

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

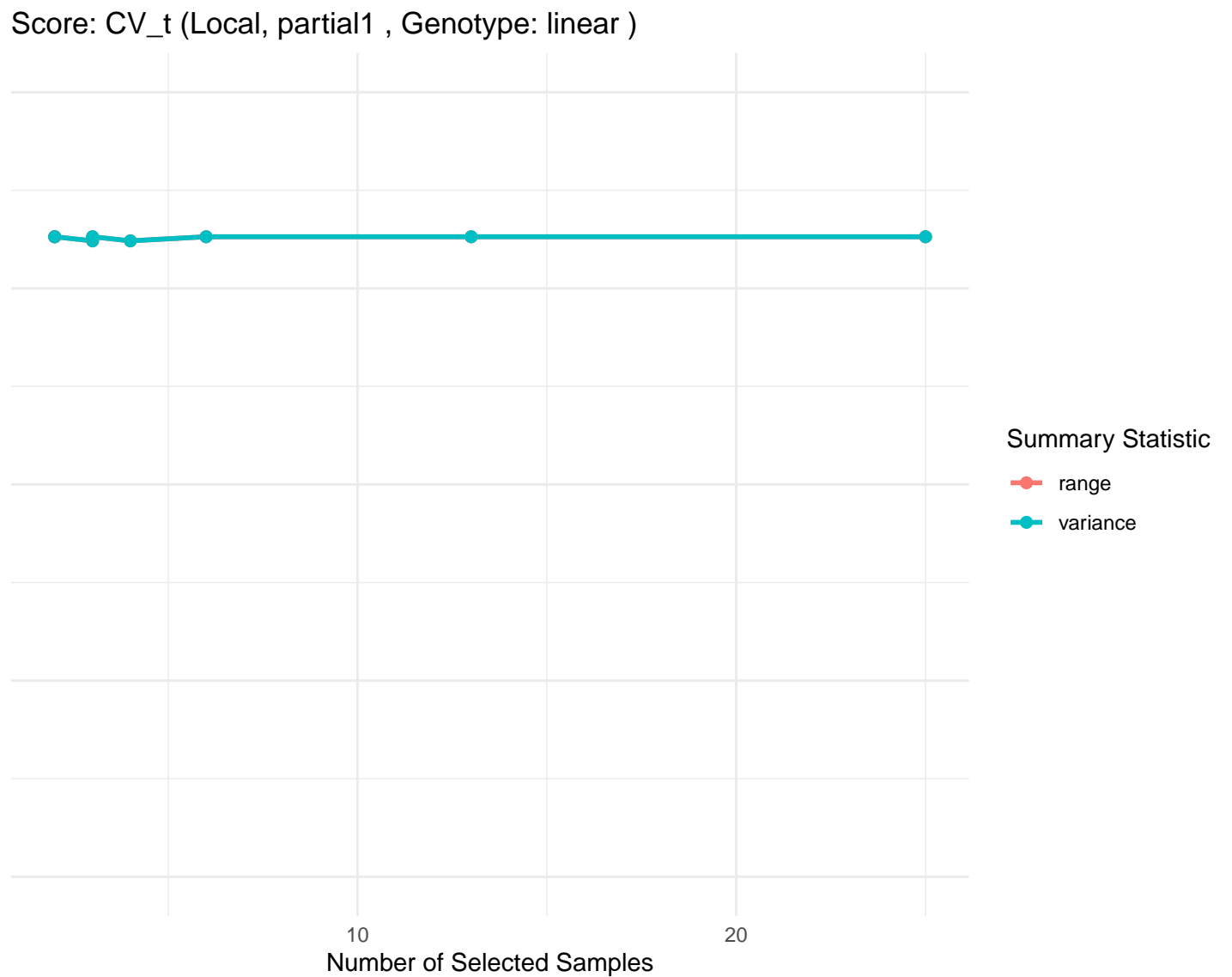
Kendall Correlation

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

1.0

0.5

0.0

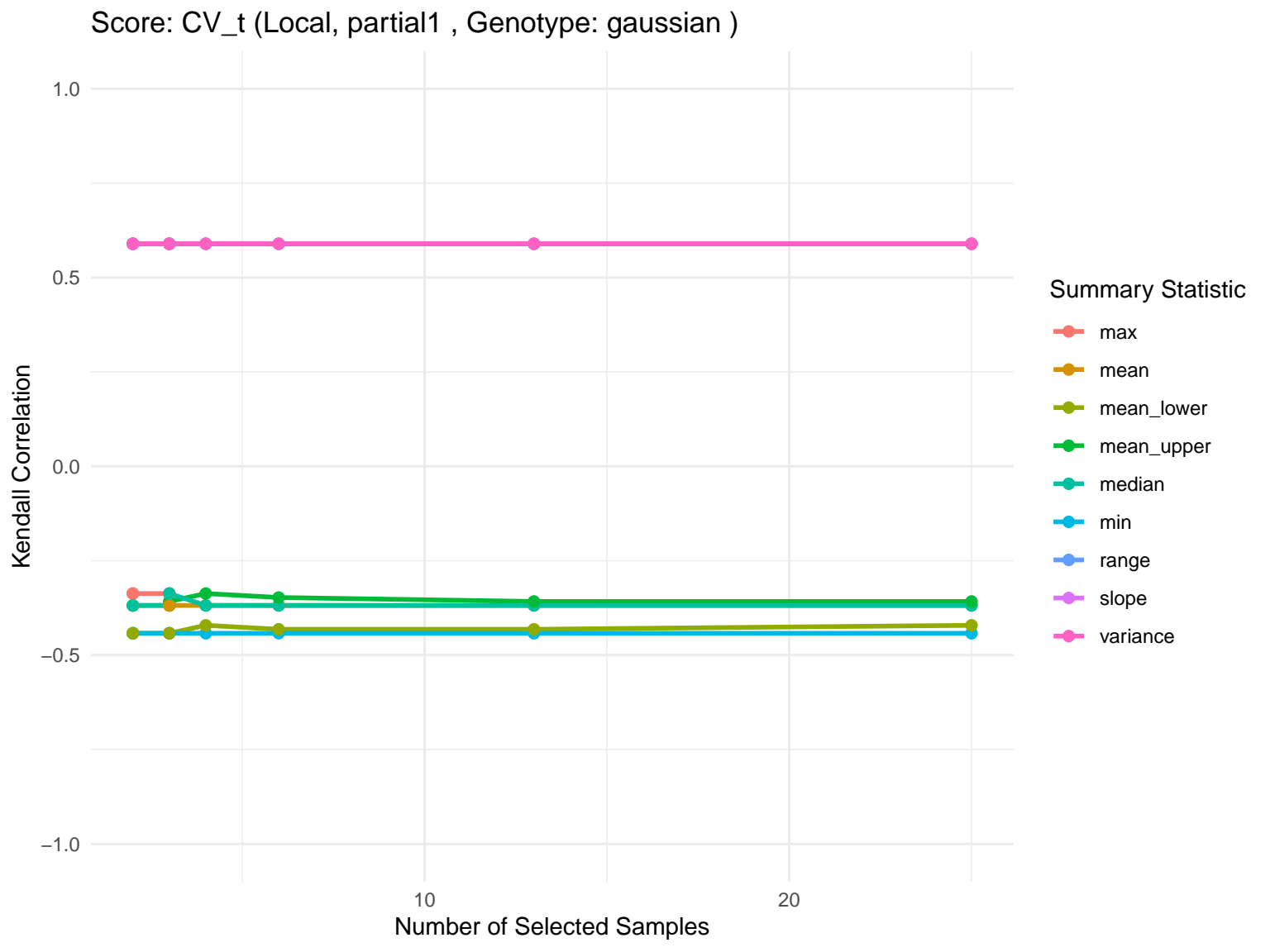
-0.5

-1.0

Number of Selected Samples

10

20



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



Summary Statistic

max

mean

mean_lower

mean_upper

median

min

range

slope

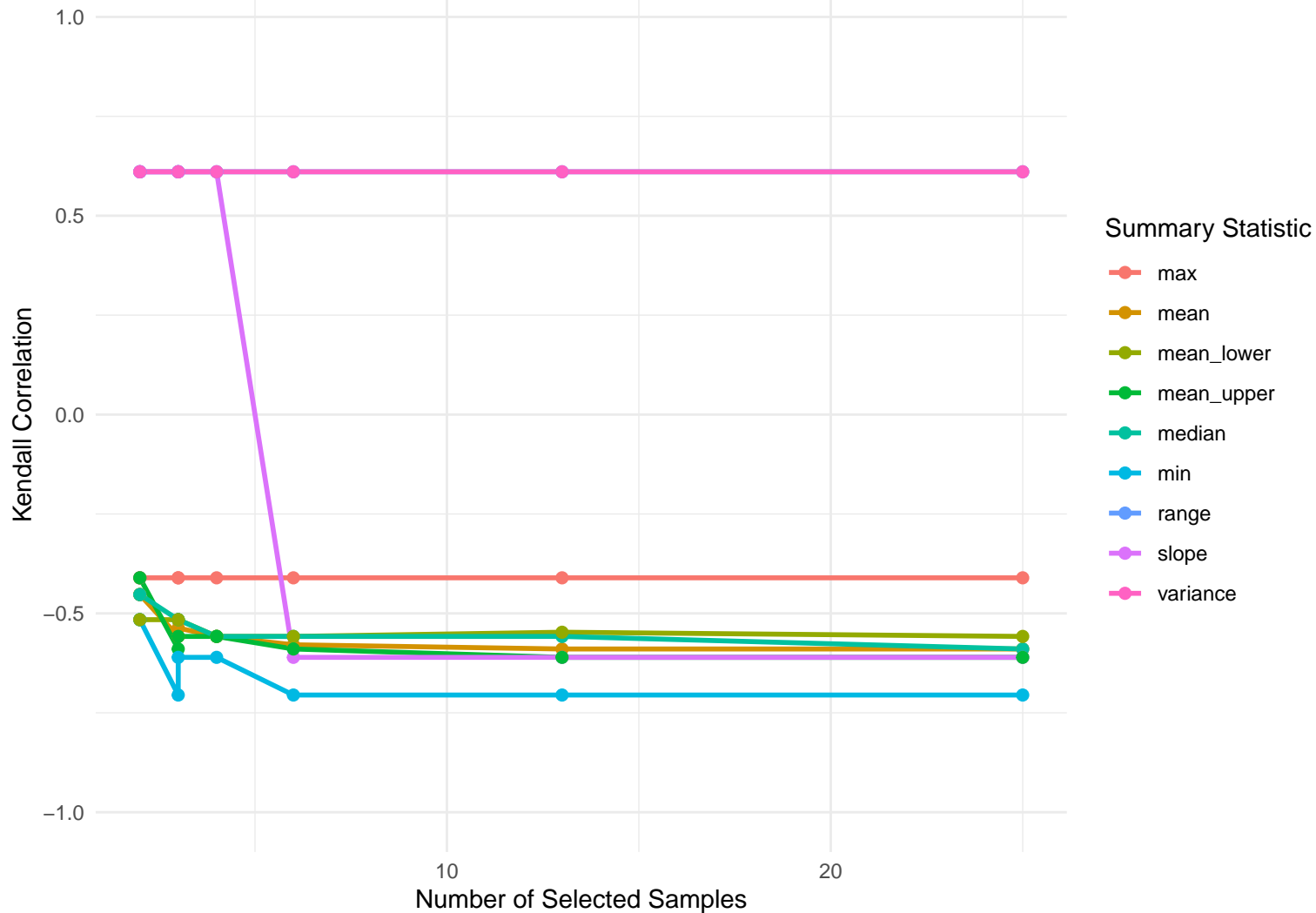
variance

Number of Selected Samples

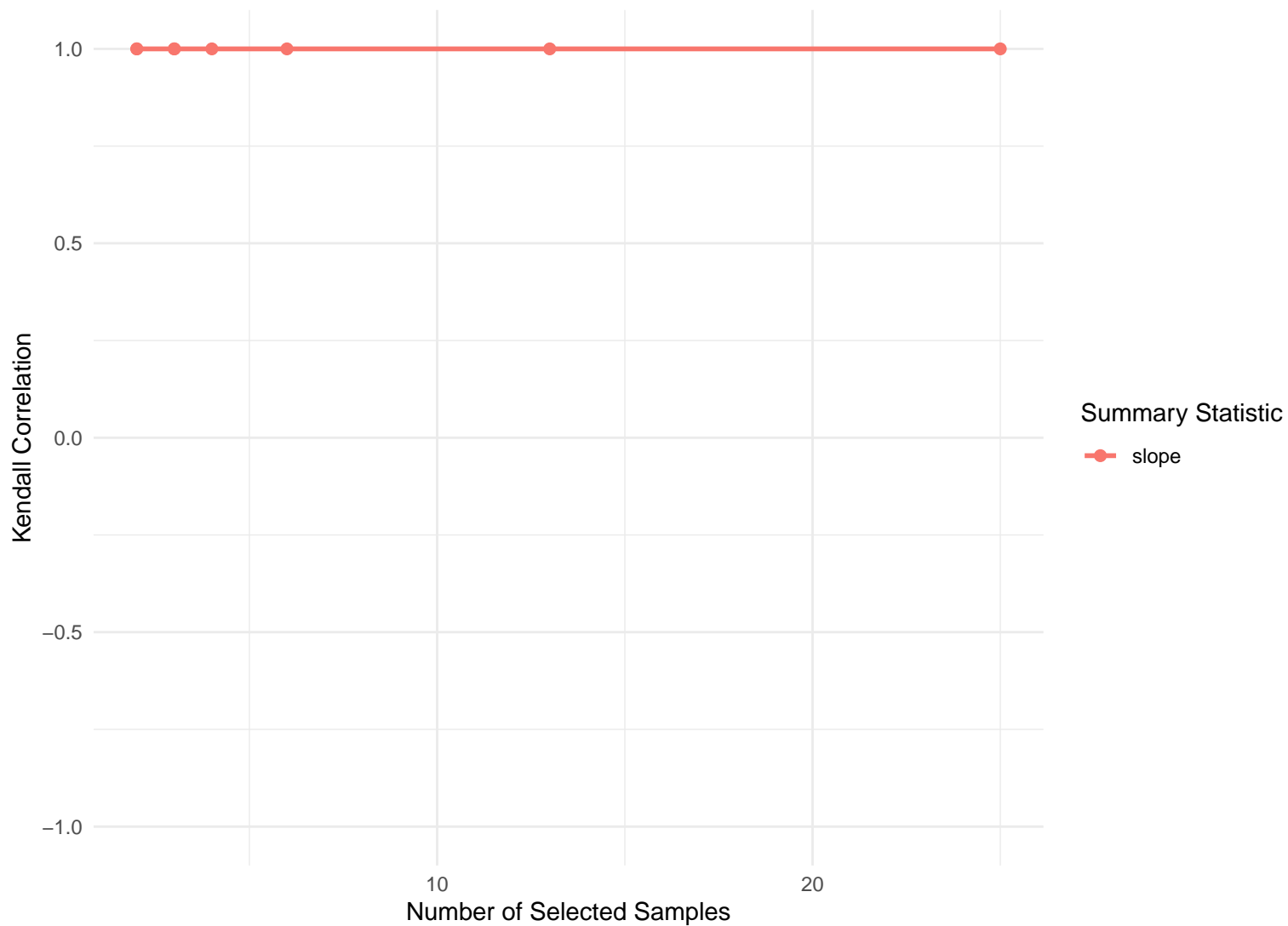
10

20

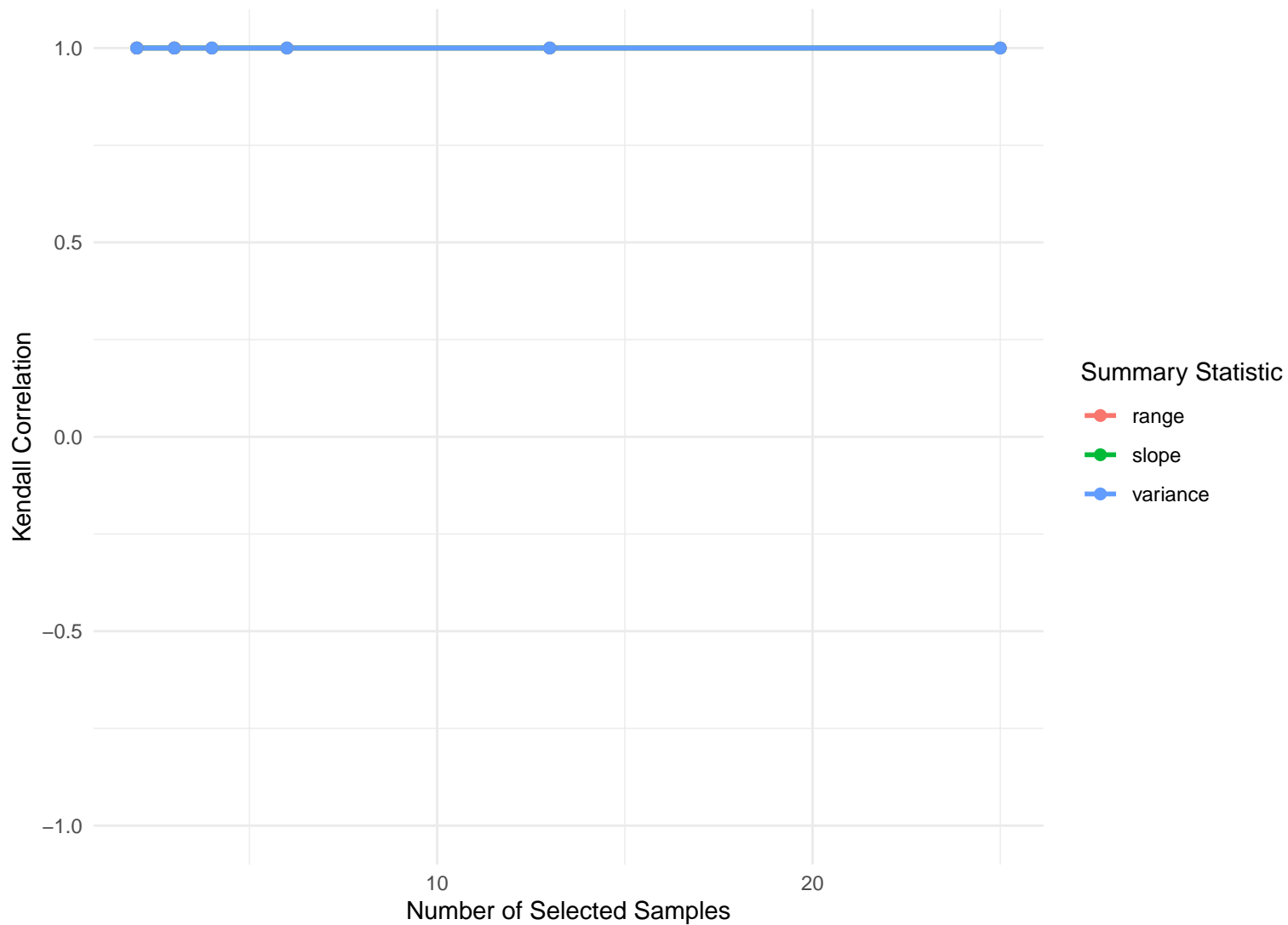
Score: CV_t (Local, partial1 , Genotype: wave)



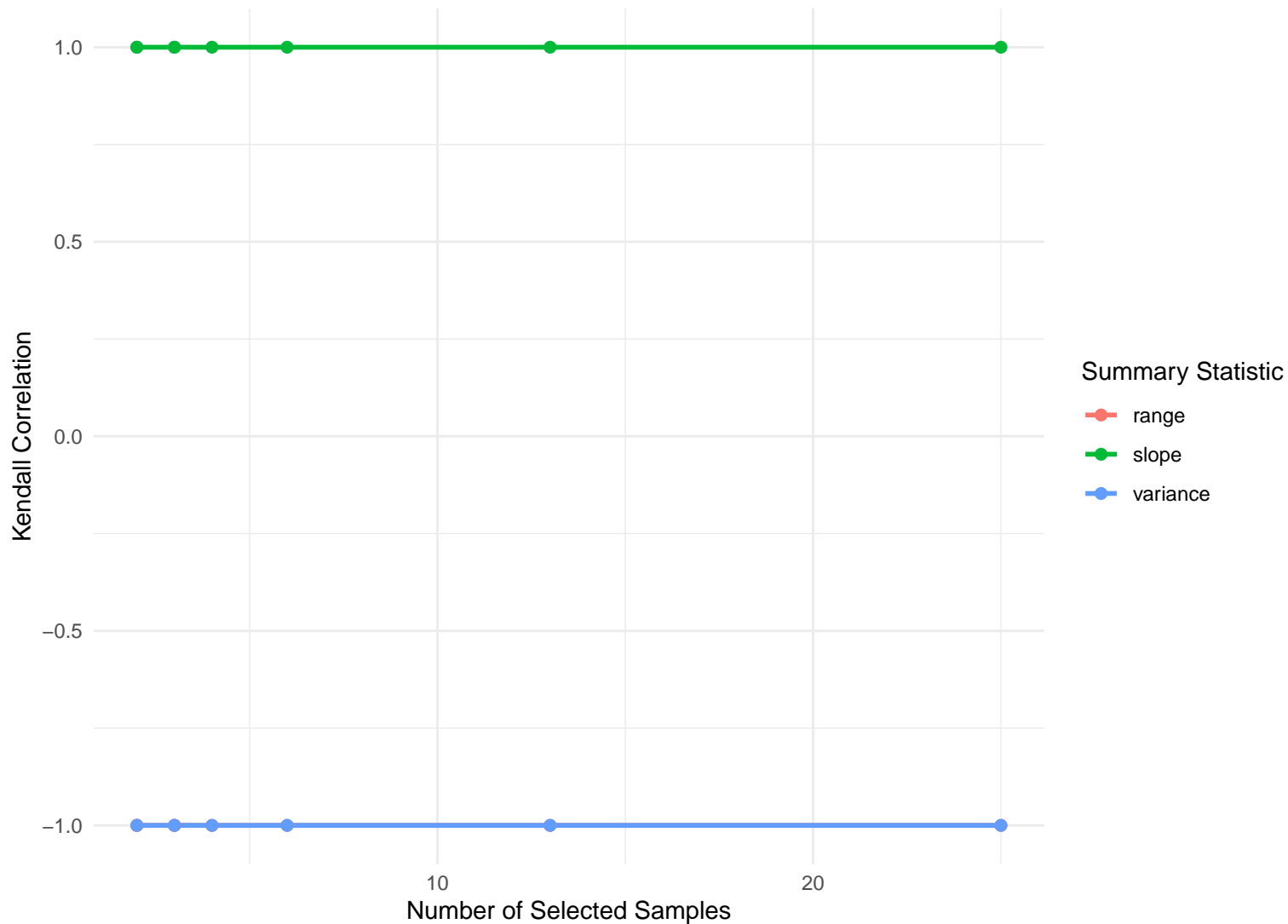
Score: RN (Local, partial1 , Genotype: linear)



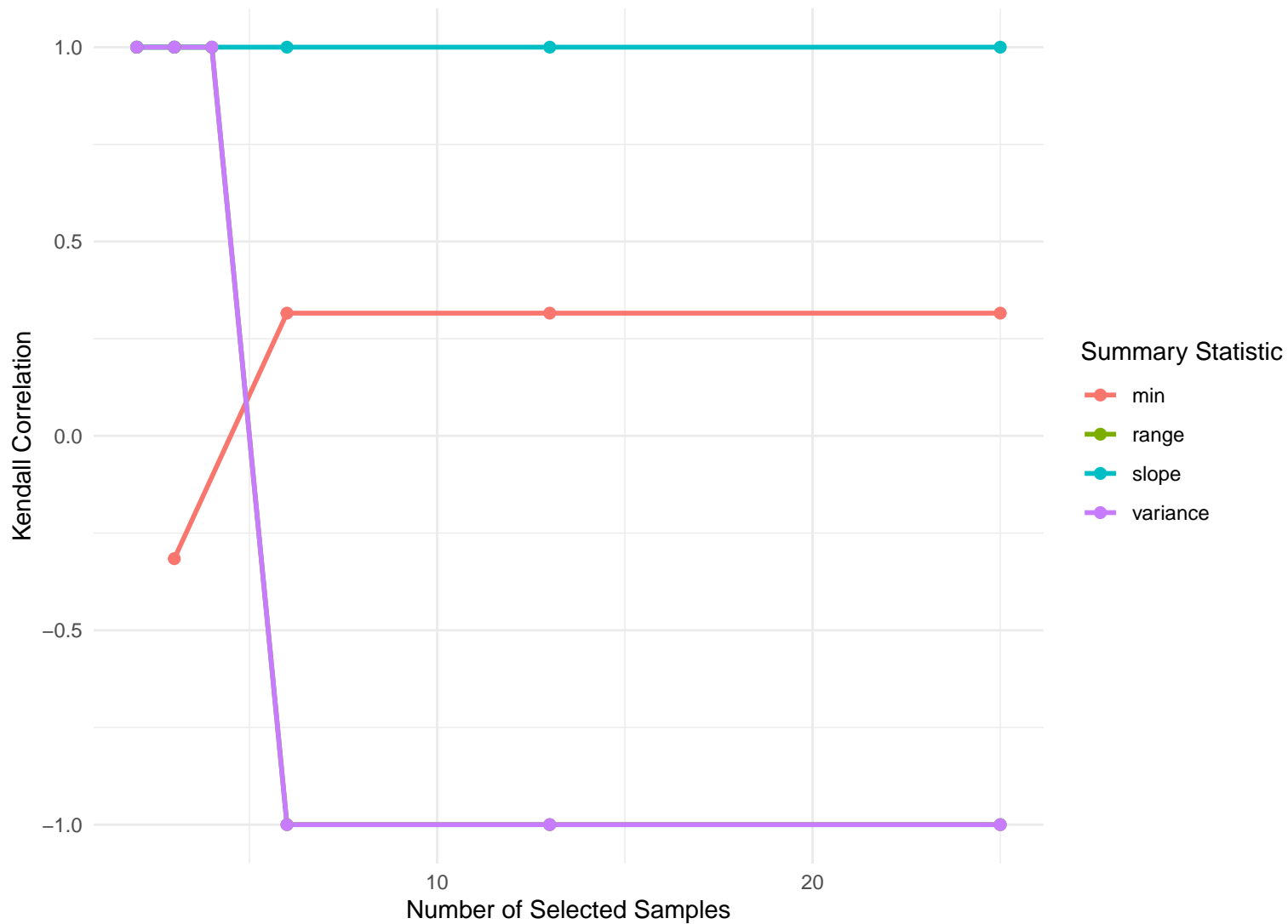
Score: RN (Local, partial1 , Genotype: gaussian)



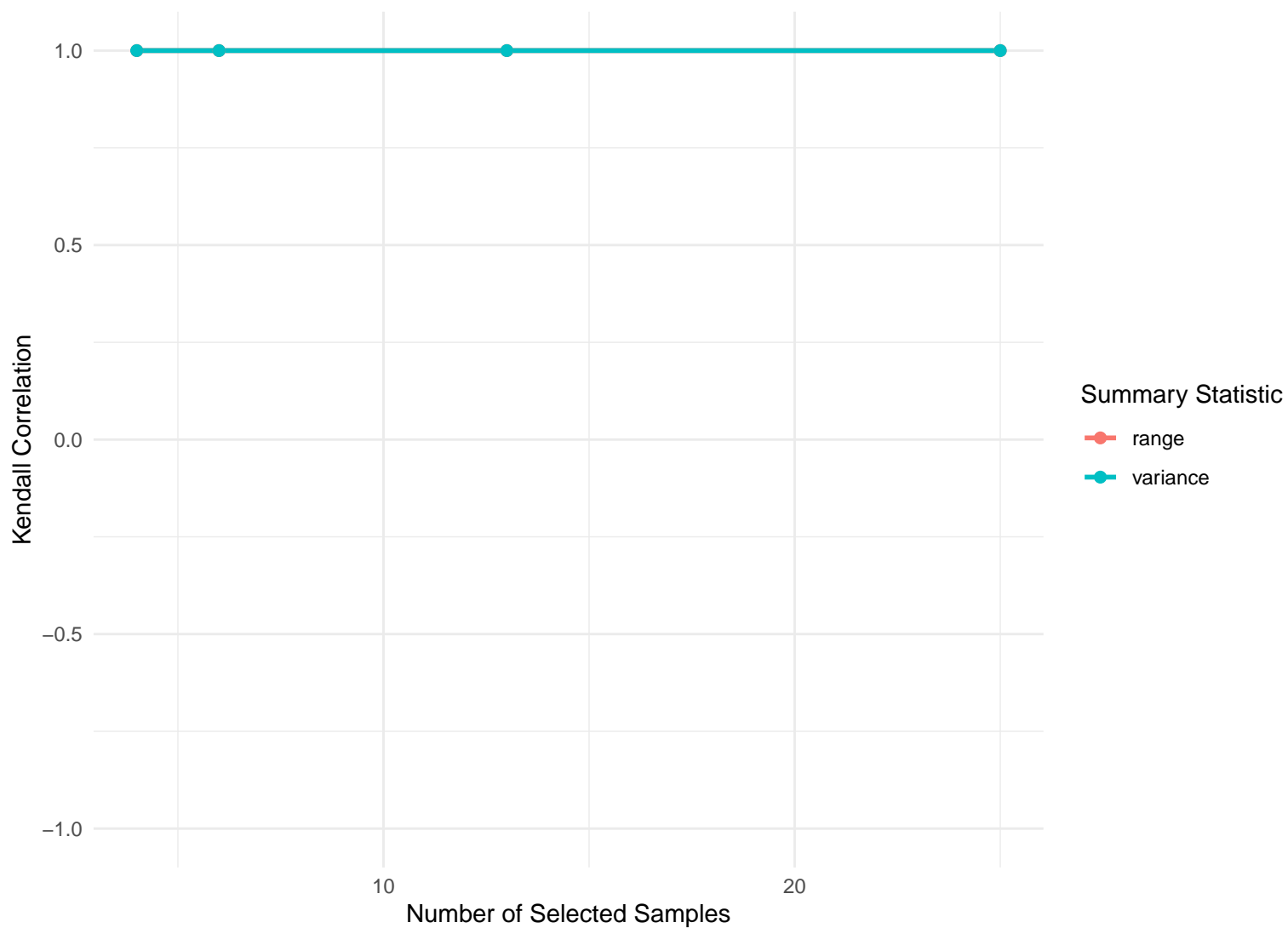
Score: RN (Local, partial1 , Genotype: sinusoidal)



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

20

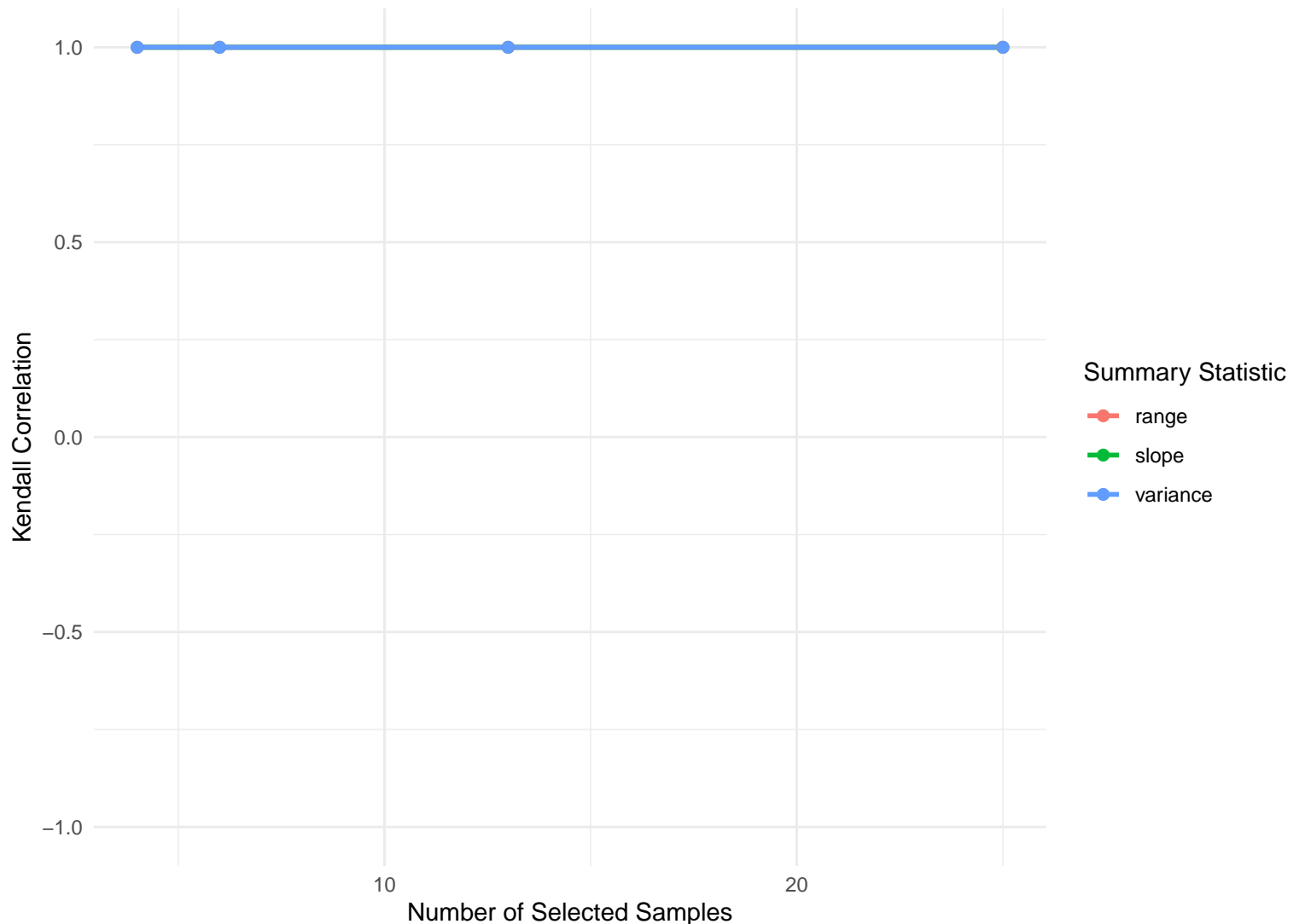
Number of Selected Samples

Summary Statistic

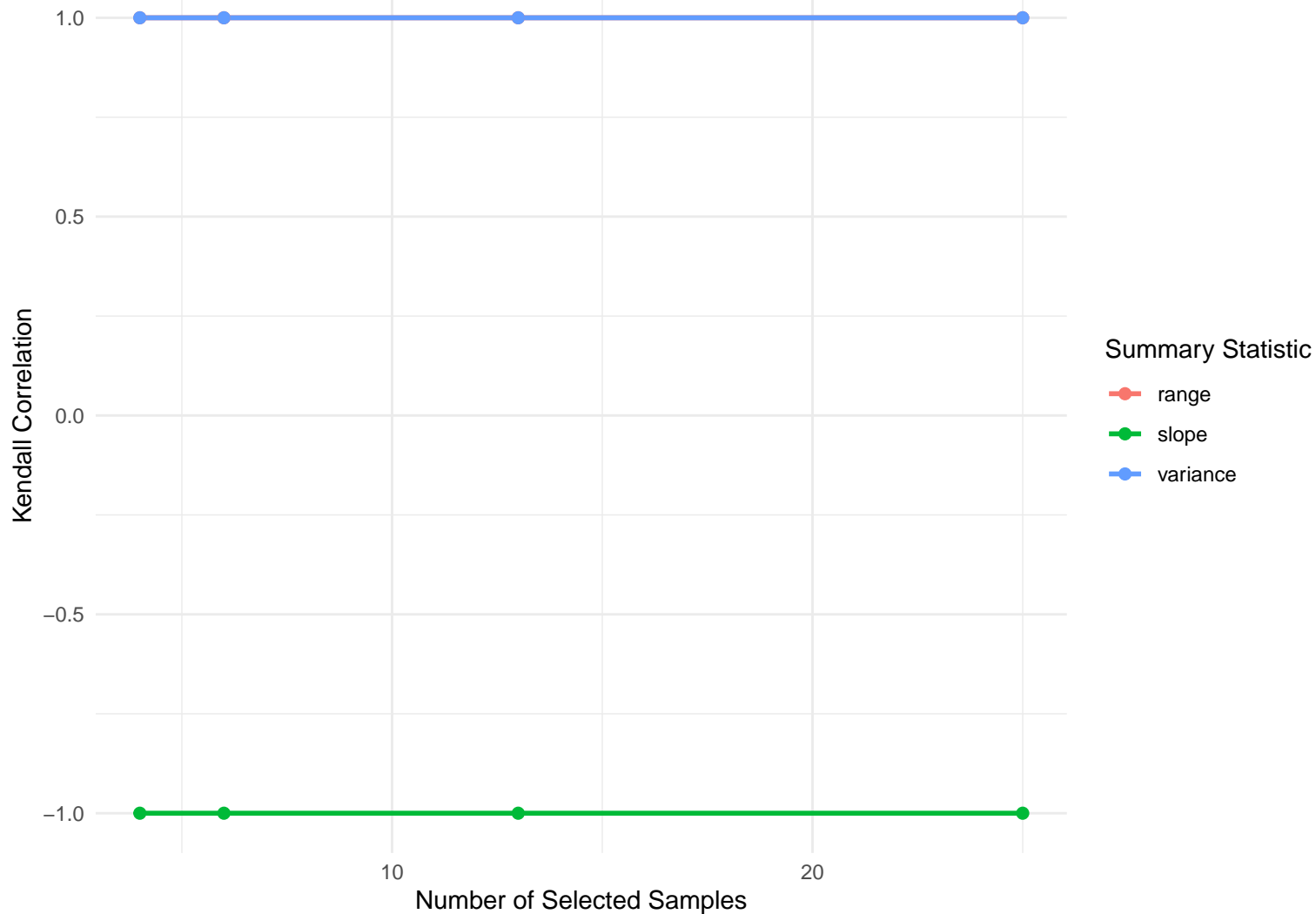
range

slope

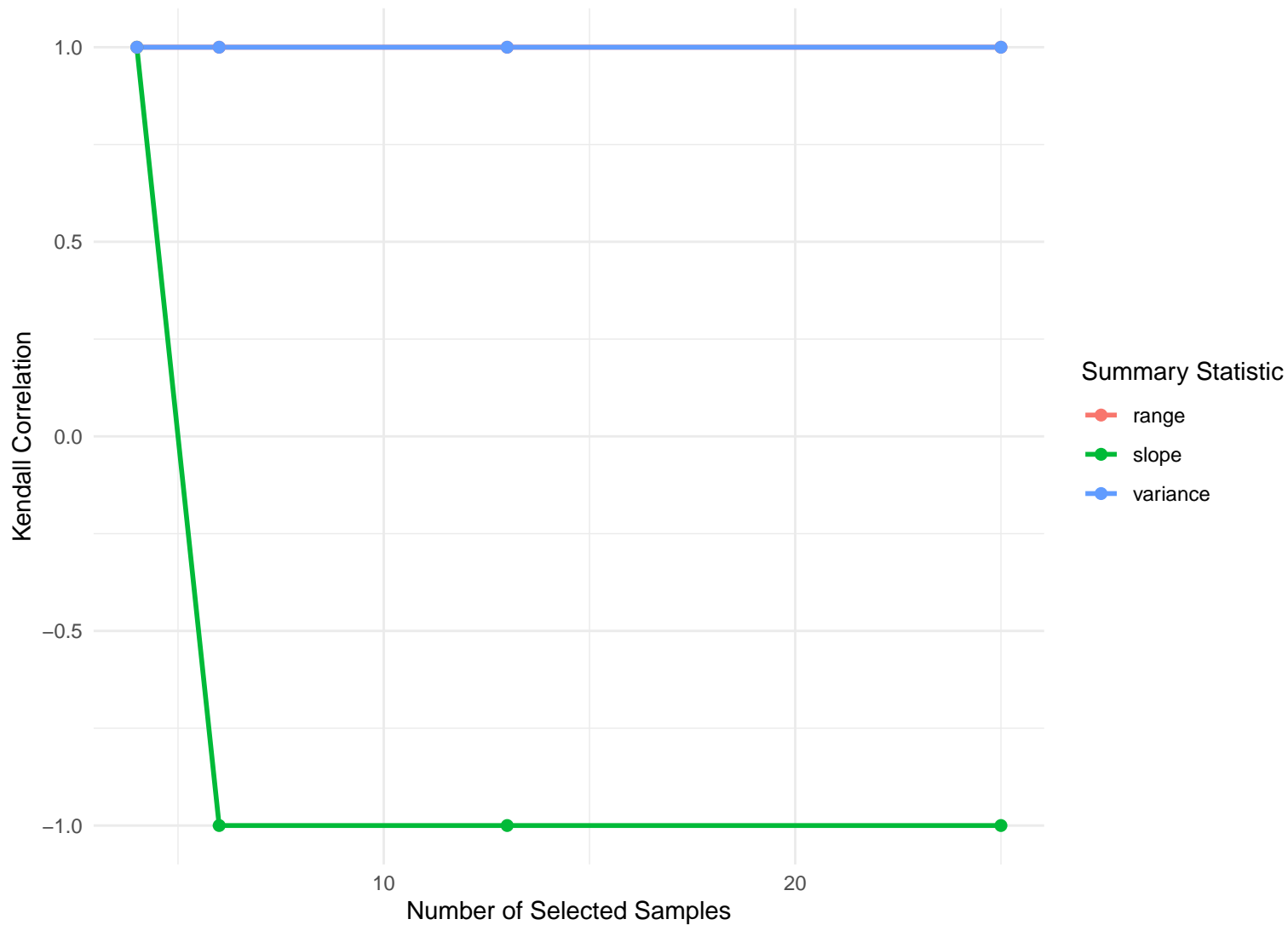
variance



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



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



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

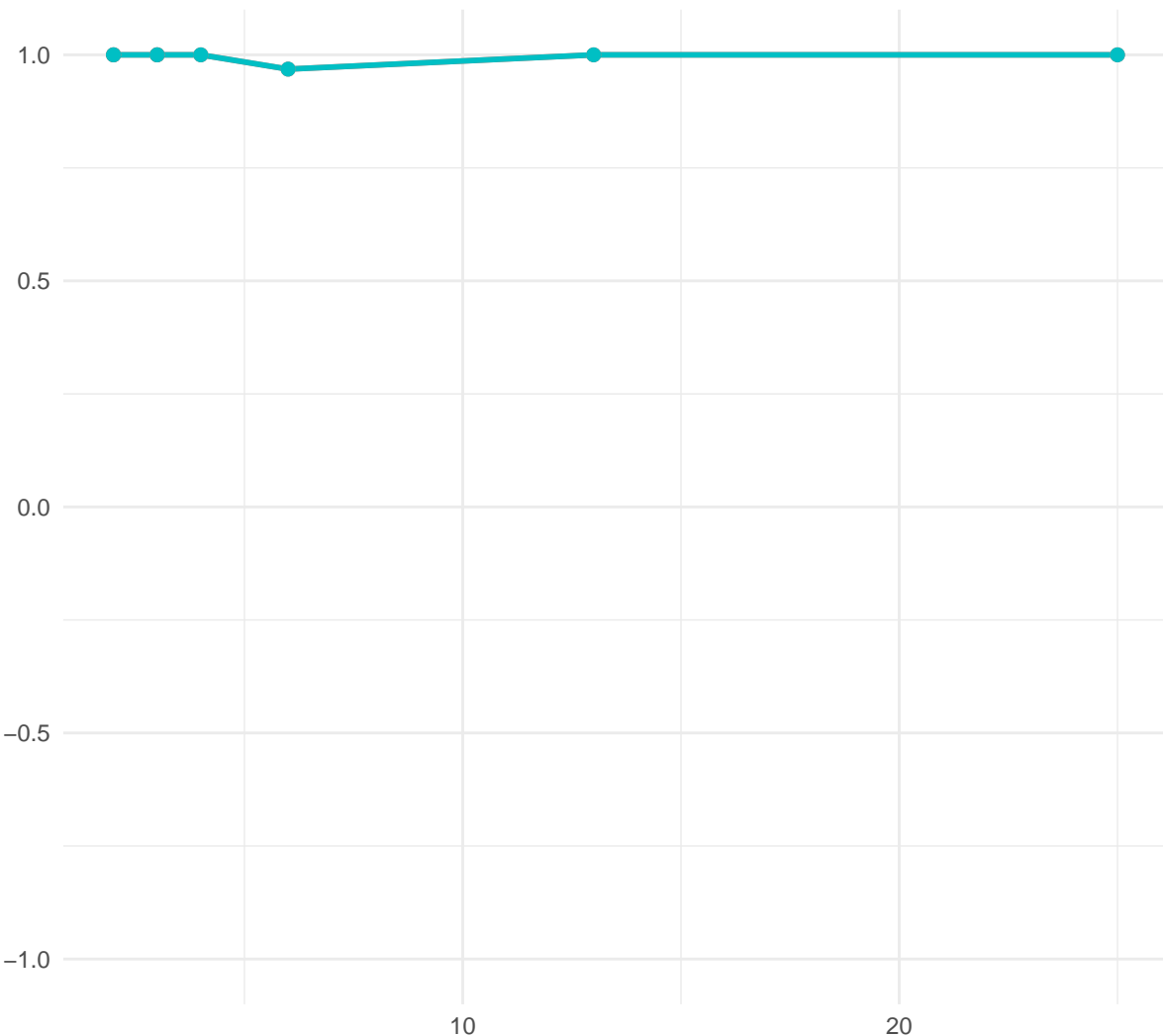
Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

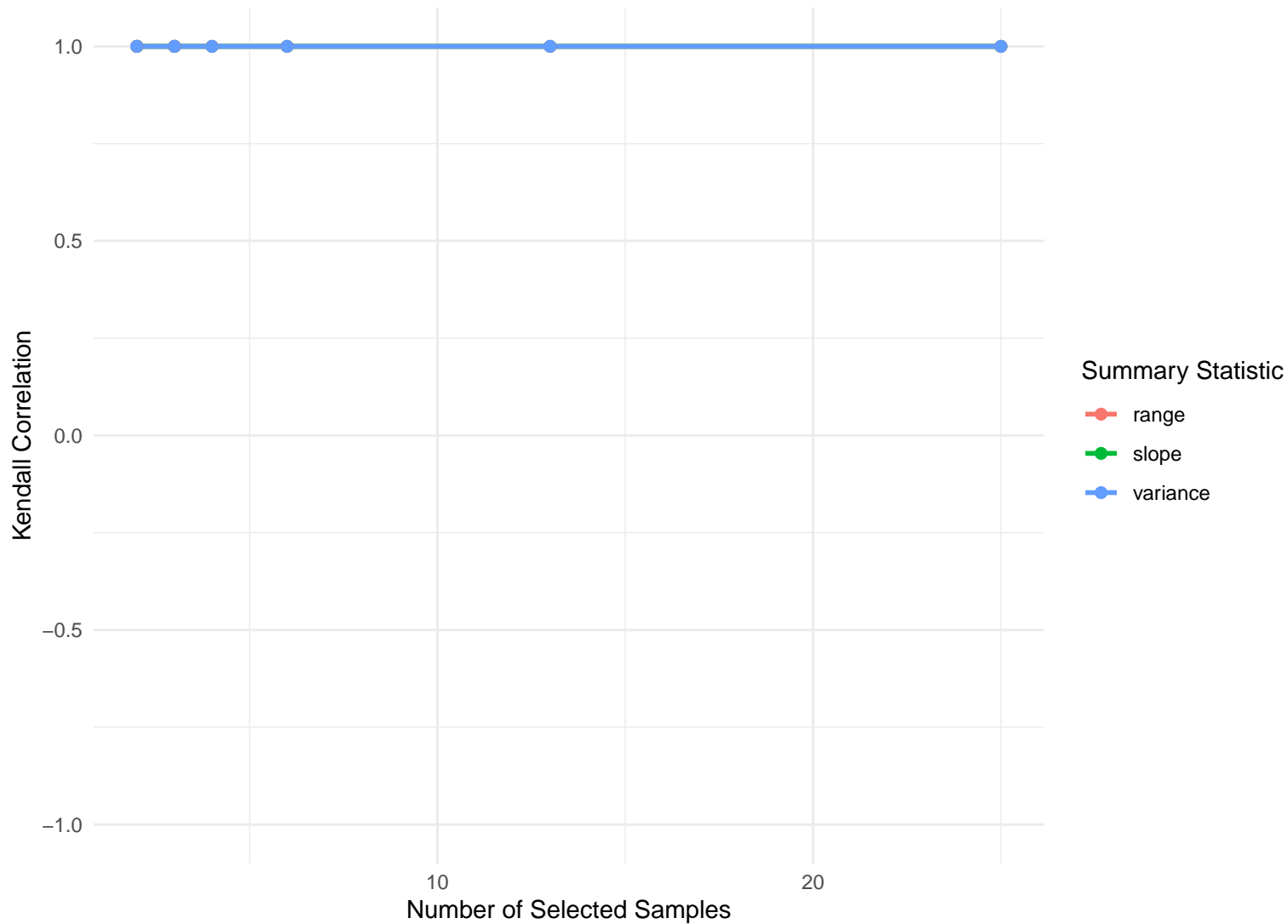
Number of Selected Samples

Summary Statistic

range
variance



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



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

Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

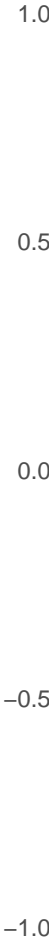
Number of Selected Samples

Summary Statistic

- range
- slope
- variance

10

20

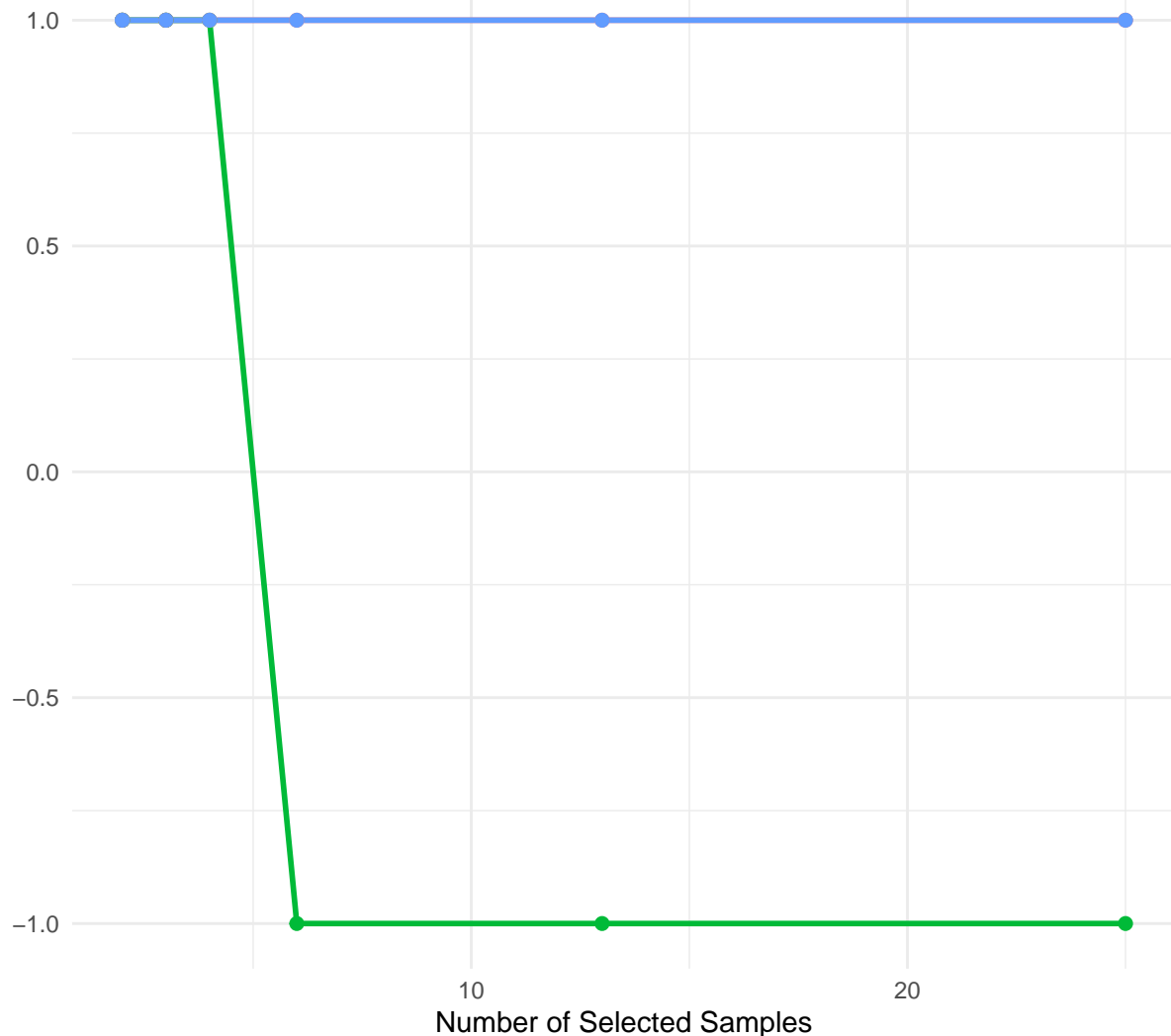


Score: D_slope (Local, partial1 , Genotype: wave)

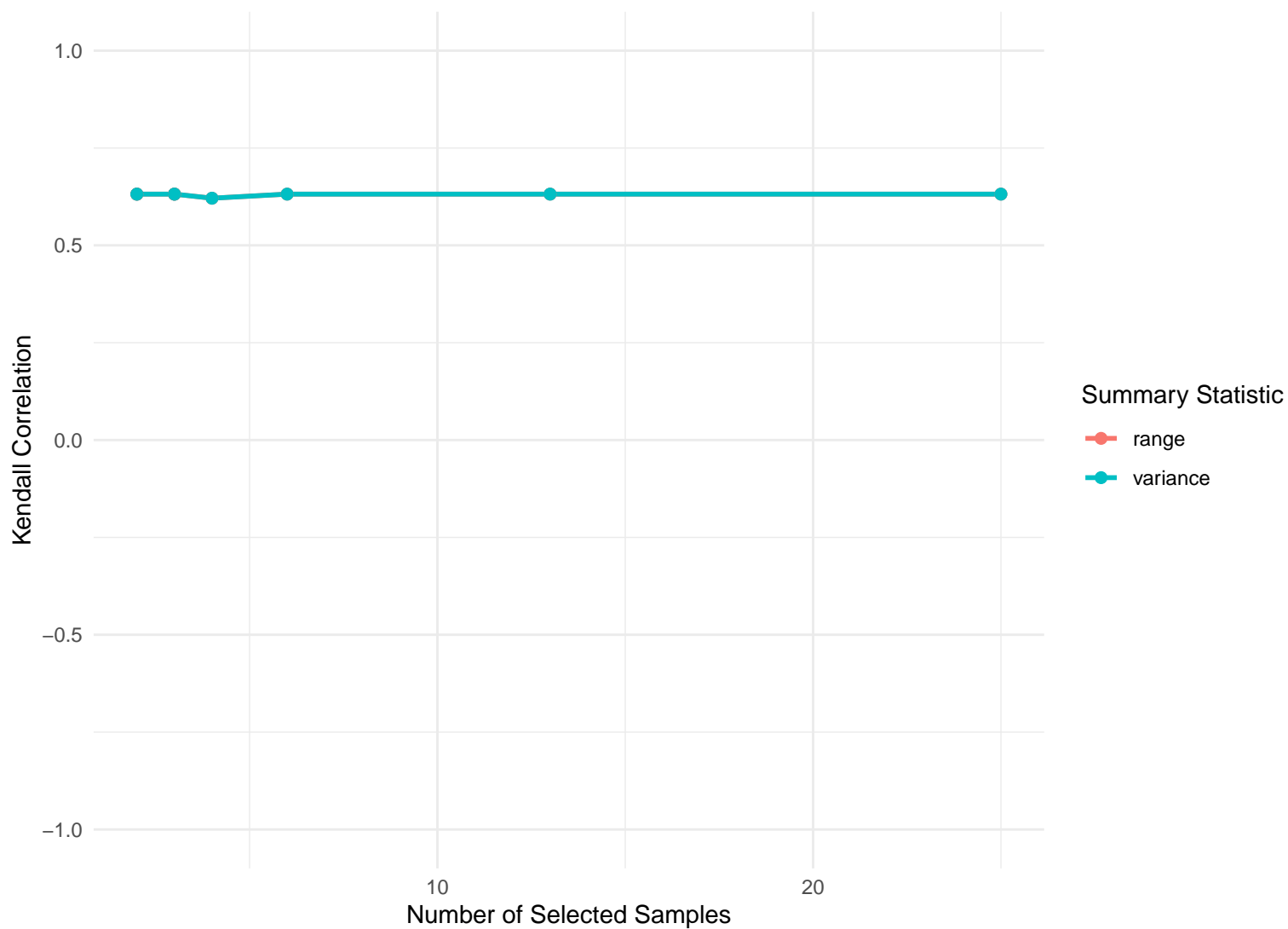
Kendall Correlation

Summary Statistic

- range
- slope
- variance



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

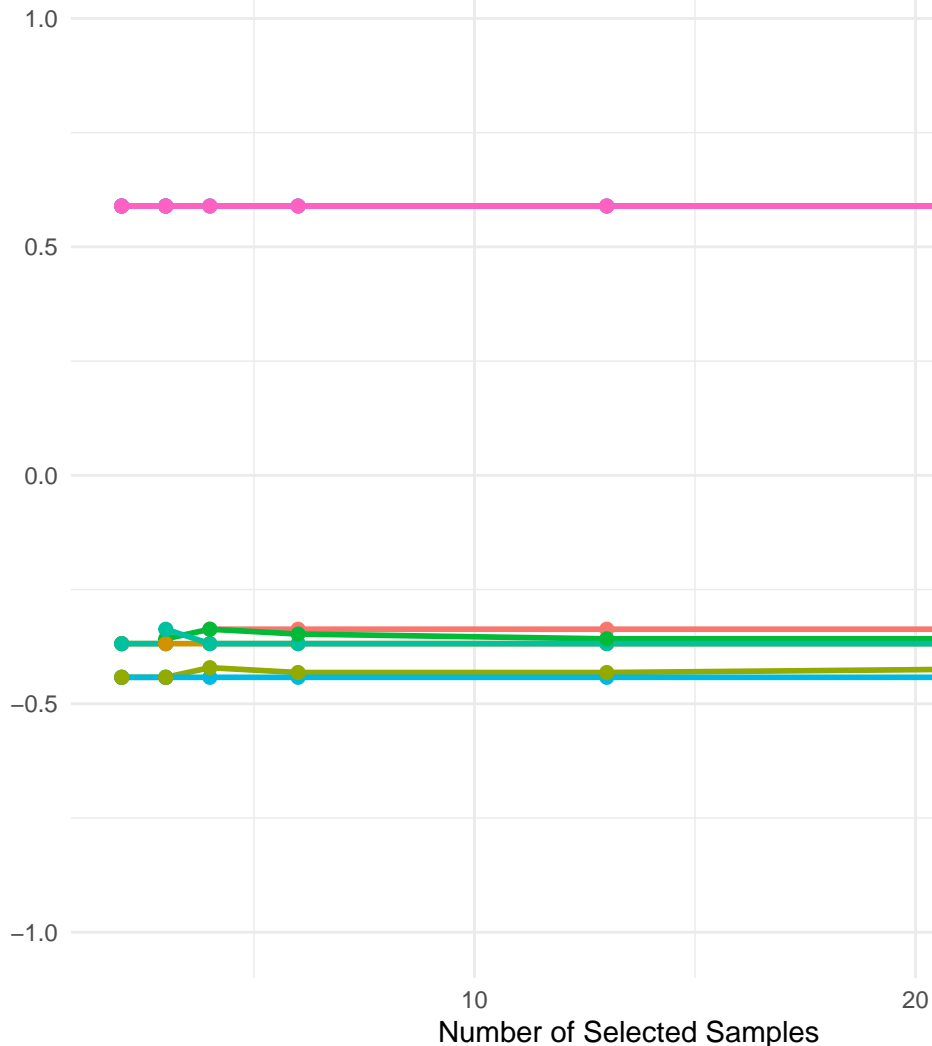


Score: RC (Local, partial1 , Genotype: gaussian)

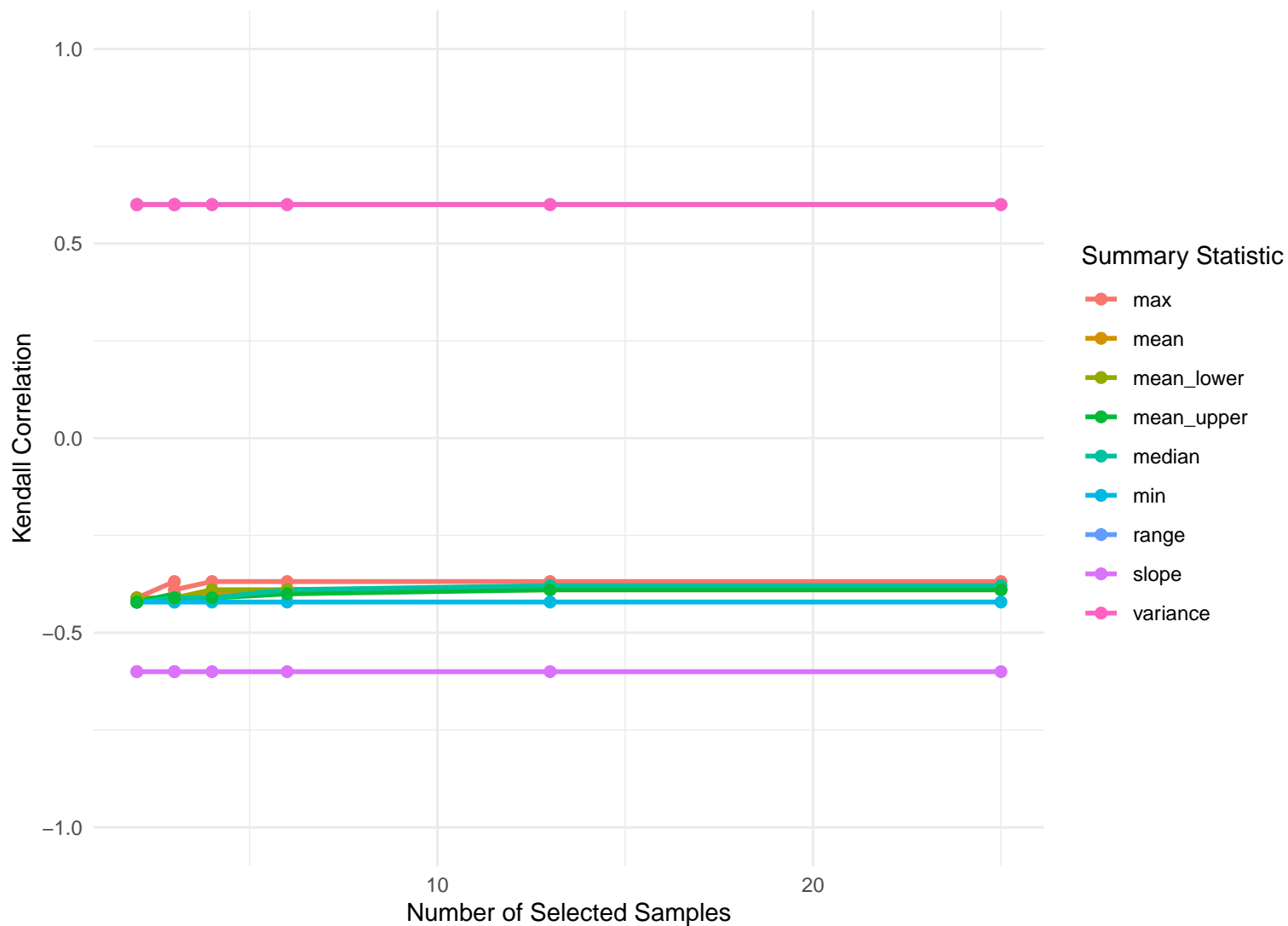
Kendall Correlation

Summary Statistic

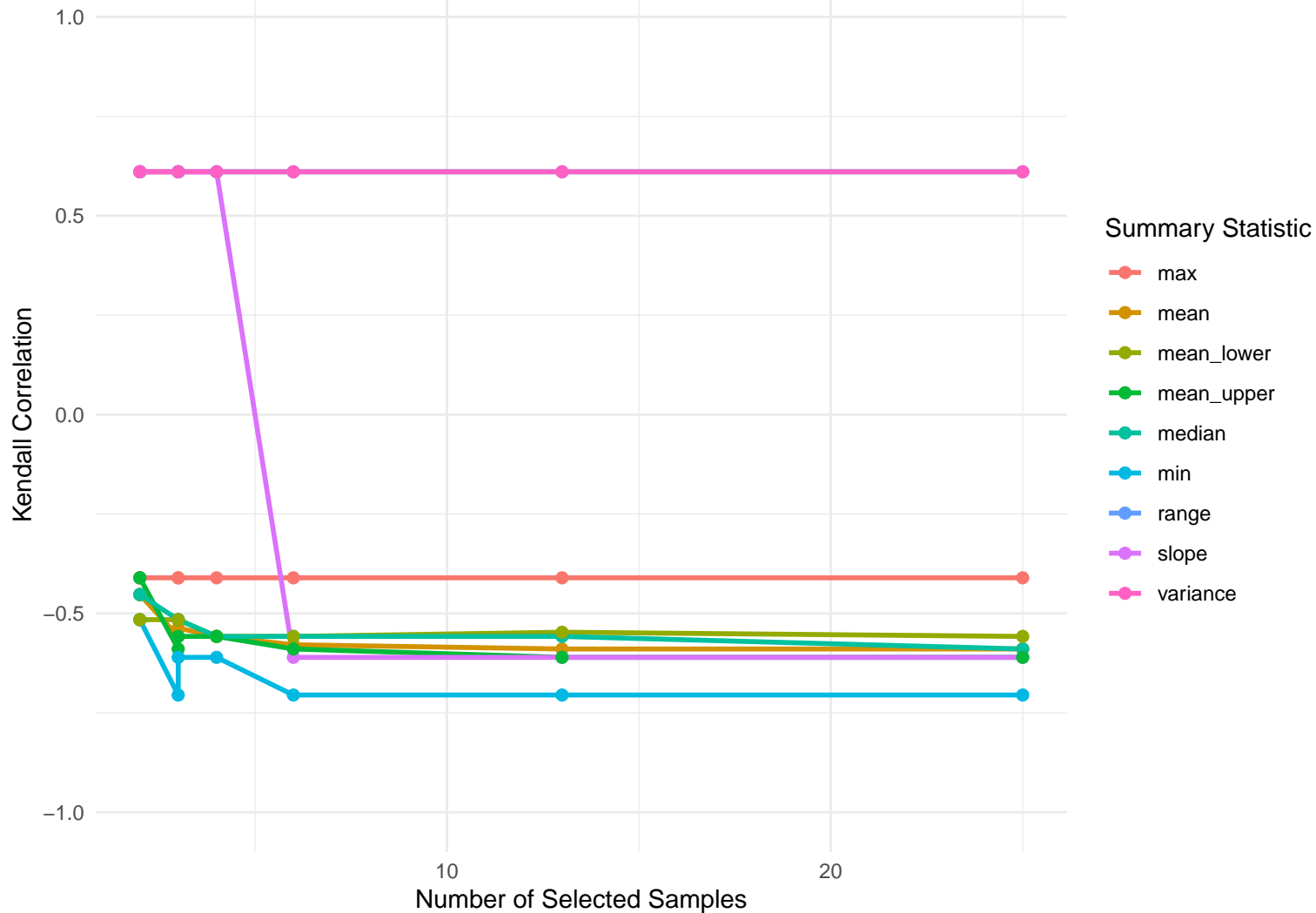
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



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

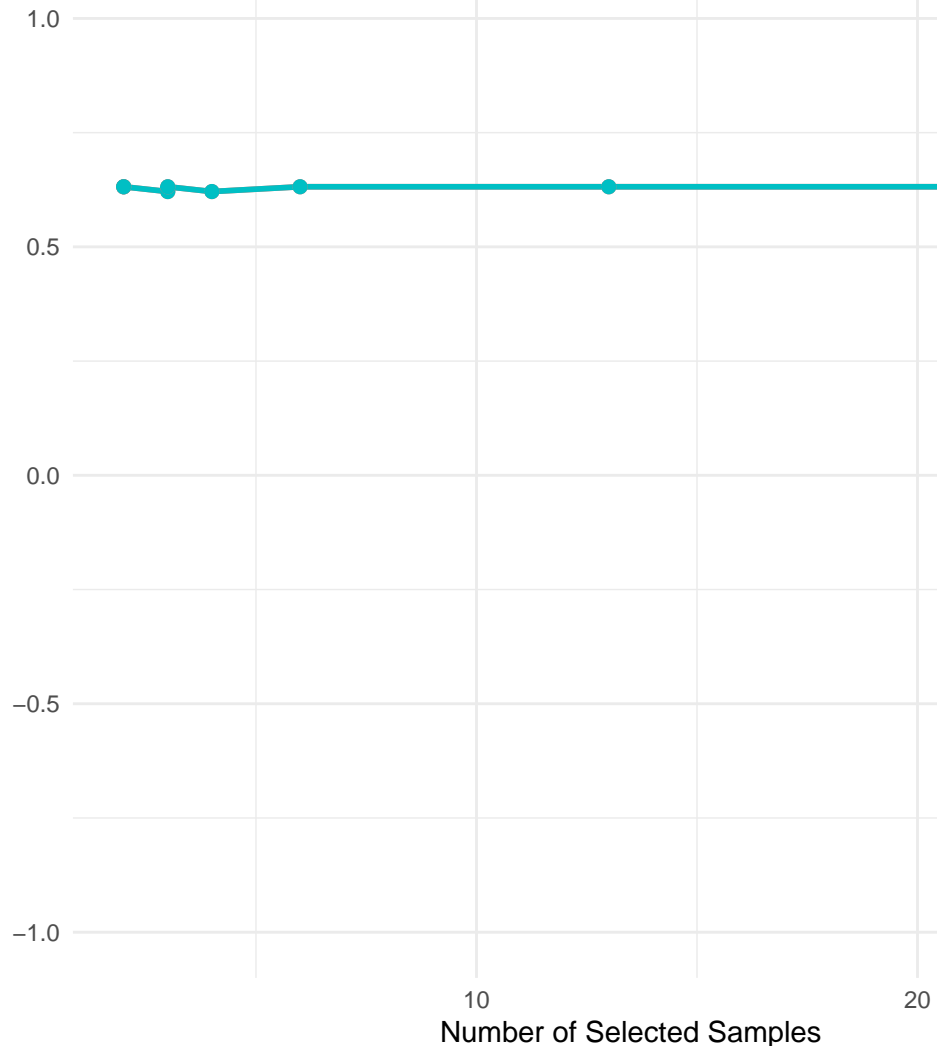


Score: RC (Local, partial1 , Genotype: wave)



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

Kendall Correlation



Summary Statistic

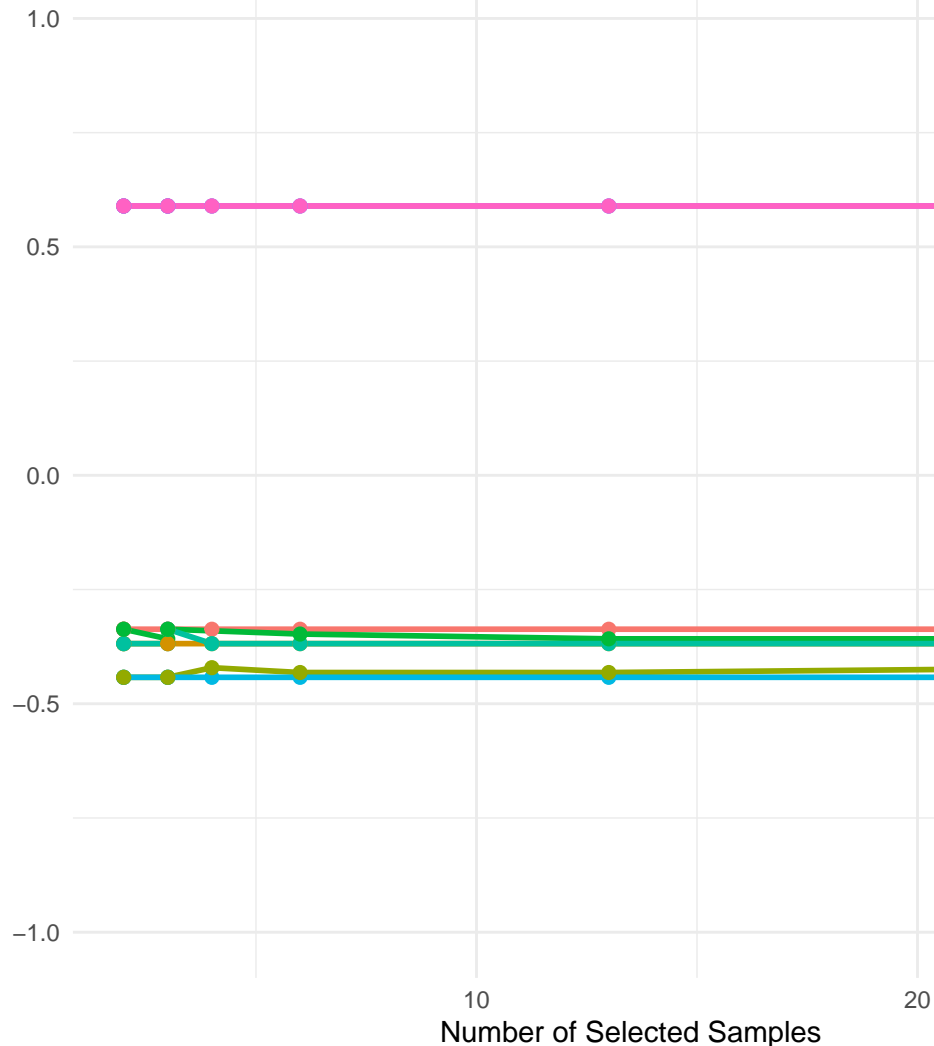
- range
- variance

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

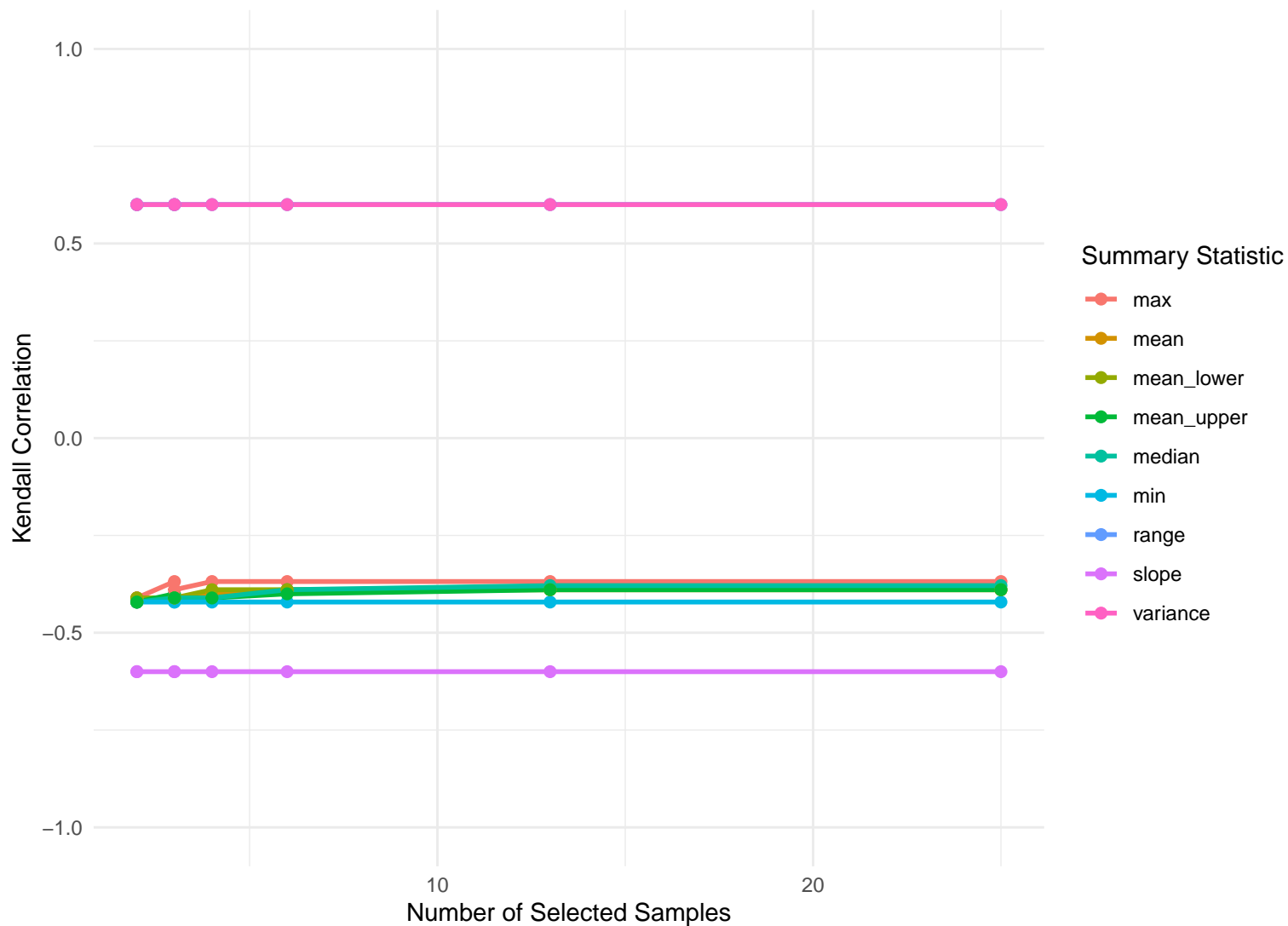
Kendall Correlation

Summary Statistic

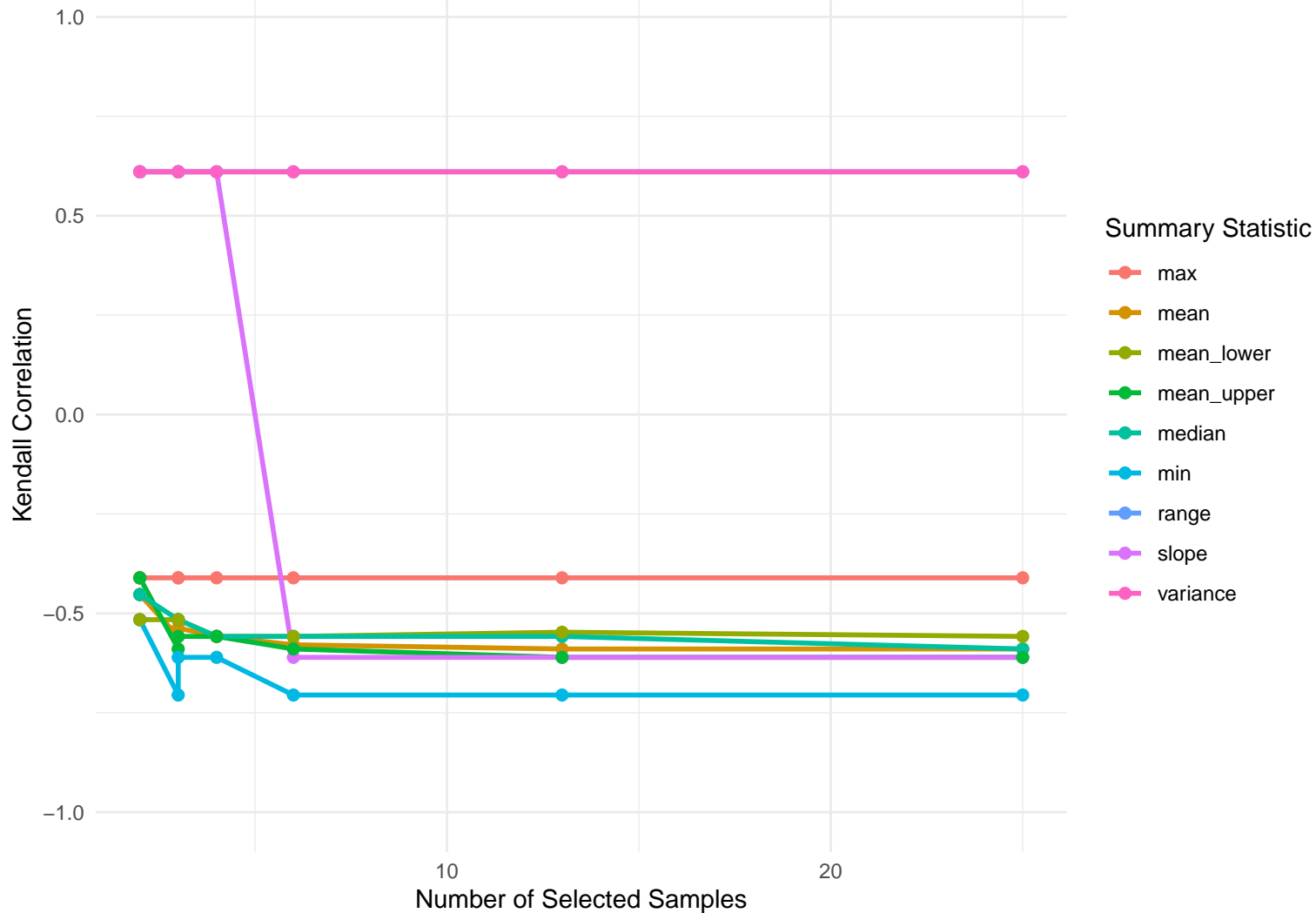
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



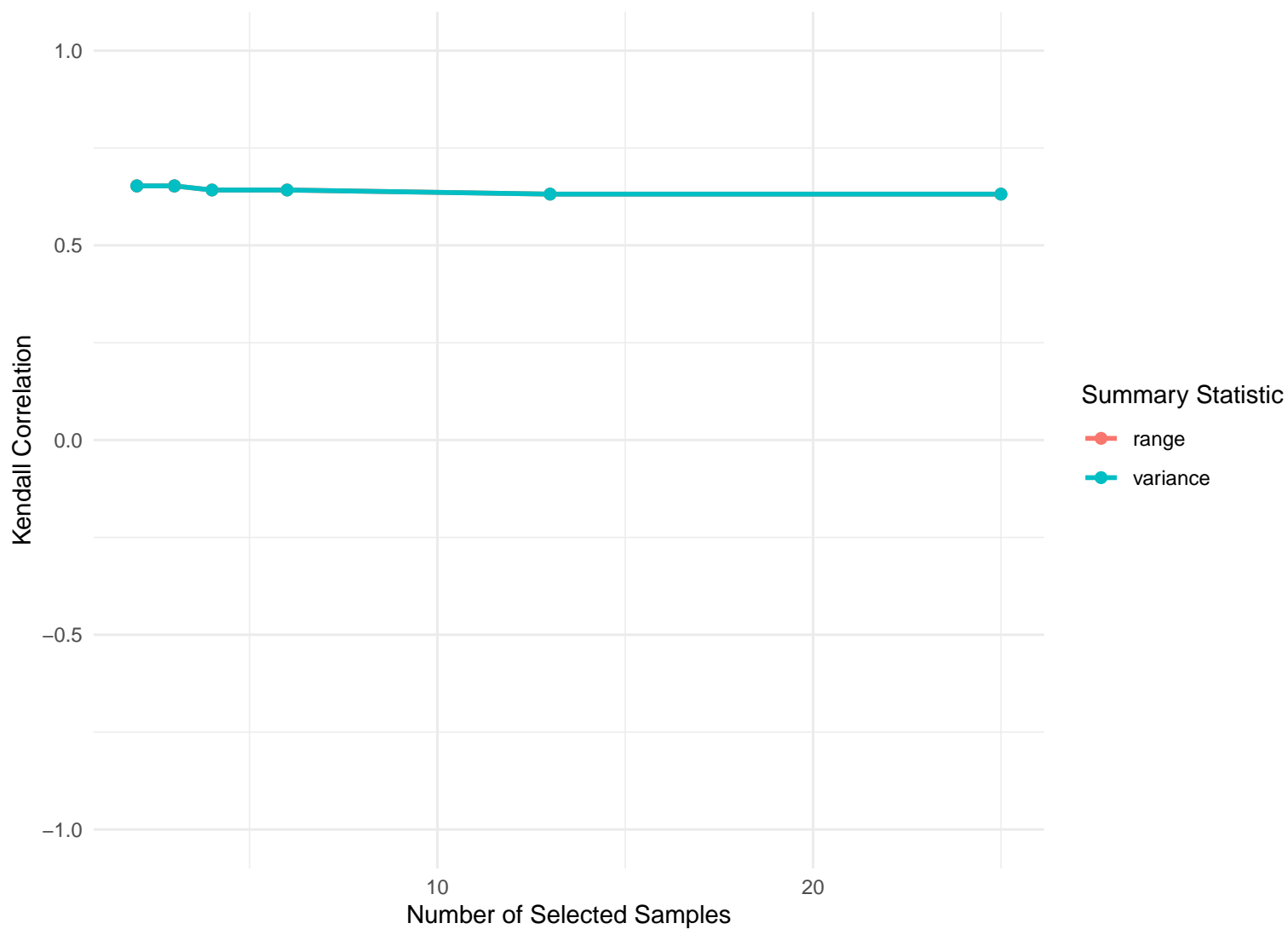
Score: gPi (Local, partial1 , Genotype: sinusoidal)



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



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



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

Kendall Correlation

Summary Statistic

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

1.0

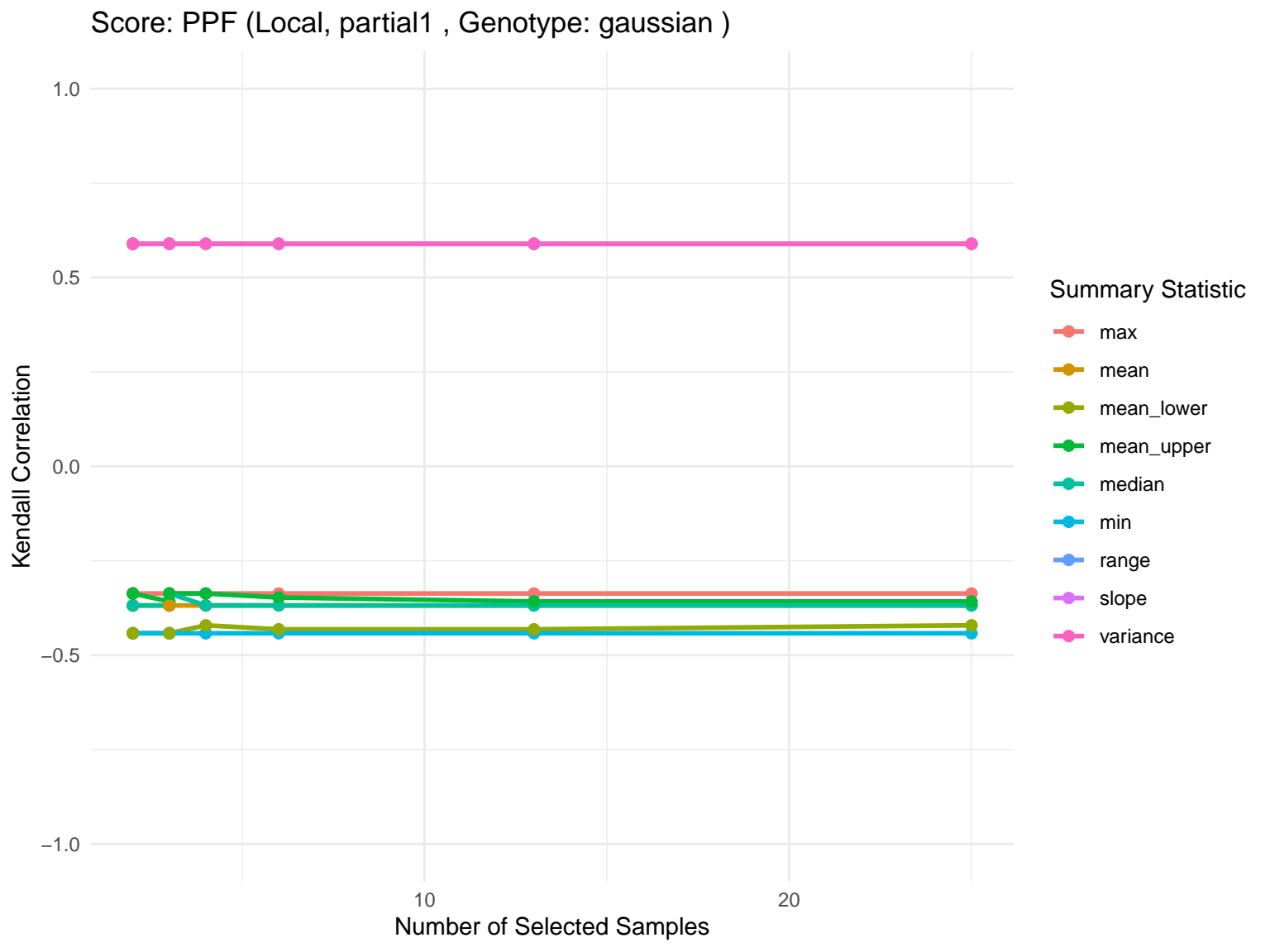
0.5

0.0

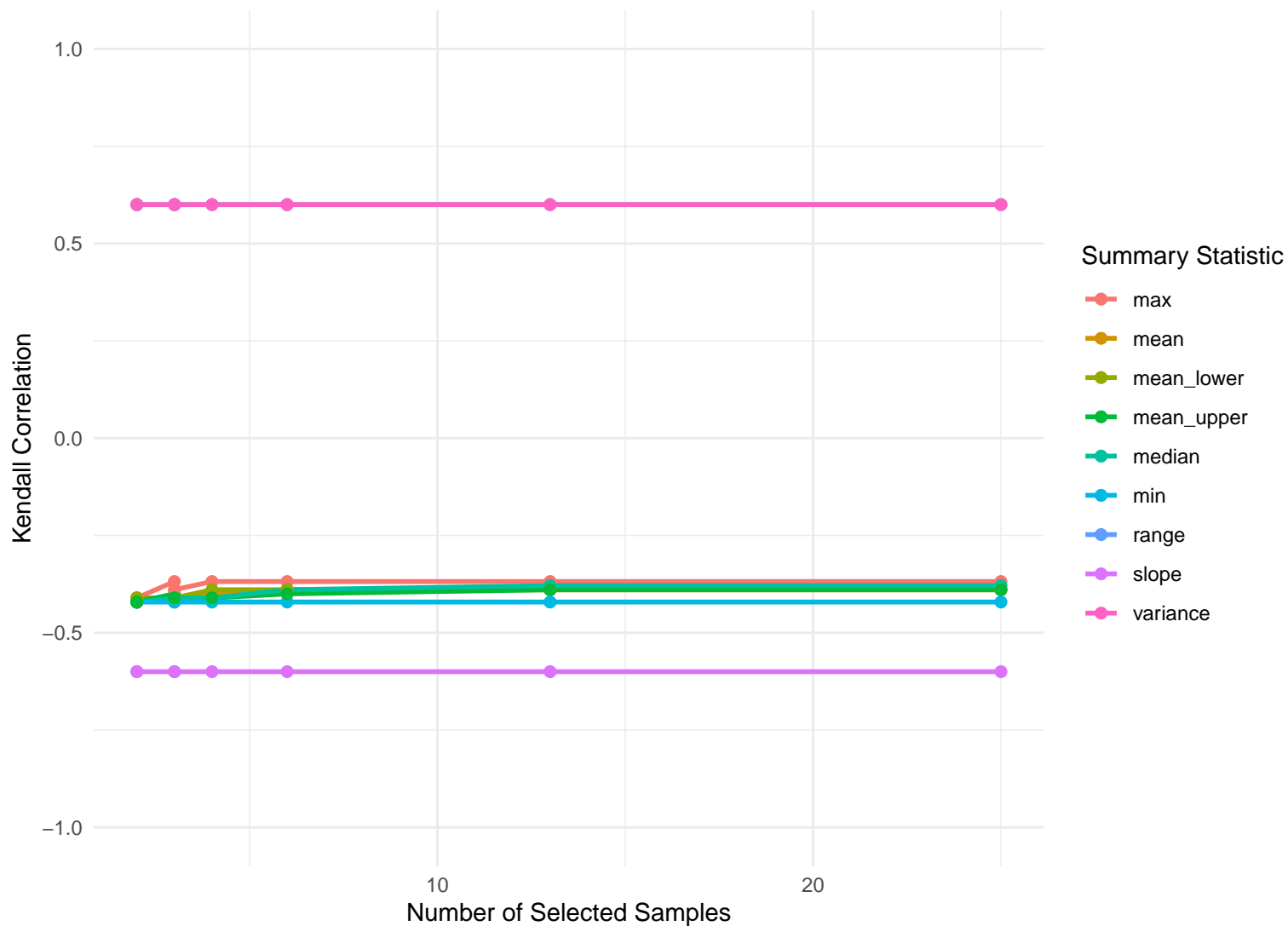
-0.5

-1.0

10
20
Number of Selected Samples



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



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

Kendall Correlation

Summary Statistic

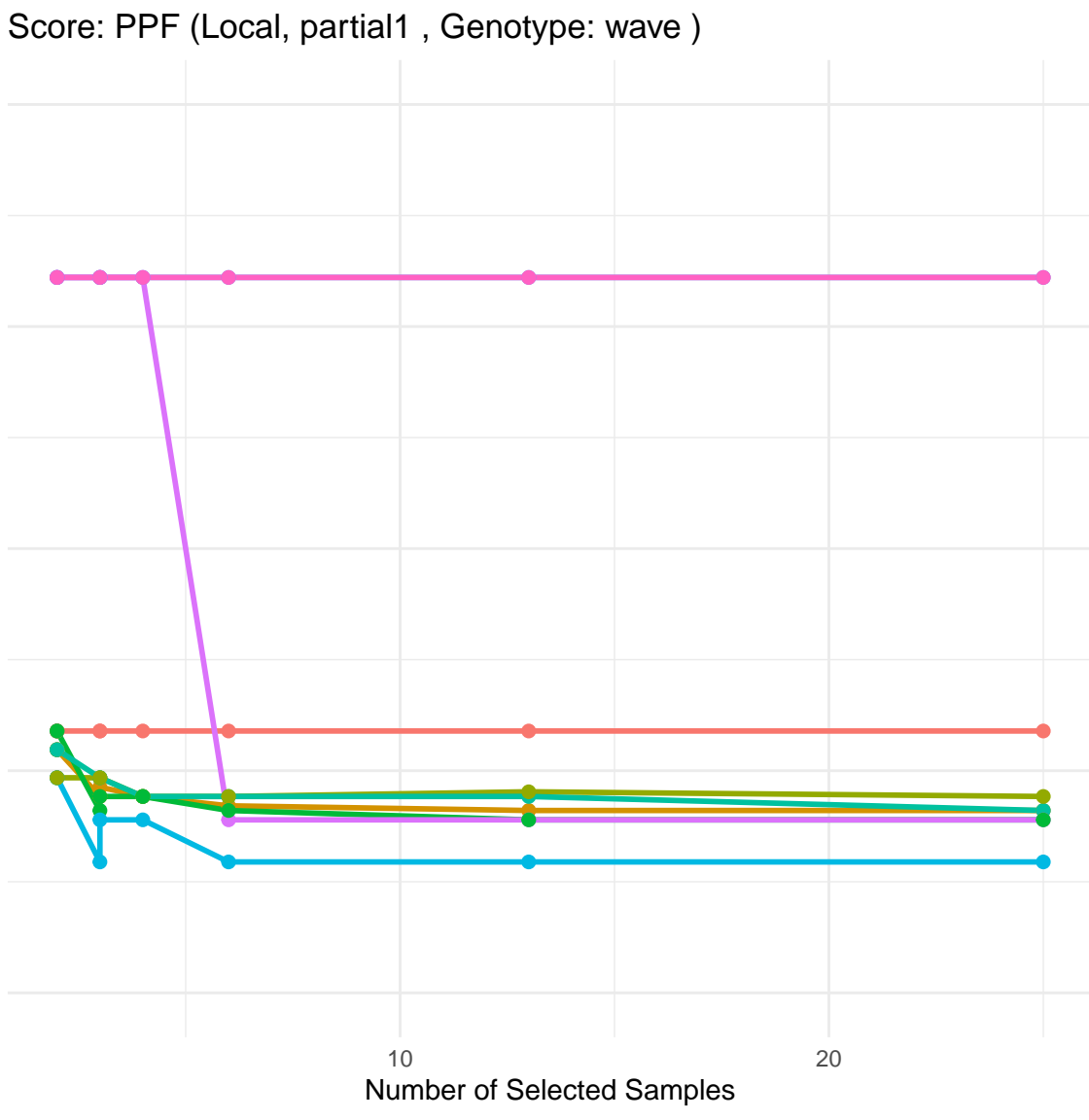
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance

1.0
0.5
0.0
-0.5
-1.0

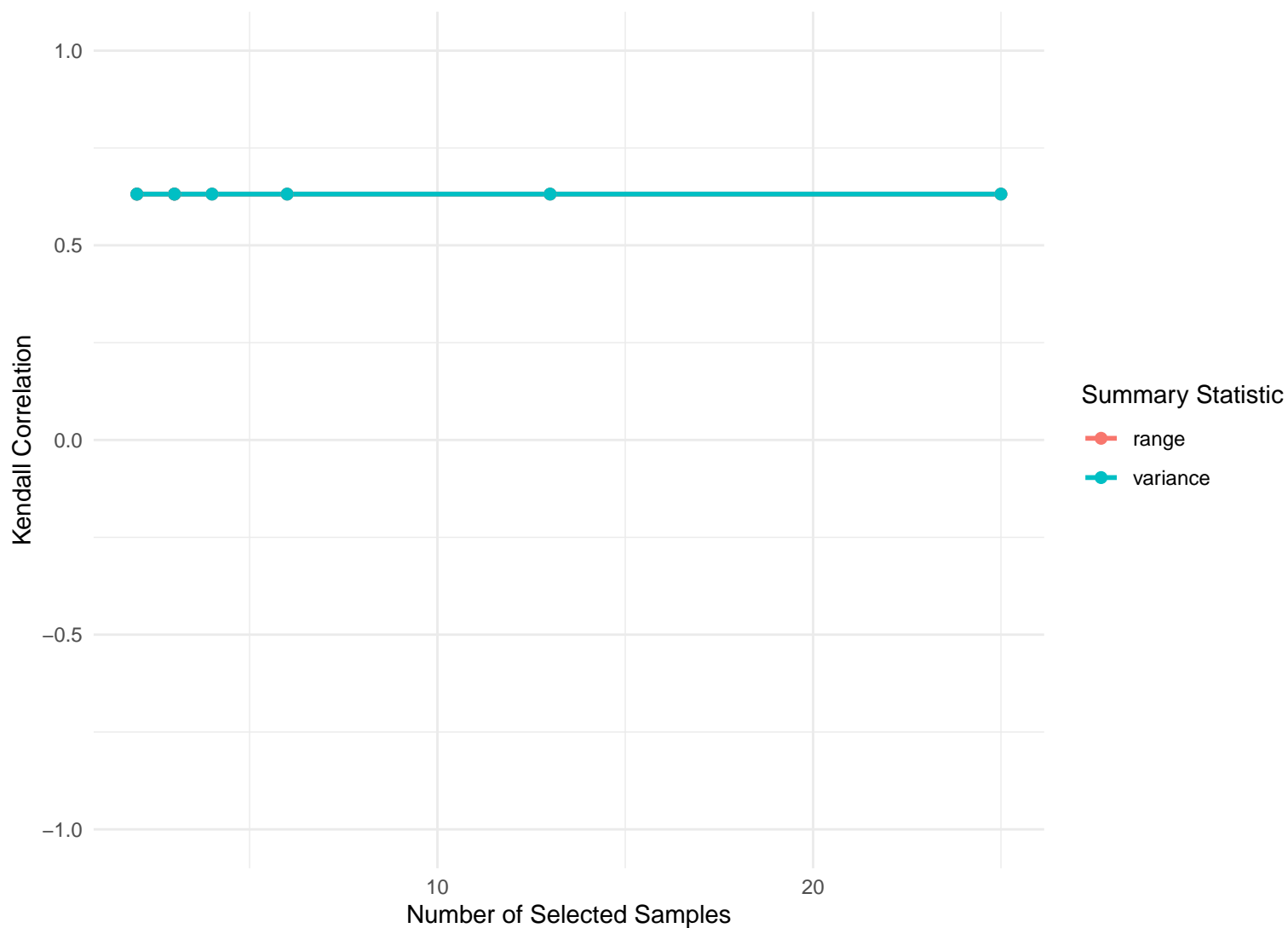
Number of Selected Samples

10

20



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

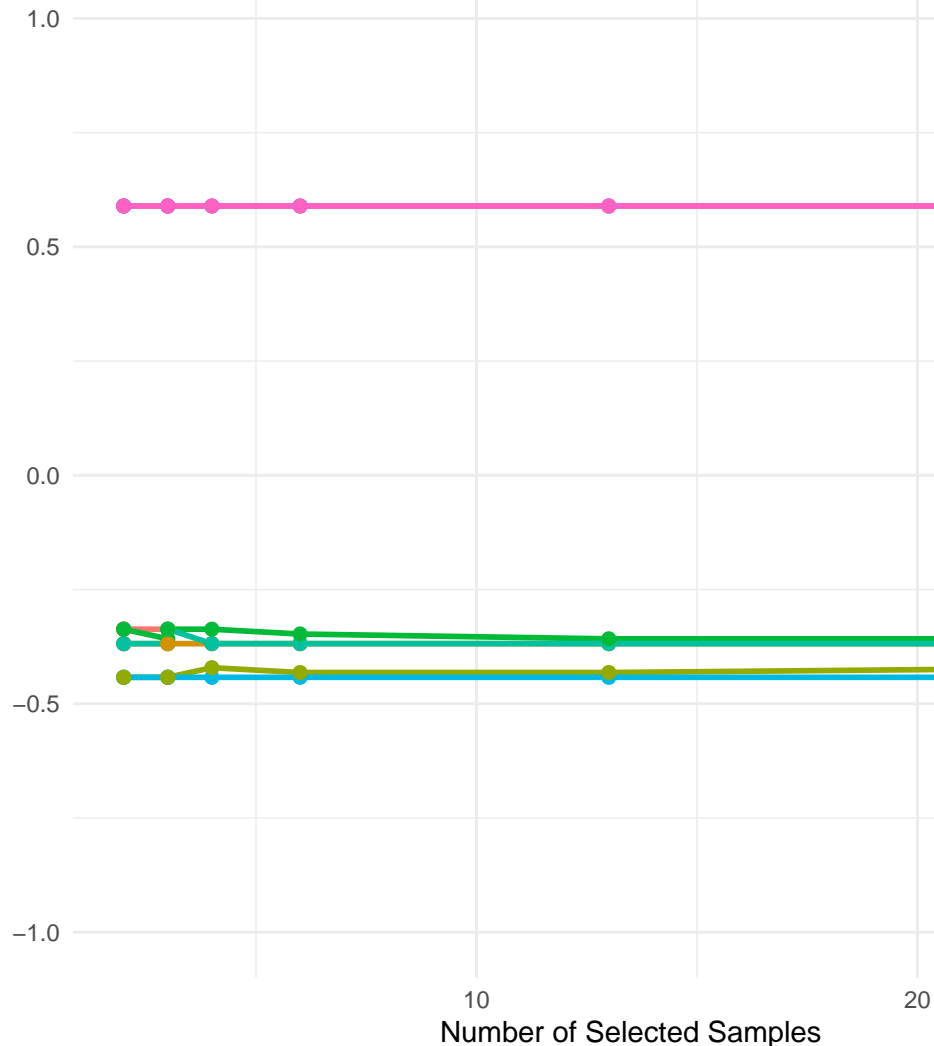


Score: PPI (Local, partial1 , Genotype: gaussian)

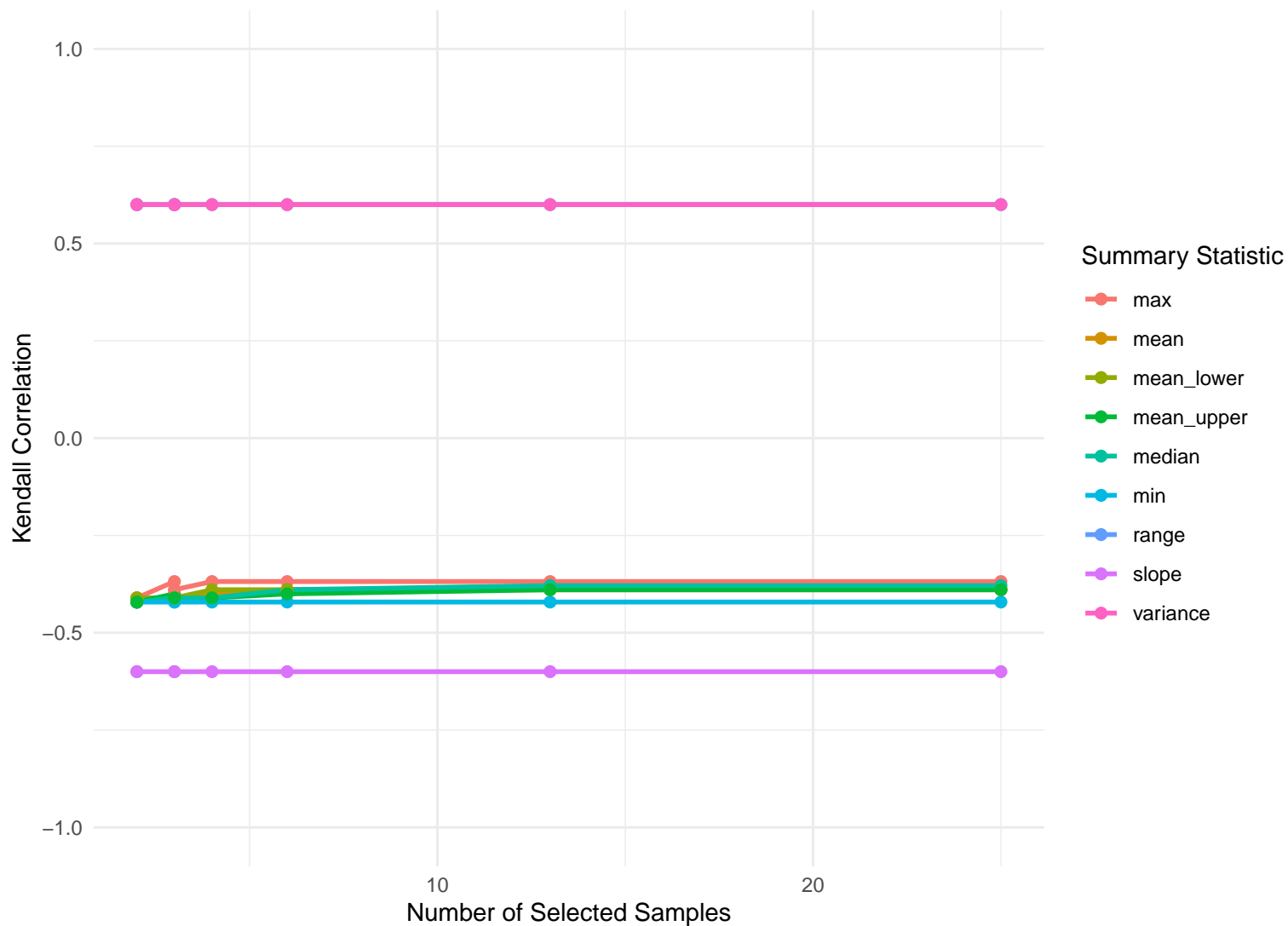
Kendall Correlation

Summary Statistic

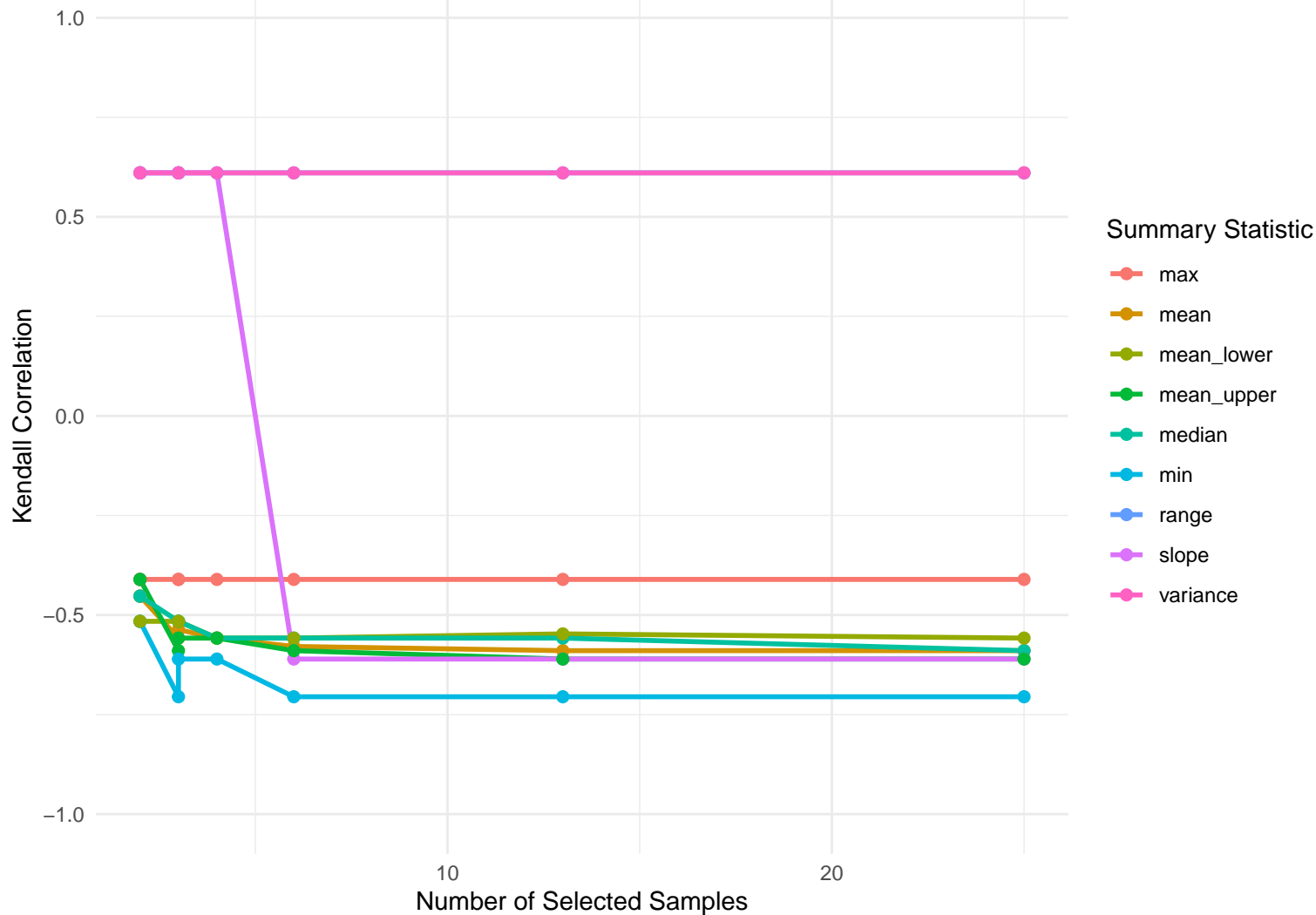
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



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

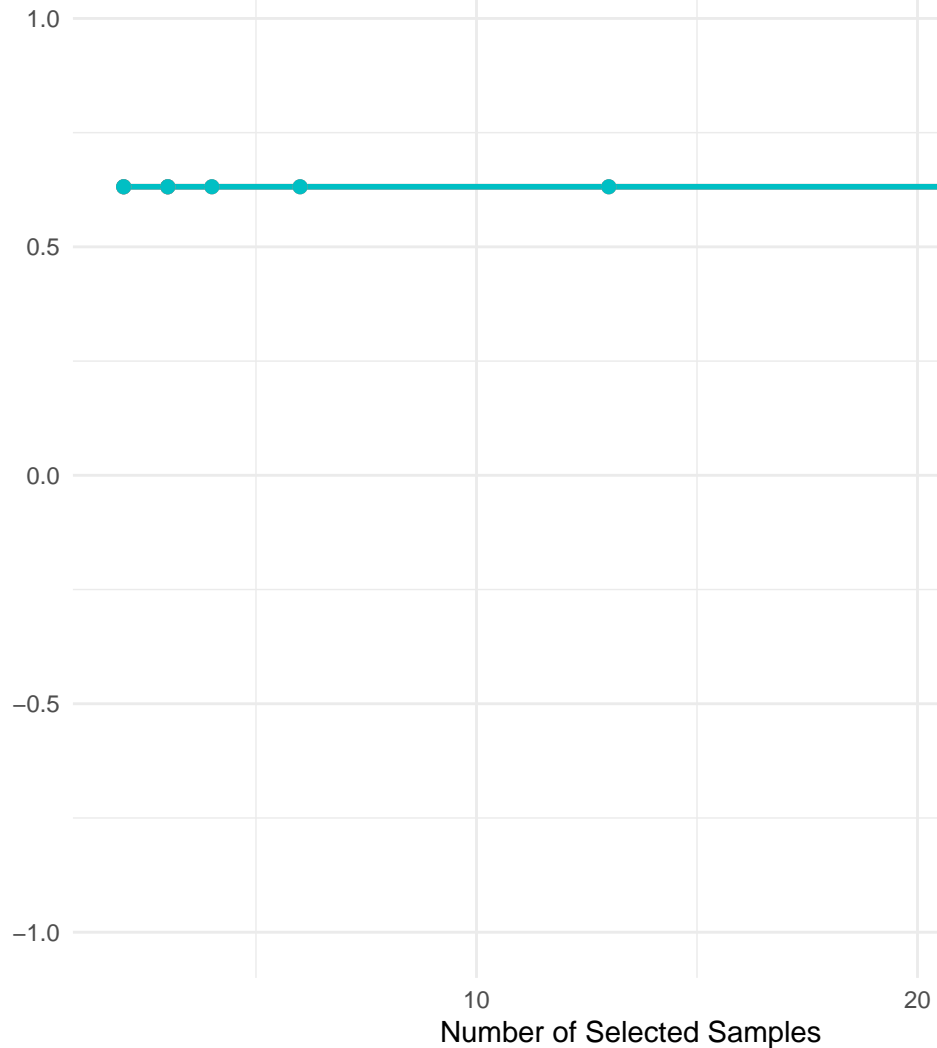


Score: PPI (Local, partial1 , Genotype: wave)



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

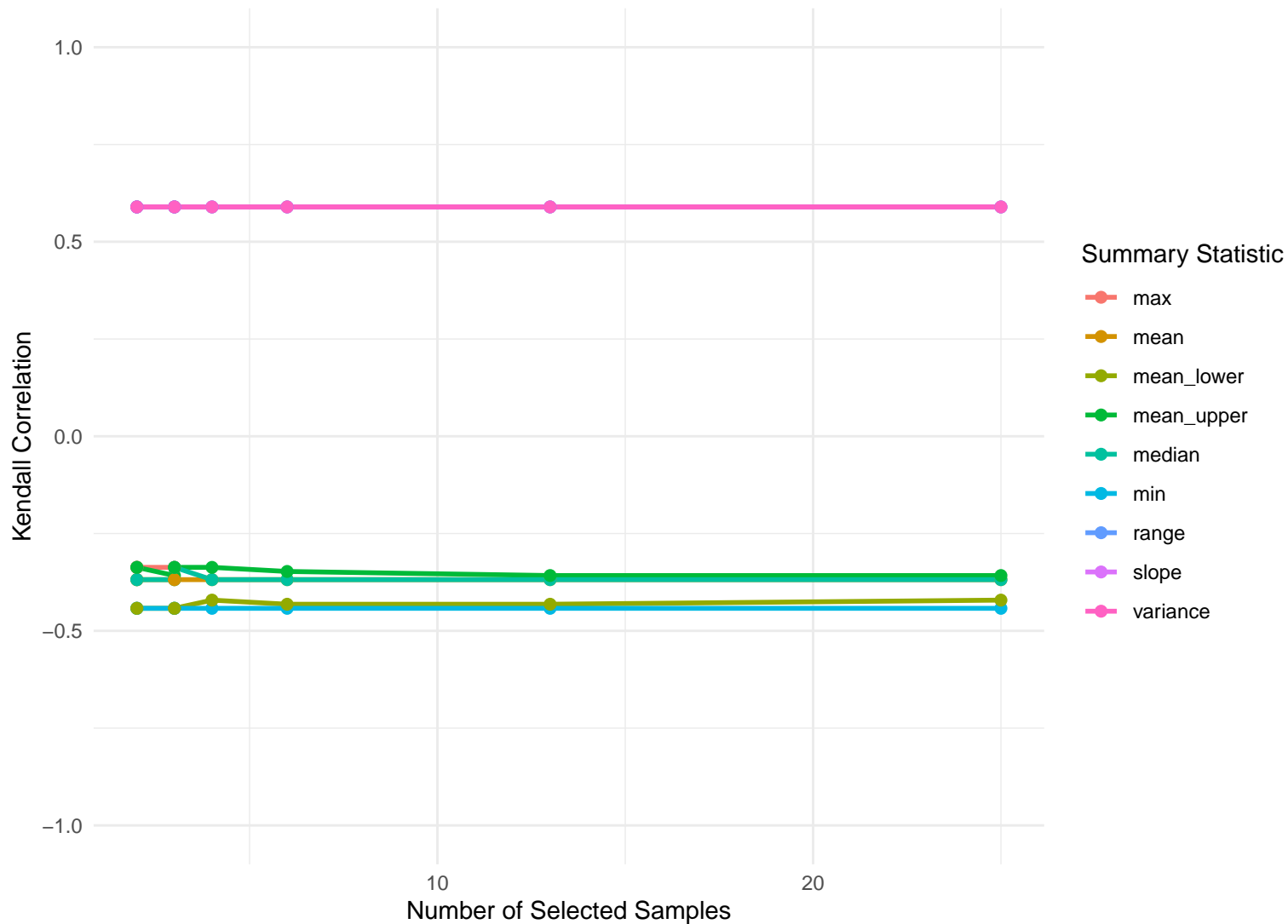
Kendall Correlation



Summary Statistic

- range
- variance

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



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



Summary Statistic

max

mean

mean_lower

mean_upper

median

min

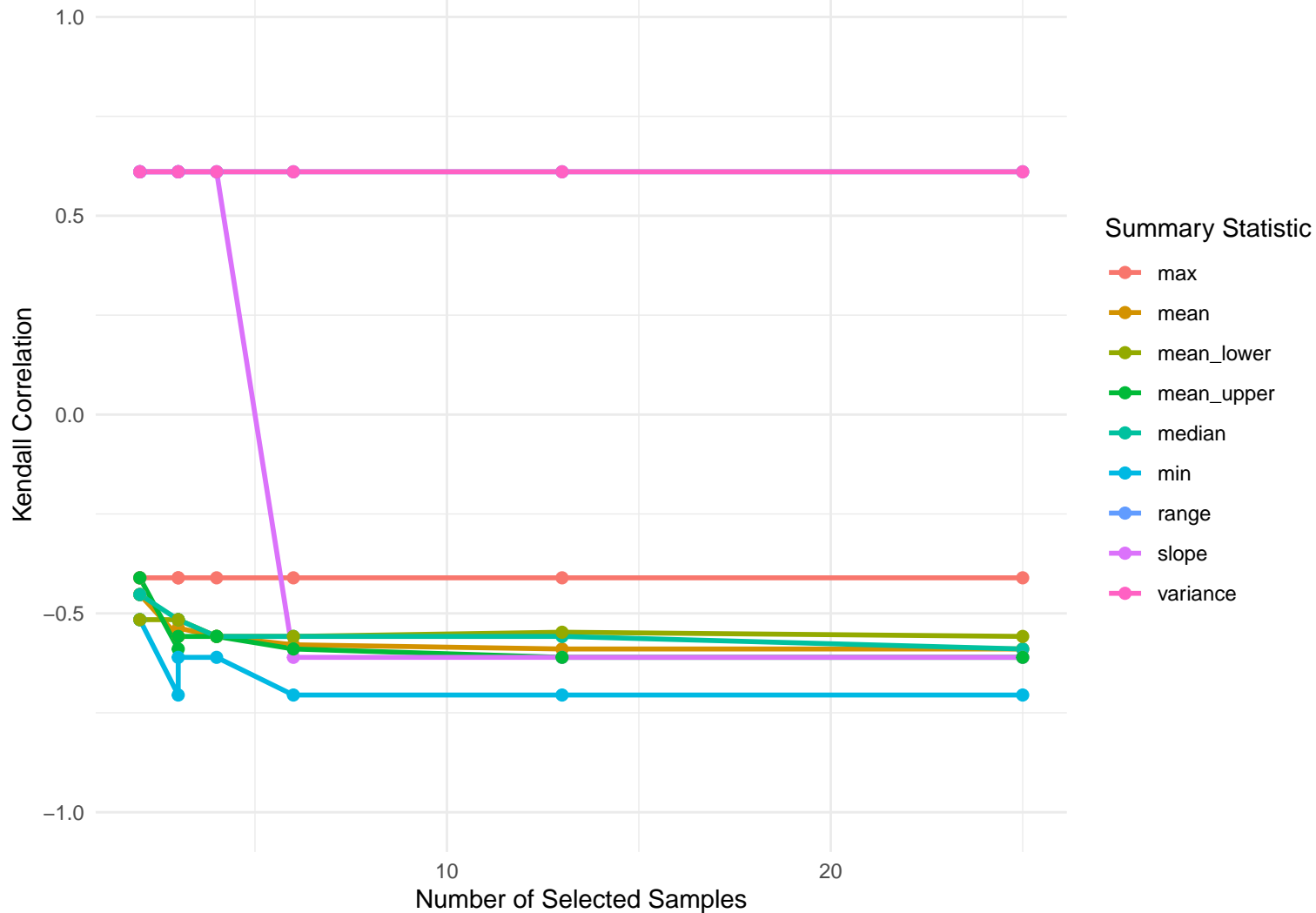
range

slope

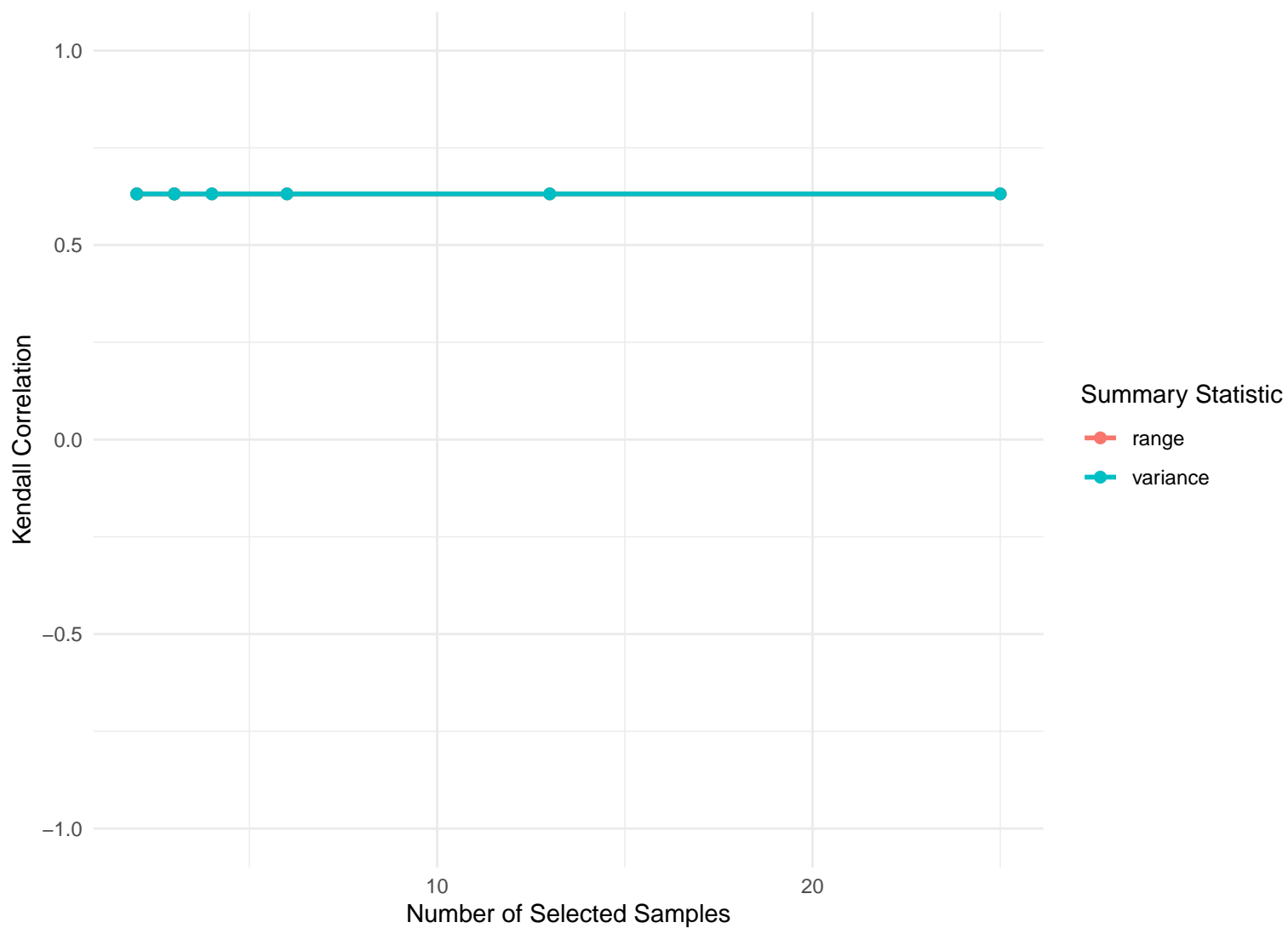
variance

Number of Selected Samples

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



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



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

Kendall Correlation

Summary Statistic

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

1.0

0.5

0.0

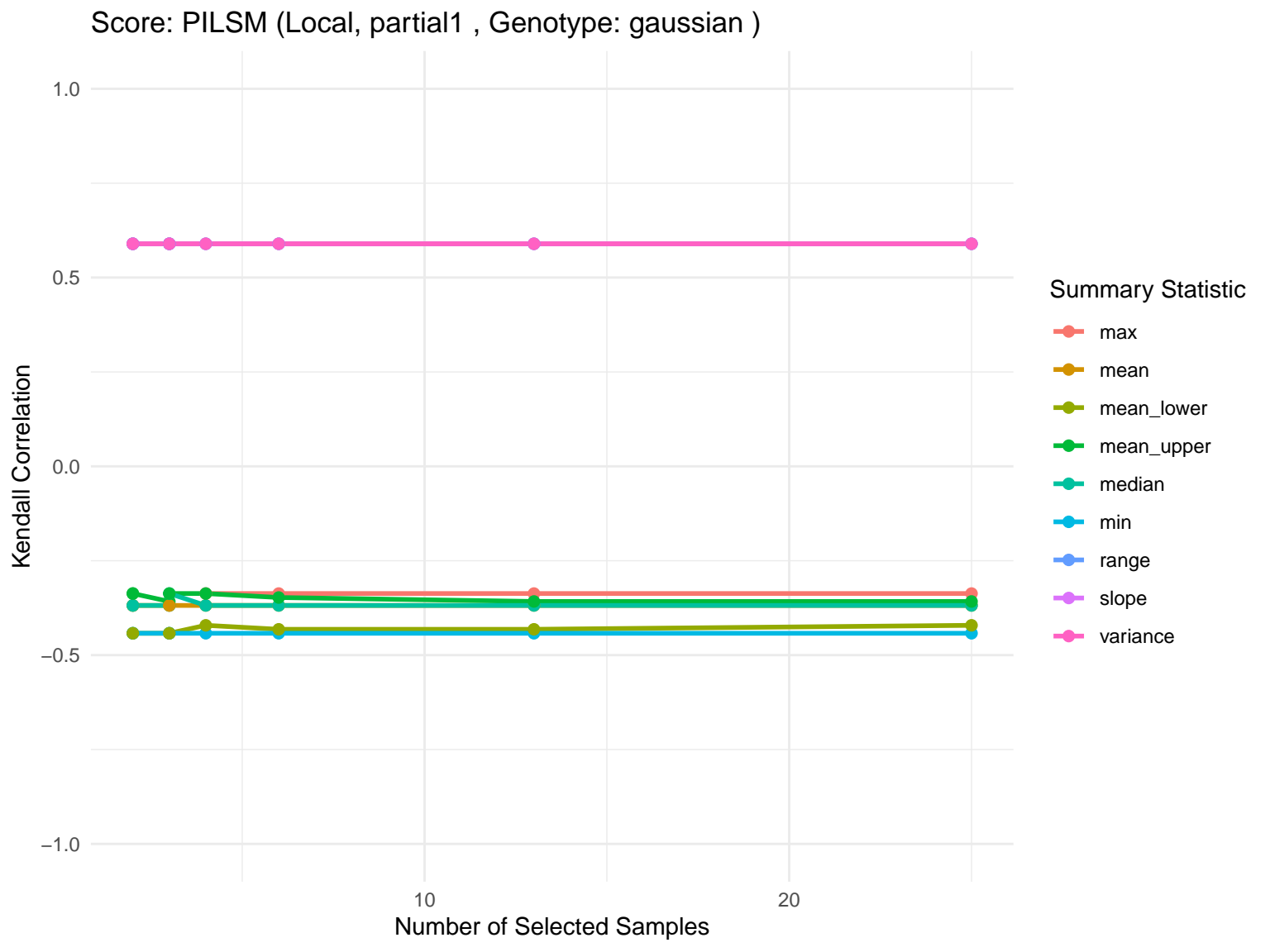
-0.5

-1.0

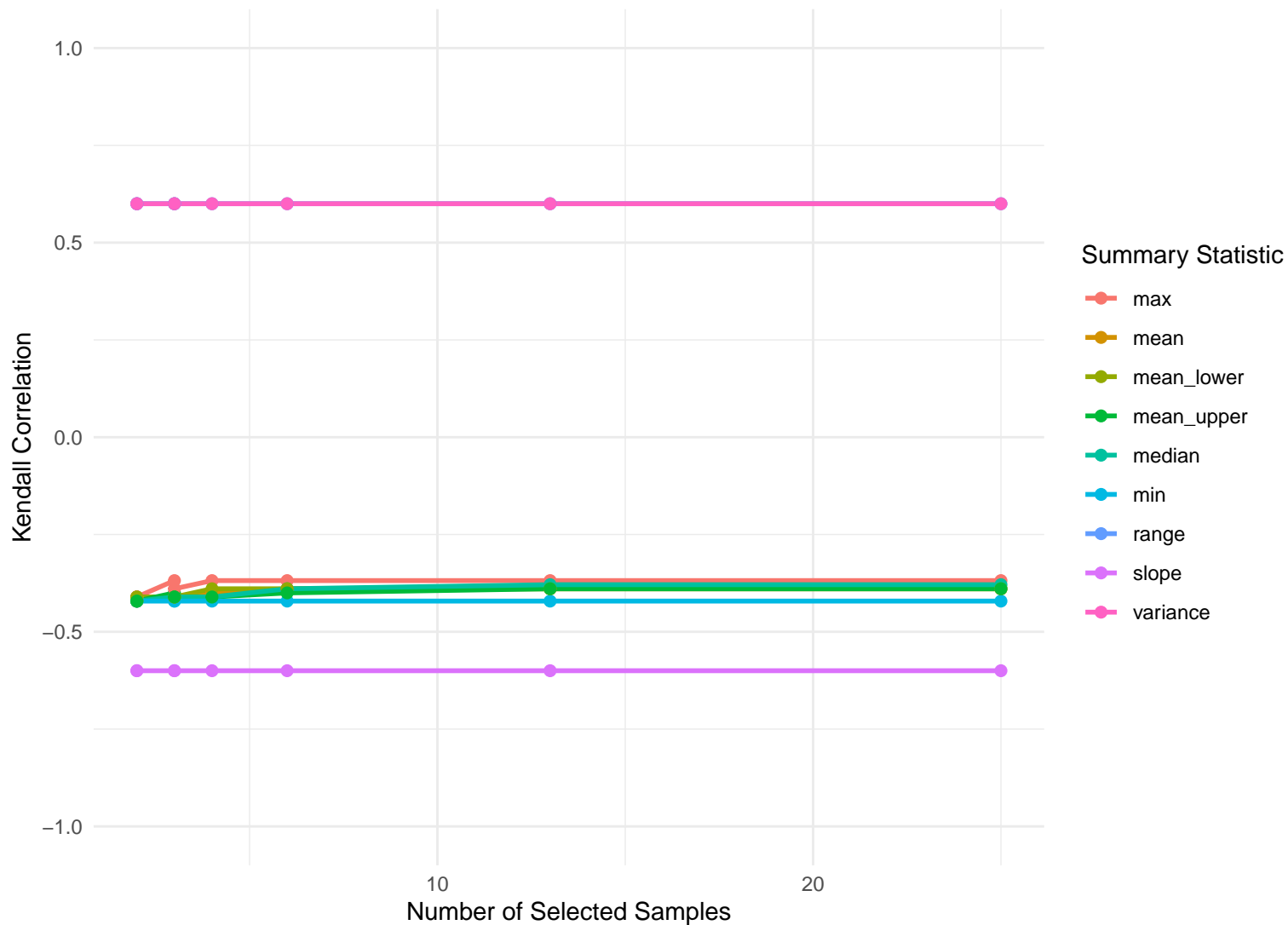
Number of Selected Samples

10

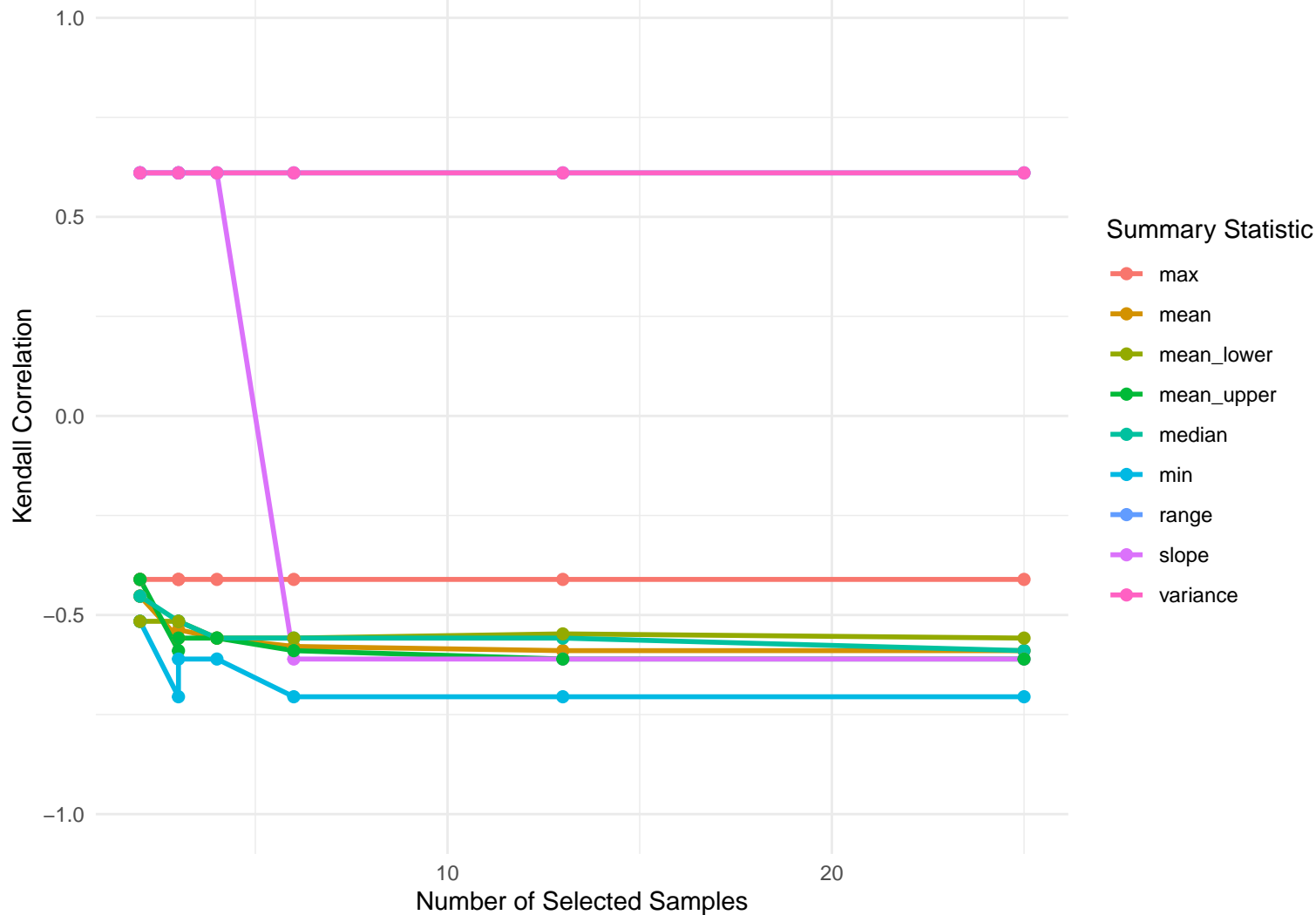
20



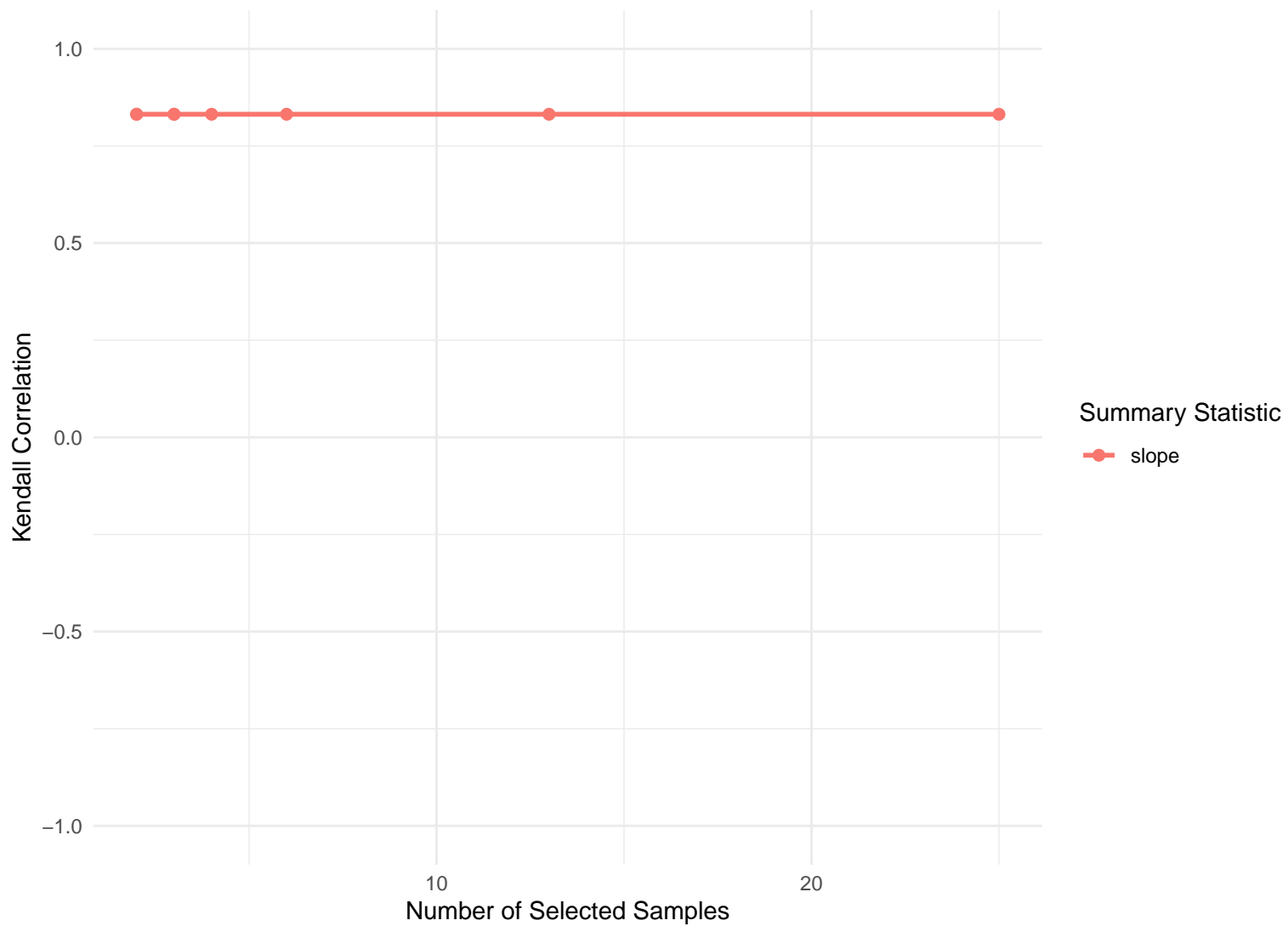
Score: PILSM (Local, partial1 , Genotype: sinusoidal)



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



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

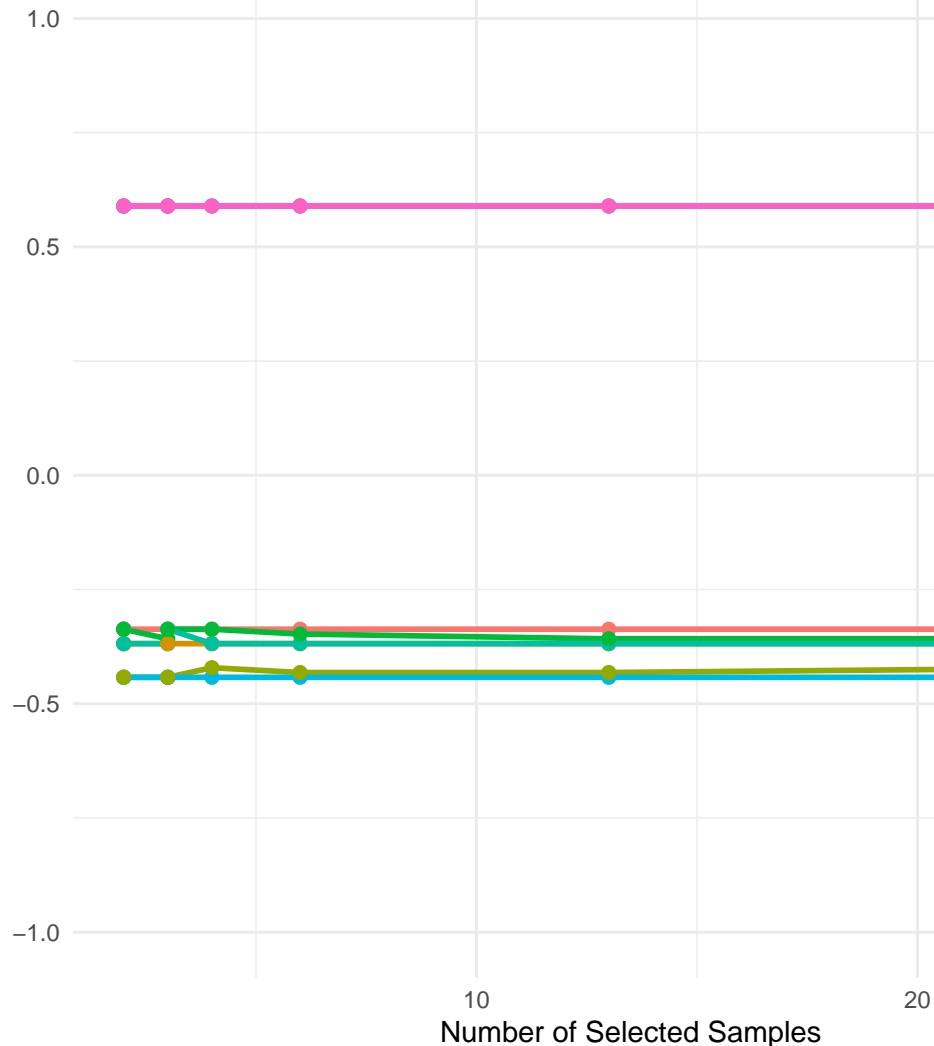


Score: RTR (Local, partial1 , Genotype: gaussian)

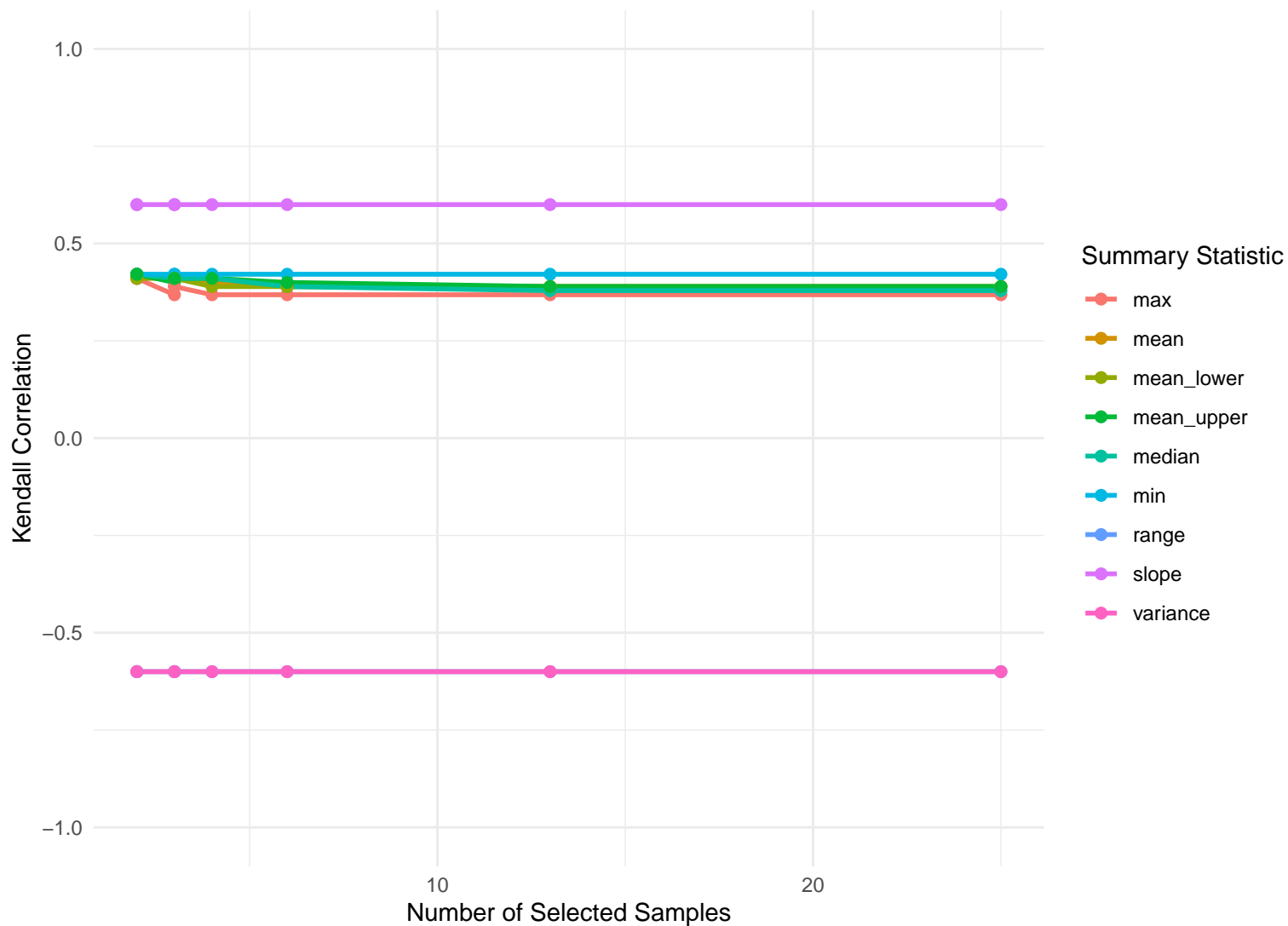
Kendall Correlation

Summary Statistic

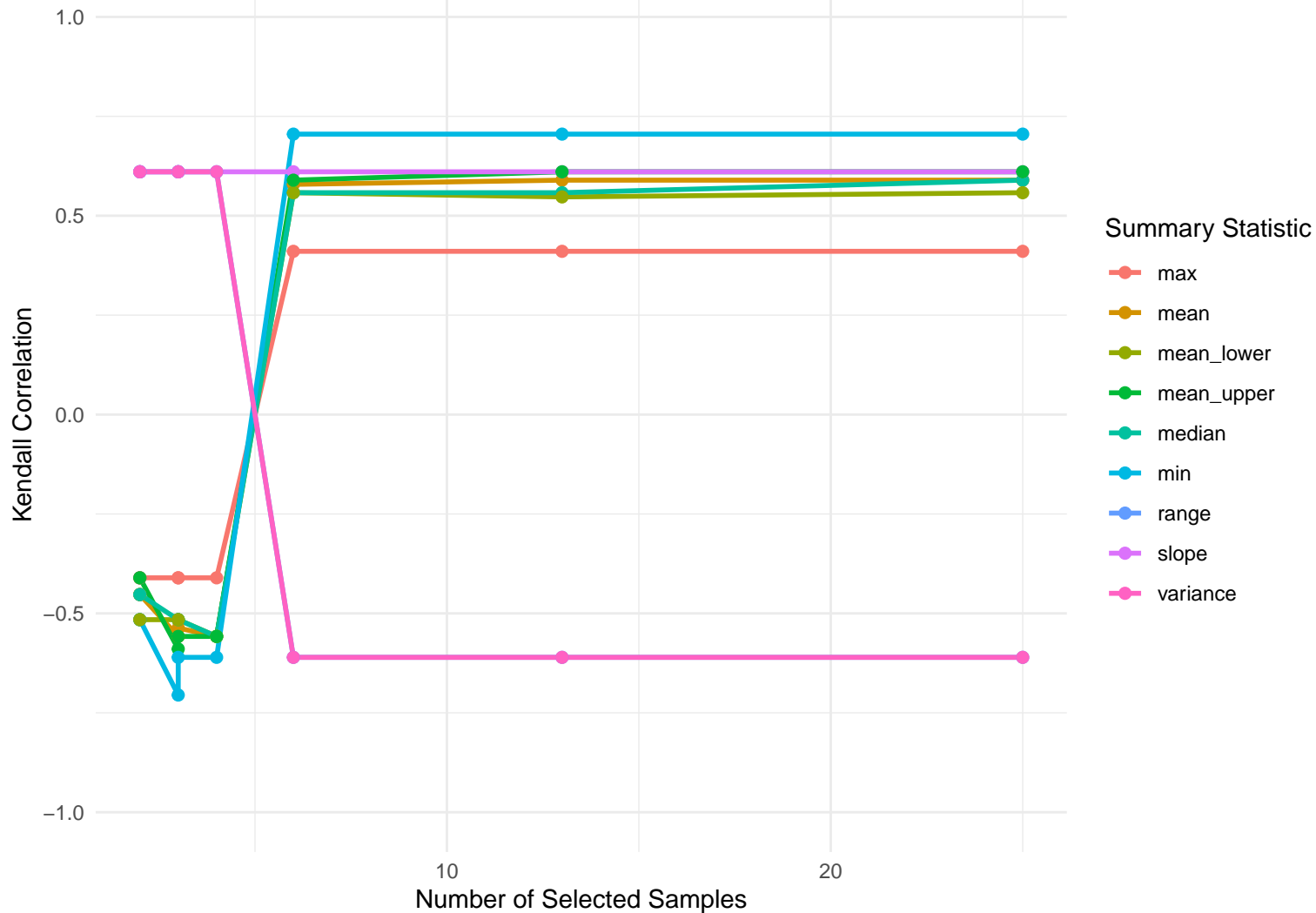
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



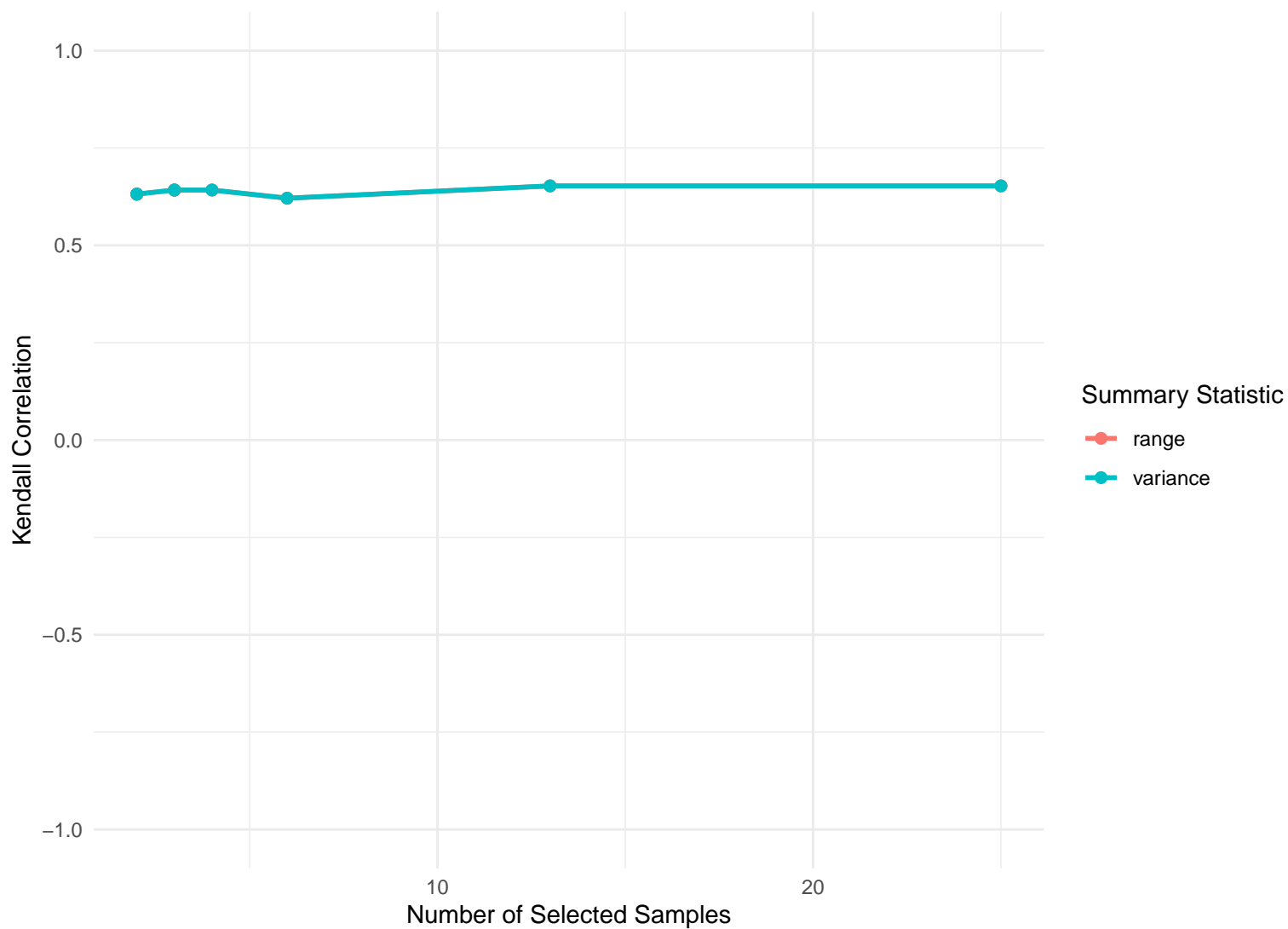
Score: RTR (Local, partial1 , Genotype: sinusoidal)



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



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

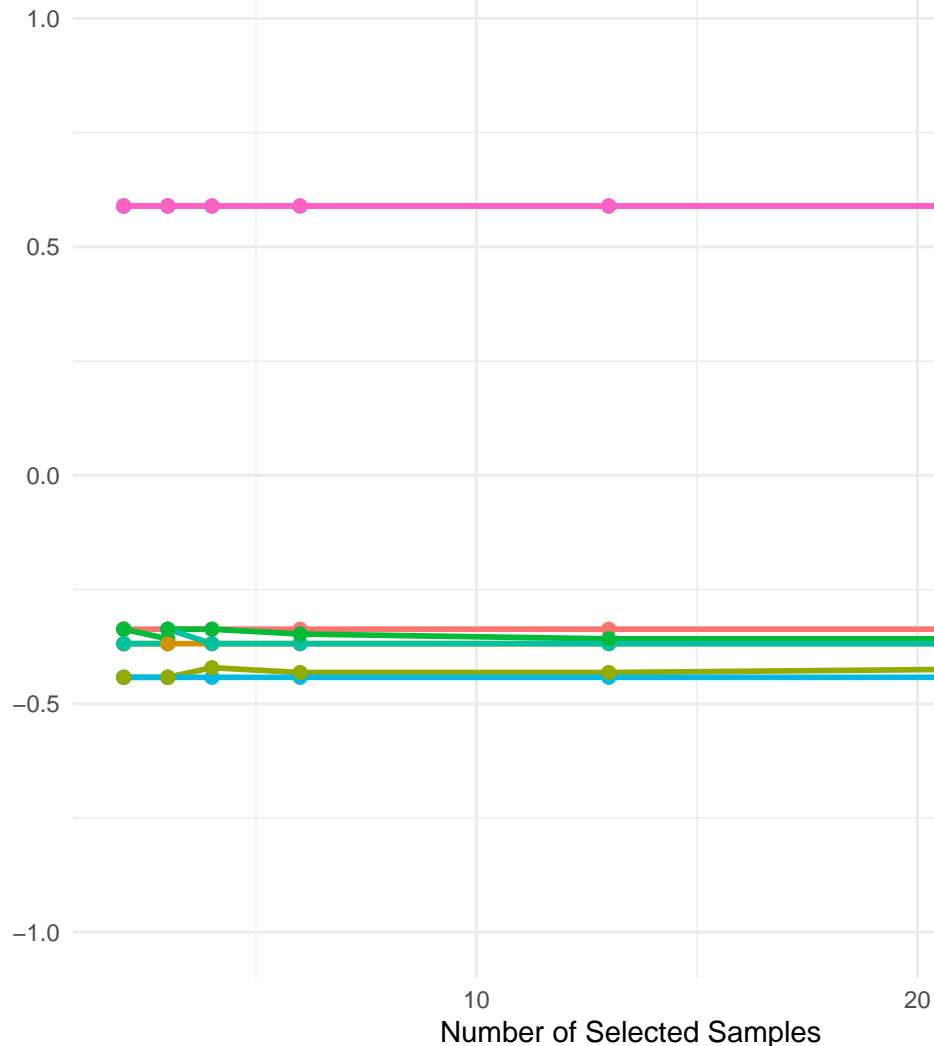


Score: PIR (Local, partial1 , Genotype: gaussian)

Kendall Correlation

Summary Statistic

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



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

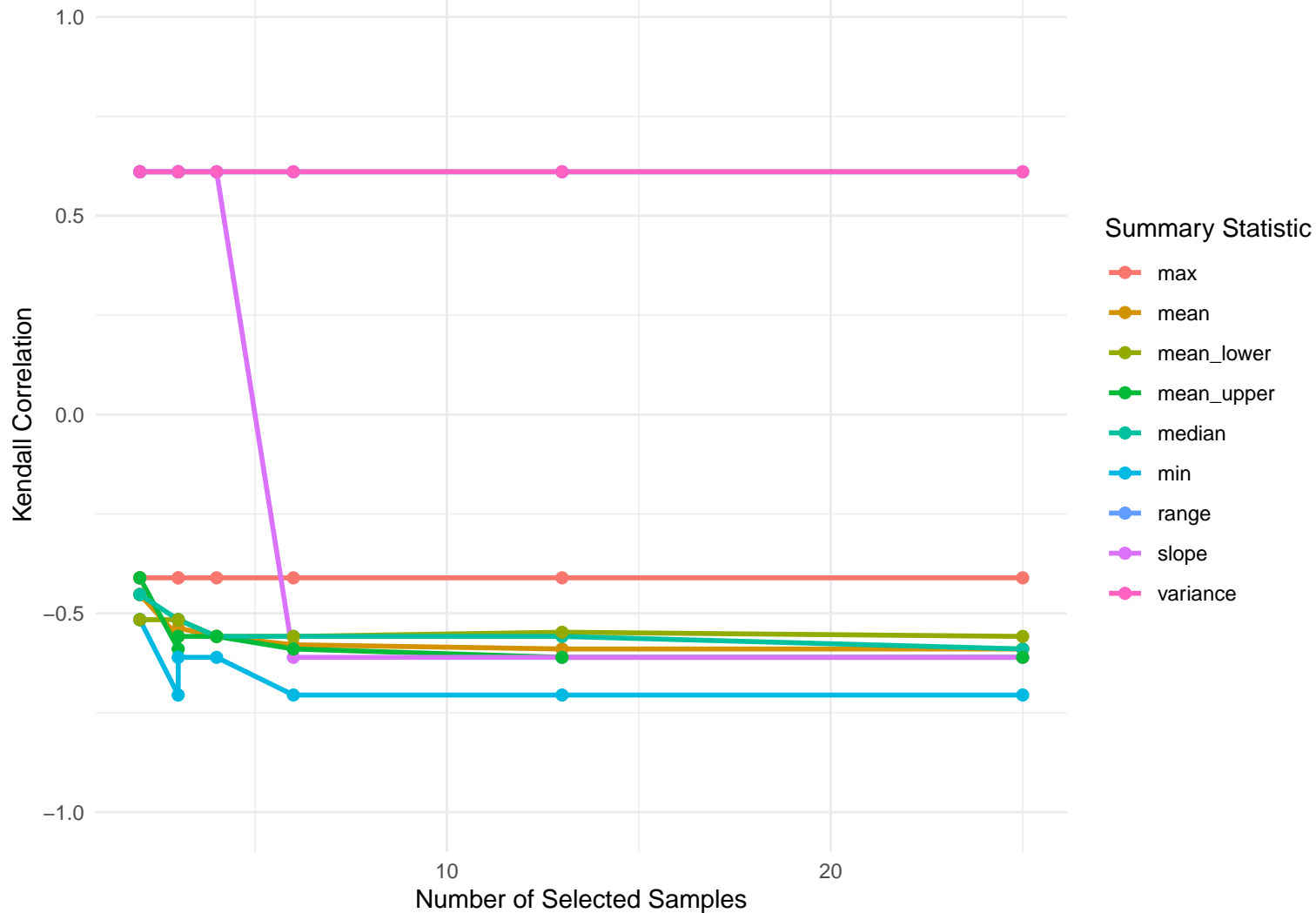


Number of Selected Samples

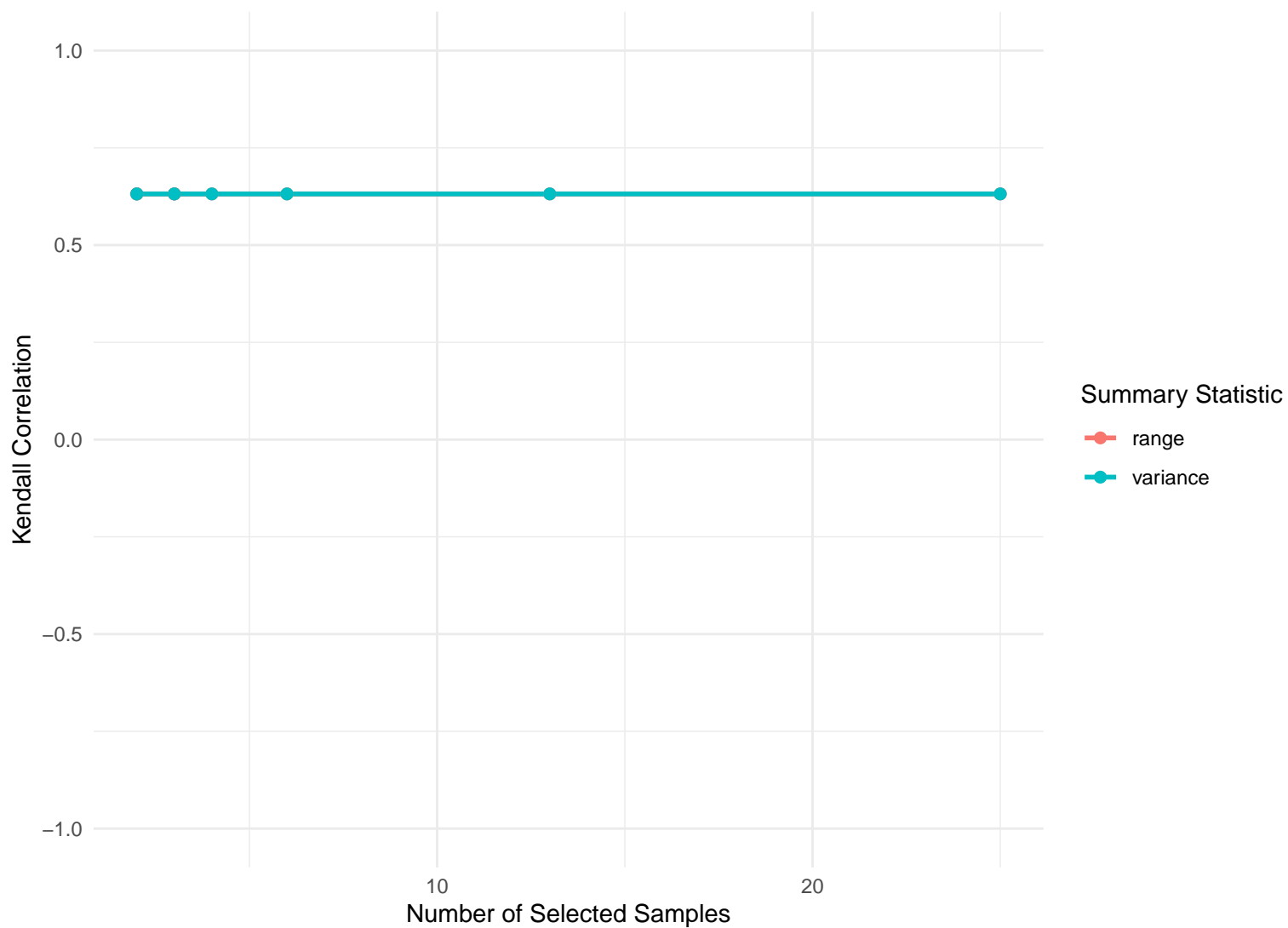
Summary Statistic

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

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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

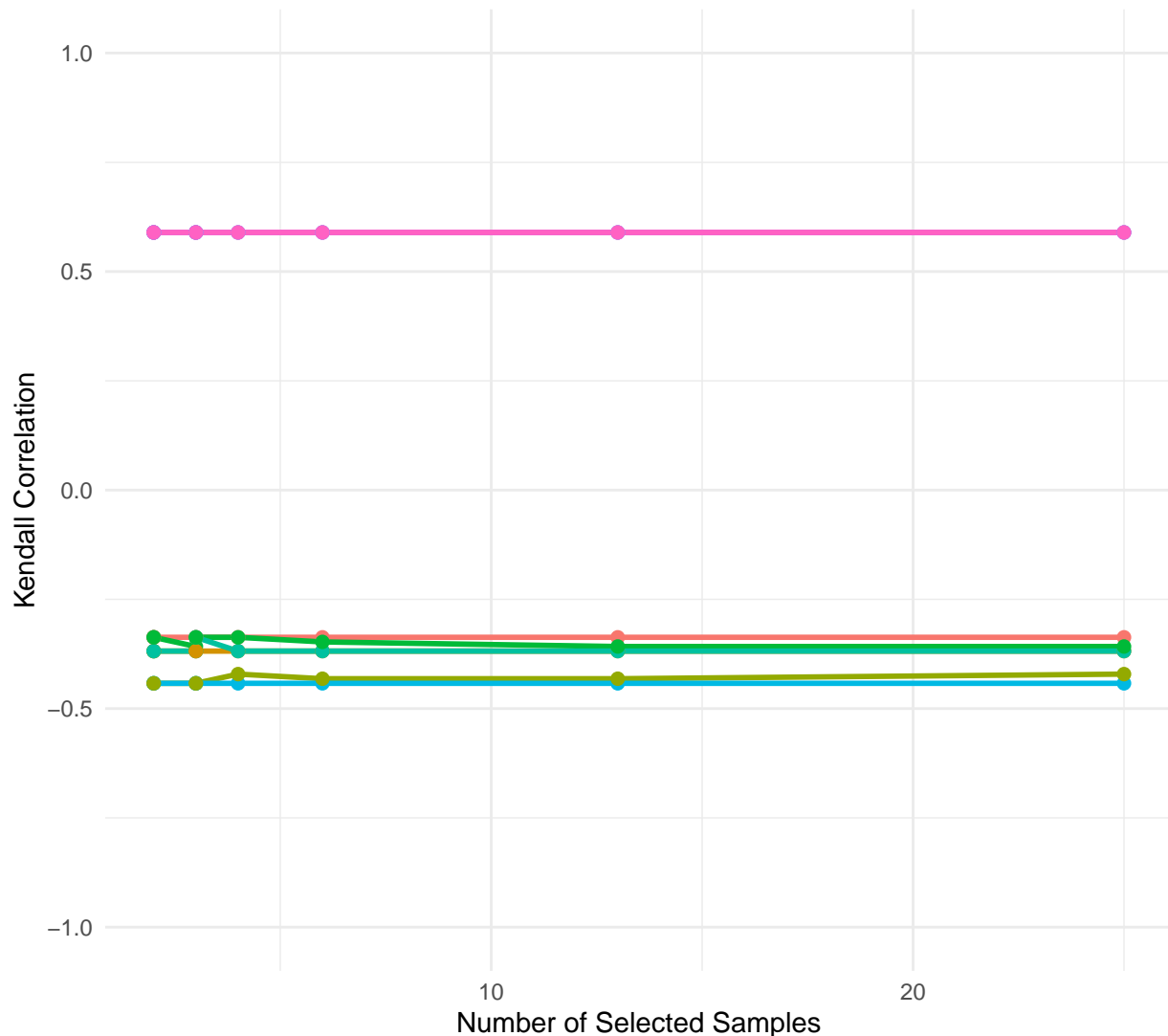
Number of Selected Samples

10

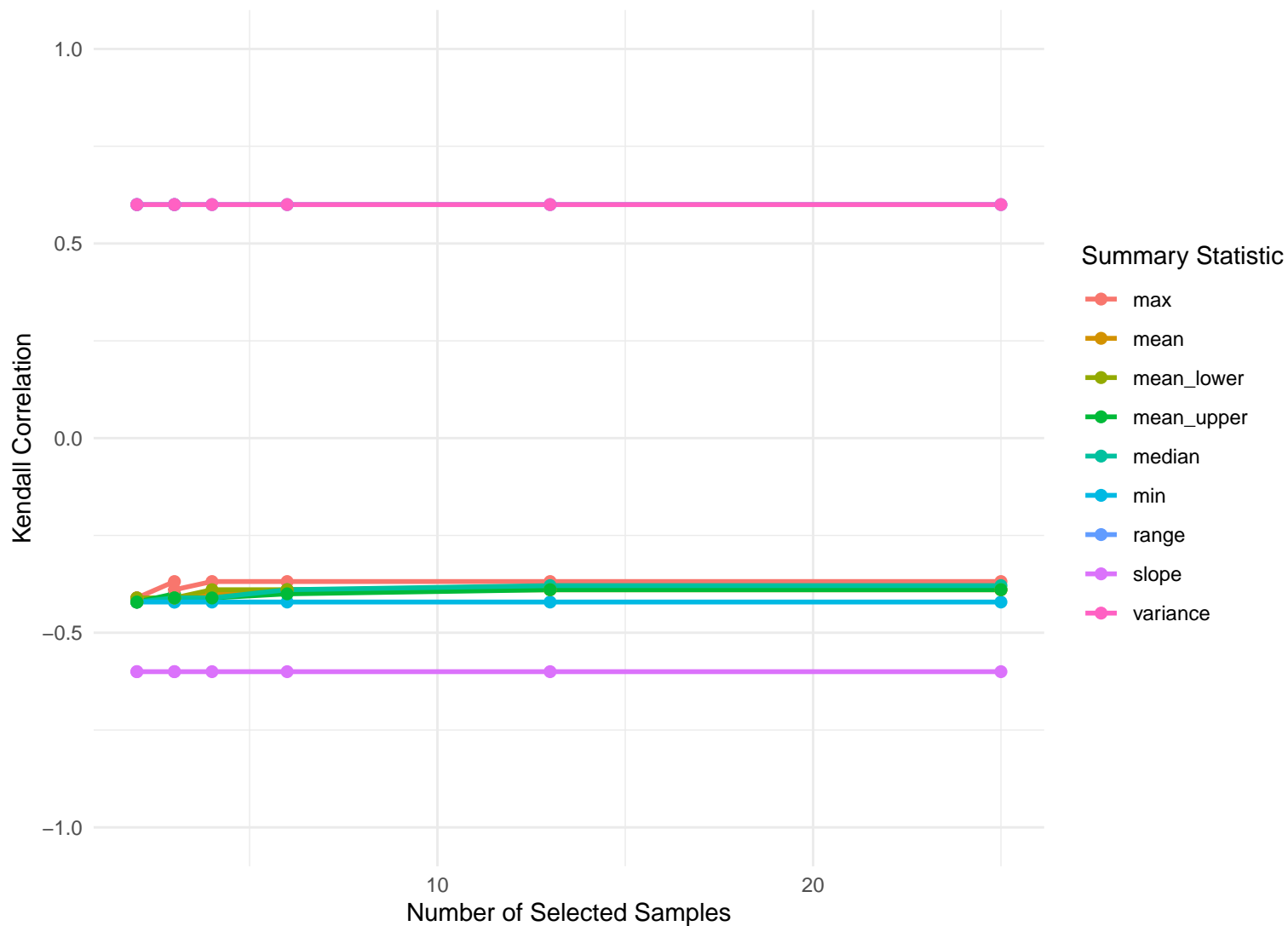
20

Summary Statistic

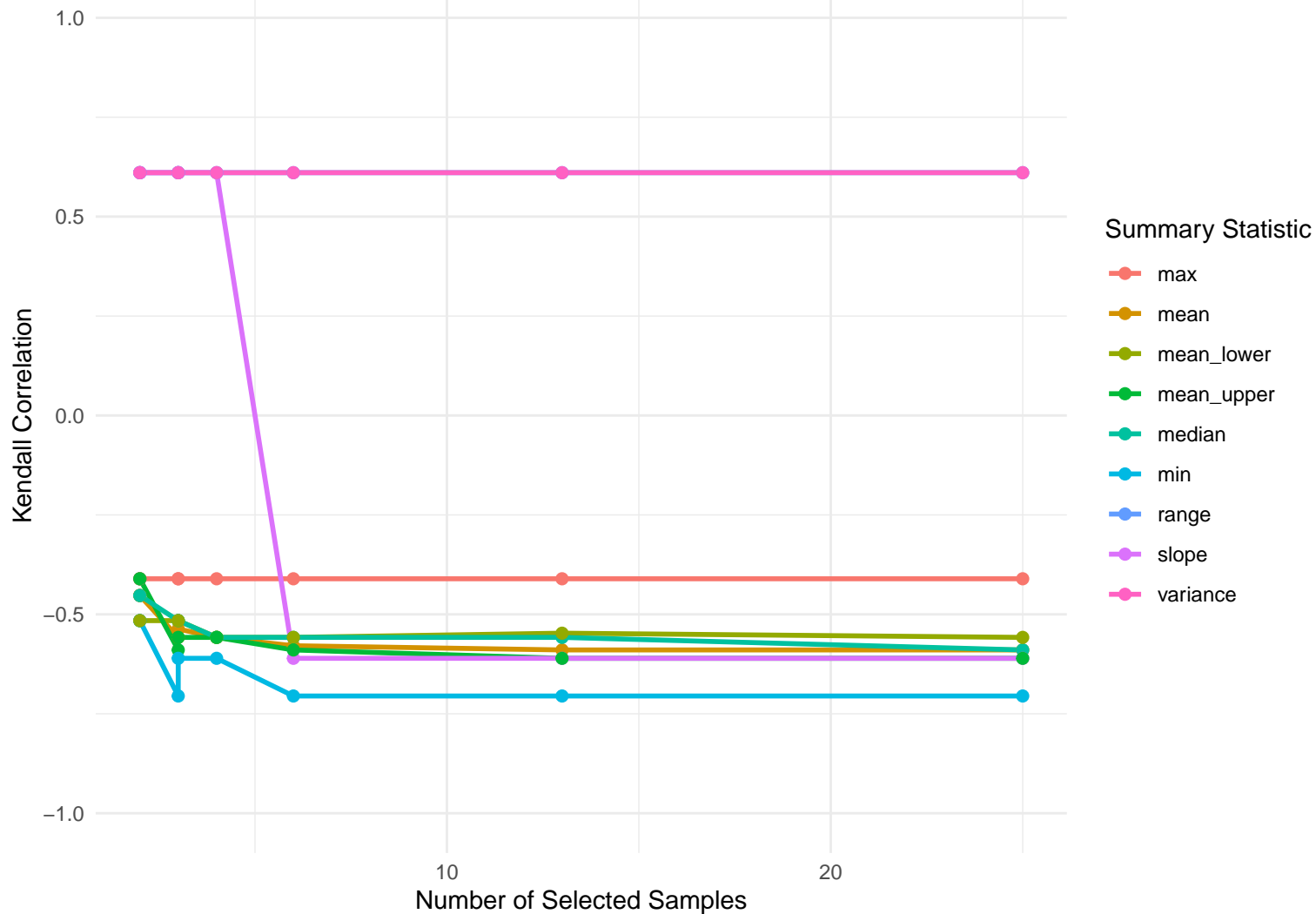
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

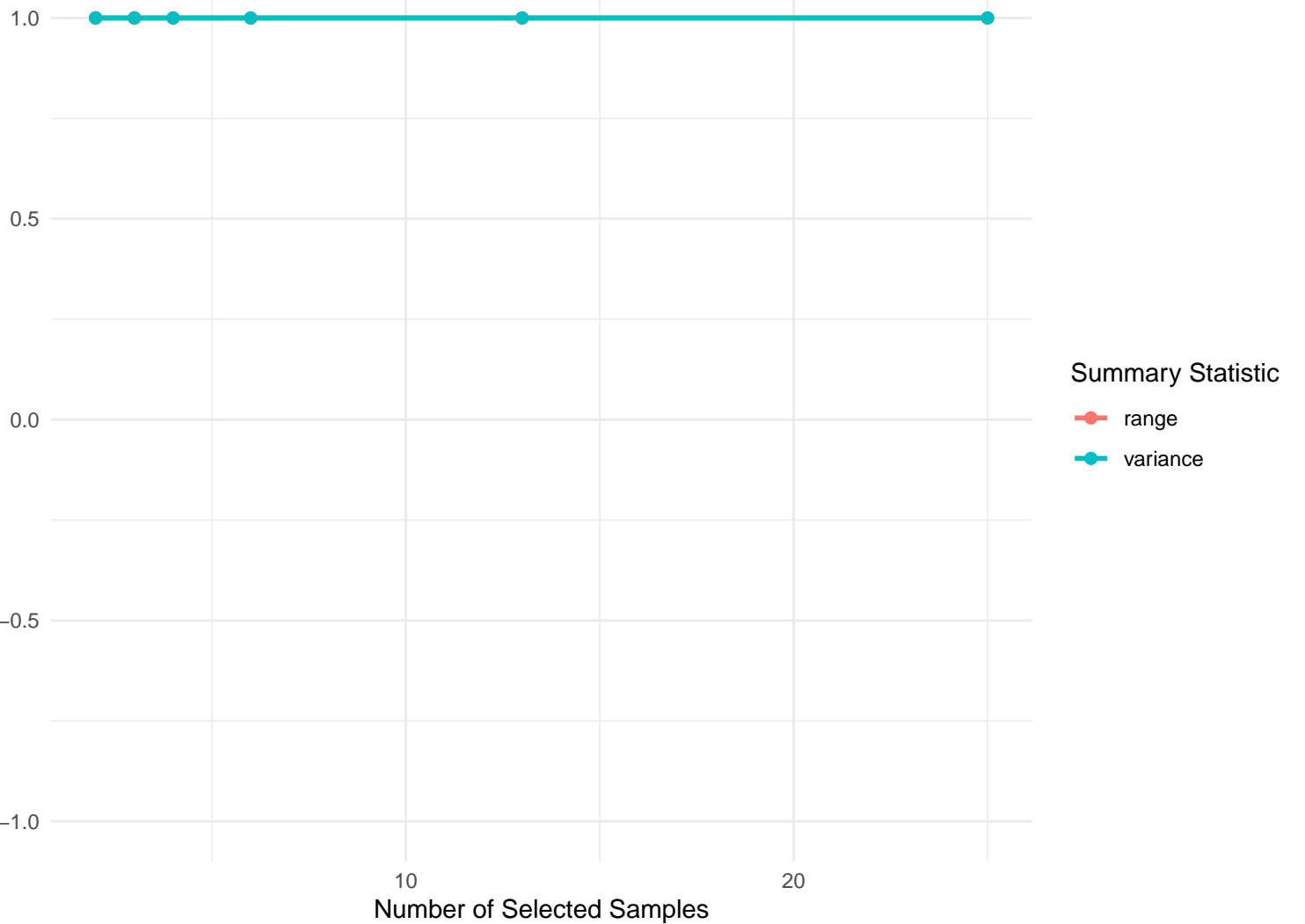
20

Number of Selected Samples

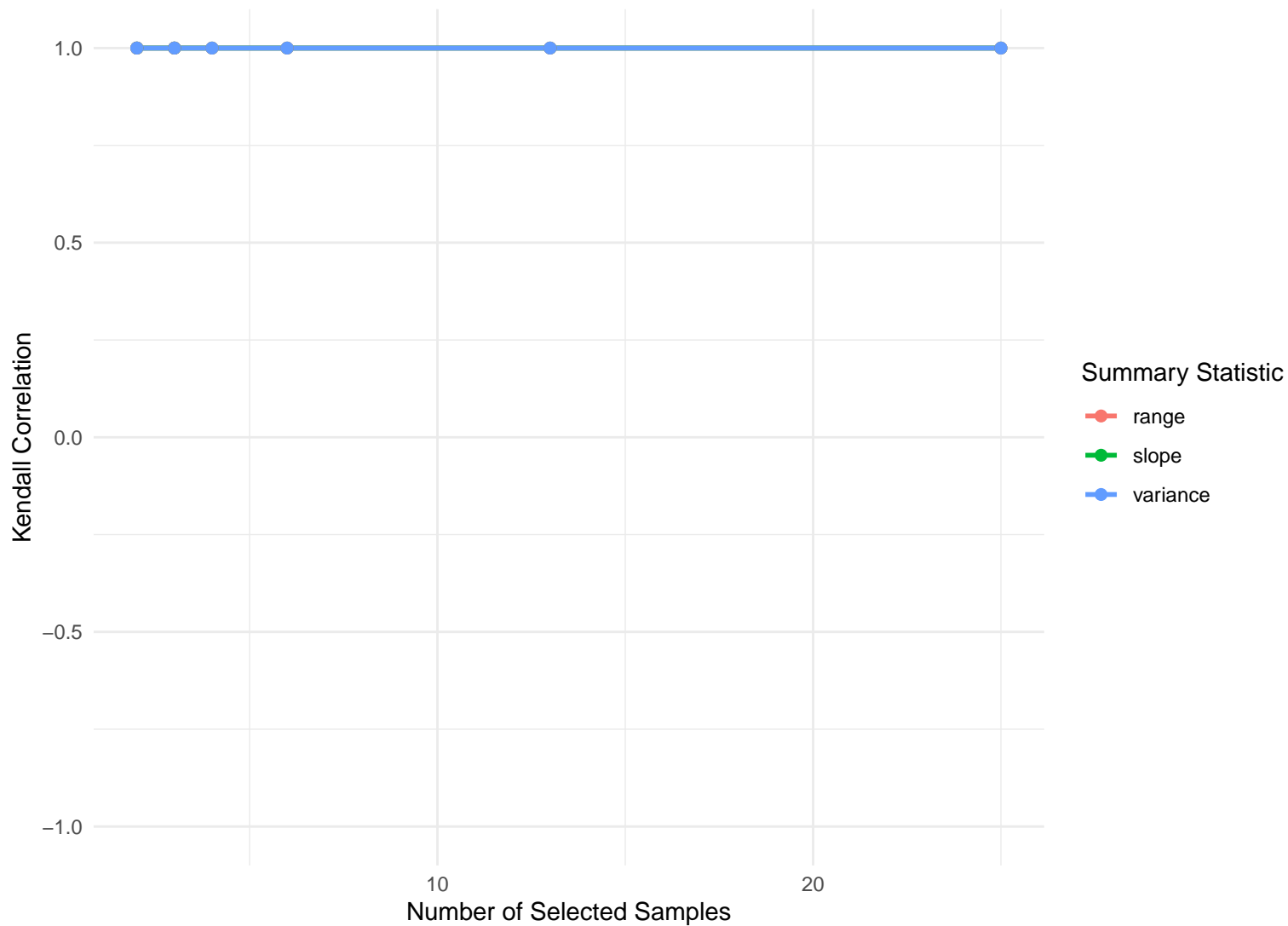
Summary Statistic

range

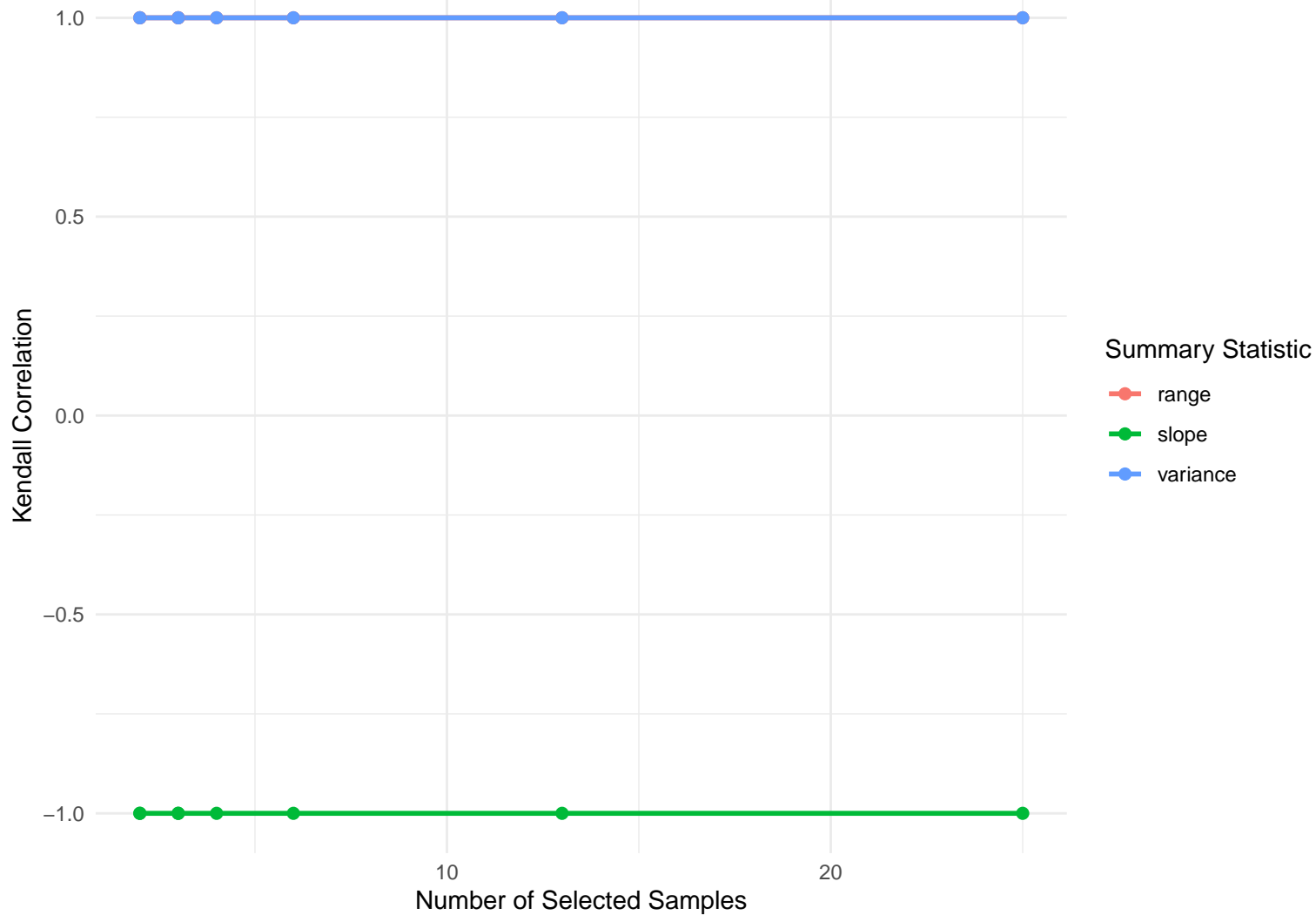
variance



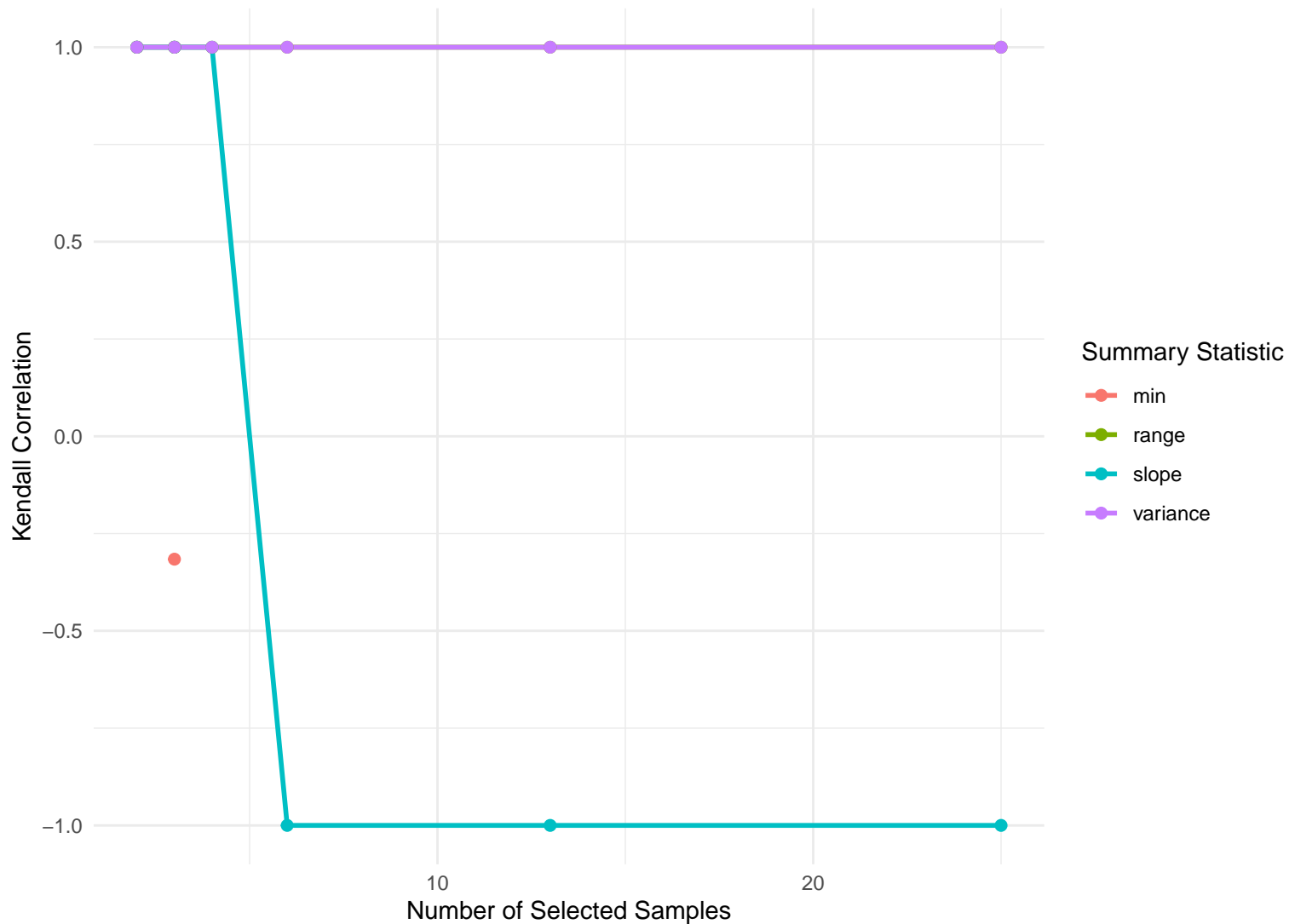
Score: ESPI (Local, partial1 , Genotype: gaussian)



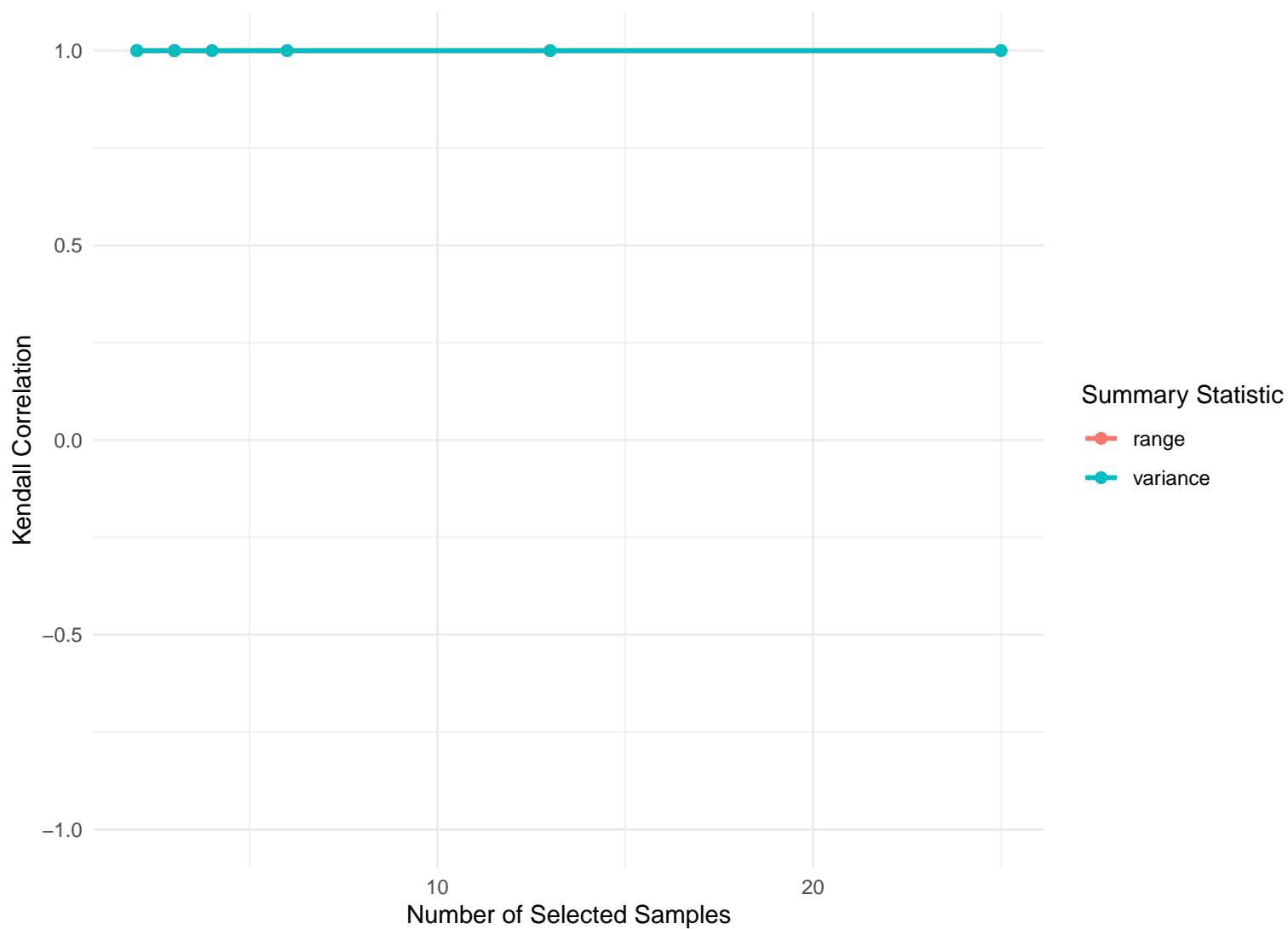
Score: ESPI (Local, partial1 , Genotype: sinusoidal)



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

20

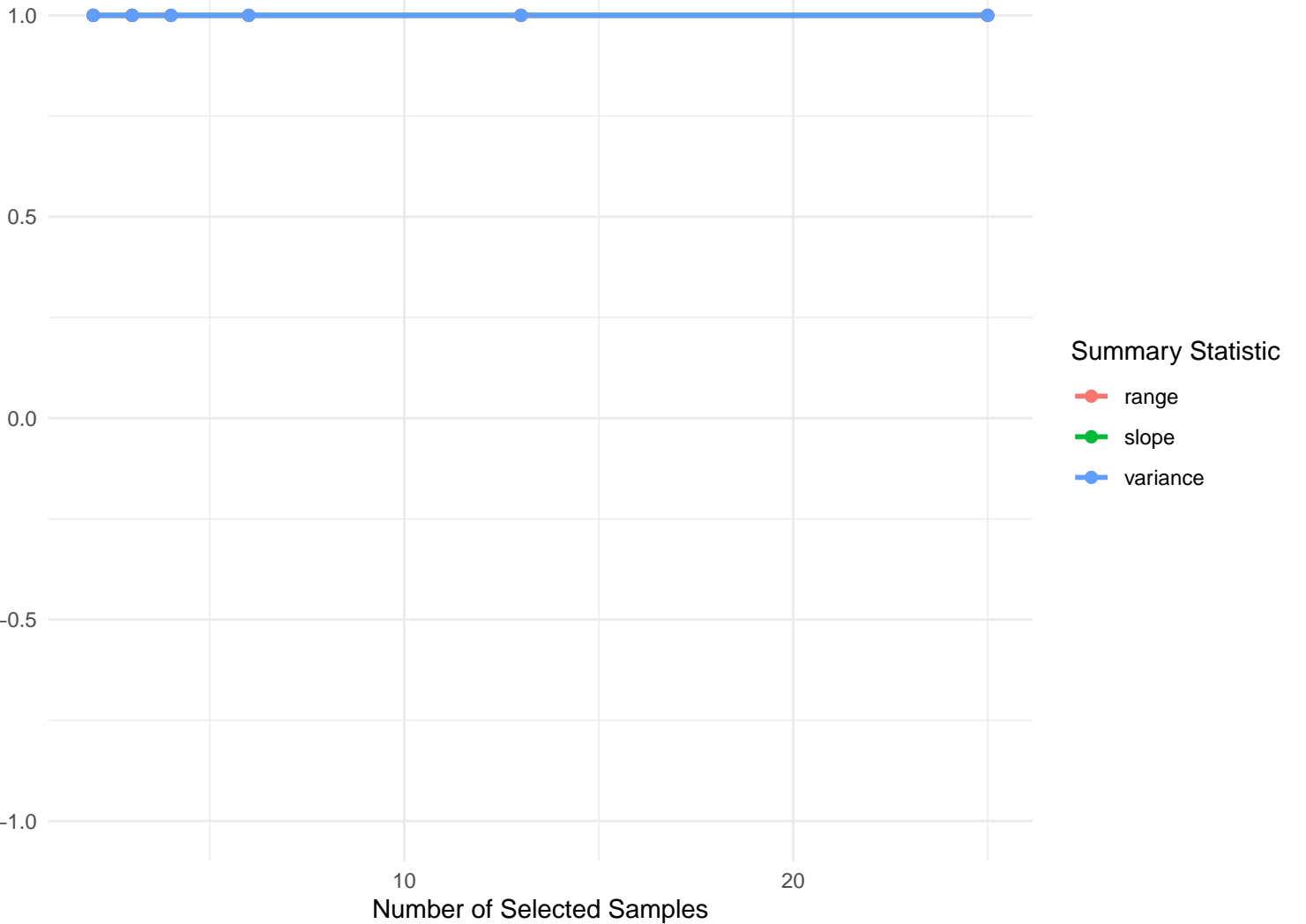
Number of Selected Samples

Summary Statistic

range

slope

variance



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

Kendall Correlation

Summary Statistic

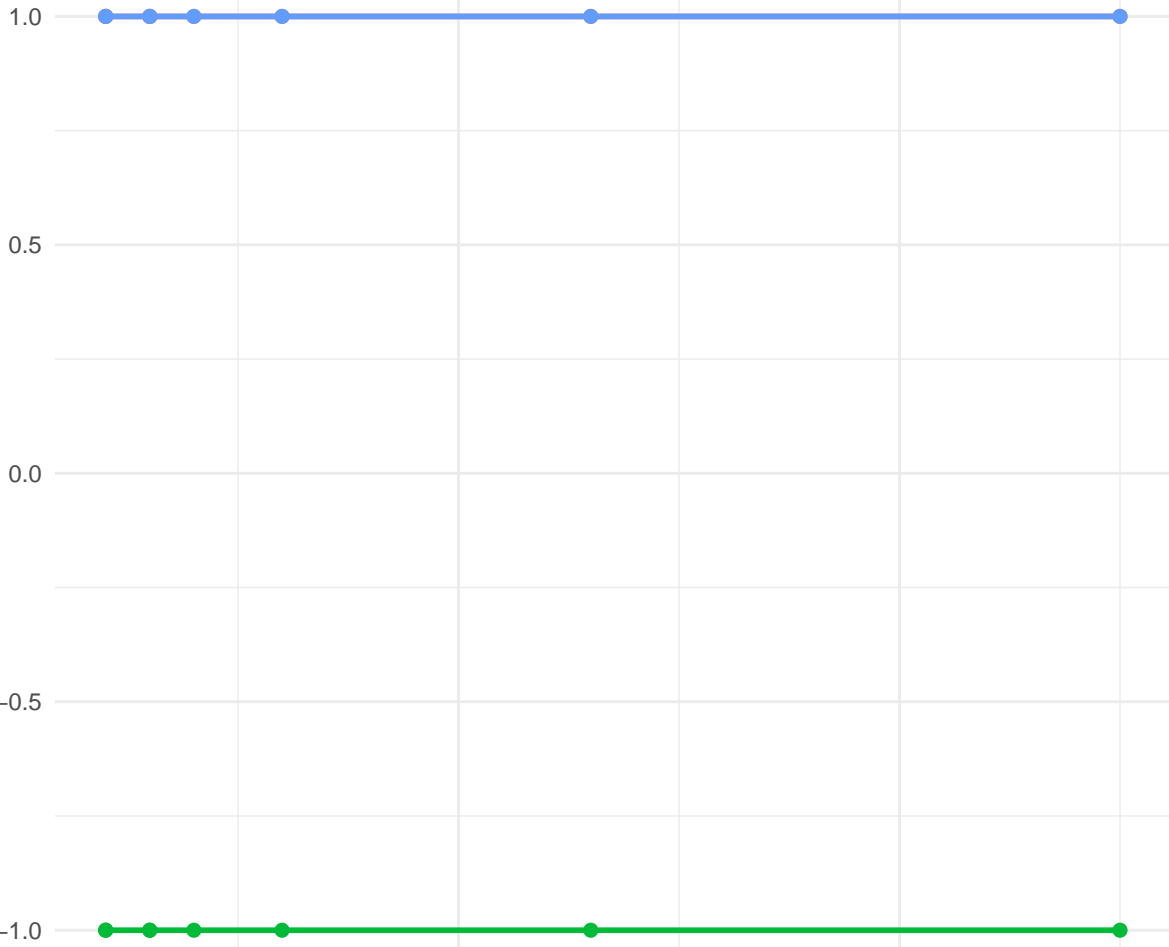
- range
- slope
- variance

1.0
0.5
0.0
-0.5
-1.0

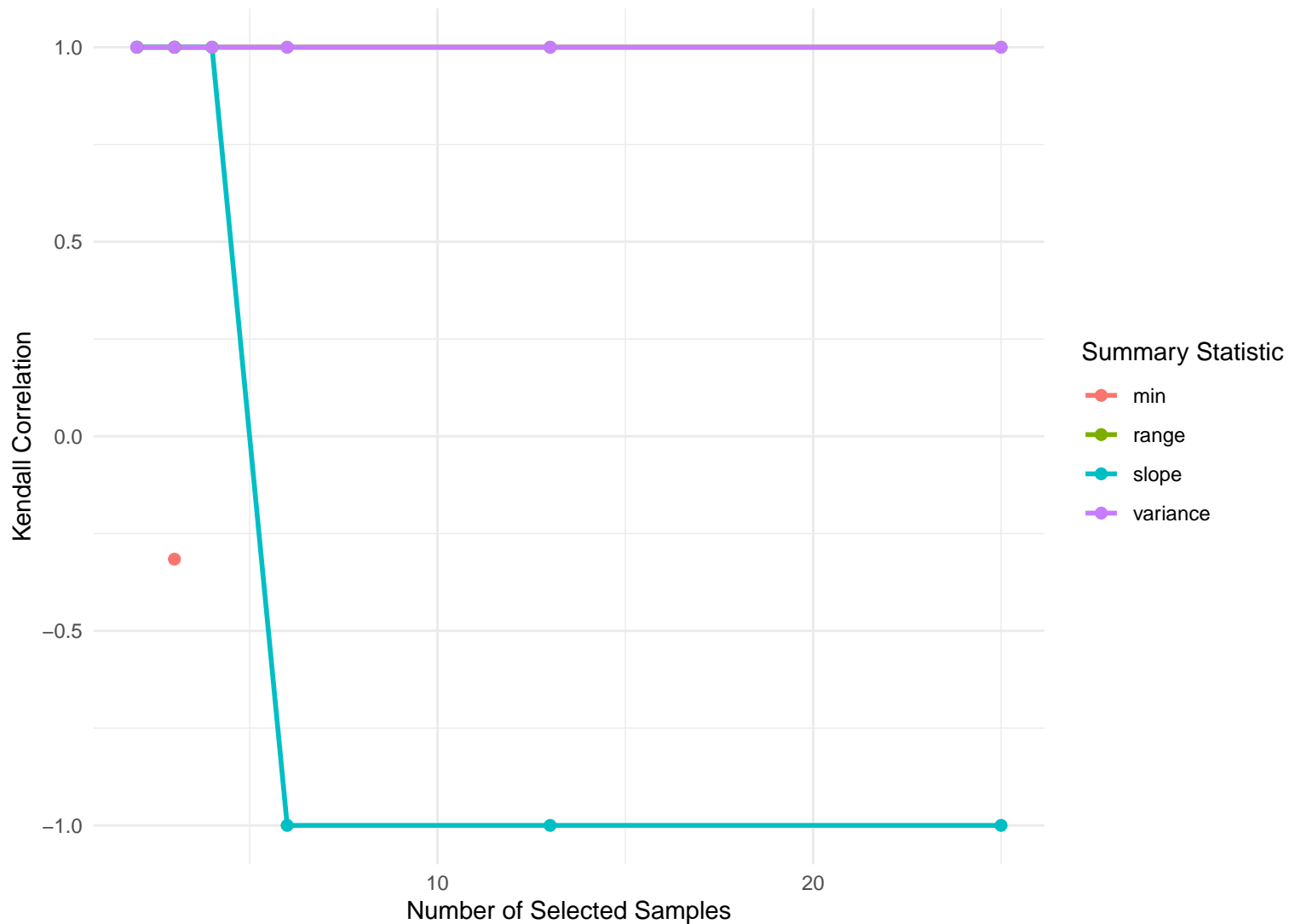
Number of Selected Samples

10

20



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

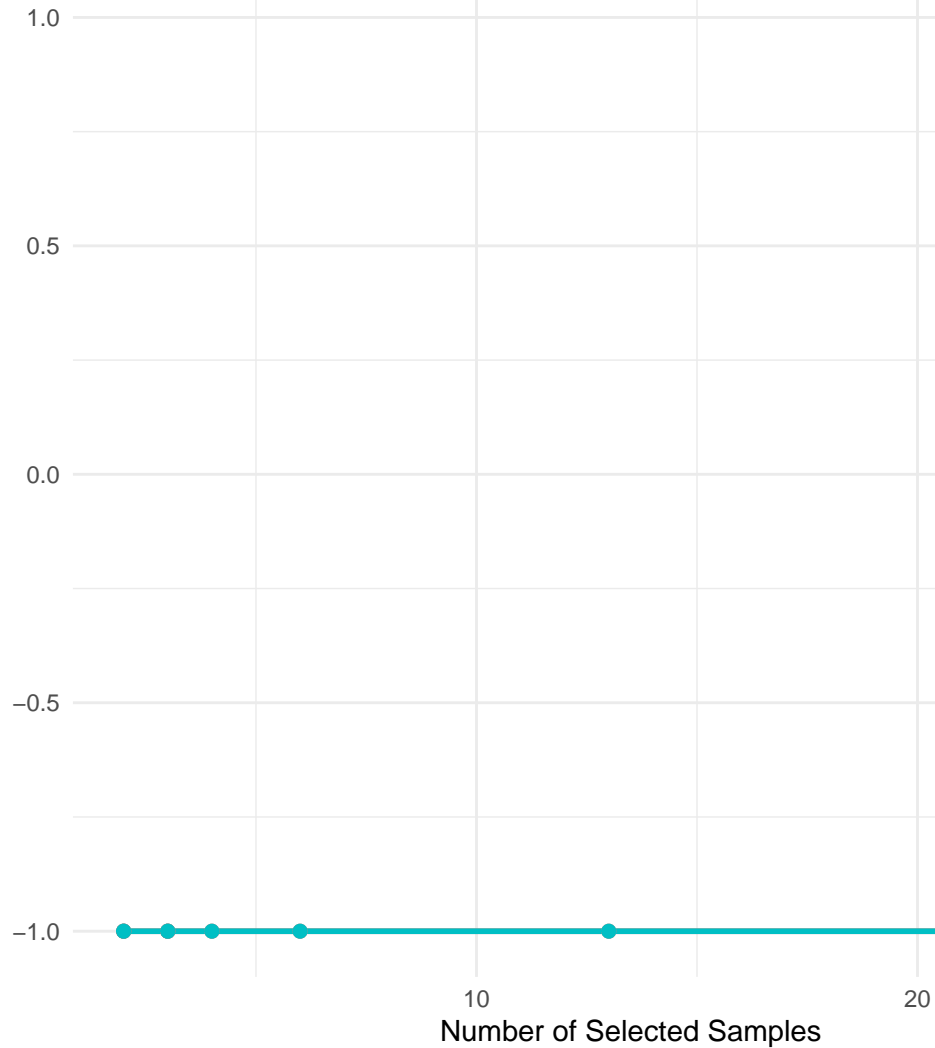


Score: PSI (Local, partial1 , Genotype: linear)

Kendall Correlation

Summary Statistic

- range
- variance

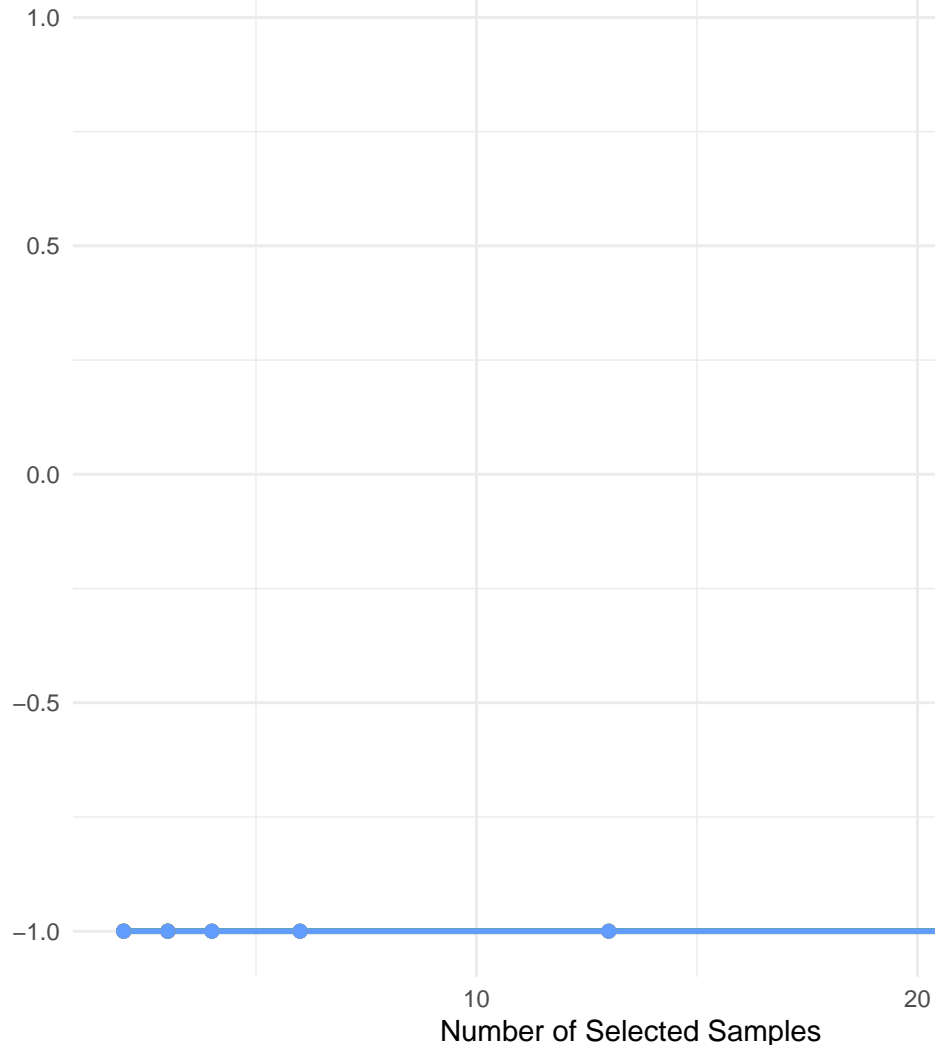


Score: PSI (Local, partial1 , Genotype: gaussian)

Kendall Correlation

Summary Statistic

- range
- slope
- variance

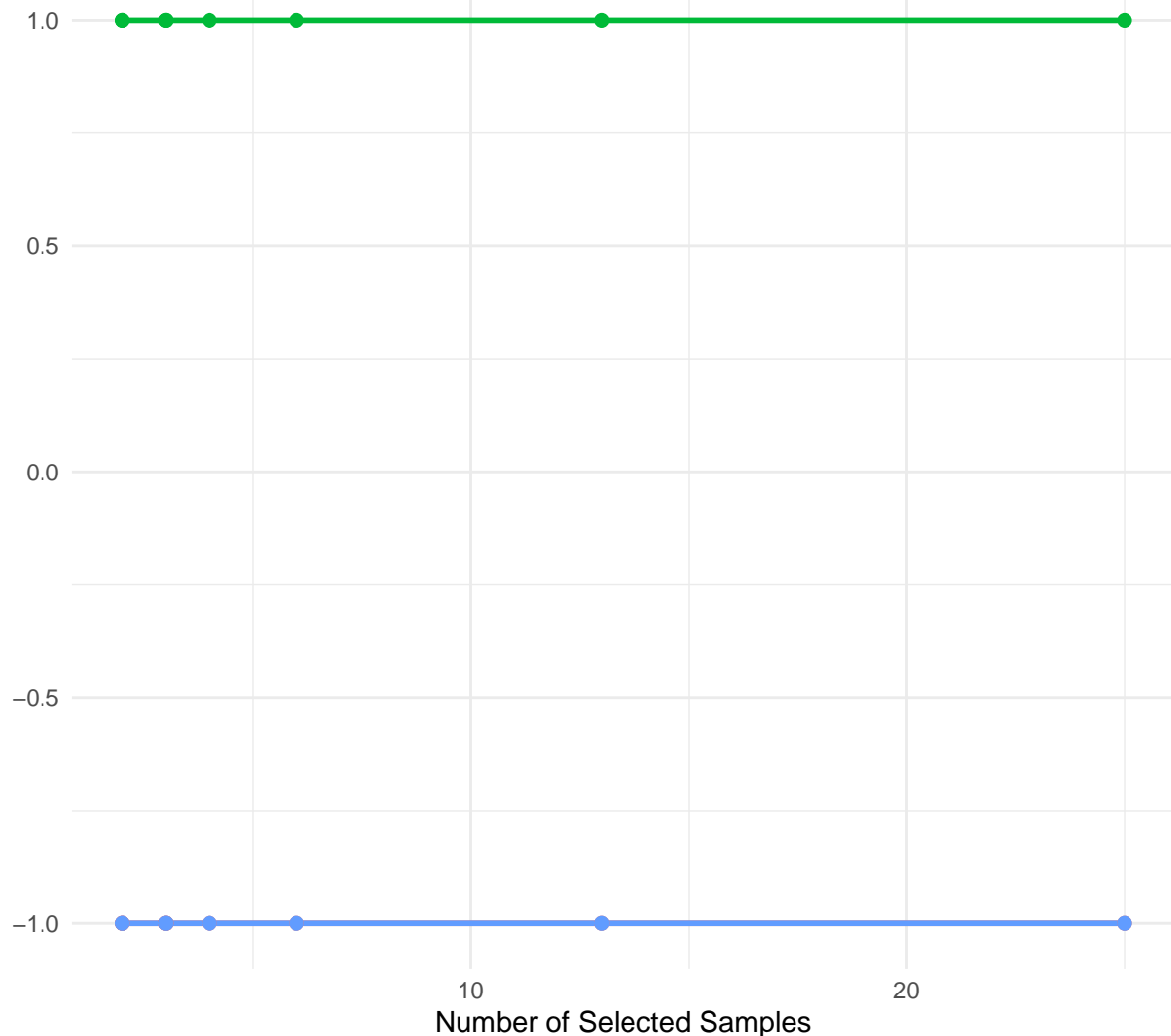


Score: PSI (Local, partial1 , Genotype: sinusoidal)

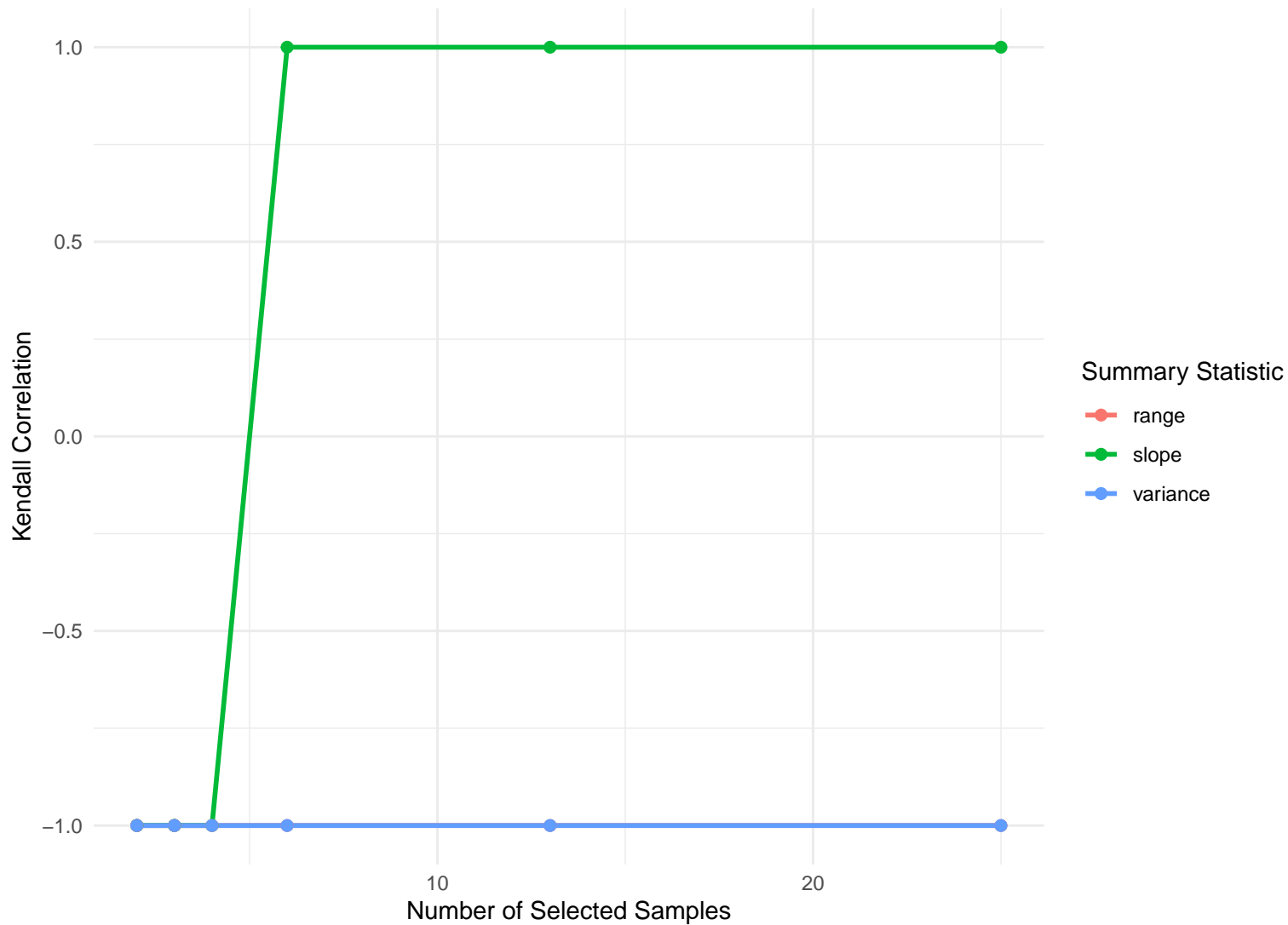
Kendall Correlation

Summary Statistic

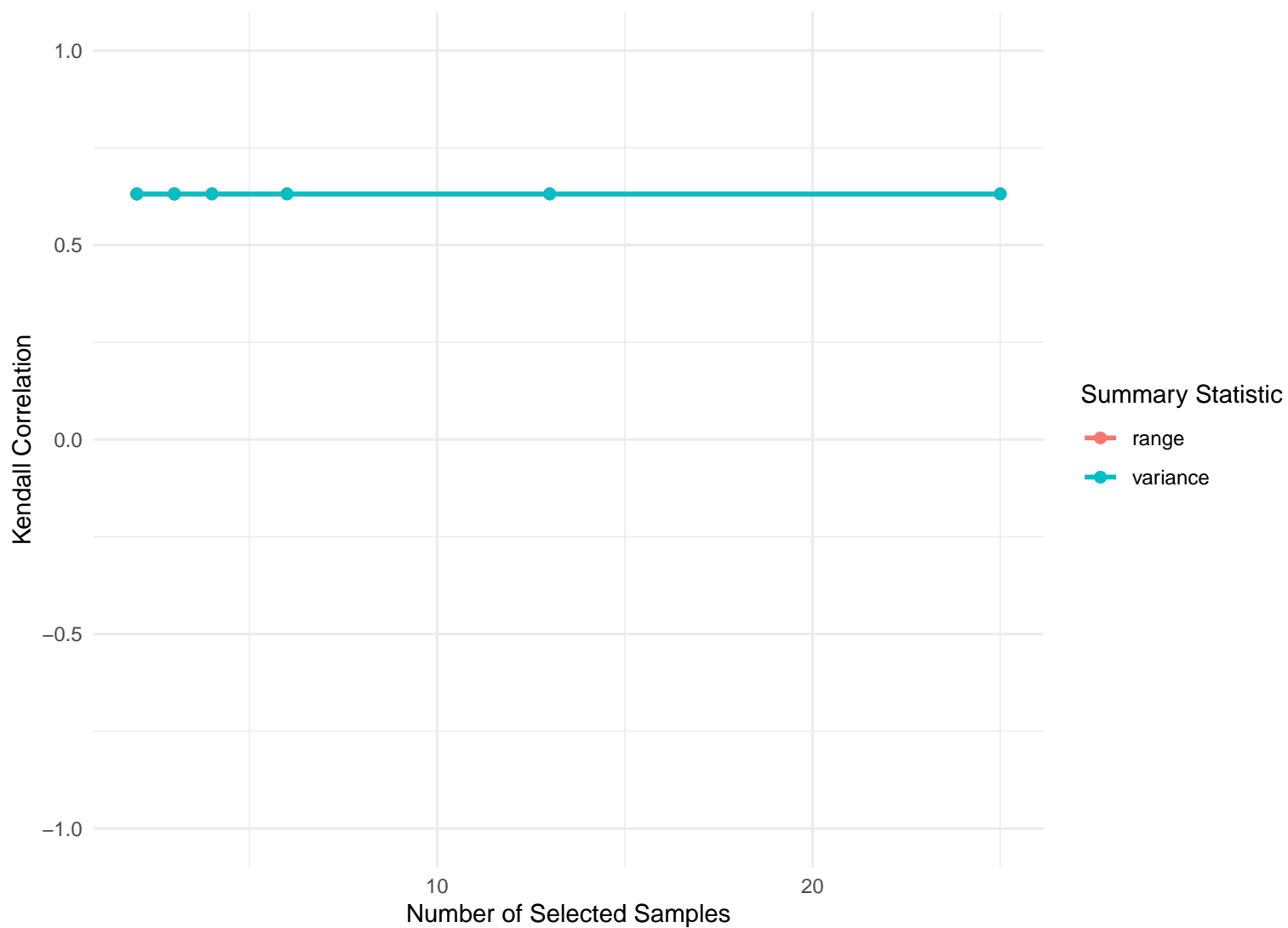
- range
- slope
- variance



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



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

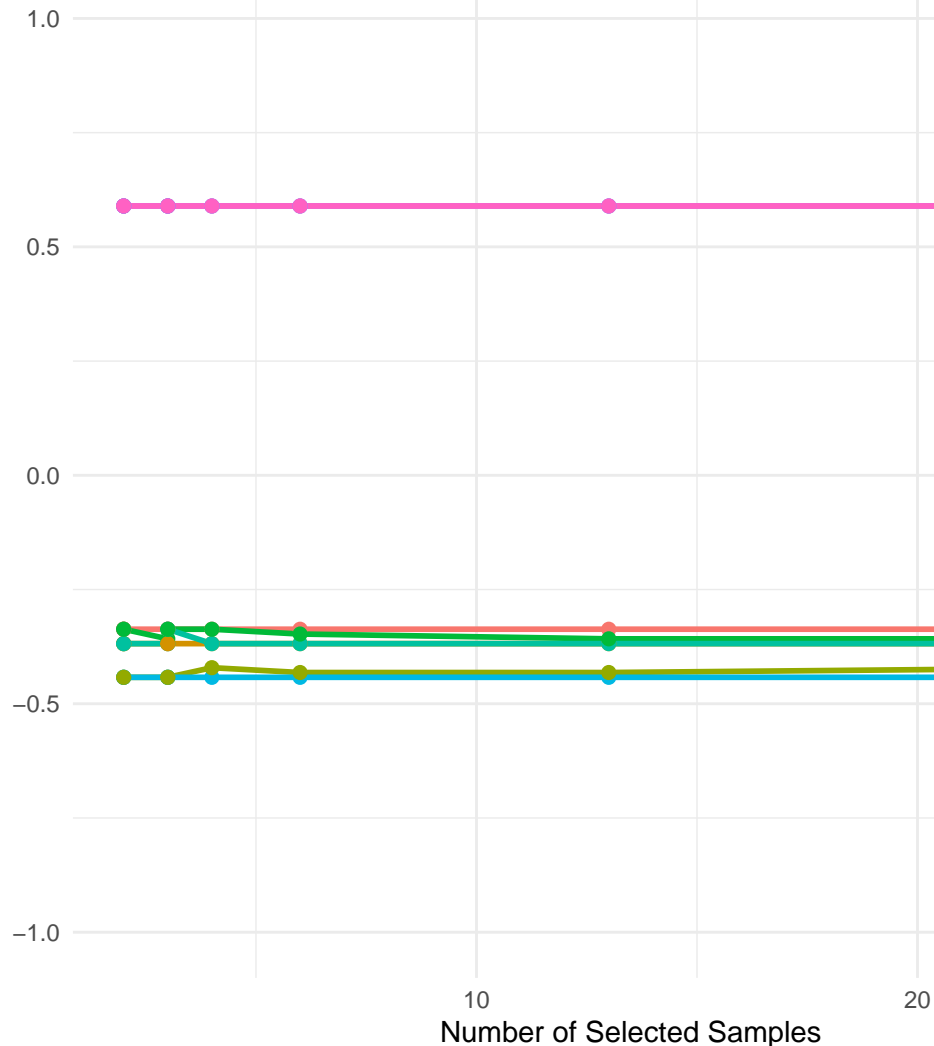


Score: RPI (Local, partial1 , Genotype: gaussian)

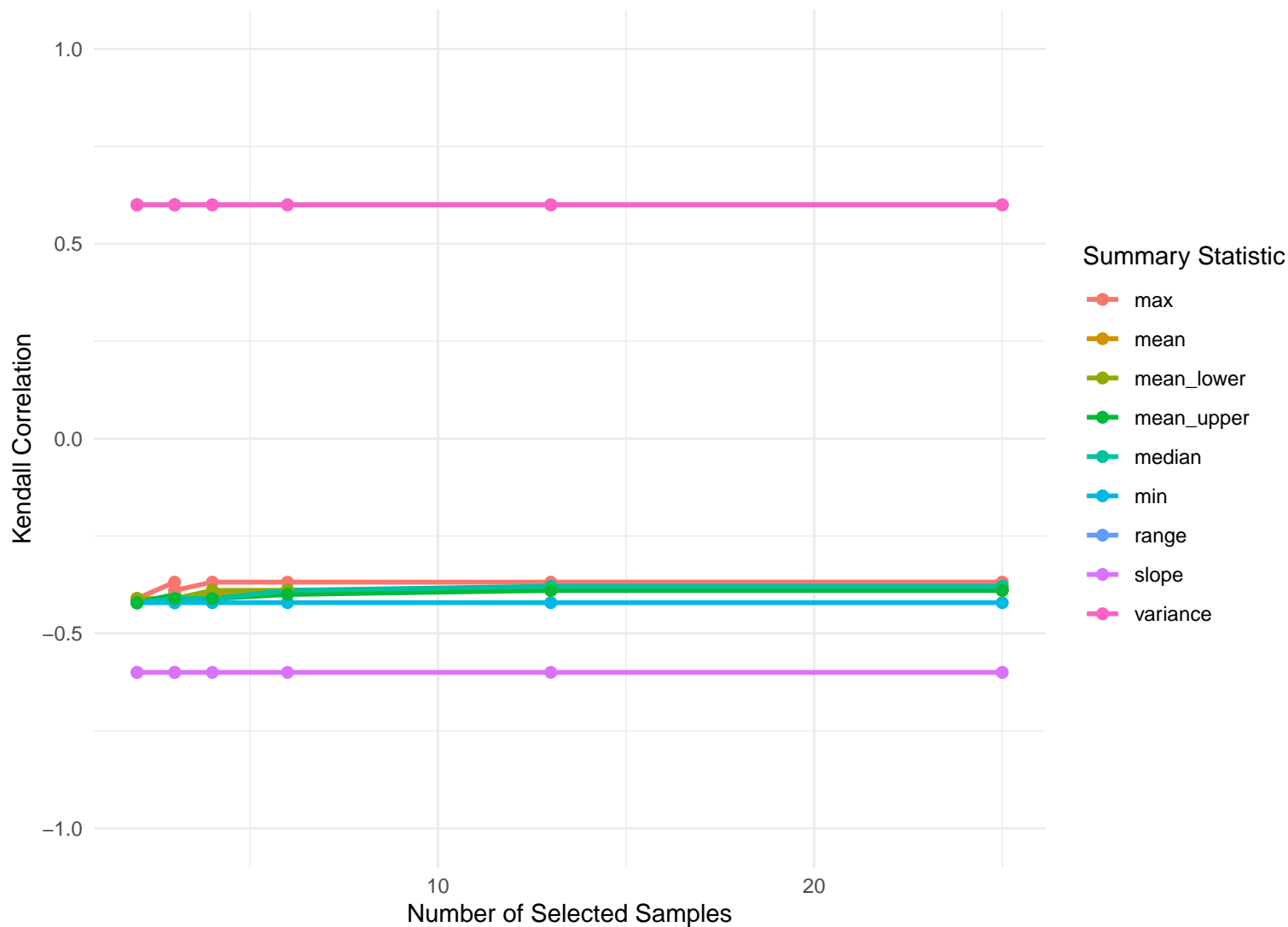
Kendall Correlation

Summary Statistic

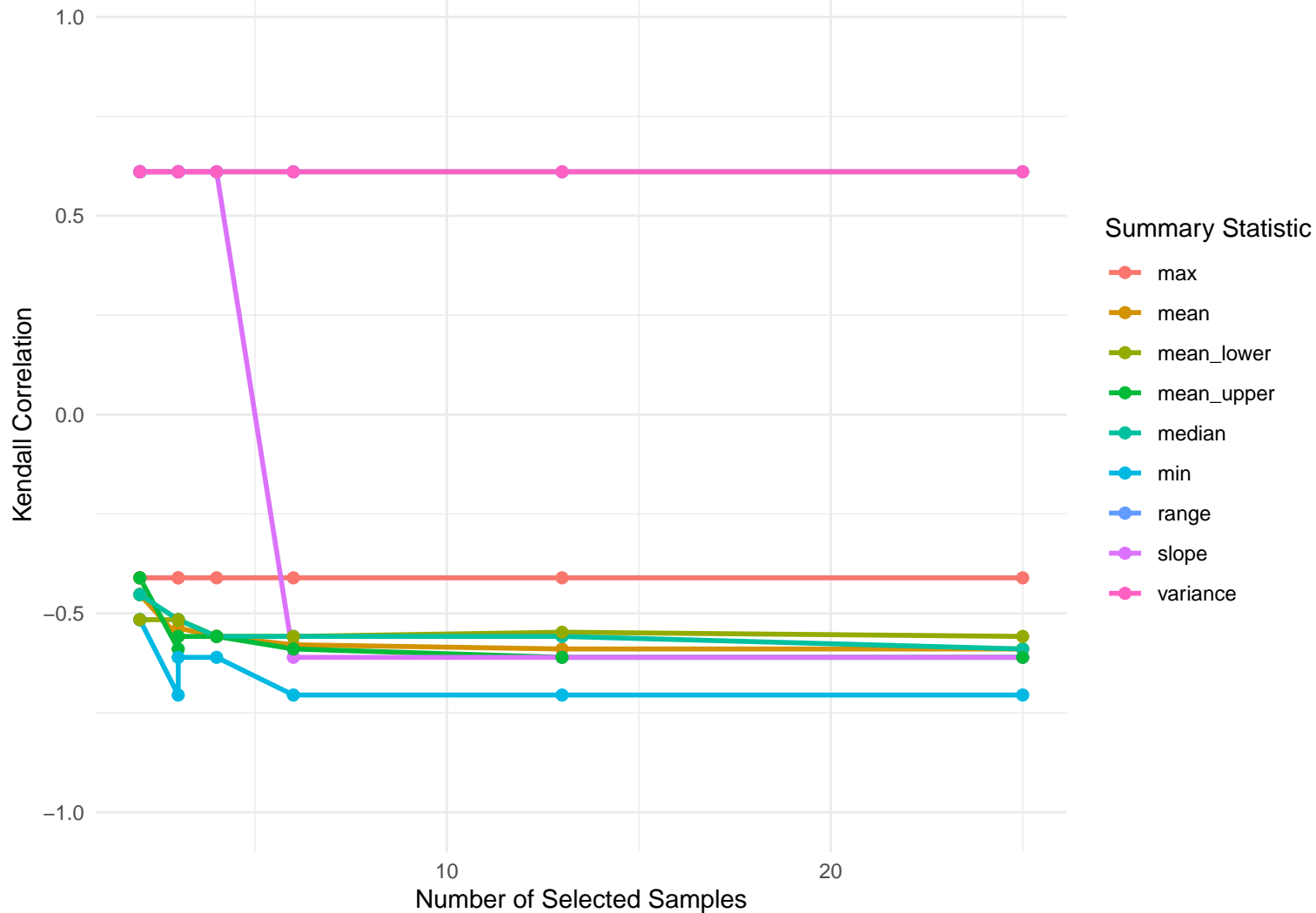
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

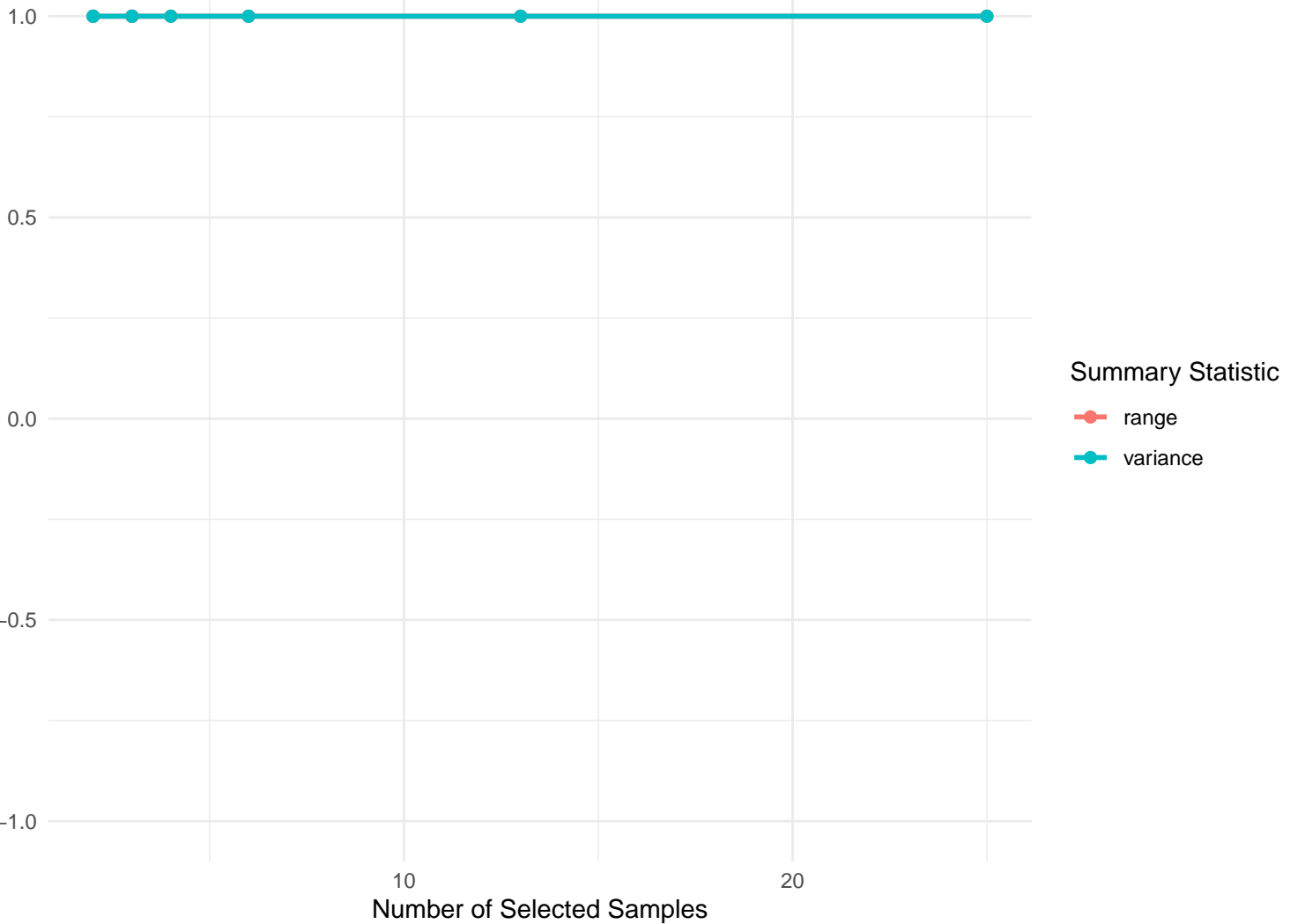
20

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

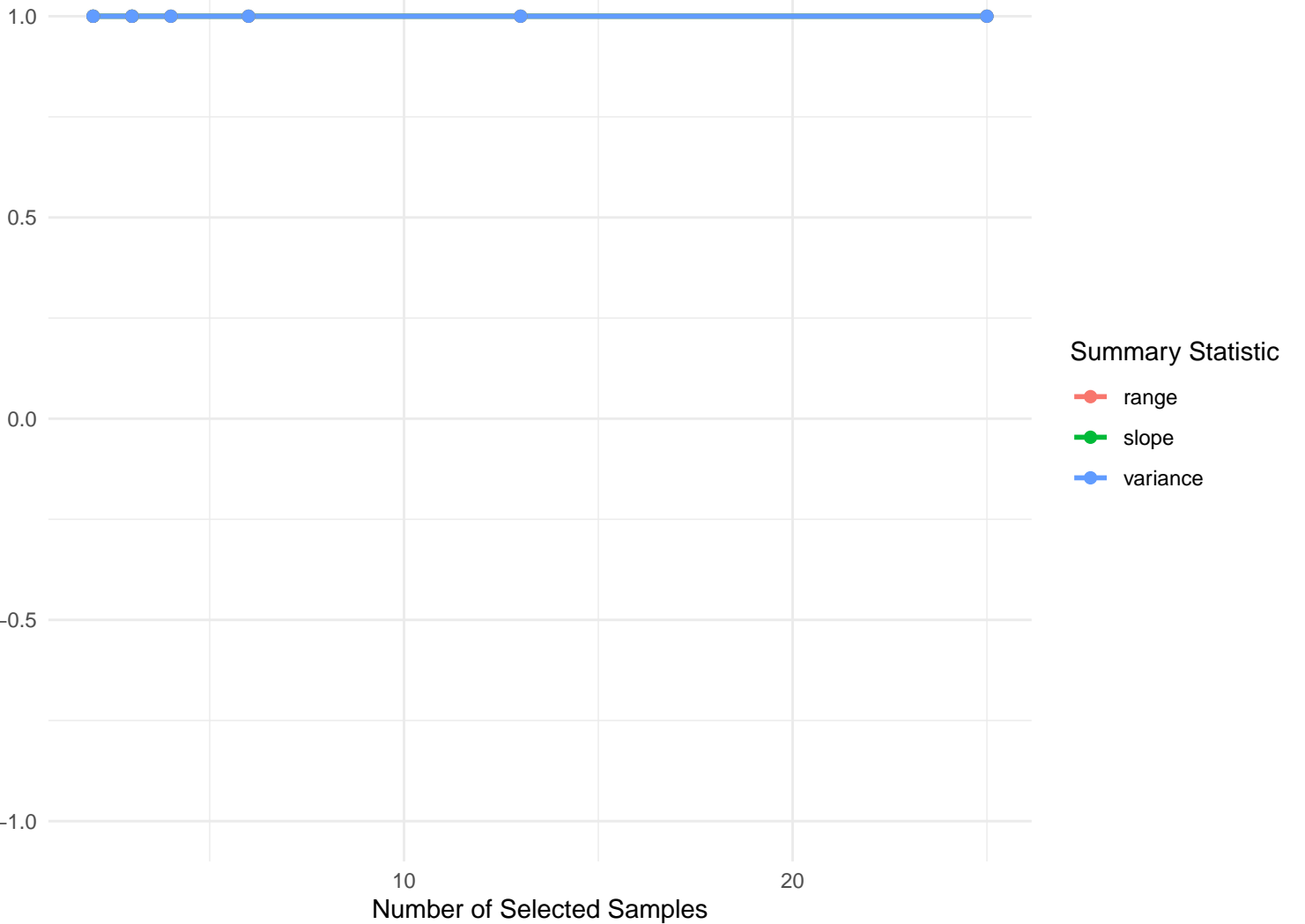
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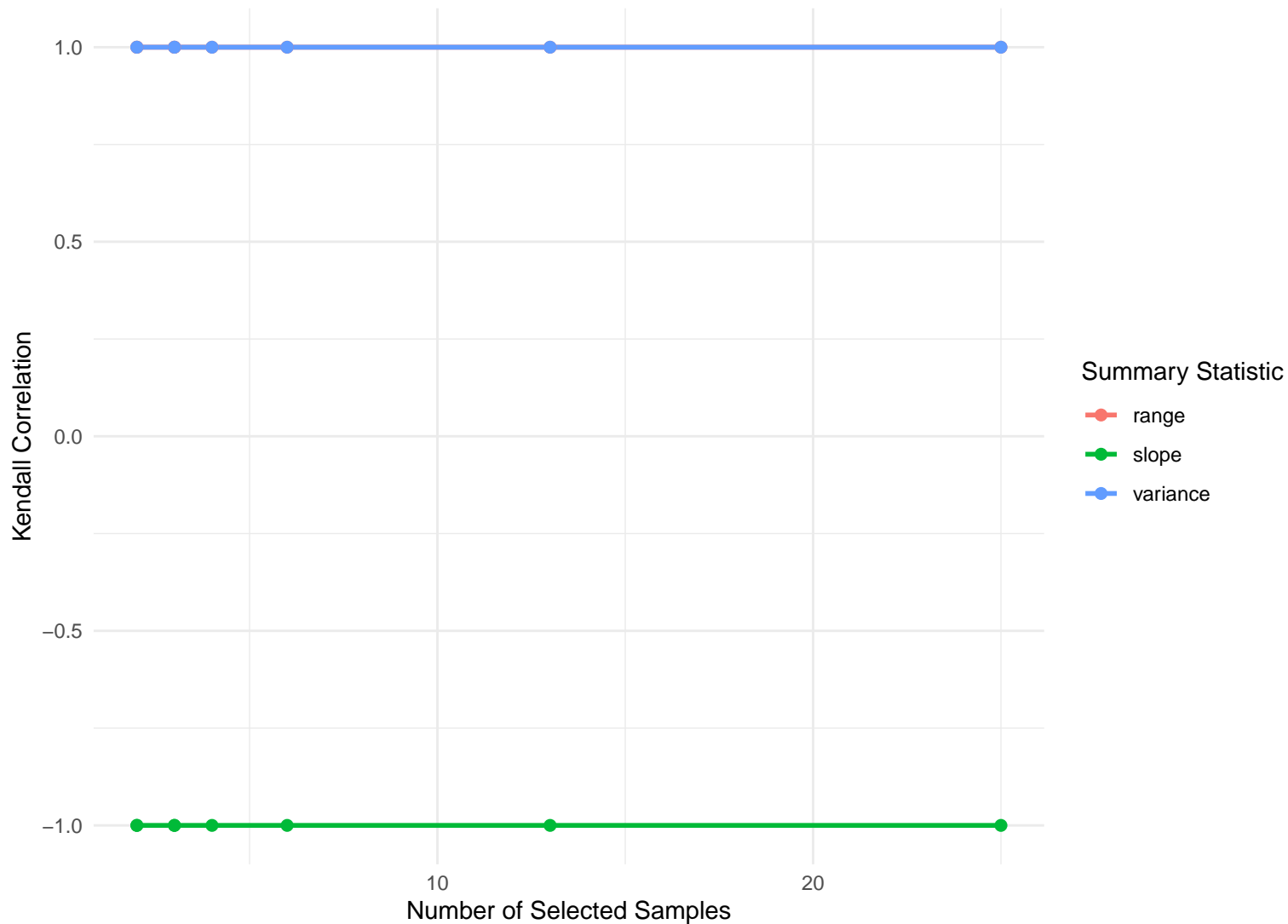
Number of Selected Samples

Summary Statistic

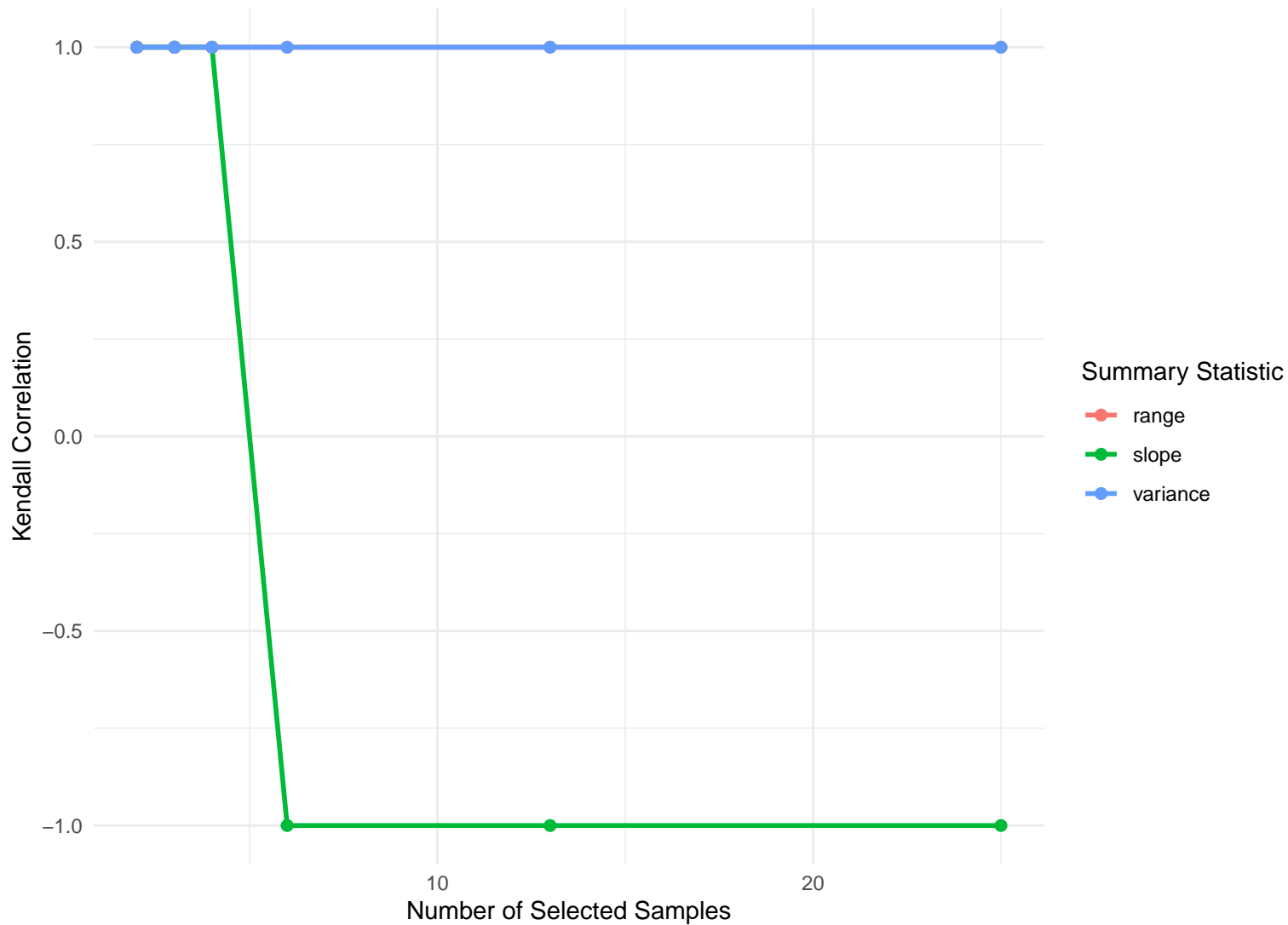
- range
- slope
- variance



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

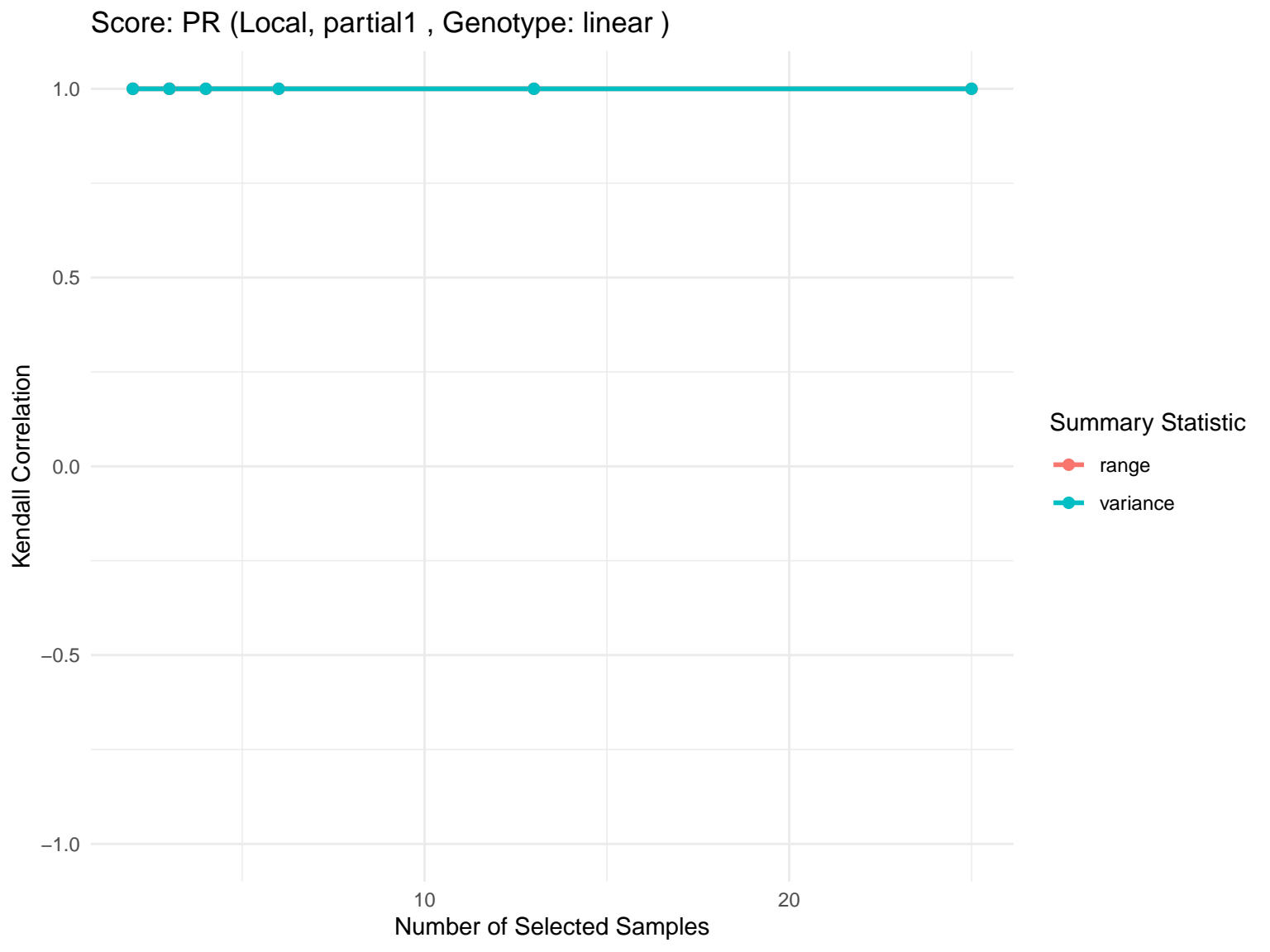
20

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

10

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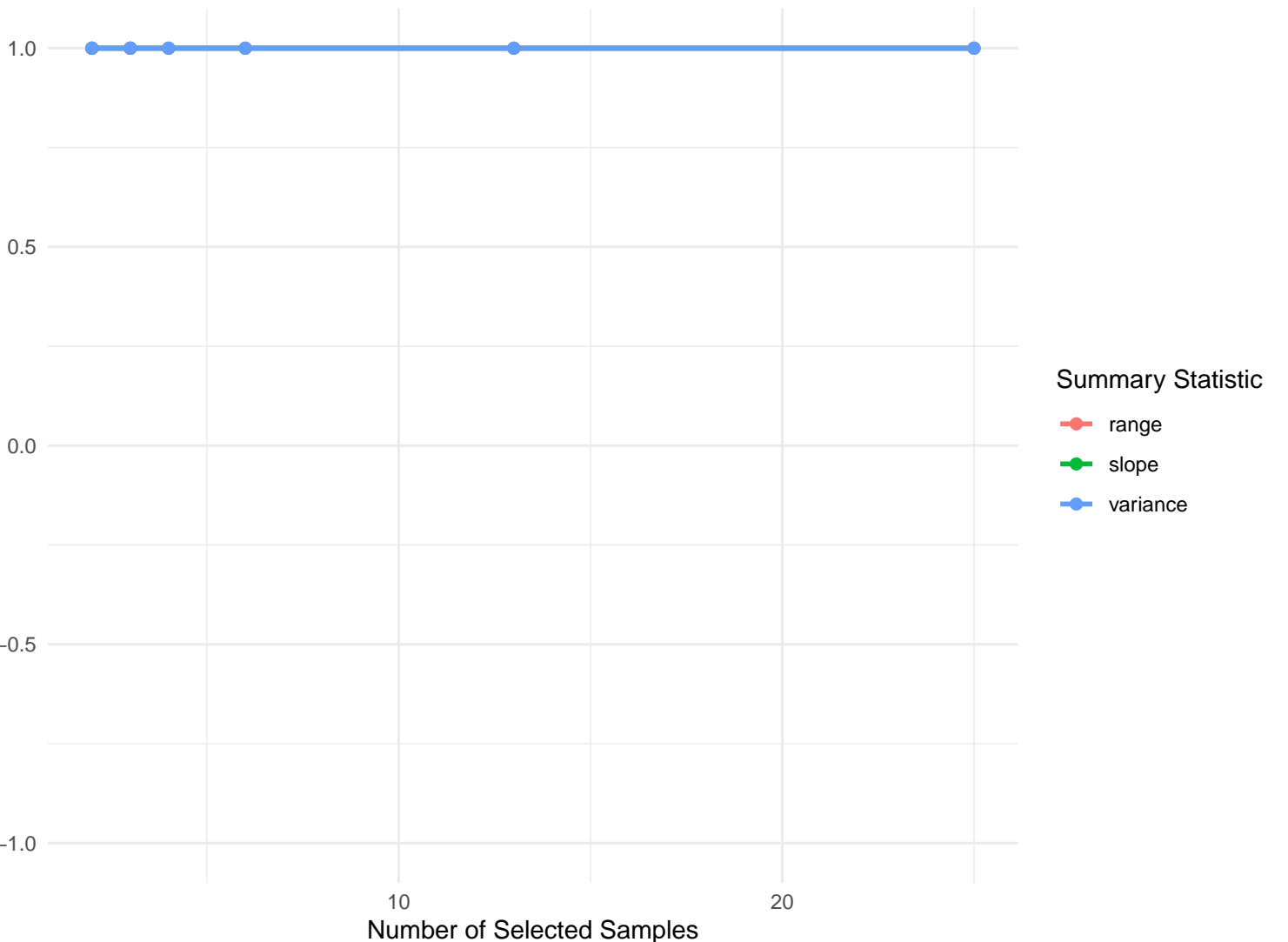
Number of Selected Samples

Summary Statistic

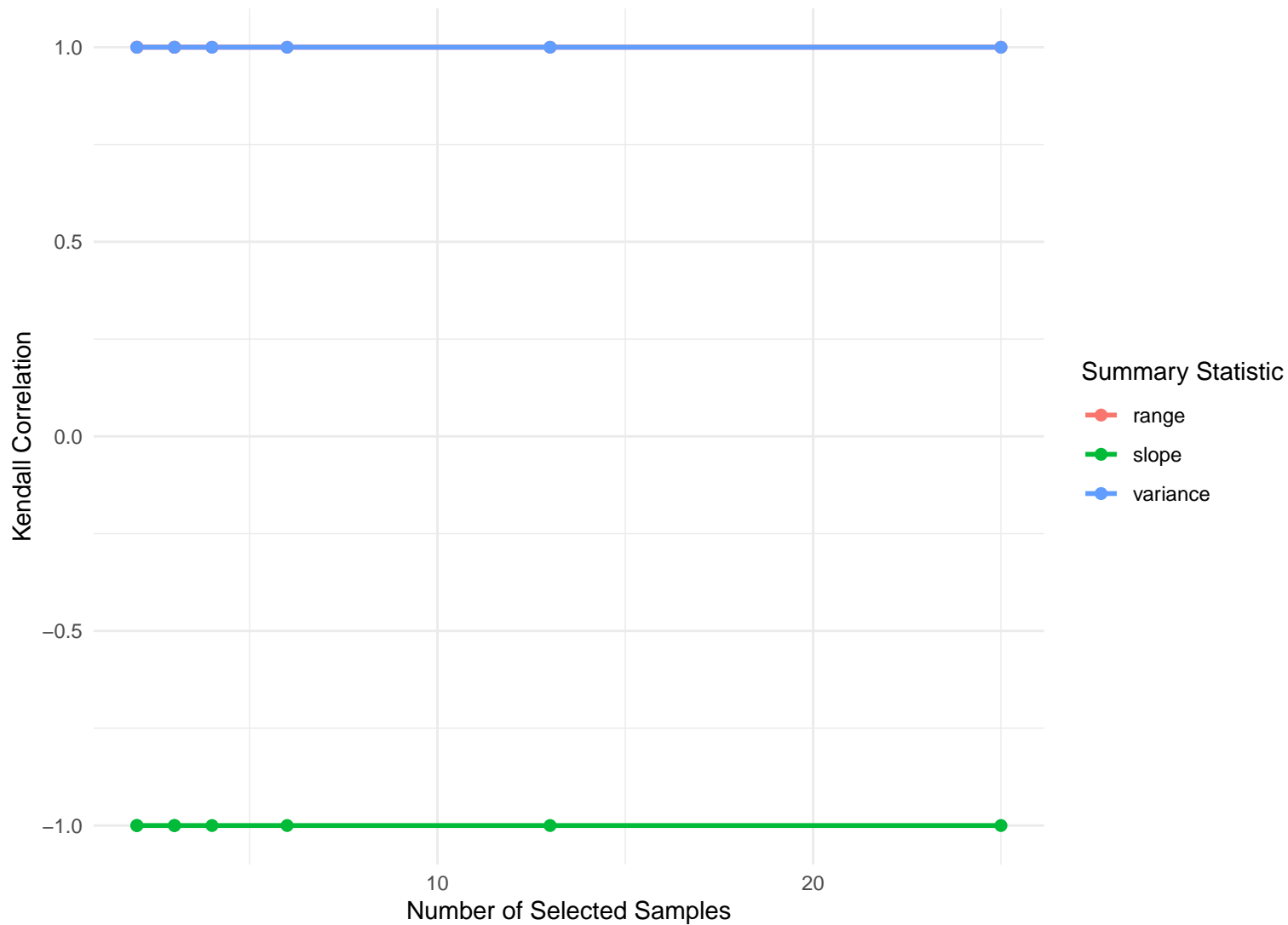
range

slope

variance



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

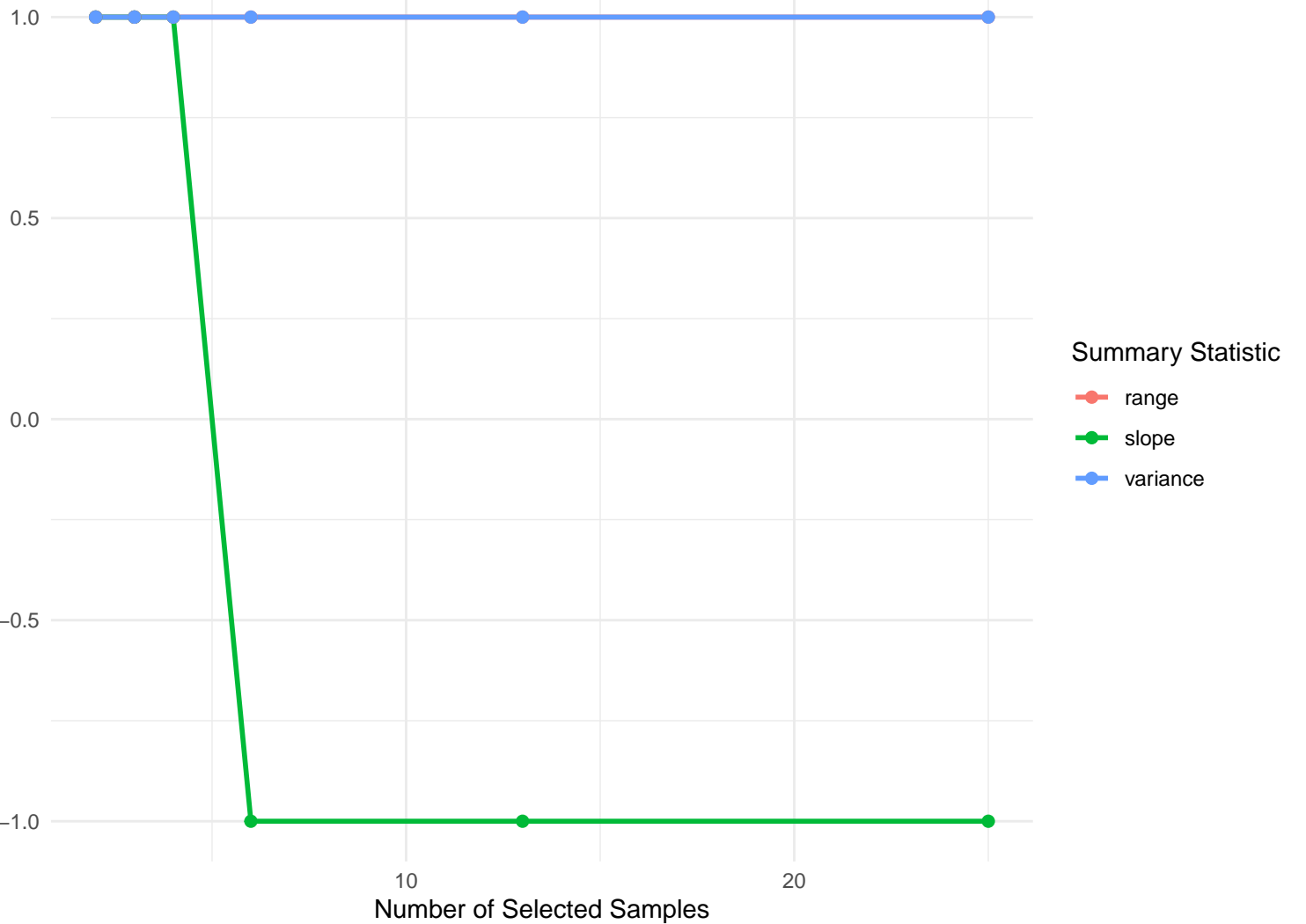
Number of Selected Samples

10

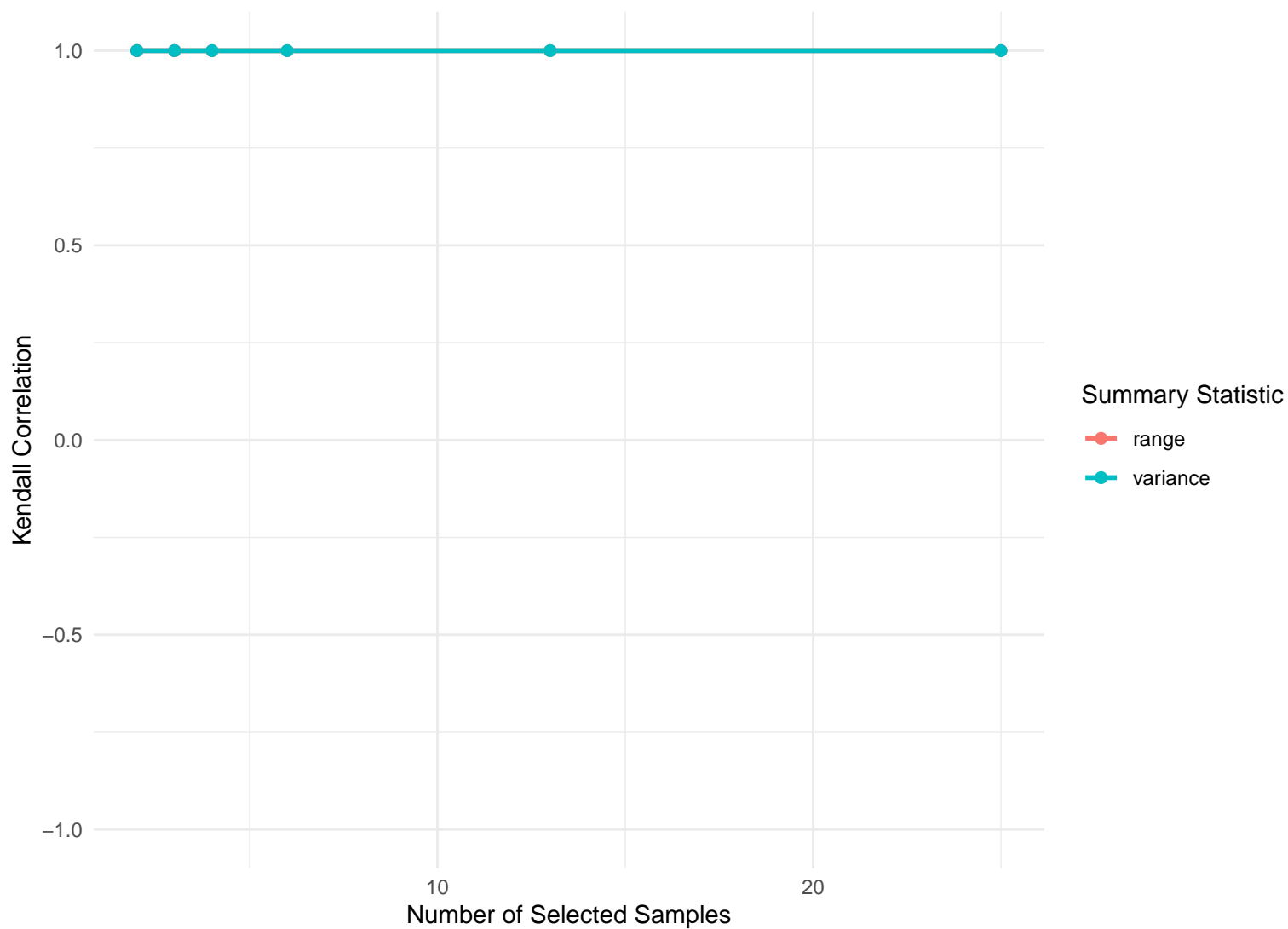
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Summary Statistic

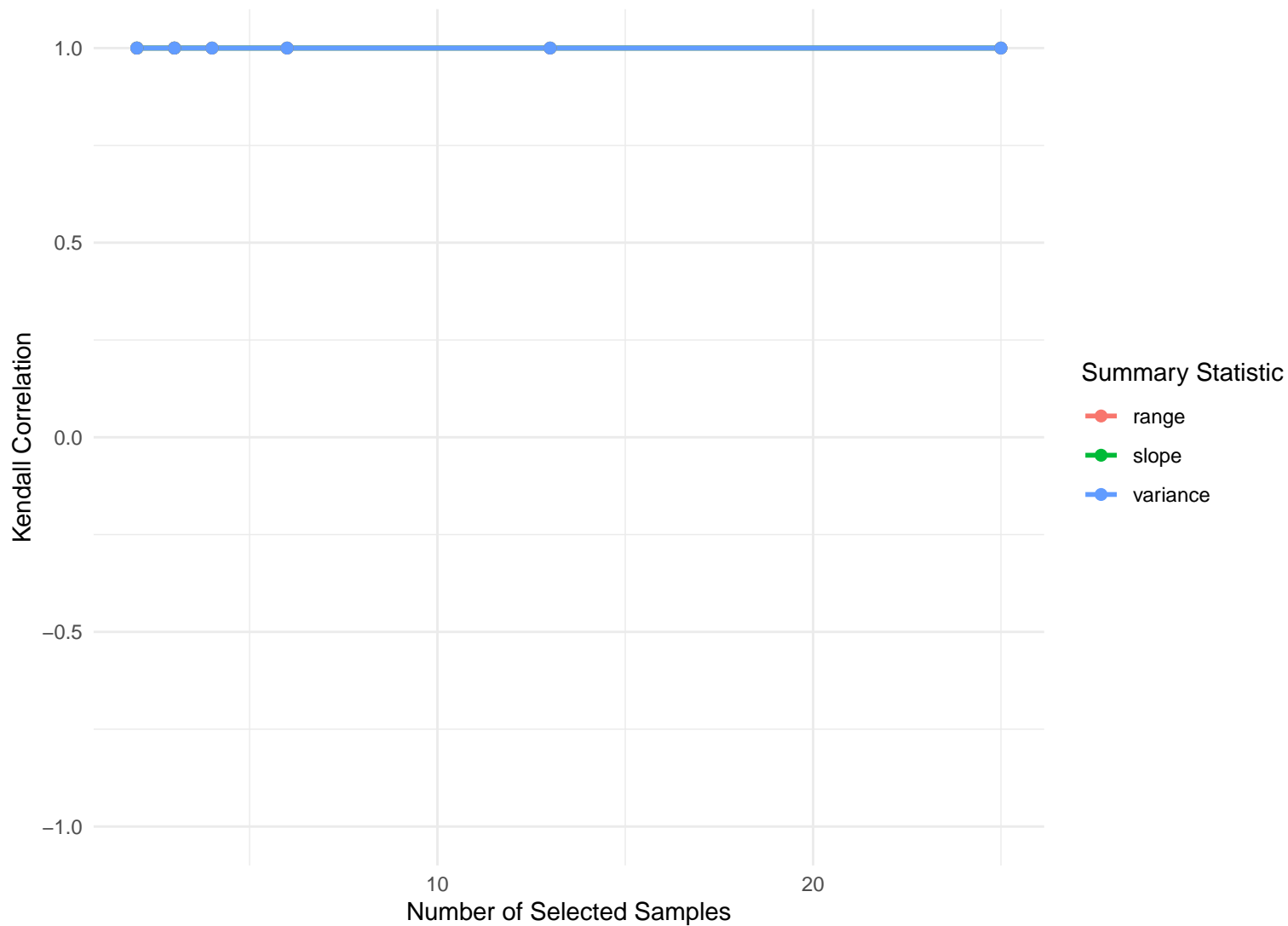
- range
- slope
- variance



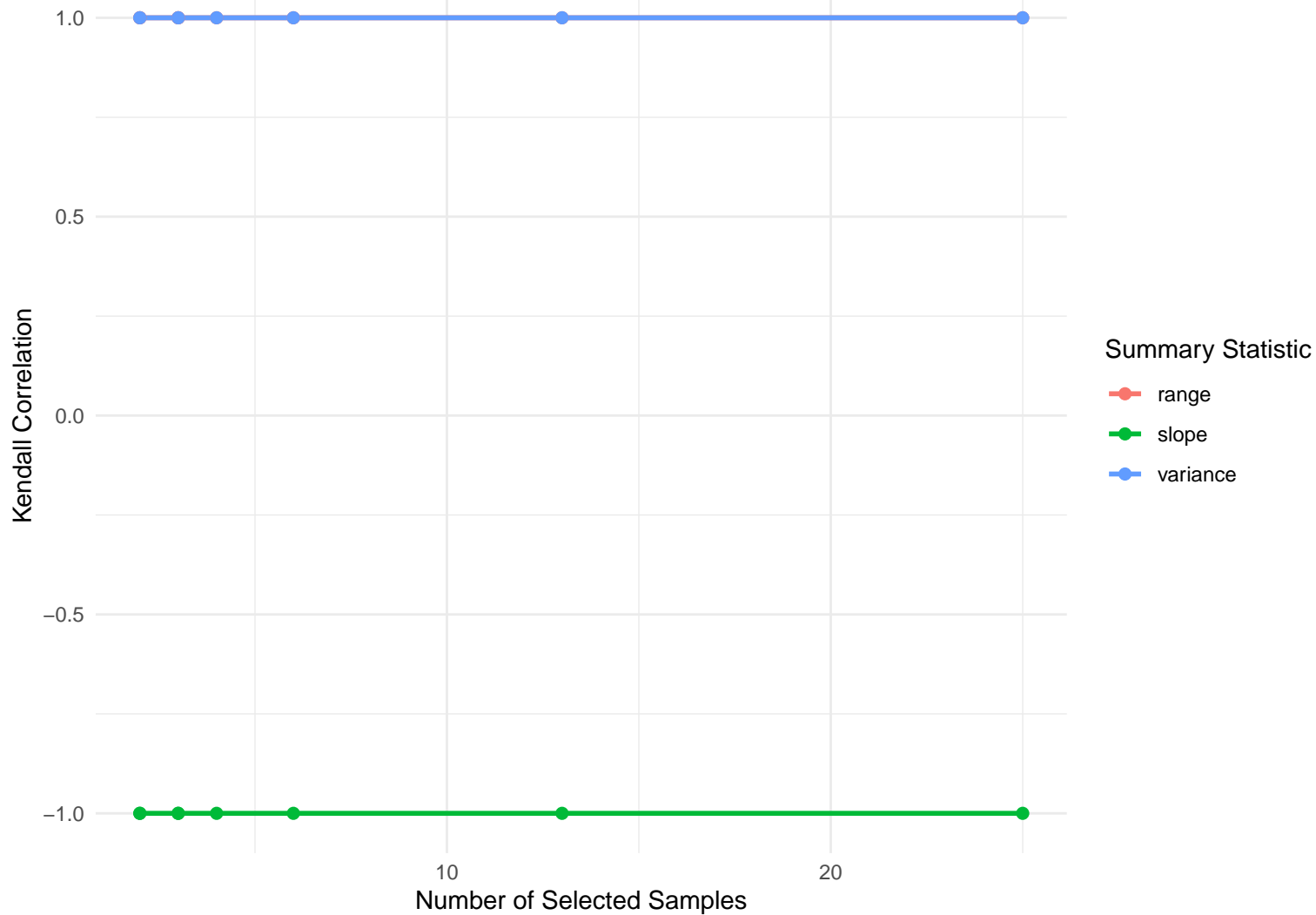
Score: NRW (Local, partial1 , Genotype: linear)



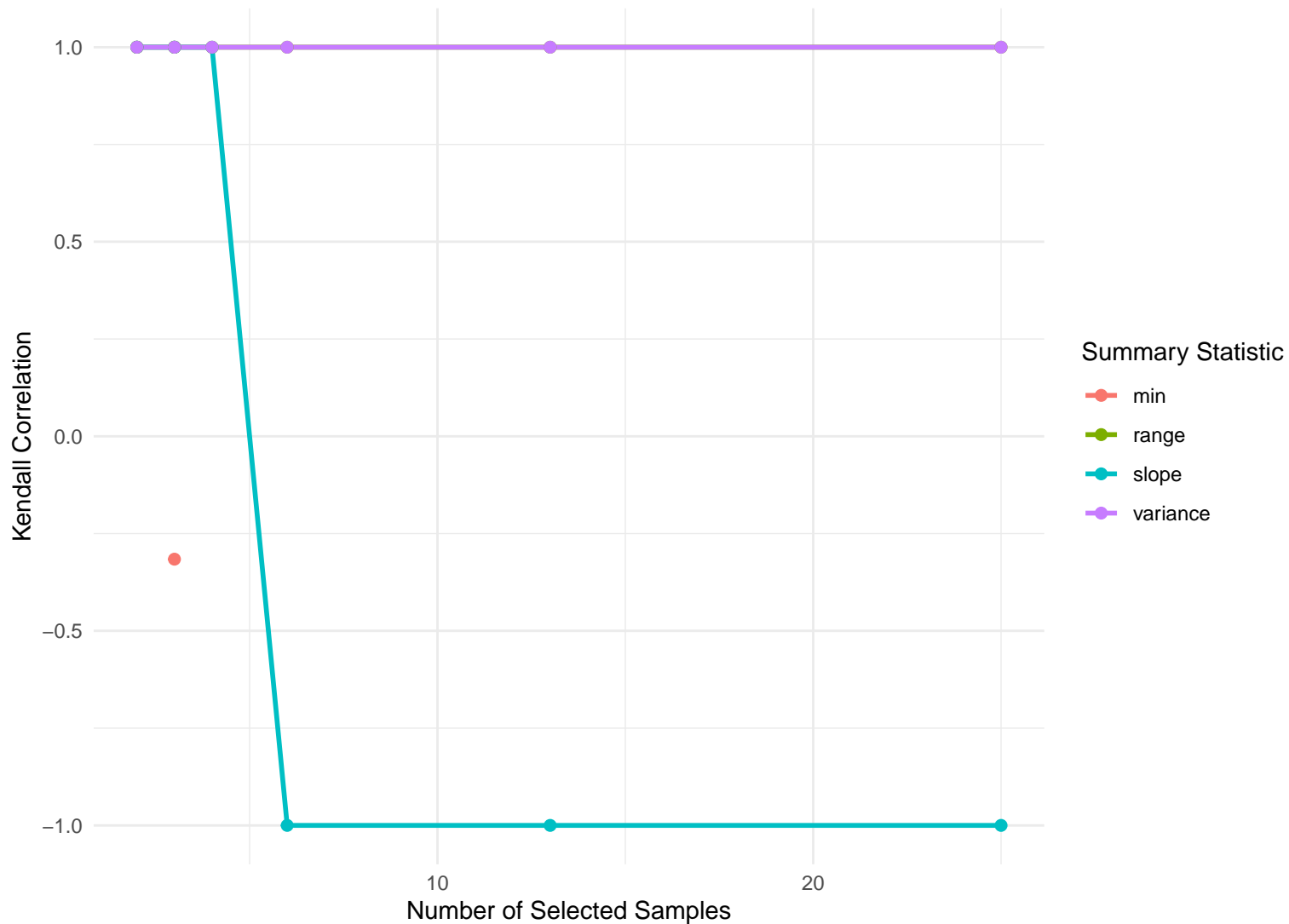
Score: NRW (Local, partial1 , Genotype: gaussian)



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



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



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

Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

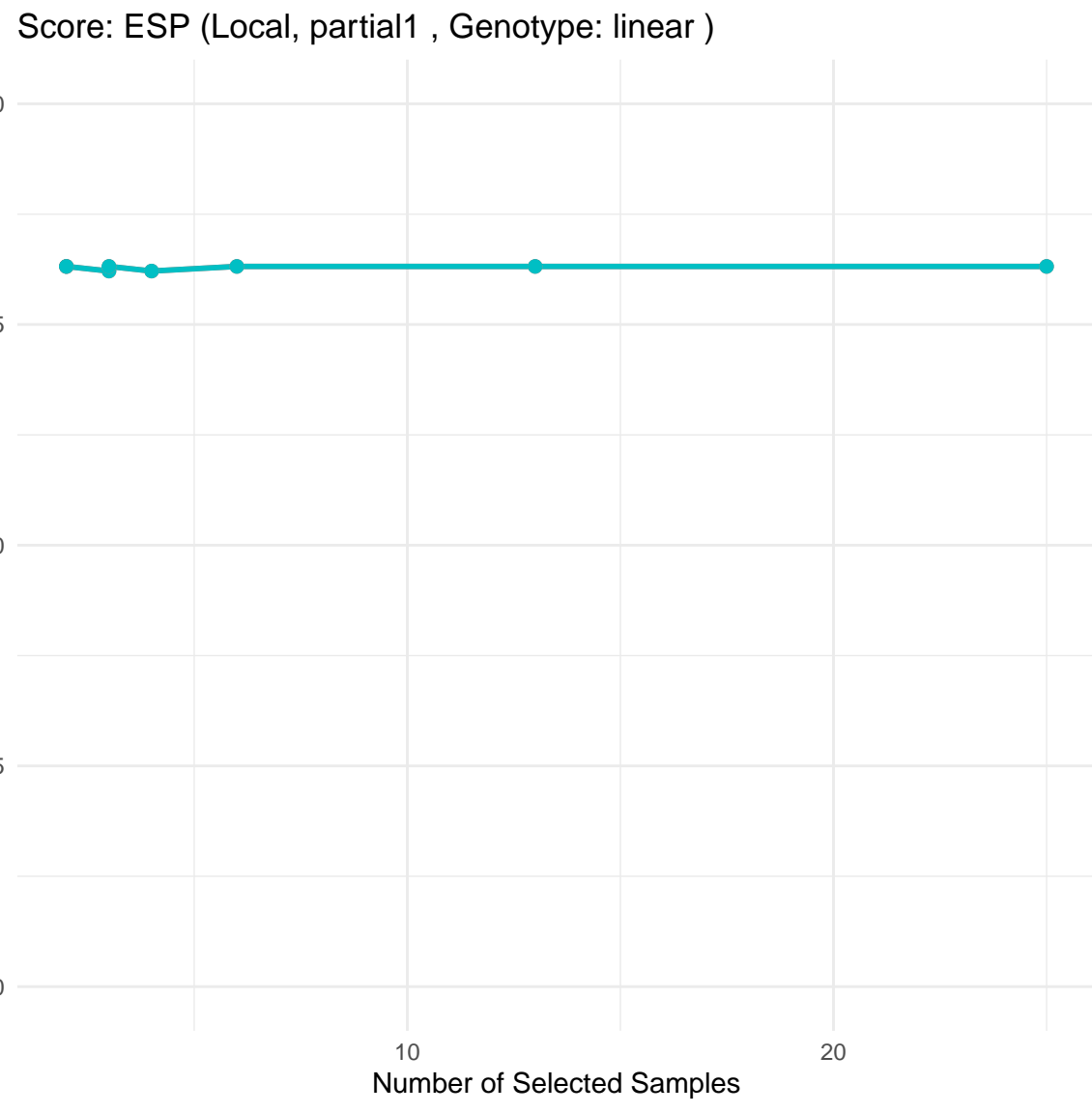
Number of Selected Samples

Summary Statistic

range
variance

10

20

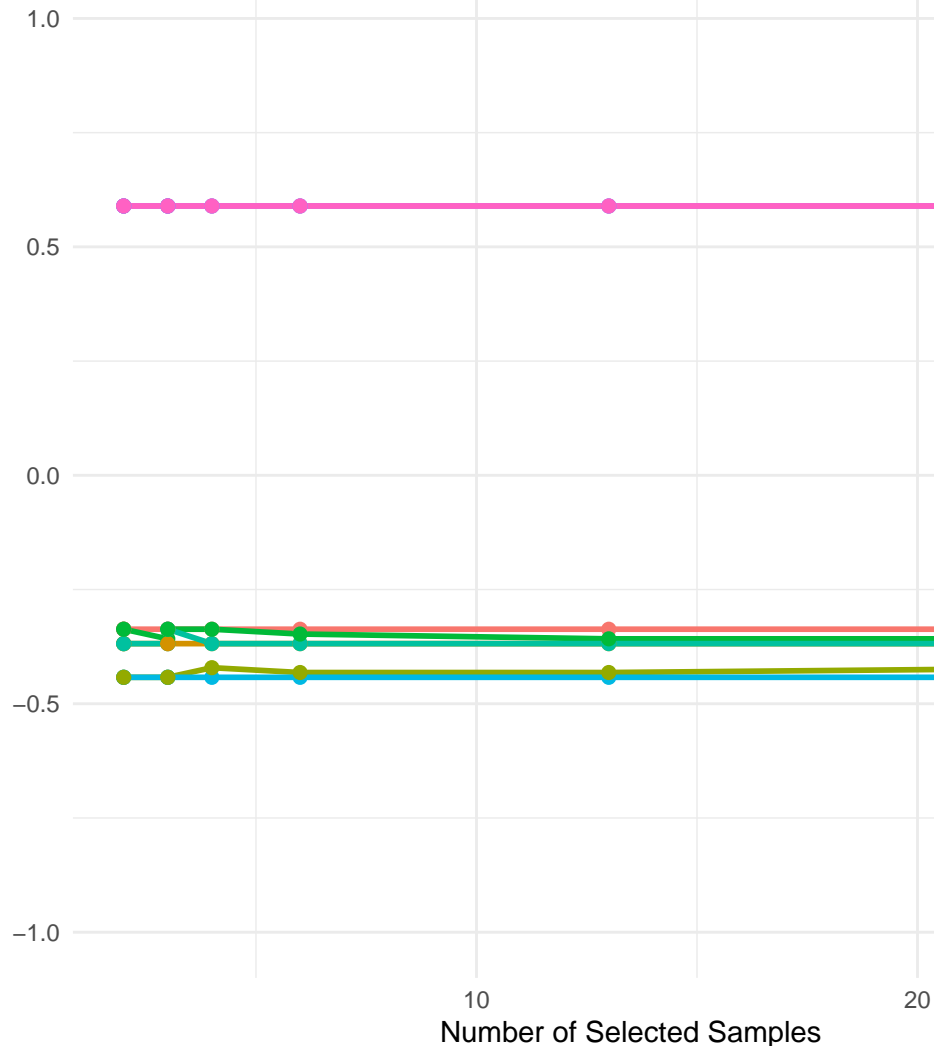


Score: ESP (Local, partial1 , Genotype: gaussian)

Kendall Correlation

Summary Statistic

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



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



Number of Selected Samples

Summary Statistic

max

mean

mean_lower

mean_upper

median

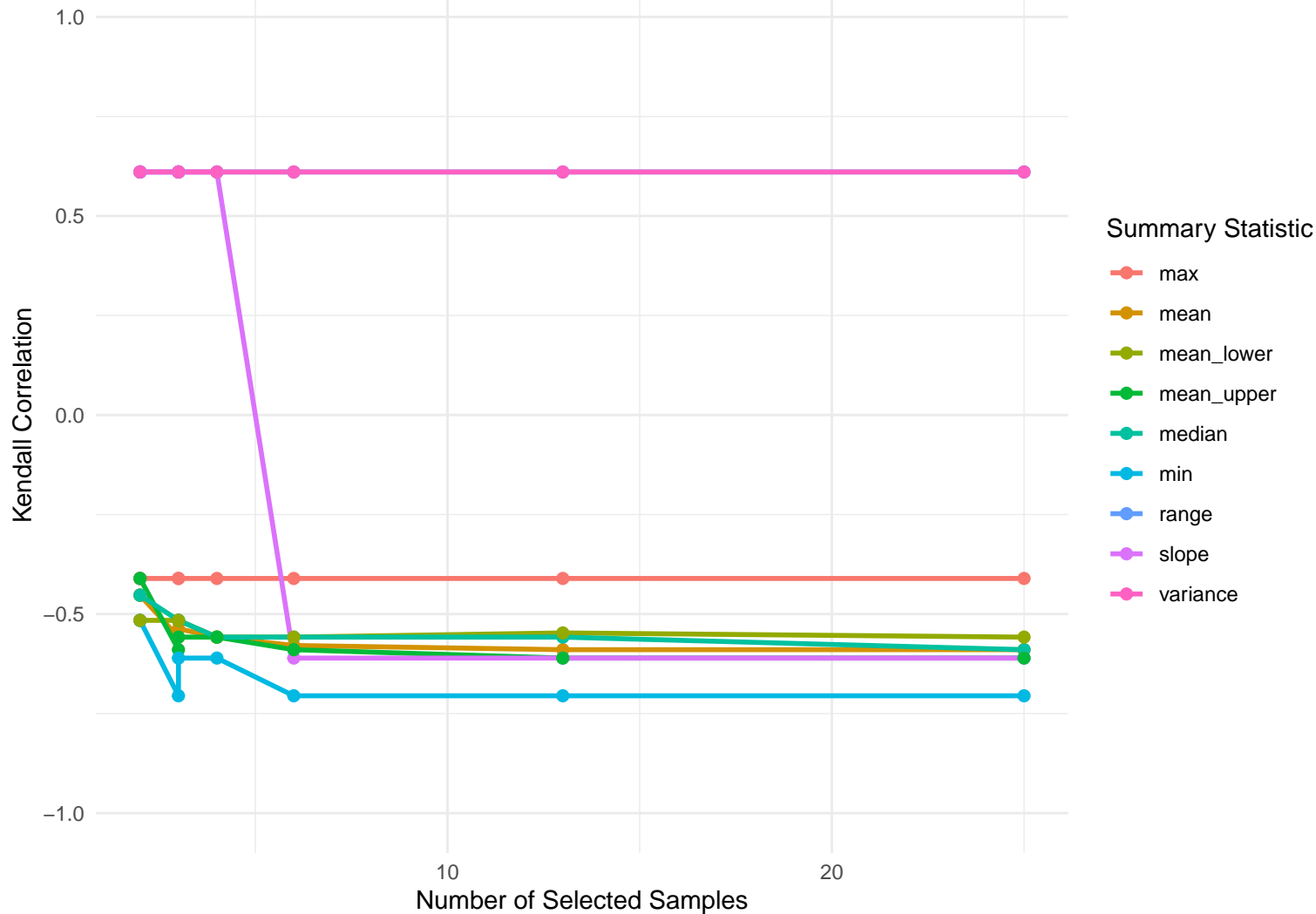
min

range

slope

variance

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



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

Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

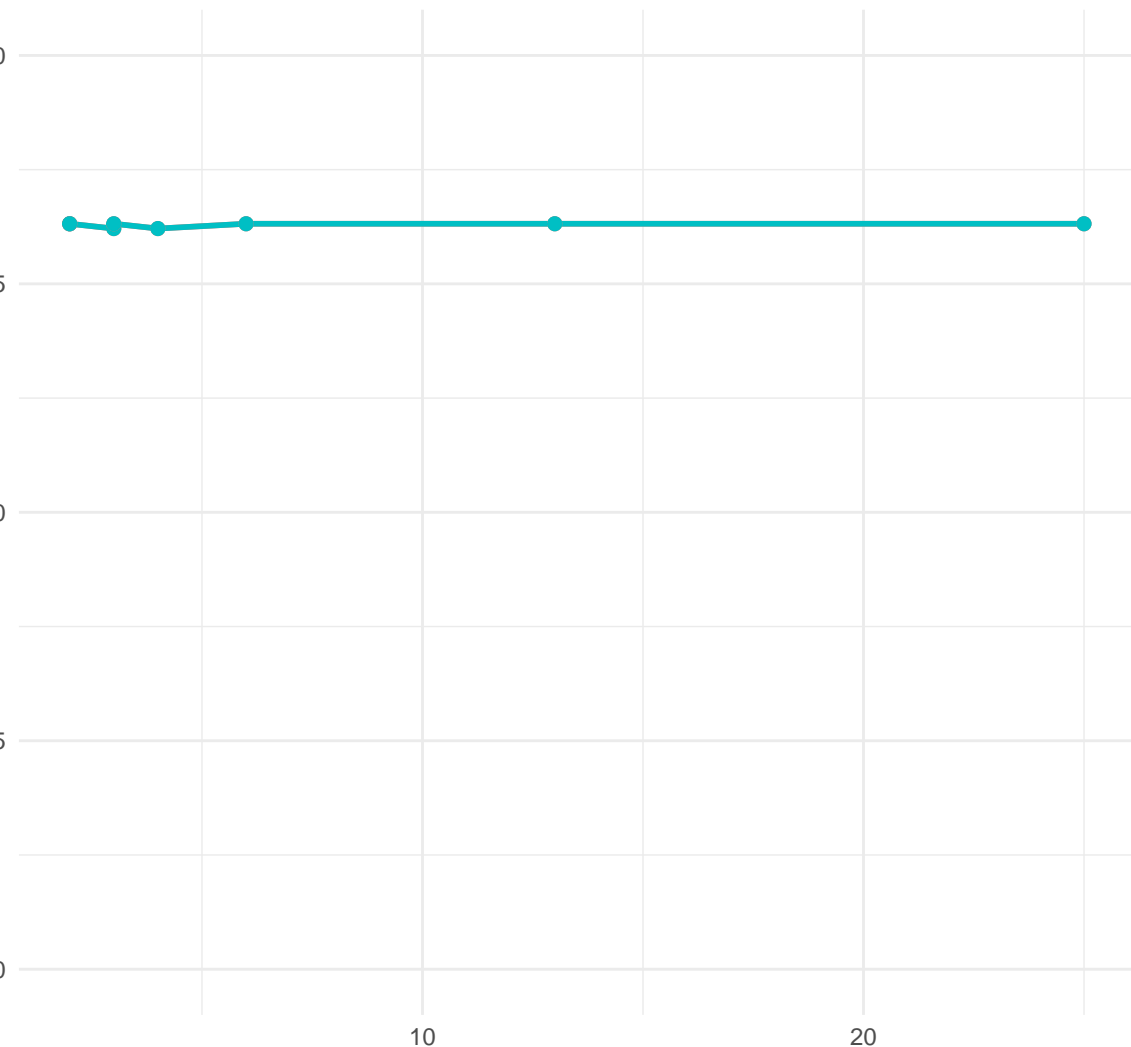
Number of Selected Samples

10

20

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

1.0

0.5

0.0

-0.5

-1.0

10
20
Number of Selected Samples

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

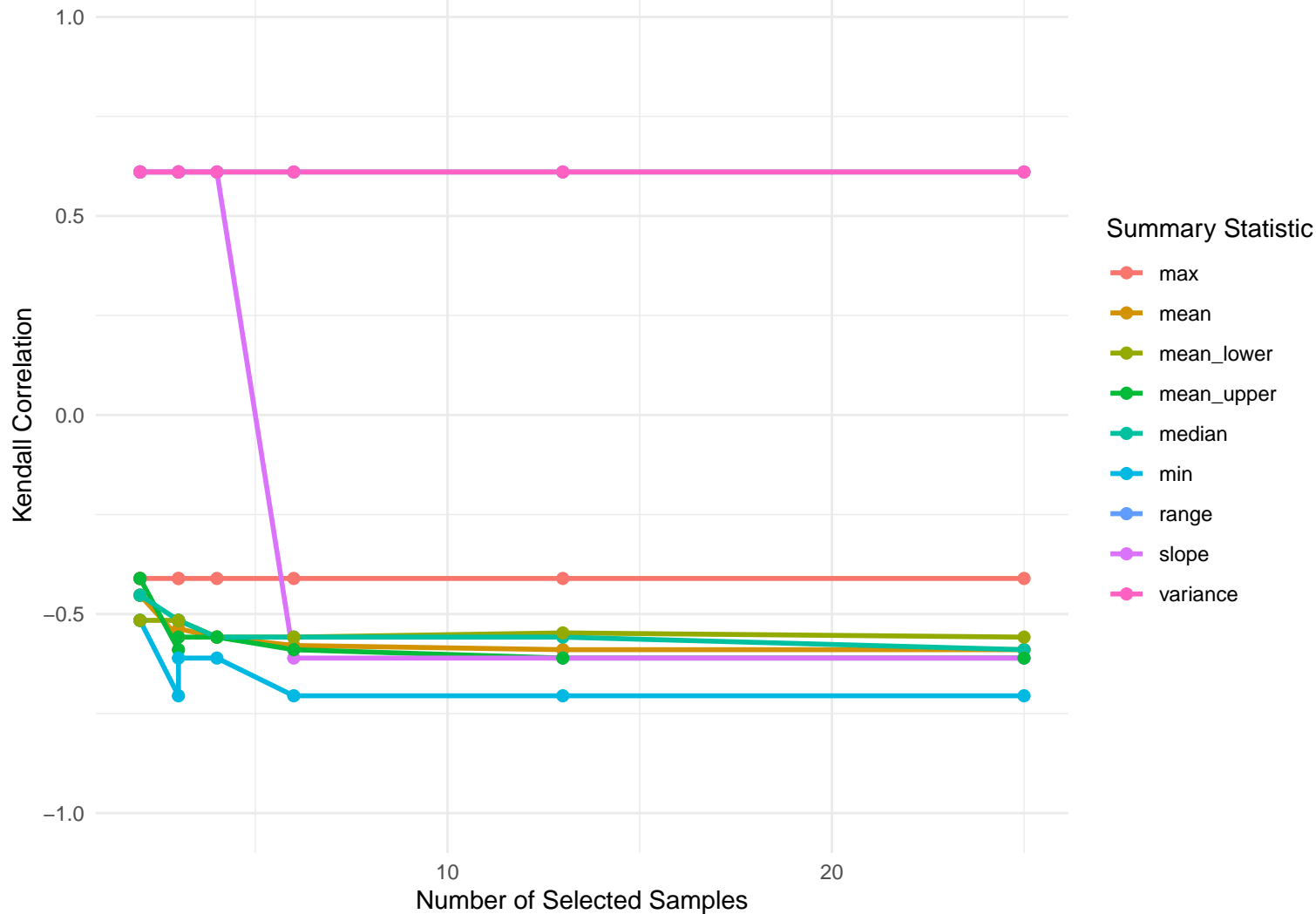


Number of Selected Samples

Summary Statistic

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

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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

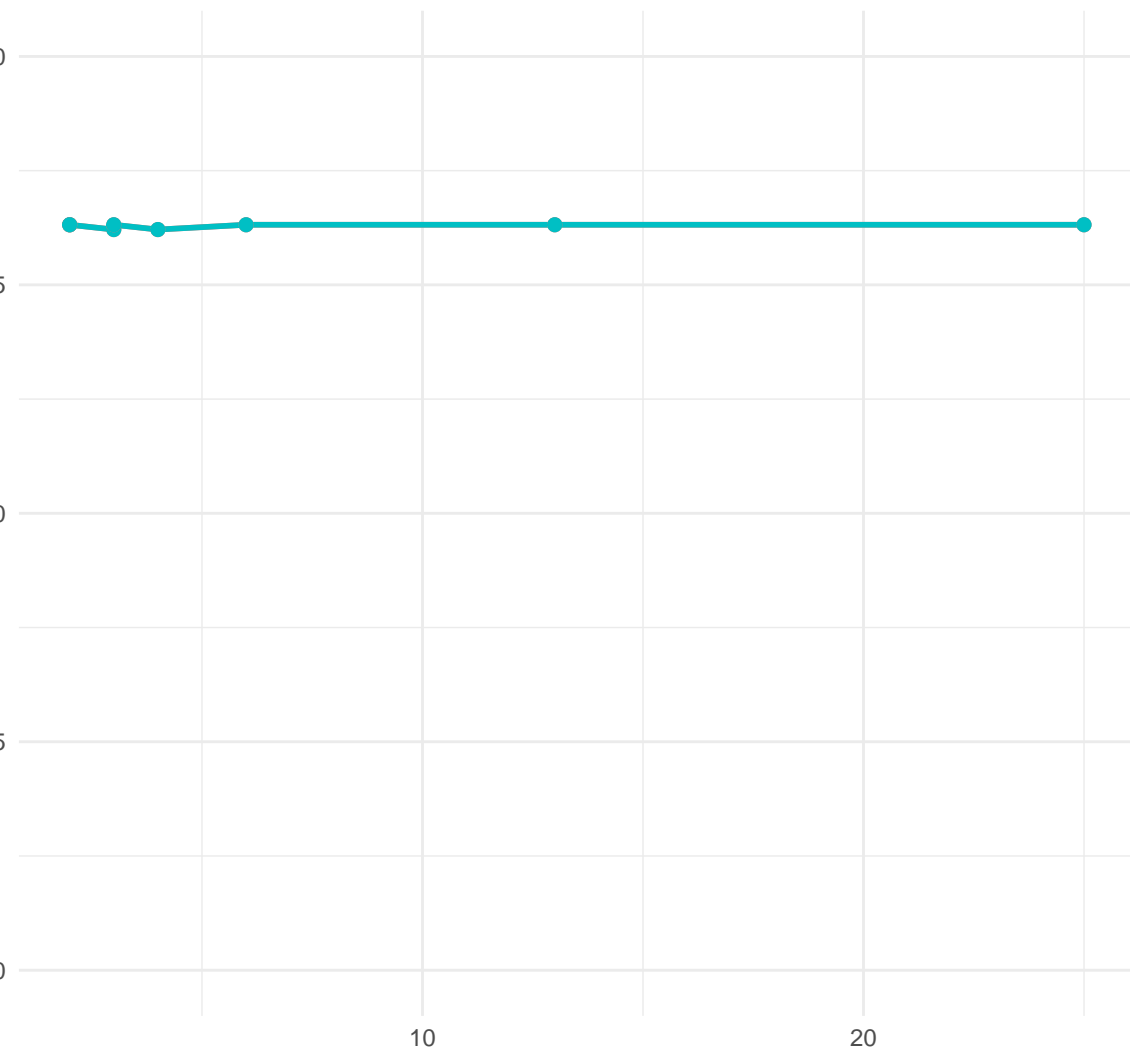
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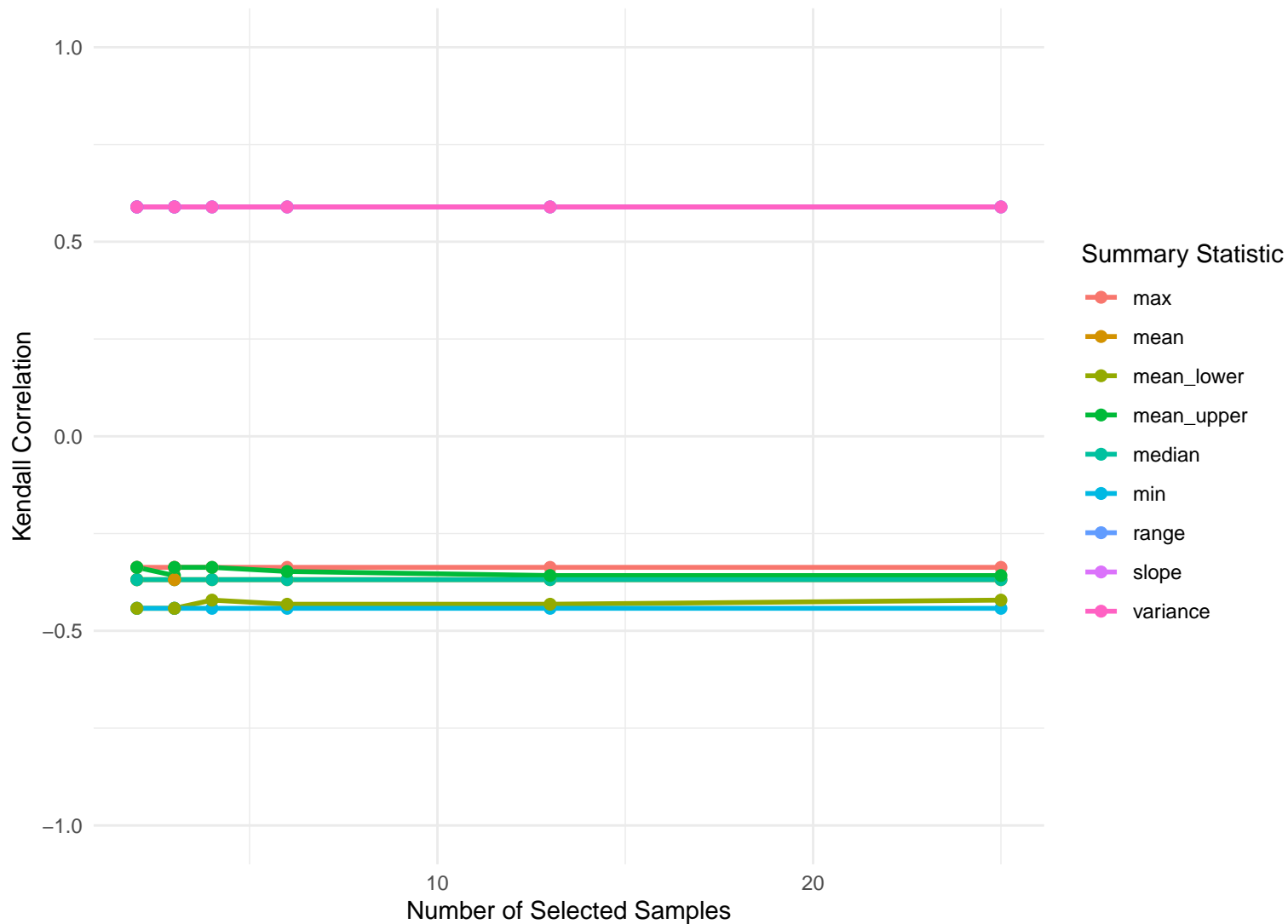
Summary Statistic

range

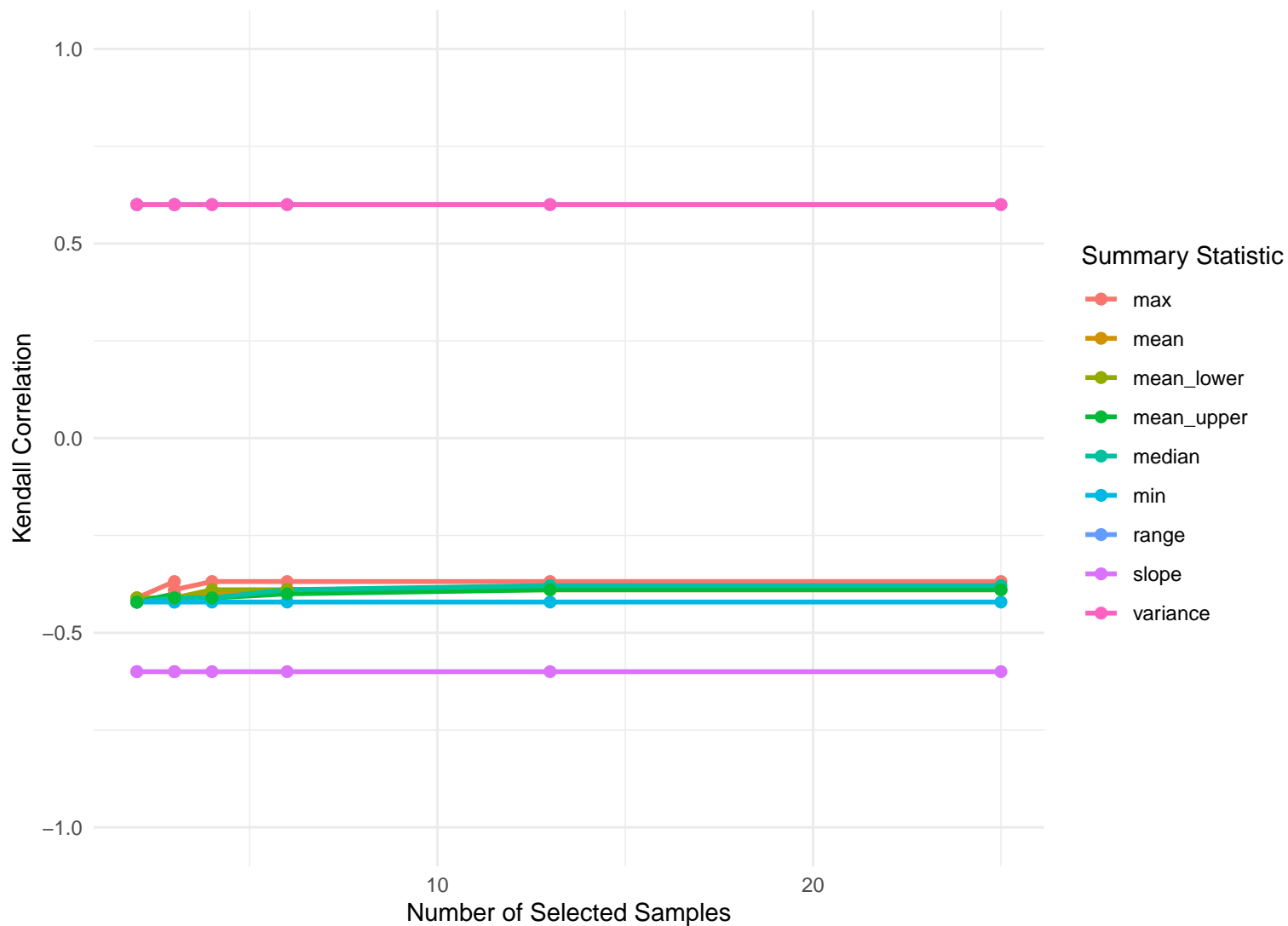
variance



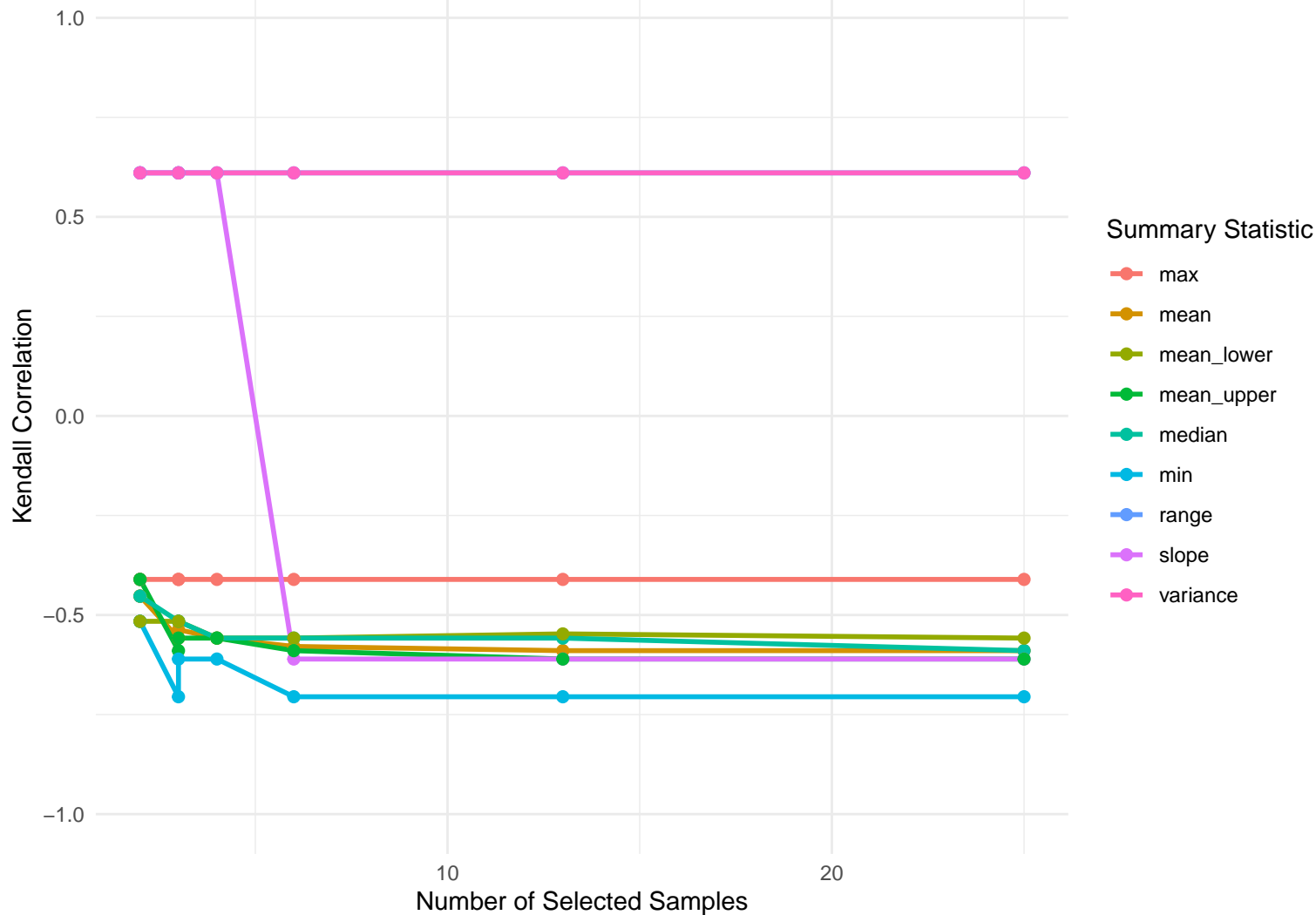
Score: PFI (Local, partial1 , Genotype: gaussian)



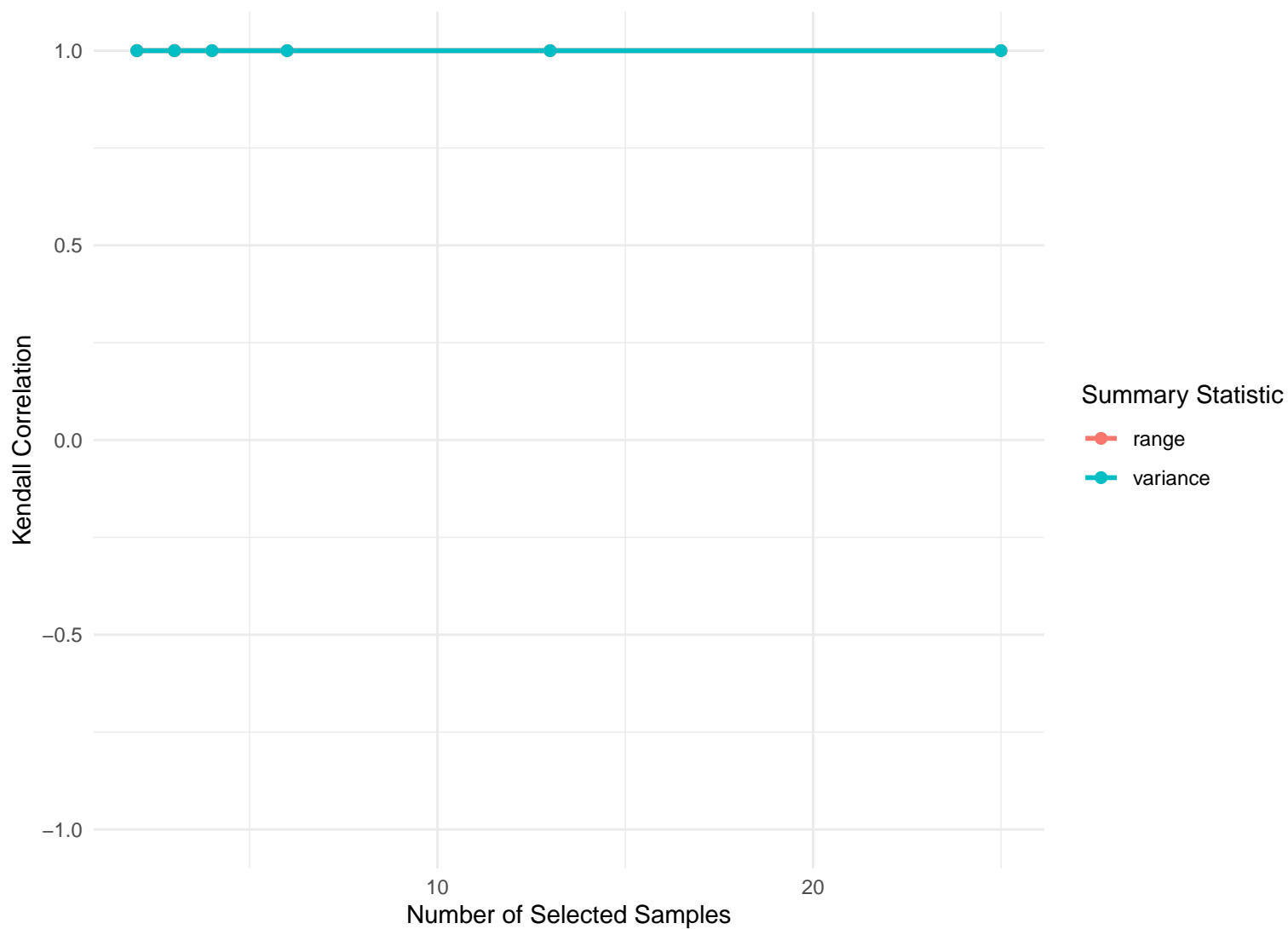
Score: PFI (Local, partial1 , Genotype: sinusoidal)



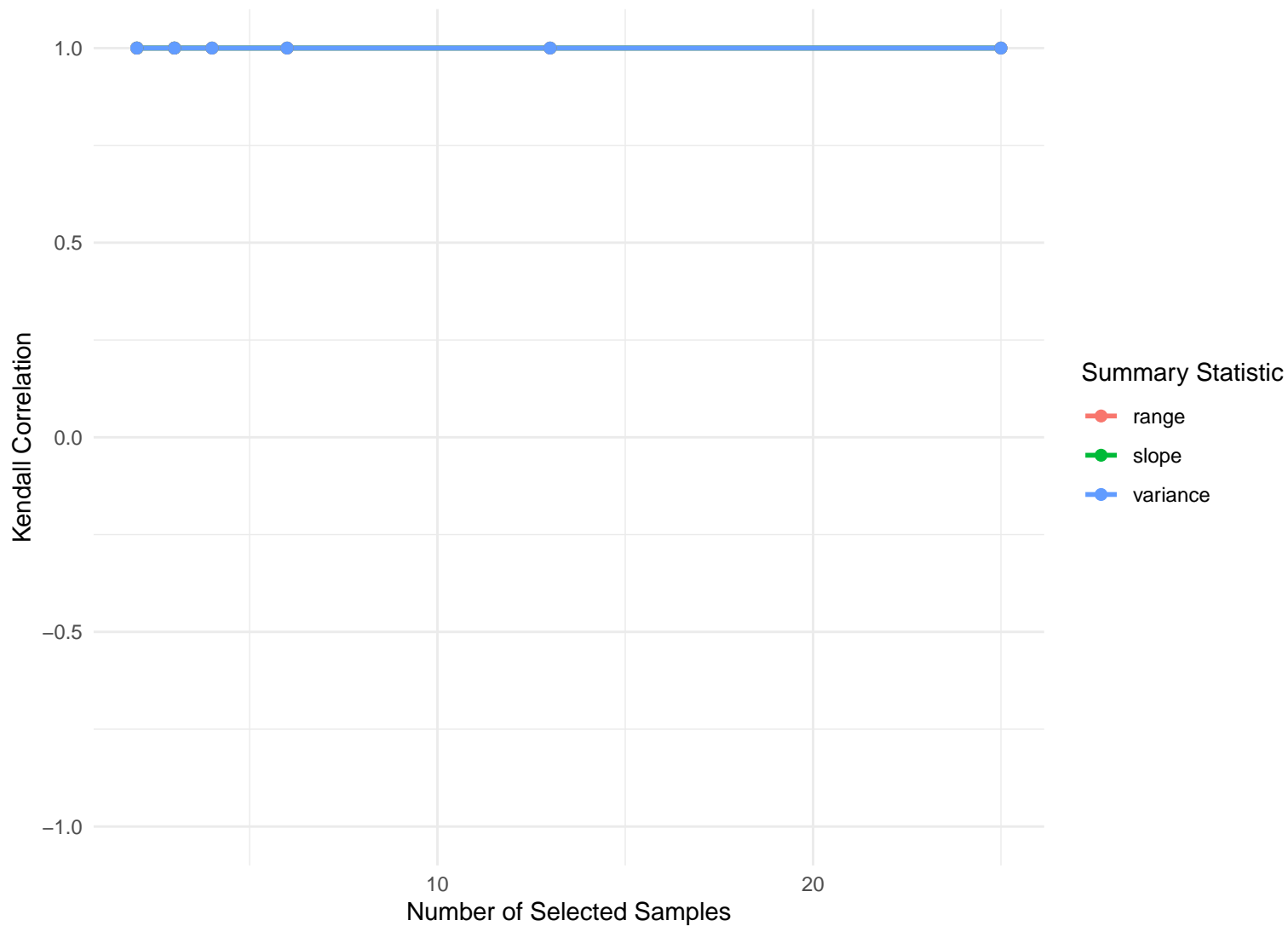
Score: PFI (Local, partial1 , Genotype: wave)



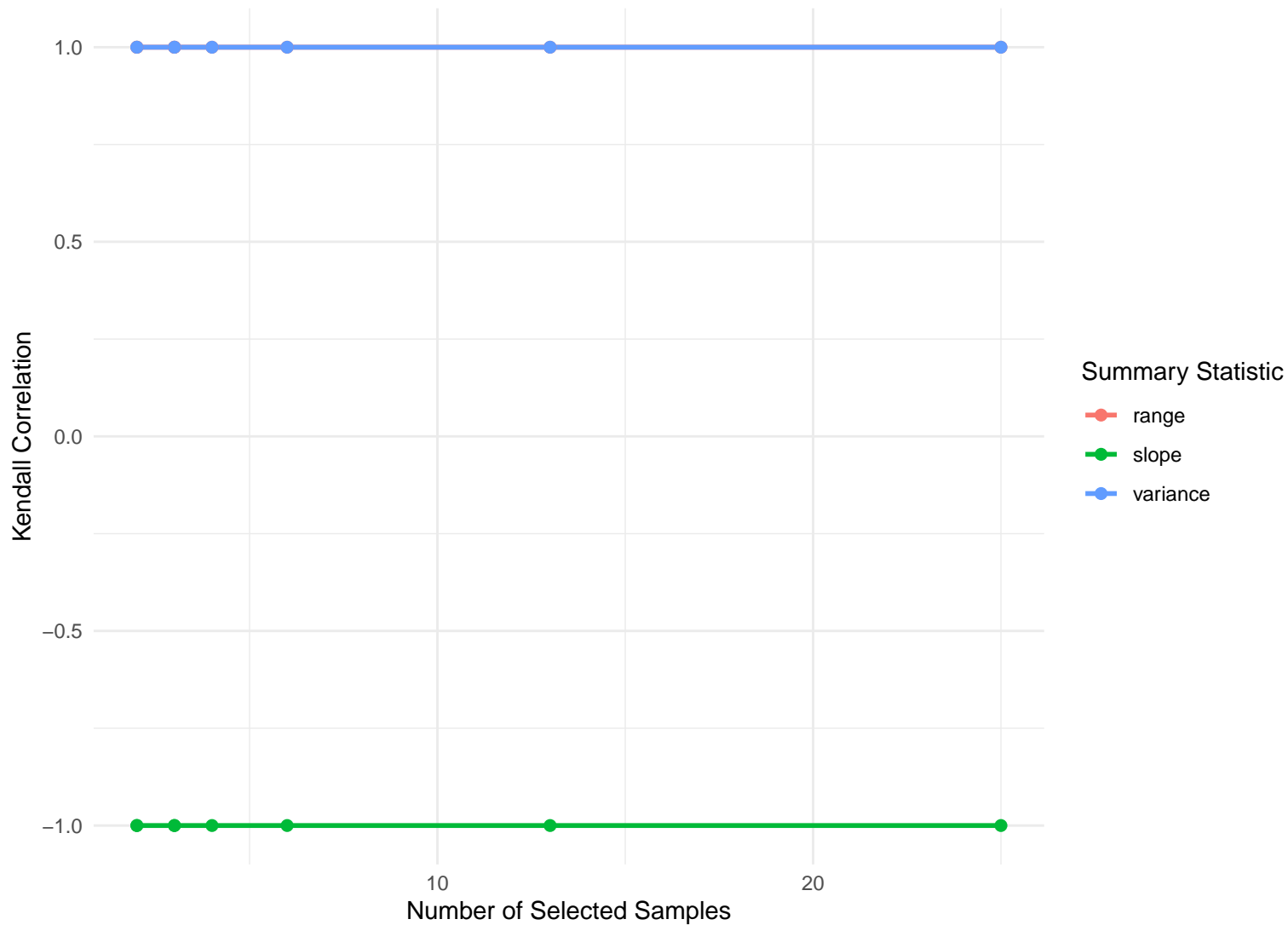
Score: APC (Local, partial1 , Genotype: linear)



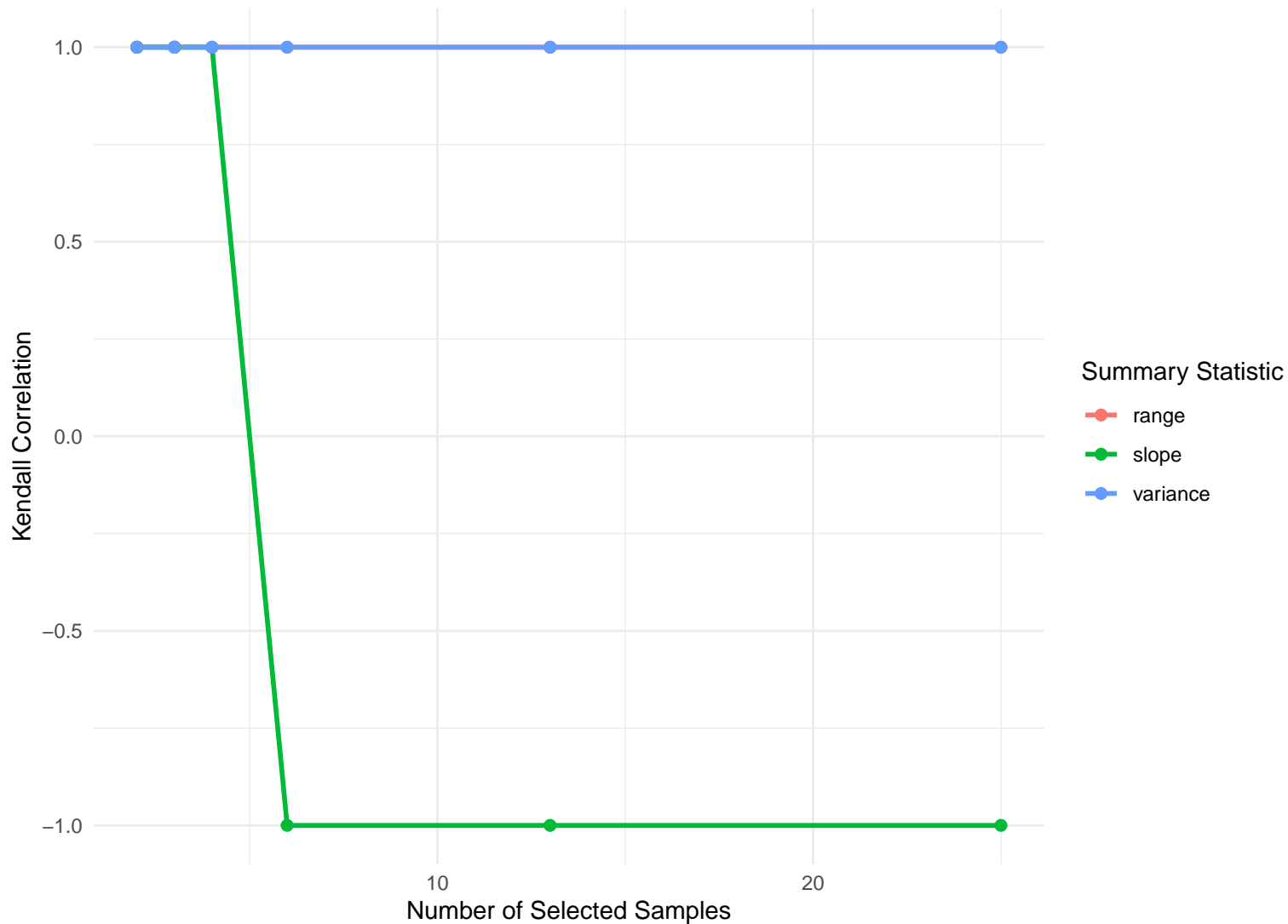
Score: APC (Local, partial1 , Genotype: gaussian)



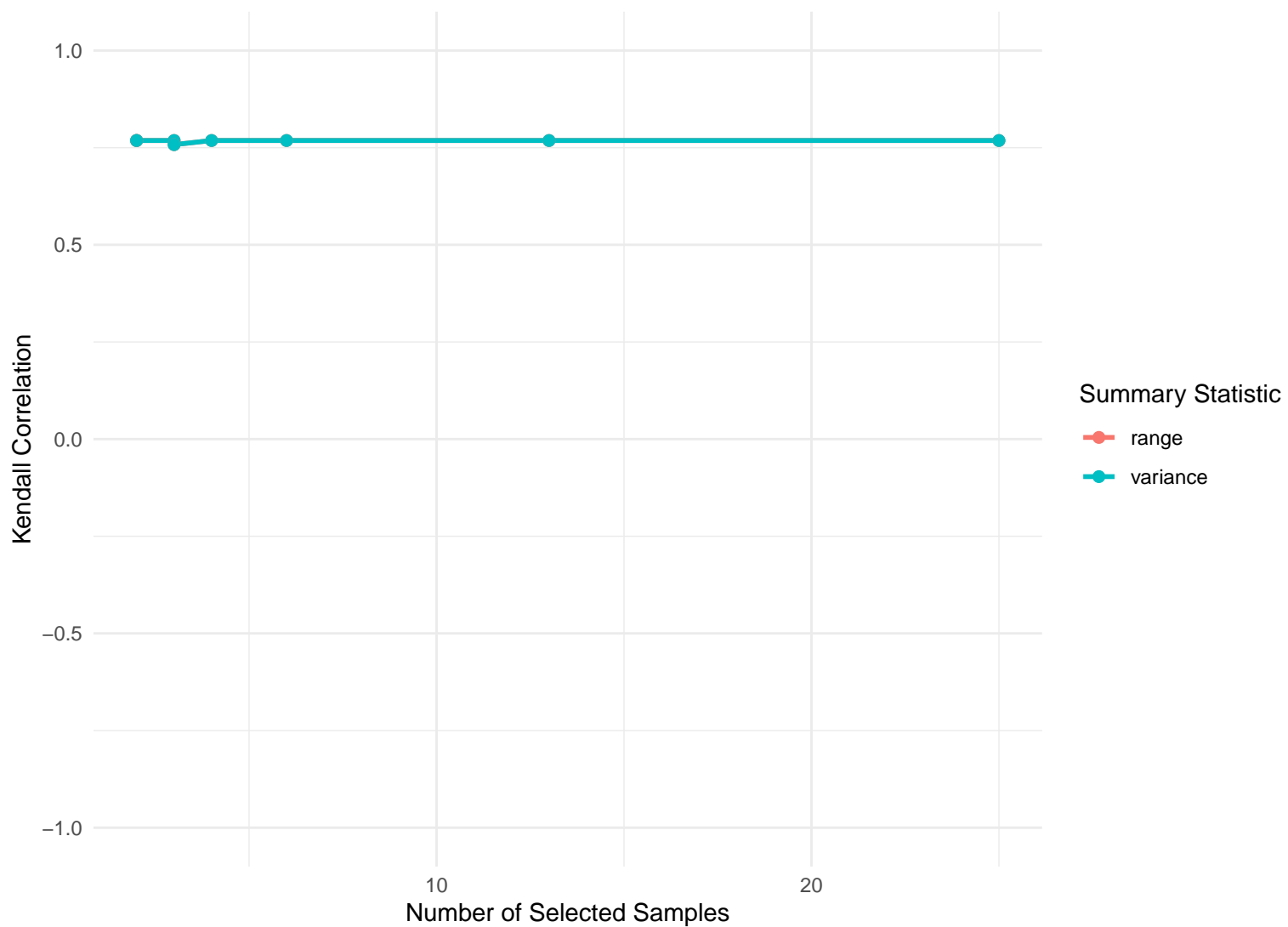
Score: APC (Local, partial1 , Genotype: sinusoidal)



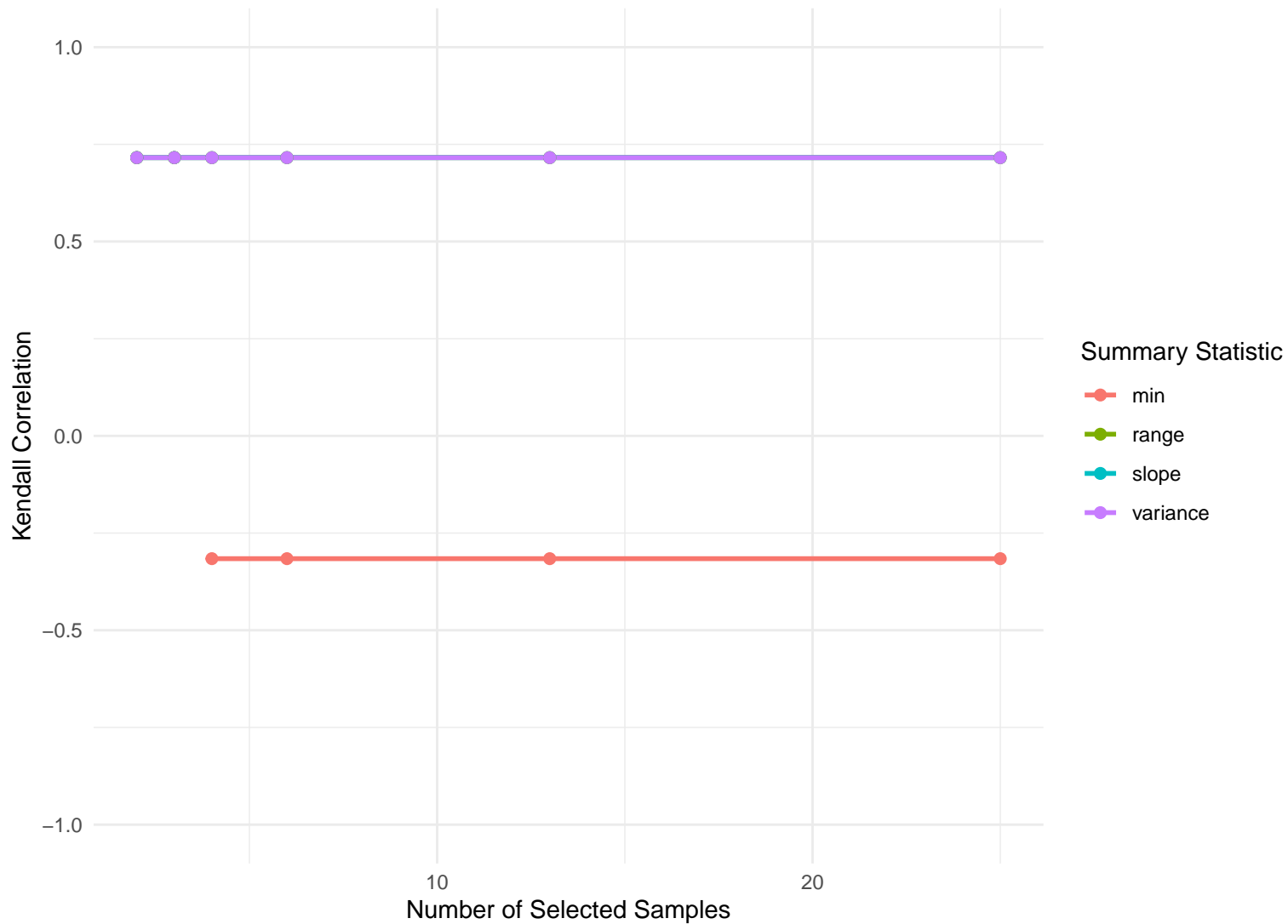
Score: APC (Local, partial1 , Genotype: wave)



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

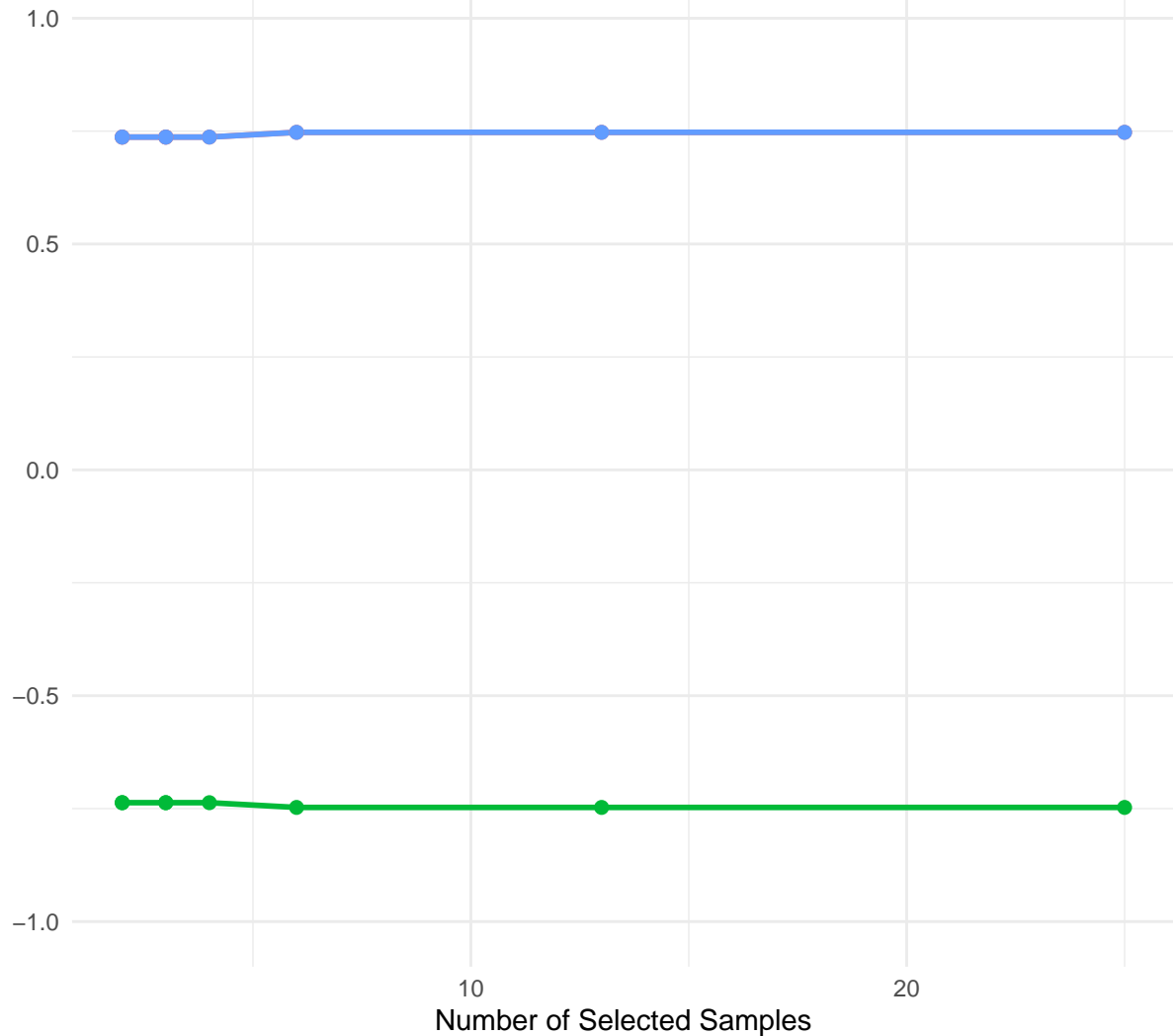


Score: SI (Local, partial1 , Genotype: gaussian)



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

Kendall Correlation



Summary Statistic

- range
- slope
- variance

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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

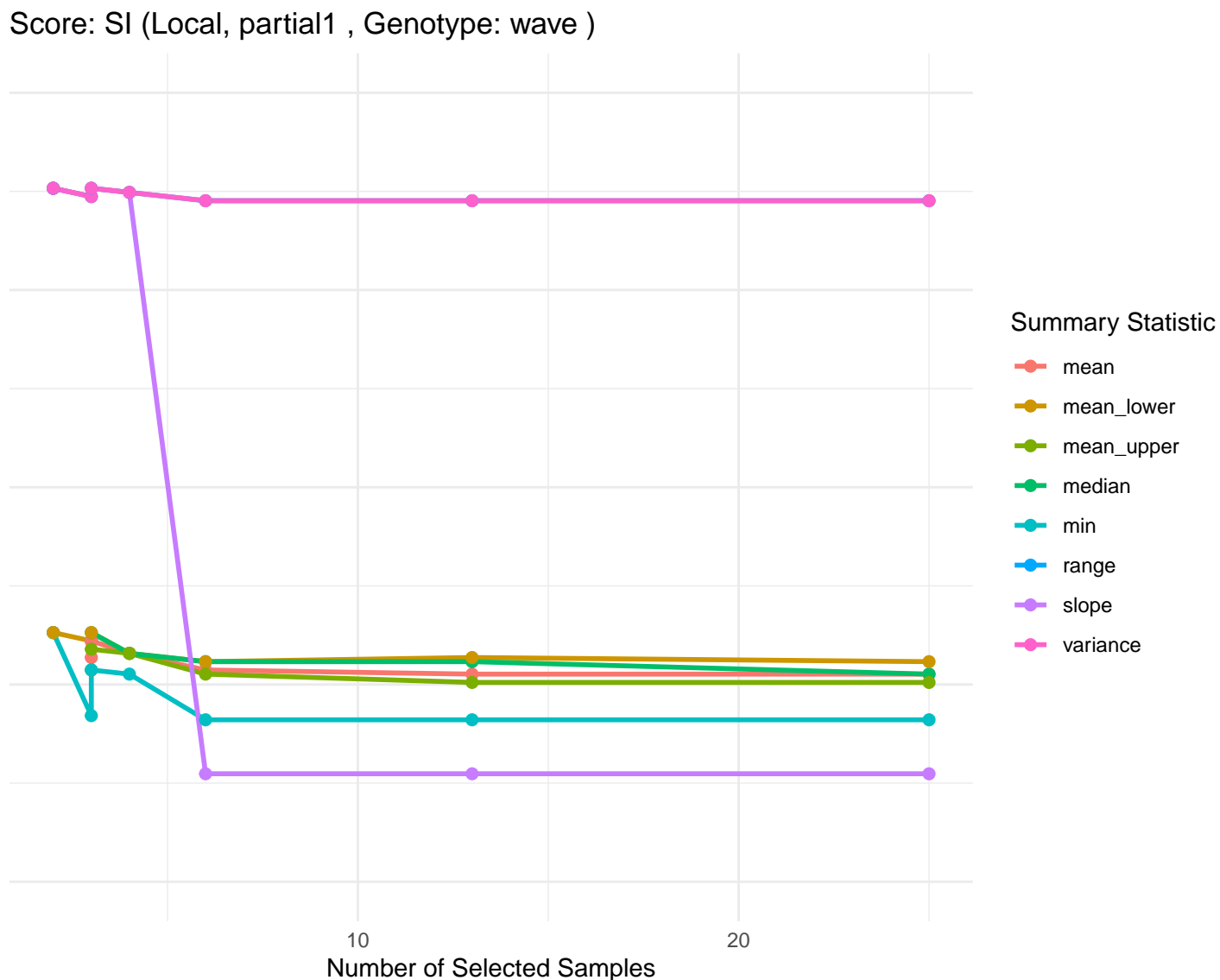
Number of Selected Samples

10

20

Summary Statistic

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

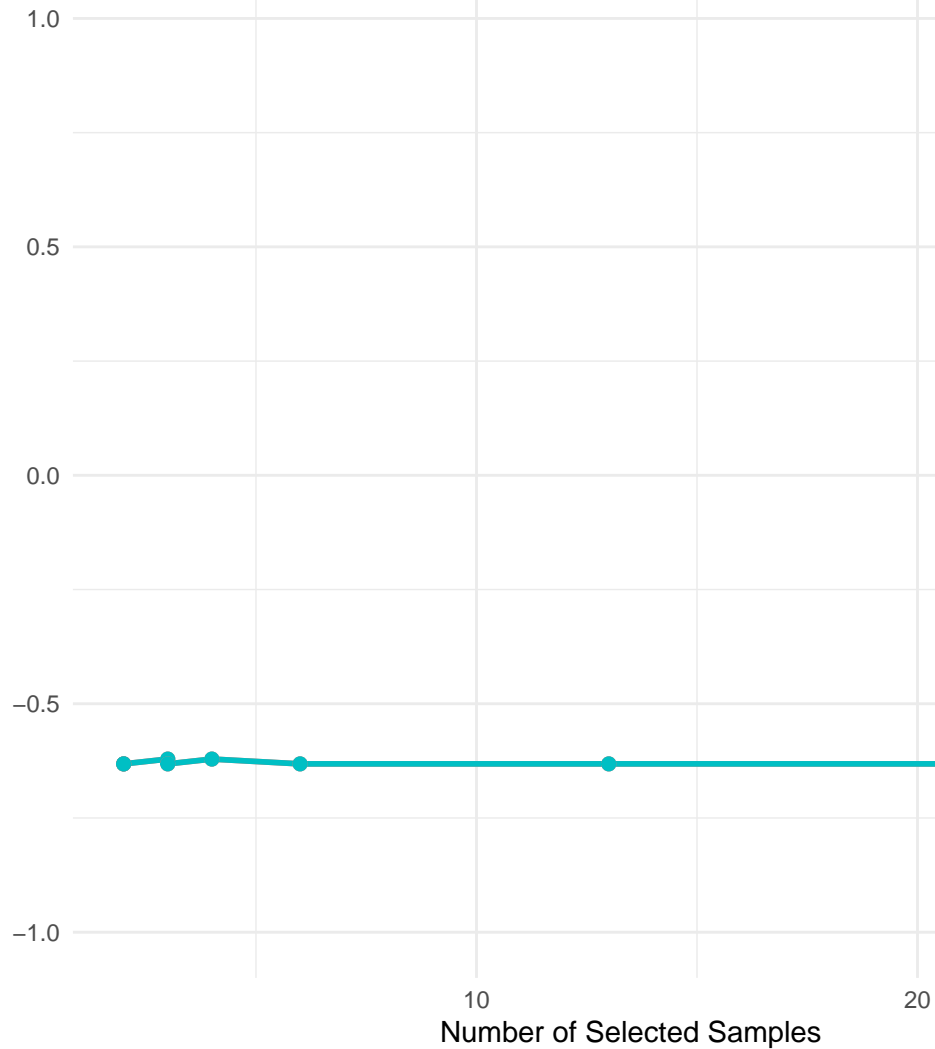


Score: RSI (Local, partial1 , Genotype: linear)

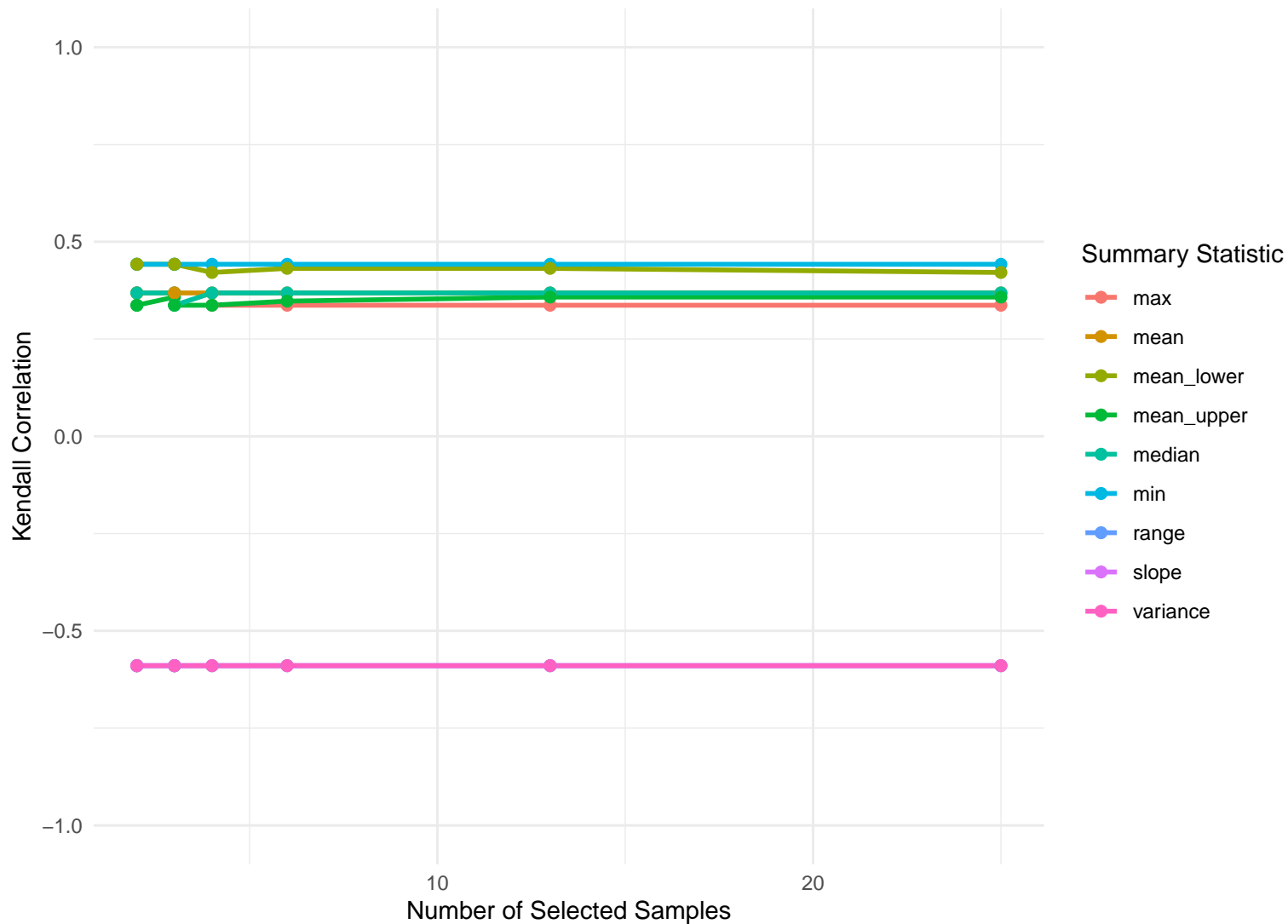
Kendall Correlation

Summary Statistic

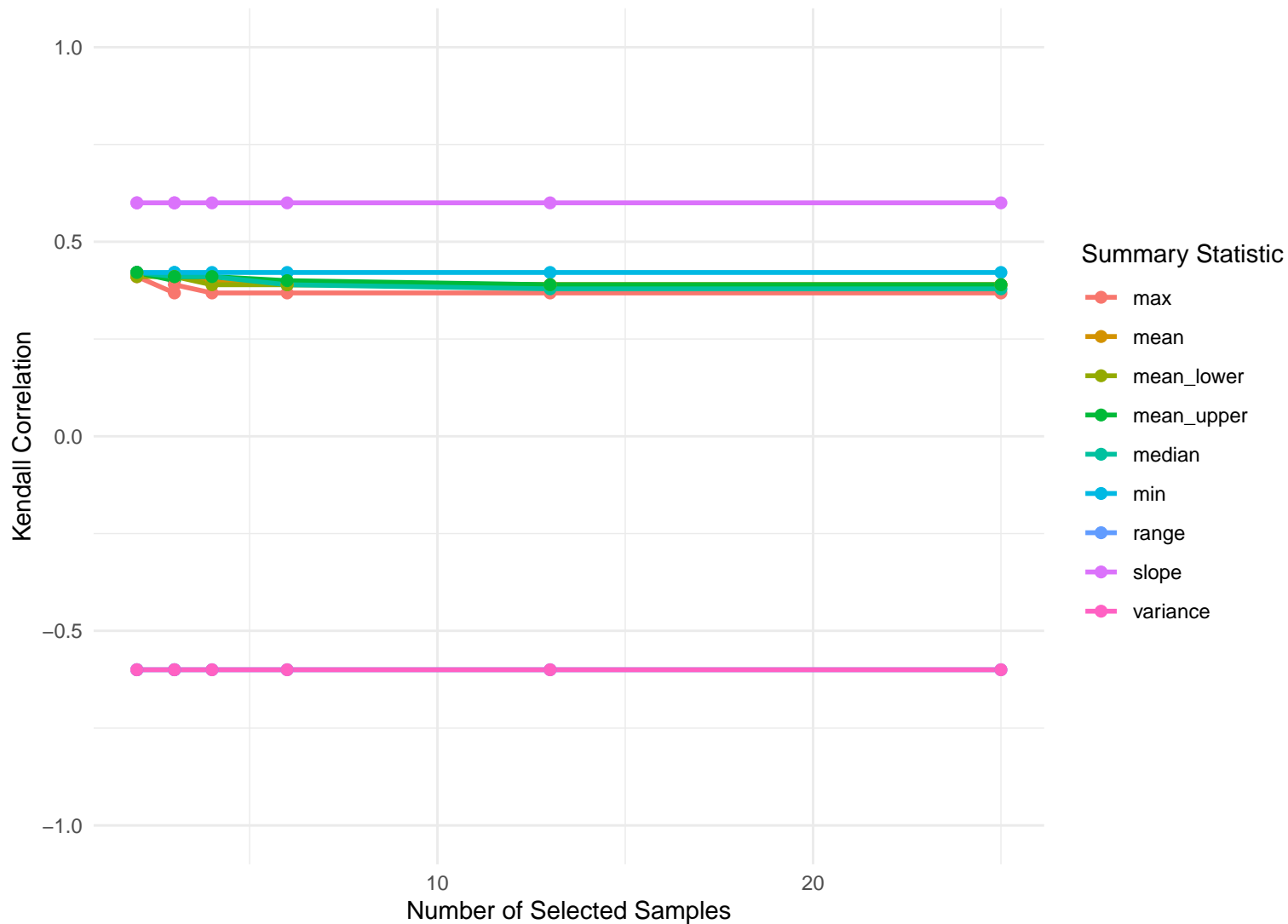
range
variance



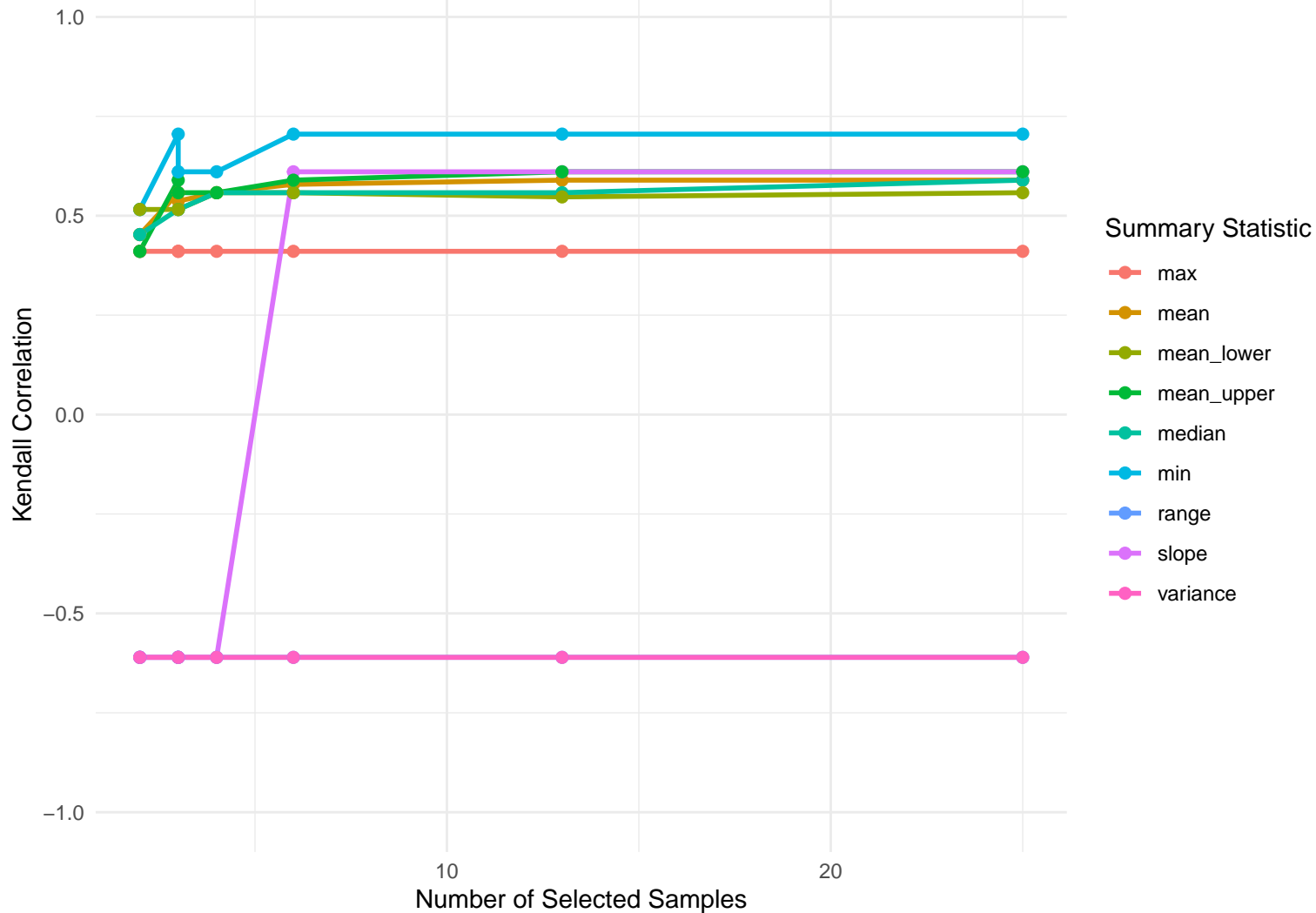
Score: RSI (Local, partial1 , Genotype: gaussian)



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

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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

10

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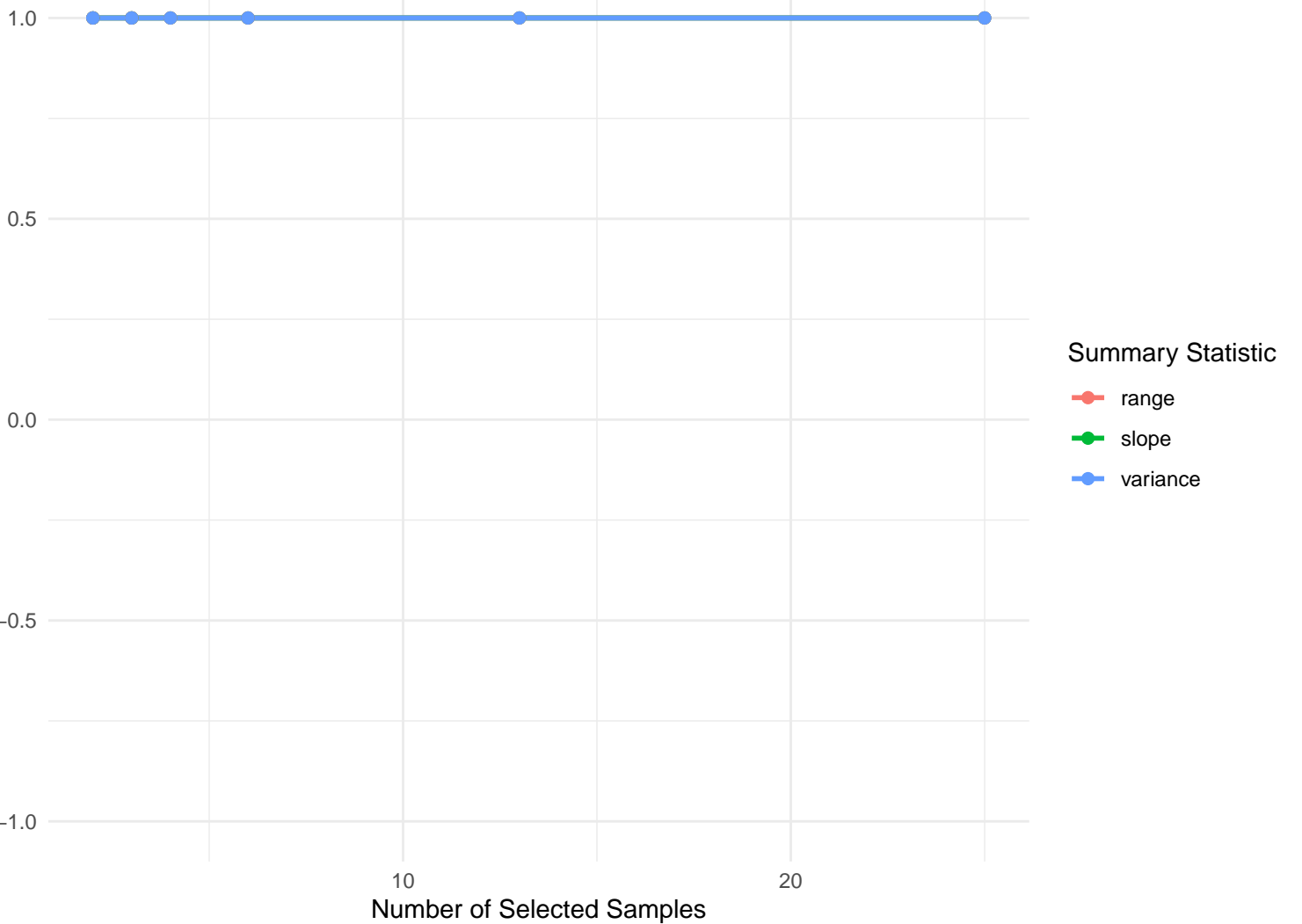
Number of Selected Samples

Summary Statistic

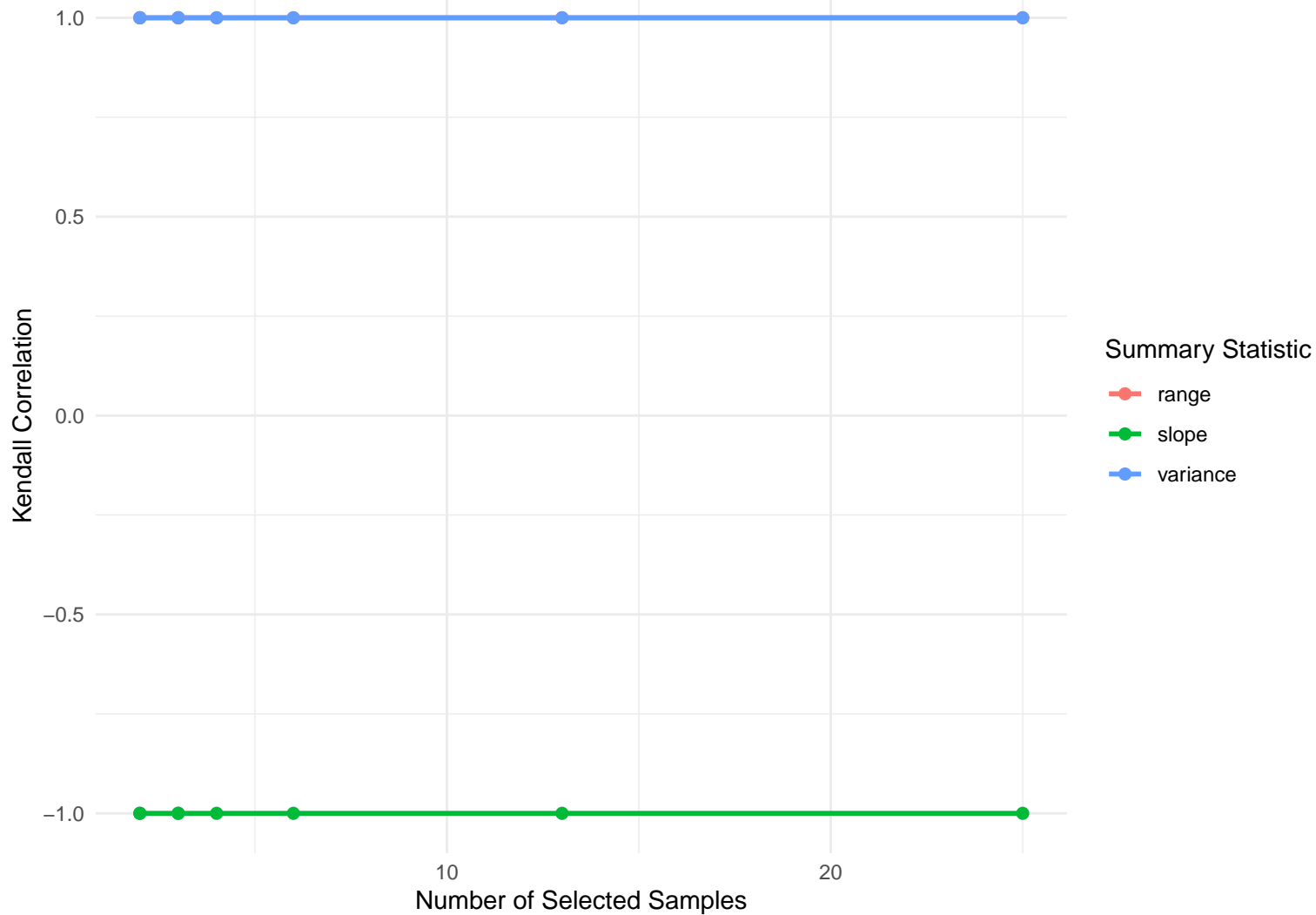
range

slope

variance



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

10

20

Summary Statistic

- range
- slope
- variance

