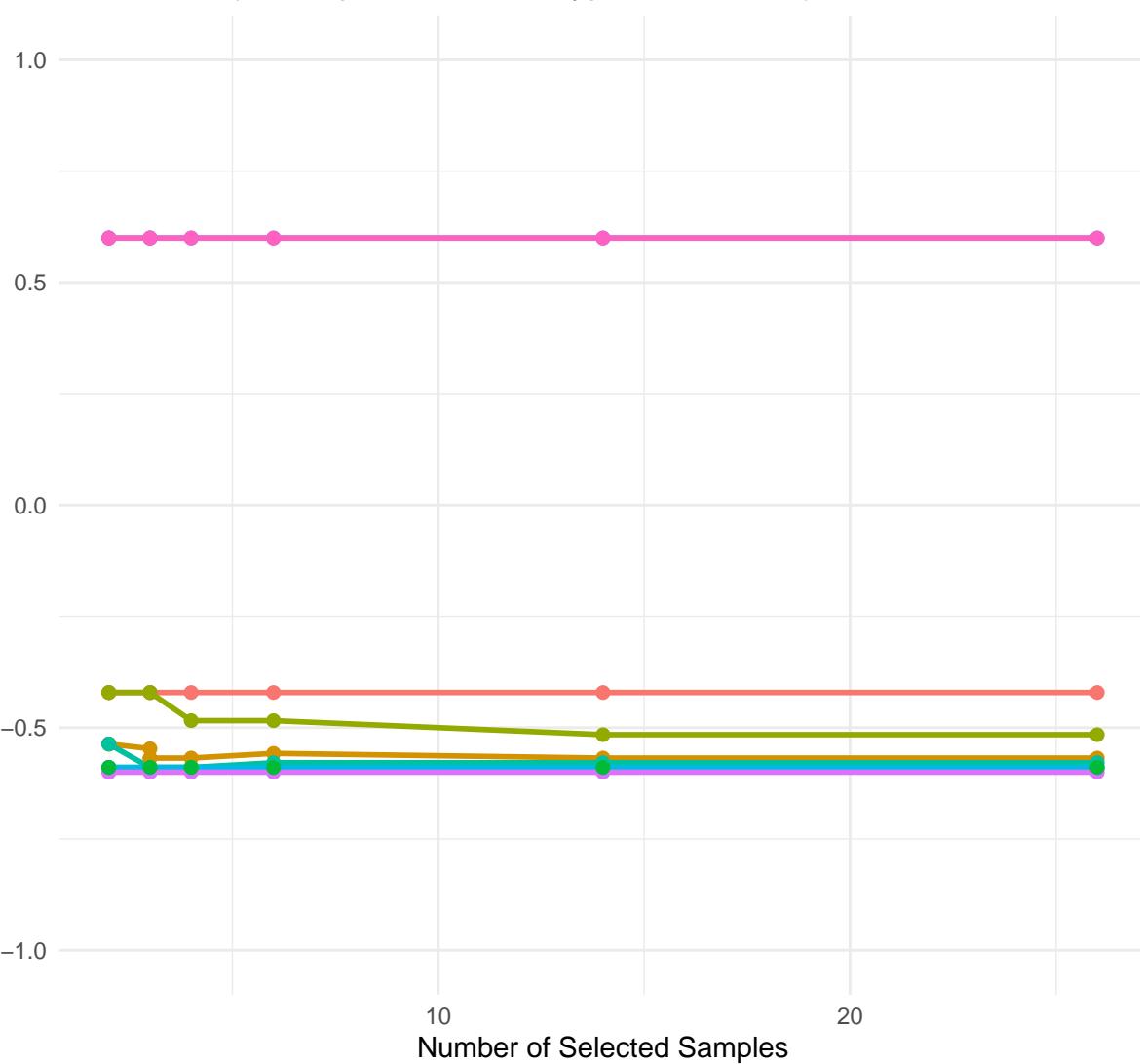


Summary Statistic

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

Score: PPi (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

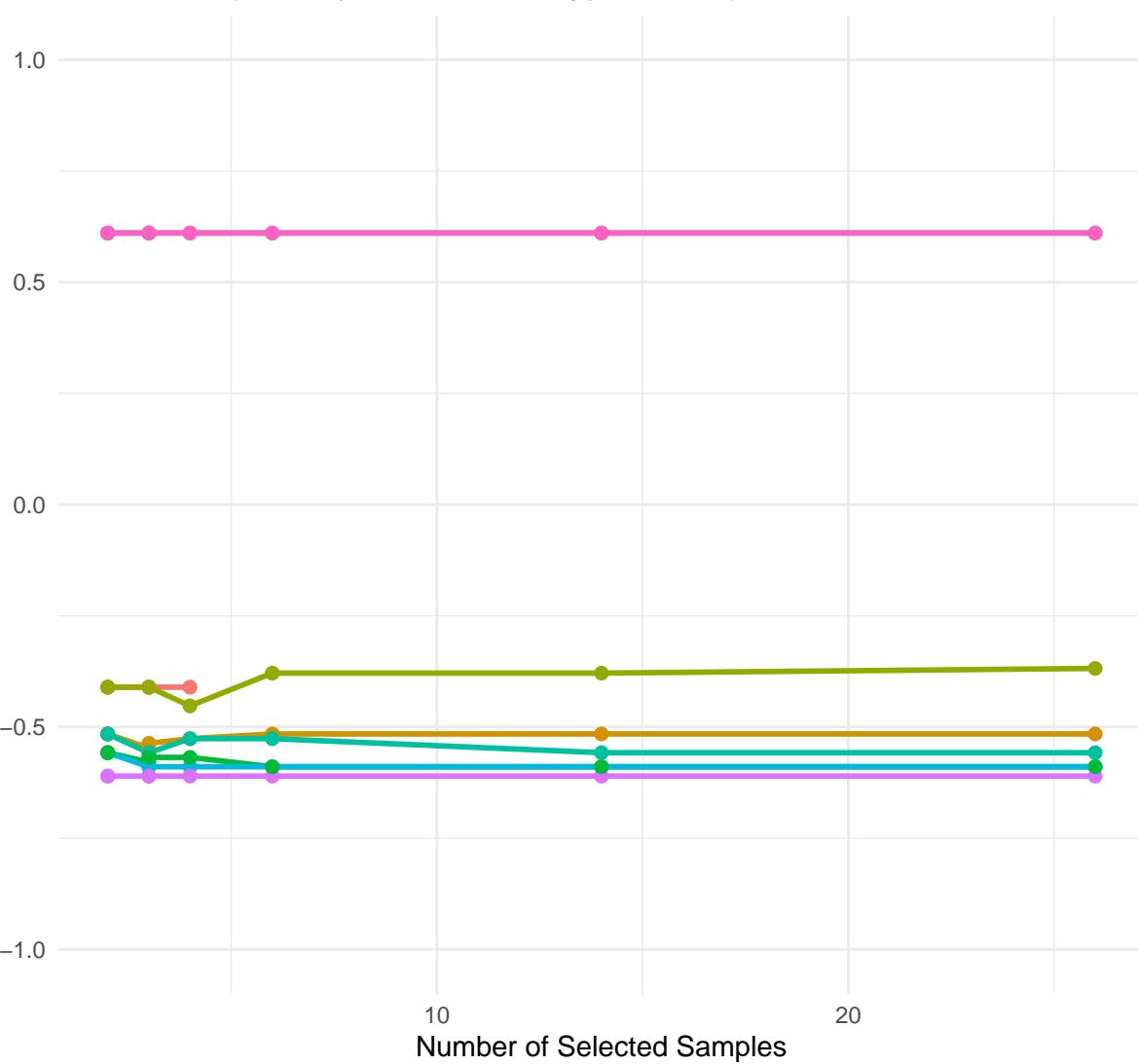


Summary Statistic

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

Score: PPi (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: Plmd (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

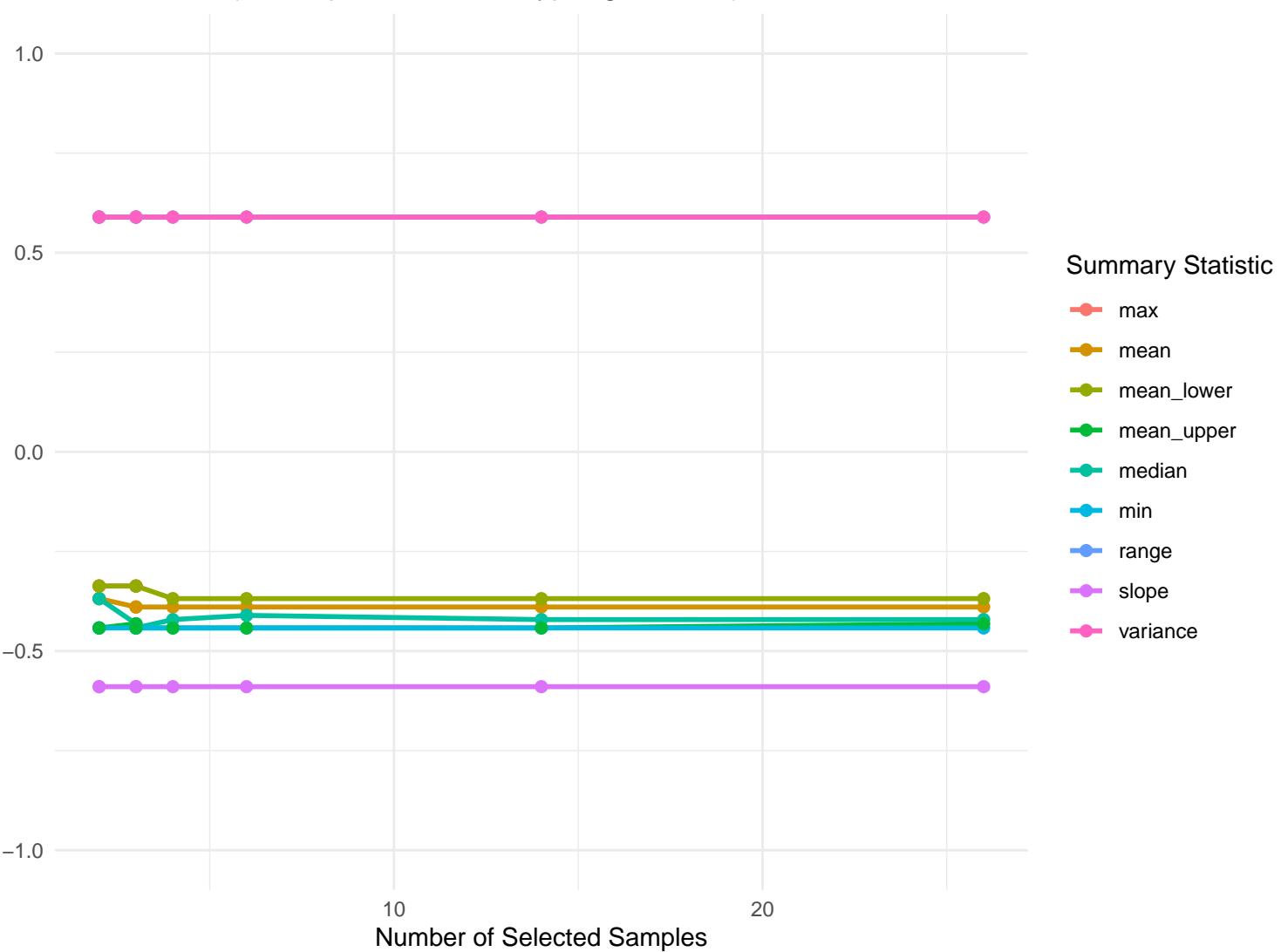
range

variance



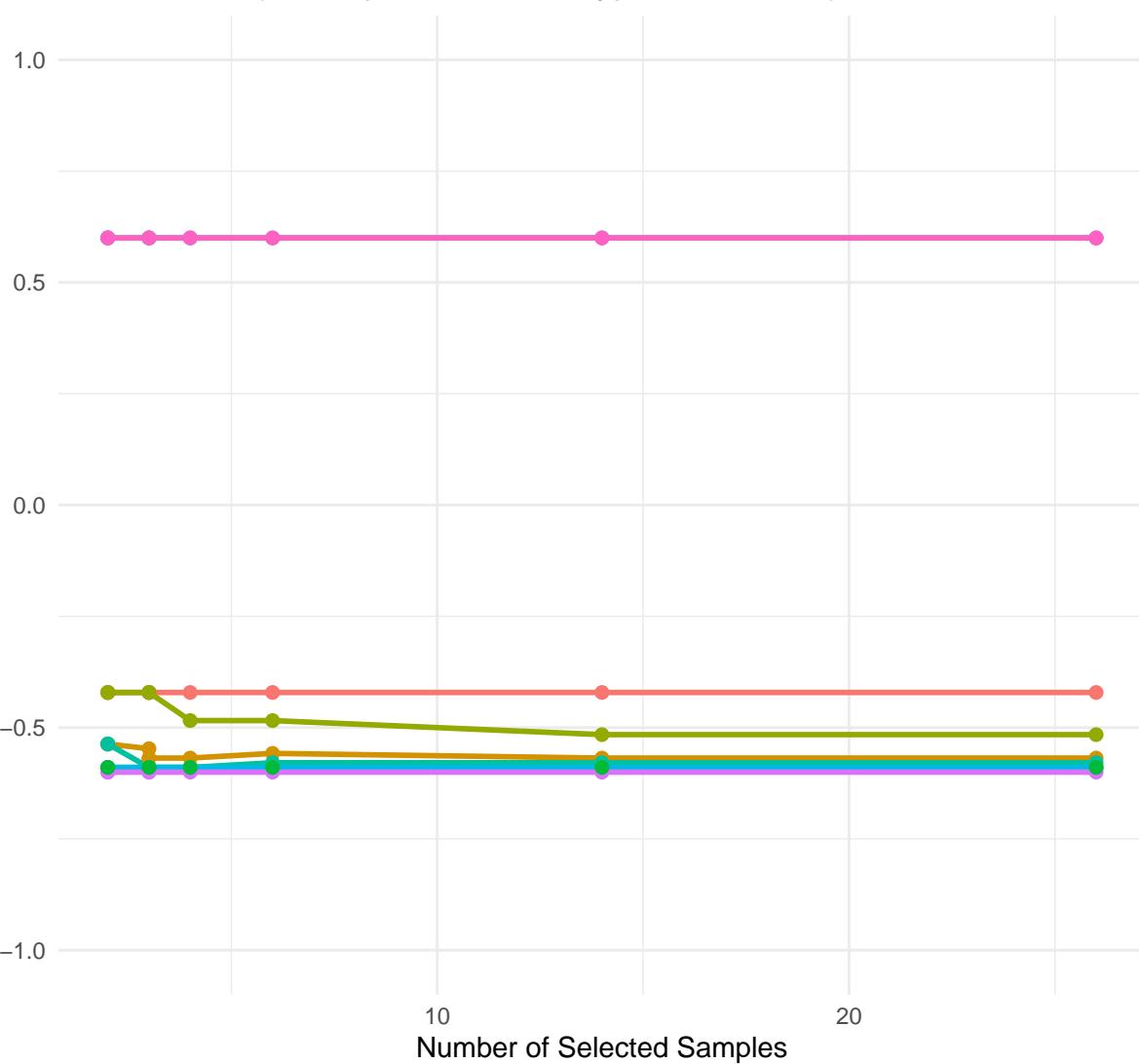
Score: Plmd (Local, partial2 , Genotype: gaussian)

Kendall Correlation



Score: Plmd (Local, partial2 , Genotype: sinusoidal)

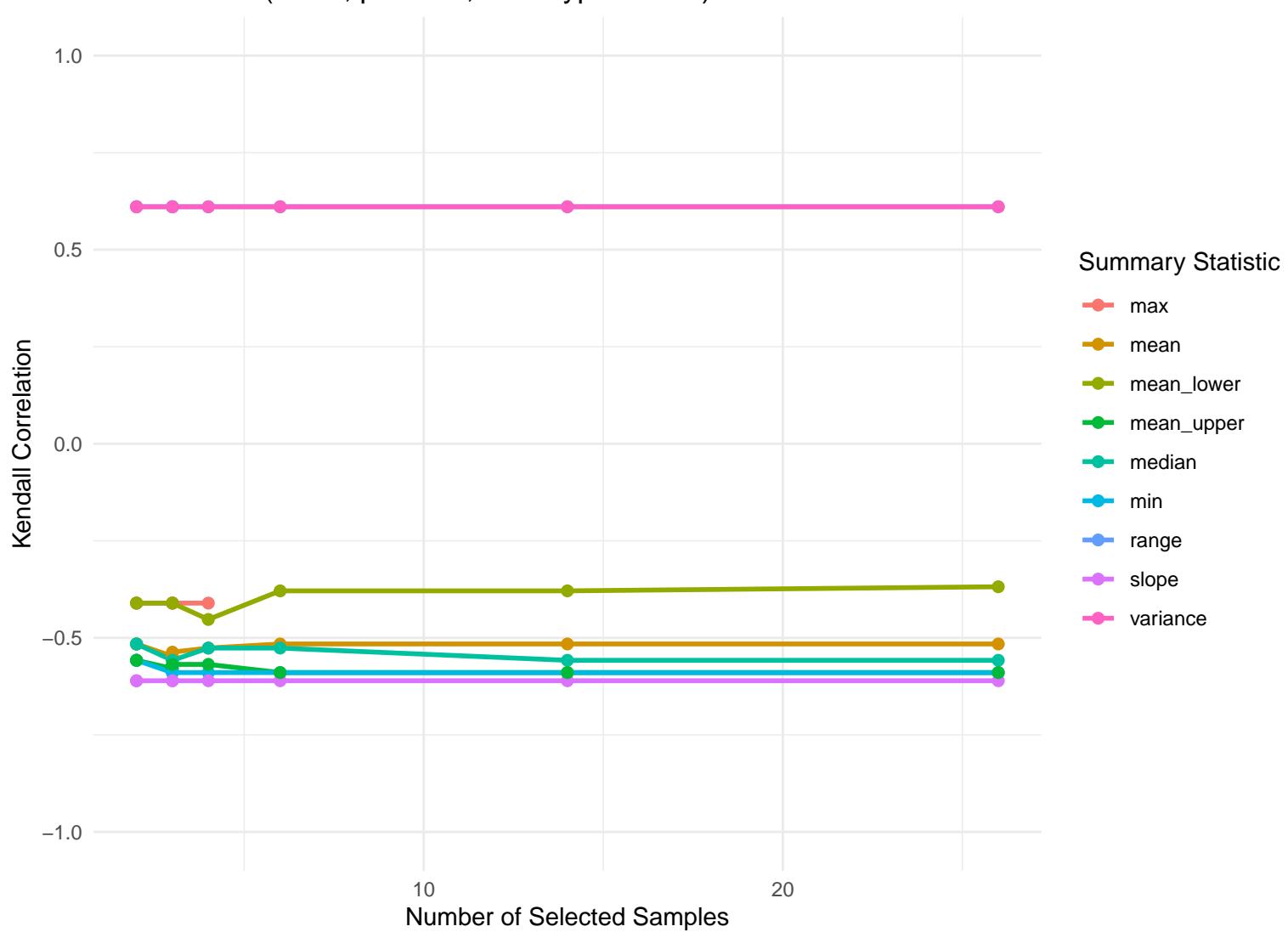
Kendall Correlation



Summary Statistic

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

Score: Plmd (Local, partial2 , Genotype: wave)



Score: PILSM (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

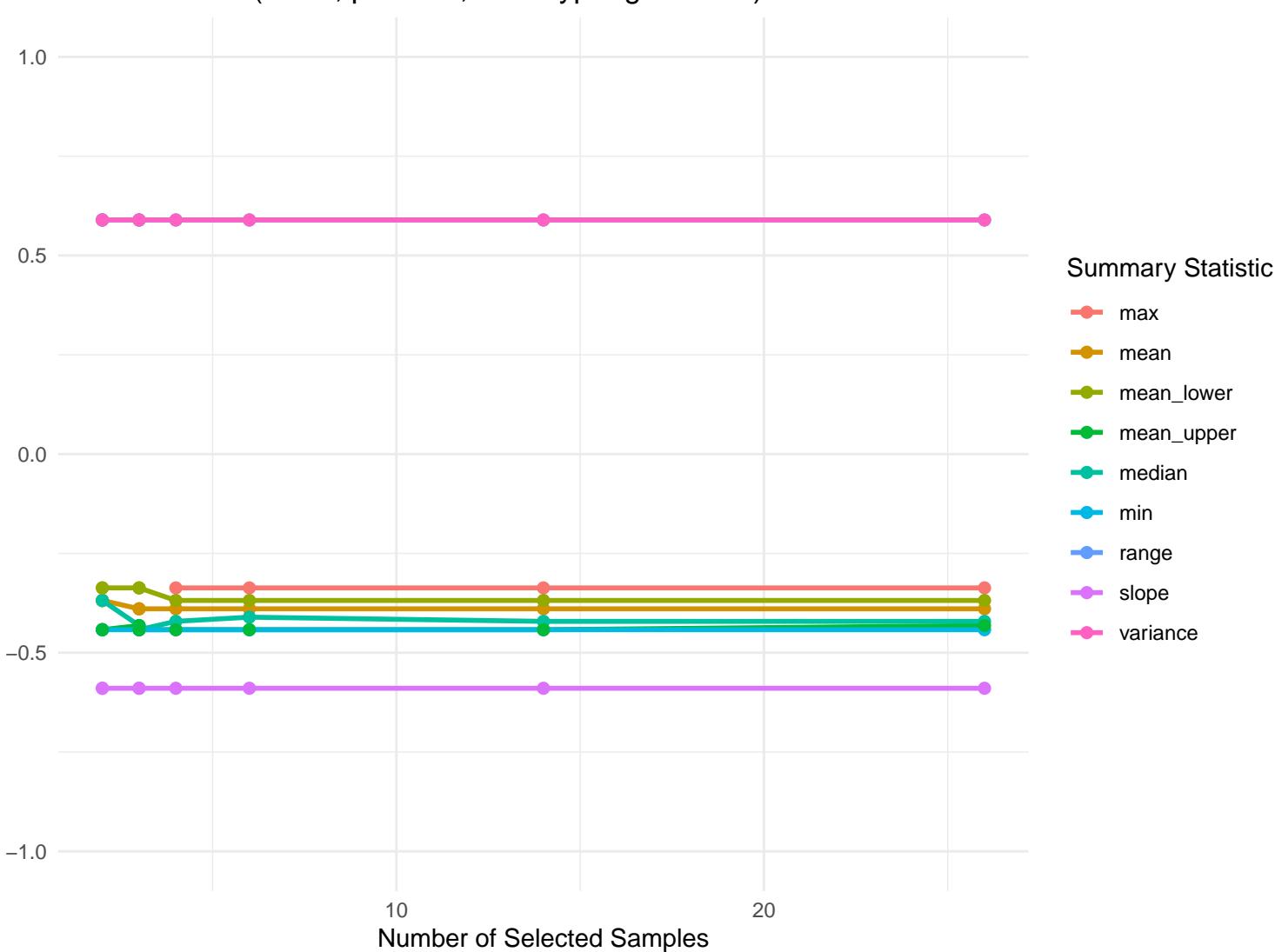
Summary Statistic

range

variance

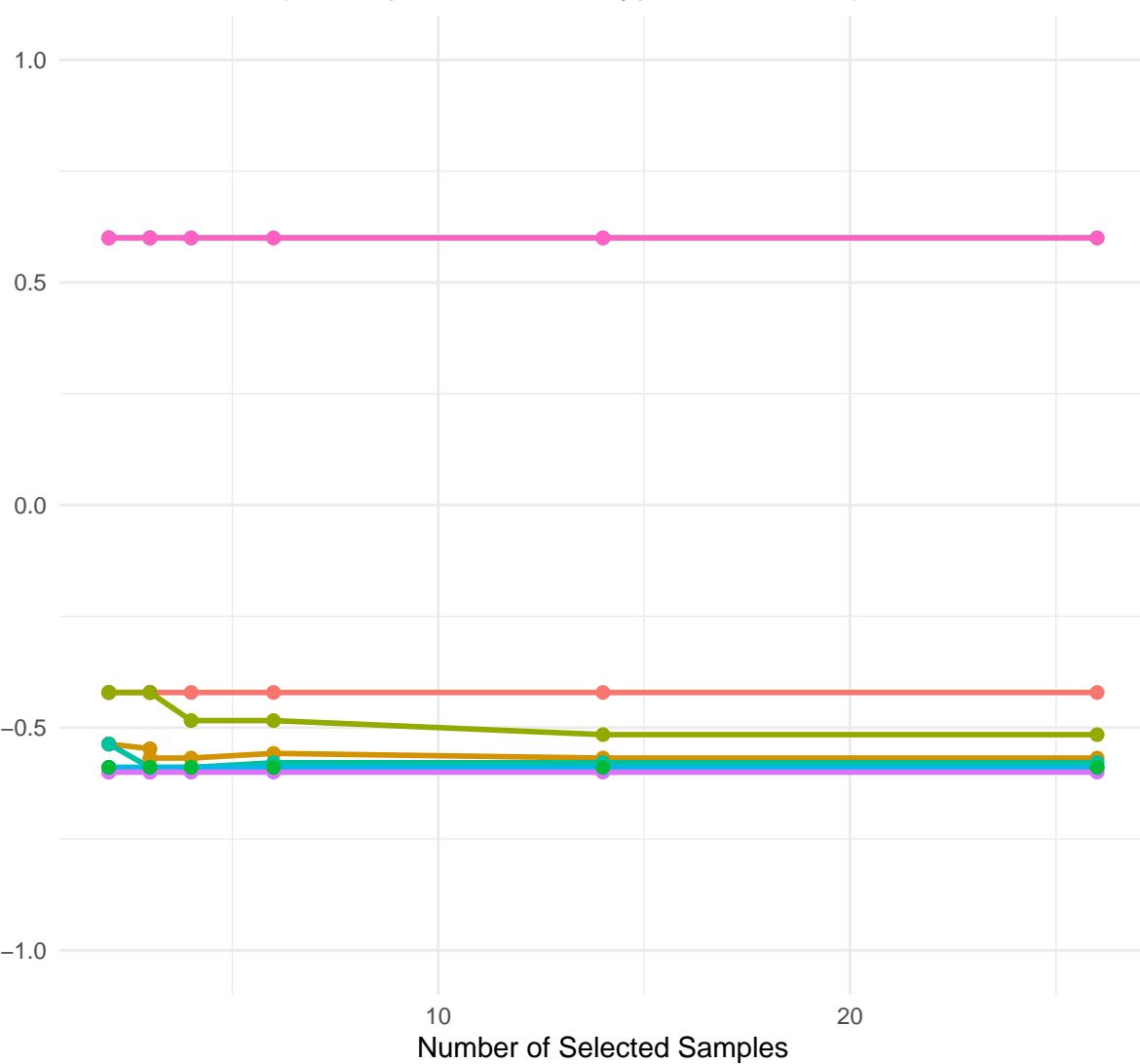


Score: PILSM (Local, partial2 , Genotype: gaussian)



Score: PILSM (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

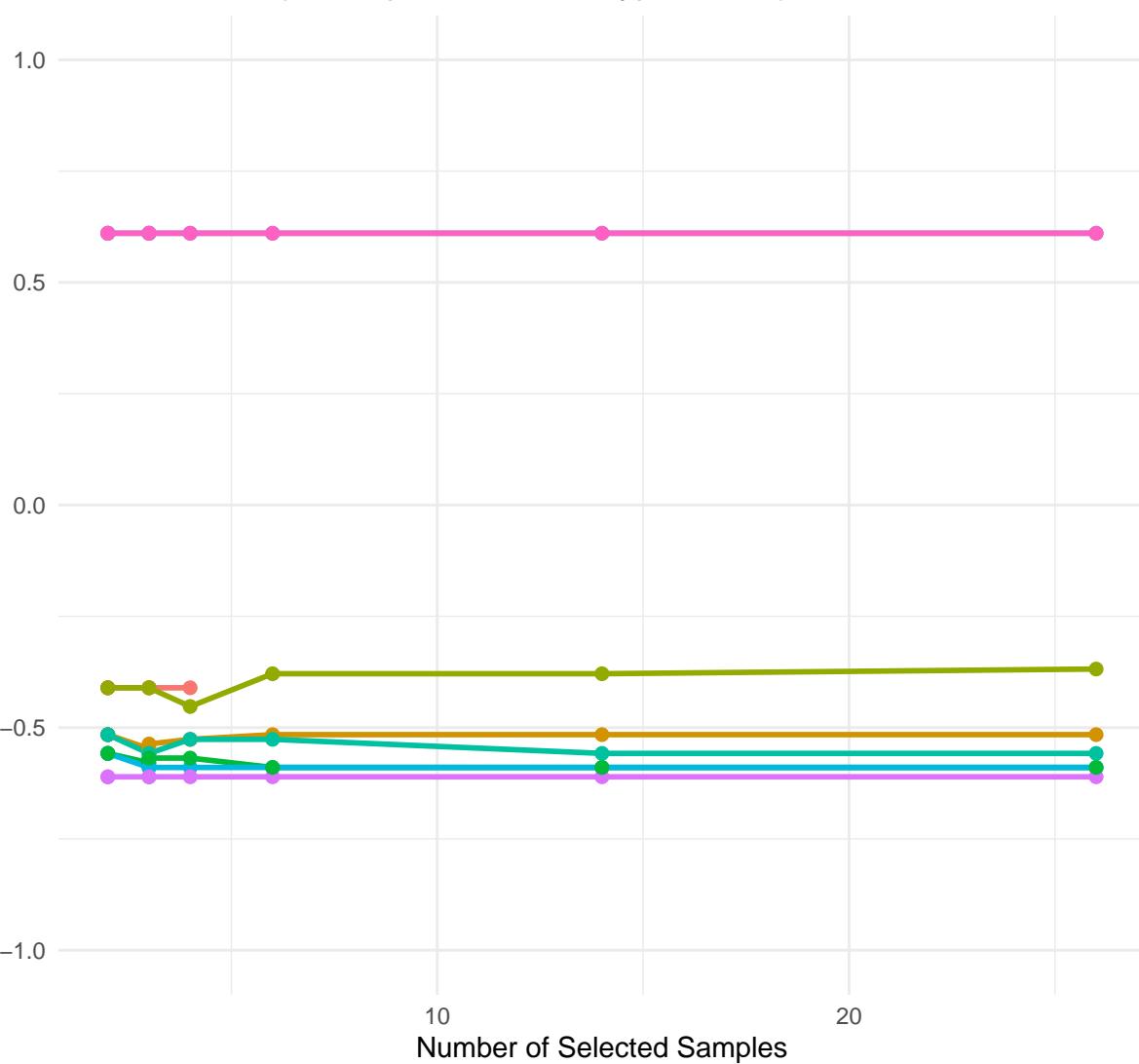


Summary Statistic

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

Score: PILSM (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: RTR (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

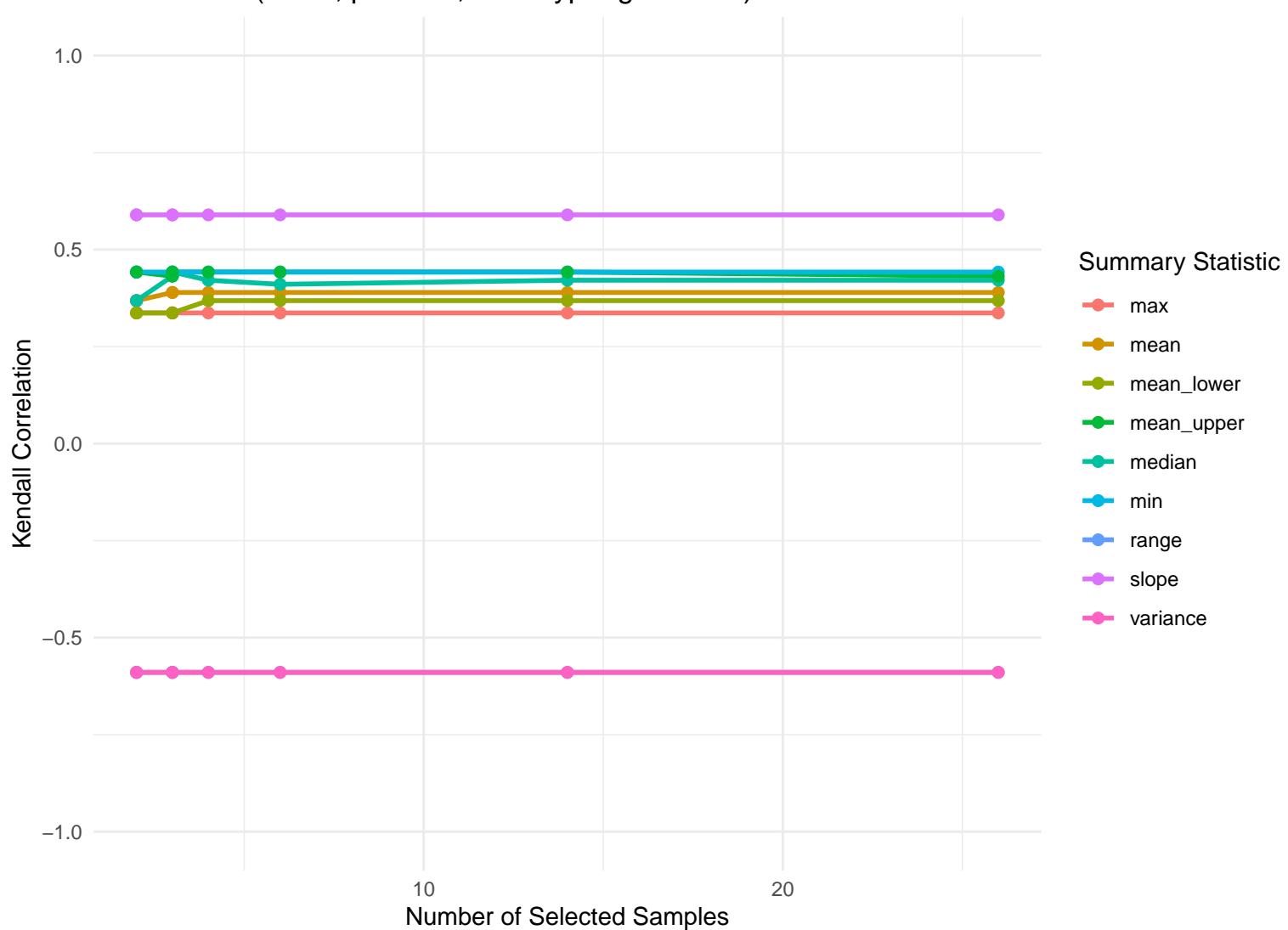
mean_upper

median

slope



Score: RTR (Local, partial2 , Genotype: gaussian)



Score: RTR (Local, partial2 , 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

0.55 0.55 0.55 0.55 0.55 0.55 0.55 0.55 0.55

0.52 0.52 0.52 0.52 0.52 0.52 0.52 0.52 0.52

0.48 0.48 0.48 0.48 0.48 0.48 0.48 0.48 0.48

0.45 0.45 0.45 0.45 0.45 0.45 0.45 0.45 0.45

0.42 0.42 0.42 0.42 0.42 0.42 0.42 0.42 0.42

0.40 0.40 0.40 0.40 0.40 0.40 0.40 0.40 0.40

0.38 0.38 0.38 0.38 0.38 0.38 0.38 0.38 0.38

0.36 0.36 0.36 0.36 0.36 0.36 0.36 0.36 0.36

0.34 0.34 0.34 0.34 0.34 0.34 0.34 0.34 0.34

0.32 0.32 0.32 0.32 0.32 0.32 0.32 0.32 0.32

0.30 0.30 0.30 0.30 0.30 0.30 0.30 0.30 0.30

0.28 0.28 0.28 0.28 0.28 0.28 0.28 0.28 0.28

0.26 0.26 0.26 0.26 0.26 0.26 0.26 0.26 0.26

0.24 0.24 0.24 0.24 0.24 0.24 0.24 0.24 0.24

0.22 0.22 0.22 0.22 0.22 0.22 0.22 0.22 0.22

0.20 0.20 0.20 0.20 0.20 0.20 0.20 0.20 0.20

0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18

0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16

0.14 0.14 0.14 0.14 0.14 0.14 0.14 0.14 0.14

0.12 0.12 0.12 0.12 0.12 0.12 0.12 0.12 0.12

0.10 0.10 0.10 0.10 0.10 0.10 0.10 0.10 0.10

0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.08

0.06 0.06 0.06 0.06 0.06 0.06 0.06 0.06 0.06

0.04 0.04 0.04 0.04 0.04 0.04 0.04 0.04 0.04

0.02 0.02 0.02 0.02 0.02 0.02 0.02 0.02 0.02

0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.00

-0.02 -0.02 -0.02 -0.02 -0.02 -0.02 -0.02 -0.02 -0.02

-0.04 -0.04 -0.04 -0.04 -0.04 -0.04 -0.04 -0.04 -0.04

-0.06 -0.06 -0.06 -0.06 -0.06 -0.06 -0.06 -0.06 -0.06

-0.08 -0.08 -0.08 -0.08 -0.08 -0.08 -0.08 -0.08 -0.08

-0.10 -0.10 -0.10 -0.10 -0.10 -0.10 -0.10 -0.10 -0.10

Score: RTR (Local, partial2 , Genotype: wave)

Kendall Correlation

1.0

0.5

0.0

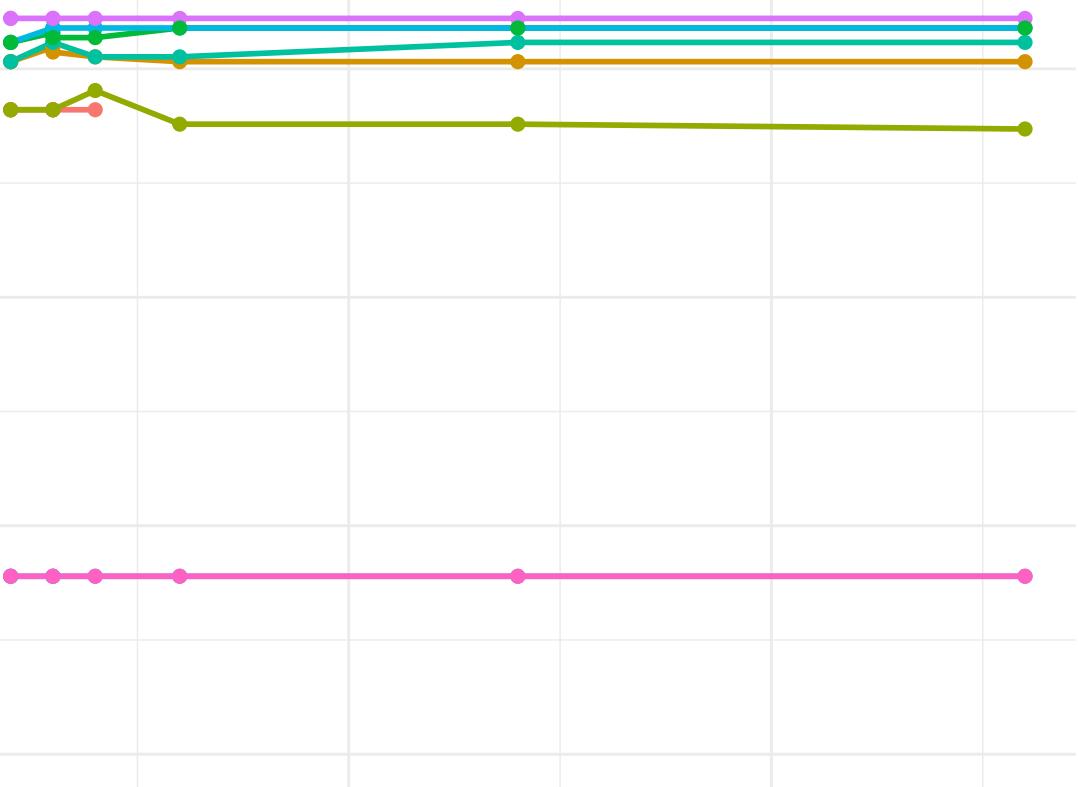
-0.5

-1.0

Number of Selected Samples

Summary Statistic

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



Score: PIR (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

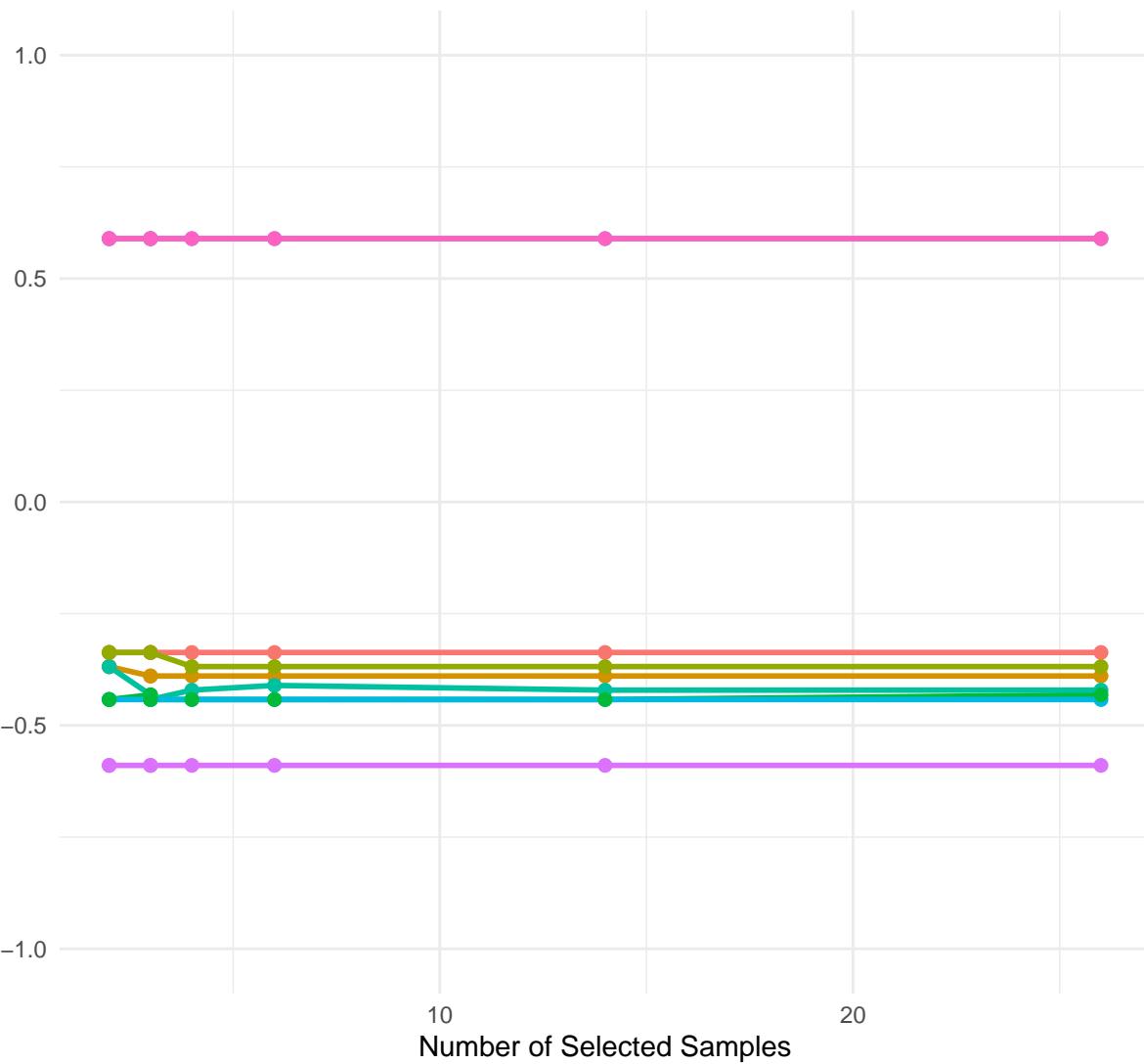
range

variance



Score: PIR (Local, partial2 , Genotype: gaussian)

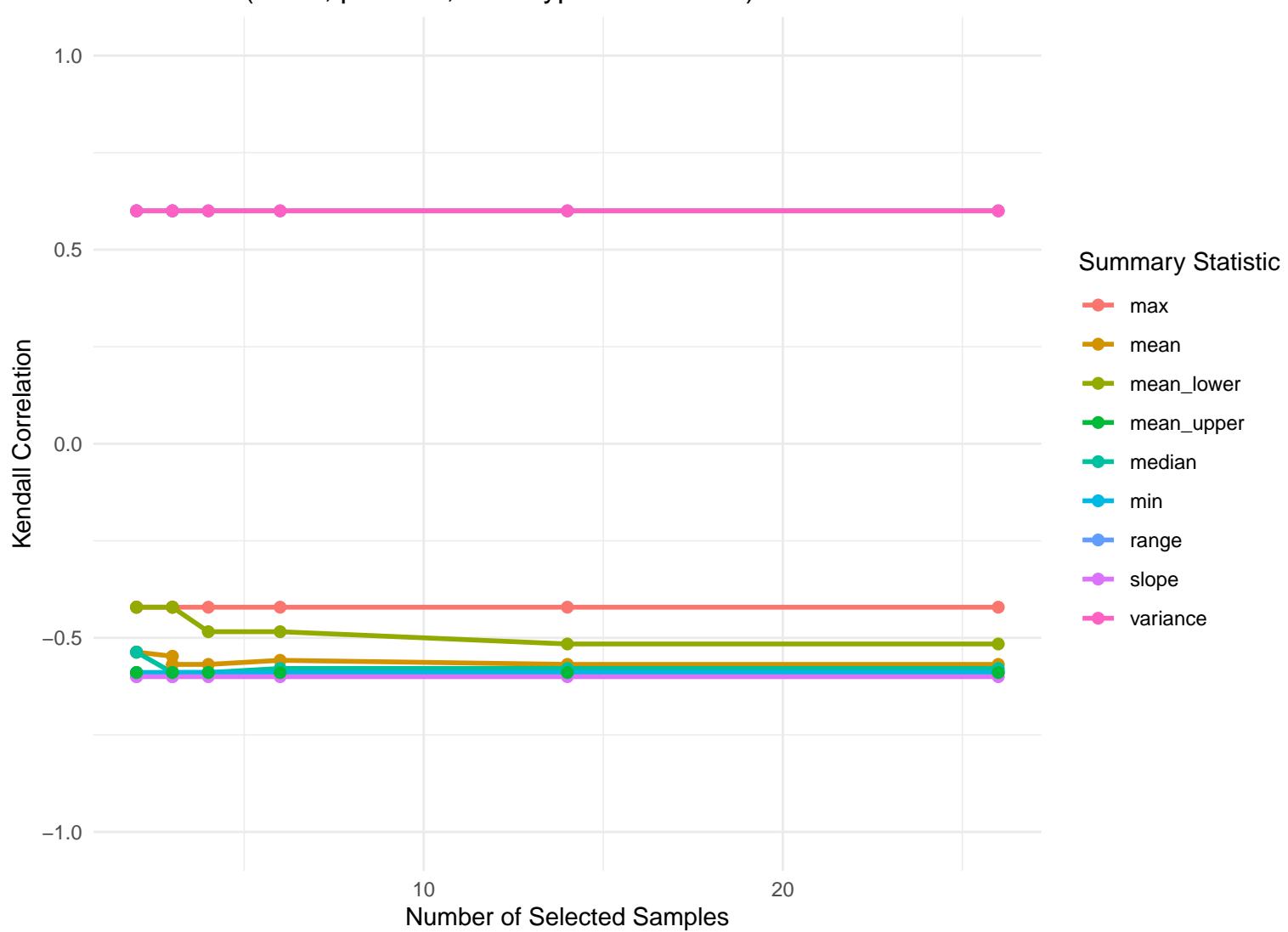
Kendall Correlation



Summary Statistic

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

Score: PIR (Local, partial2 , Genotype: sinusoidal)



Score: PIR (Local, partial2 , Genotype: wave)

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

-0.55 -0.55 -0.55 -0.55 -0.55 -0.55 -0.55

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

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

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

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

Score: RDPI (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

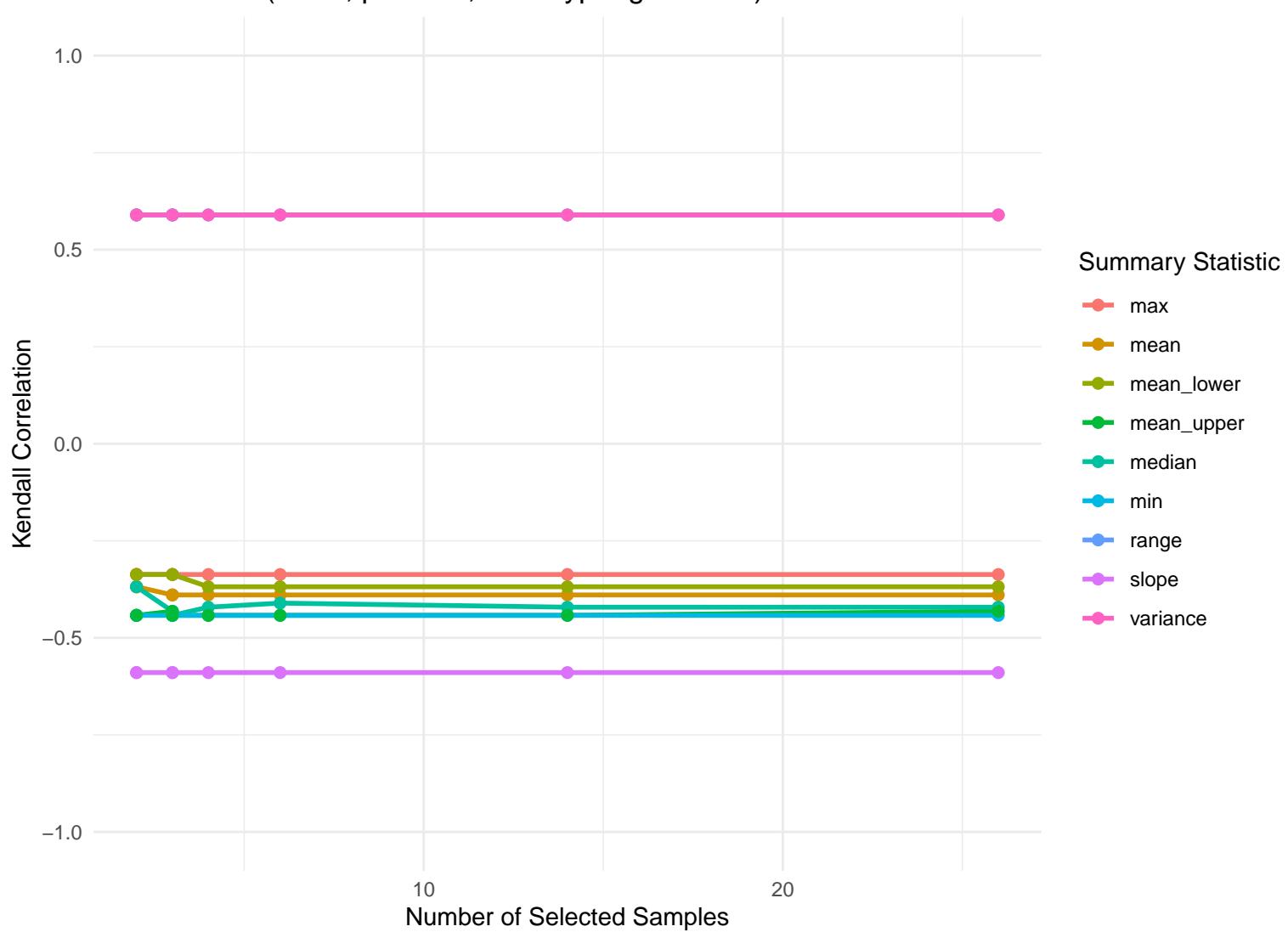
Summary Statistic

range

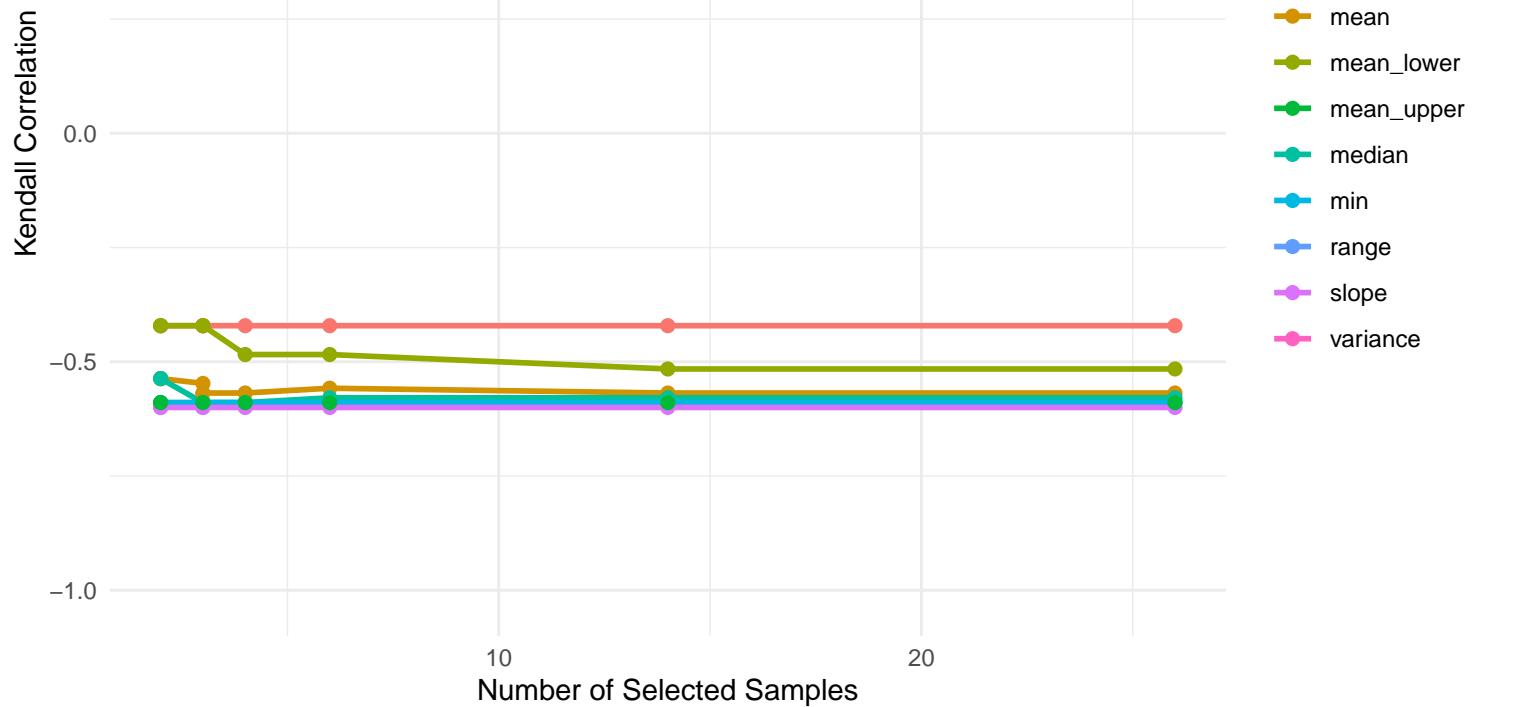
variance



Score: RDPI (Local, partial2 , Genotype: gaussian)

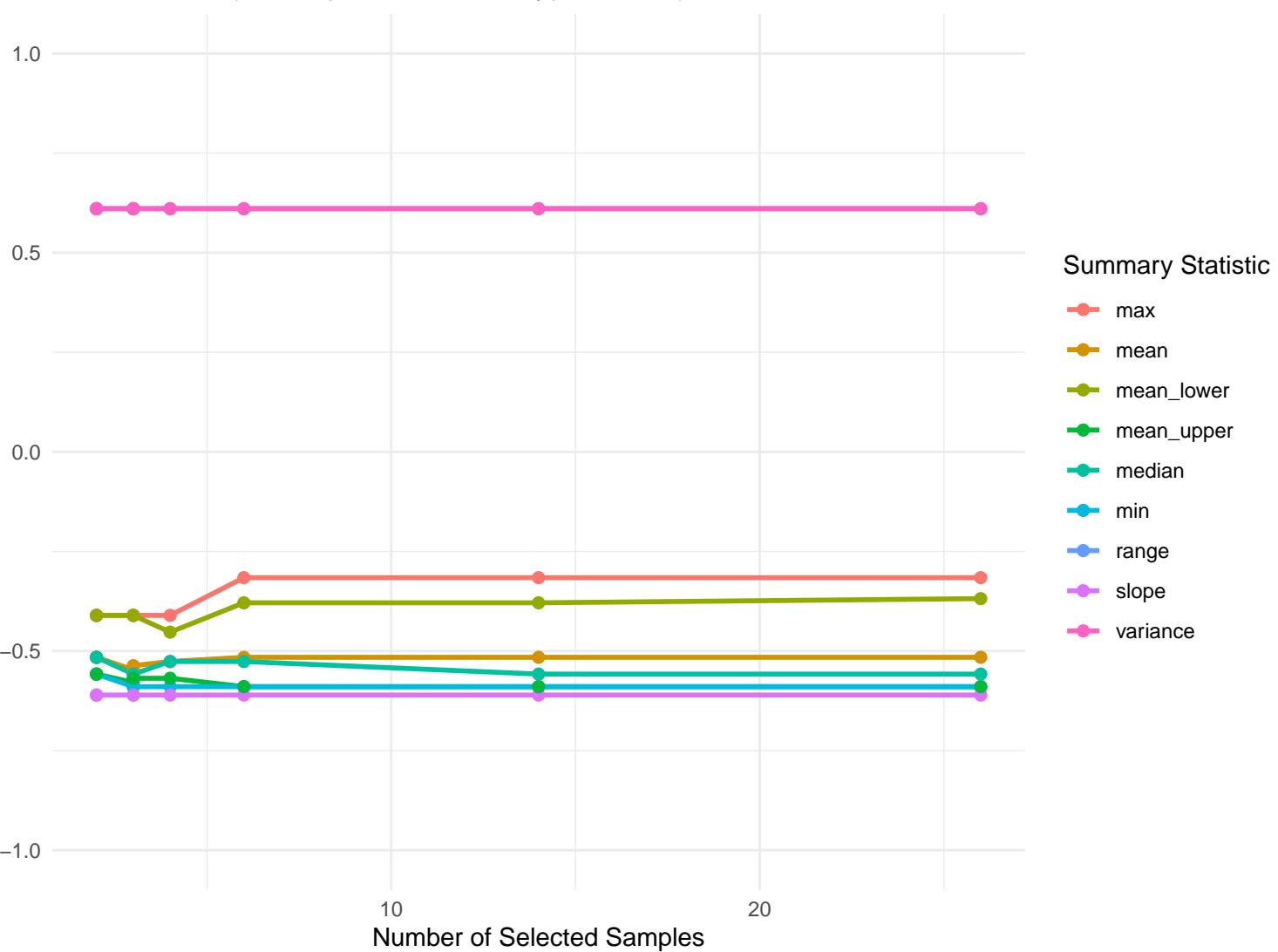


Score: RDPI (Local, partial2 , Genotype: sinusoidal)



Score: RDPI (Local, partial2 , Genotype: wave)

Kendall Correlation



Score: ESPI (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

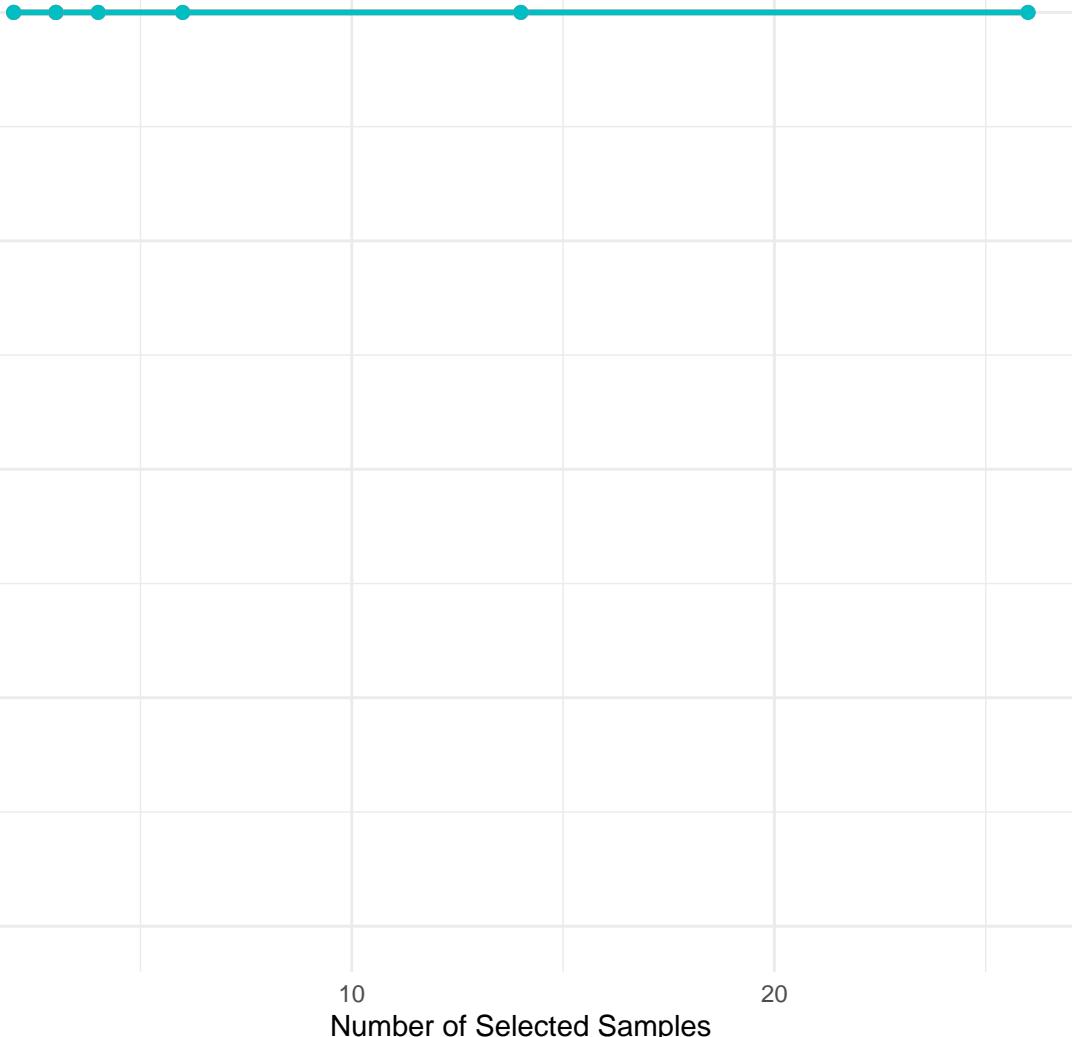
-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: ESPI (Local, partial2 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

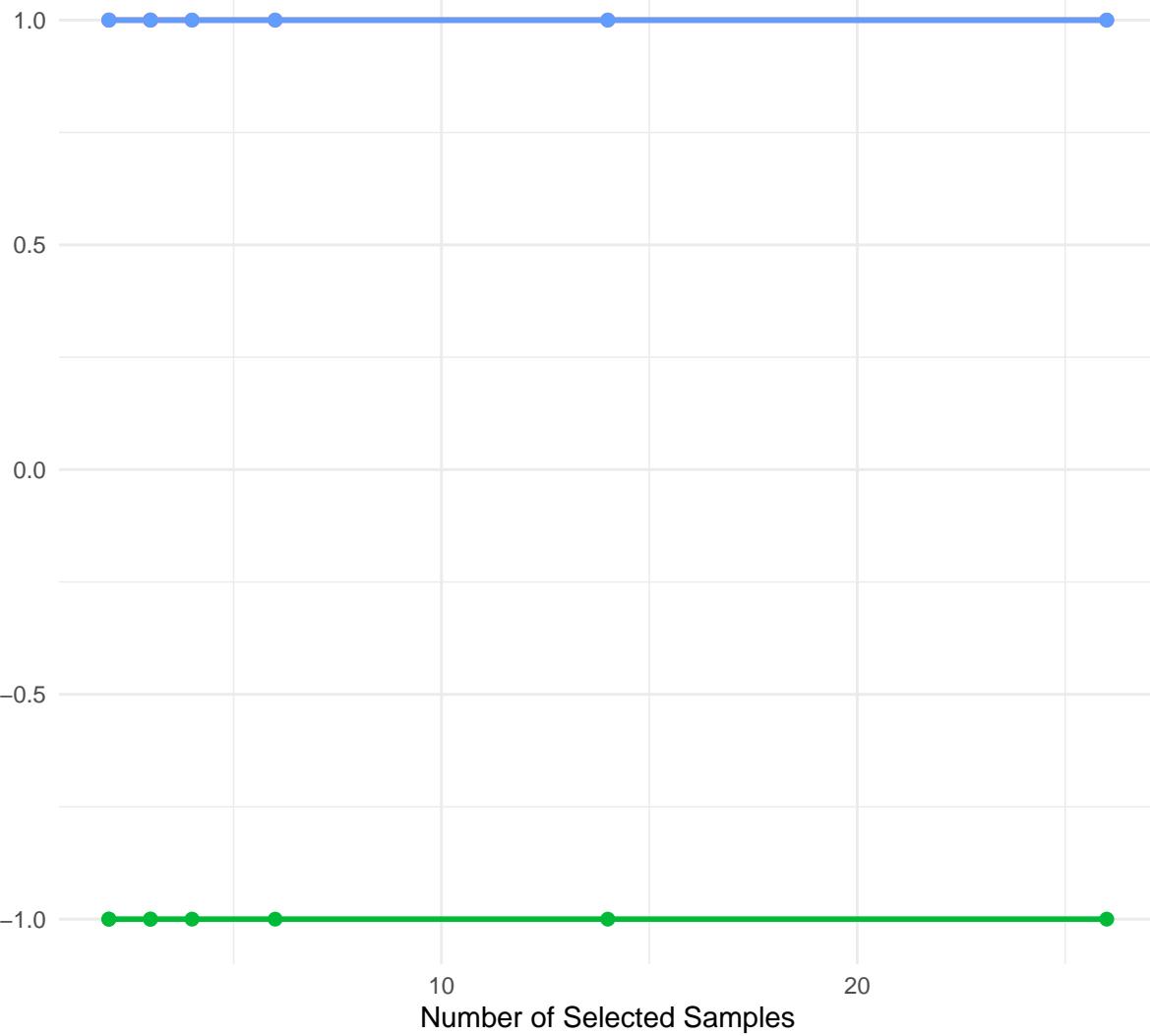
slope

variance



Score: ESPI (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

range

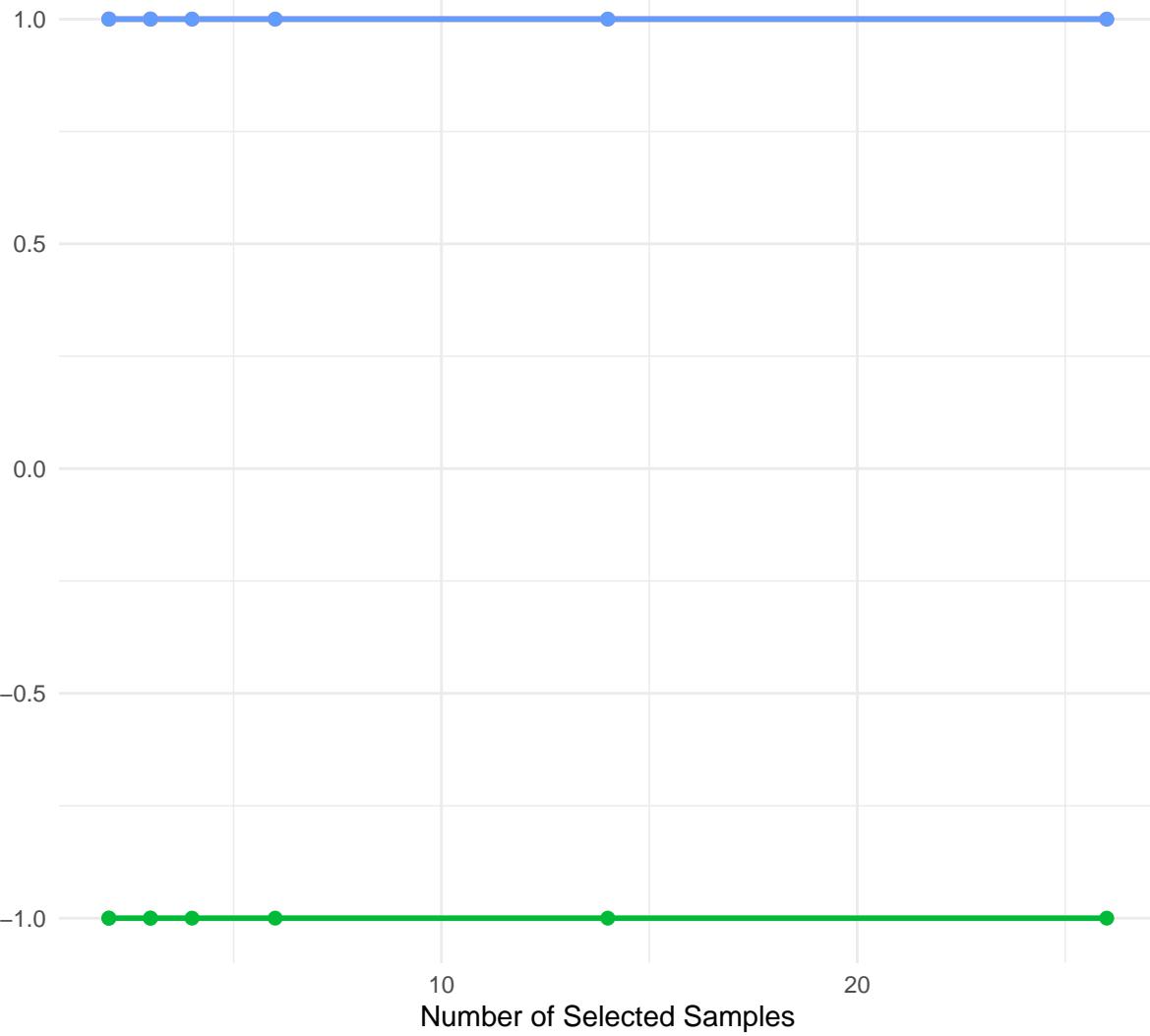
slope

variance

Number of Selected Samples

Score: ESPI (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

range

slope

variance

Number of Selected Samples

Score: ESPIID (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

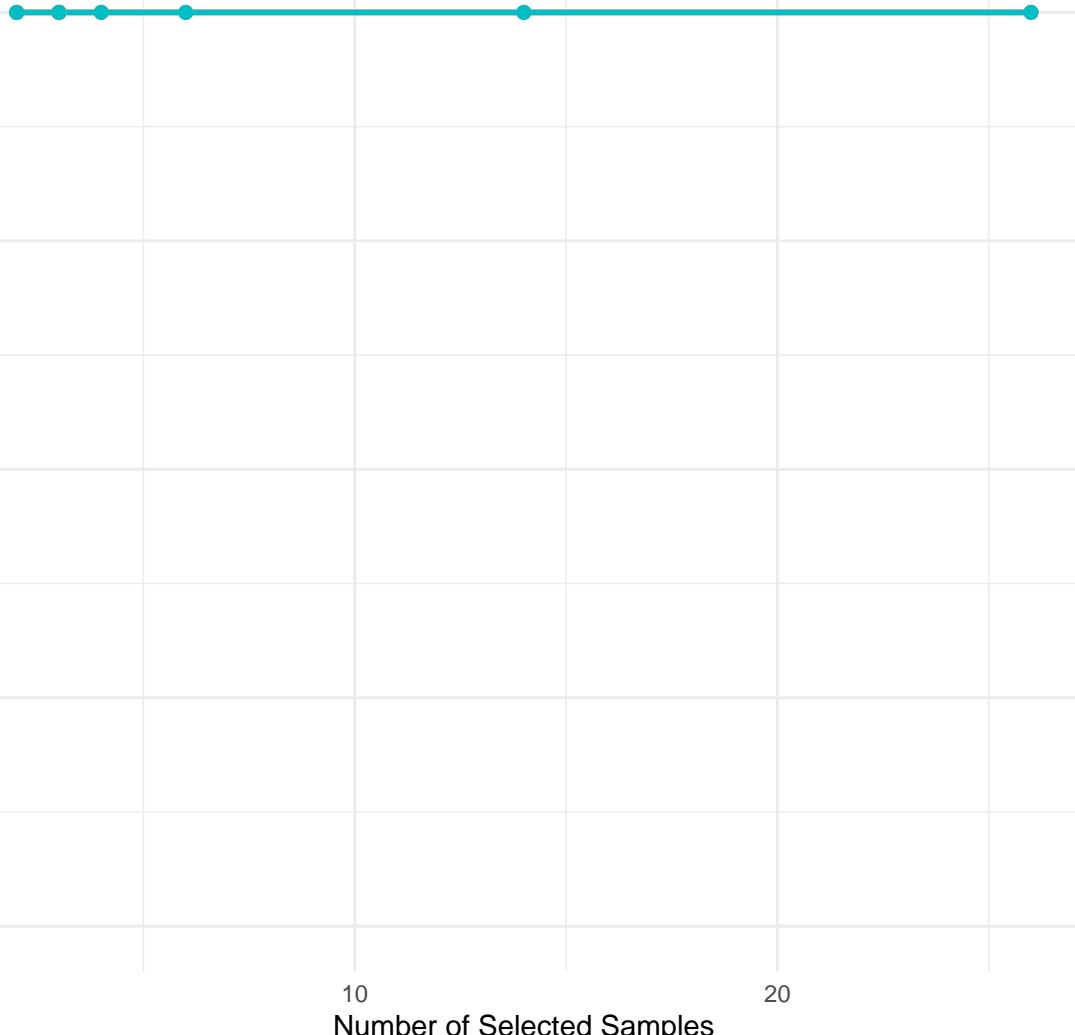
-1.0

Number of Selected Samples

Summary Statistic

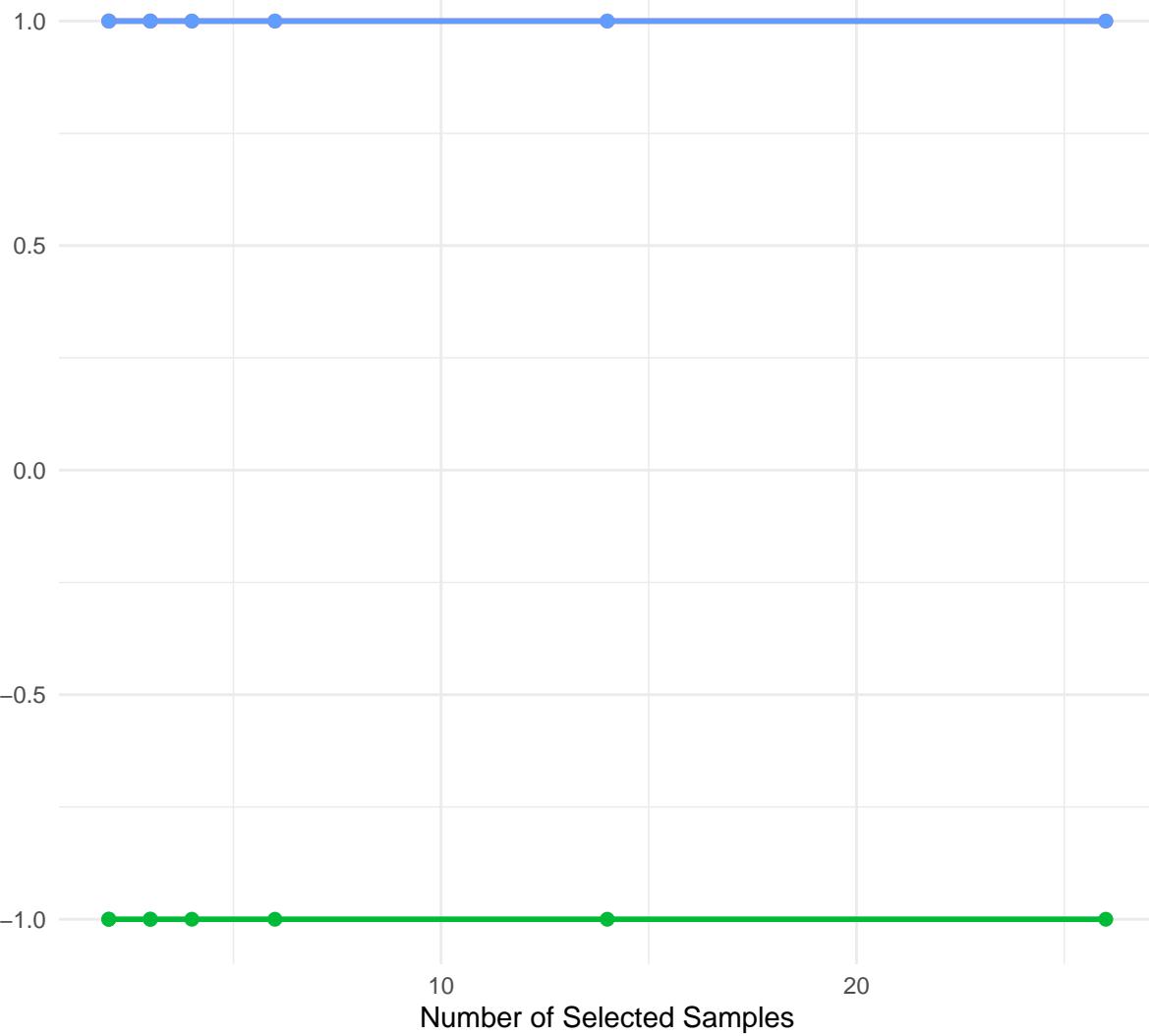
range

variance



Score: ESPIID (Local, partial2 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

range

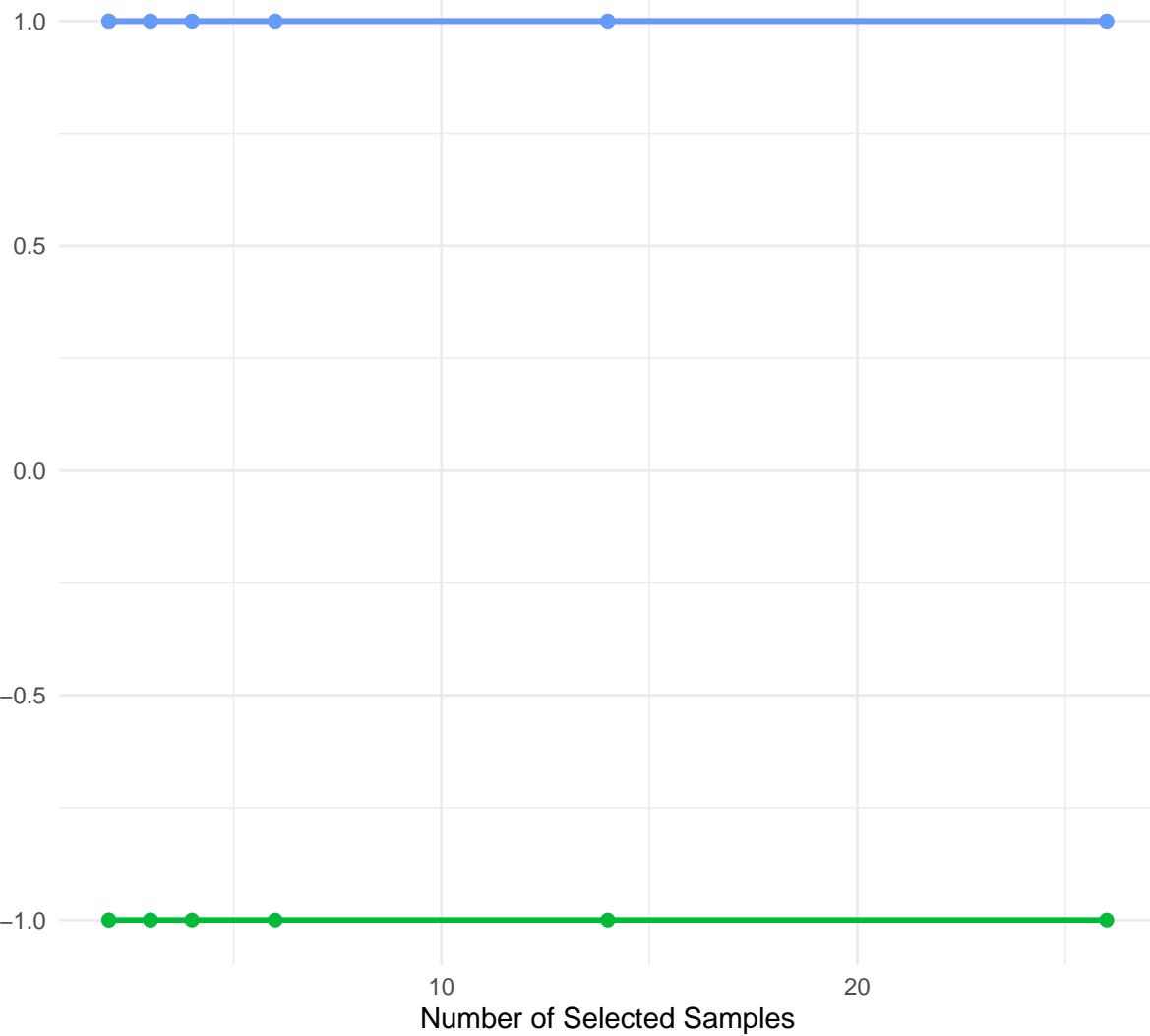
slope

variance

Number of Selected Samples

Score: ESPIID (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

range

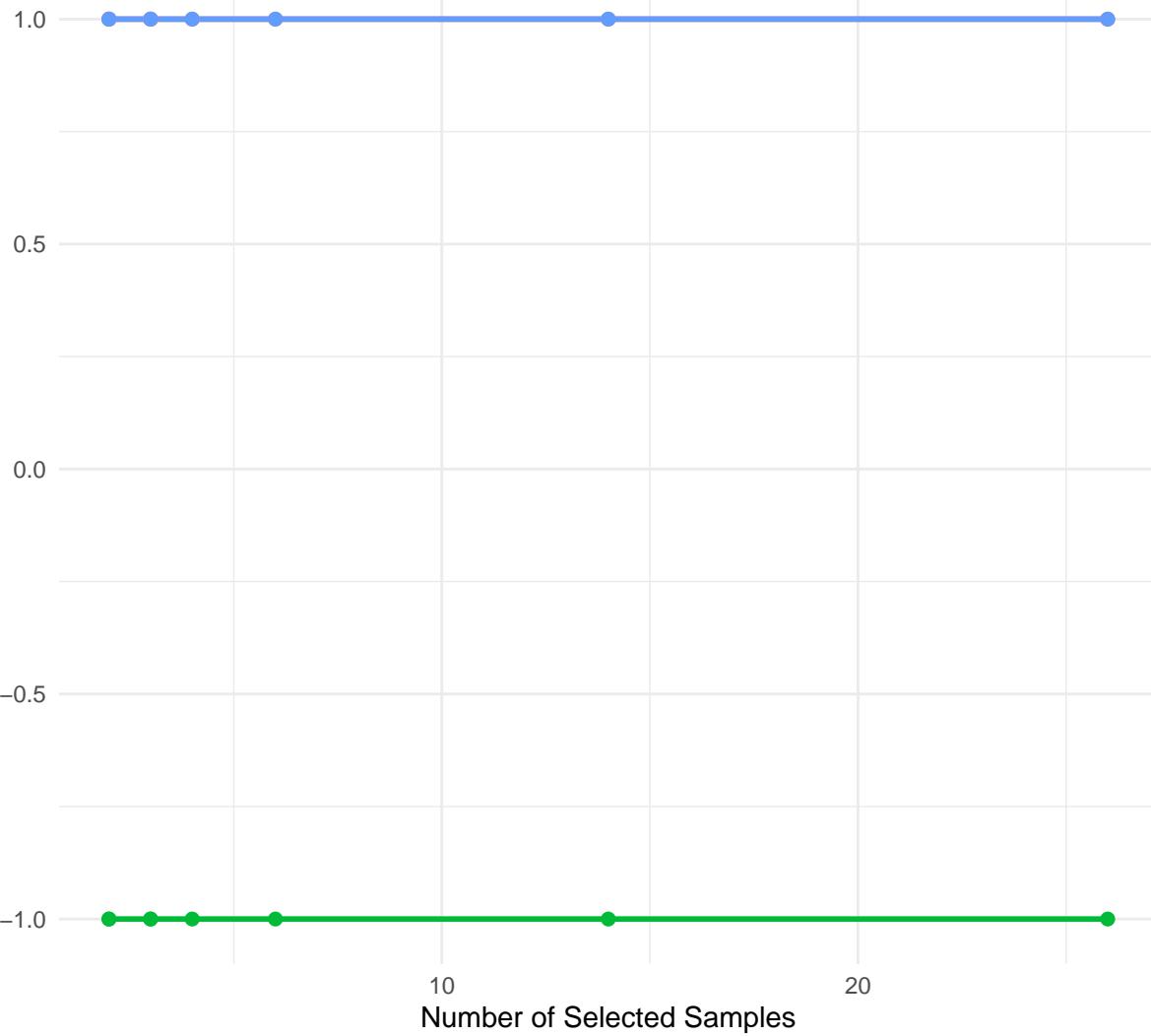
slope

variance

Number of Selected Samples

Score: ESPIID (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

range

slope

variance

Number of Selected Samples

Score: PSI (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

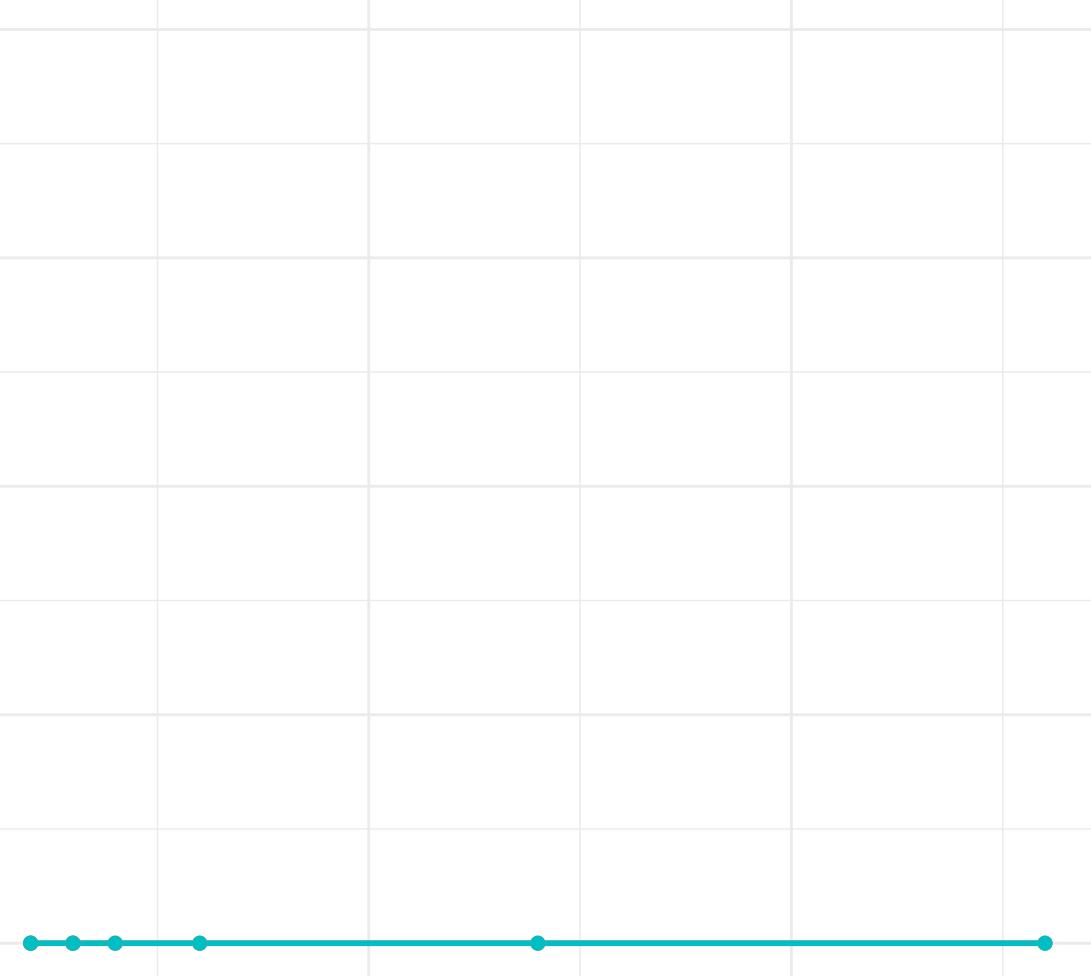
-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: PSI (Local, partial2 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

slope

variance



Score: PSI (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

slope

variance



Score: PSI (Local, partial2 , Genotype: wave)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

slope

variance



Score: RPI (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

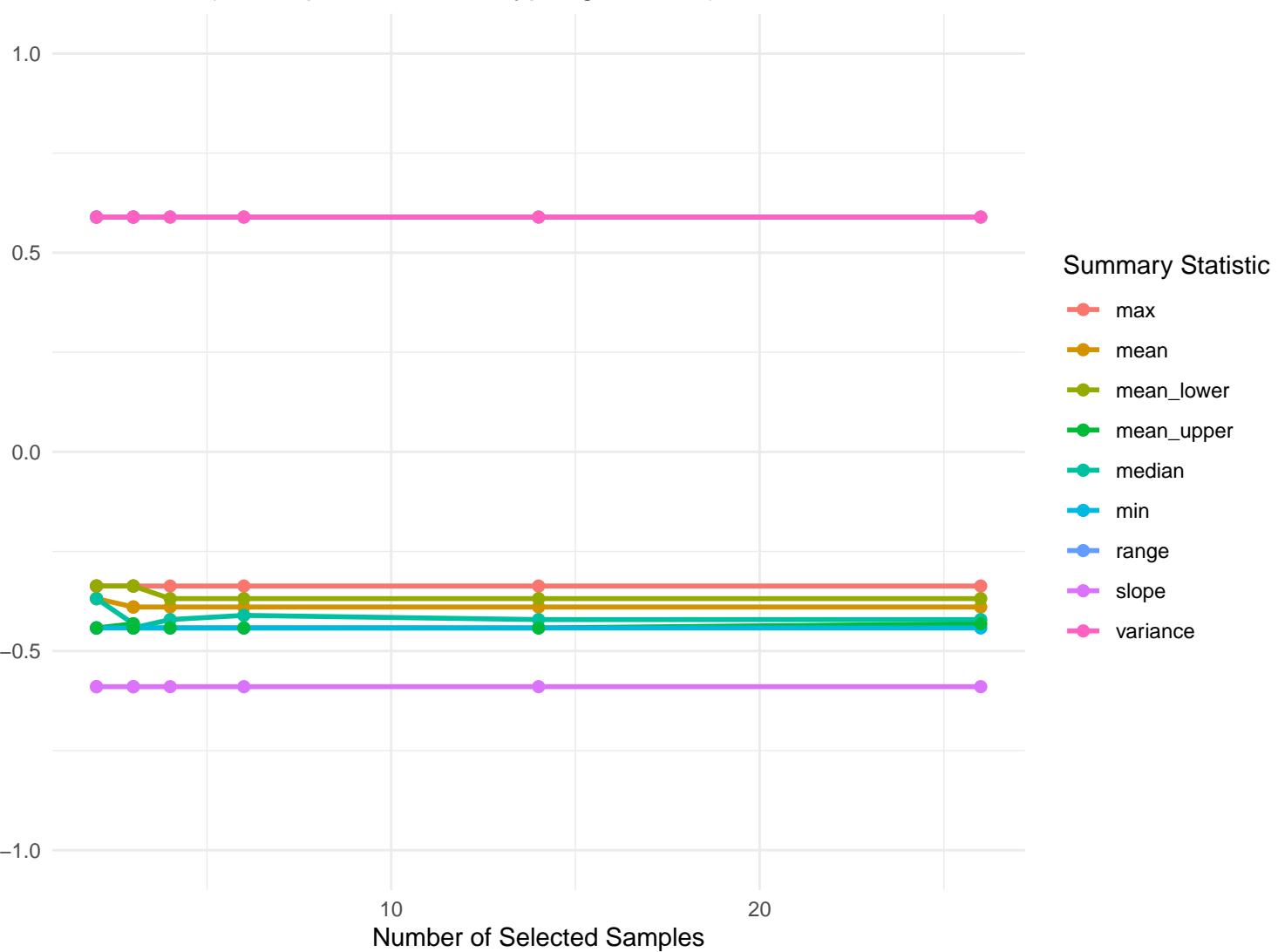
range

variance



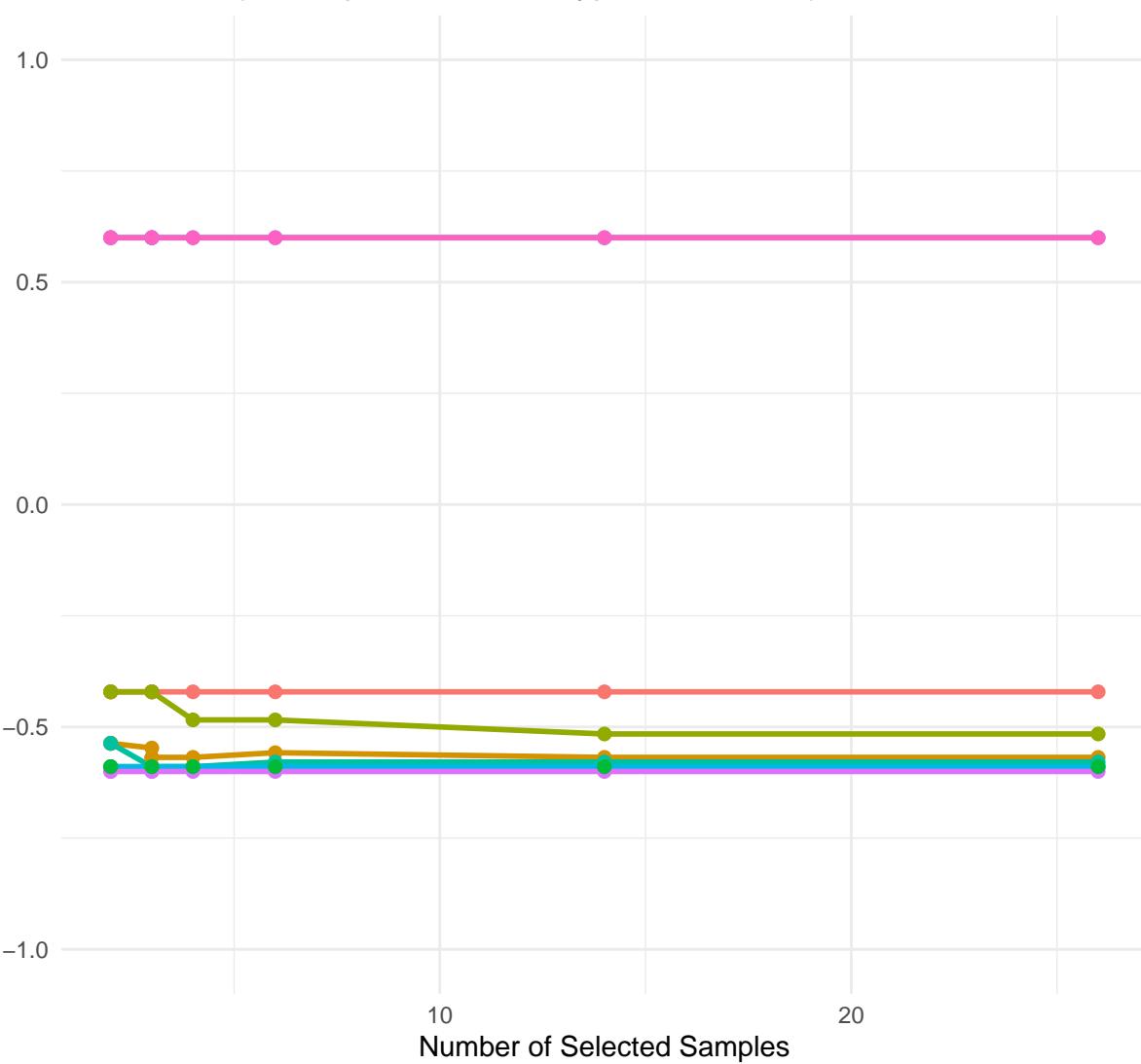
Score: RPI (Local, partial2 , Genotype: gaussian)

Kendall Correlation



Score: RPI (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

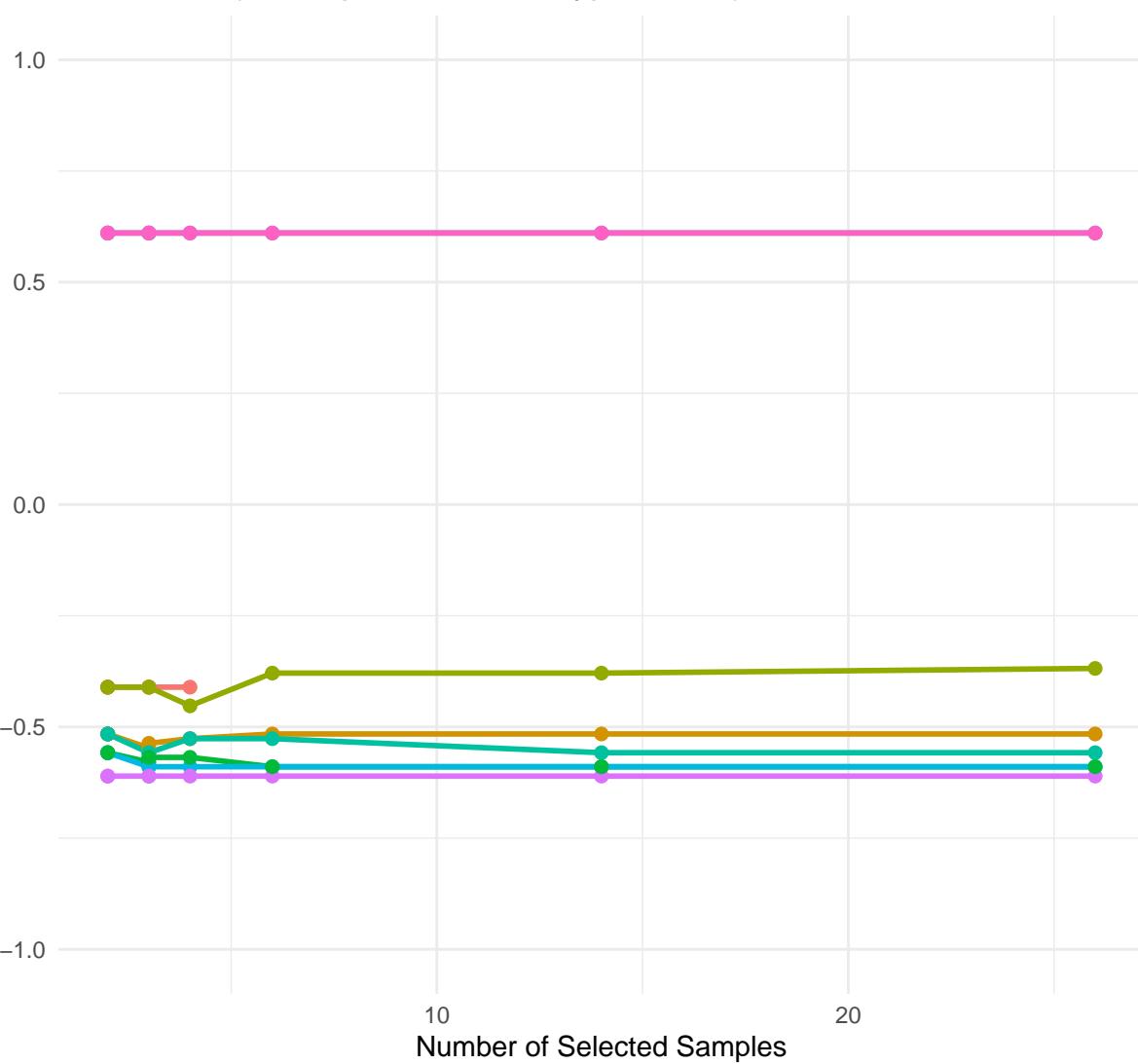


Summary Statistic

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

Score: RPI (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: PQ (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

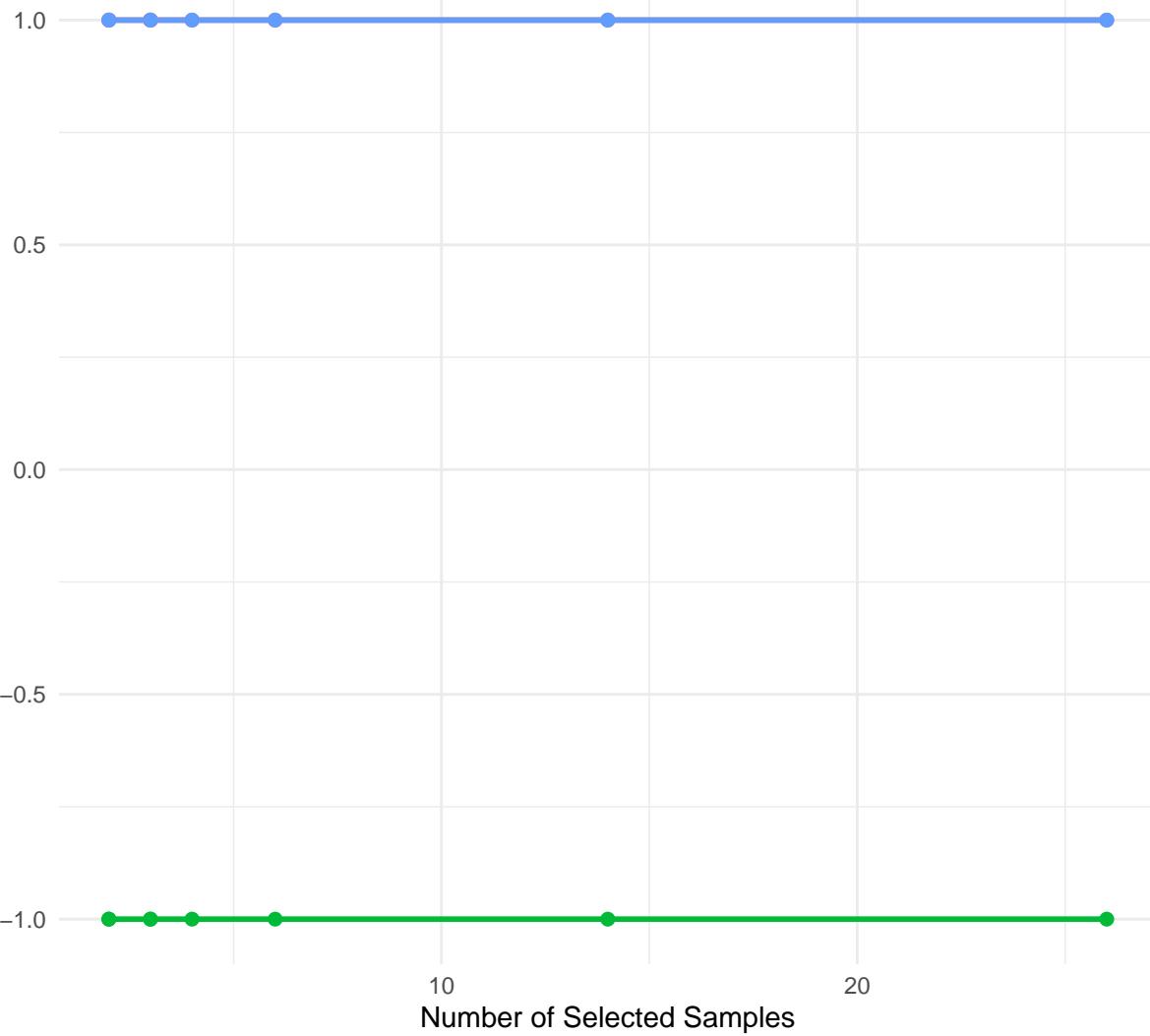
range

variance



Score: PQ (Local, partial2 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

range

slope

variance

Number of Selected Samples

Score: PQ (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Summary Statistic

range

slope

variance

Number of Selected Samples

Score: PQ (Local, partial2 , Genotype: wave)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Summary Statistic

range

slope

variance

Number of Selected Samples

Score: PR (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

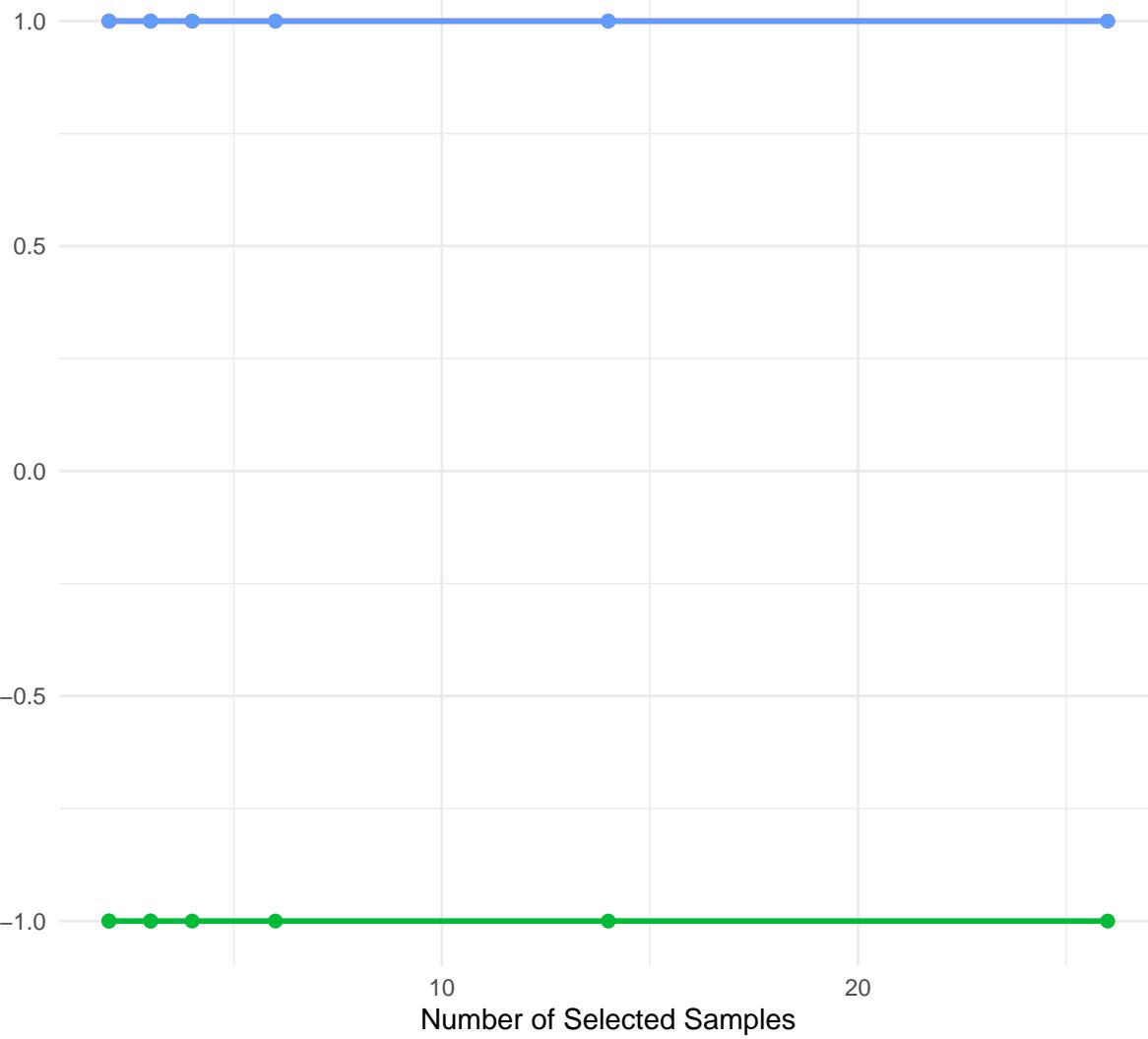
range

variance



Score: PR (Local, partial2 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

range

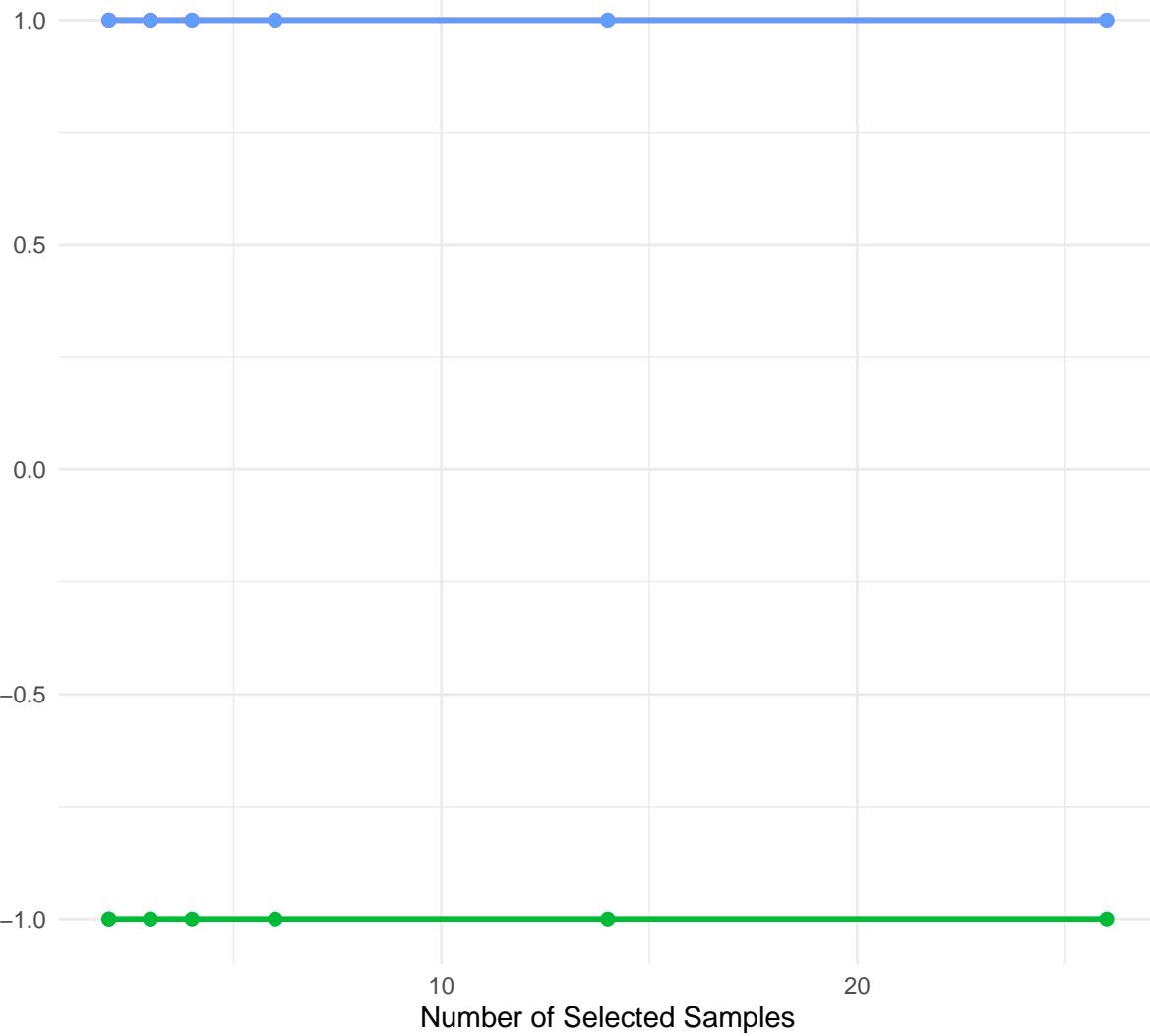
slope

variance

Number of Selected Samples

Score: PR (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

range

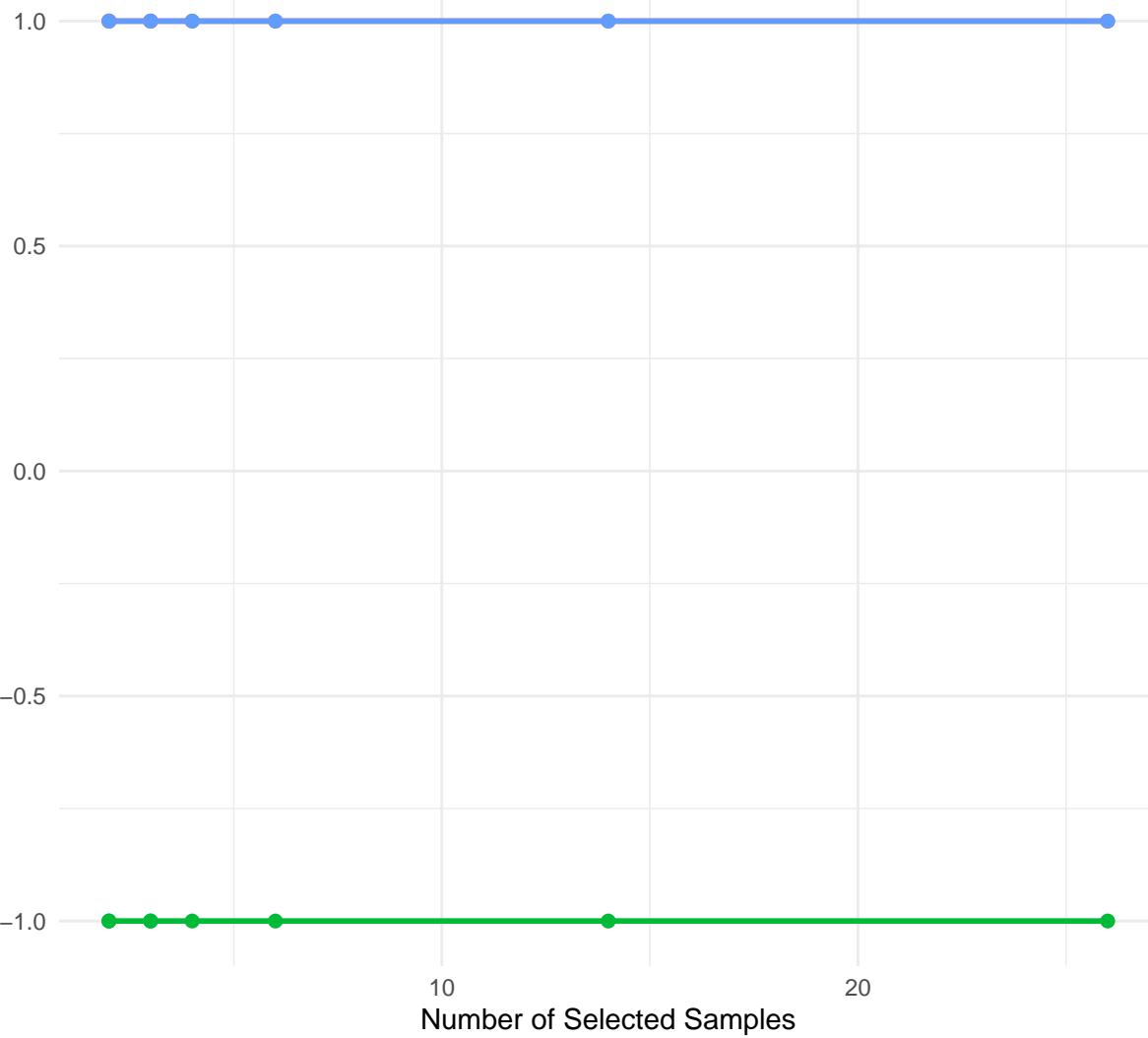
slope

variance

Number of Selected Samples

Score: PR (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

range

slope

variance

Number of Selected Samples

Score: NRW (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

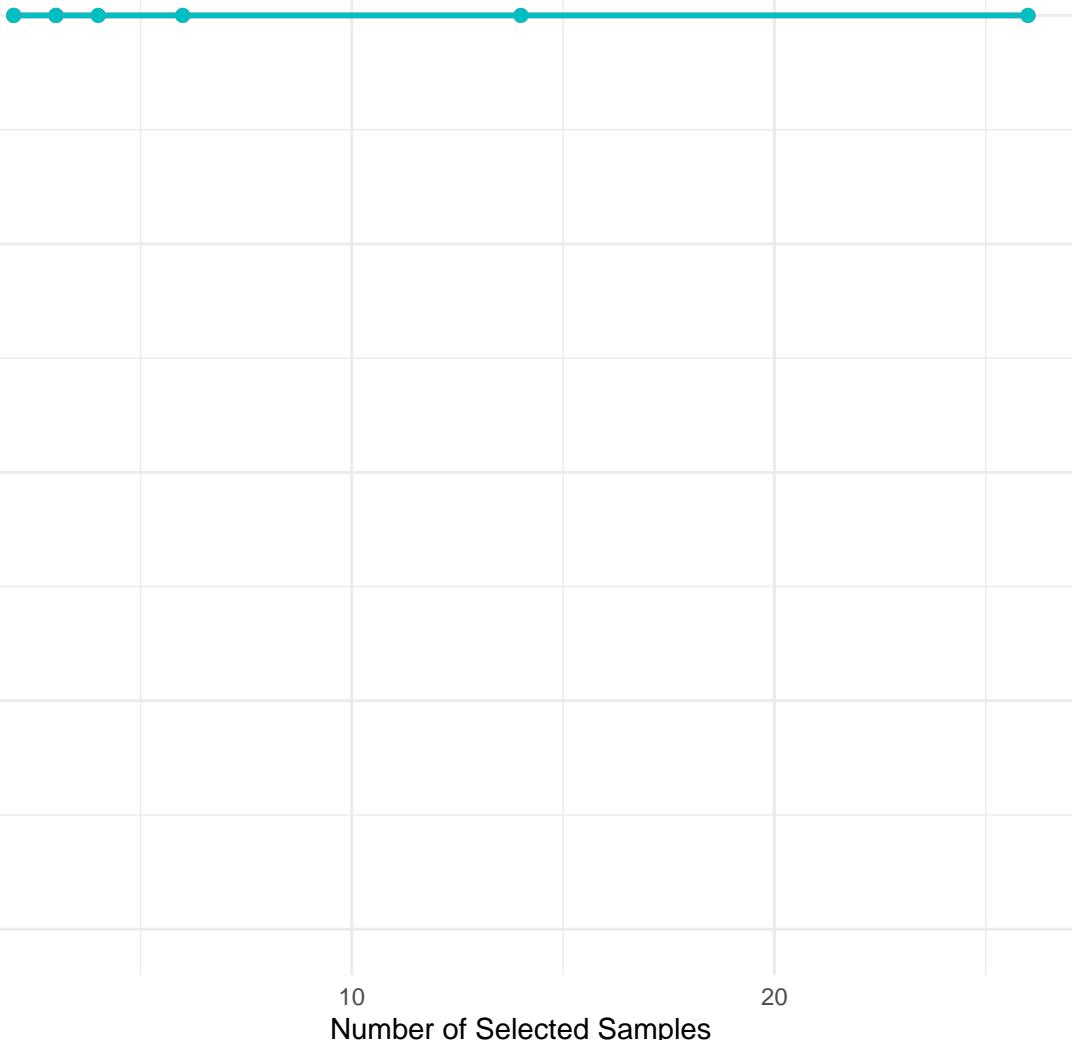
-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: NRW (Local, partial2 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

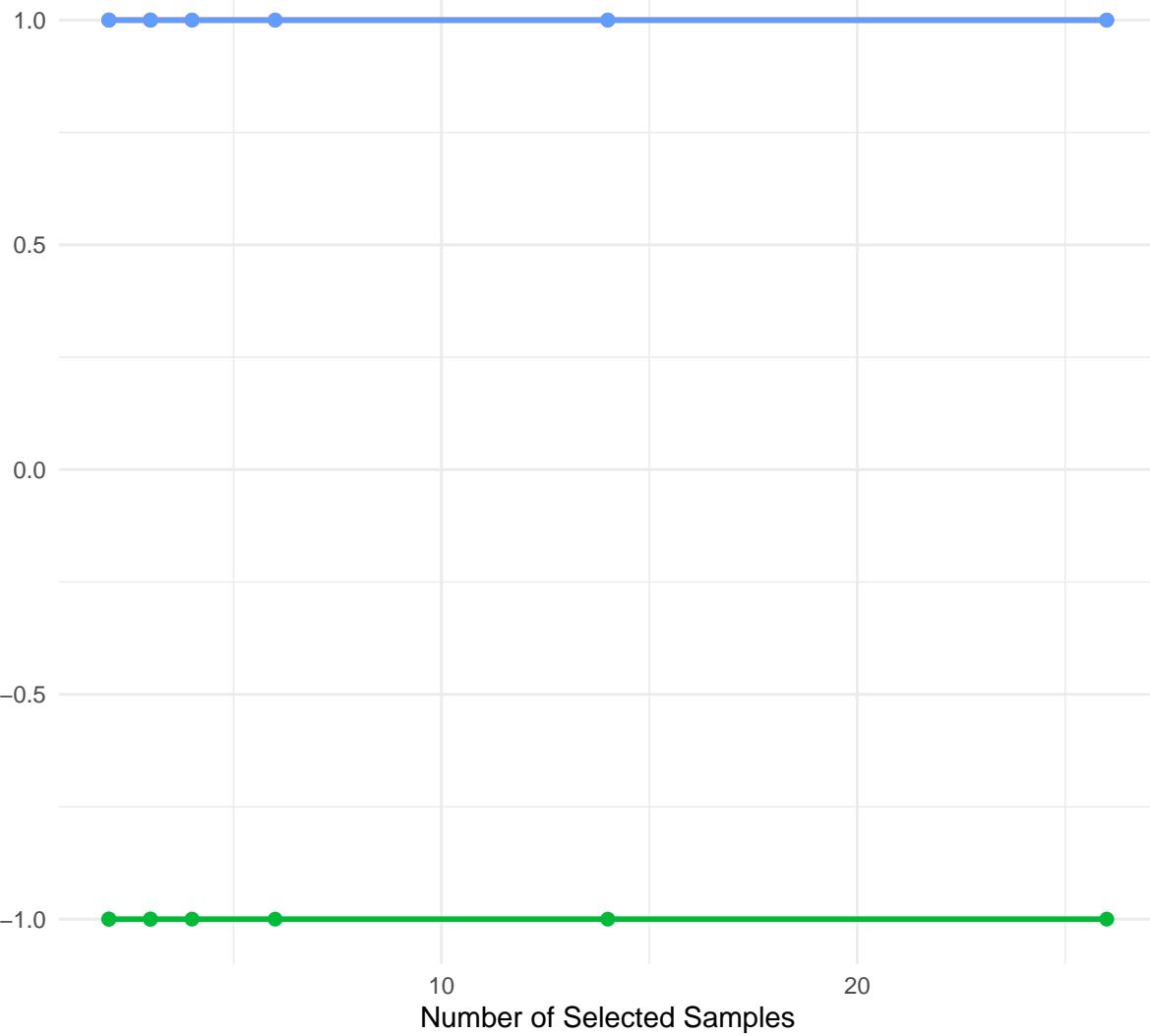
slope

variance



Score: NRW (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

range

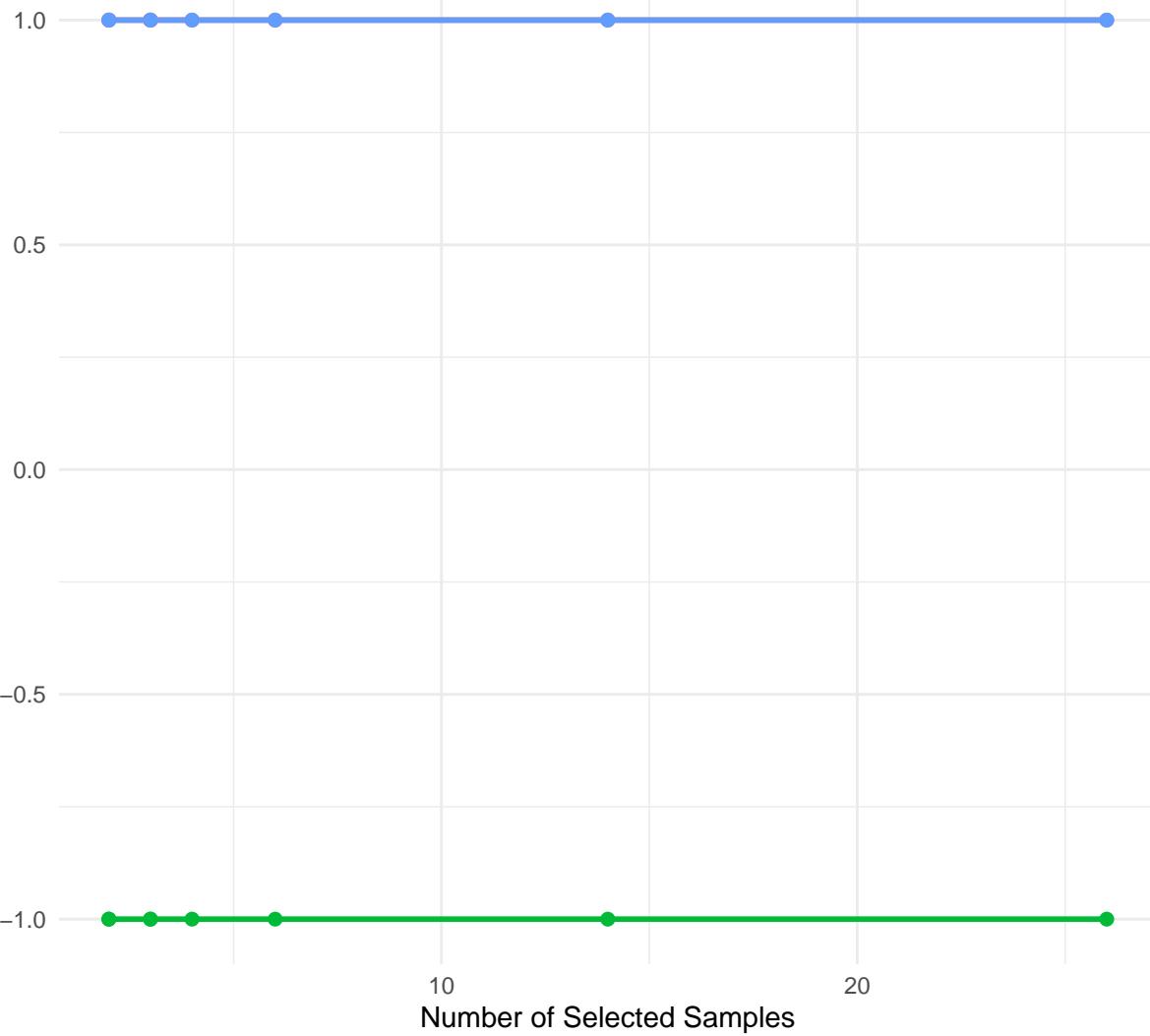
slope

variance

Number of Selected Samples

Score: NRW (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

range

slope

variance

Number of Selected Samples

Score: ESP (Local, partial2 , 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, partial2 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

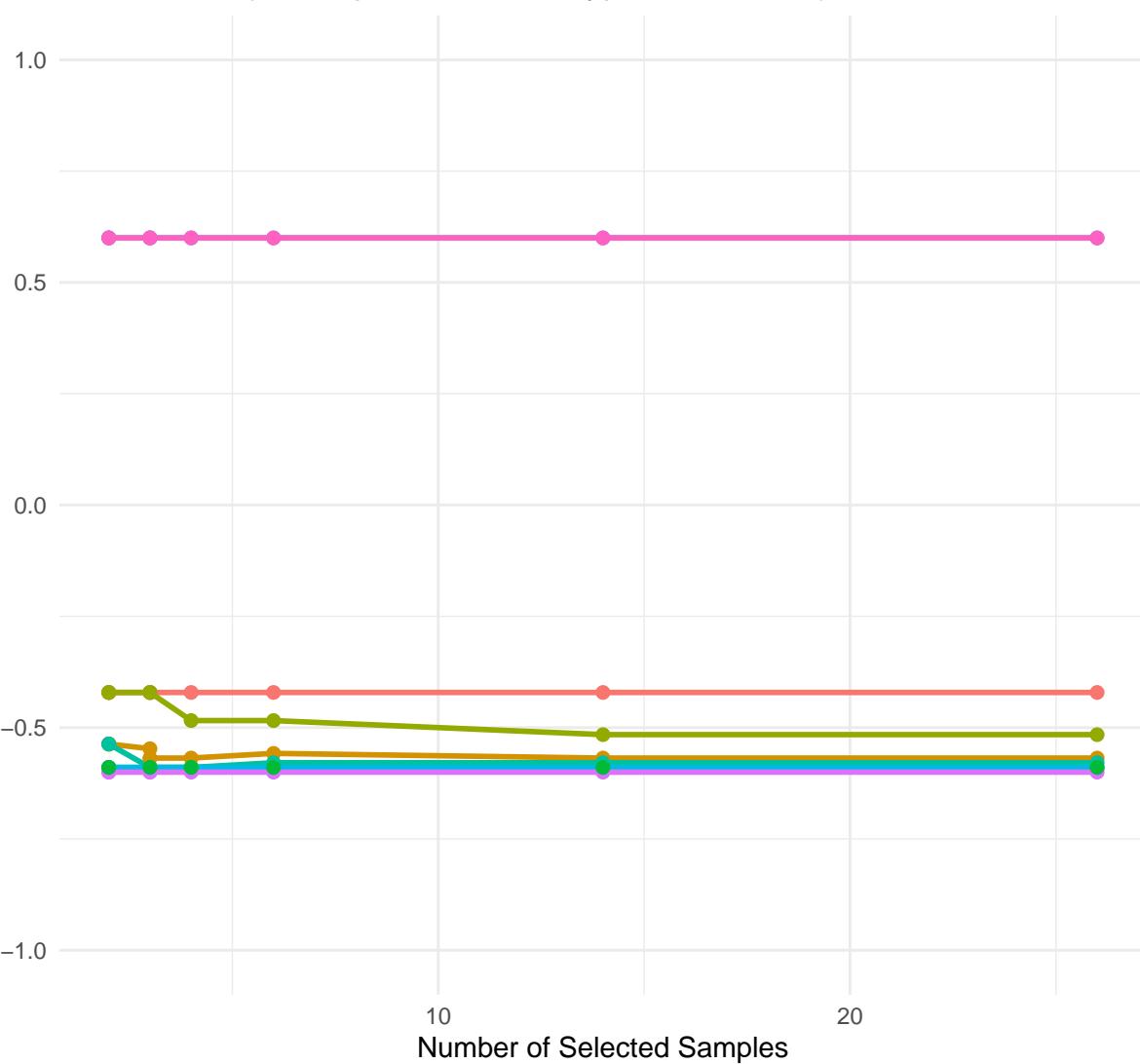
Summary Statistic

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



Score: ESP (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

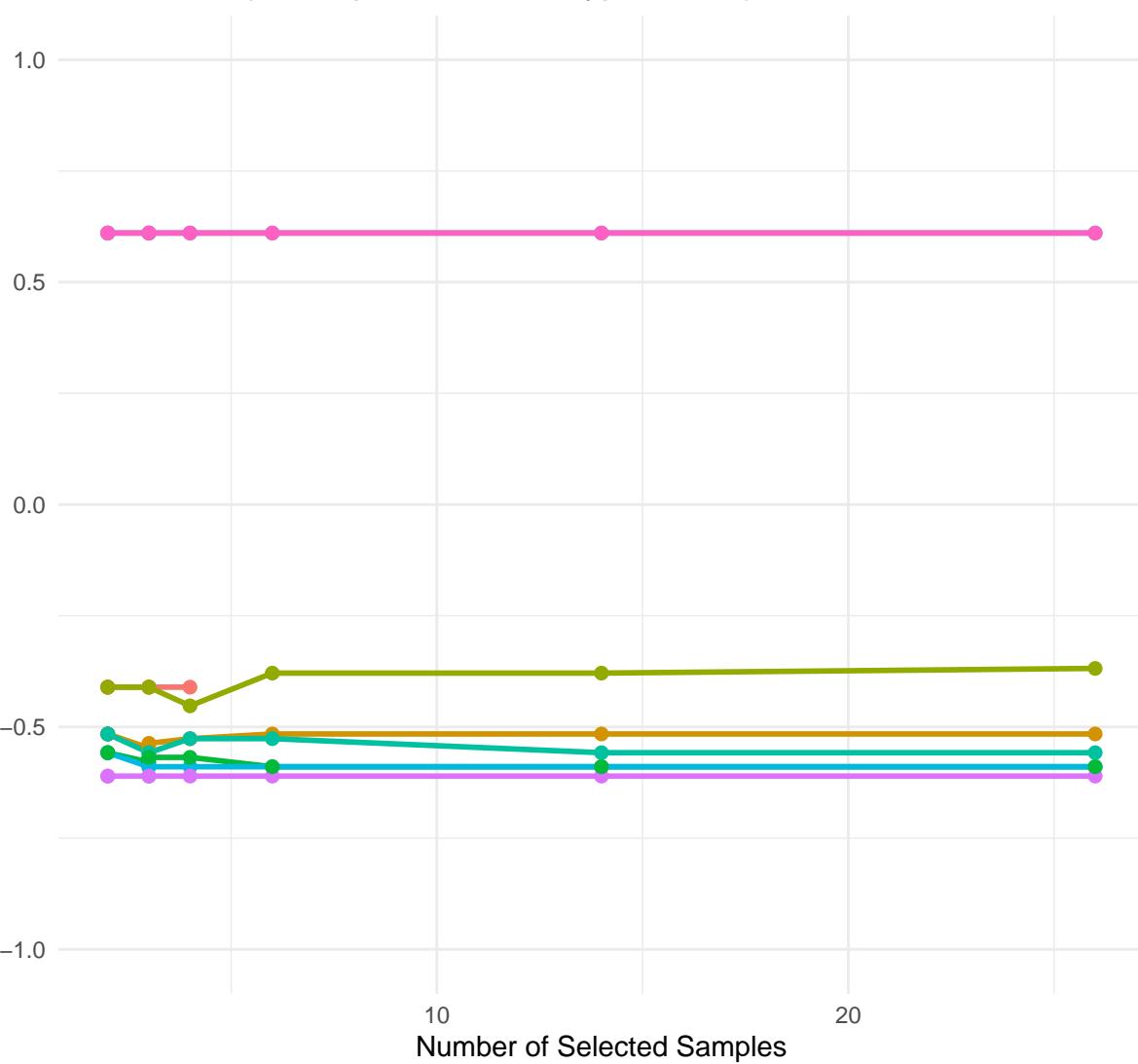


Summary Statistic

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

Score: ESP (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: CEV (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

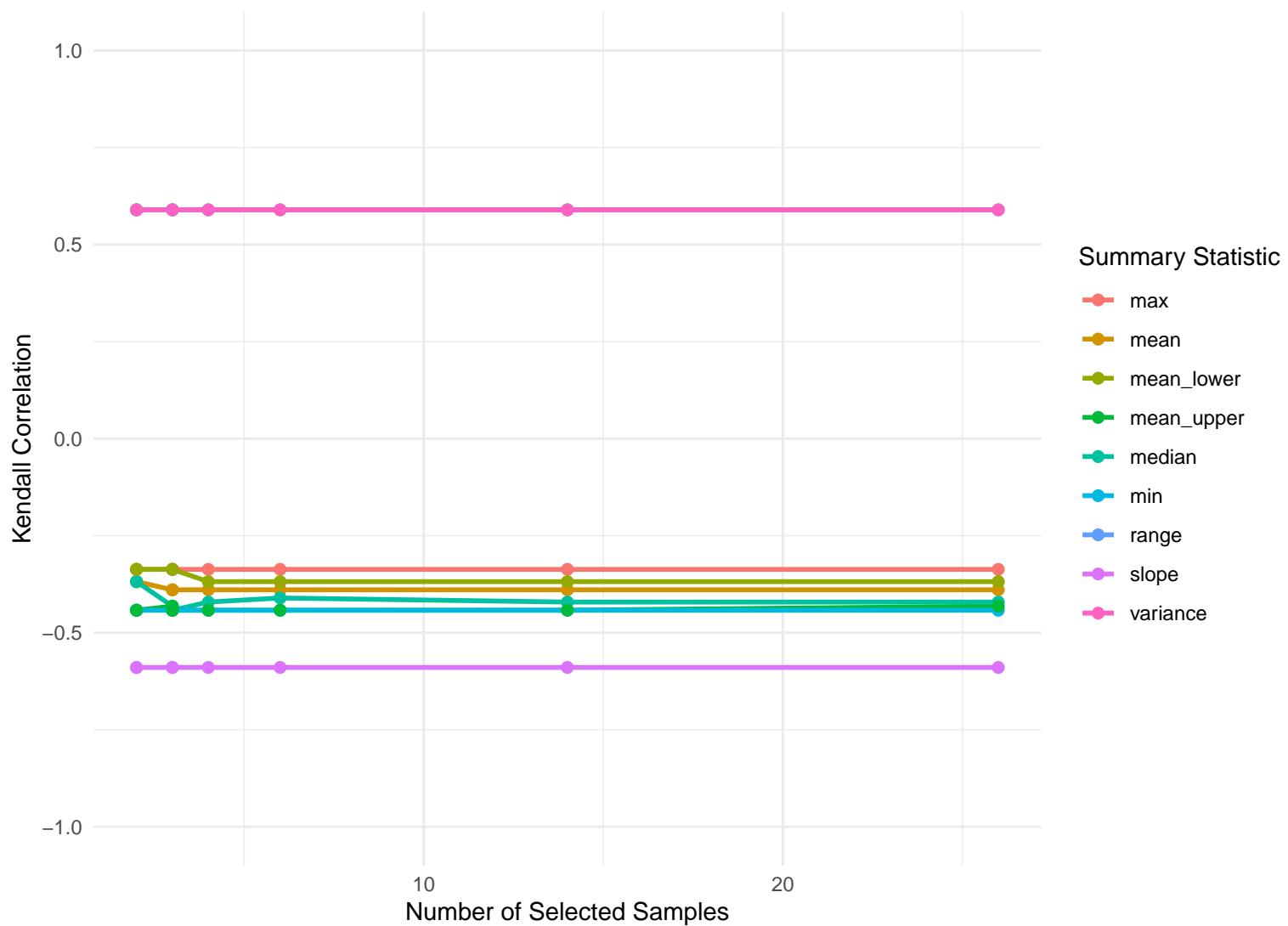
Summary Statistic

range

variance



Score: CEV (Local, partial2 , Genotype: gaussian)



Score: CEV (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

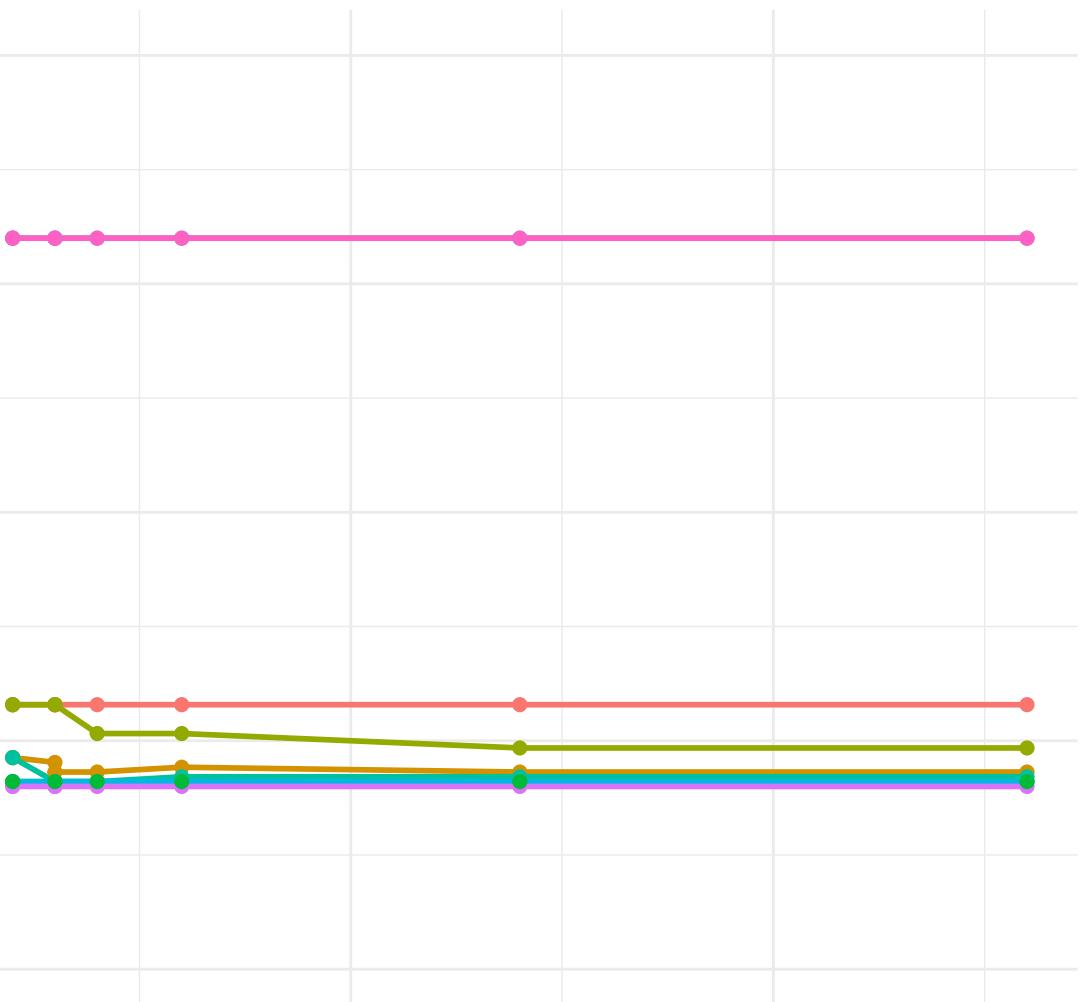
-0.5

-1.0

Number of Selected Samples

Summary Statistic

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



Score: CEV (Local, partial2 , Genotype: wave)

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

-0.45 -0.45 -0.45 -0.45 -0.45 -0.45 -0.45

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

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

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

Score: PFI (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

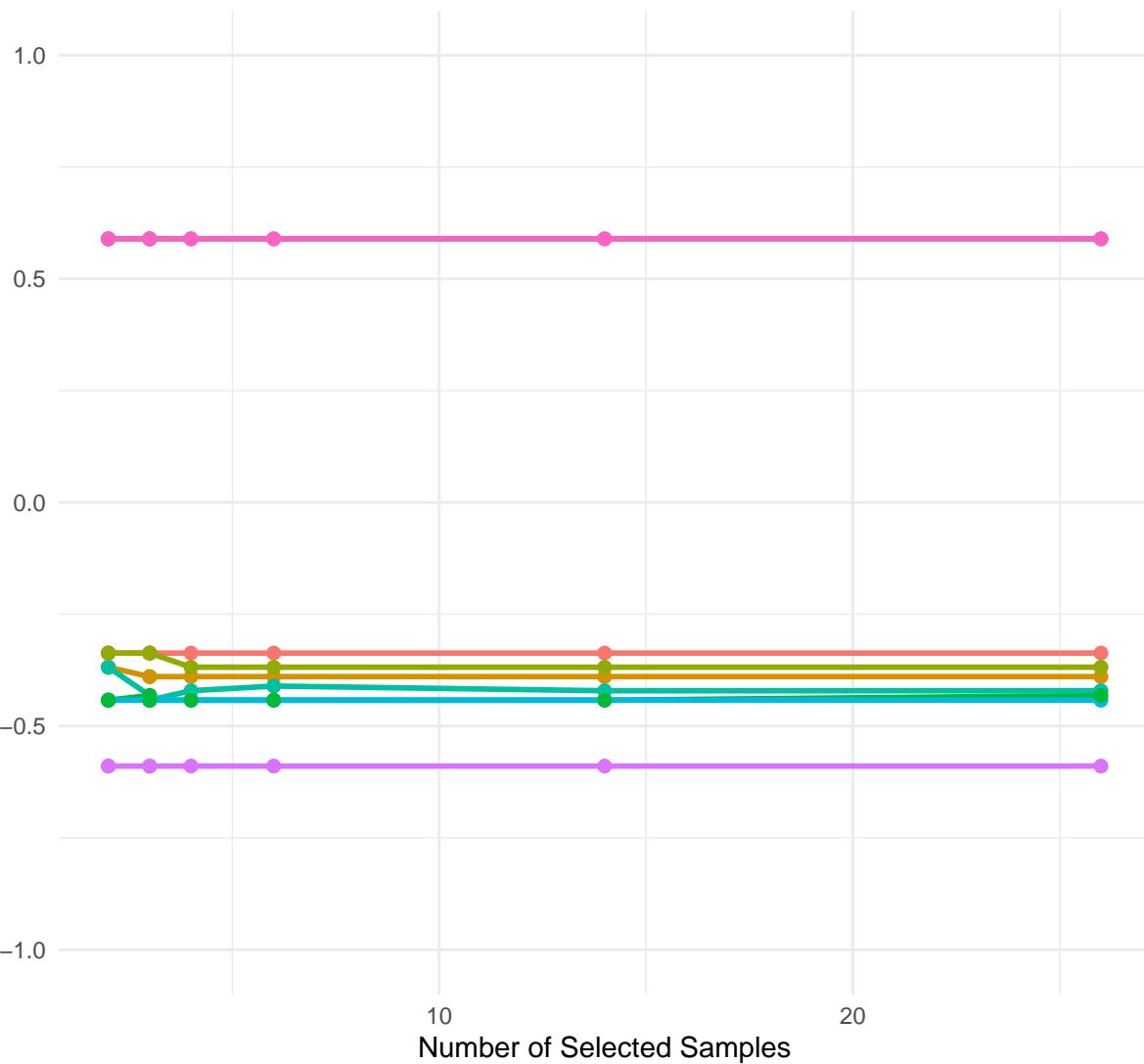
range

variance



Score: PFI (Local, partial2 , Genotype: gaussian)

Kendall Correlation

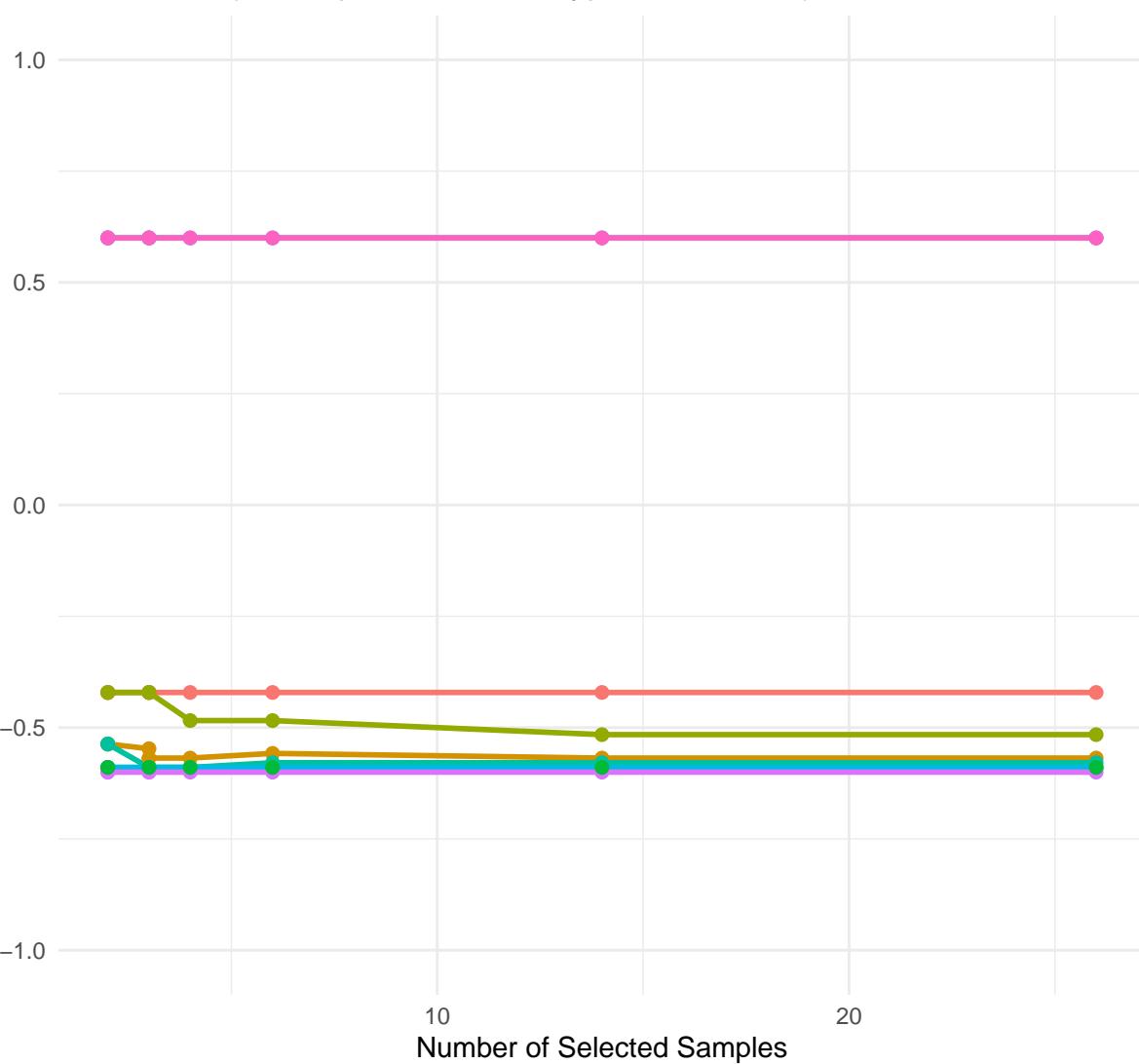


Summary Statistic

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

Score: PFI (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

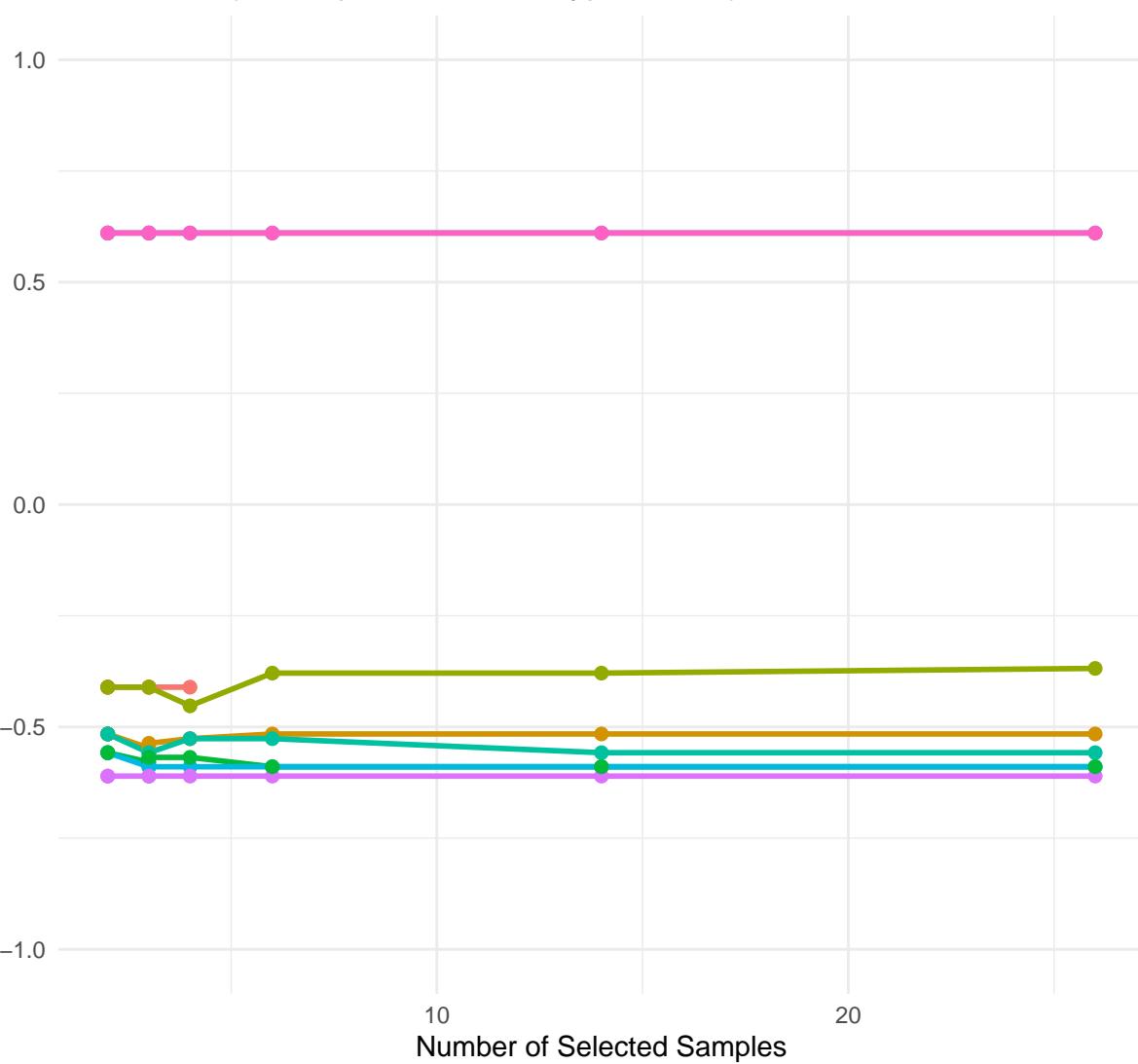


Summary Statistic

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

Score: PFI (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: APC (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

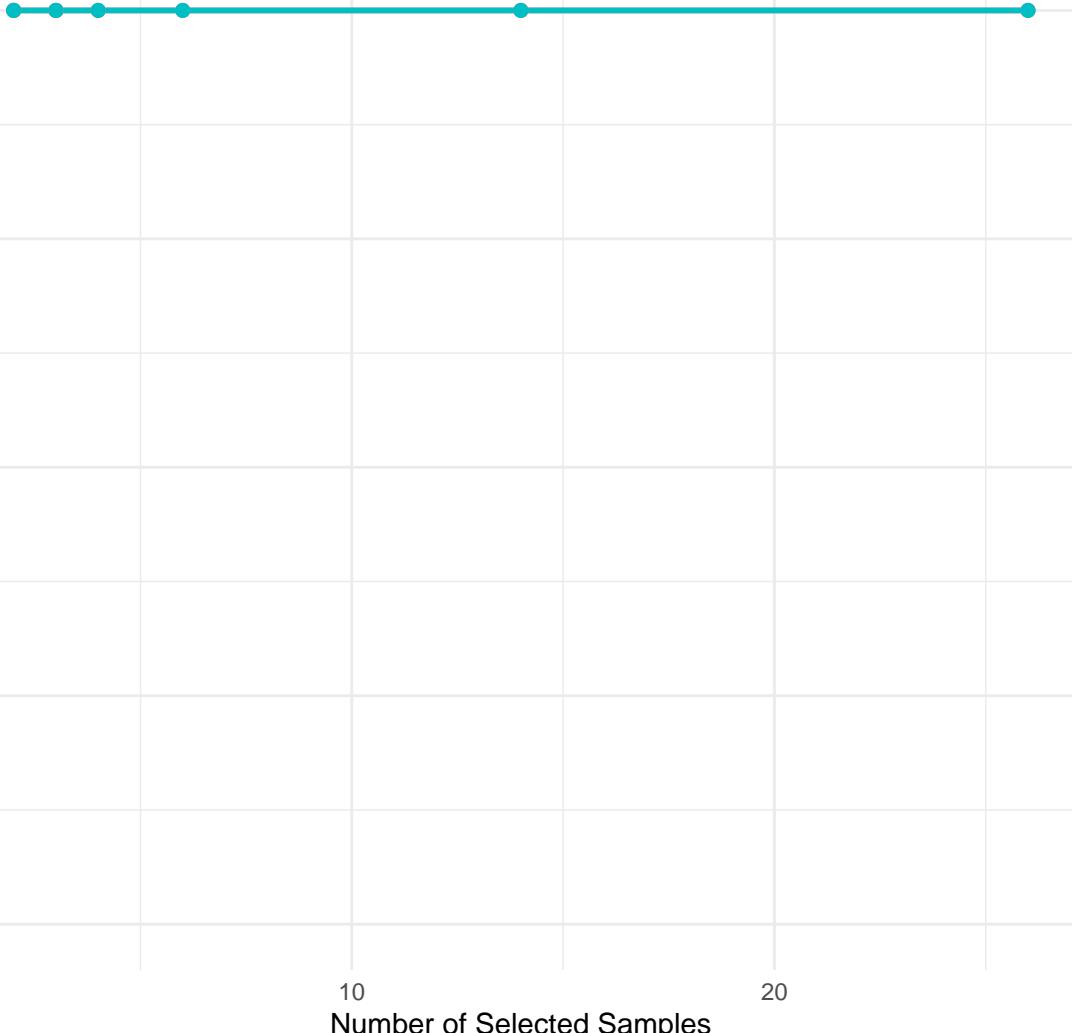
-1.0

Number of Selected Samples

Summary Statistic

range

variance



Score: APC (Local, partial2 , Genotype: gaussian)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

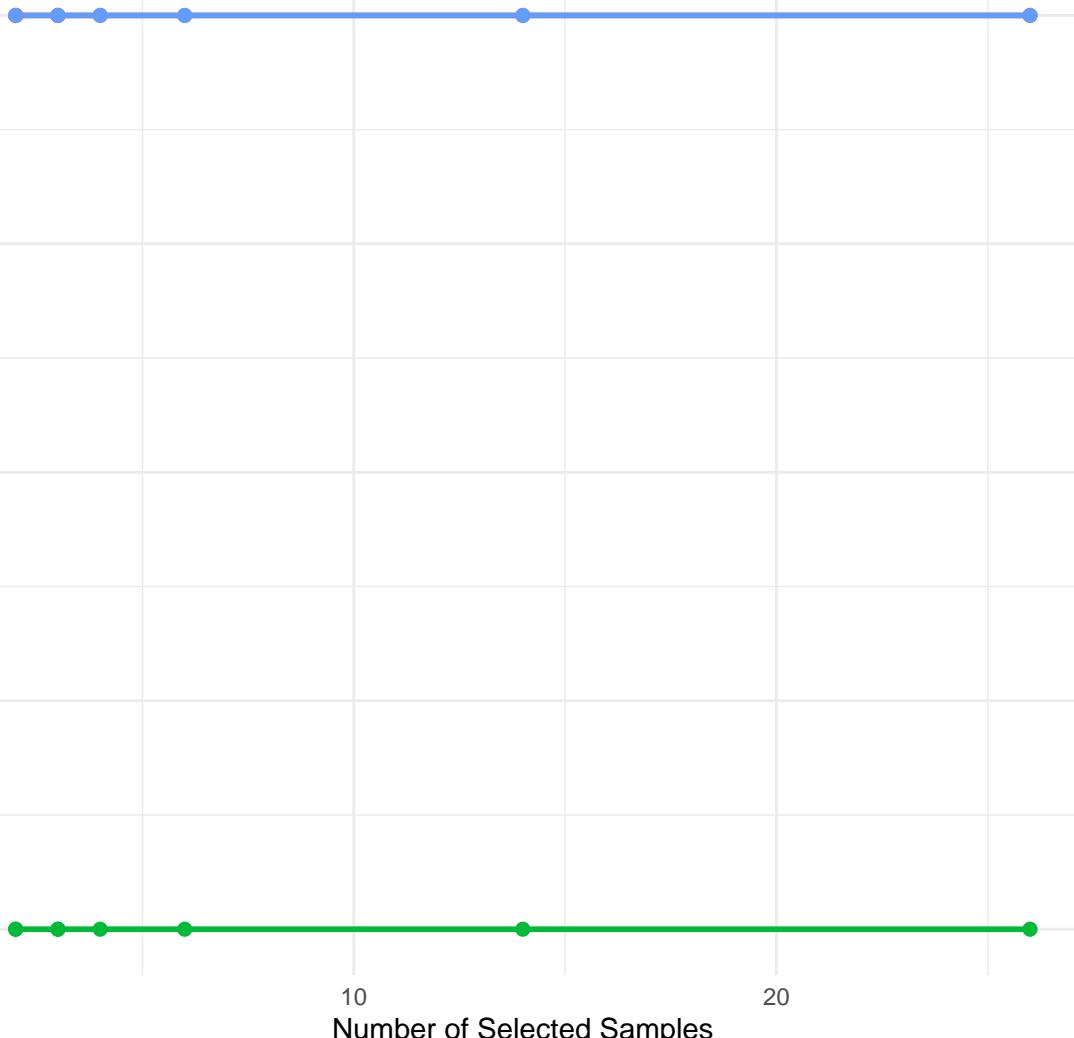
Number of Selected Samples

Summary Statistic

range

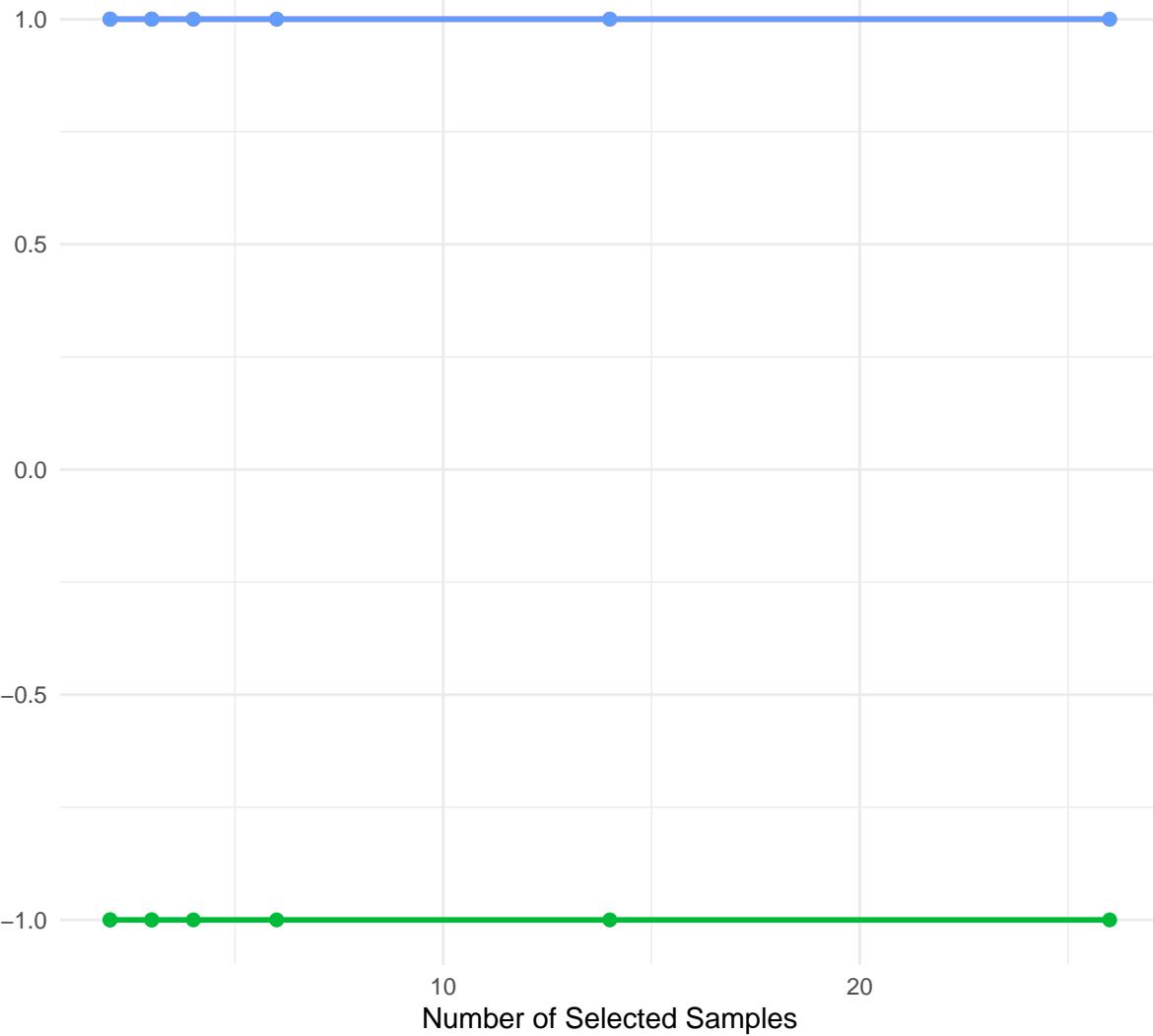
slope

variance



Score: APC (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation



Summary Statistic

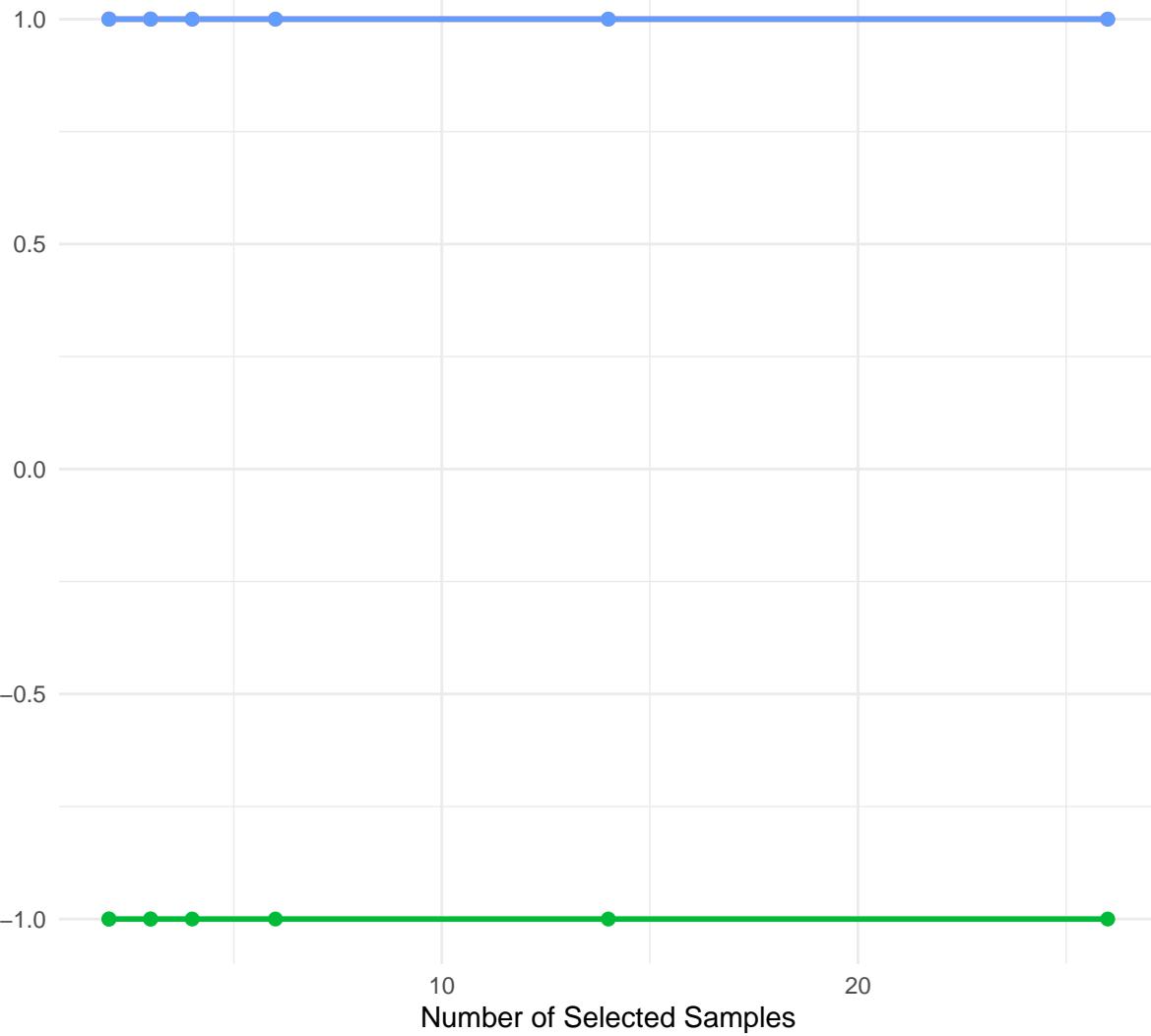
range

slope

variance

Score: APC (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

range

slope

variance

Number of Selected Samples

Score: SI (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

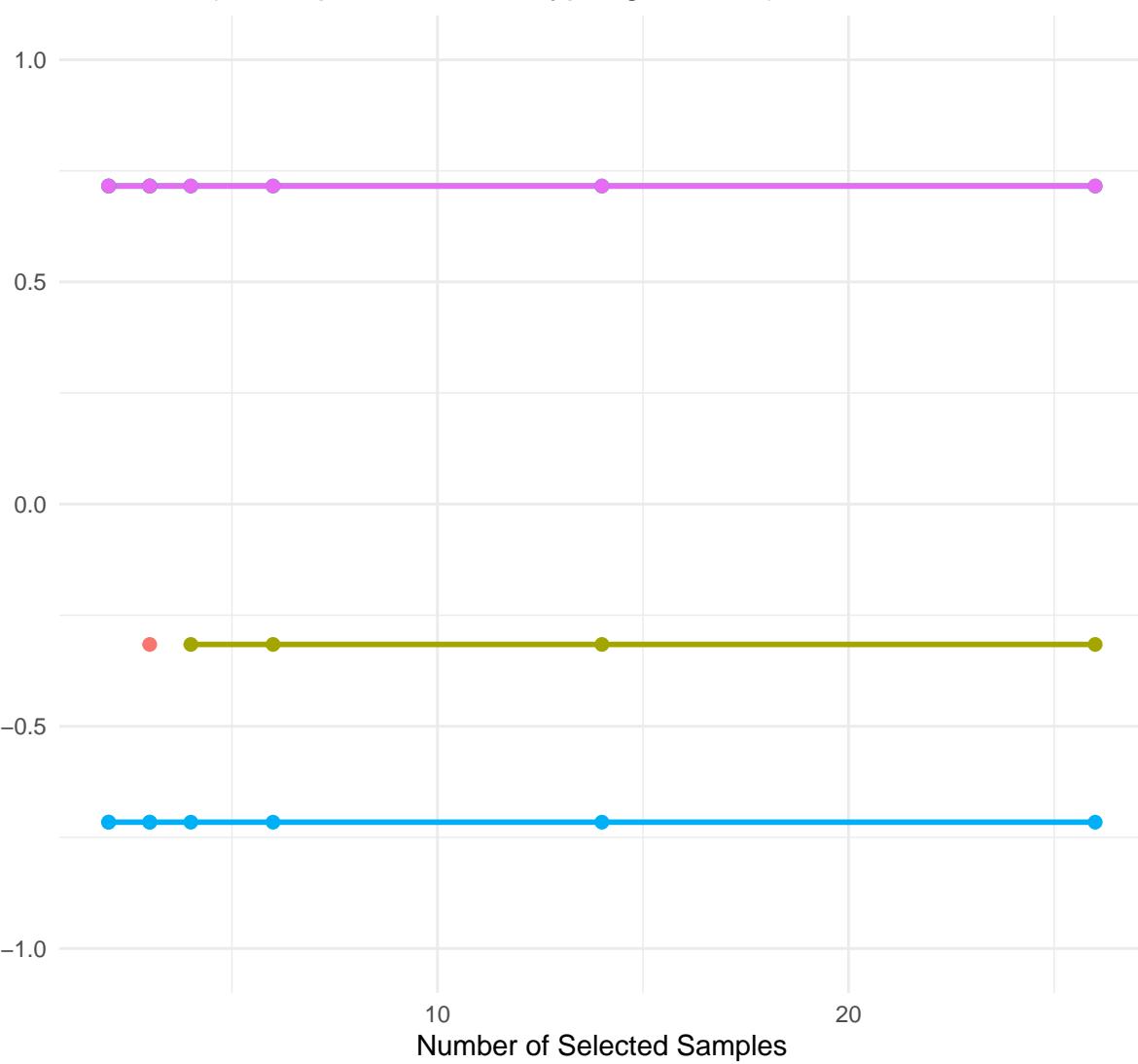
range

variance



Score: SI (Local, partial2 , Genotype: gaussian)

Kendall Correlation

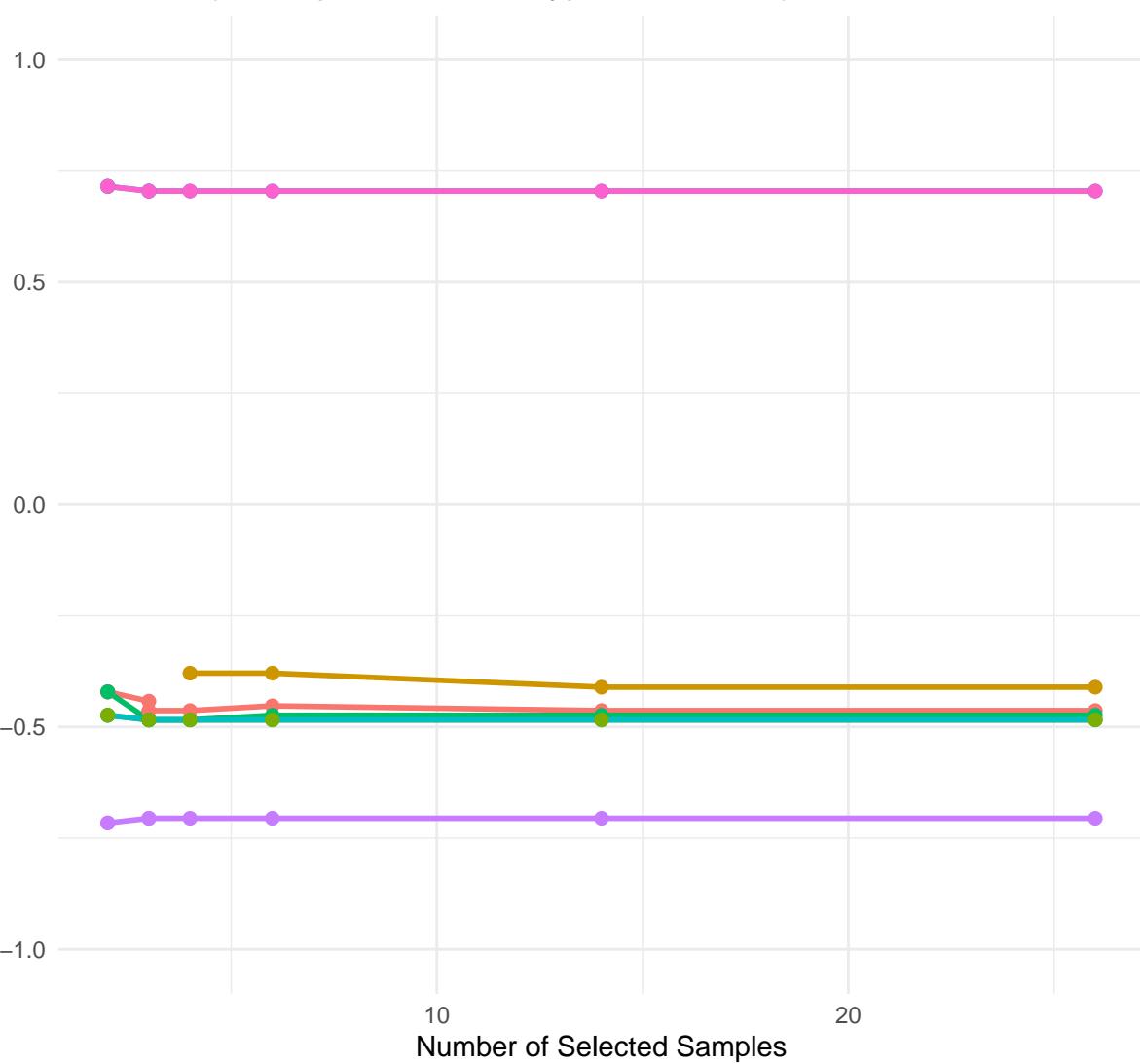


Summary Statistic

- median
- min
- range
- slope
- variance

Score: SI (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

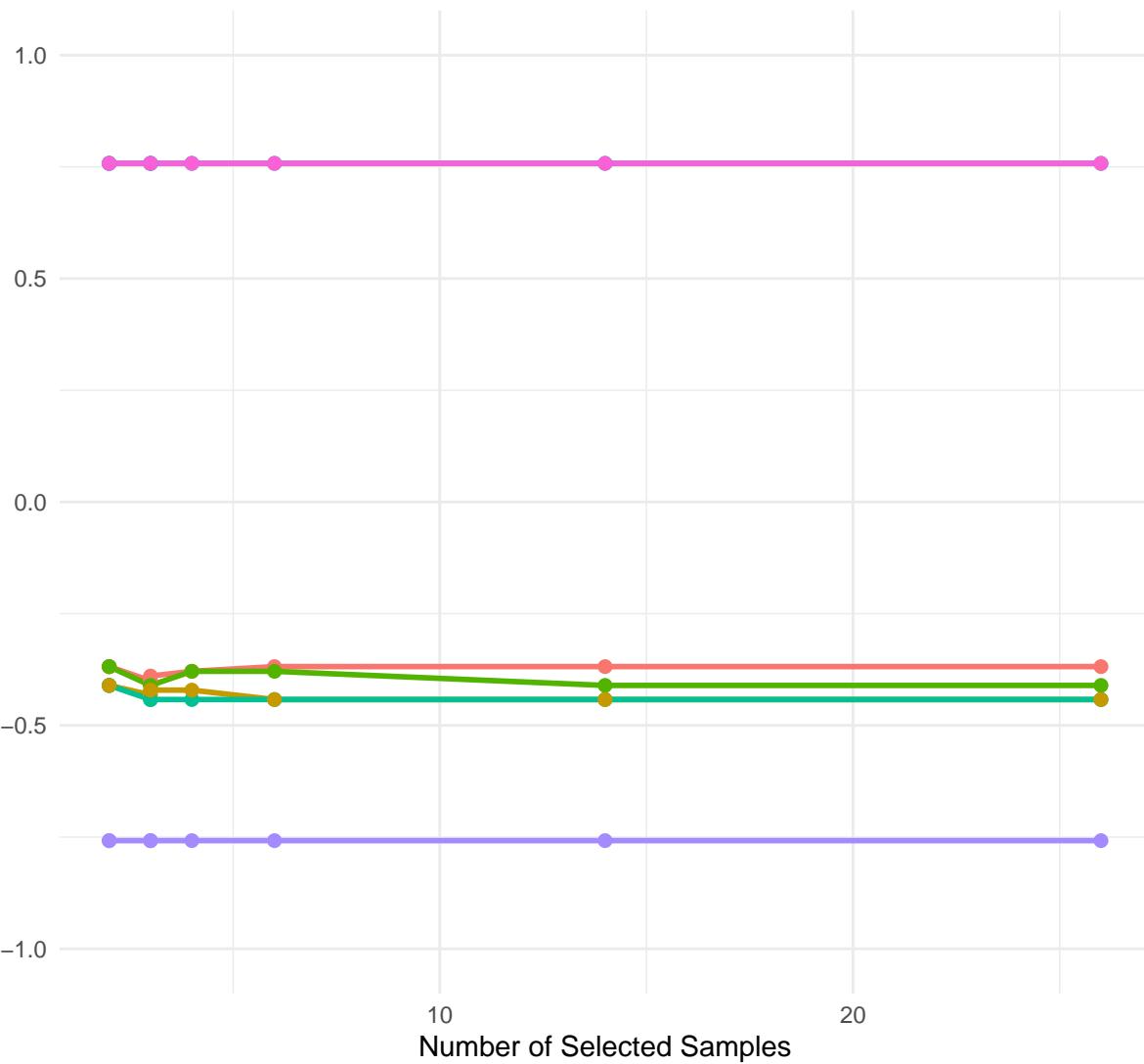


Summary Statistic

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

Score: SI (Local, partial2 , Genotype: wave)

Kendall Correlation



Summary Statistic

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

Score: RSI (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

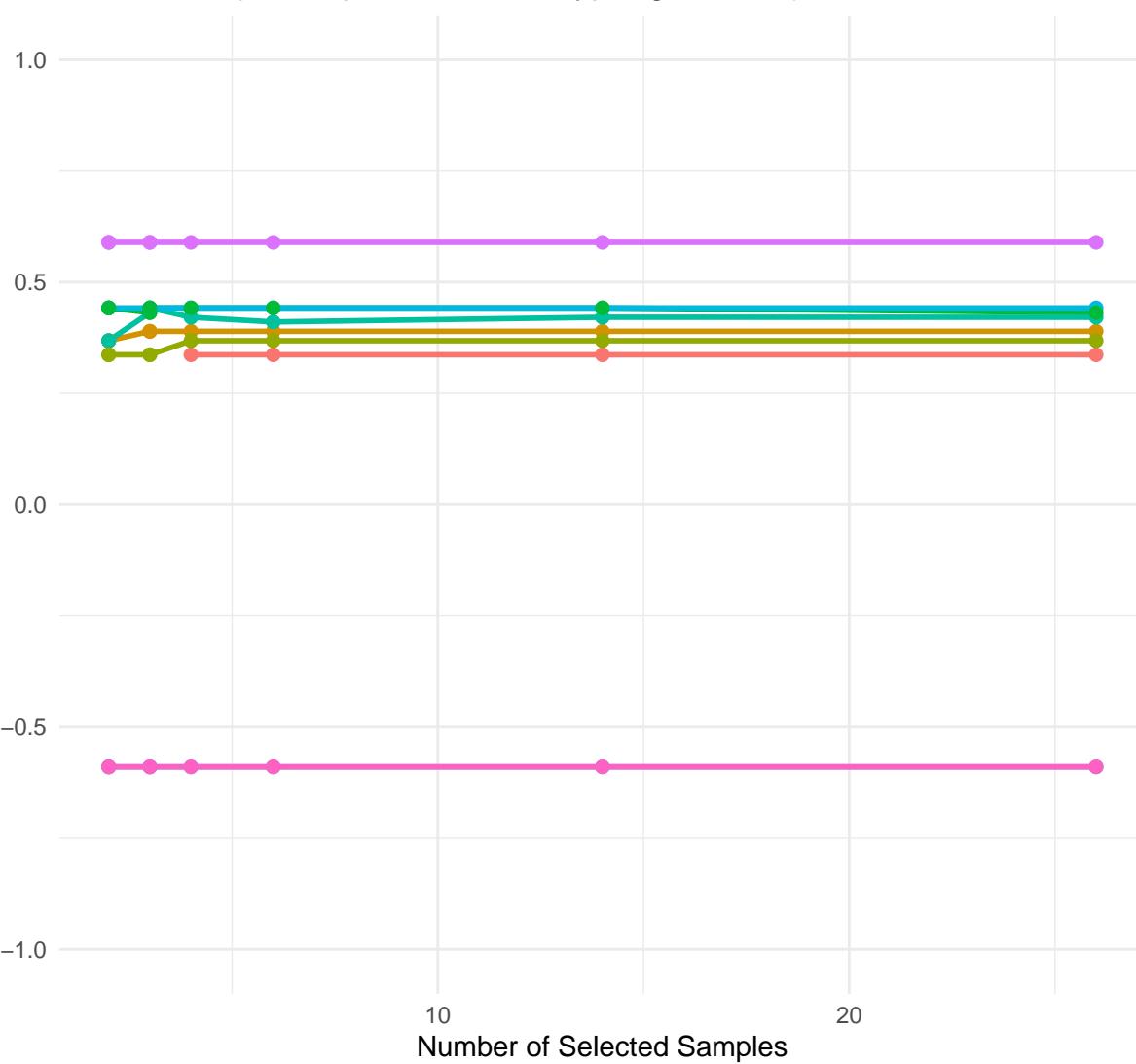
range

variance



Score: RSI (Local, partial2 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

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

Score: RSI (Local, partial2 , 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

0.55 0.55 0.55 0.55 0.55 0.55 0.55 0.55 0.55

0.52 0.52 0.52 0.52 0.52 0.52 0.52 0.52 0.52

0.50 0.50 0.50 0.50 0.50 0.50 0.50 0.50 0.50

0.48 0.48 0.48 0.48 0.48 0.48 0.48 0.48 0.48

0.46 0.46 0.46 0.46 0.46 0.46 0.46 0.46 0.46

0.44 0.44 0.44 0.44 0.44 0.44 0.44 0.44 0.44

0.42 0.42 0.42 0.42 0.42 0.42 0.42 0.42 0.42

0.40 0.40 0.40 0.40 0.40 0.40 0.40 0.40 0.40

0.38 0.38 0.38 0.38 0.38 0.38 0.38 0.38 0.38

0.36 0.36 0.36 0.36 0.36 0.36 0.36 0.36 0.36

0.34 0.34 0.34 0.34 0.34 0.34 0.34 0.34 0.34

0.32 0.32 0.32 0.32 0.32 0.32 0.32 0.32 0.32

0.30 0.30 0.30 0.30 0.30 0.30 0.30 0.30 0.30

0.28 0.28 0.28 0.28 0.28 0.28 0.28 0.28 0.28

0.26 0.26 0.26 0.26 0.26 0.26 0.26 0.26 0.26

0.24 0.24 0.24 0.24 0.24 0.24 0.24 0.24 0.24

0.22 0.22 0.22 0.22 0.22 0.22 0.22 0.22 0.22

0.20 0.20 0.20 0.20 0.20 0.20 0.20 0.20 0.20

0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18

0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16

0.14 0.14 0.14 0.14 0.14 0.14 0.14 0.14 0.14

0.12 0.12 0.12 0.12 0.12 0.12 0.12 0.12 0.12

0.10 0.10 0.10 0.10 0.10 0.10 0.10 0.10 0.10

0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.08

0.06 0.06 0.06 0.06 0.06 0.06 0.06 0.06 0.06

0.04 0.04 0.04 0.04 0.04 0.04 0.04 0.04 0.04

0.02 0.02 0.02 0.02 0.02 0.02 0.02 0.02 0.02

0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.00

-0.02 -0.02 -0.02 -0.02 -0.02 -0.02 -0.02 -0.02 -0.02

-0.04 -0.04 -0.04 -0.04 -0.04 -0.04 -0.04 -0.04 -0.04

-0.06 -0.06 -0.06 -0.06 -0.06 -0.06 -0.06 -0.06 -0.06

-0.08 -0.08 -0.08 -0.08 -0.08 -0.08 -0.08 -0.08 -0.08

-0.10 -0.10 -0.10 -0.10 -0.10 -0.10 -0.10 -0.10 -0.10

-0.12 -0.12 -0.12 -0.12 -0.12 -0.12 -0.12 -0.12 -0.12

-0.14 -0.14 -0.14 -0.14 -0.14 -0.14 -0.14 -0.14 -0.14

-0.16 -0.16 -0.16 -0.16 -0.16 -0.16 -0.16 -0.16 -0.16

-0.18 -0.18 -0.18 -0.18 -0.18 -0.18 -0.18 -0.18 -0.18

-0.20 -0.20 -0.20 -0.20 -0.20 -0.20 -0.20 -0.20 -0.20

Score: RSI (Local, partial2 , Genotype: wave)

Kendall Correlation

1.0

0.5

0.0

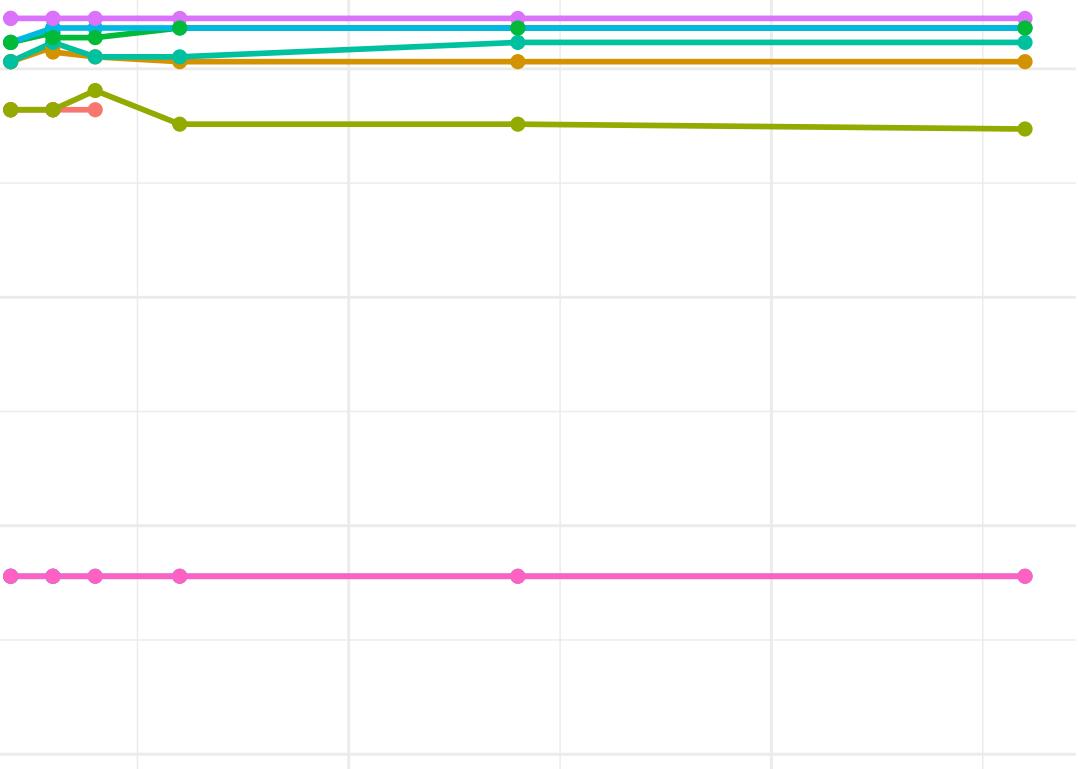
-0.5

-1.0

Number of Selected Samples

Summary Statistic

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



Score: EVS (Local, partial2 , Genotype: linear)

Kendall Correlation

1.0

0.5

0.0

-0.5

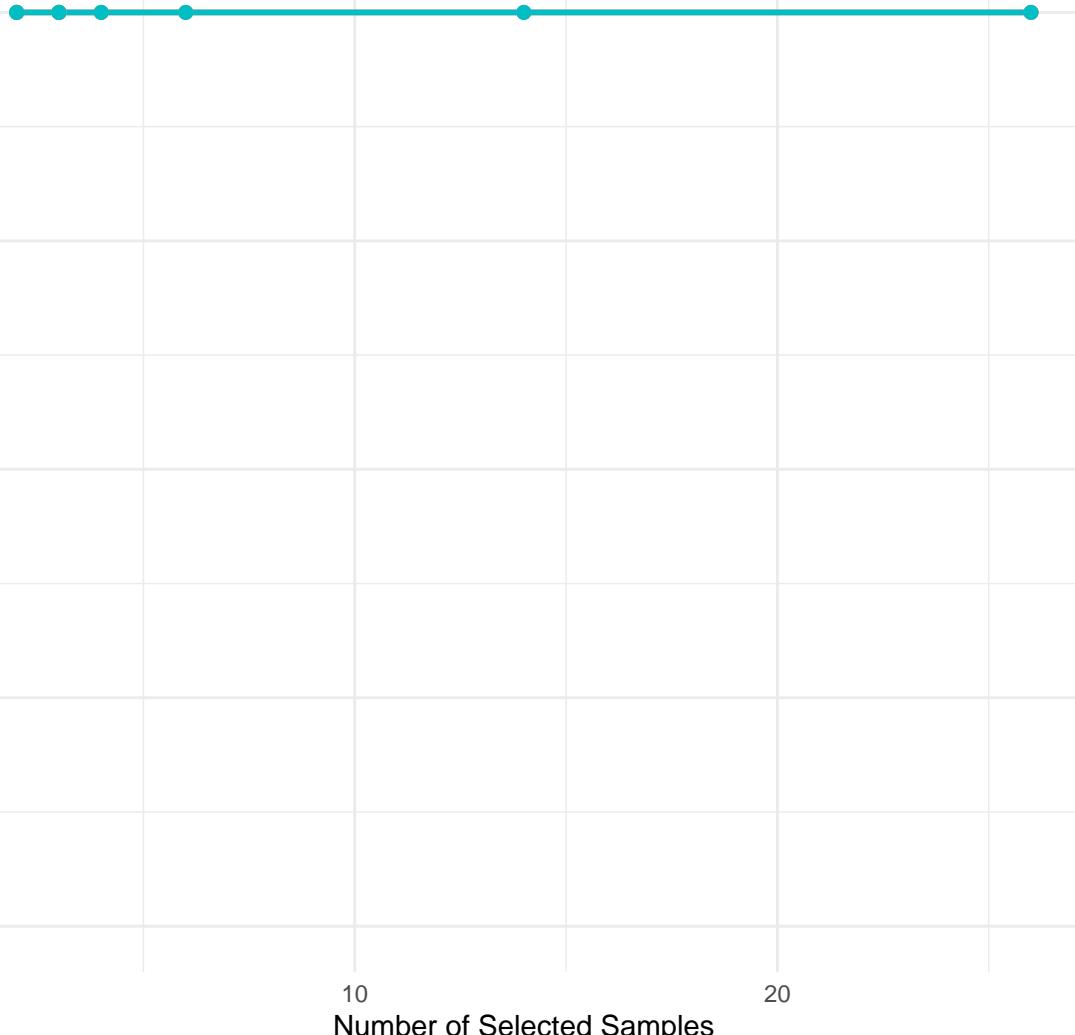
-1.0

Number of Selected Samples

Summary Statistic

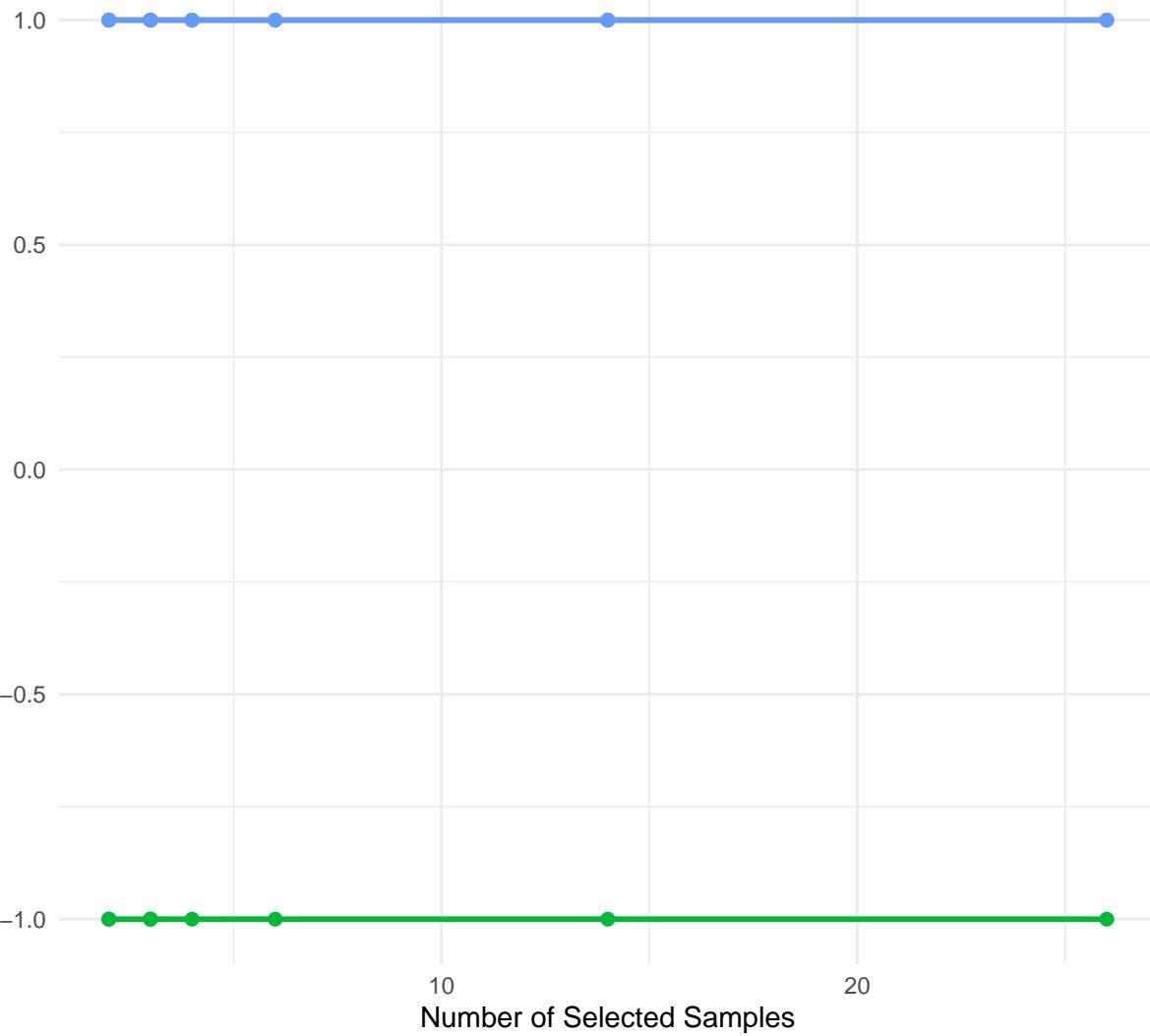
range

variance



Score: EVS (Local, partial2 , Genotype: gaussian)

Kendall Correlation



Summary Statistic

range

slope

variance

Number of Selected Samples

Score: EVS (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

Summary Statistic

range

slope

variance



Score: EVS (Local, partial2 , Genotype: wave)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Summary Statistic

range

slope

variance

Number of Selected Samples