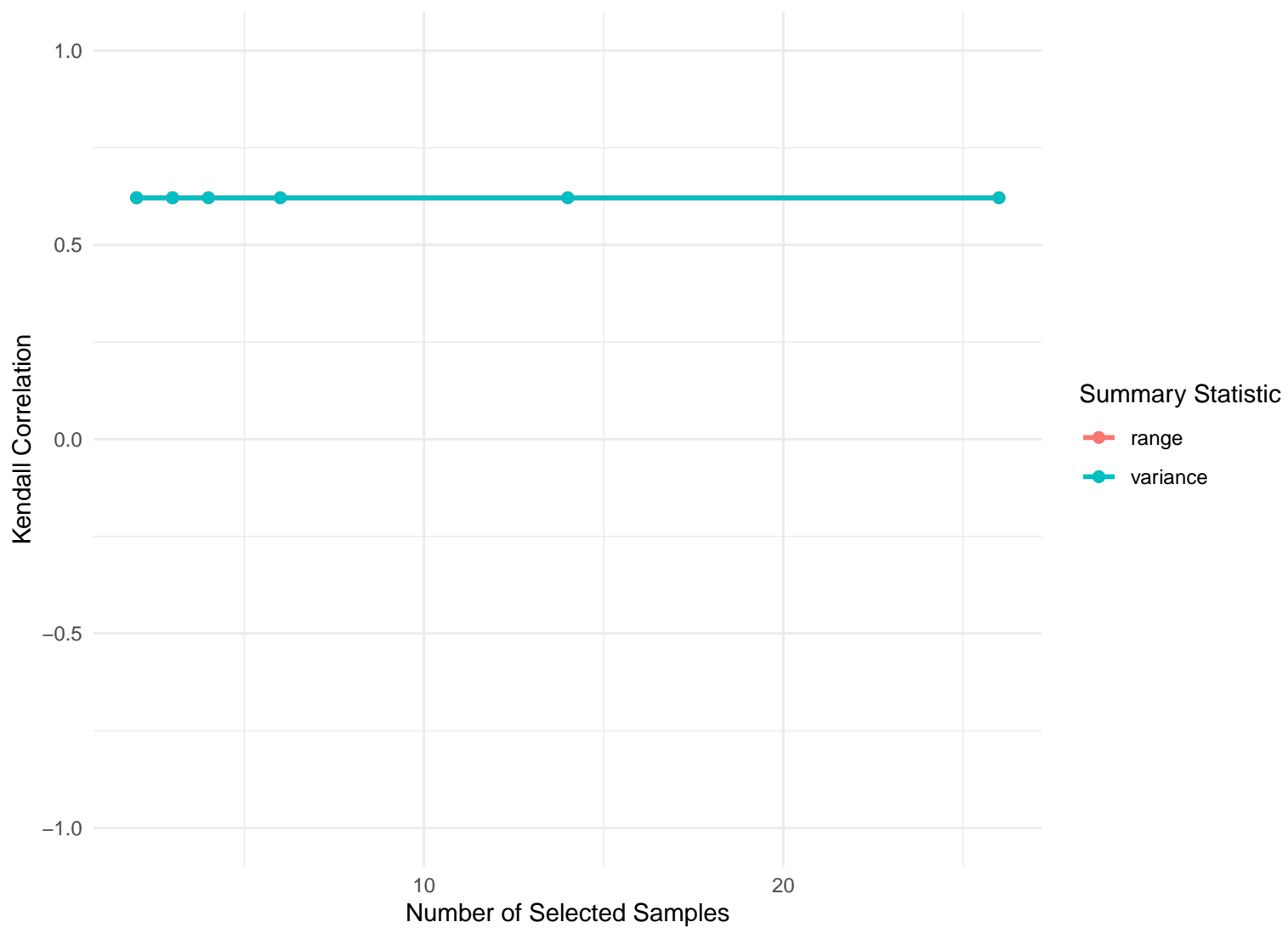


Score: CV_t (Local, partial2 , Genotype: linear)



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

Kendall Correlation

Summary Statistic

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

1.0

0.5

0.0

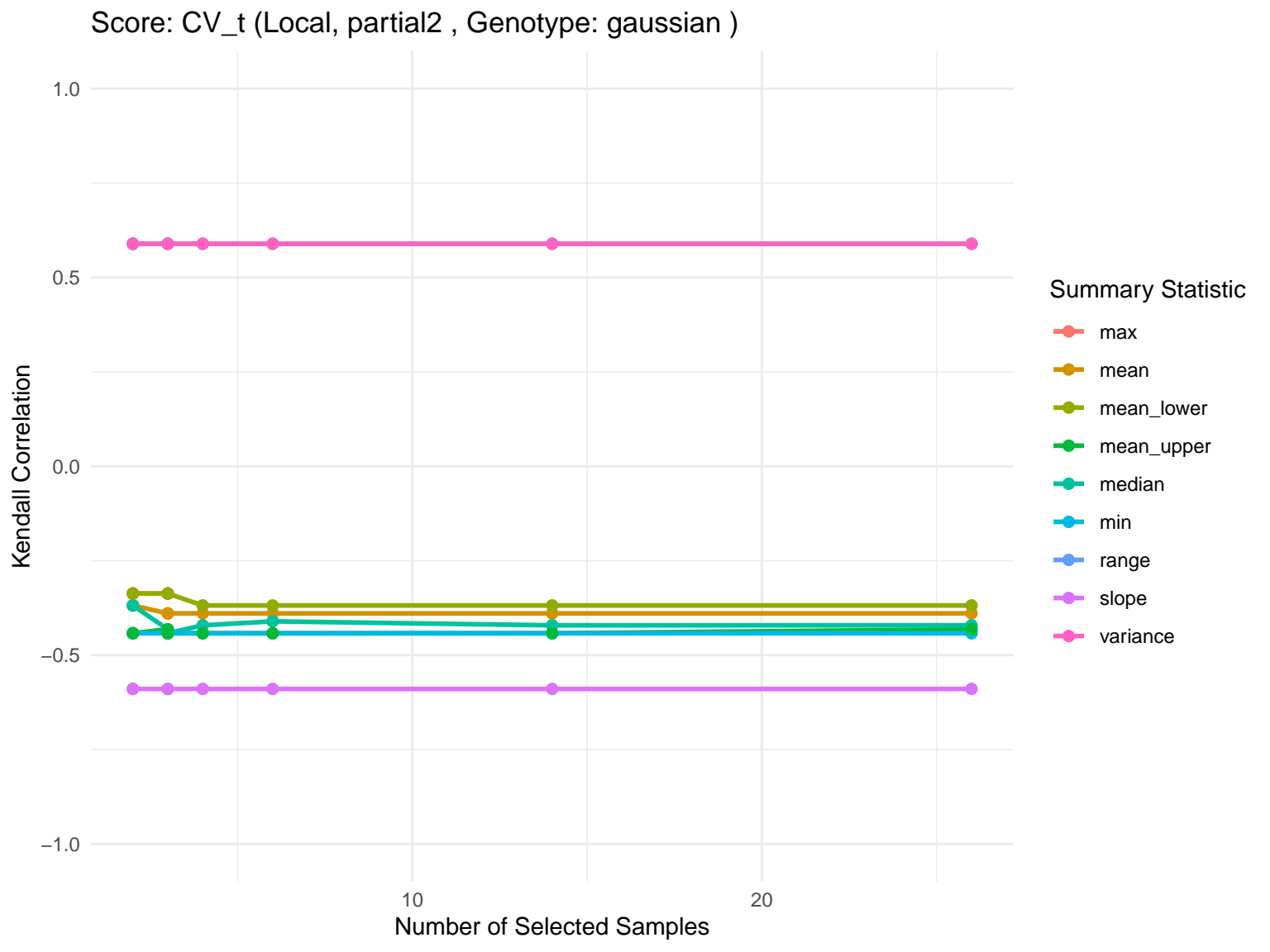
-0.5

-1.0

10

20

Number of Selected Samples

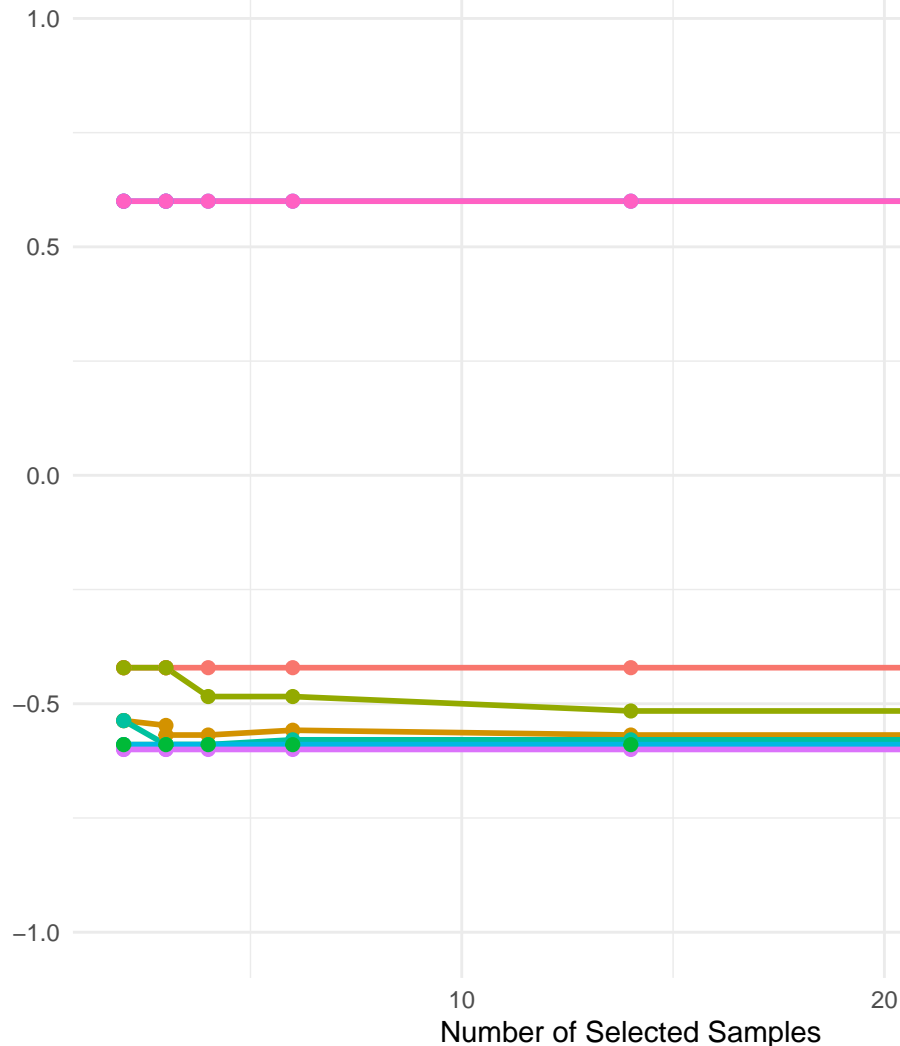


Score: CV_t (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

Summary Statistic

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



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

Kendall Correlation

Summary Statistic

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

1.0

0.5

0.0

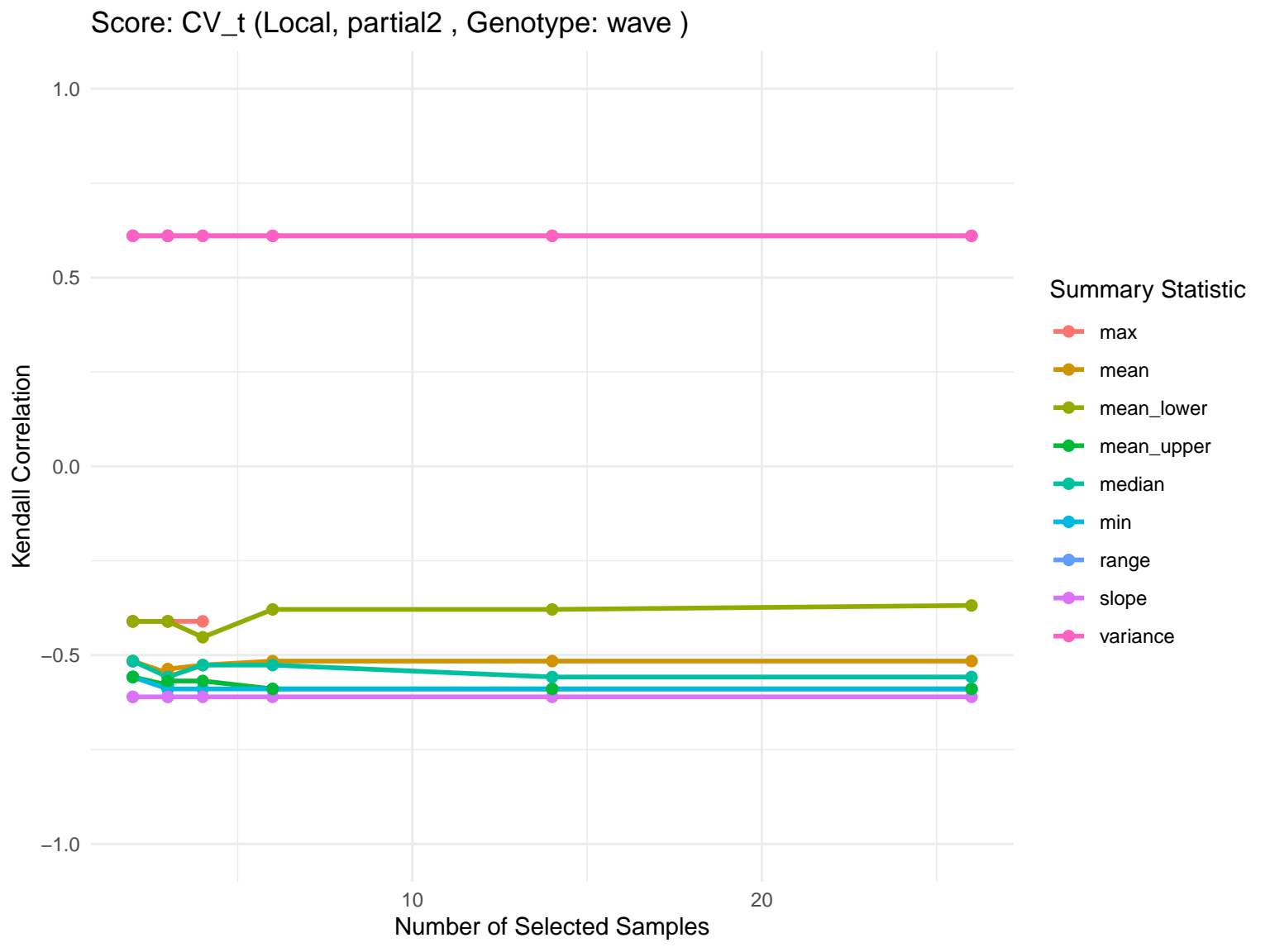
-0.5

-1.0

Number of Selected Samples

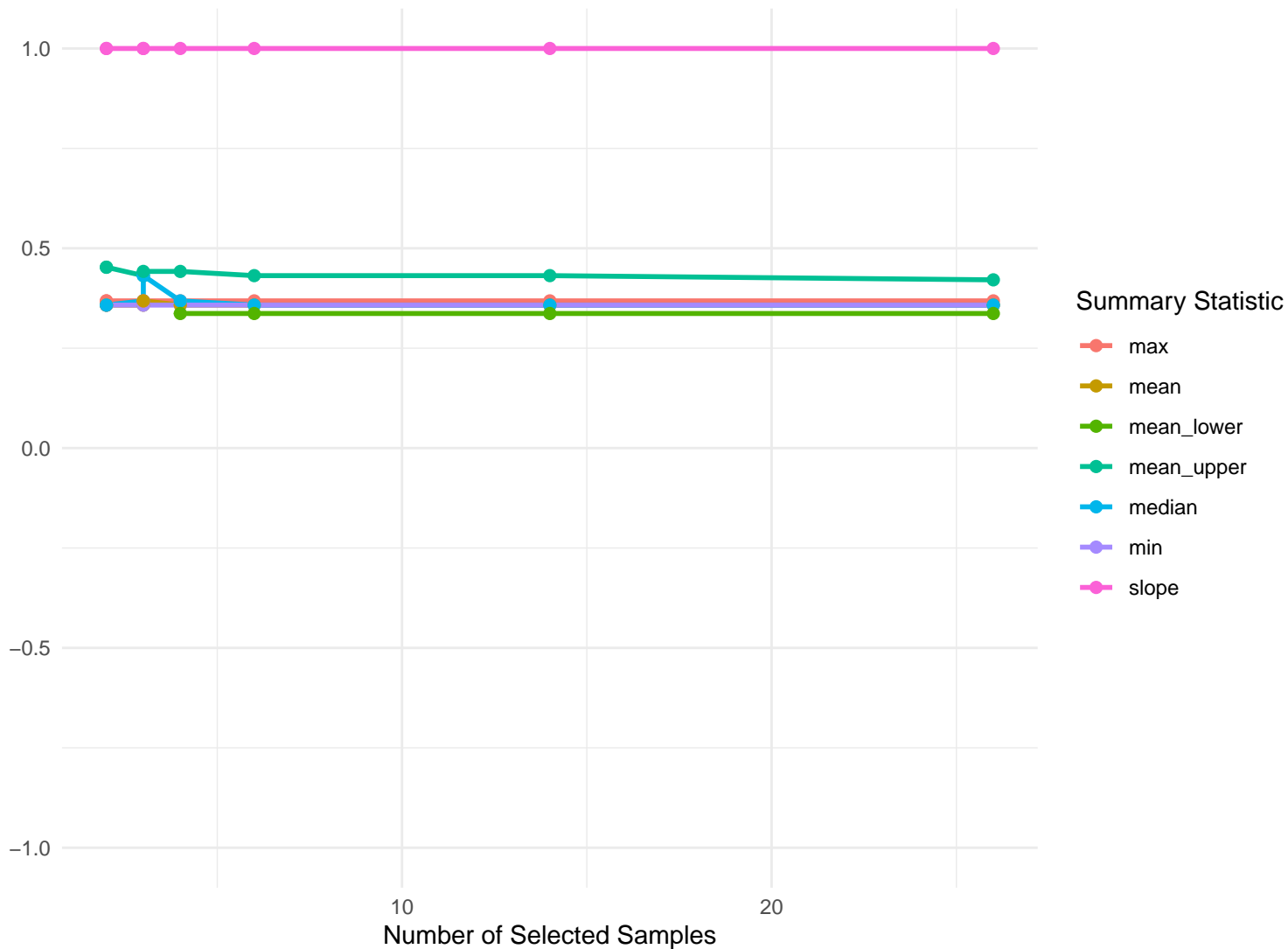
10

20



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

Kendall Correlation



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

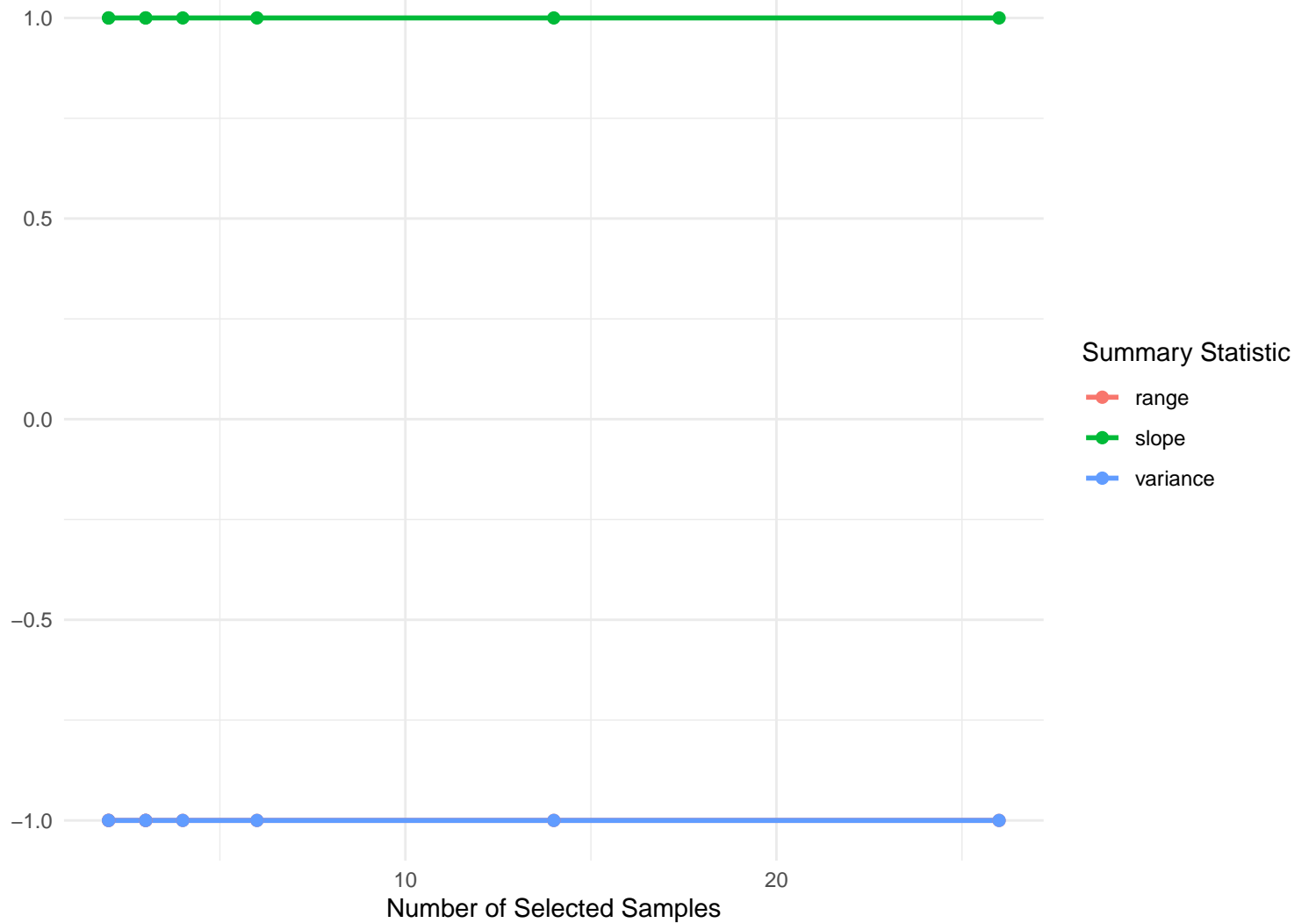
10

20

Number of Selected Samples

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

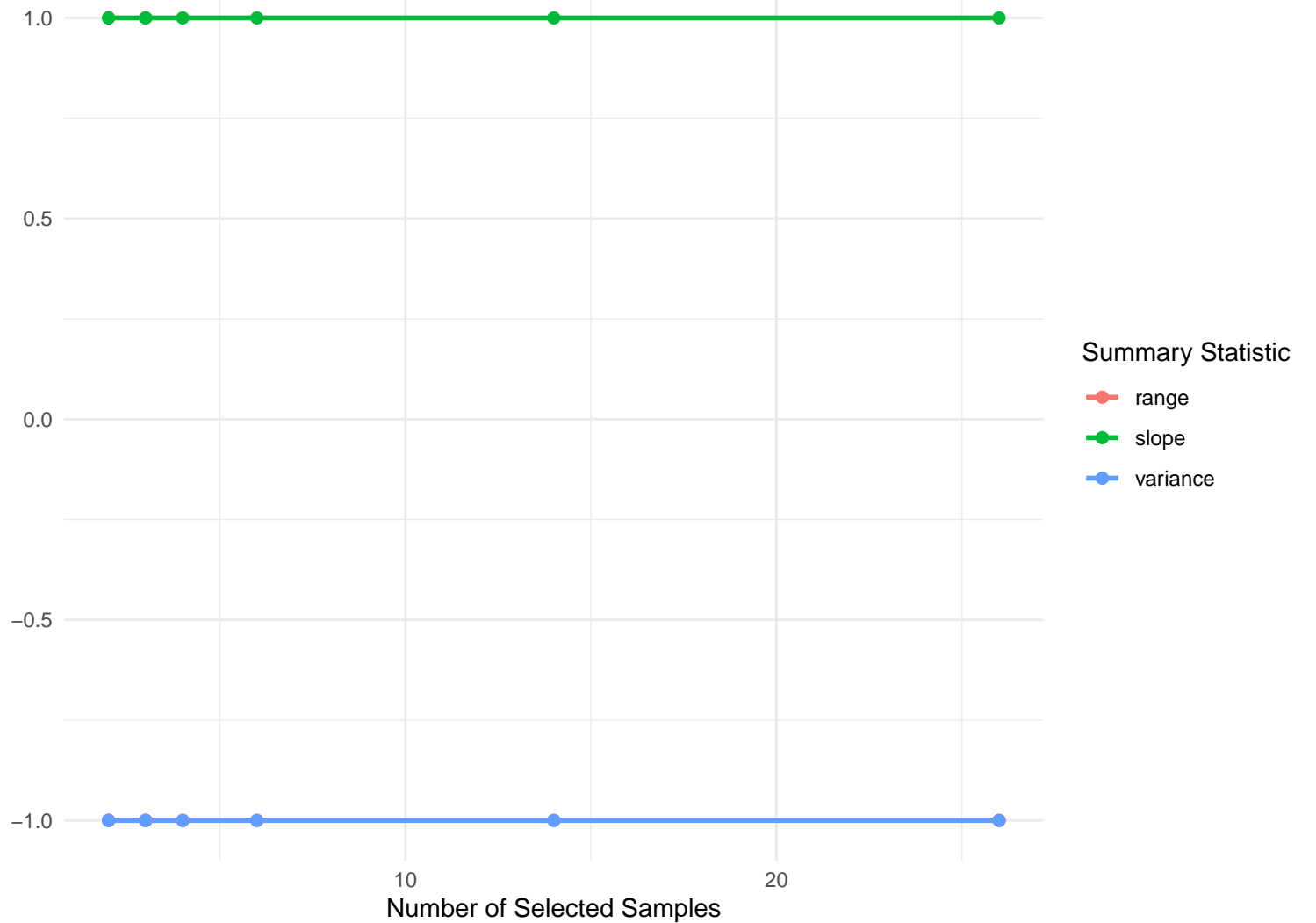
10

20

Number of Selected Samples

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

20

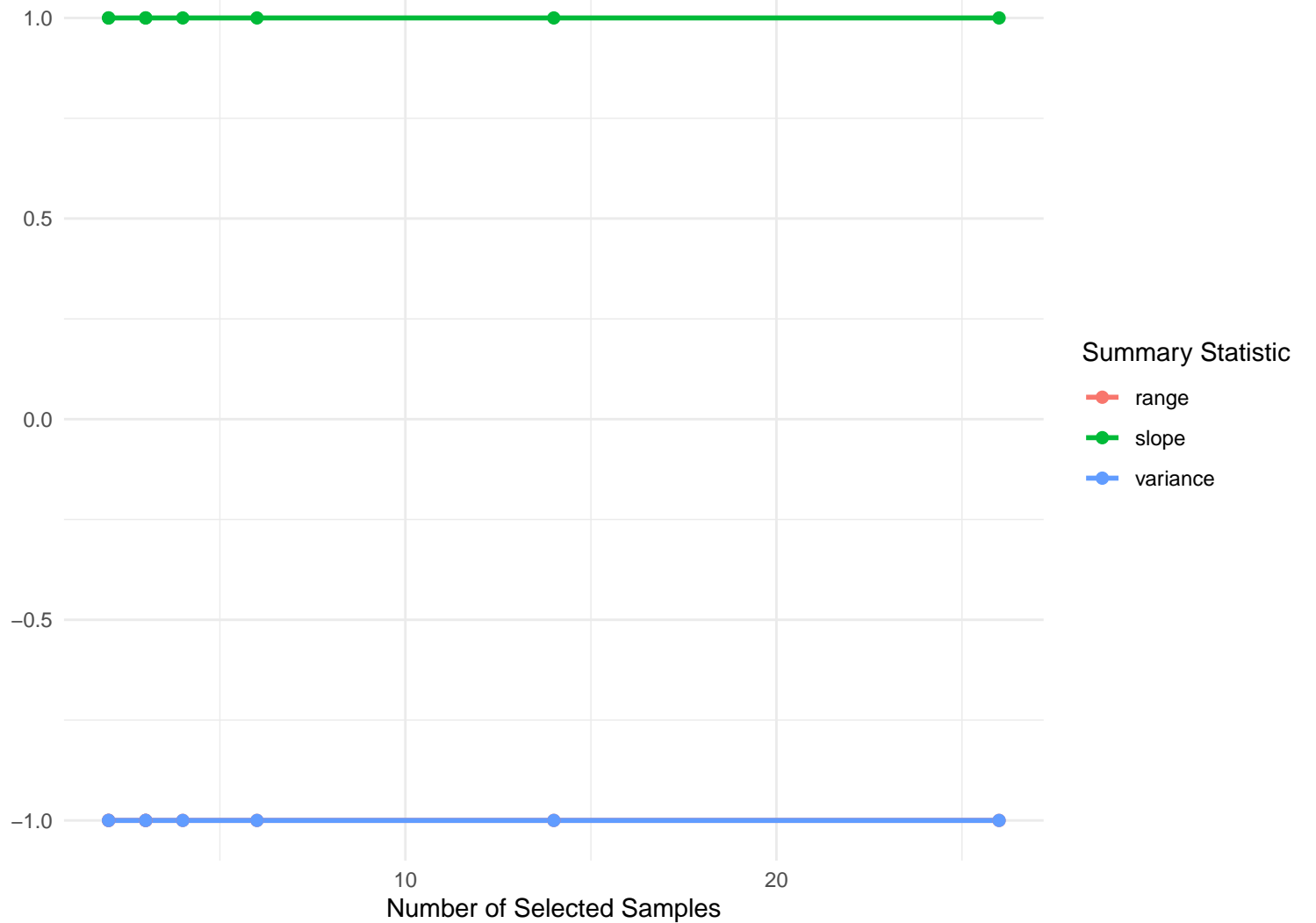
Number of Selected Samples

Summary Statistic

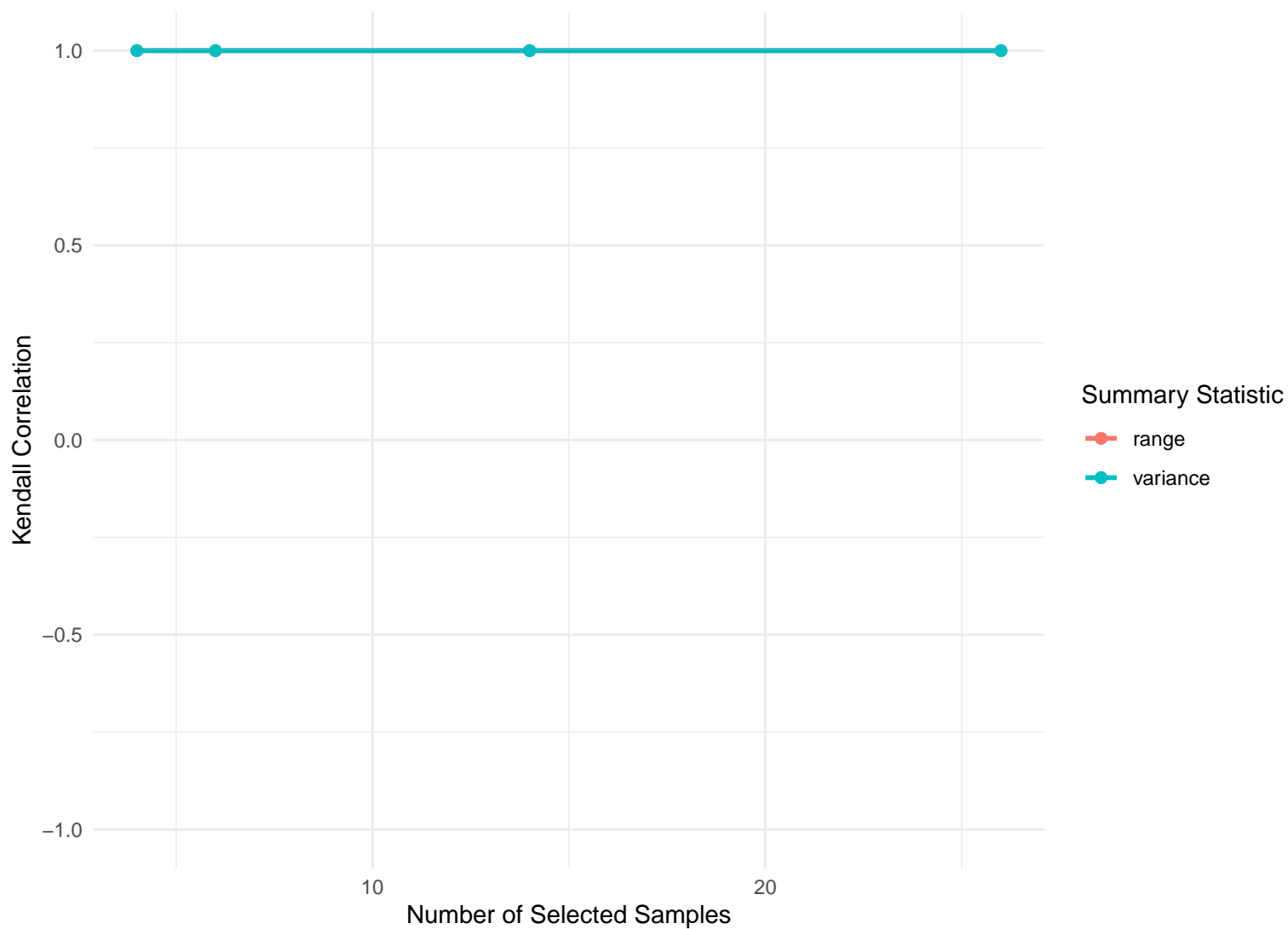
range

slope

variance



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

20

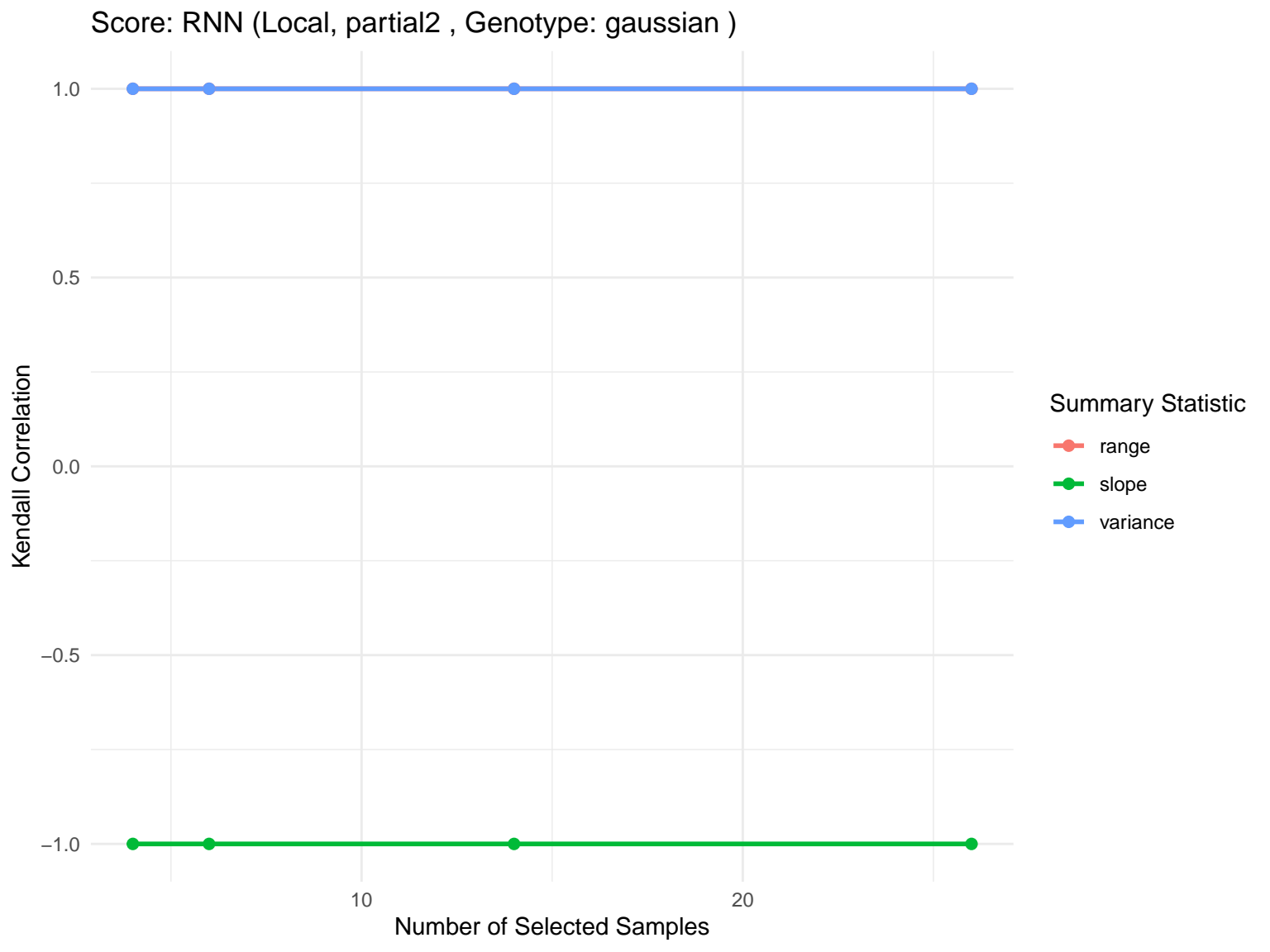
Number of Selected Samples

Summary Statistic

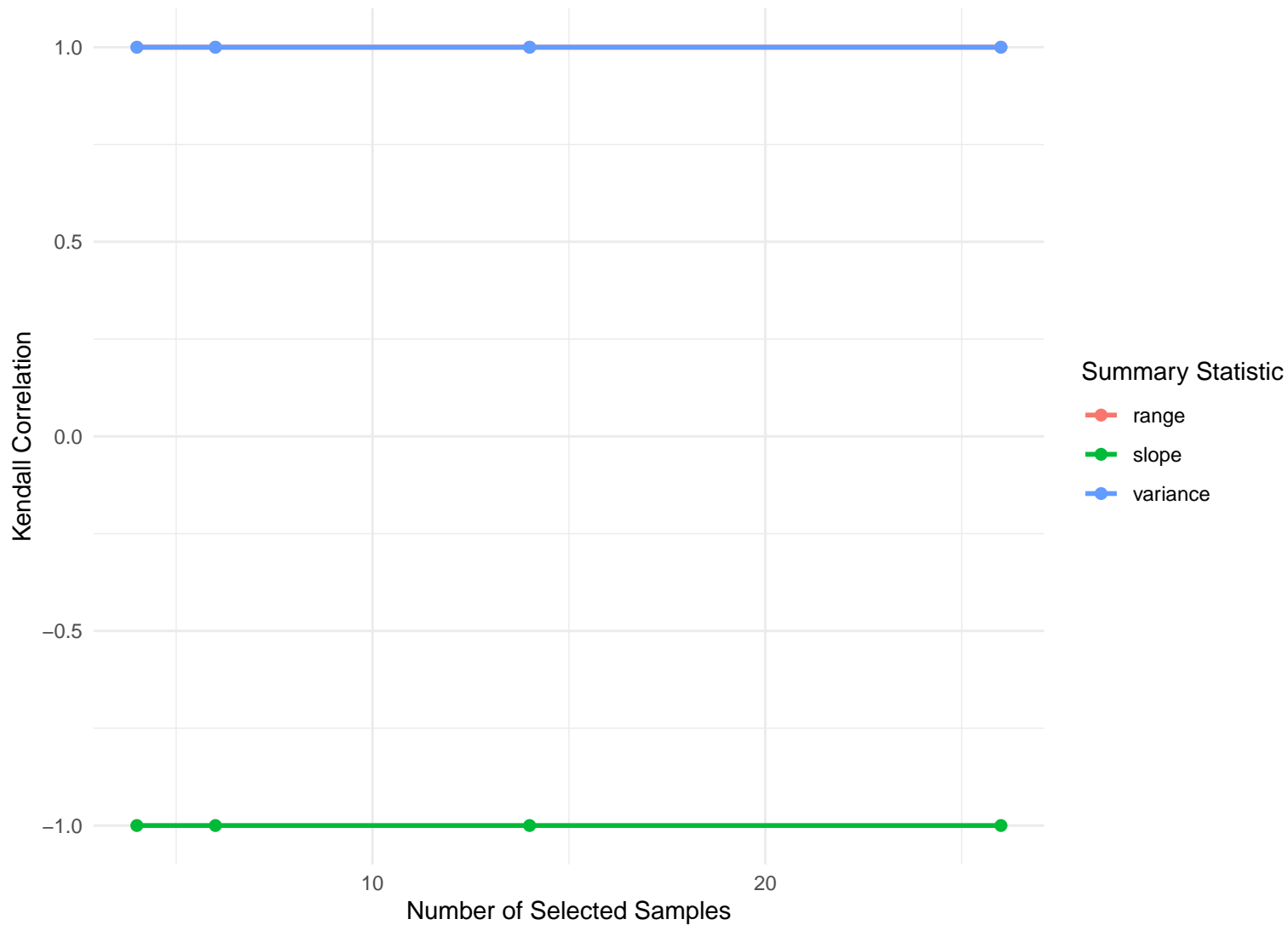
range

slope

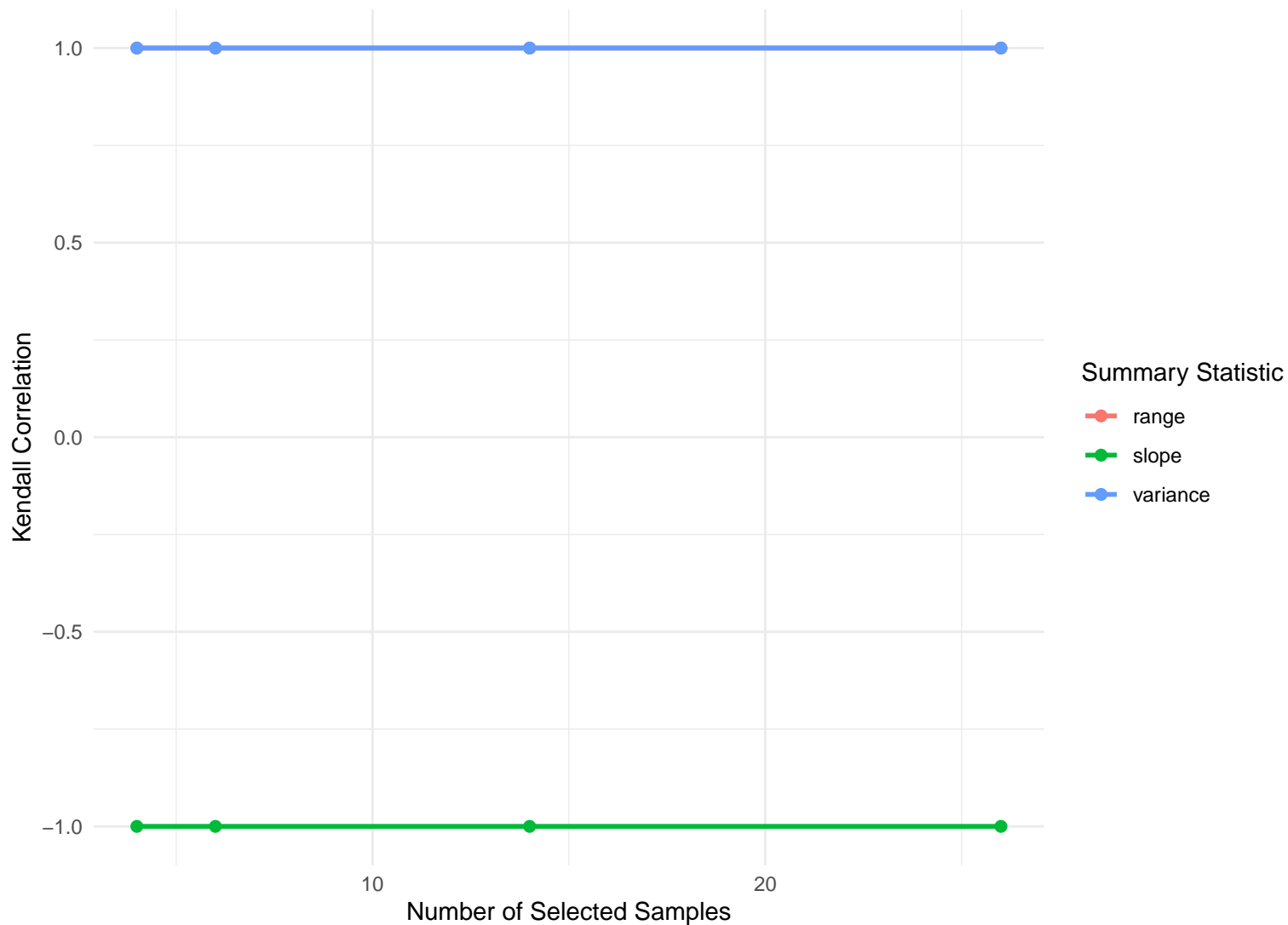
variance



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

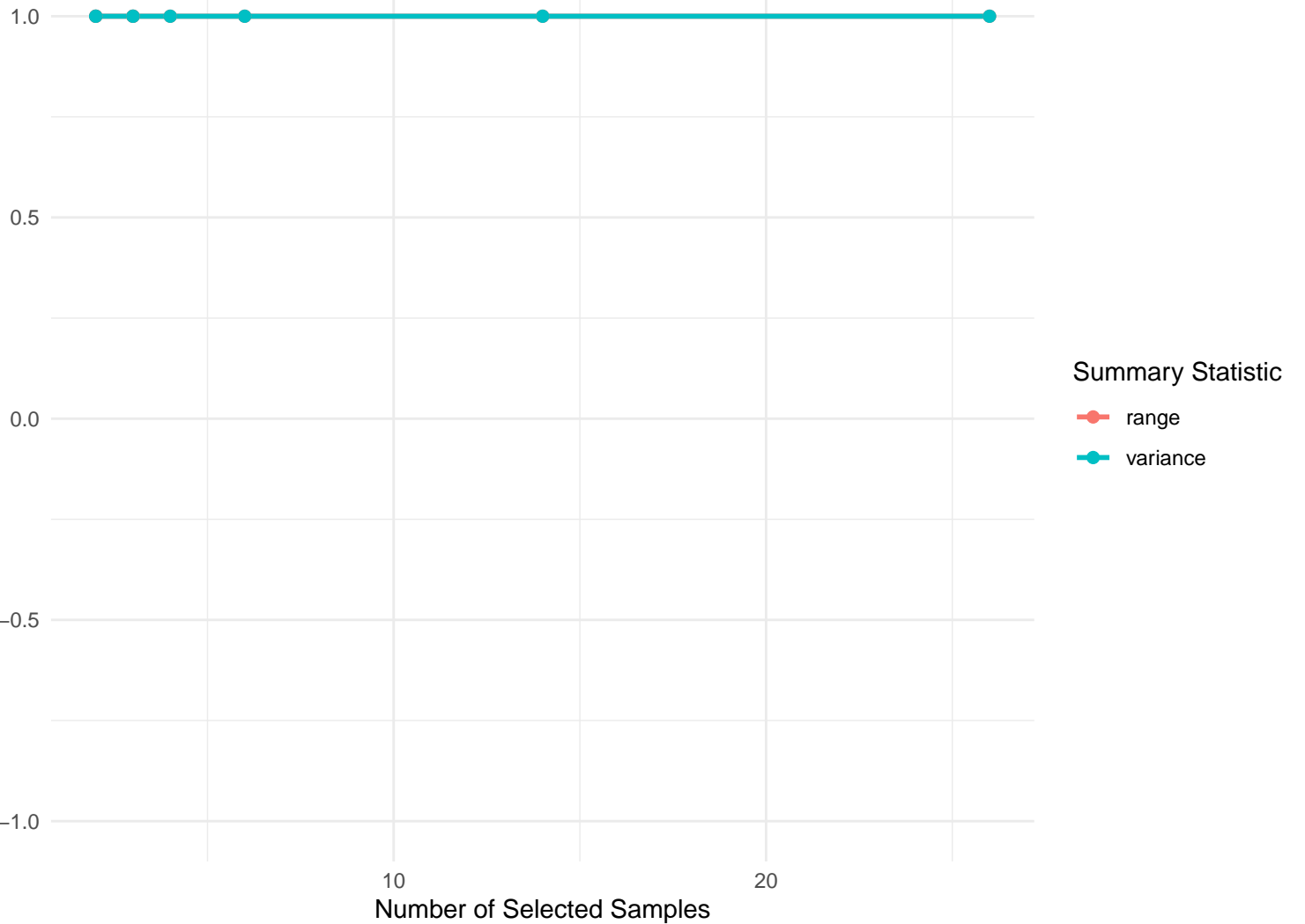
20

Number of Selected Samples

Summary Statistic

range

variance



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

Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

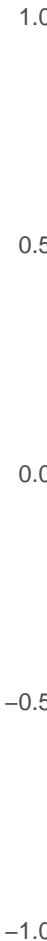
Number of Selected Samples

Summary Statistic

- range
- slope
- variance

10

20



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

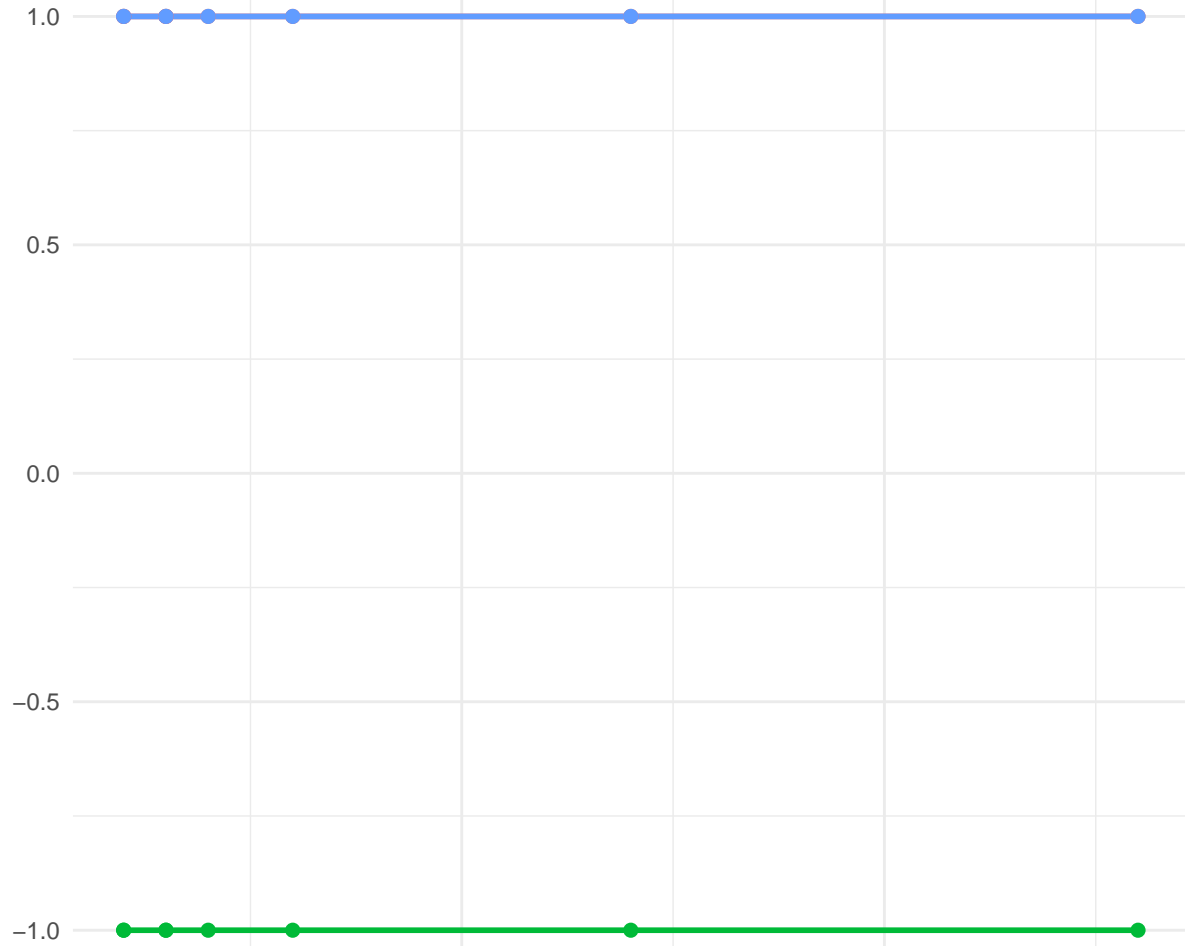
Number of Selected Samples

10

20

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

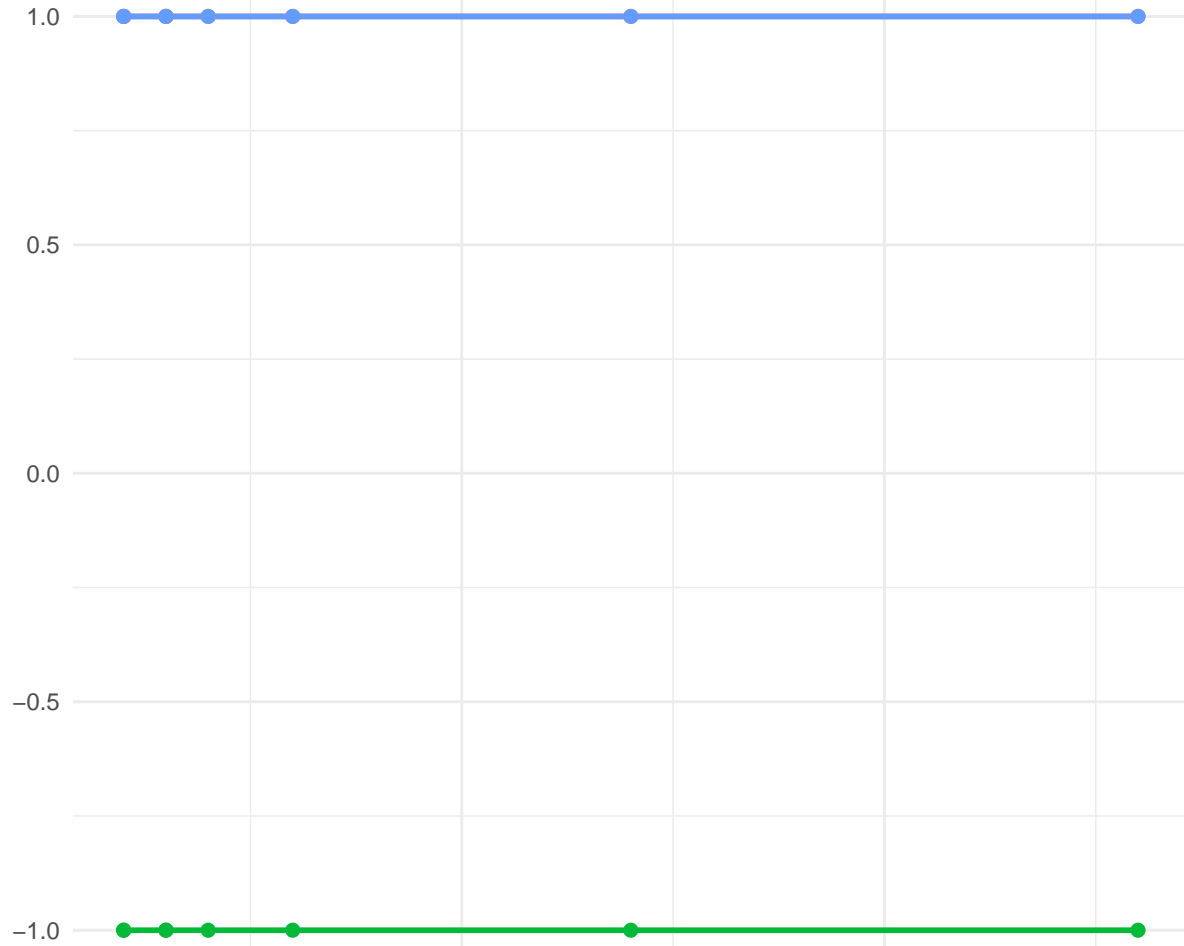
Number of Selected Samples

10

20

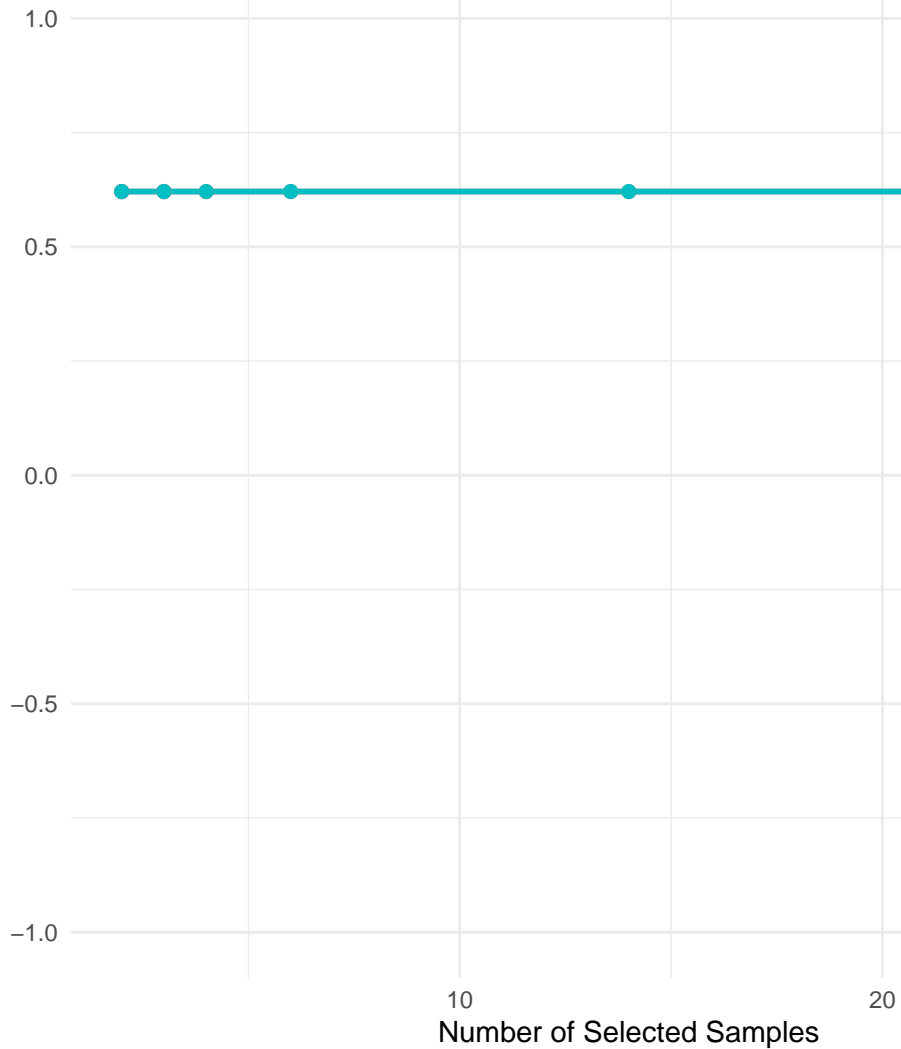
Summary Statistic

- range
- slope
- variance



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

Kendall Correlation



Summary Statistic

- range
- variance

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

Kendall Correlation

Summary Statistic

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

1.0

0.5

0.0

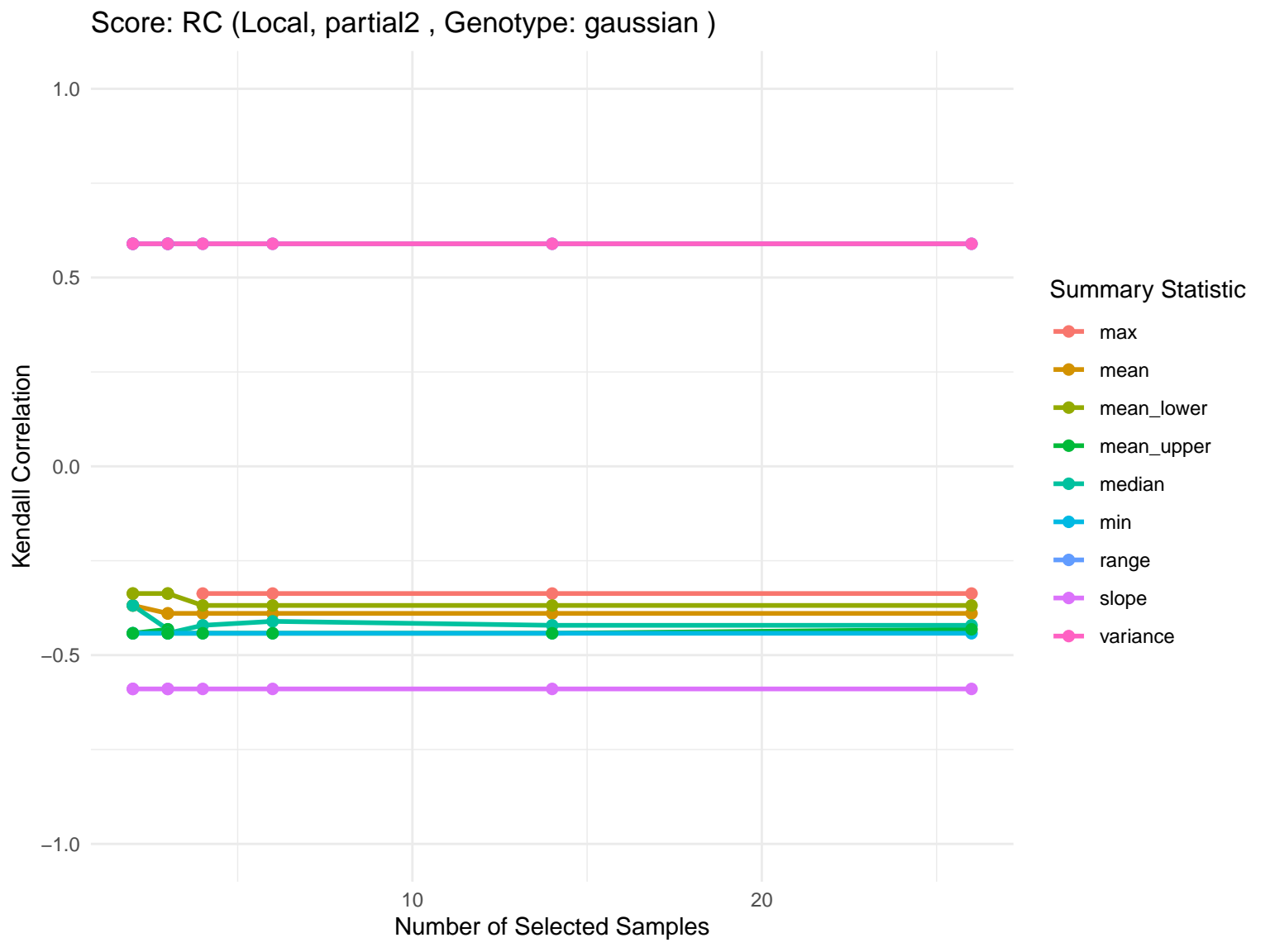
-0.5

-1.0

10

20

Number of Selected Samples

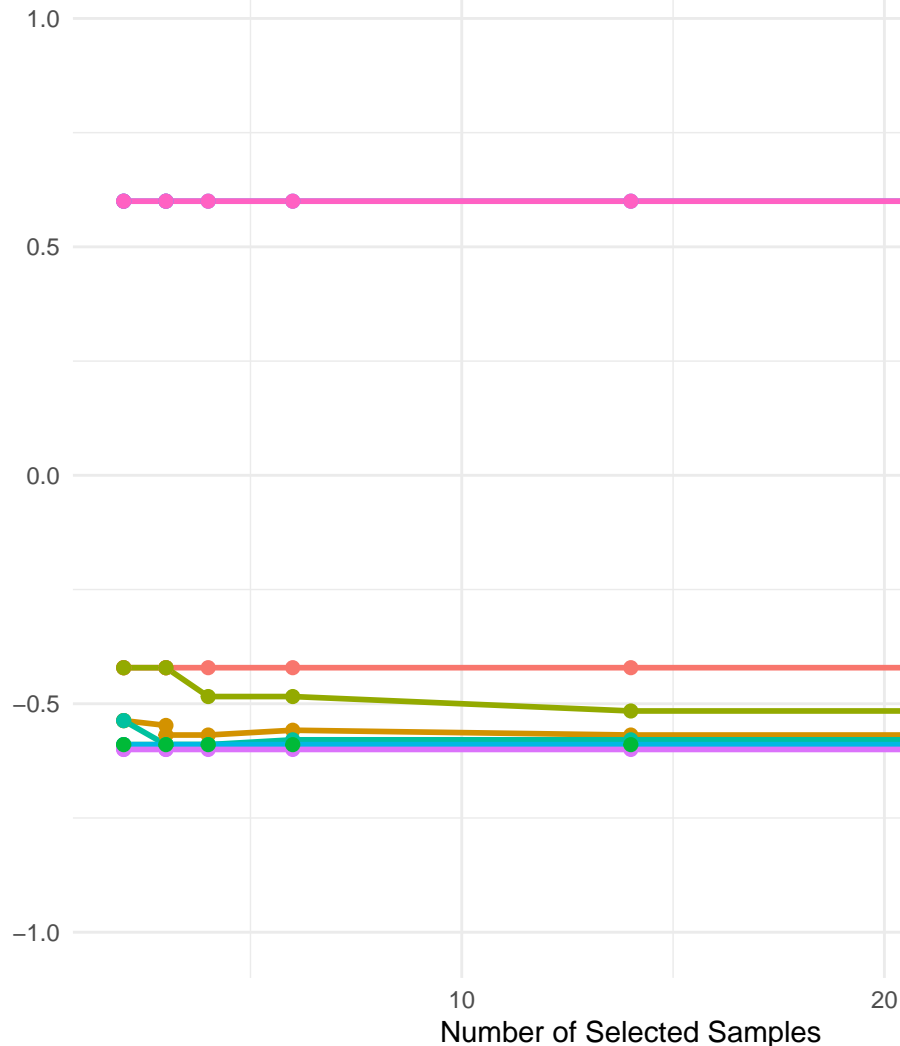


Score: RC (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

Summary Statistic

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



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

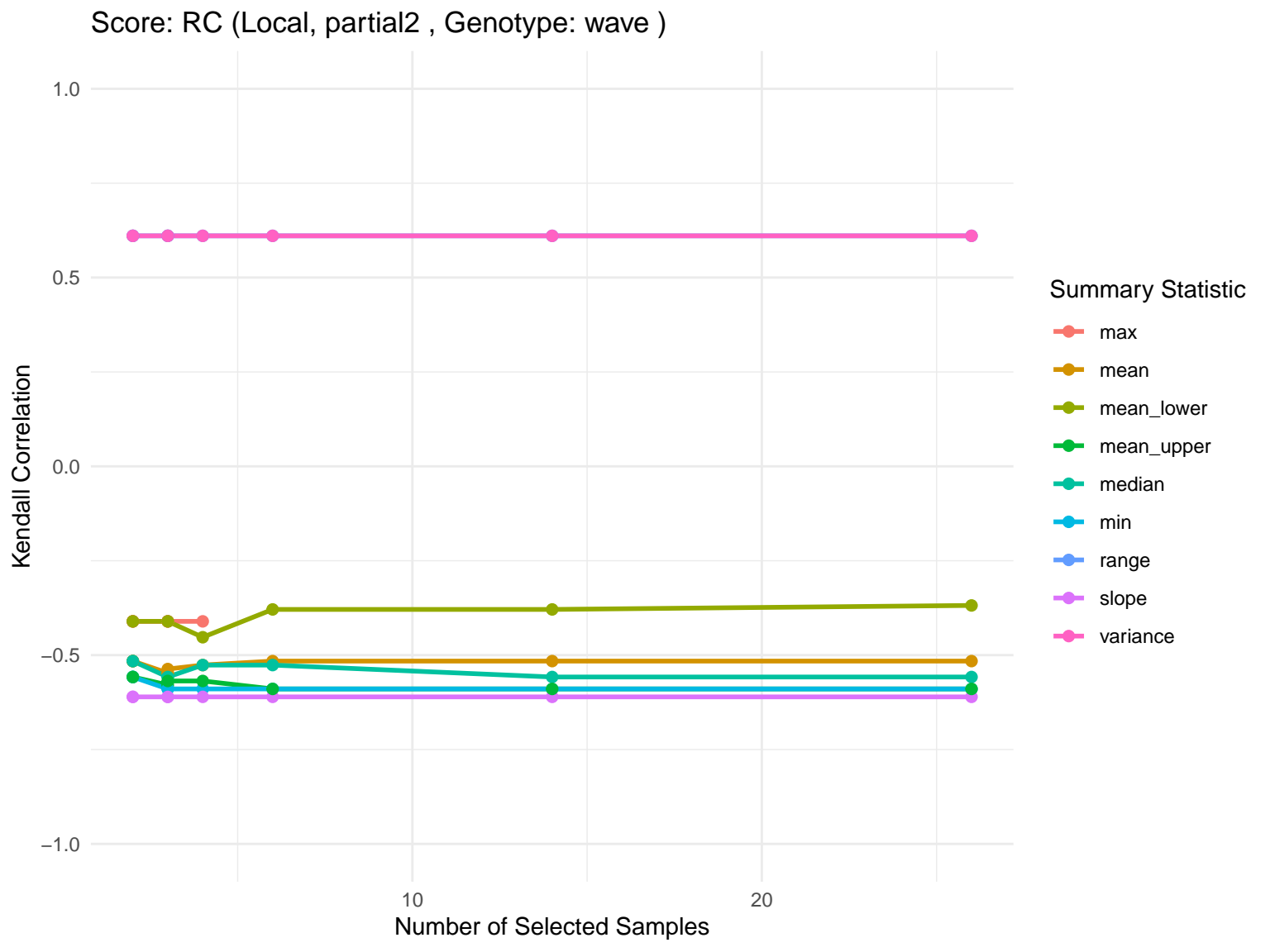
Kendall Correlation

Summary Statistic

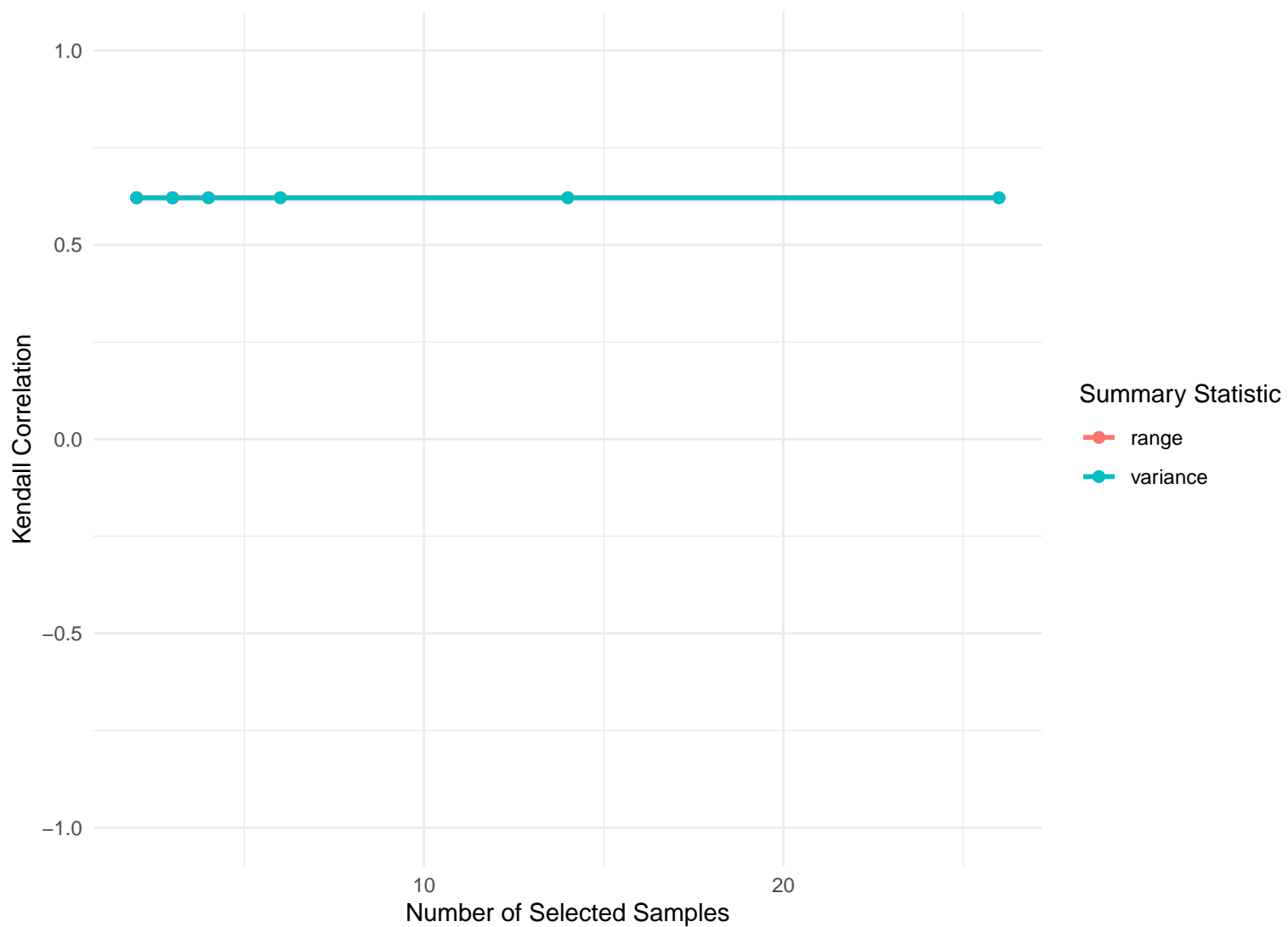
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance

1.0
0.5
0.0
-0.5
-1.0

10 20
Number of Selected Samples



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



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

Kendall Correlation

Summary Statistic

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

1.0

0.5

0.0

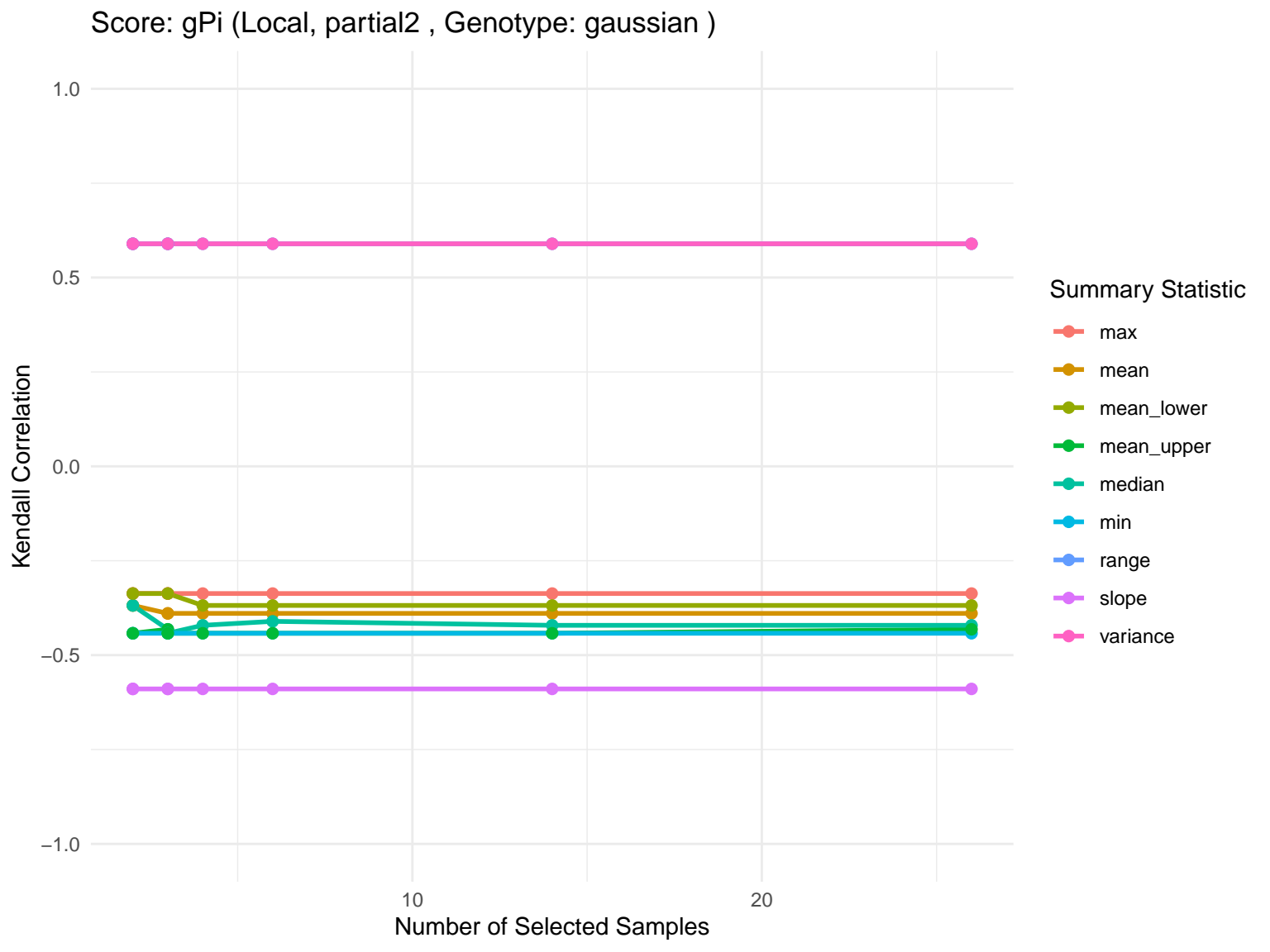
-0.5

-1.0

10

20

Number of Selected Samples

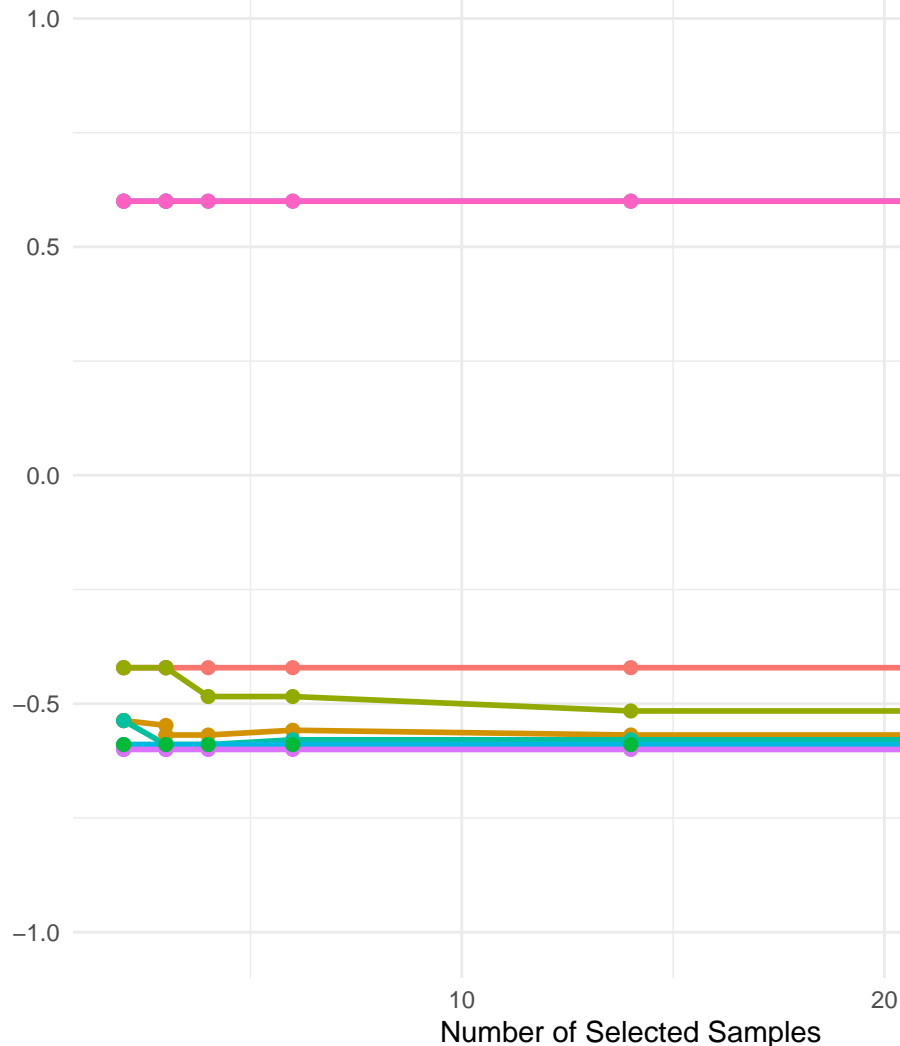


Score: gPi (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

Summary Statistic

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

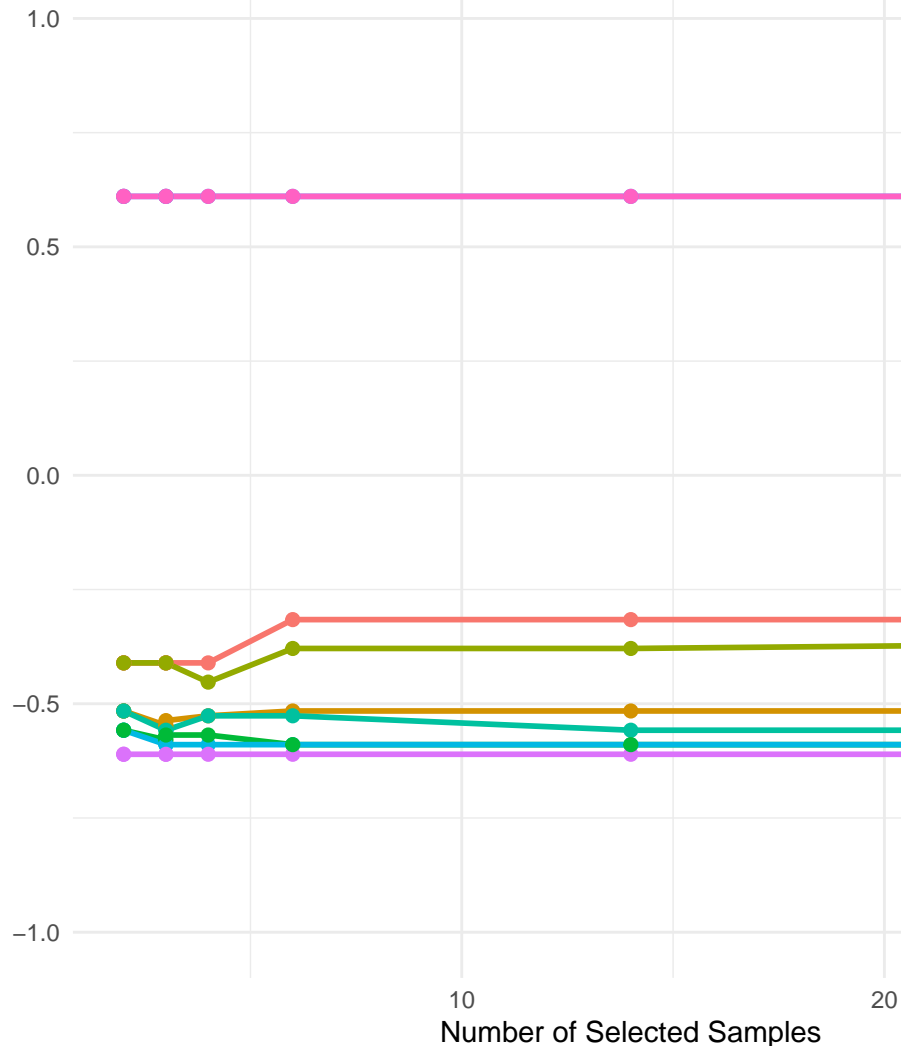


Score: gPi (Local, partial2 , Genotype: wave)

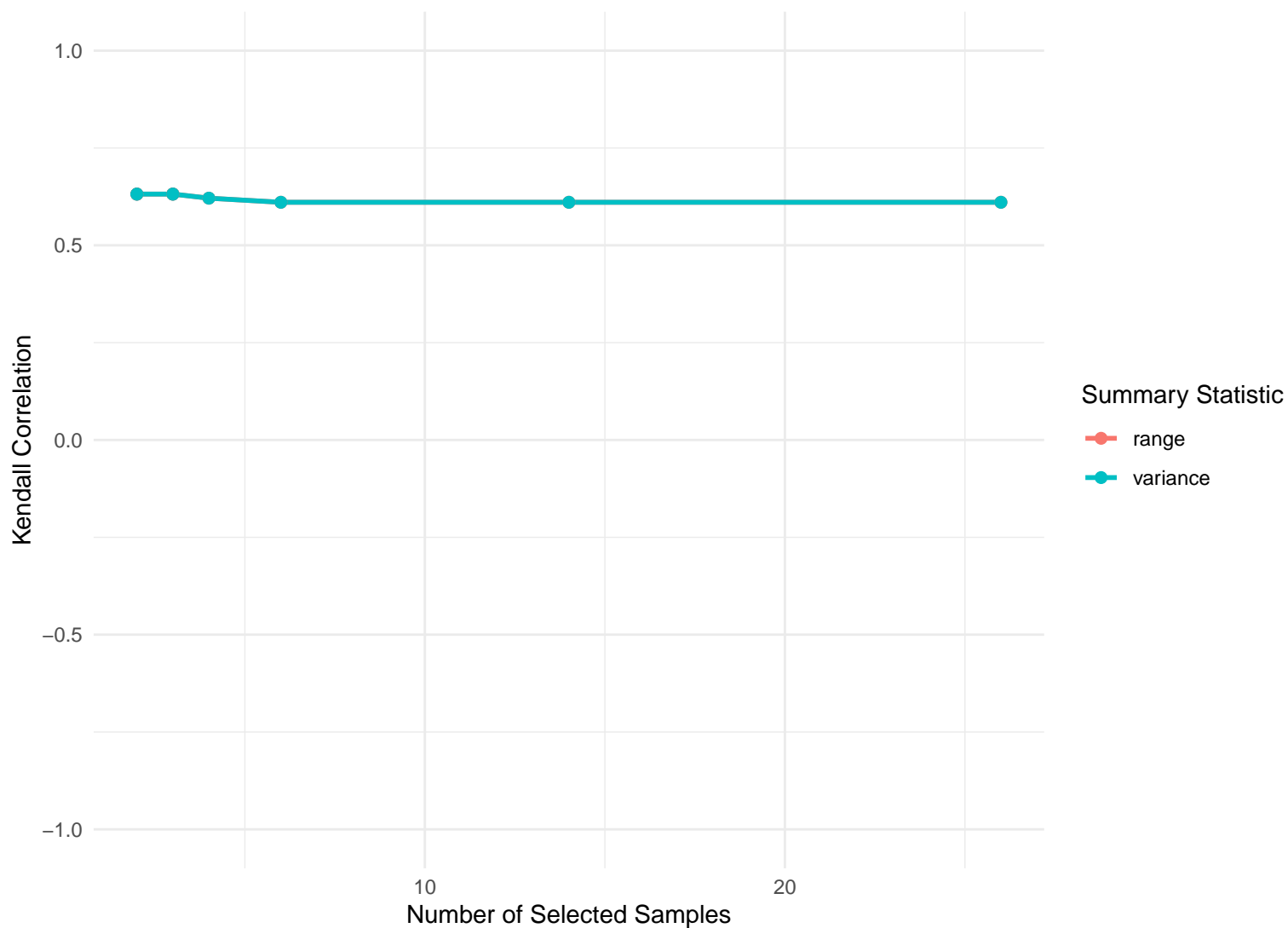
Kendall Correlation

Summary Statistic

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



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



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

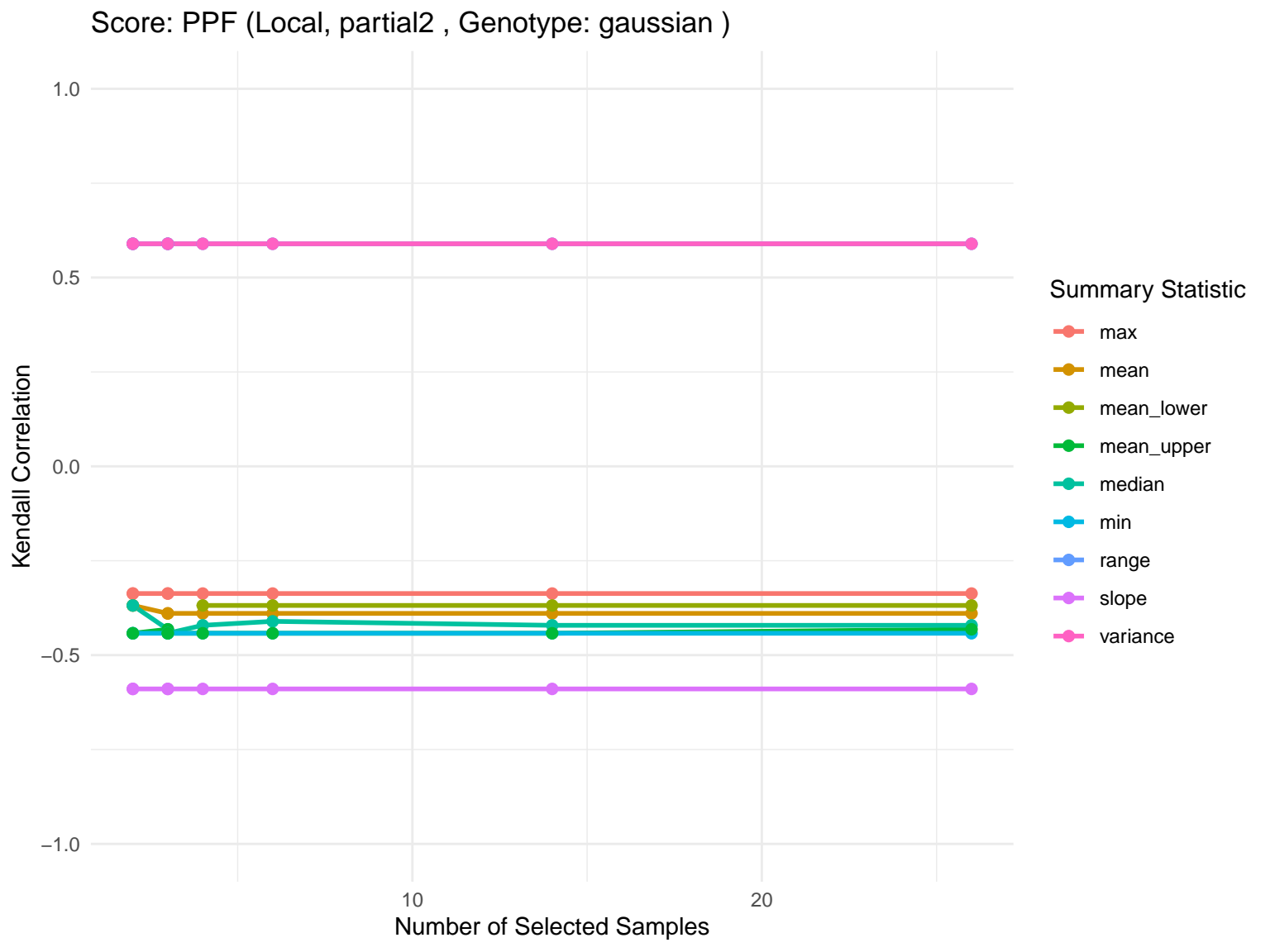
Kendall Correlation

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, partial2 , Genotype: sinusoidal)



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

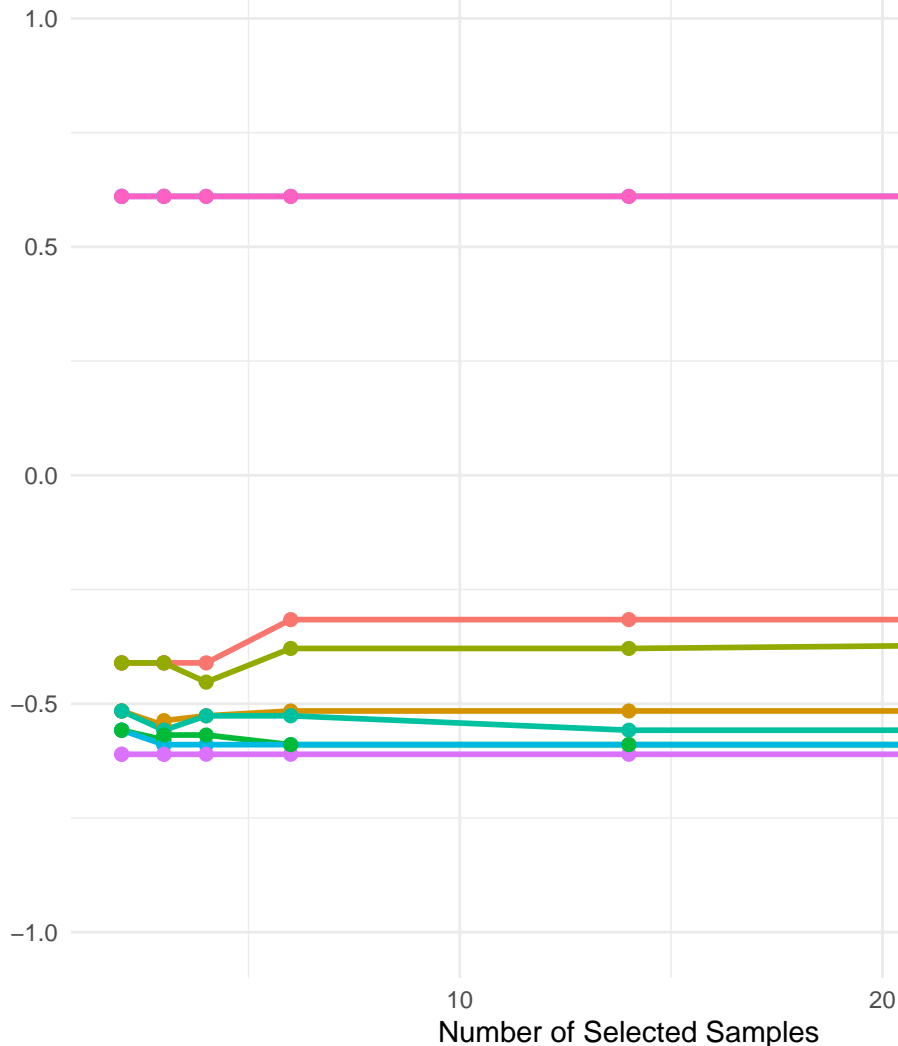
variance

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

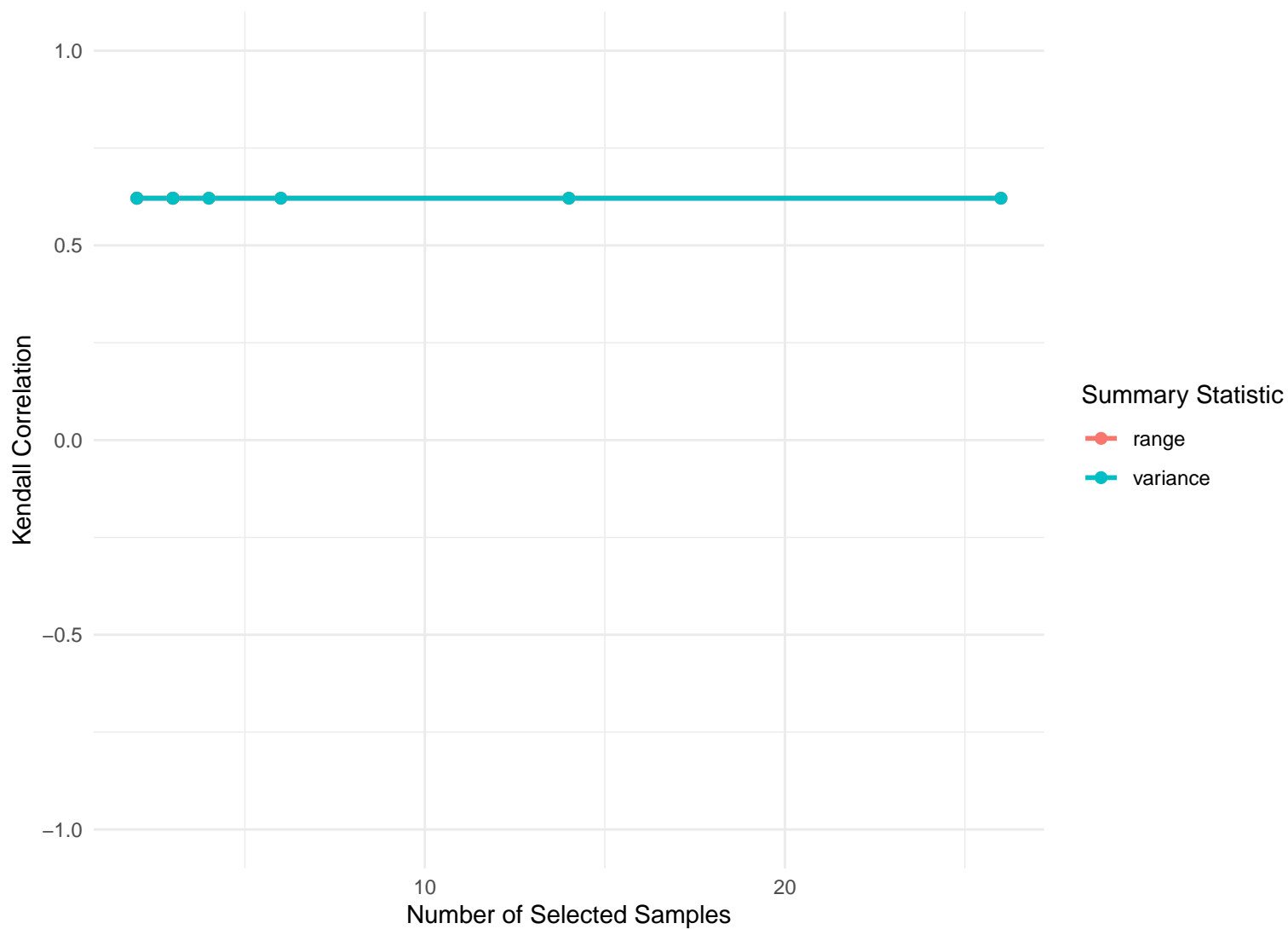
Kendall Correlation

Summary Statistic

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



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

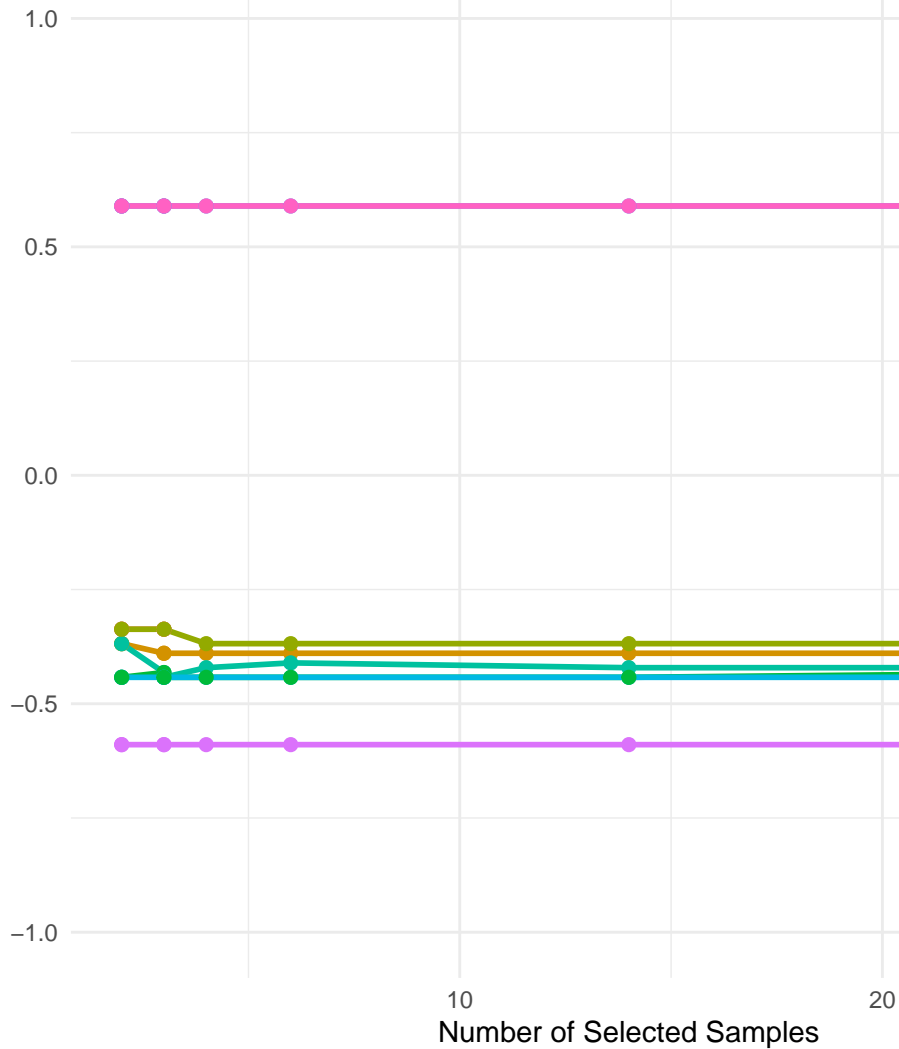


Score: PPI (Local, partial2 , Genotype: gaussian)

Kendall Correlation

Summary Statistic

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

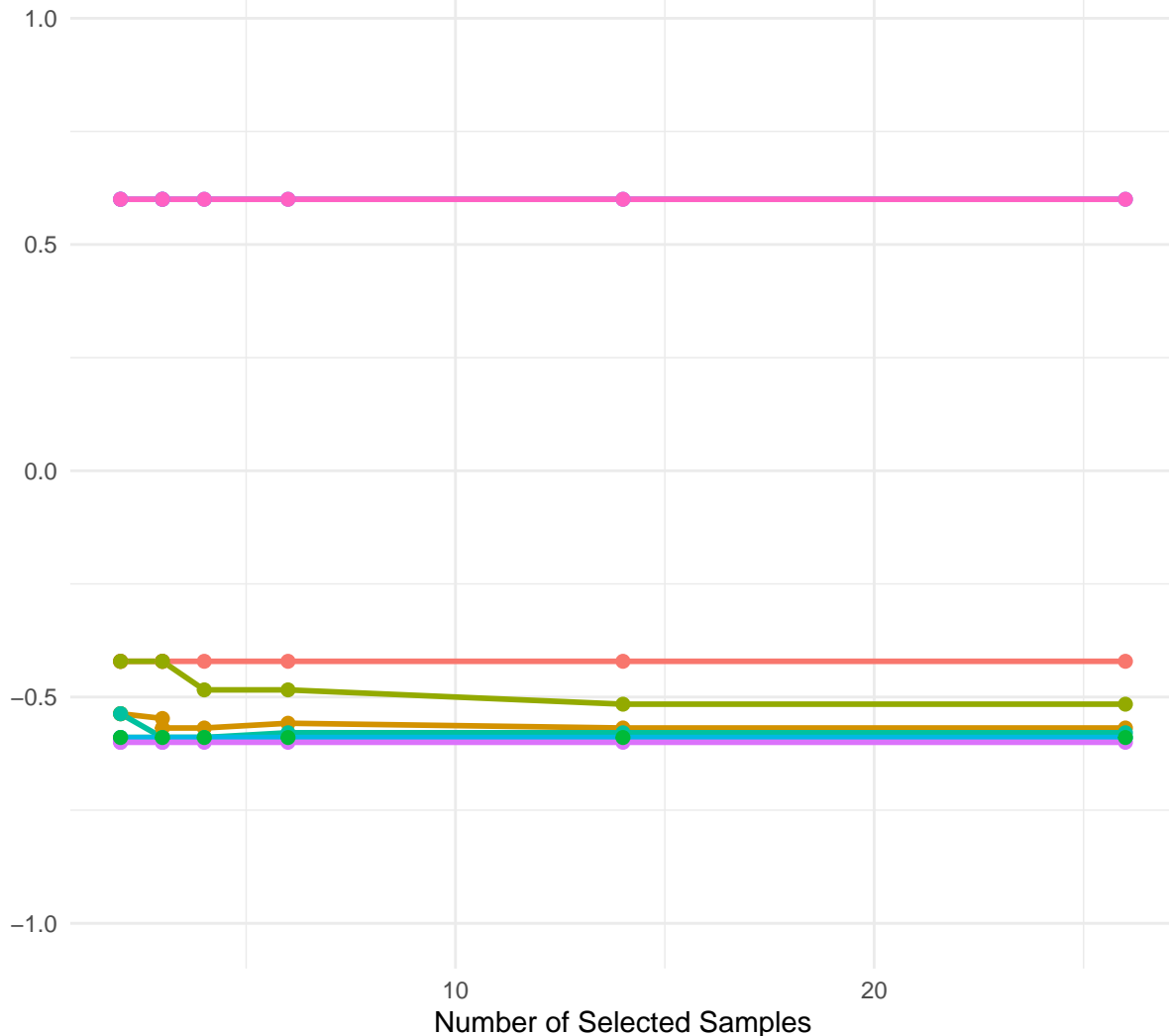


Score: PPI (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

Summary Statistic

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

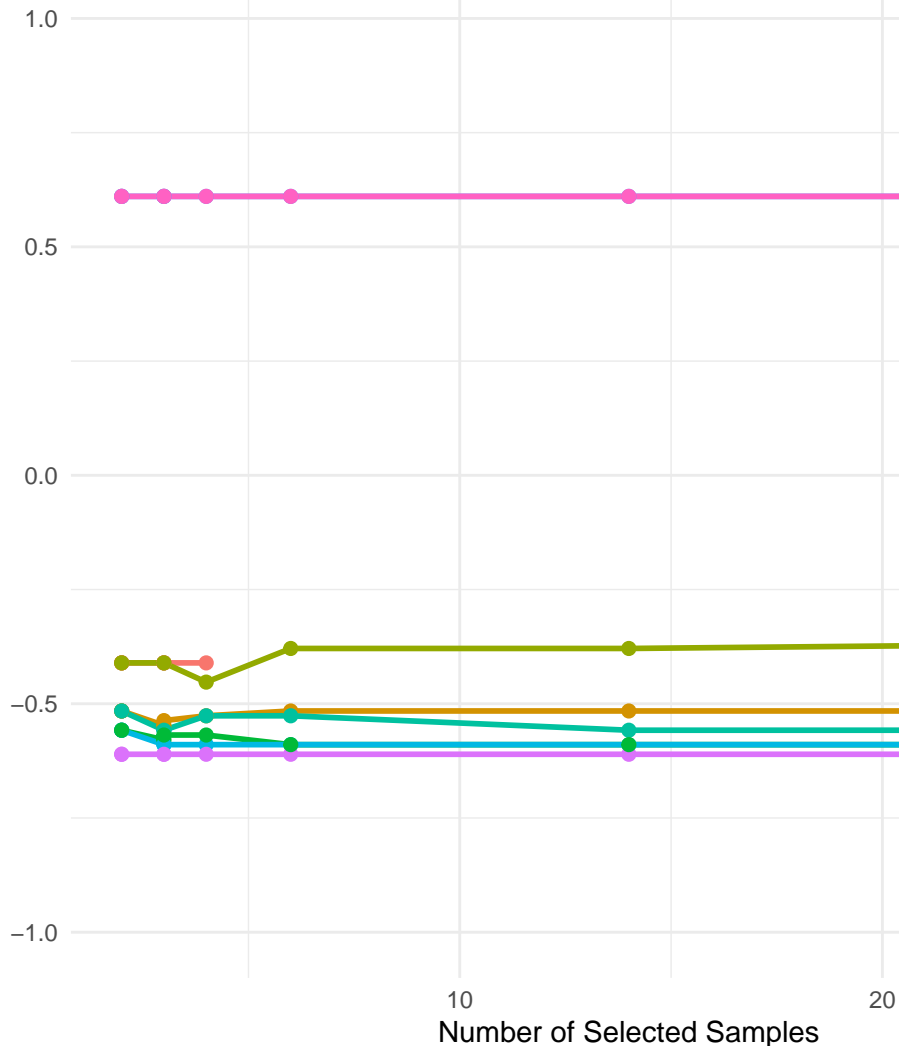


Score: PPI (Local, partial2 , Genotype: wave)

Kendall Correlation

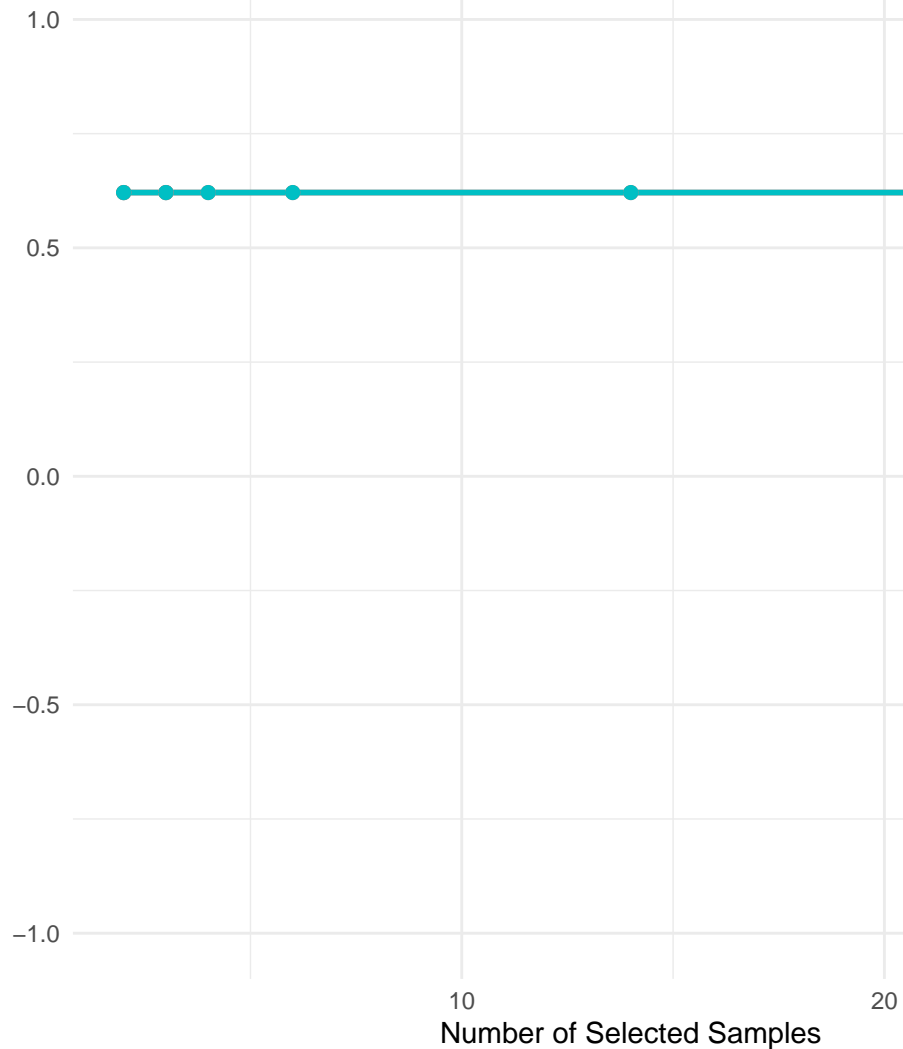
Summary Statistic

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



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

Kendall Correlation



Summary Statistic

- range
- variance

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

Kendall Correlation

Summary Statistic

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

1.0

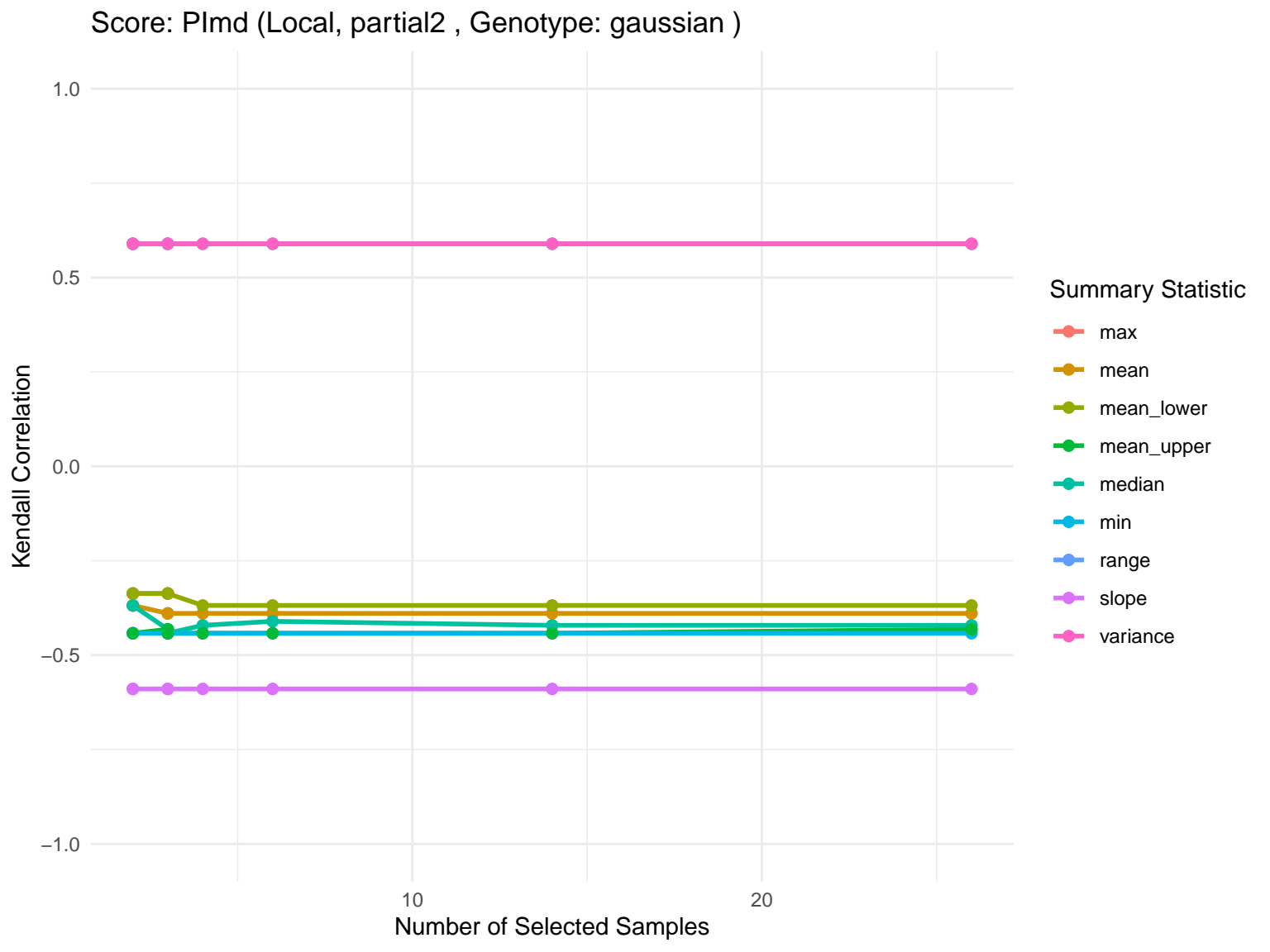
0.5

0.0

-0.5

-1.0

10
20
Number of Selected Samples

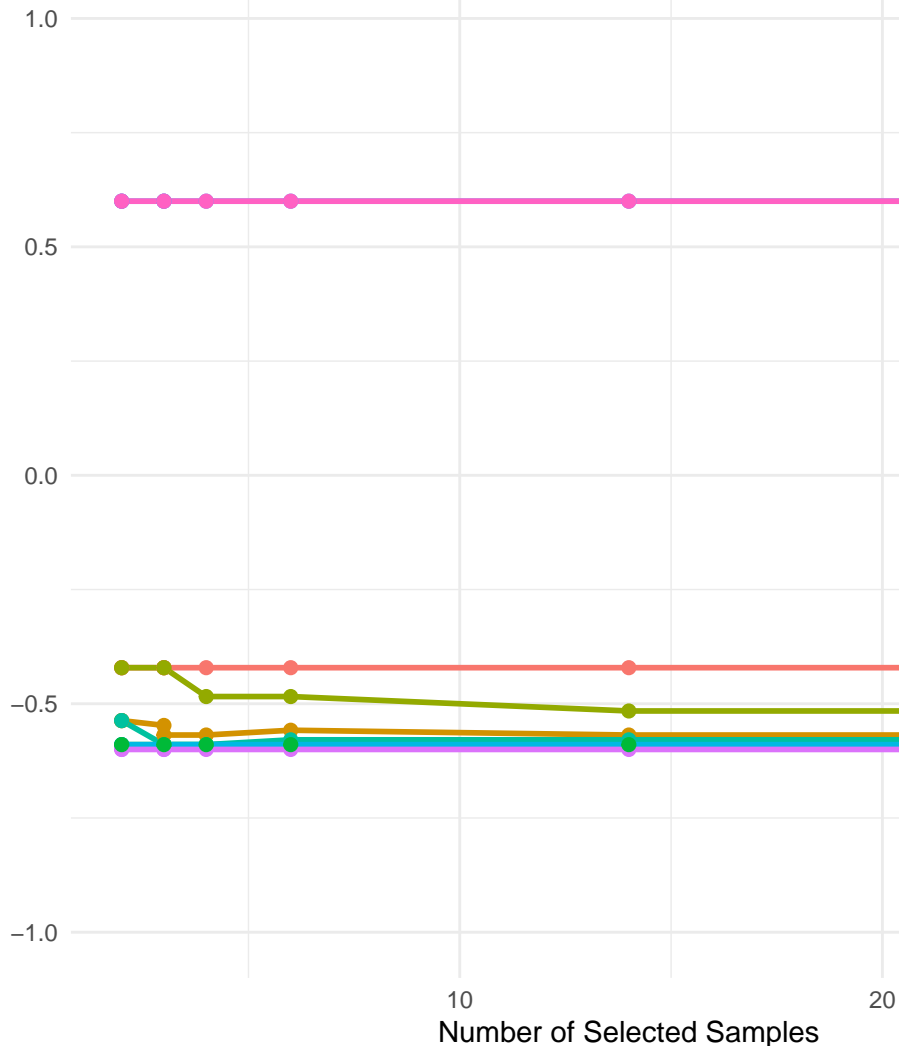


Score: Plmd (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

Summary Statistic

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

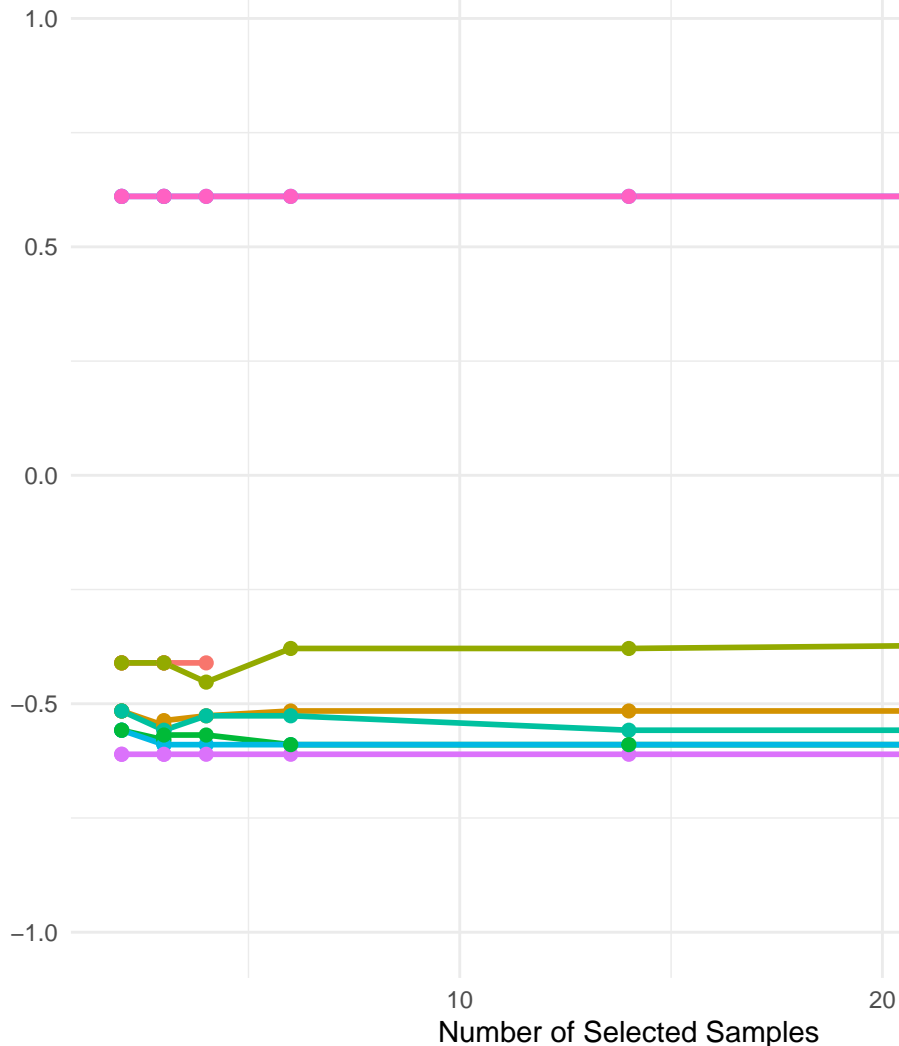


Score: Plmd (Local, partial2 , Genotype: wave)

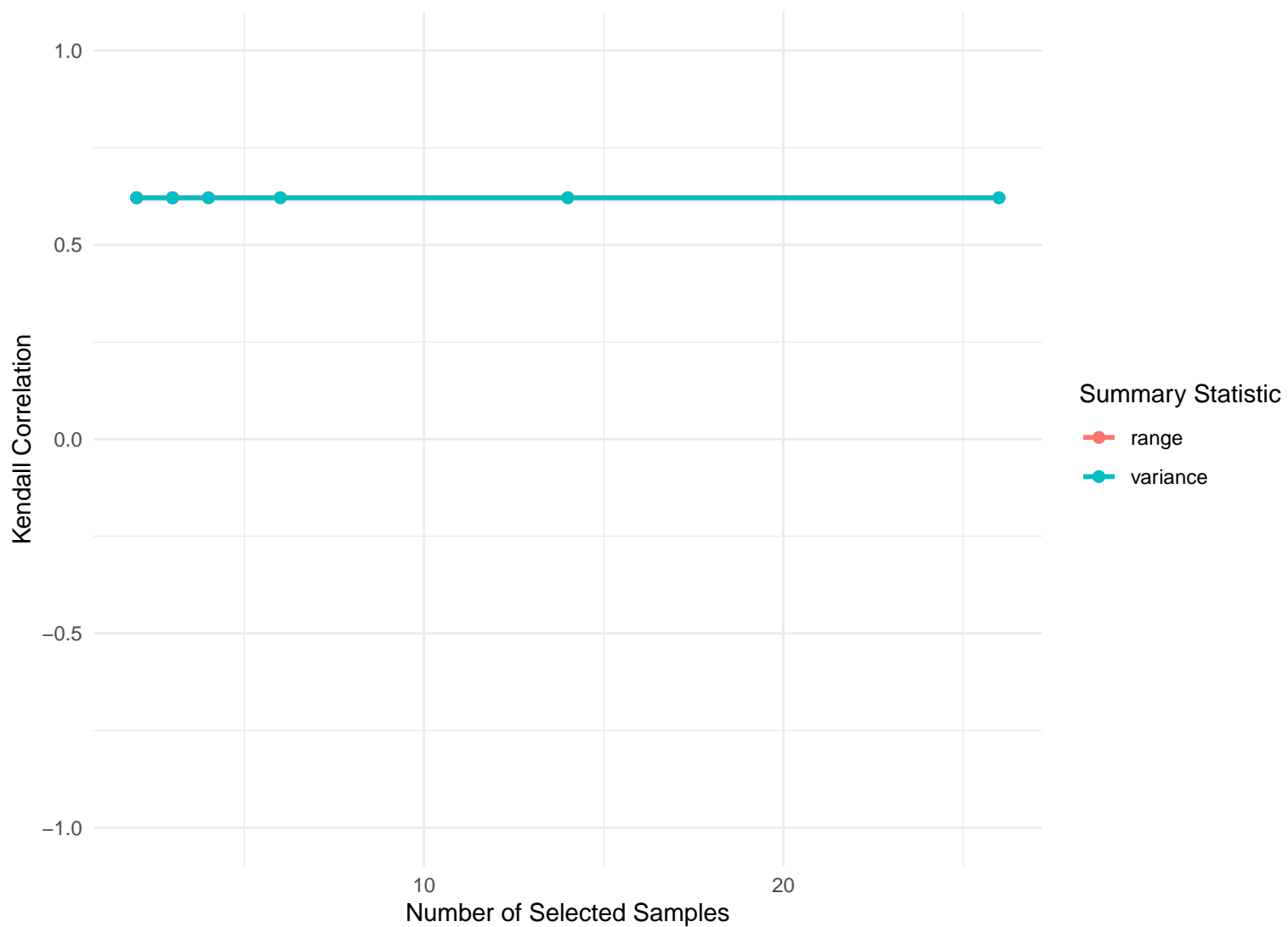
Kendall Correlation

Summary Statistic

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



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



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

Kendall Correlation

Summary Statistic

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

1.0

0.5

0.0

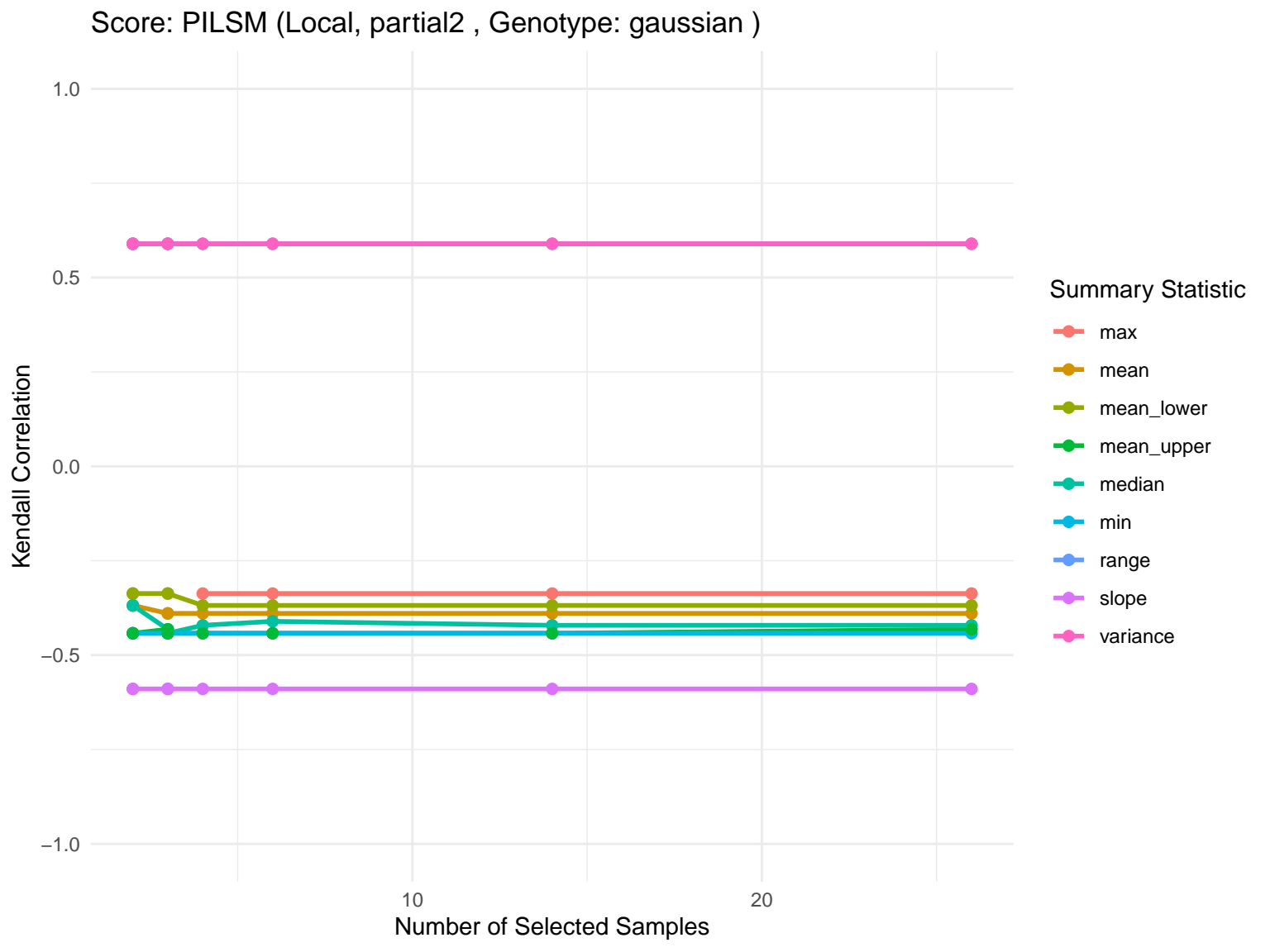
-0.5

-1.0

10

20

Number of Selected Samples



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

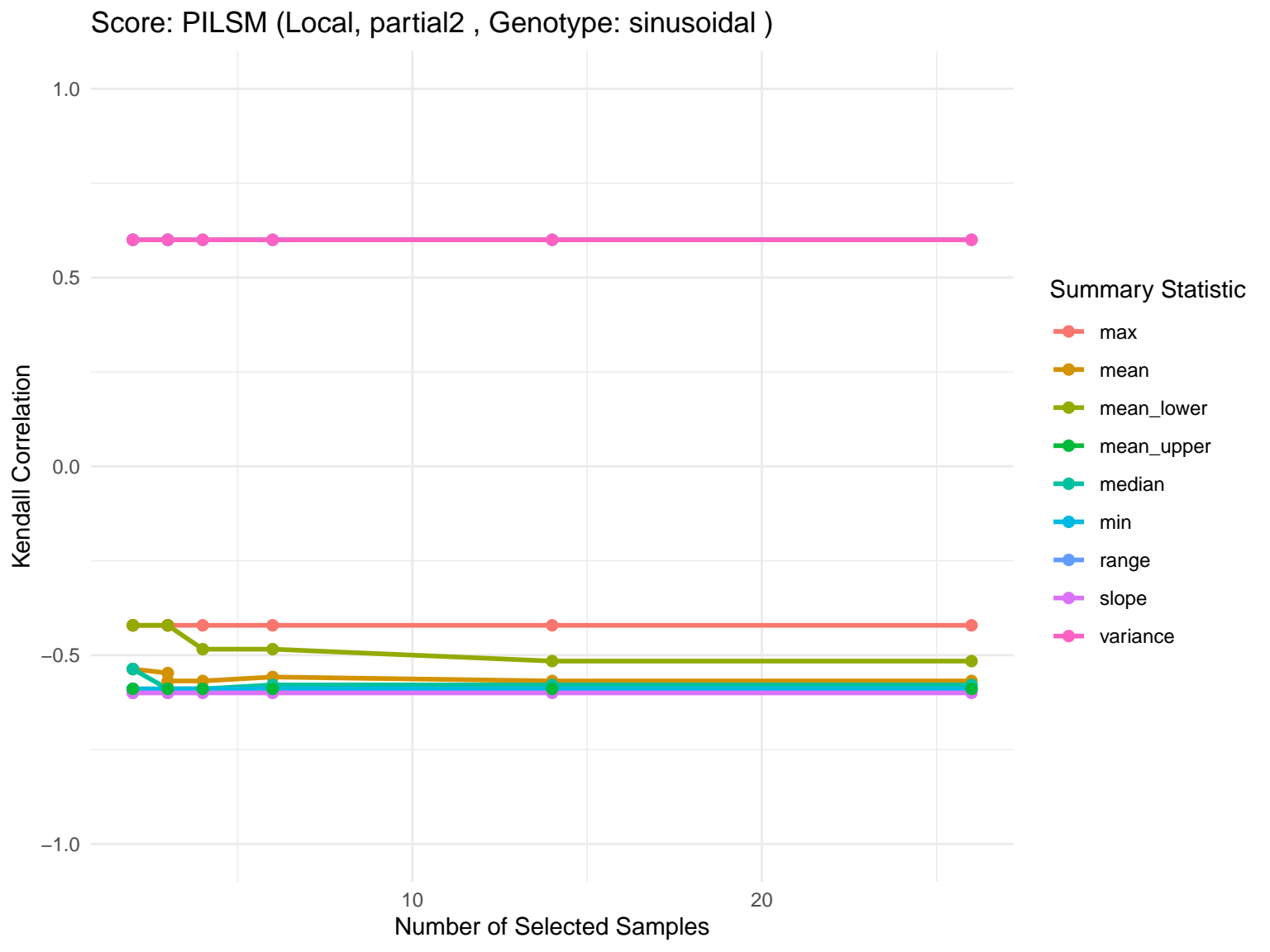
Kendall Correlation

Summary Statistic

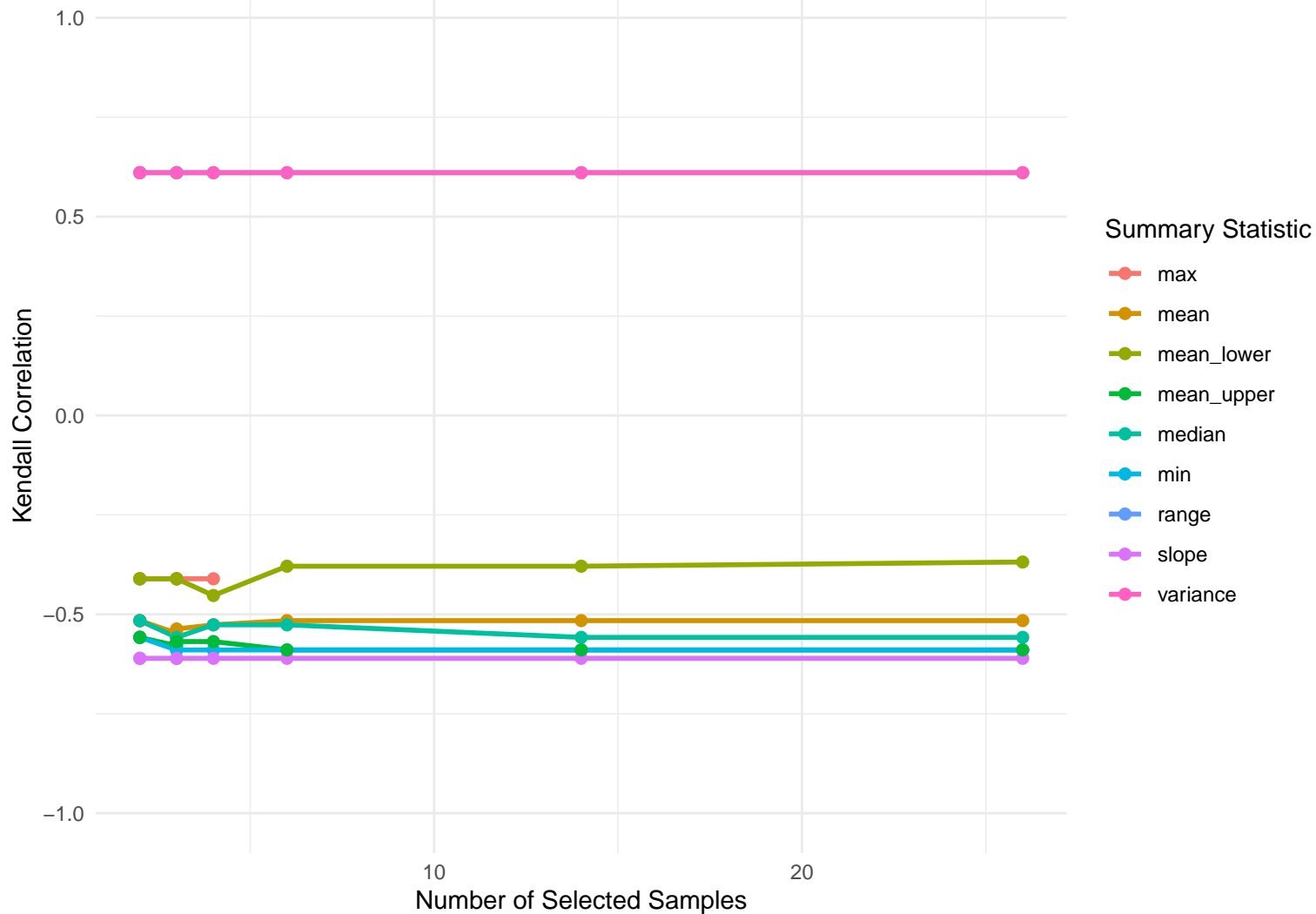
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance

1.0
0.5
0.0
-0.5
-1.0

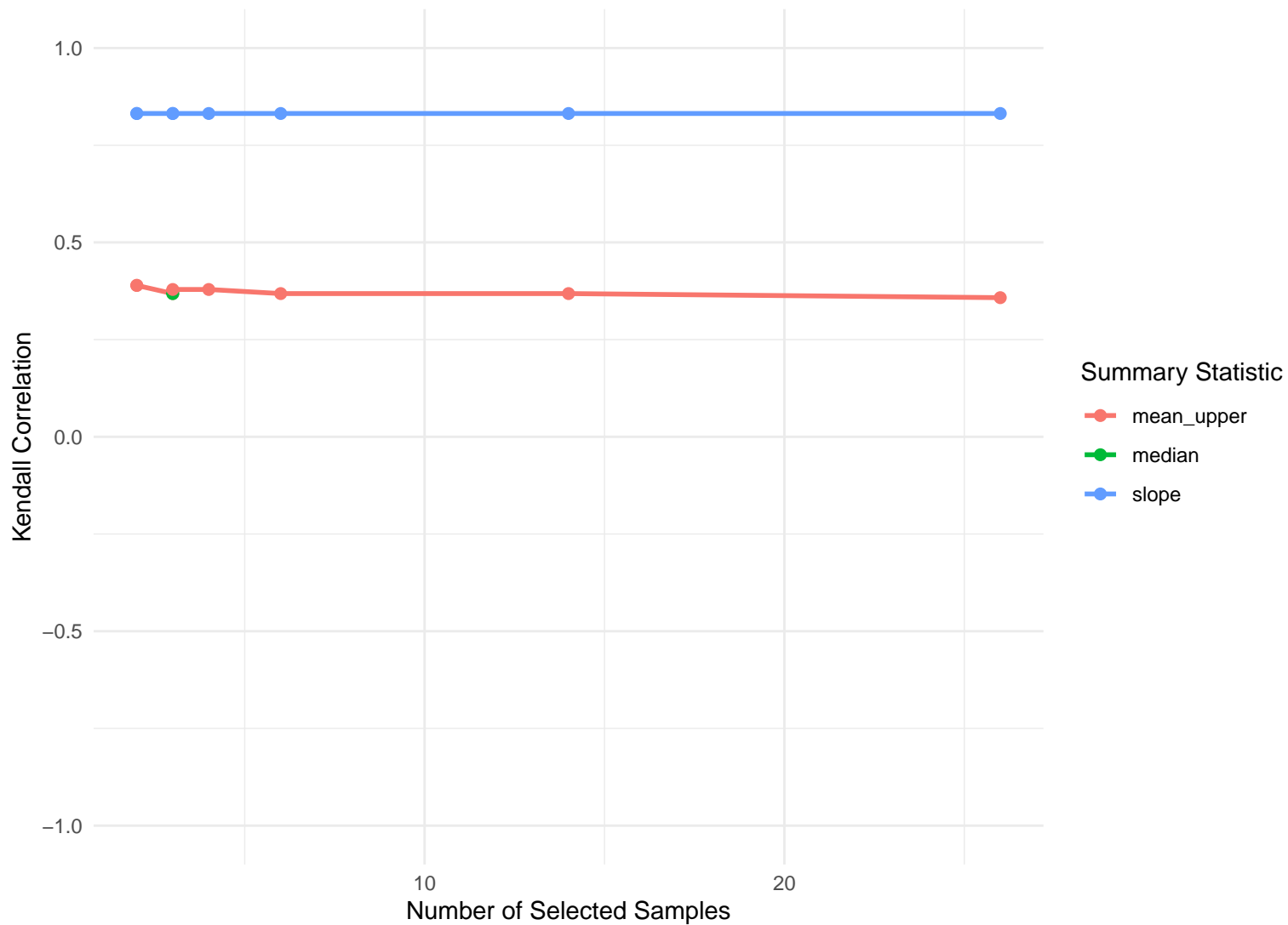
10 20
Number of Selected Samples



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

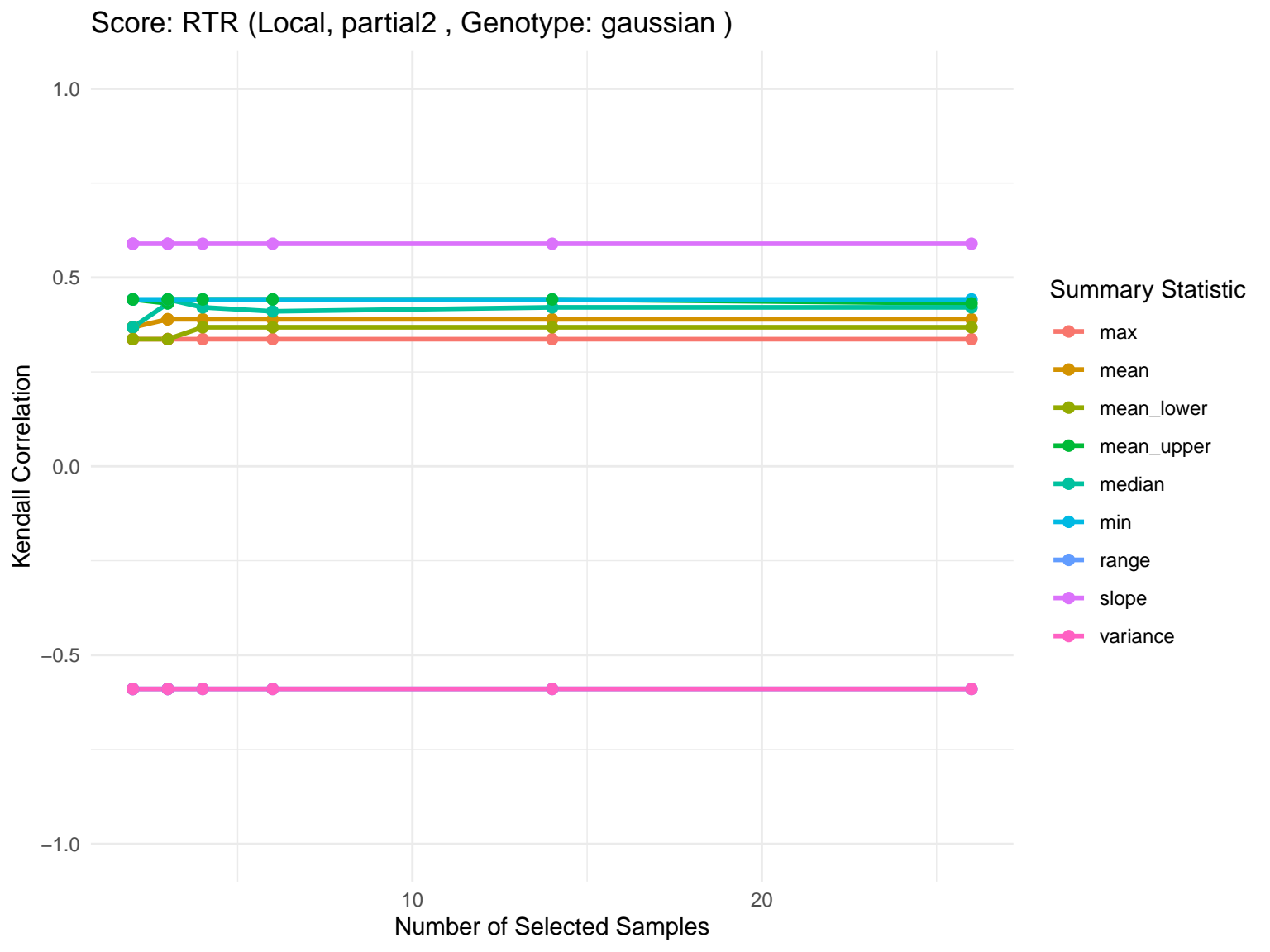
10

20

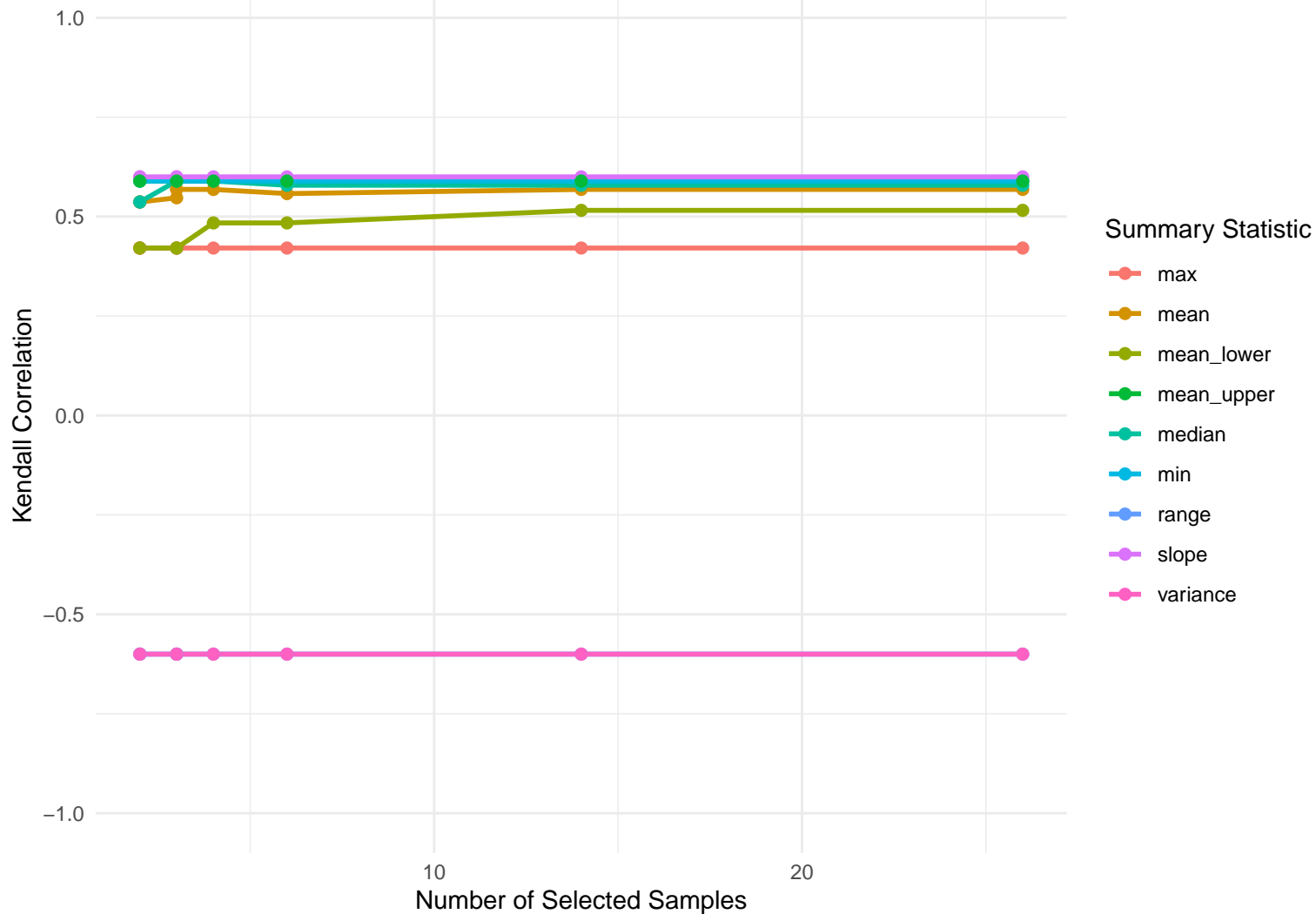
Number of Selected Samples

Summary Statistic

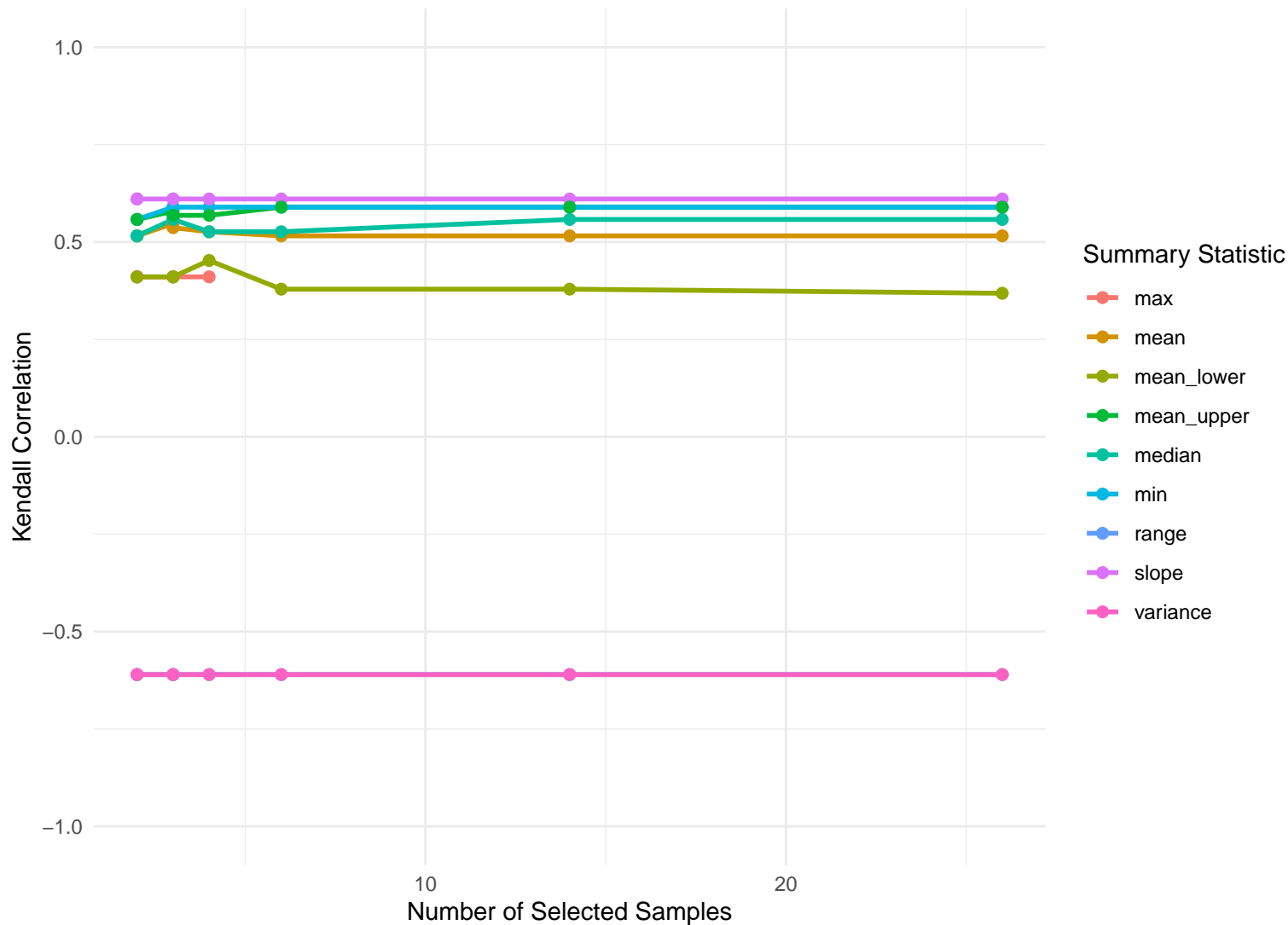
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



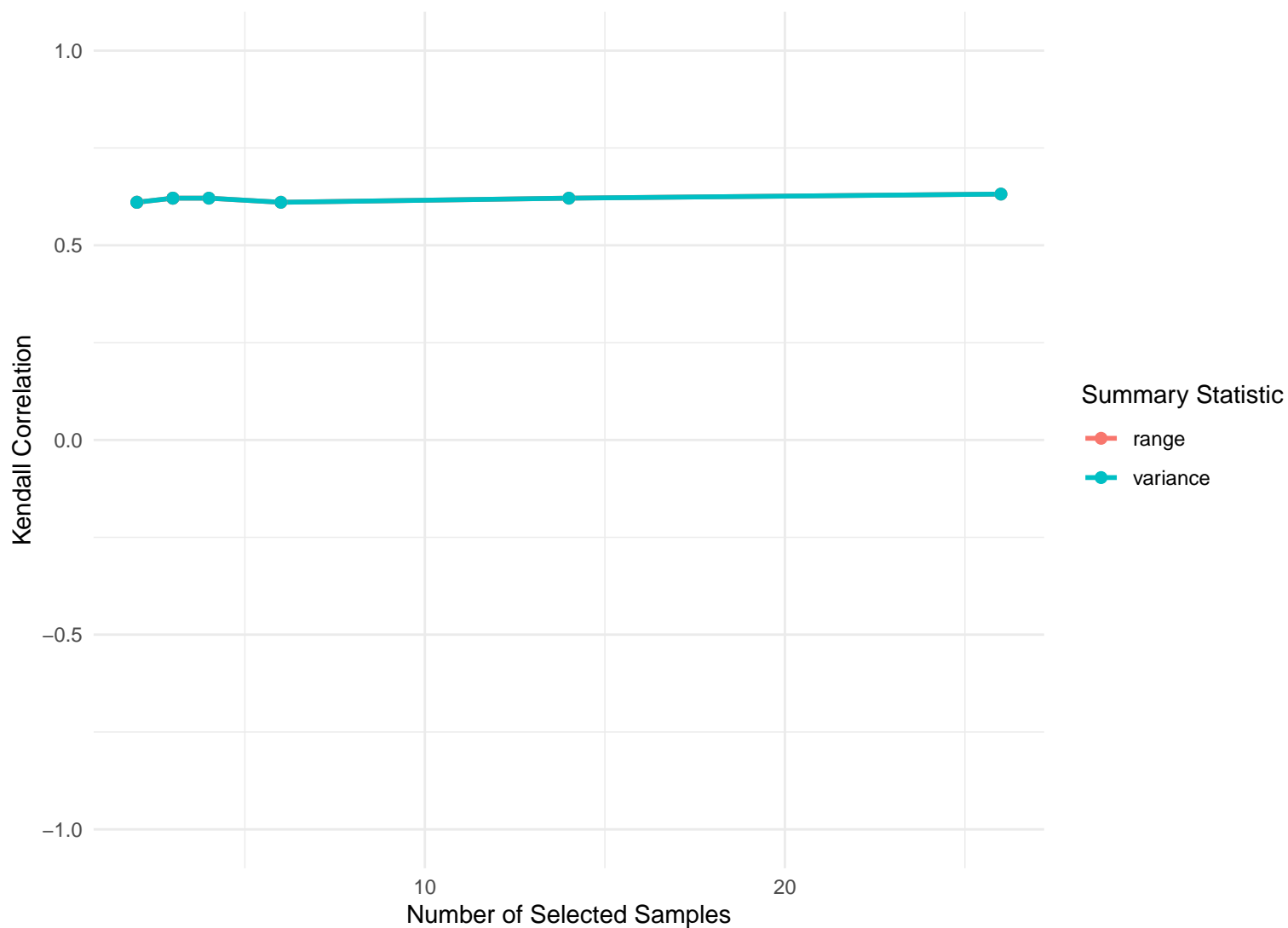
Score: RTR (Local, partial2 , Genotype: sinusoidal)



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



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

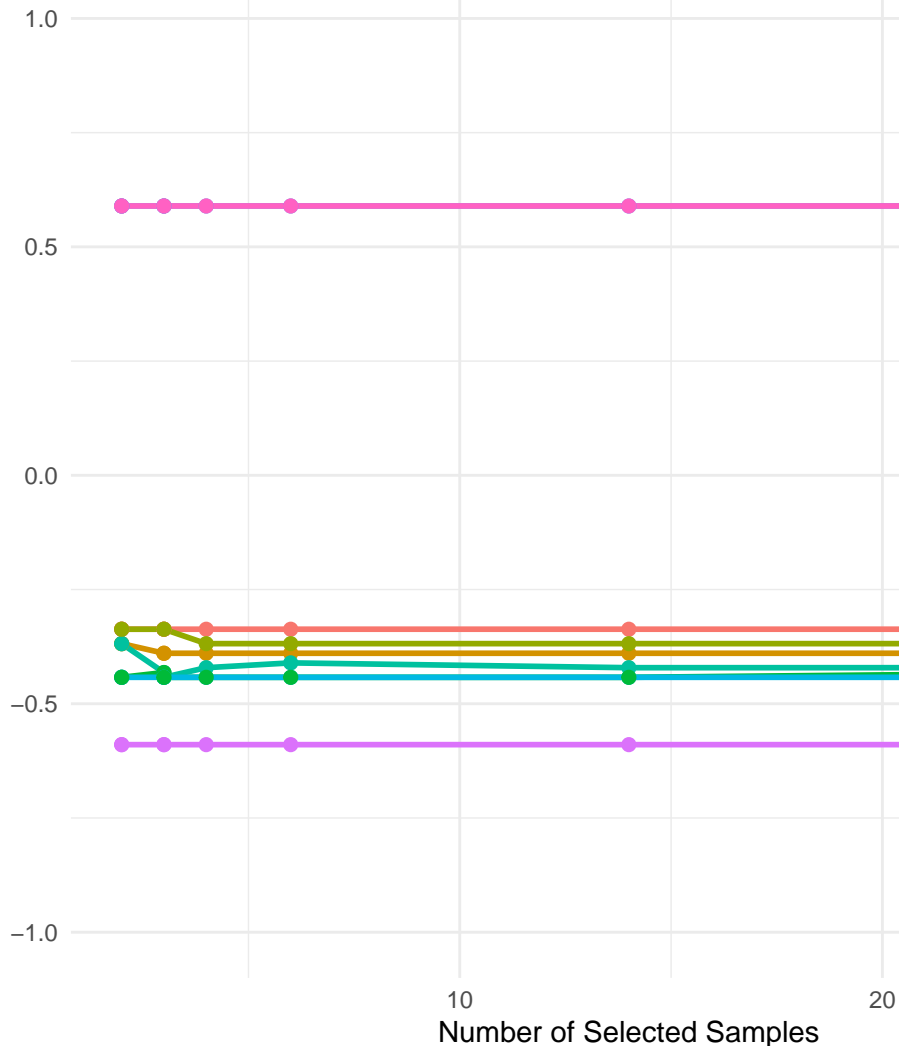


Score: PIR (Local, partial2 , Genotype: gaussian)

Kendall Correlation

Summary Statistic

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

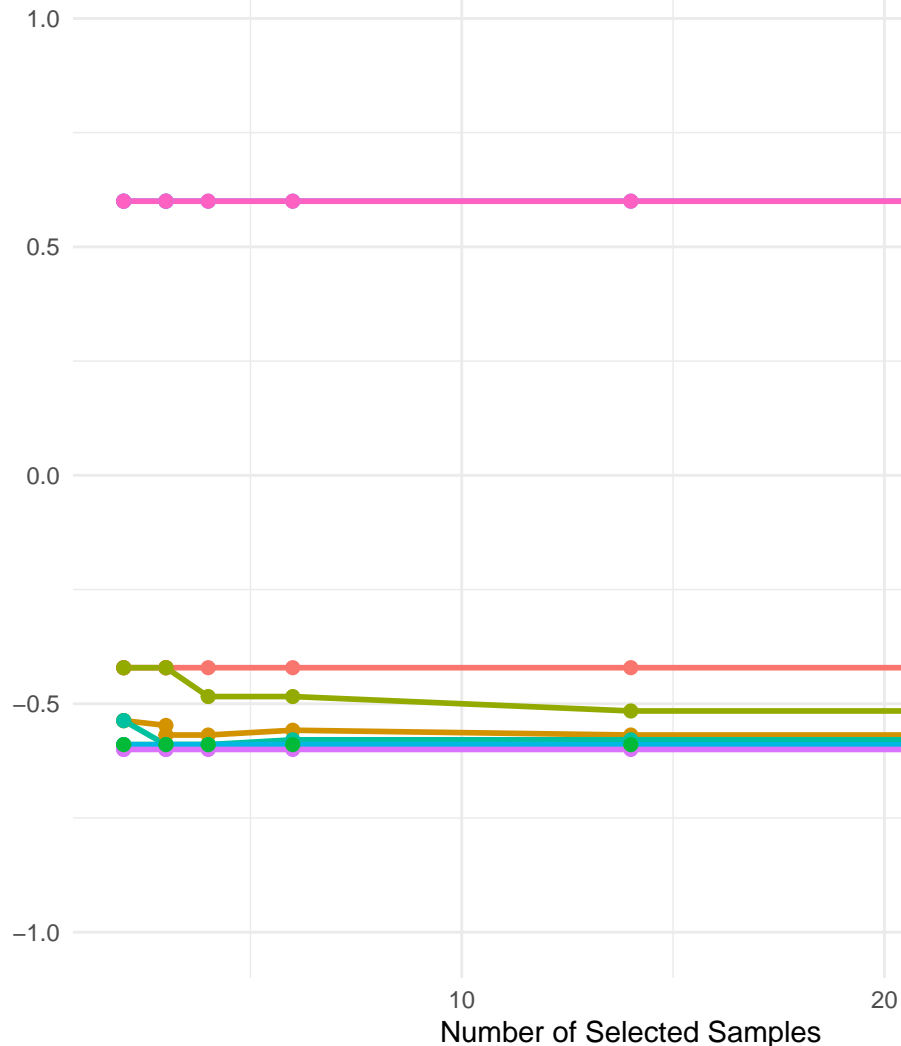


Score: PIR (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

Summary Statistic

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



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

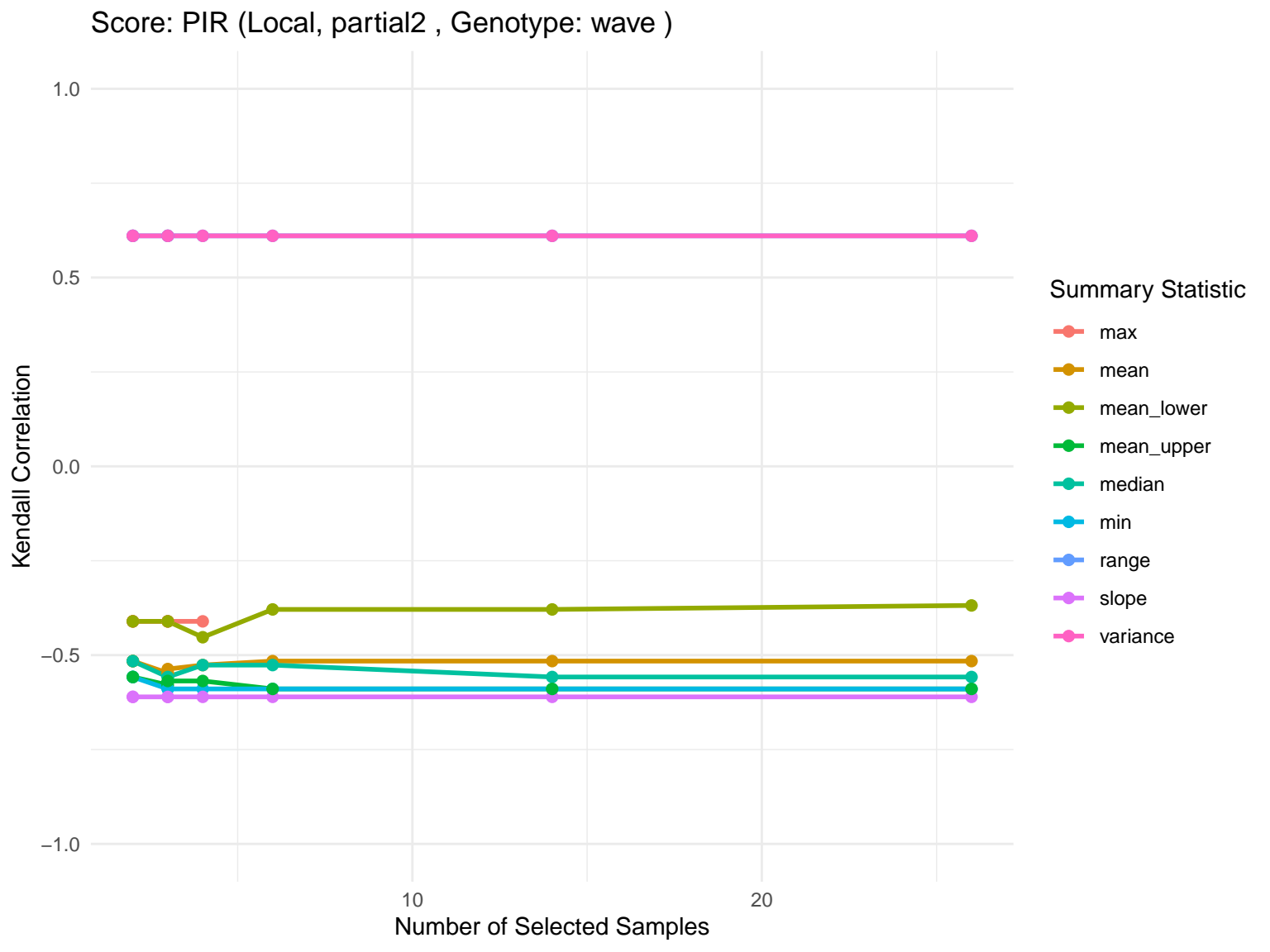
Kendall Correlation

Summary Statistic

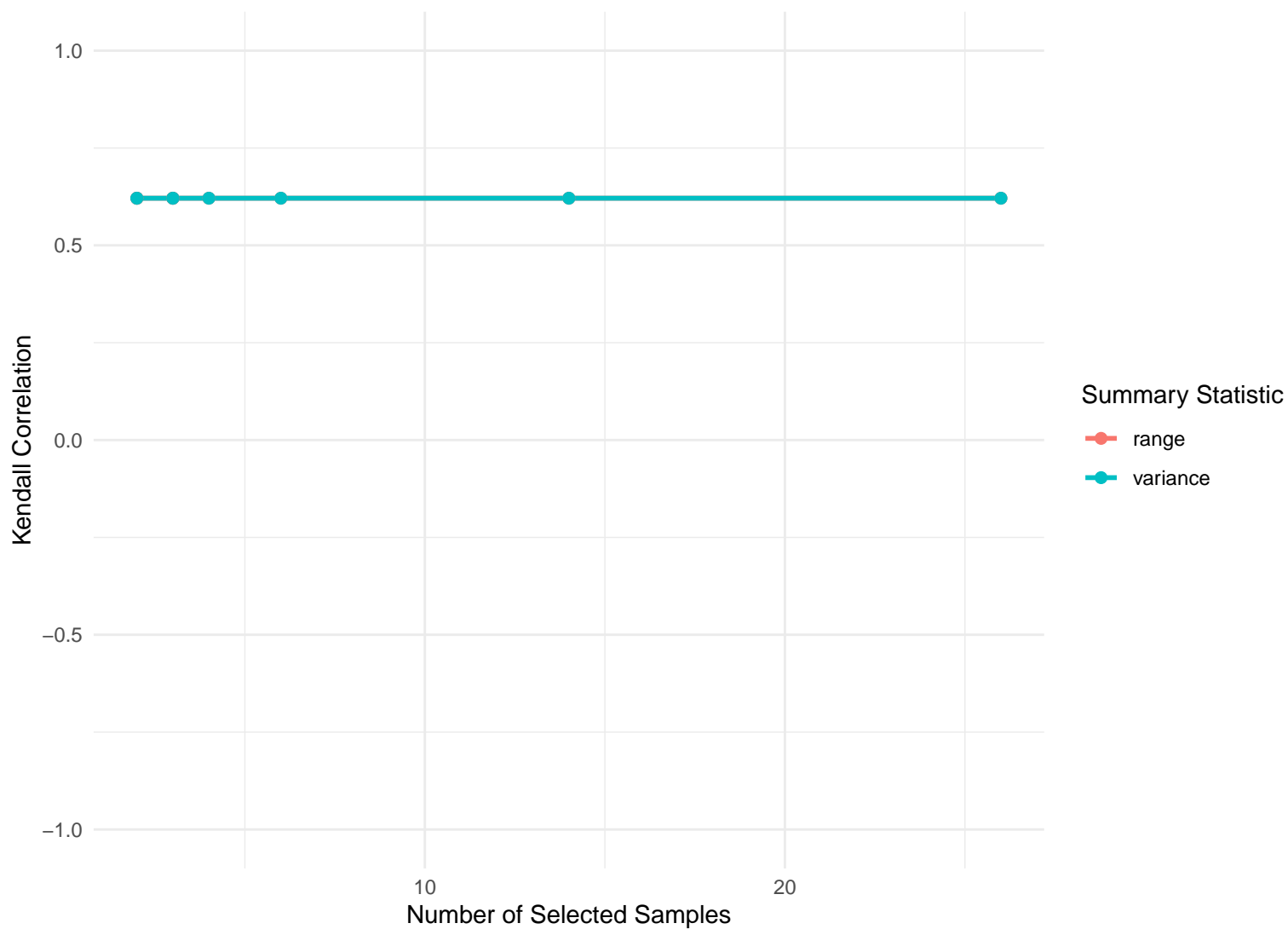
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance

1.0
0.5
0.0
-0.5
-1.0

10 20
Number of Selected Samples



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



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

Kendall Correlation

Summary Statistic

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

1.0

0.5

0.0

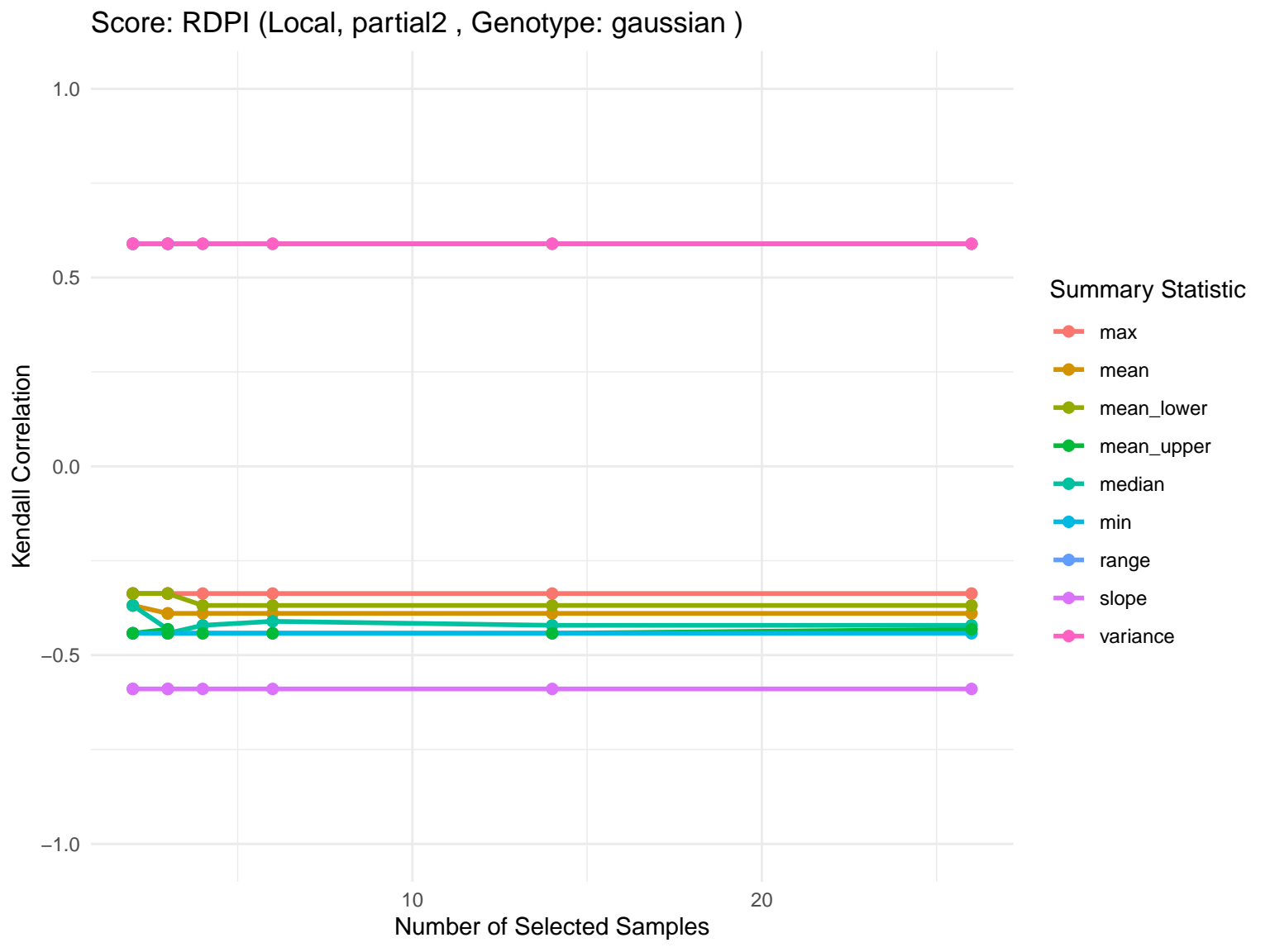
-0.5

-1.0

10

20

Number of Selected Samples



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

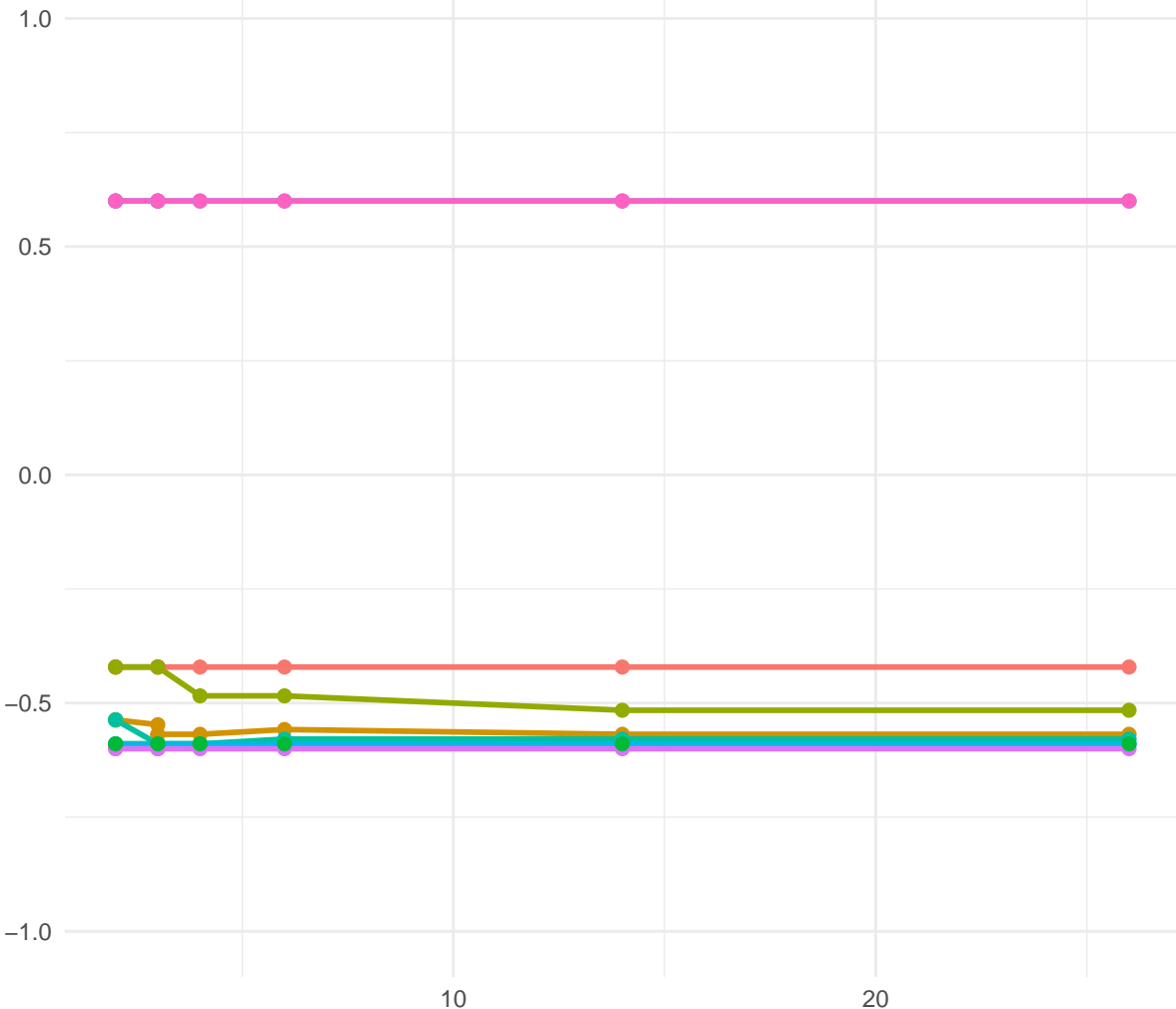
Kendall Correlation

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

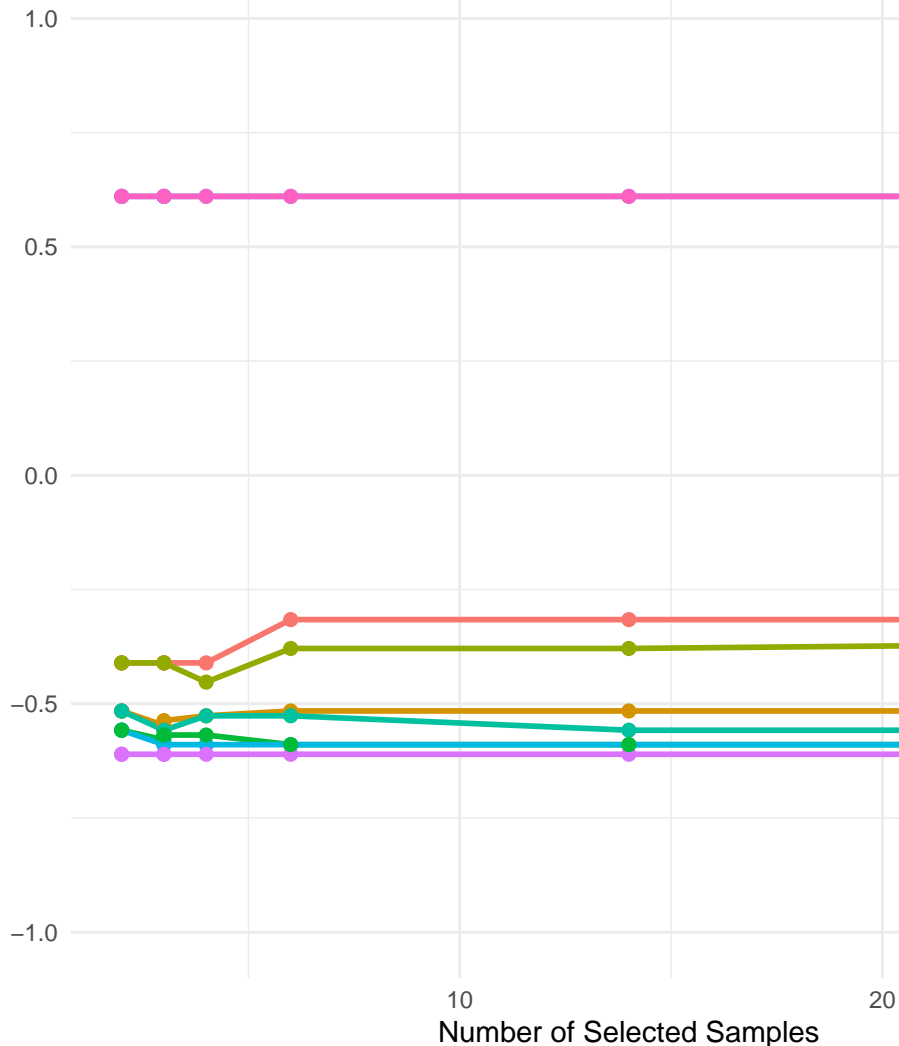


Score: RDPI (Local, partial2 , Genotype: wave)

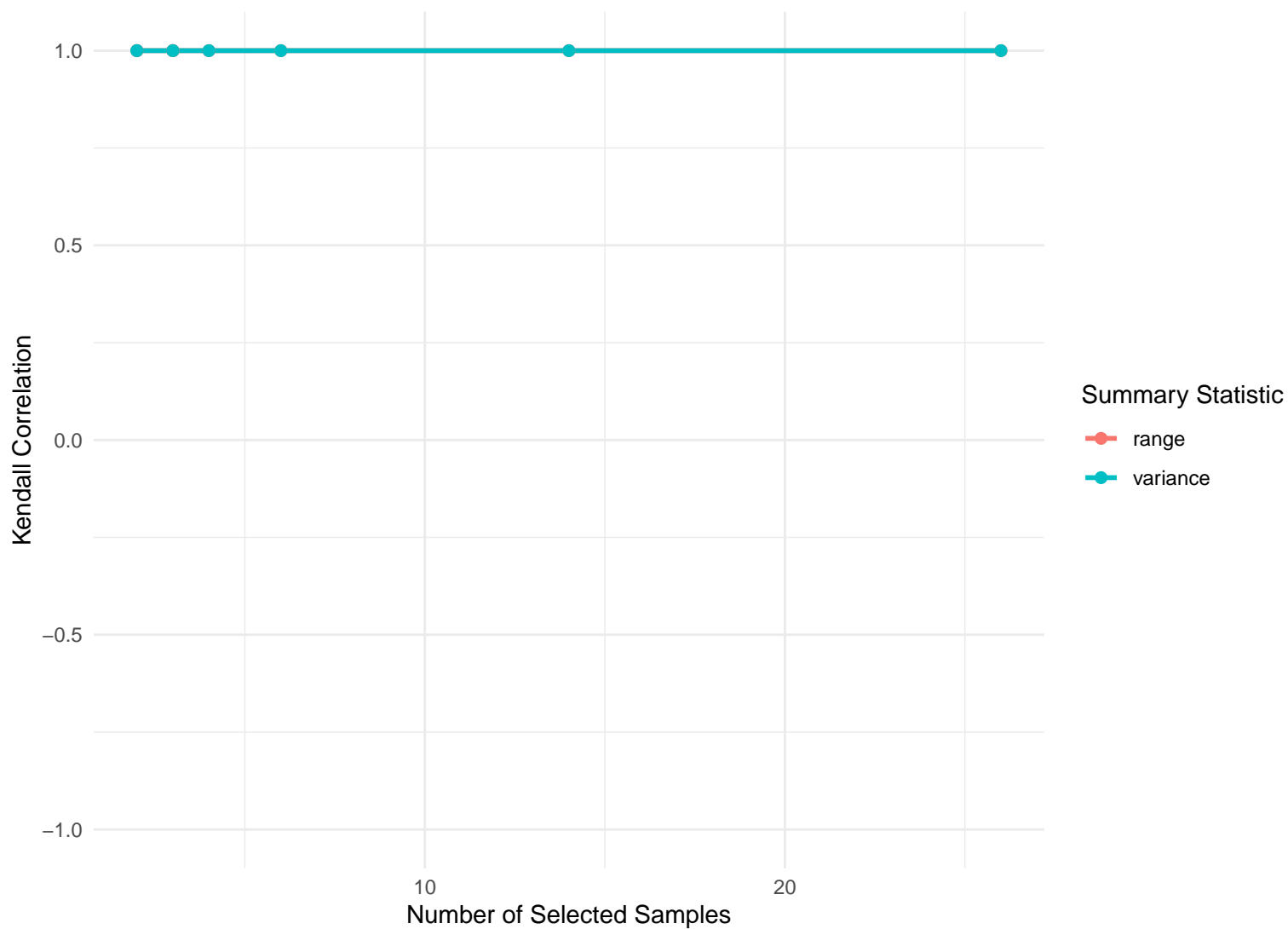
Kendall Correlation

Summary Statistic

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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

10

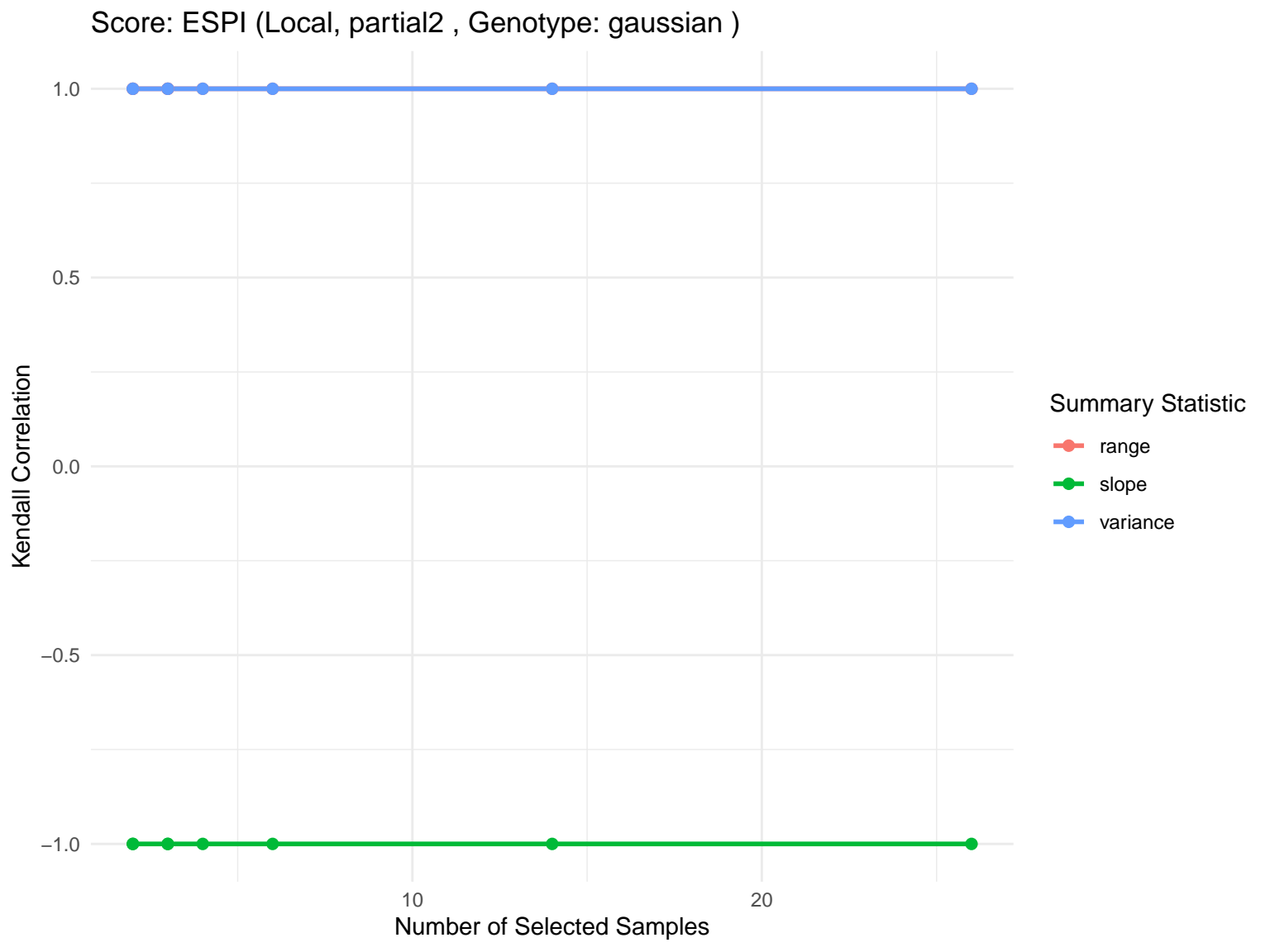
20

Summary Statistic

range

slope

variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

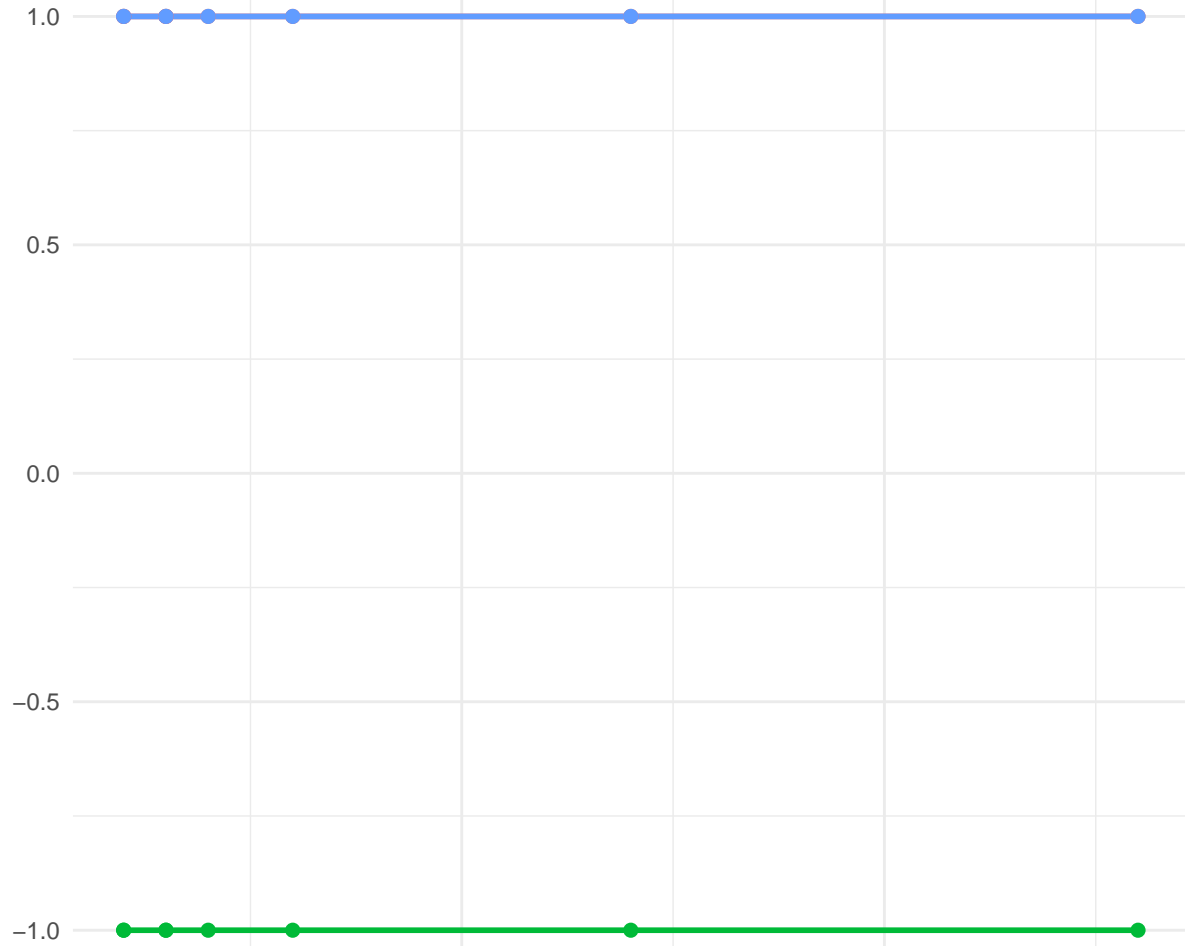
Number of Selected Samples

10

20

Summary Statistic

- range
- slope
- variance

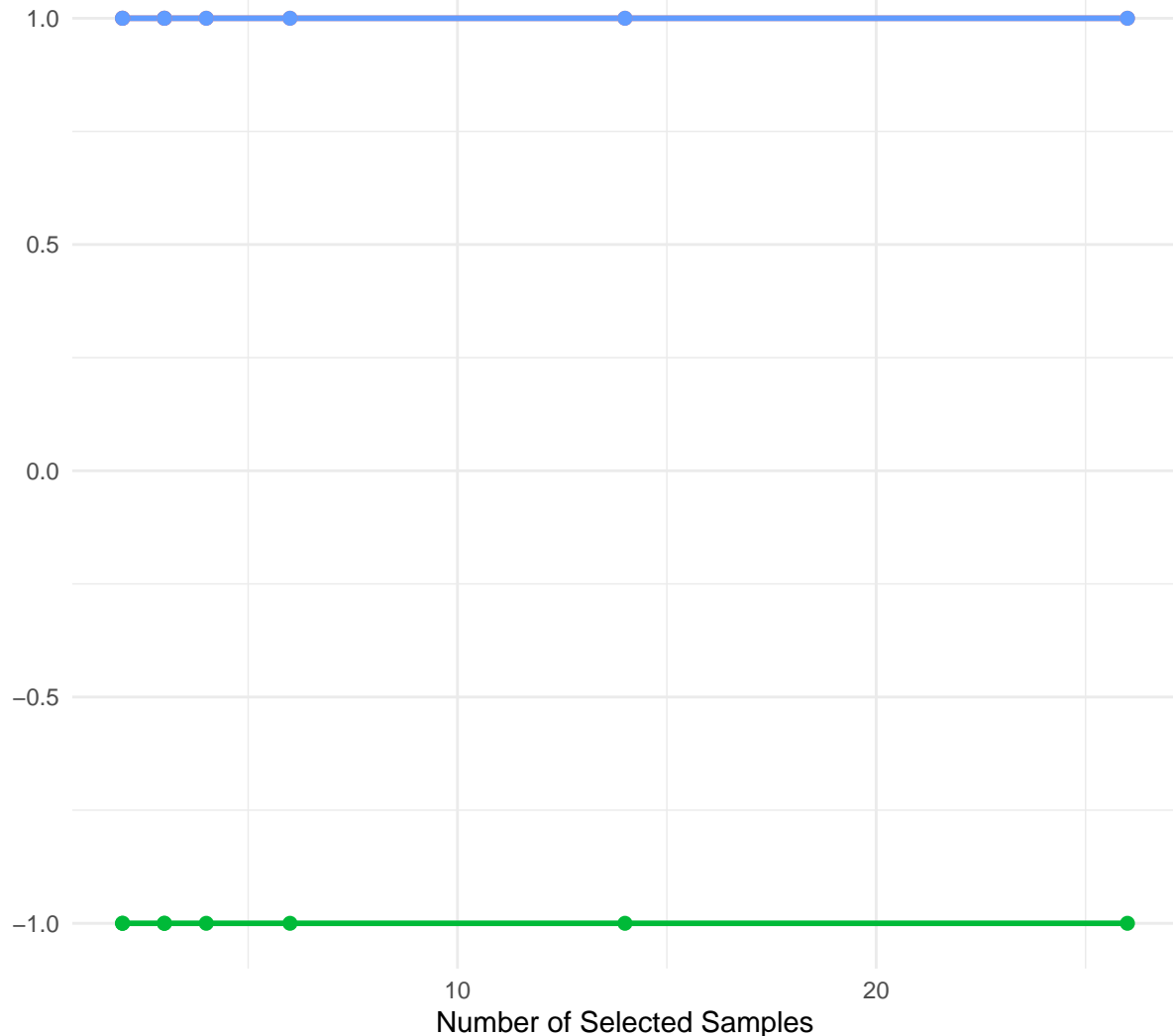


Score: ESPI (Local, partial2 , Genotype: wave)

Kendall Correlation

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

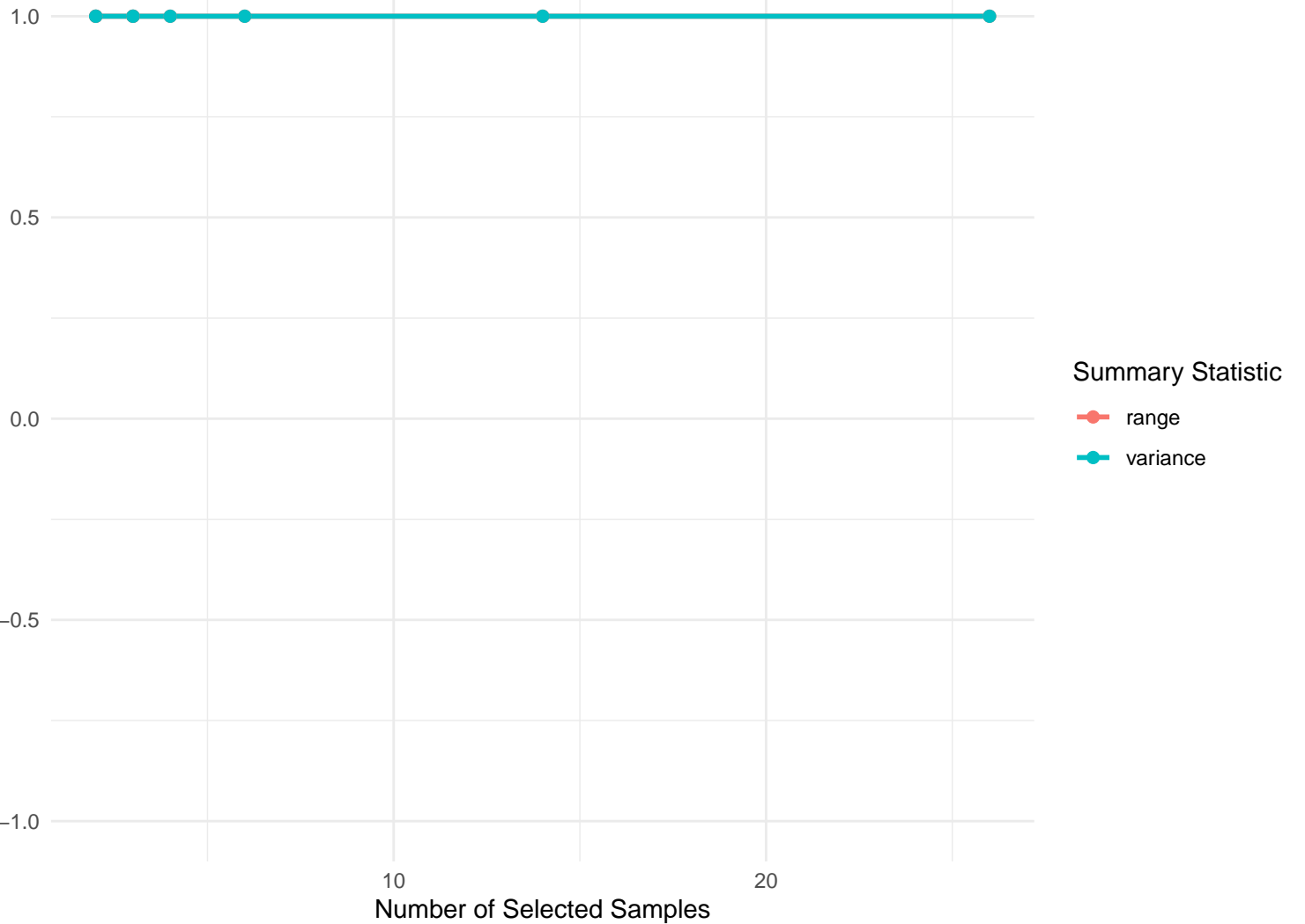
20

Number of Selected Samples

Summary Statistic

range

variance



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

Kendall Correlation

Summary Statistic

- range
- slope
- variance

1.0
0.5
0.0
-0.5
-1.0

Number of Selected Samples

10

20

1.0
0.5
0.0
-0.5
-1.0

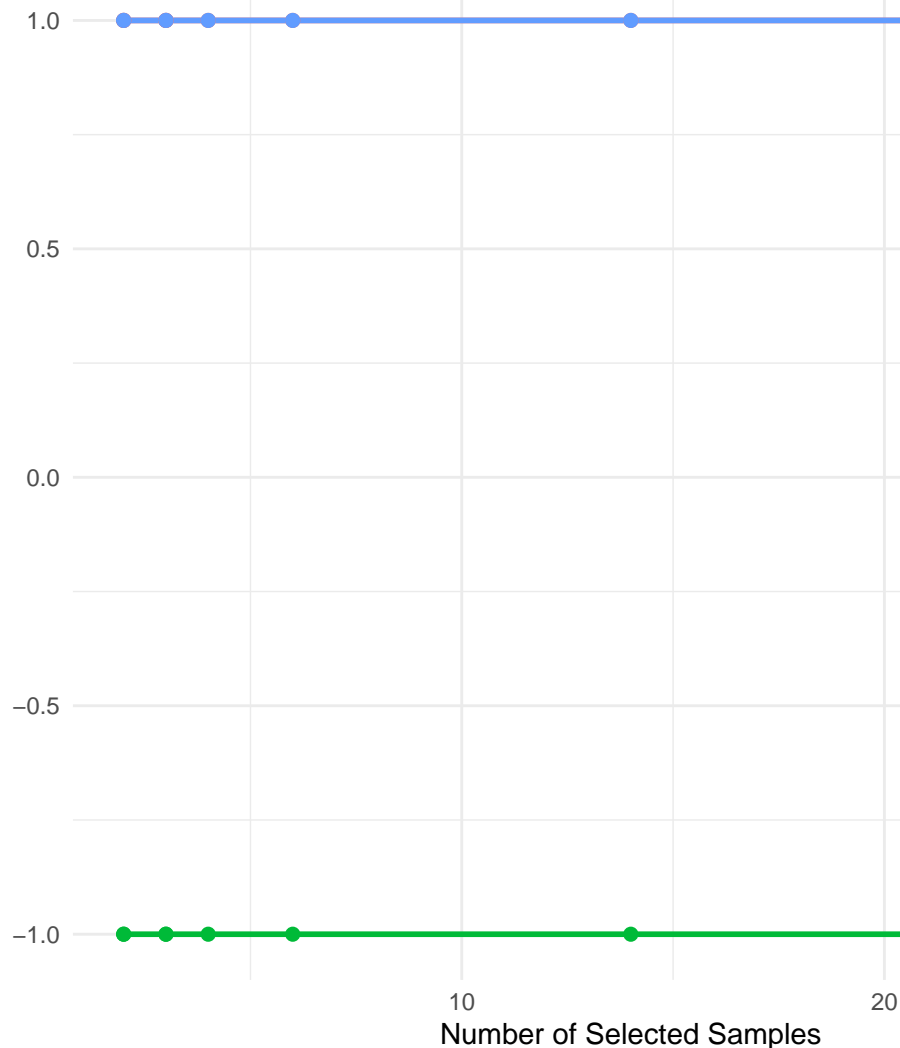


Score: ESPIID (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

Summary Statistic

- range
- slope
- variance

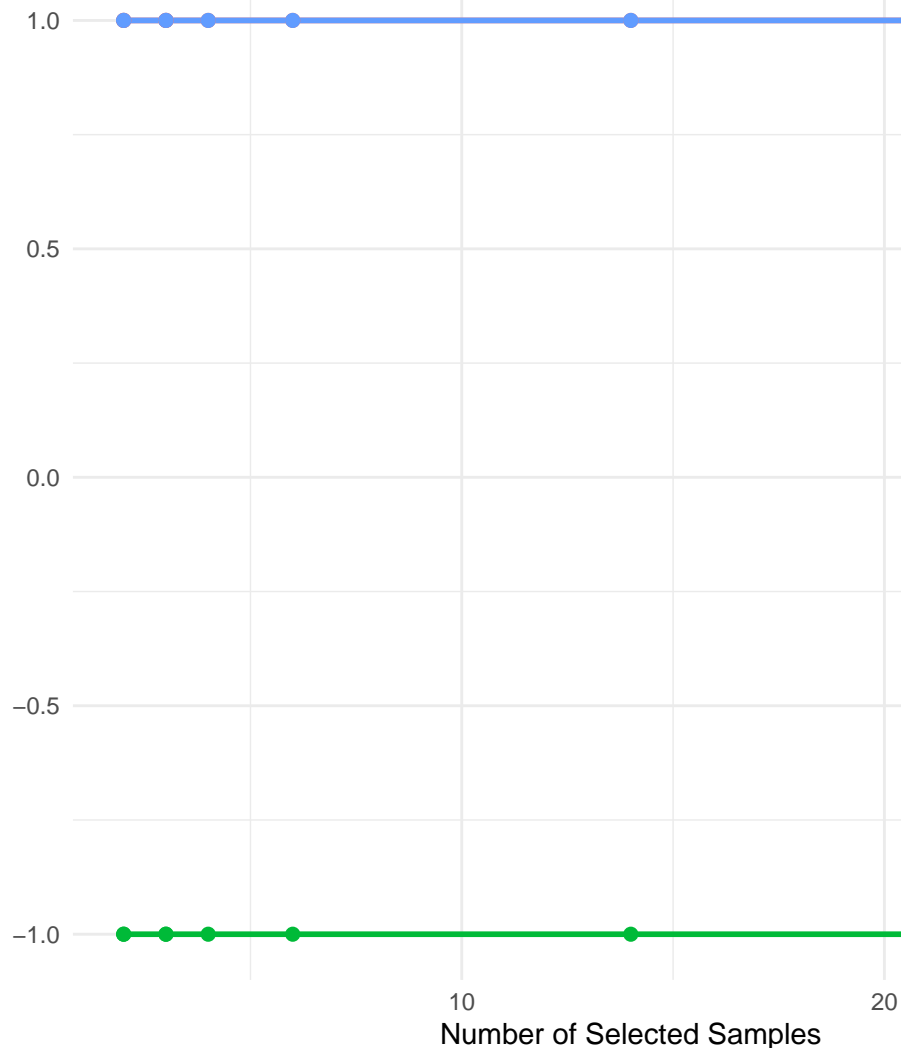


Score: ESPIID (Local, partial2 , Genotype: wave)

Kendall Correlation

Summary Statistic

- range
- slope
- variance

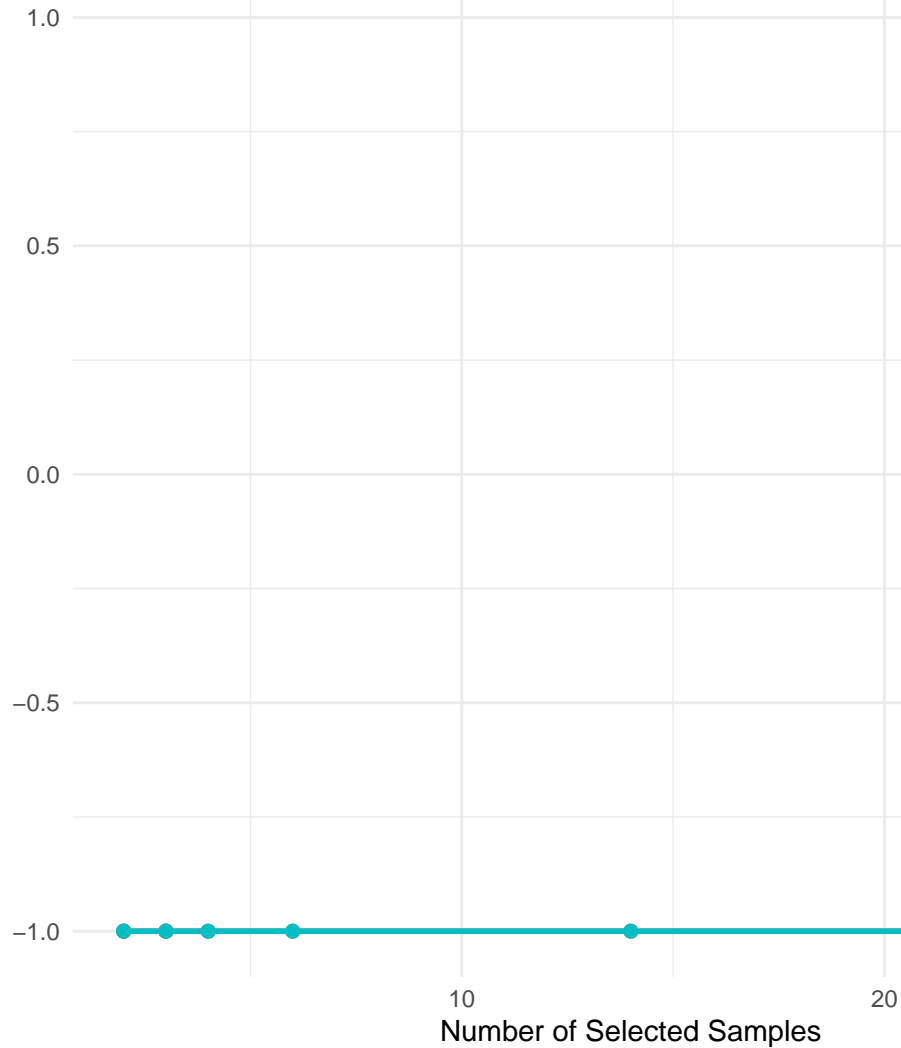


Score: PSI (Local, partial2 , Genotype: linear)

Kendall Correlation

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

10

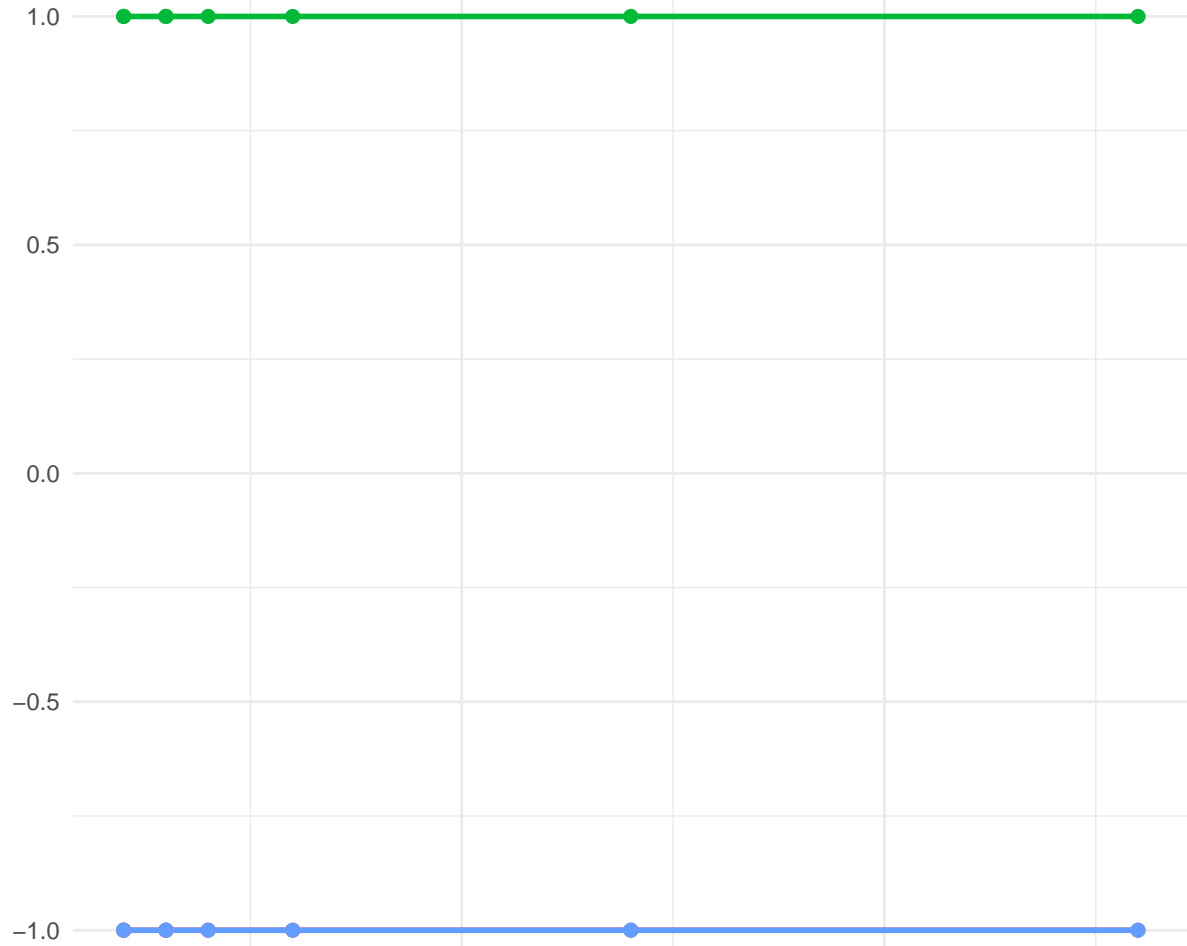
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Summary Statistic

range

slope

variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

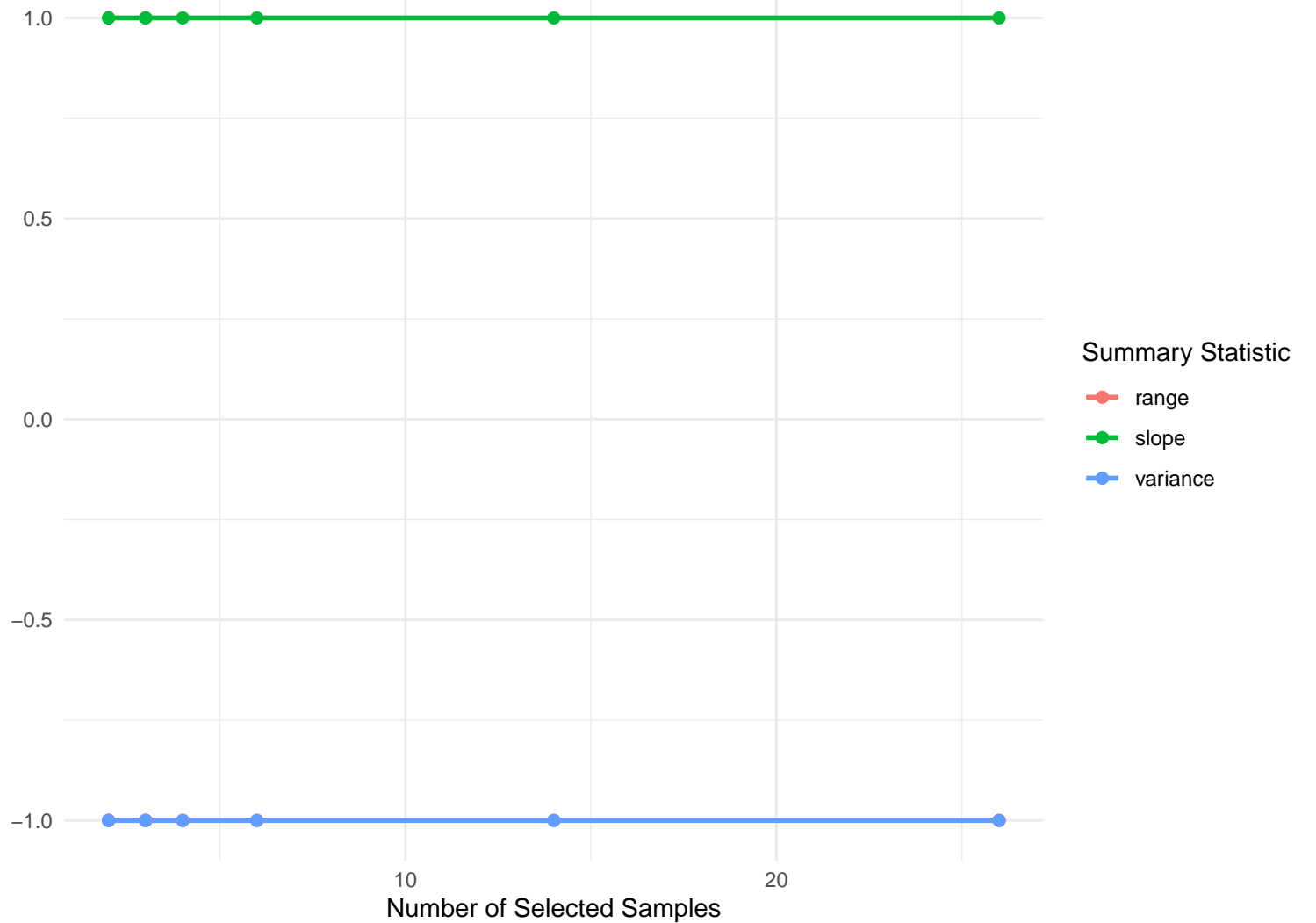
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Number of Selected Samples

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

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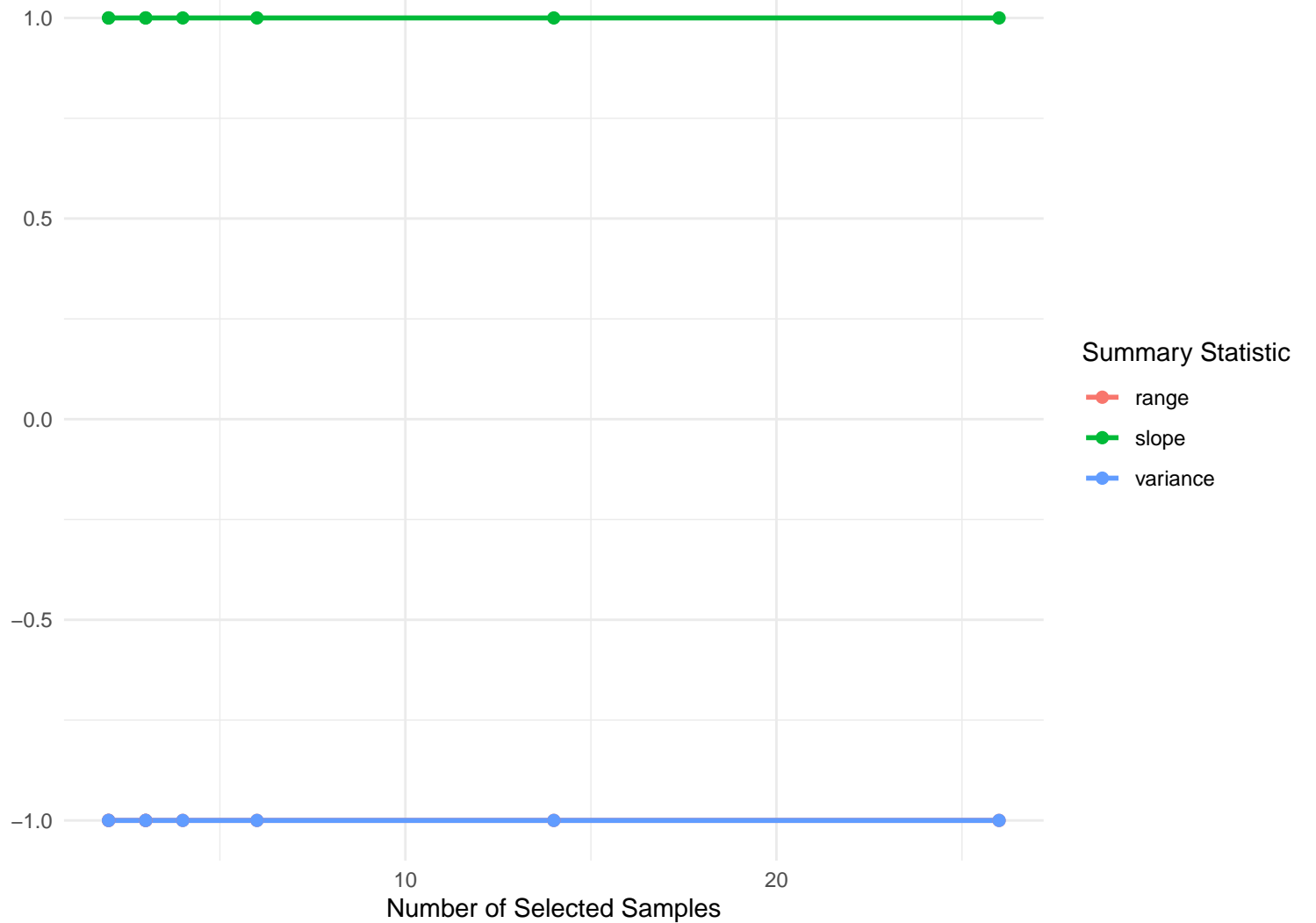
Number of Selected Samples

Summary Statistic

range

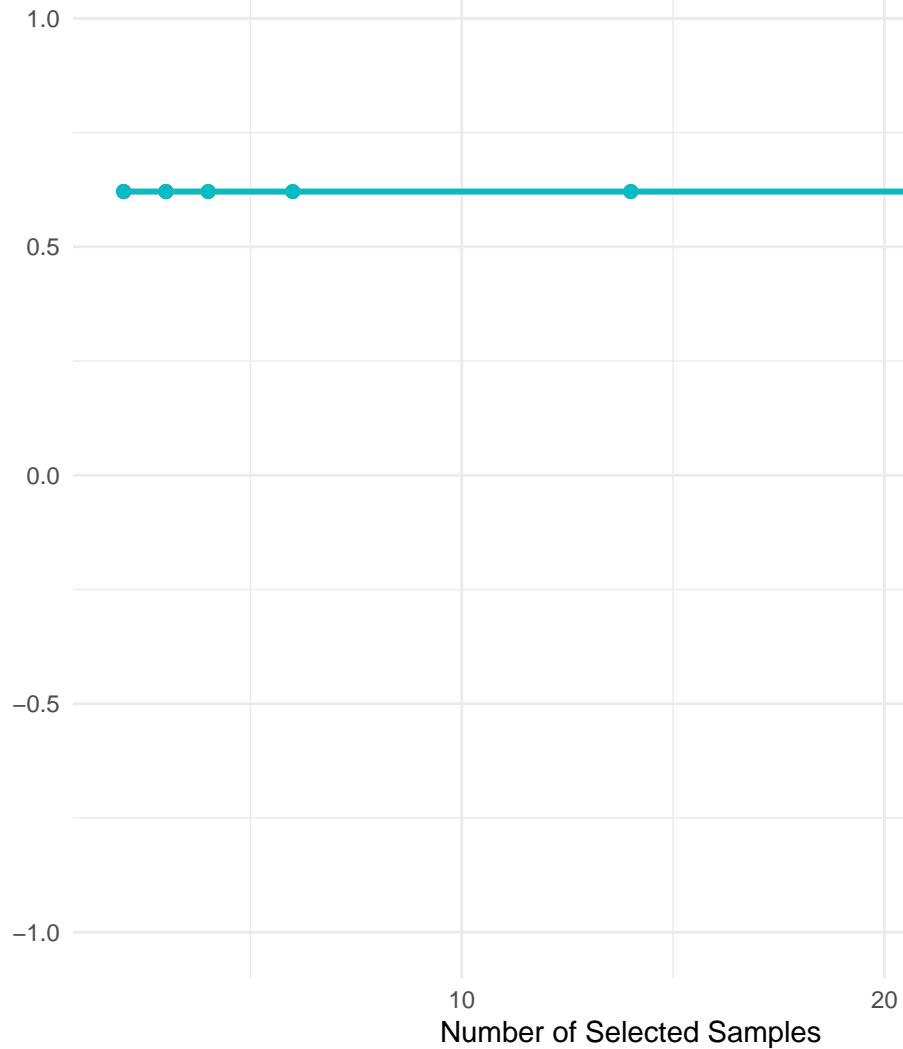
slope

variance



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

Kendall Correlation



Summary Statistic

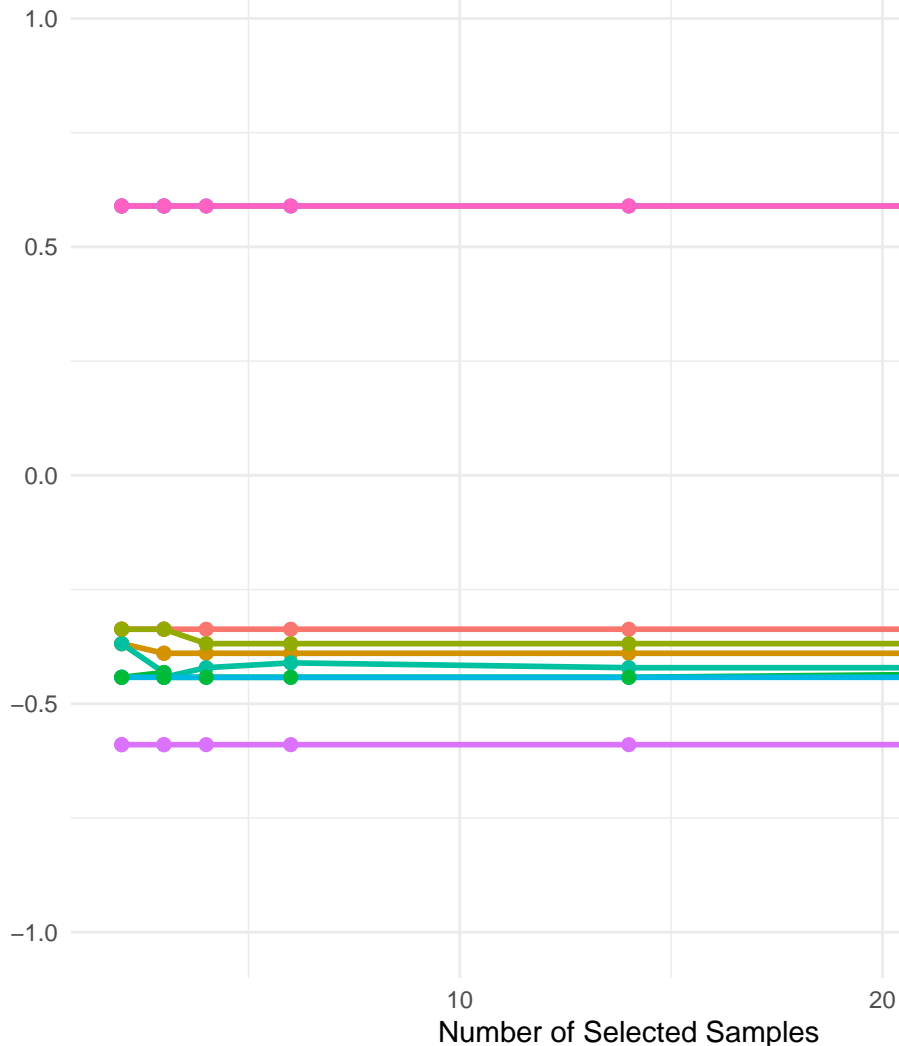
- range
- variance

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

Kendall Correlation

Summary Statistic

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

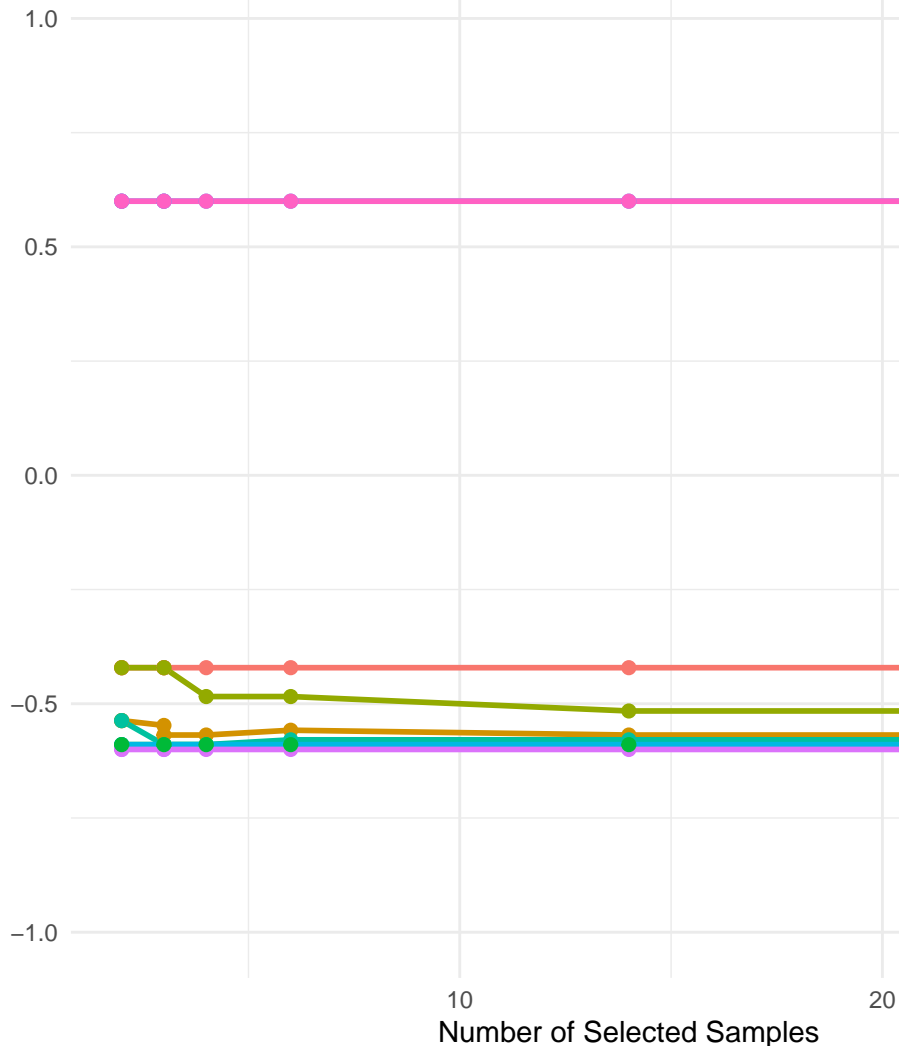


Score: RPI (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

Summary Statistic

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

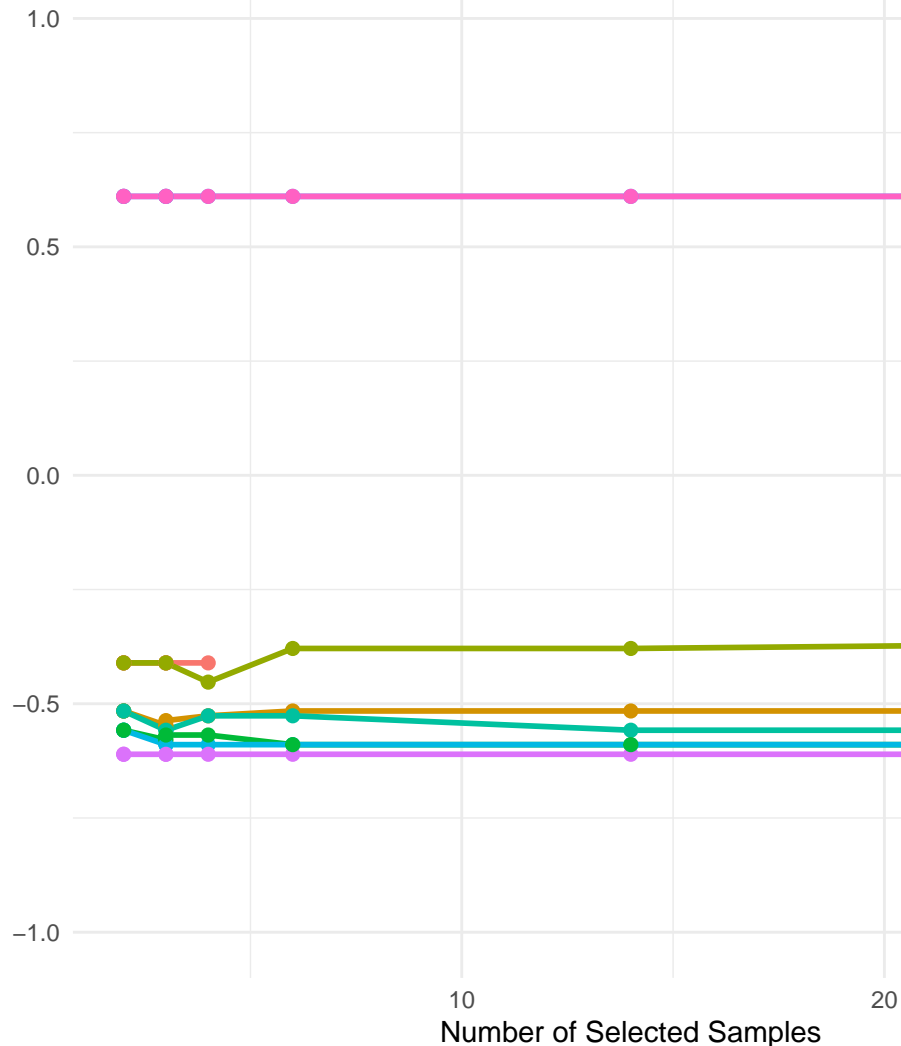


Score: RPI (Local, partial2 , Genotype: wave)

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

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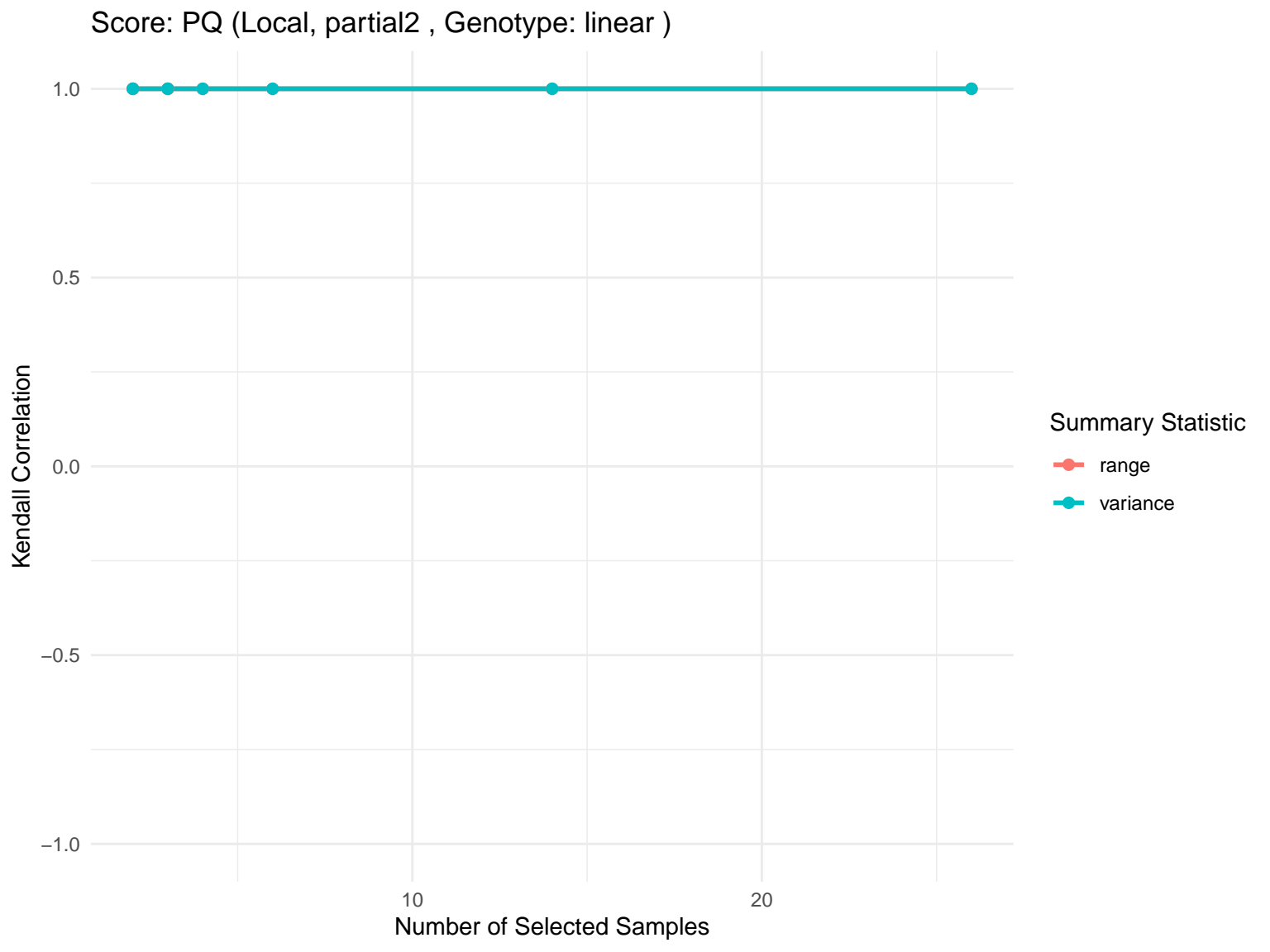
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Number of Selected Samples

Summary Statistic

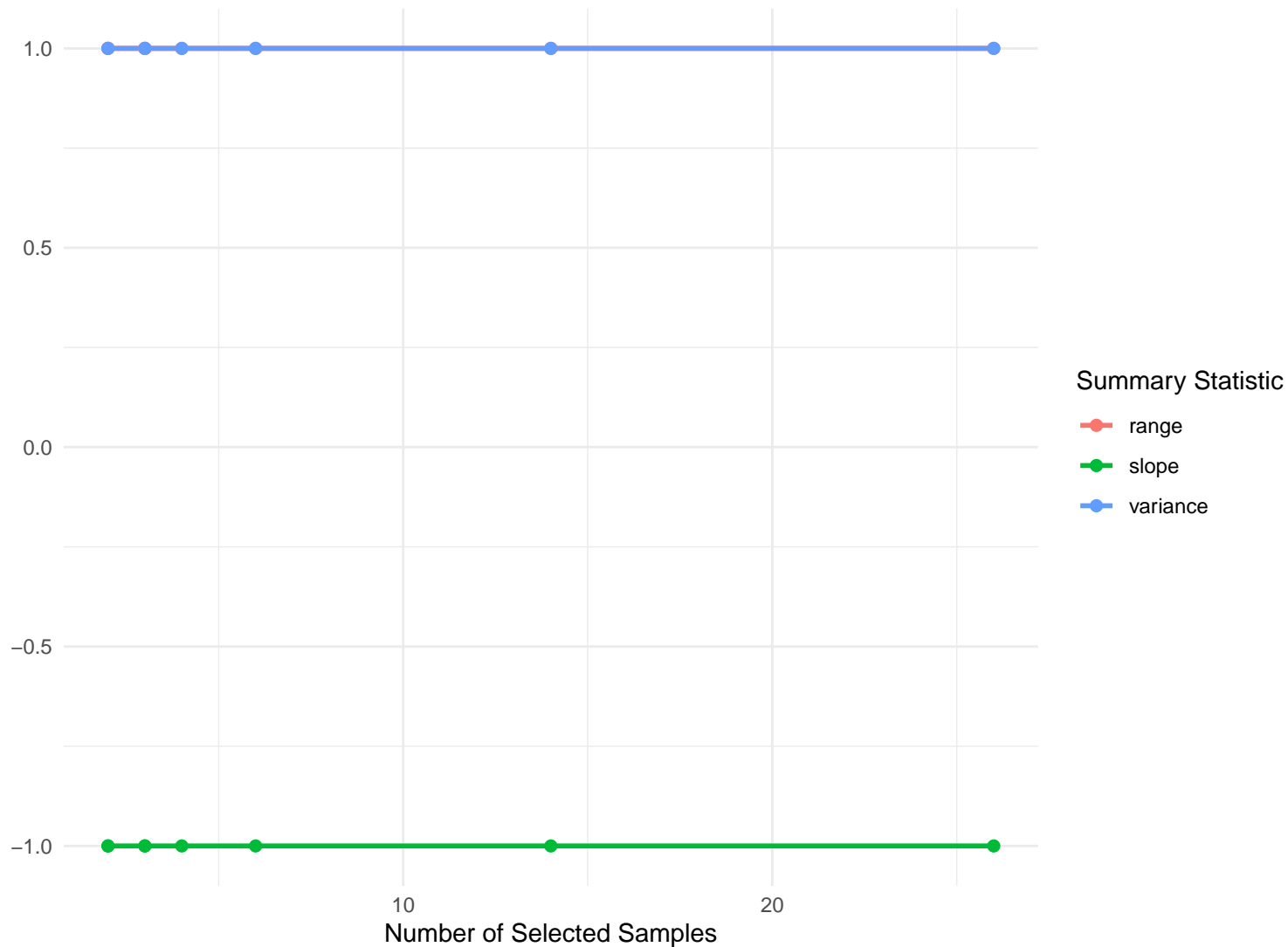
range

variance



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

Kendall Correlation



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

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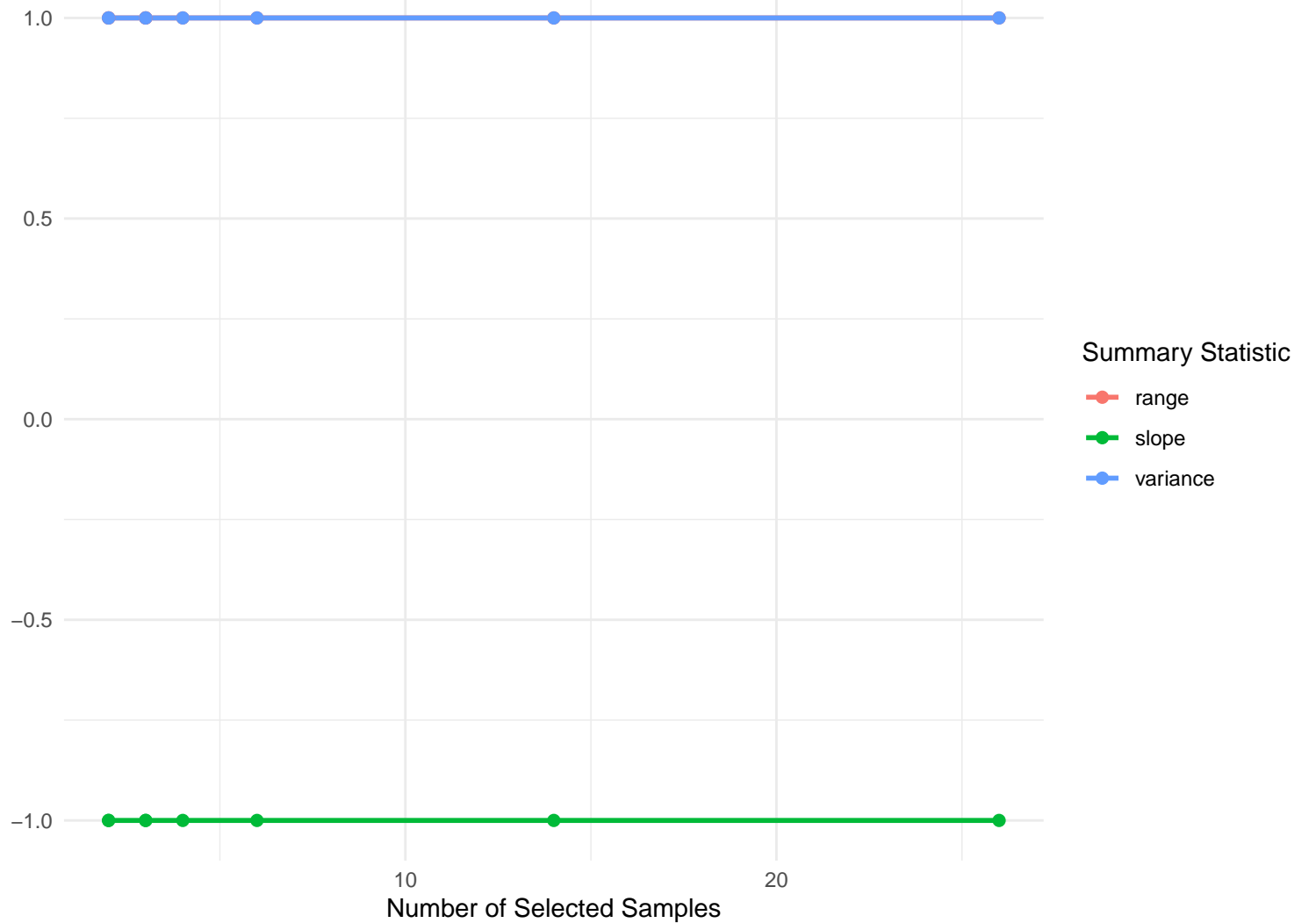
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Summary Statistic

range

slope

variance



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

Kendall Correlation

1.0

0.5

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-0.5

-1.0

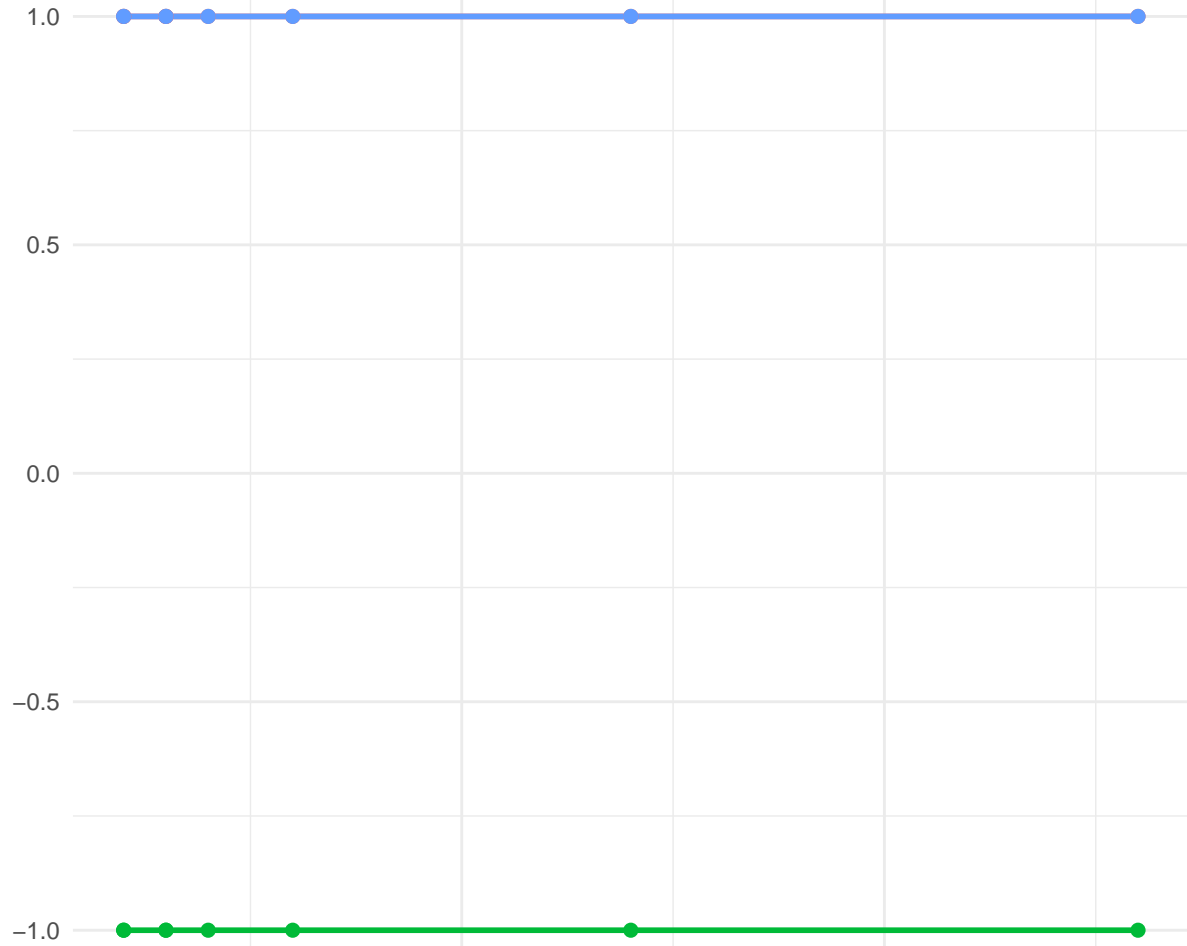
Number of Selected Samples

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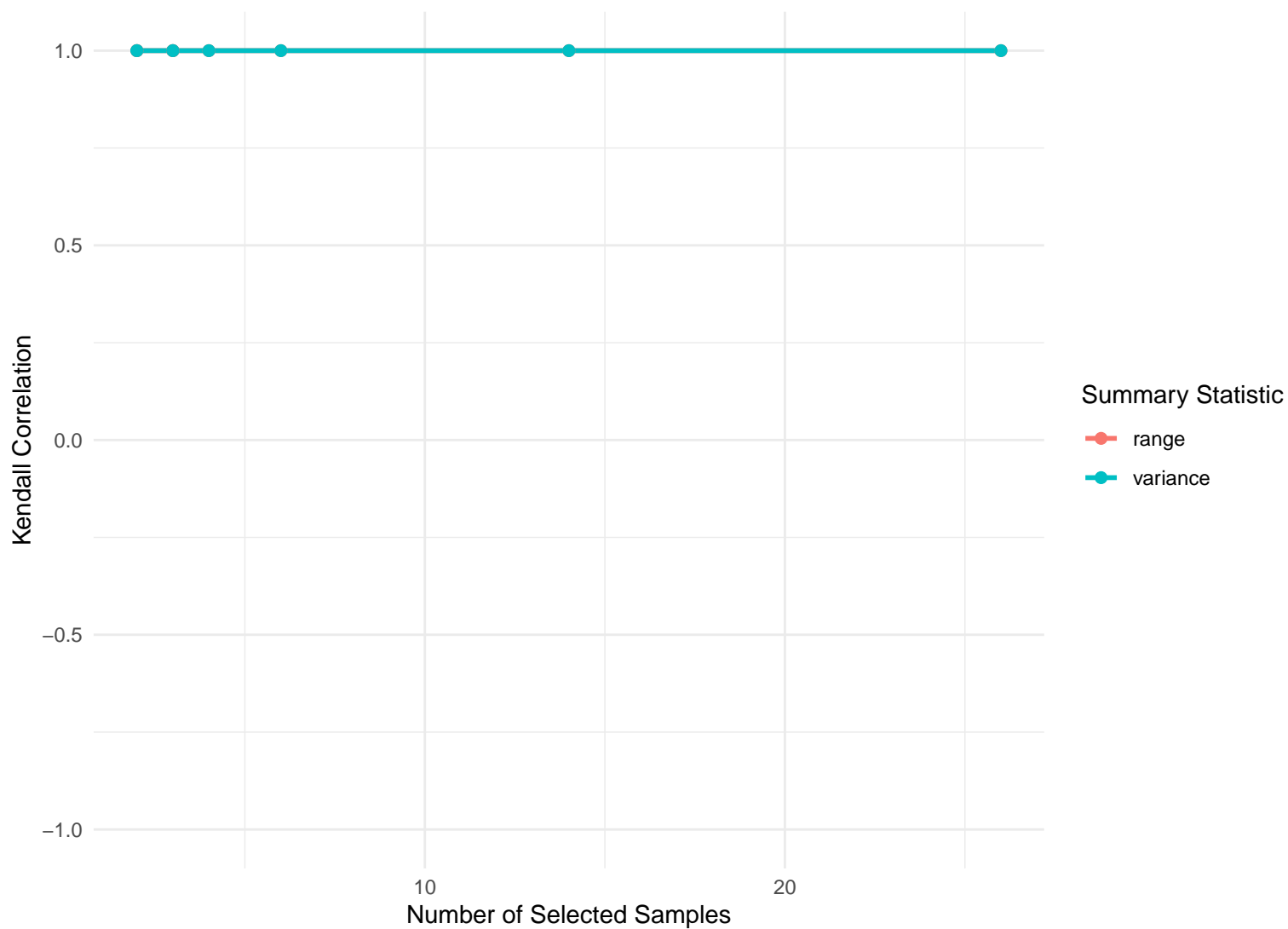
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Summary Statistic

- range
- slope
- variance

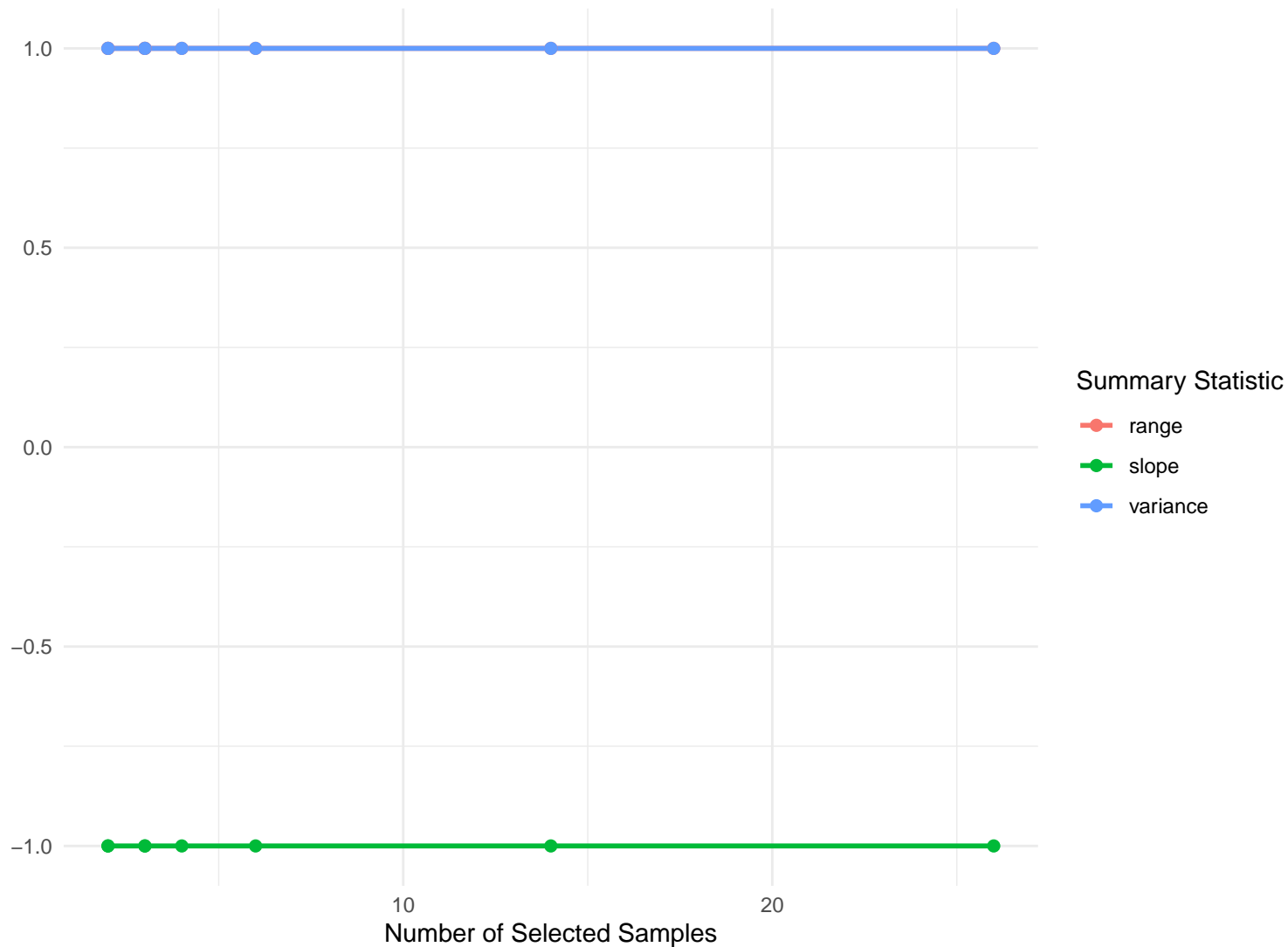


Score: PR (Local, partial2 , Genotype: linear)



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

Kendall Correlation



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

Kendall Correlation

Summary Statistic

- range
- slope
- variance

1.0
0.5
0.0
-0.5
-1.0

Number of Selected Samples

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1.0
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Score: PR (Local, partial2 , Genotype: wave)

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

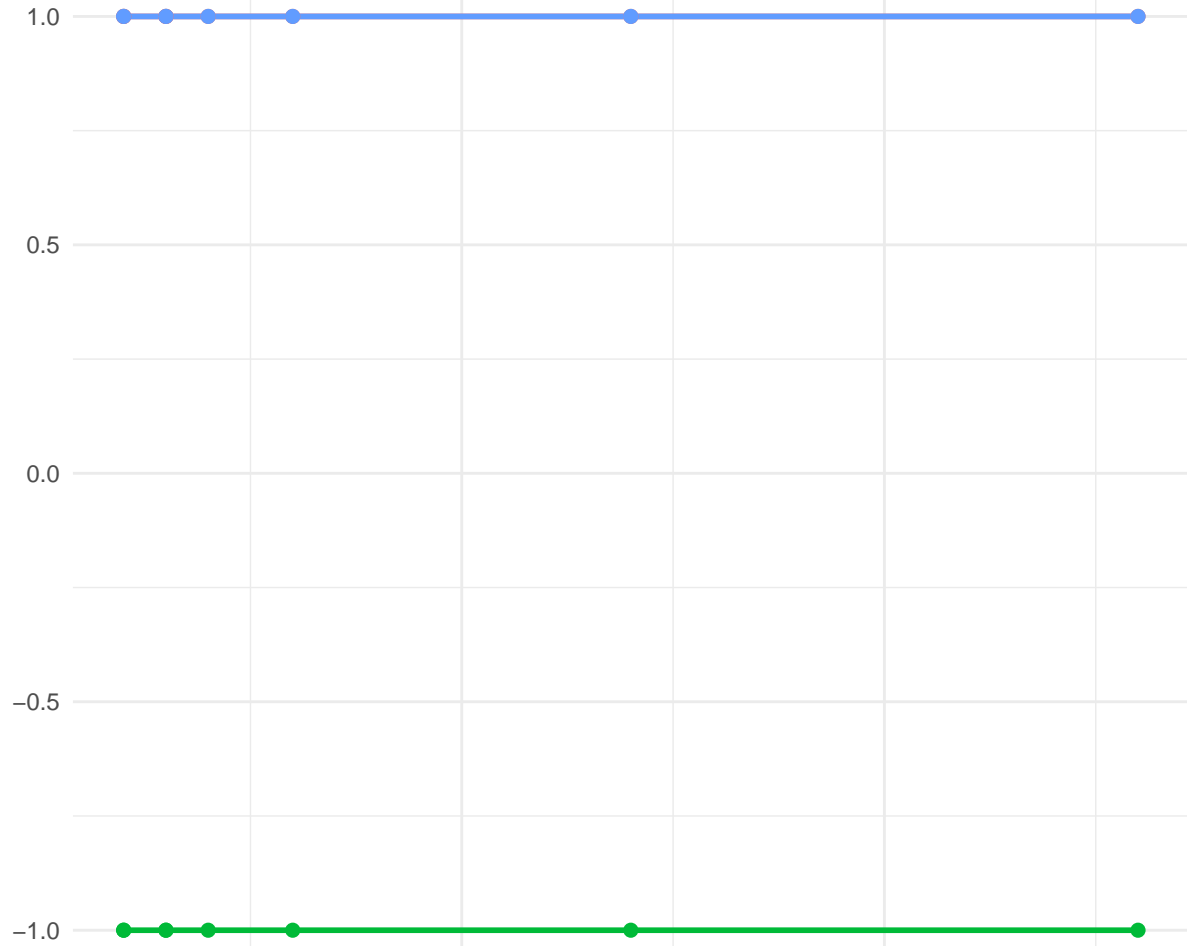
Number of Selected Samples

10

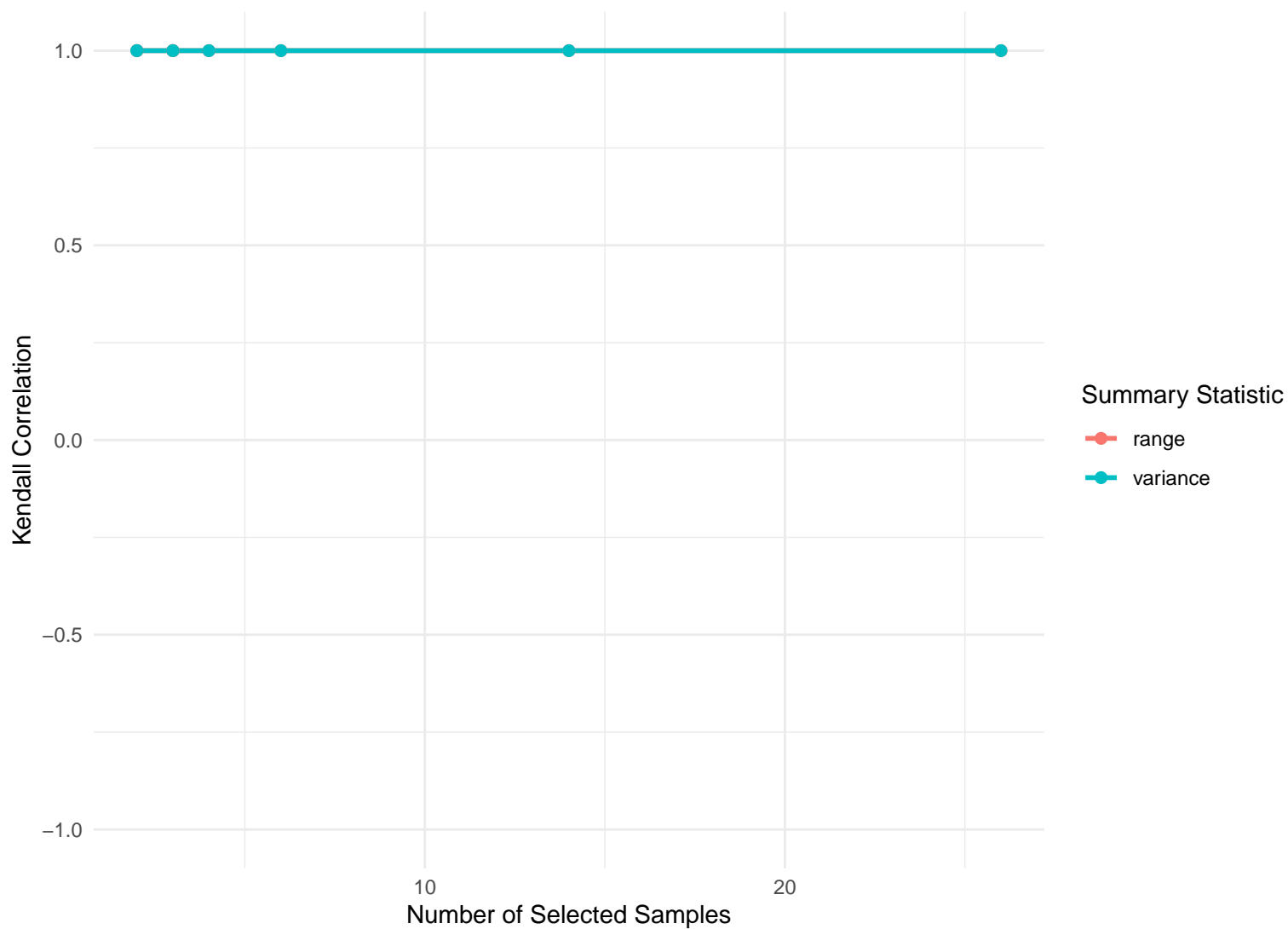
20

Summary Statistic

- range
- slope
- variance



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

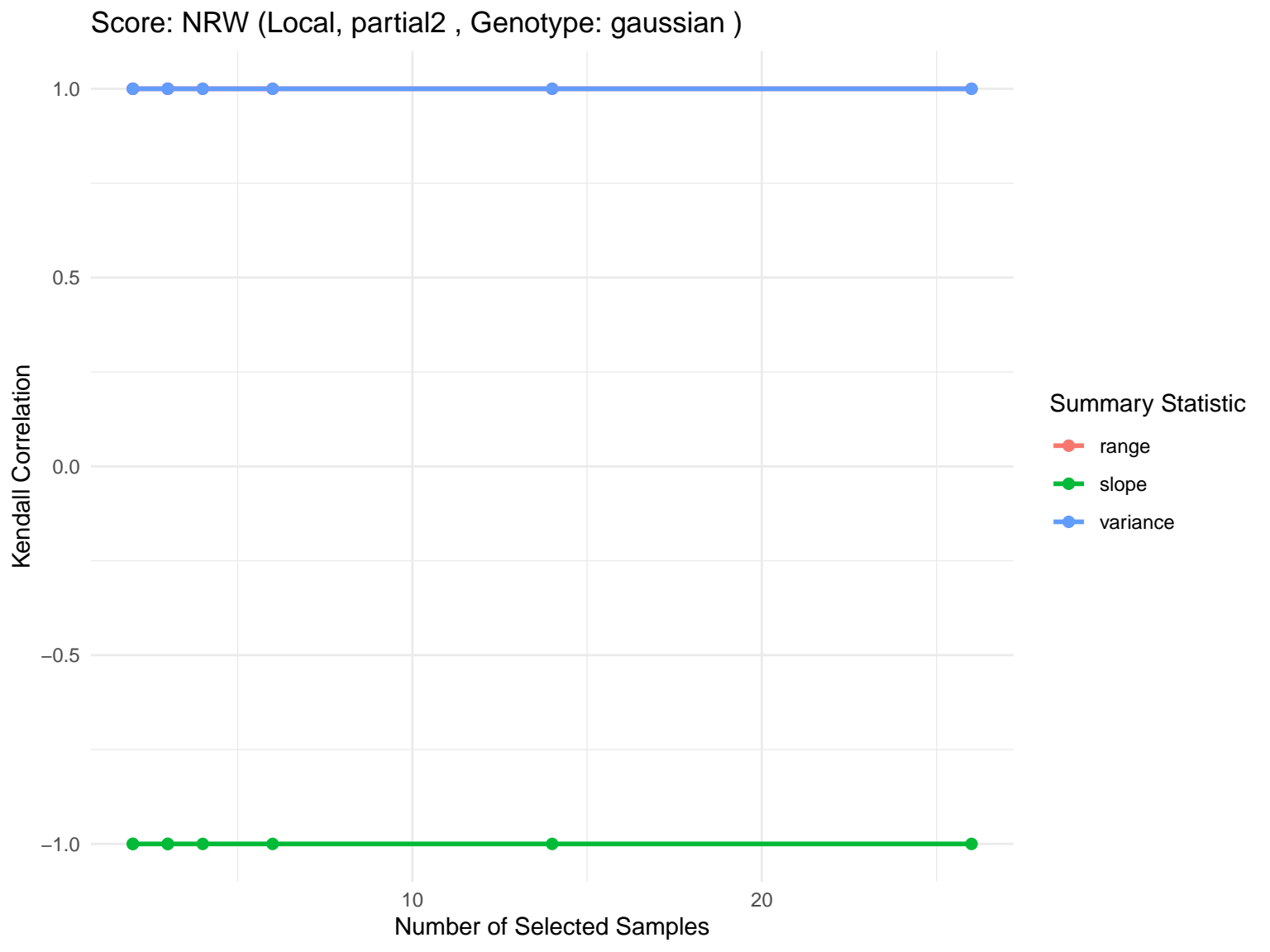
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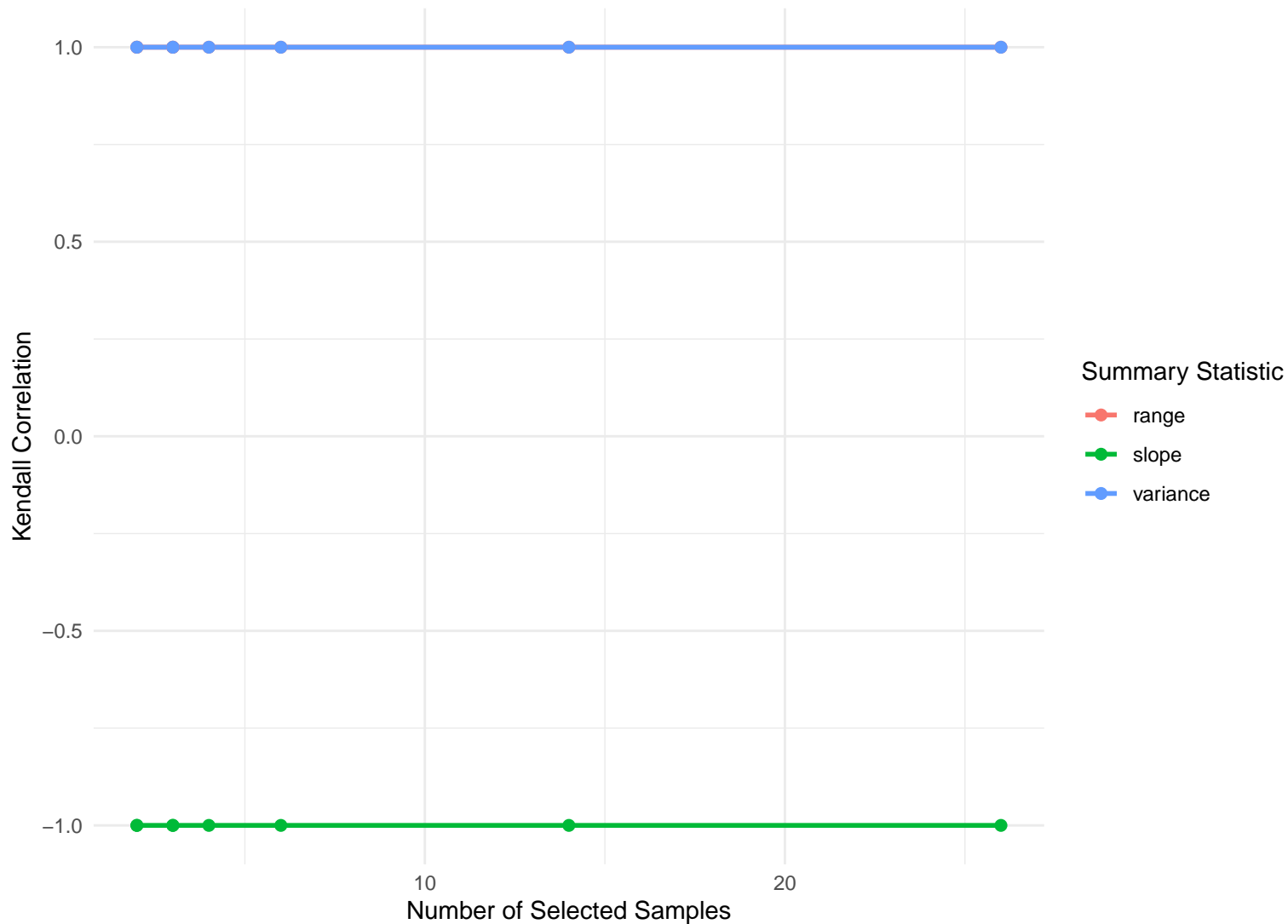
Number of Selected Samples

Summary Statistic

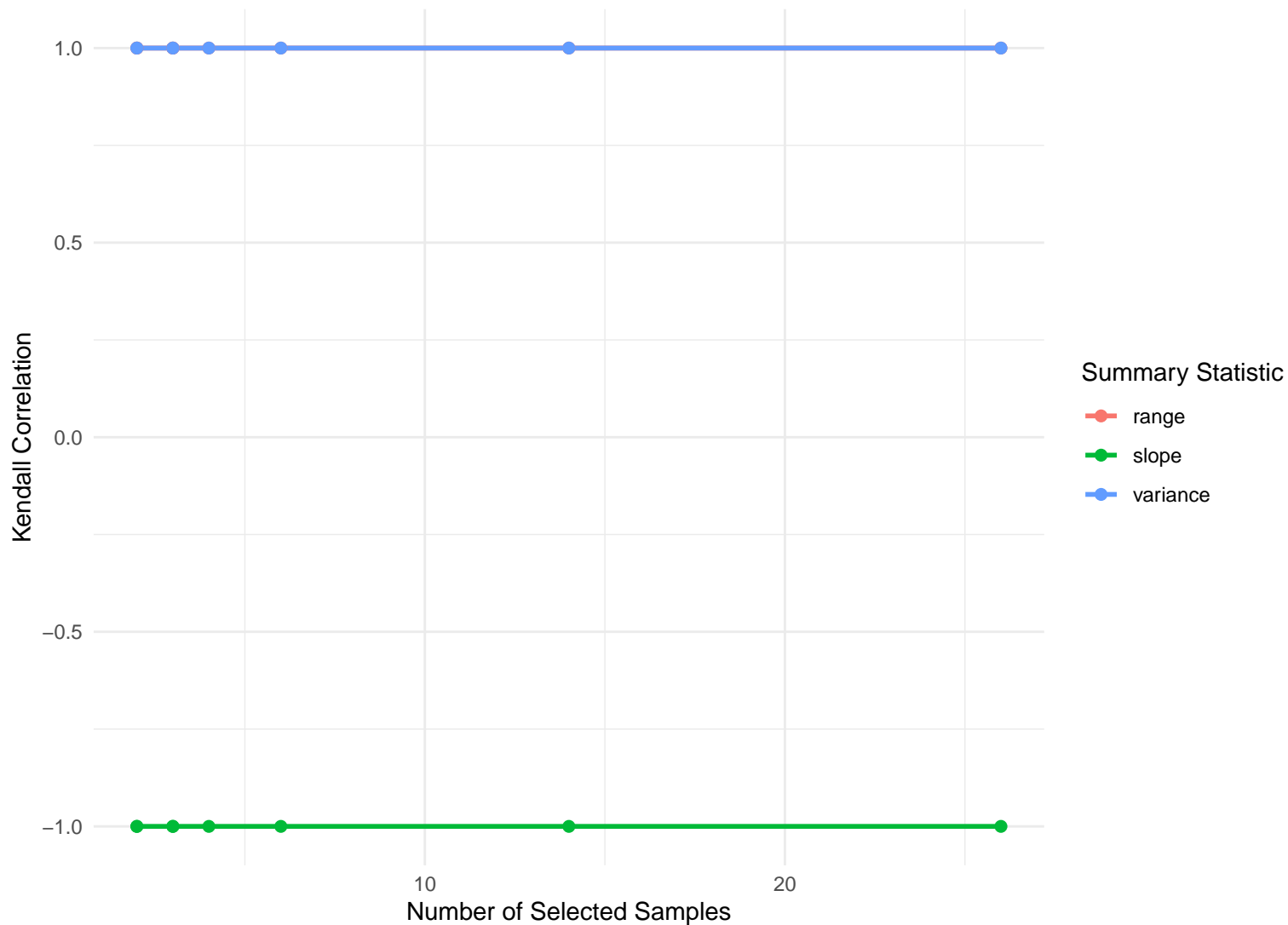
- range
- slope
- variance



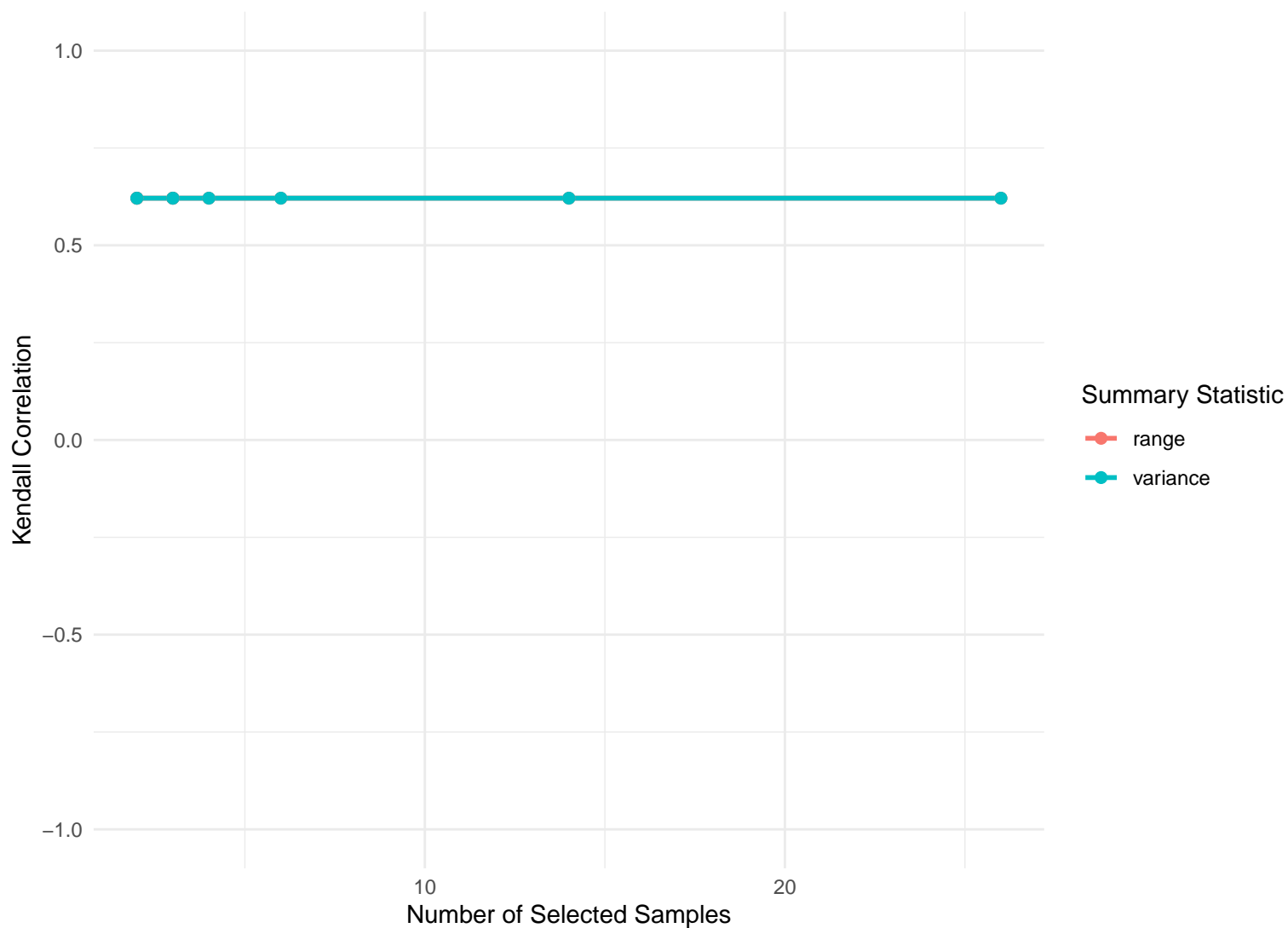
Score: NRW (Local, partial2 , Genotype: sinusoidal)



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



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



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

Kendall Correlation

Summary Statistic

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

1.0

0.5

0.0

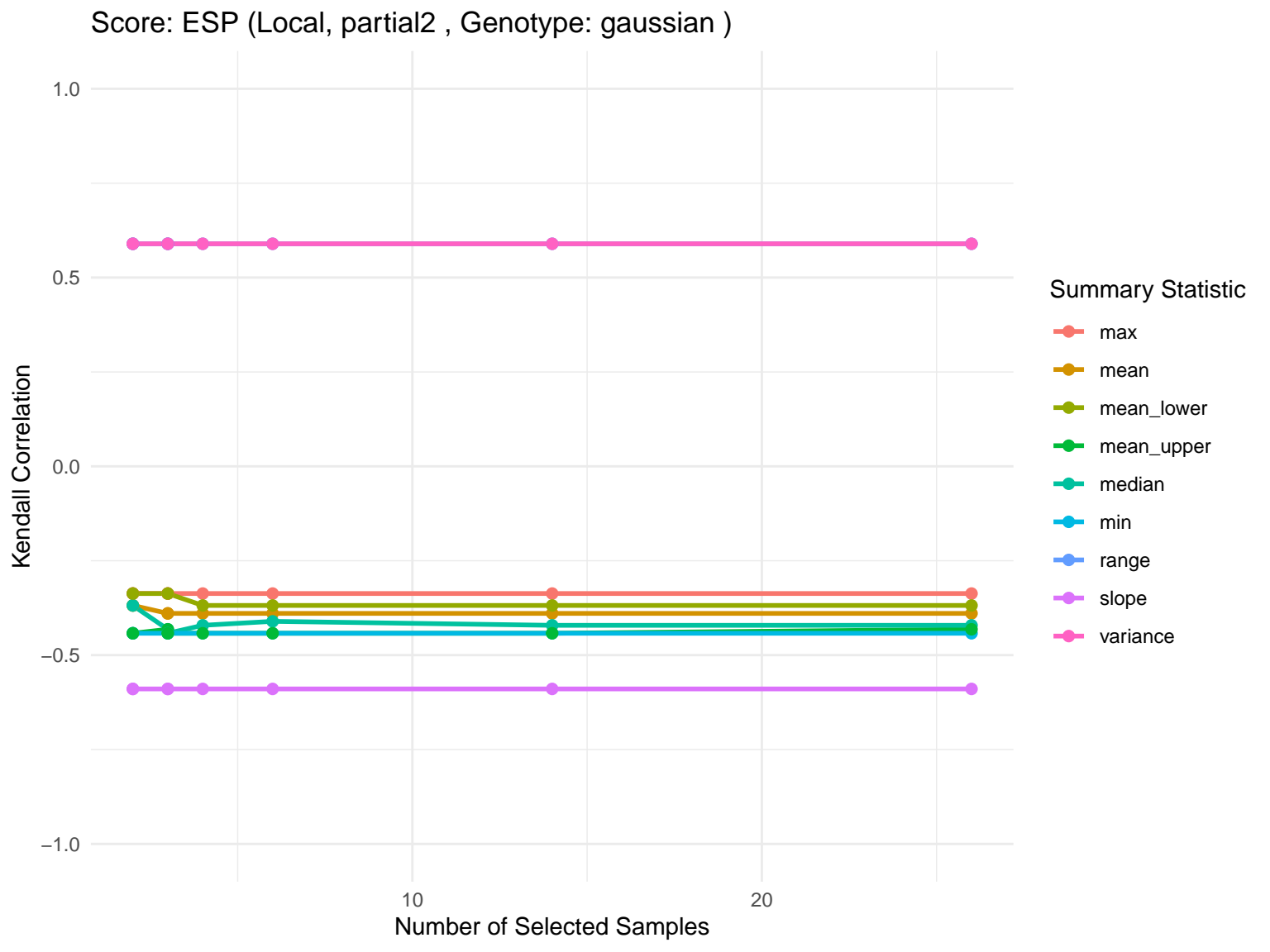
-0.5

-1.0

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Number of Selected Samples

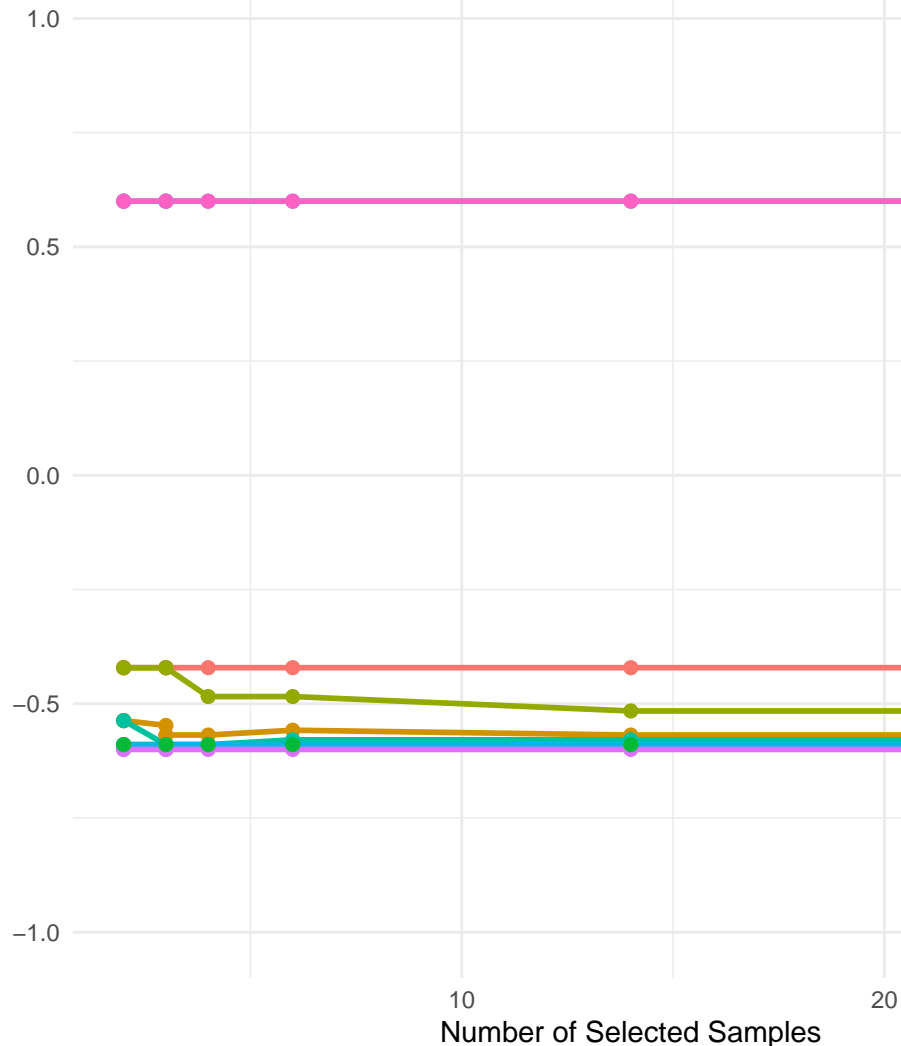


Score: ESP (Local, partial2 , Genotype: sinusoidal)

Kendall Correlation

Summary Statistic

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



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

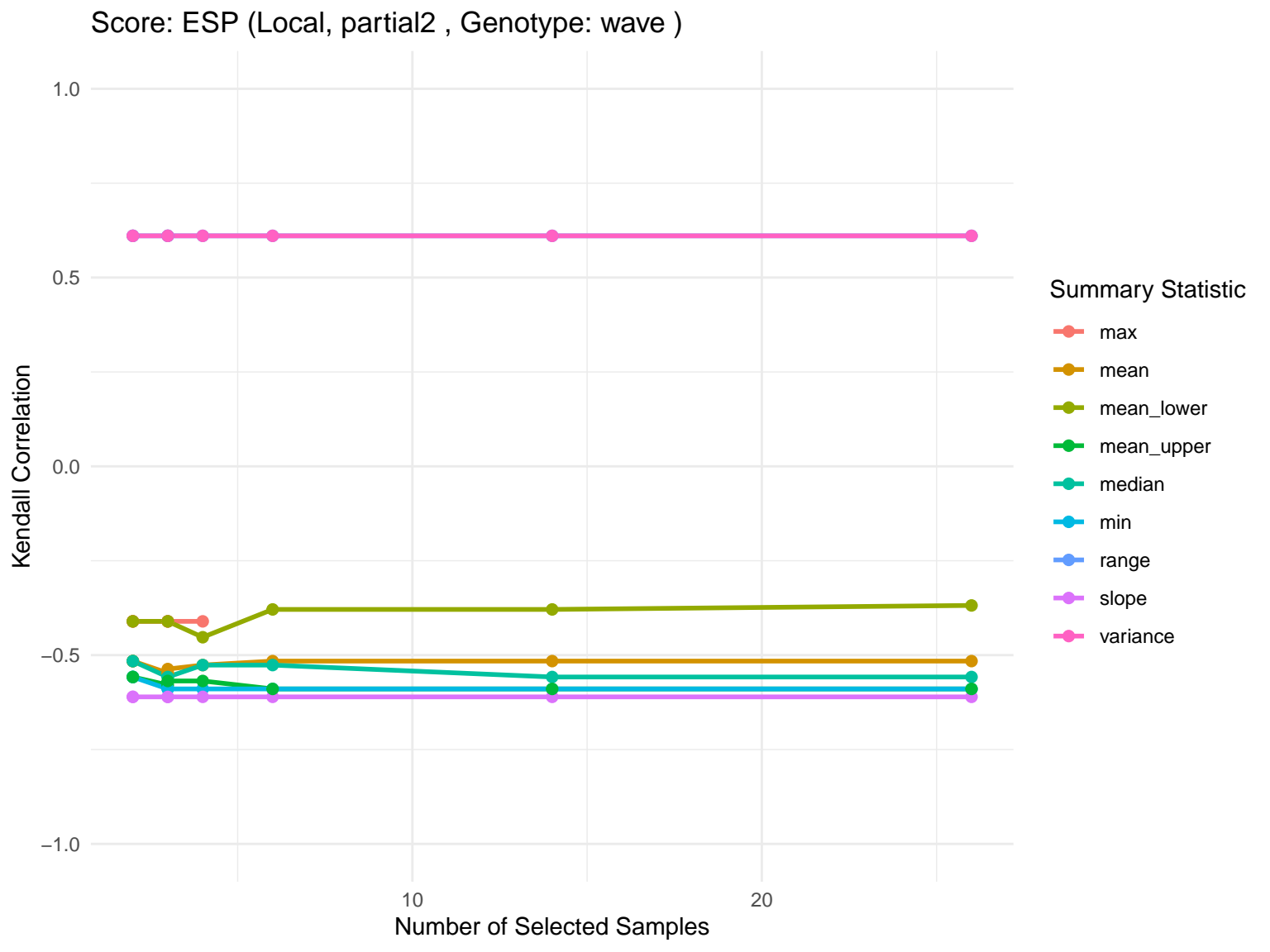
Kendall Correlation

Summary Statistic

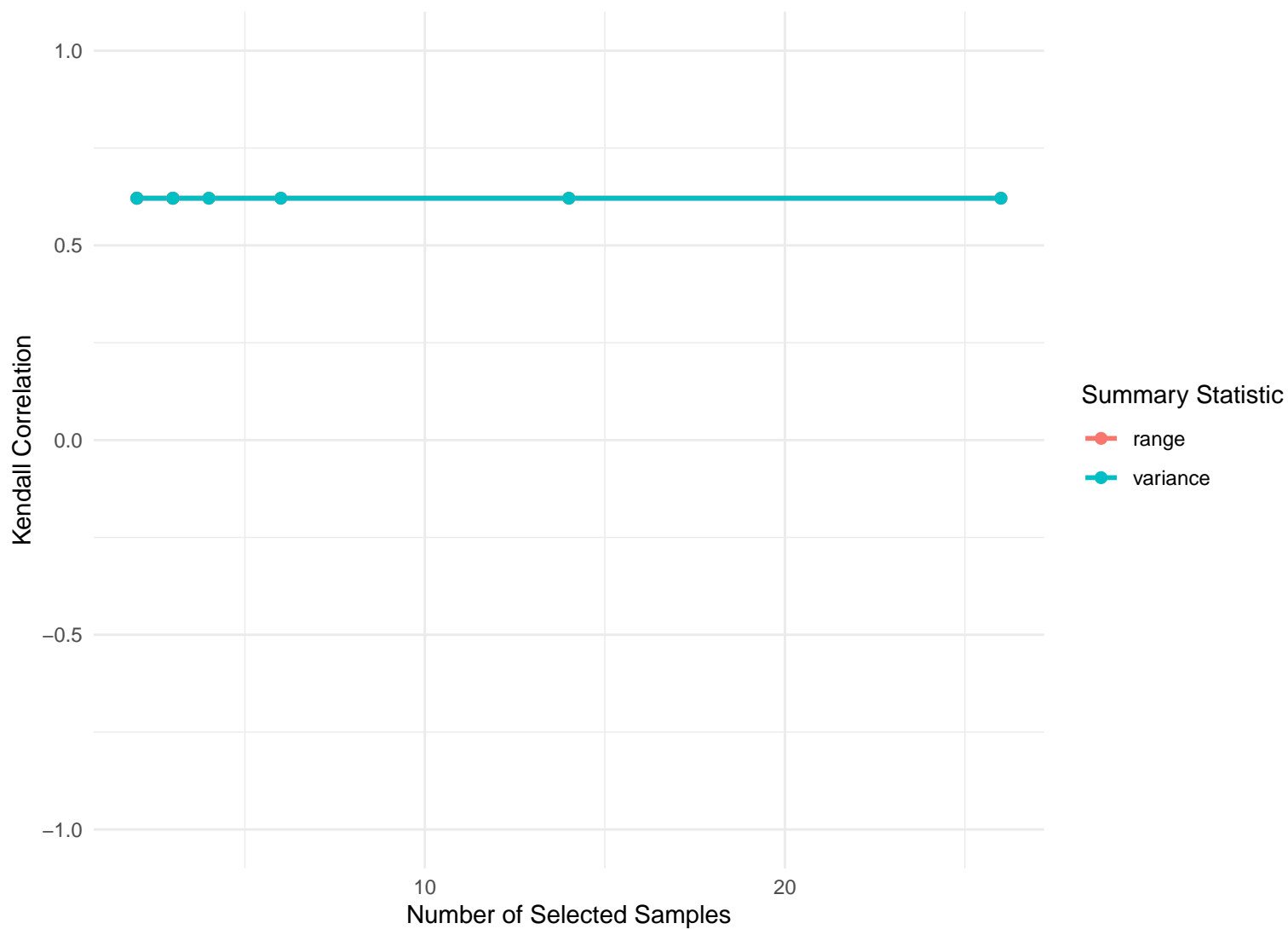
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance

1.0
0.5
0.0
-0.5
-1.0

10 20
Number of Selected Samples



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



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

Kendall Correlation

Summary Statistic

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

1.0

0.5

0.0

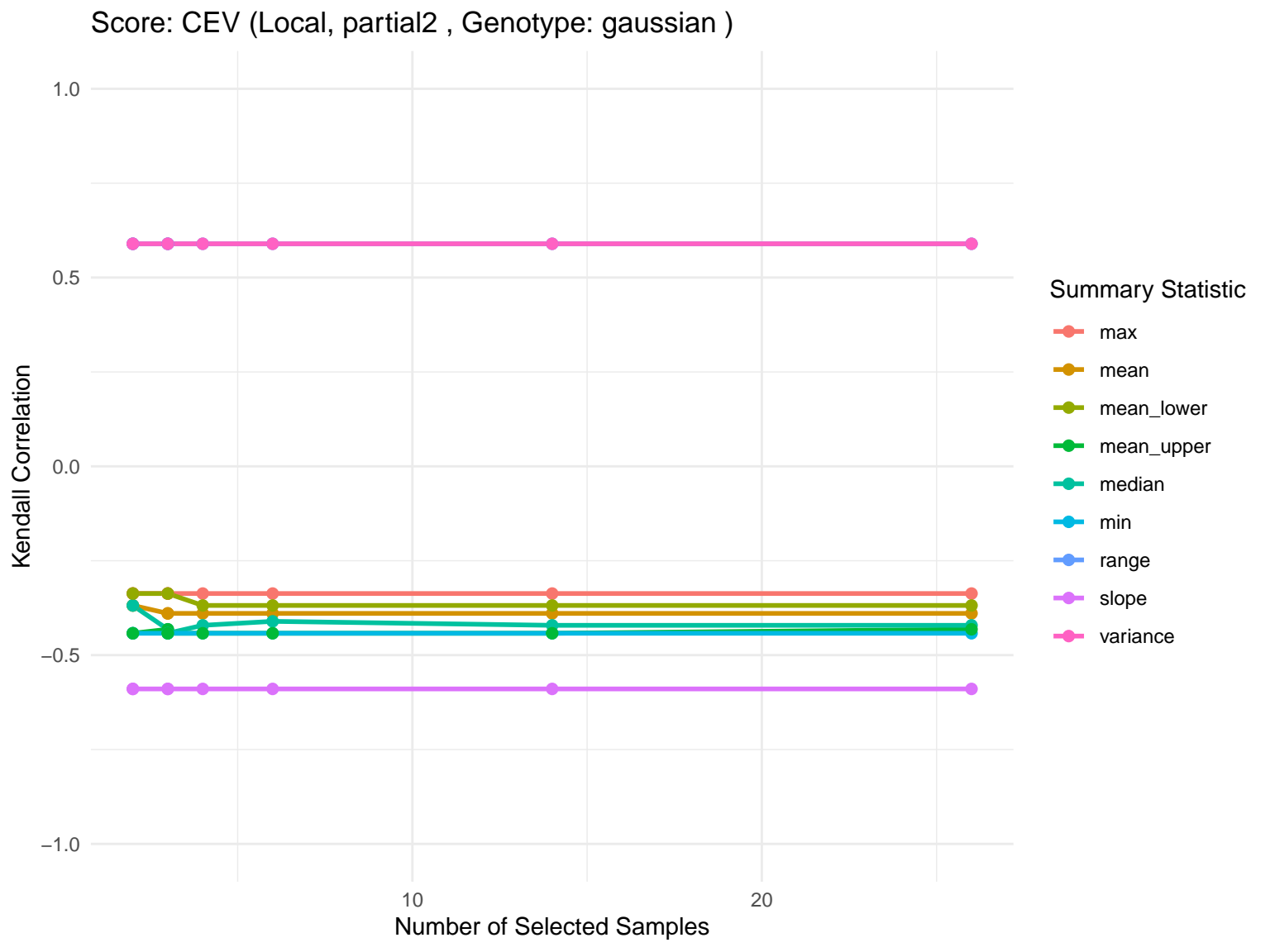
-0.5

-1.0

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Number of Selected Samples



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



1.0

0.0

-1.0

Number of Selected Samples

20

Summary Statistic

max

mean

mean_lower

mean_upper

median

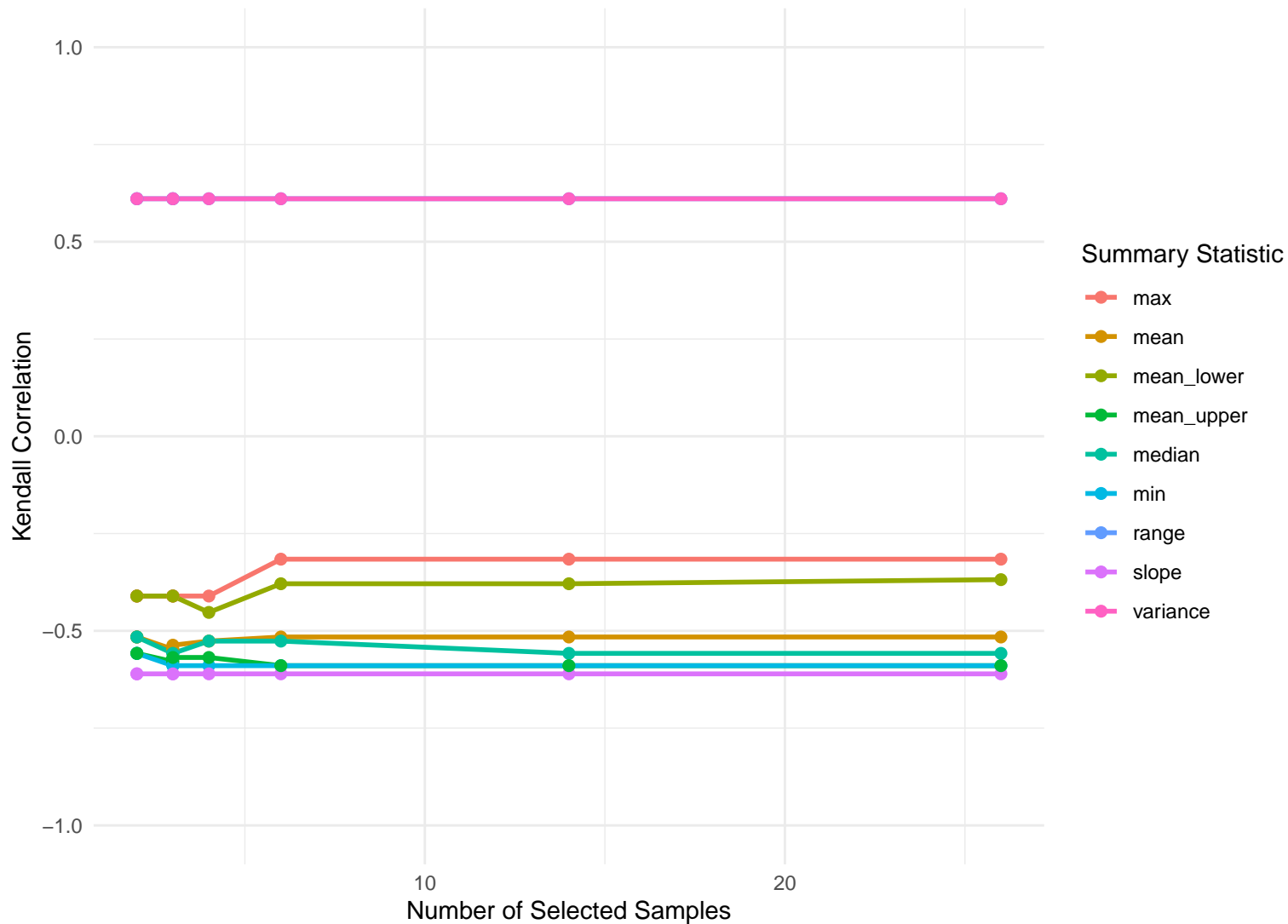
min

range

slope

variance

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



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

Kendall Correlation

1.0
0.5
0.0
-0.5
-1.0

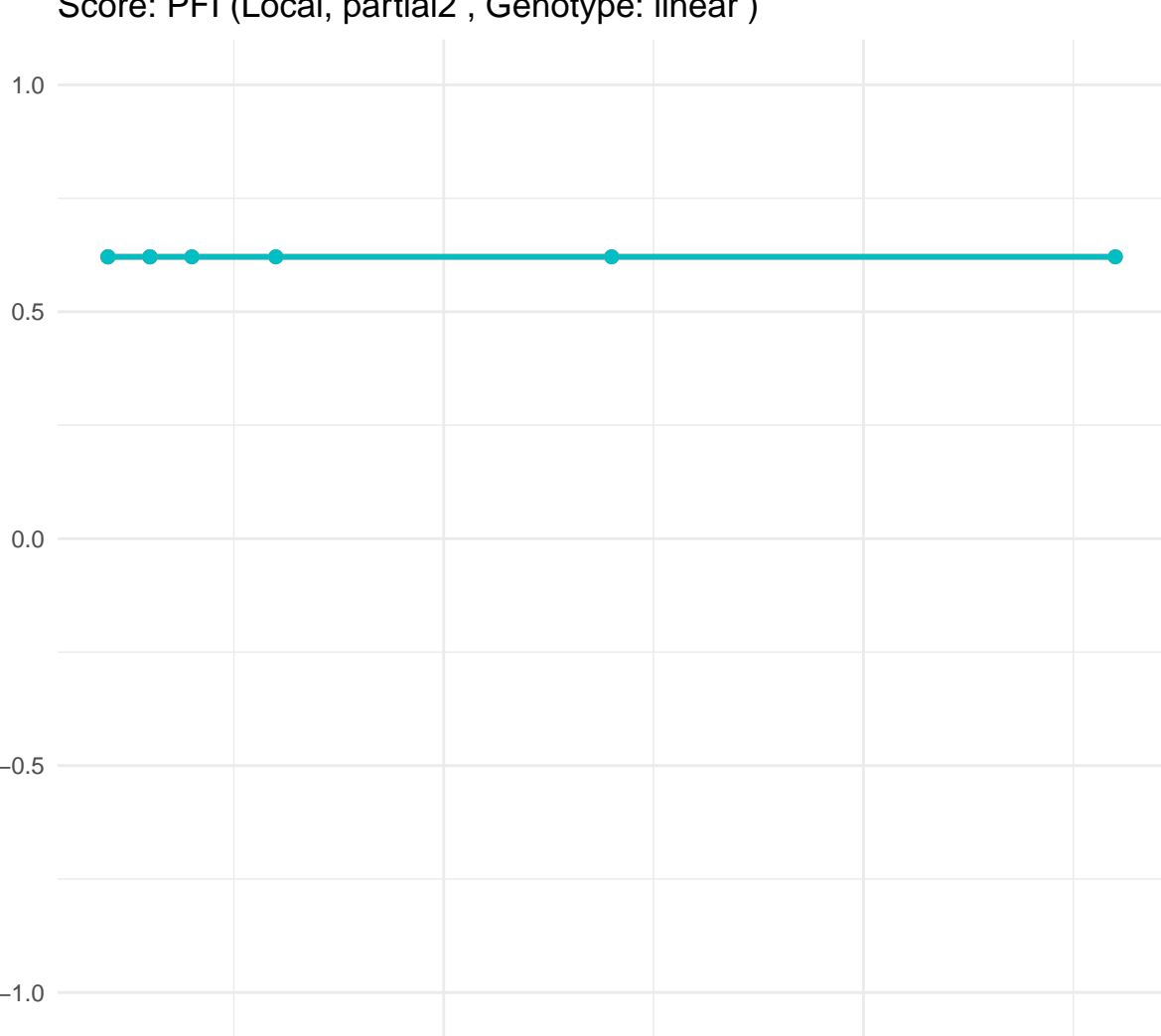
Number of Selected Samples

10

20

Summary Statistic

range
variance



Score: PFI (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

10

20

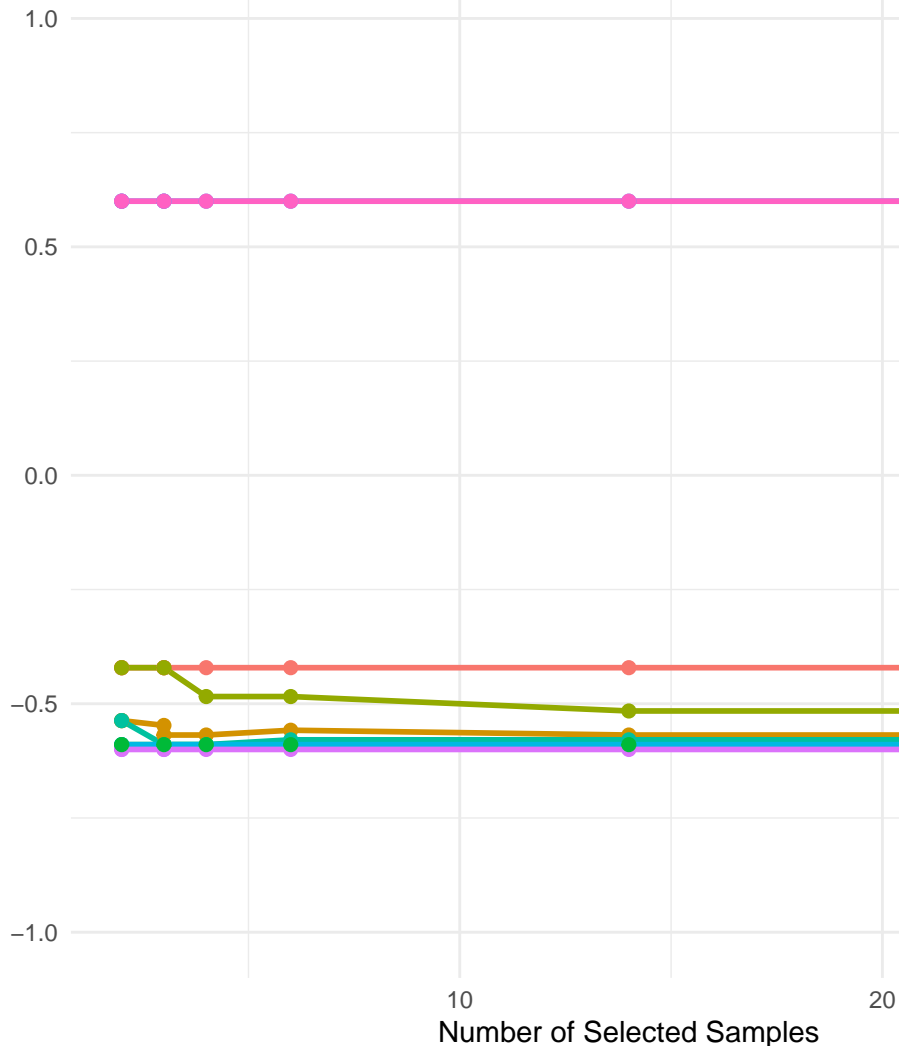
Number of Selected Samples

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)

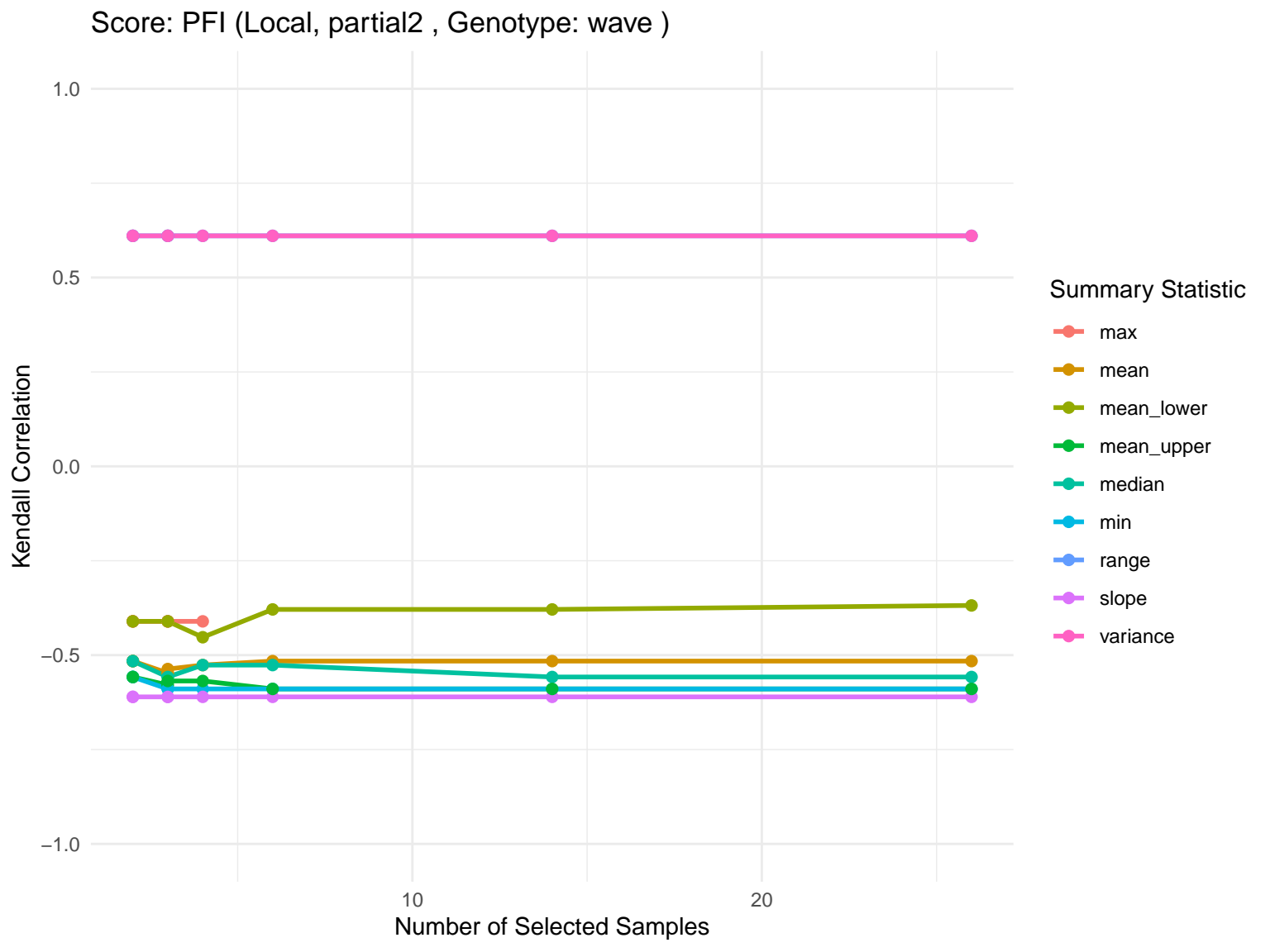
Kendall Correlation

Summary Statistic

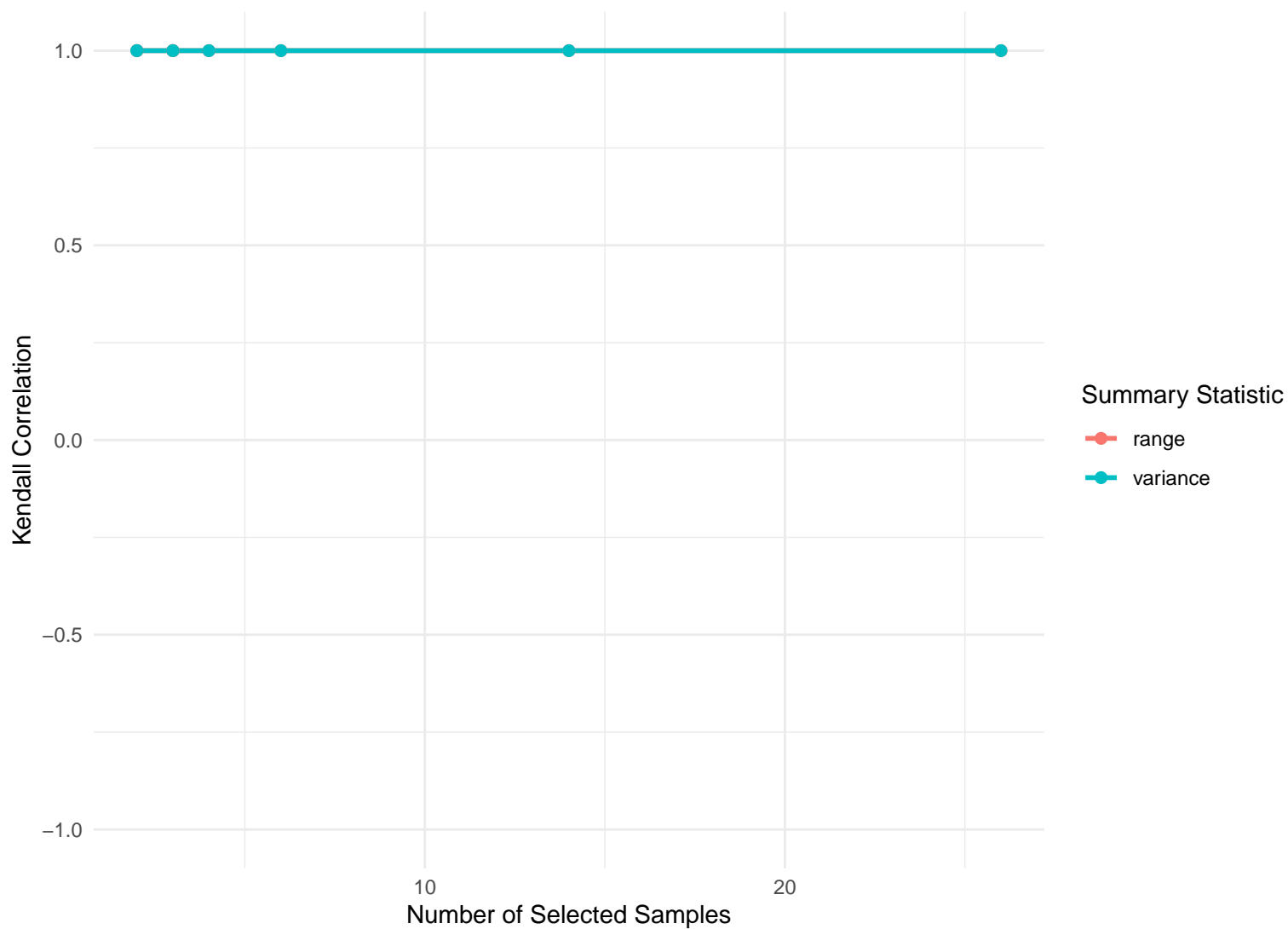
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance

1.0
0.5
0.0
-0.5
-1.0

10 20
Number of Selected Samples



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

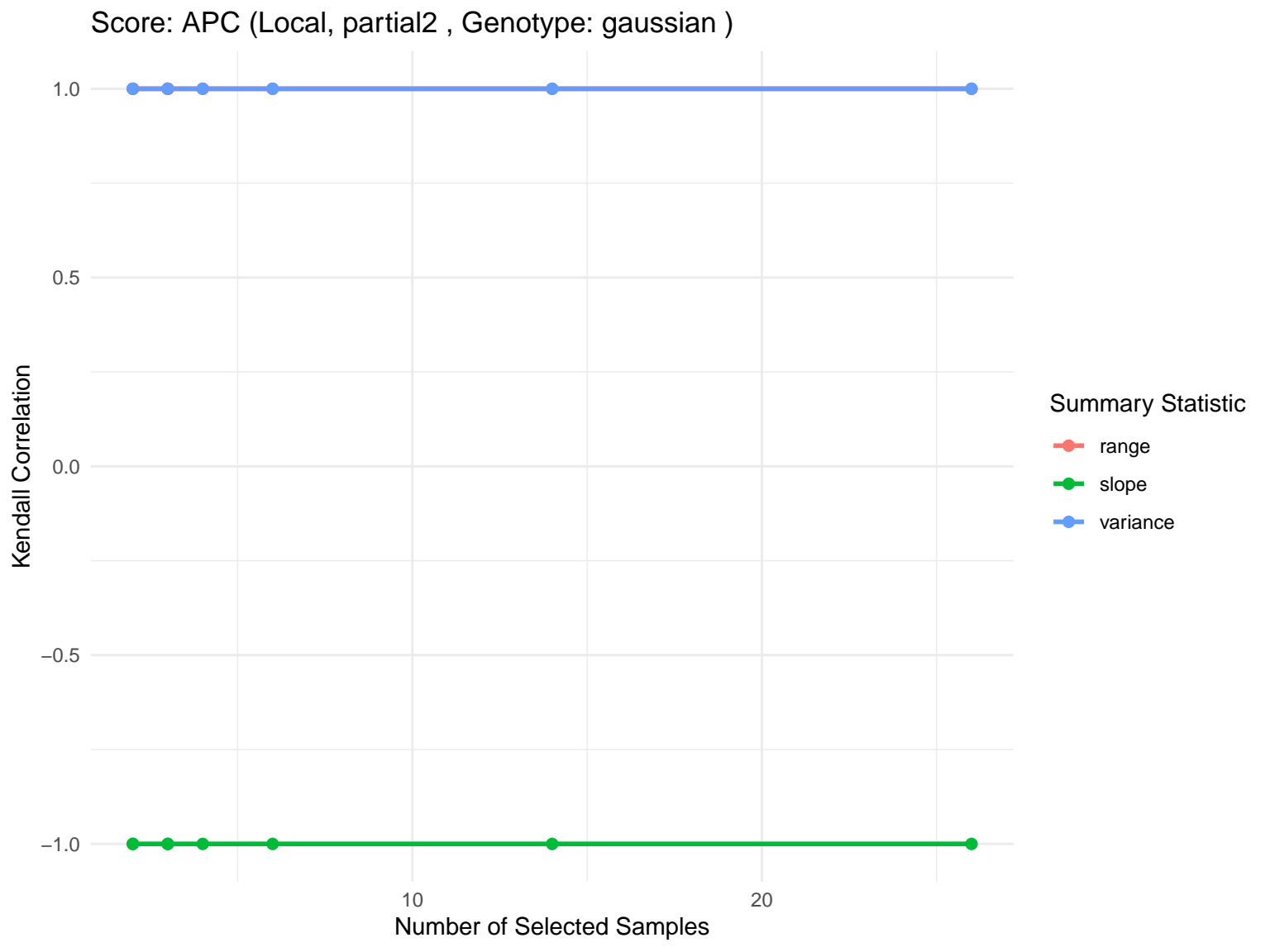
10

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Number of Selected Samples

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

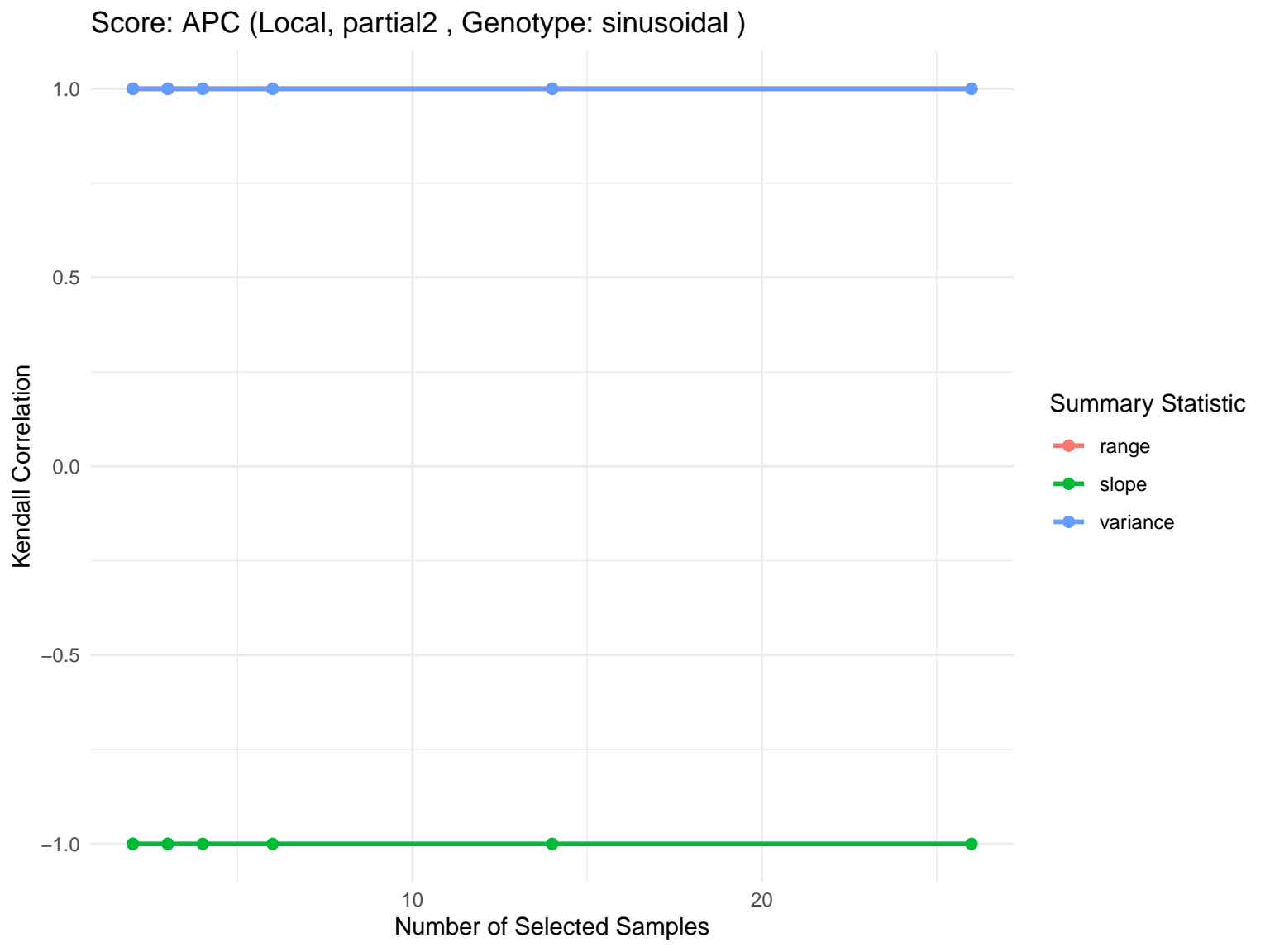
10

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Number of Selected Samples

Summary Statistic

- range
- slope
- variance

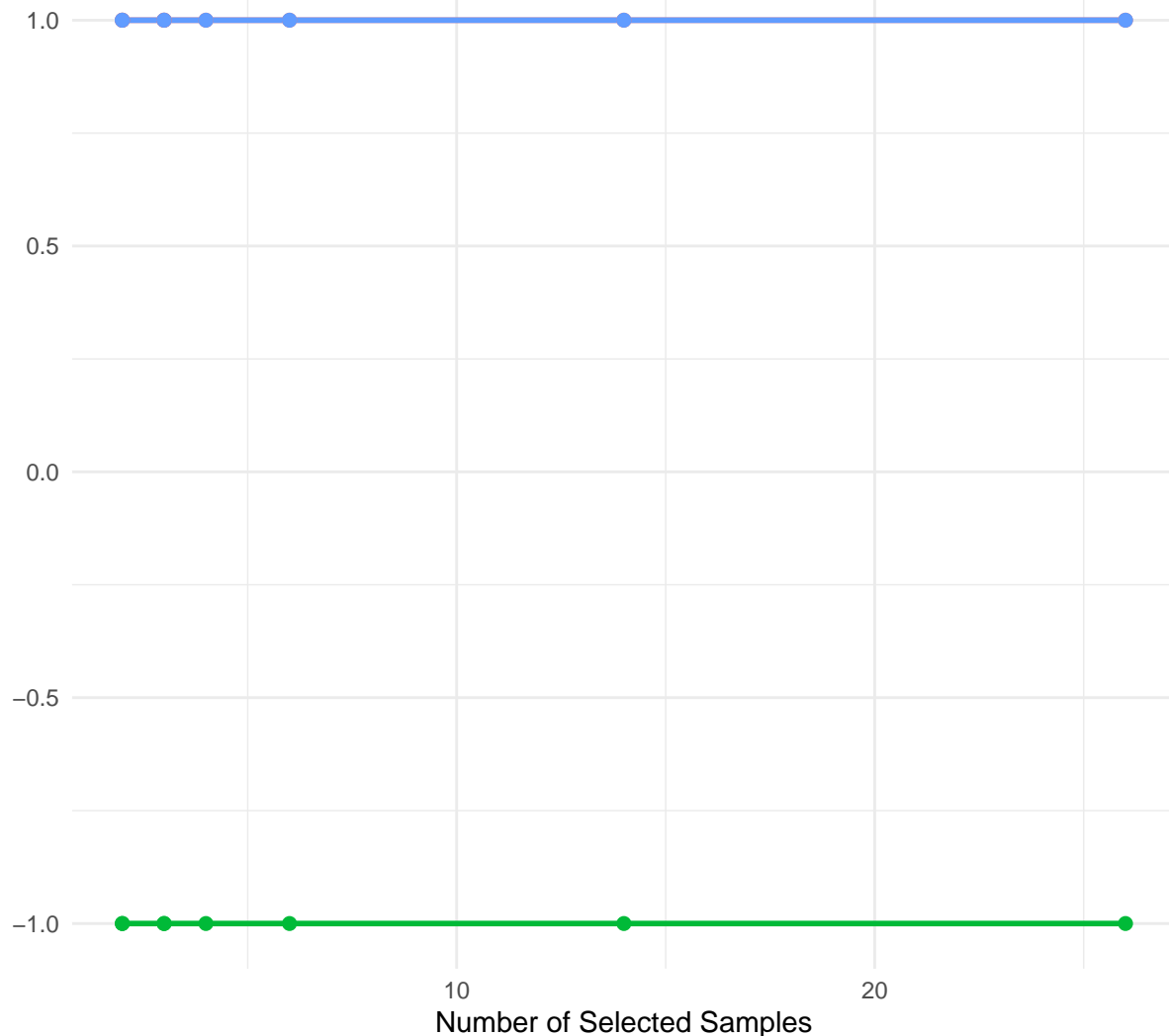


Score: APC (Local, partial2 , Genotype: wave)

Kendall Correlation

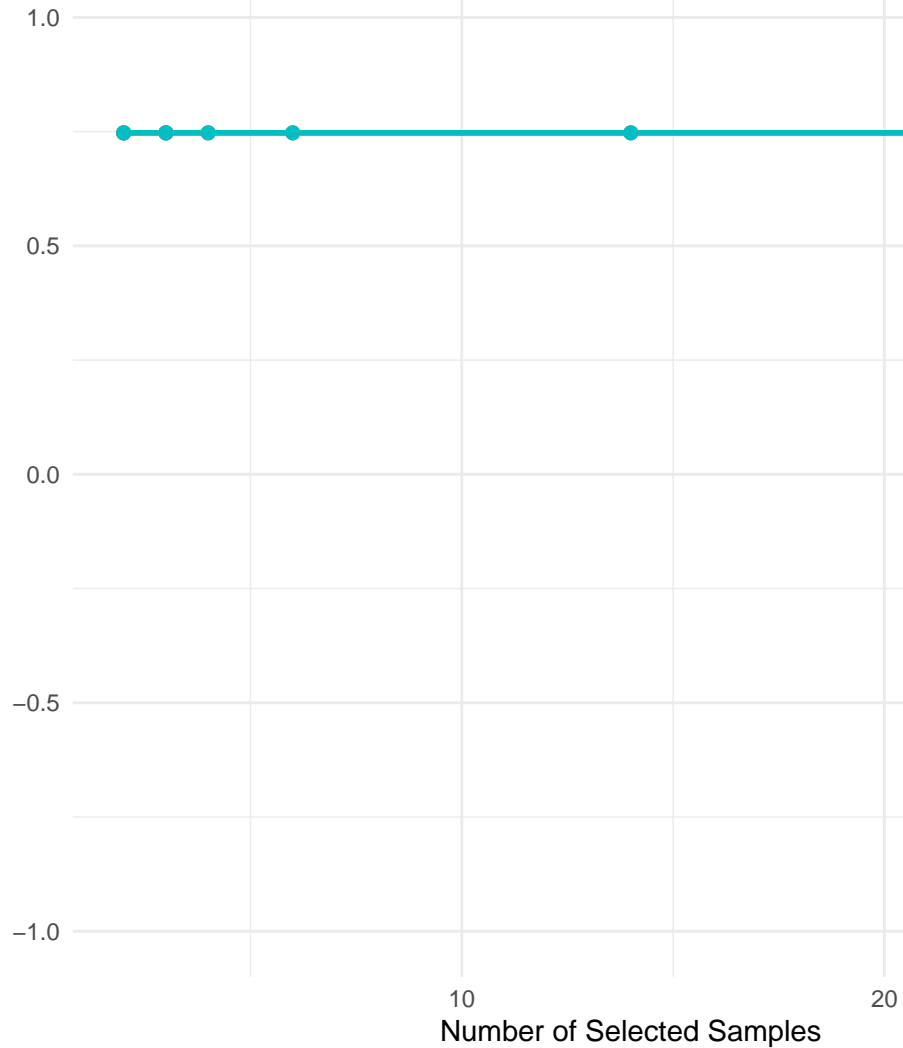
Summary Statistic

- range
- slope
- variance



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

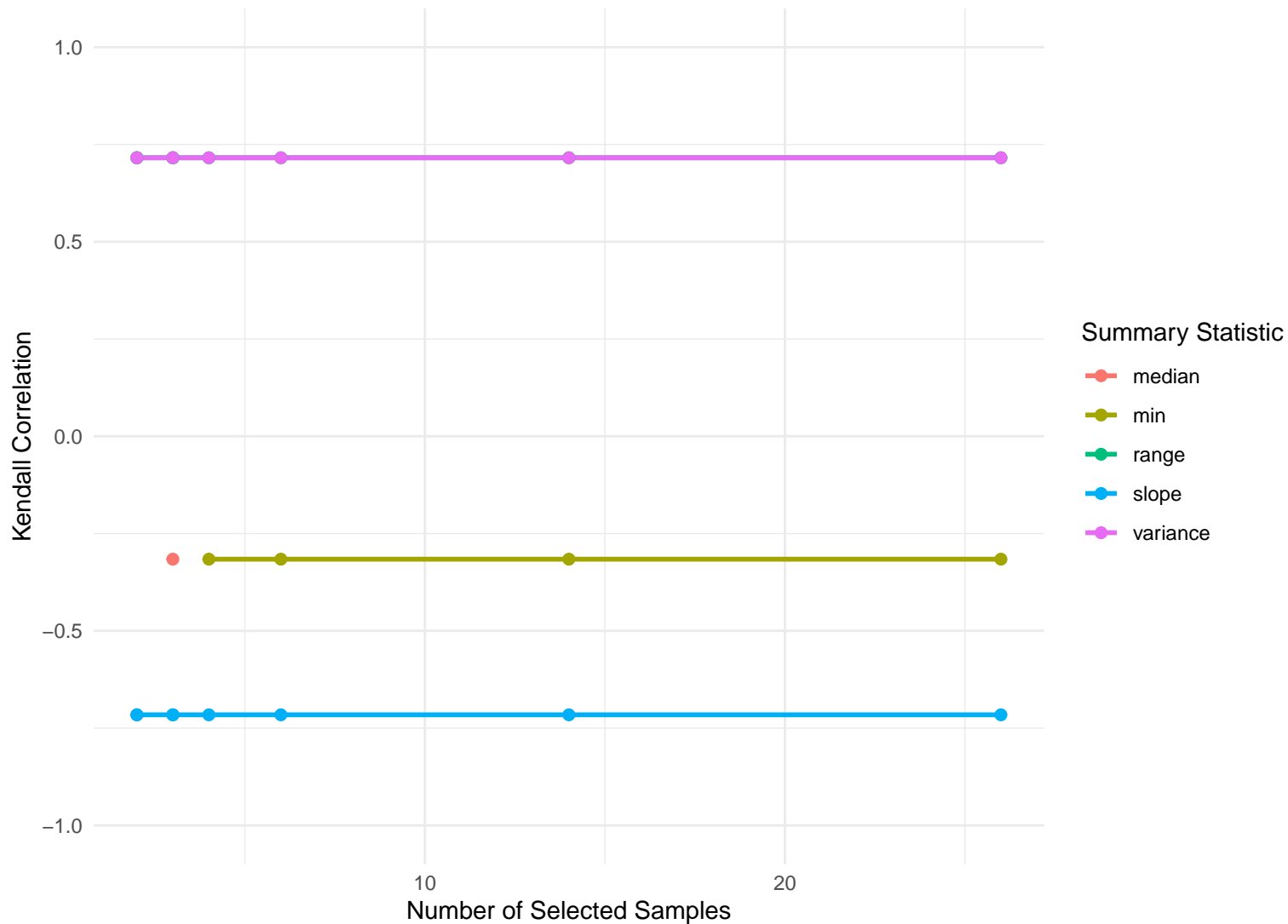
Kendall Correlation



Summary Statistic

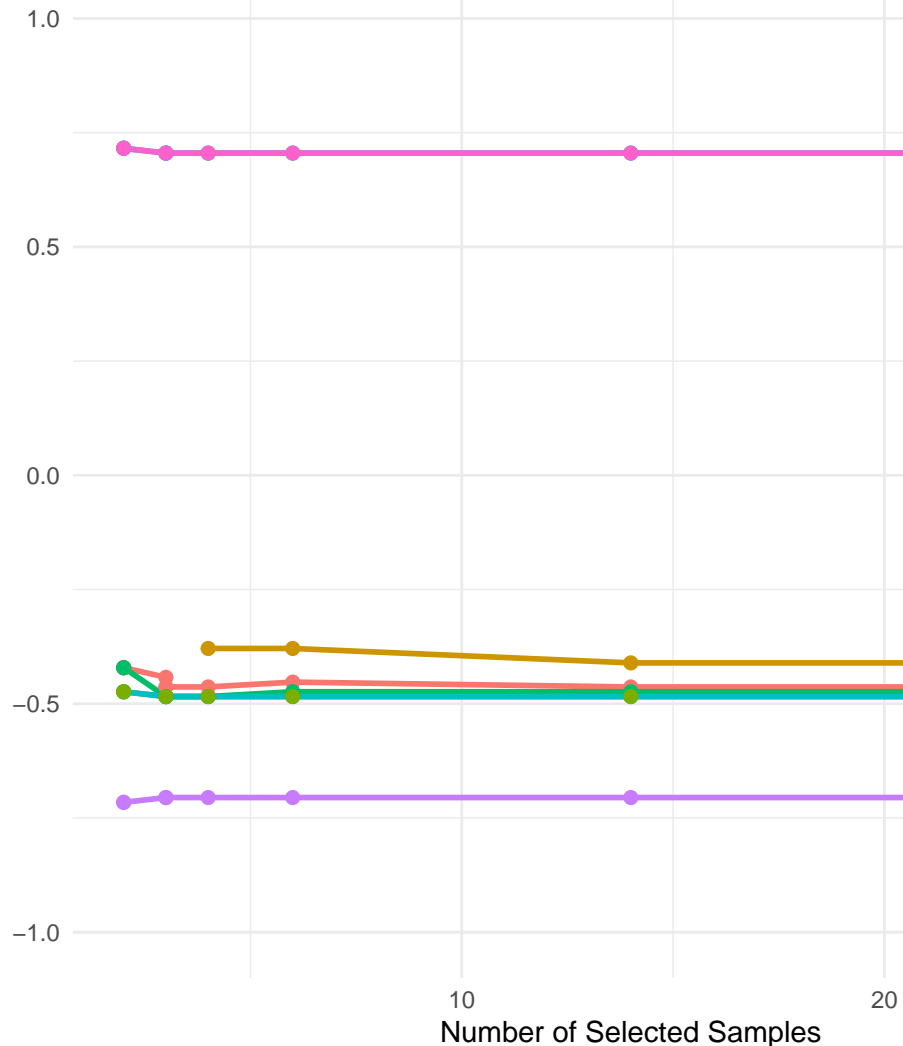
- range
- variance

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



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

Kendall Correlation



Summary Statistic

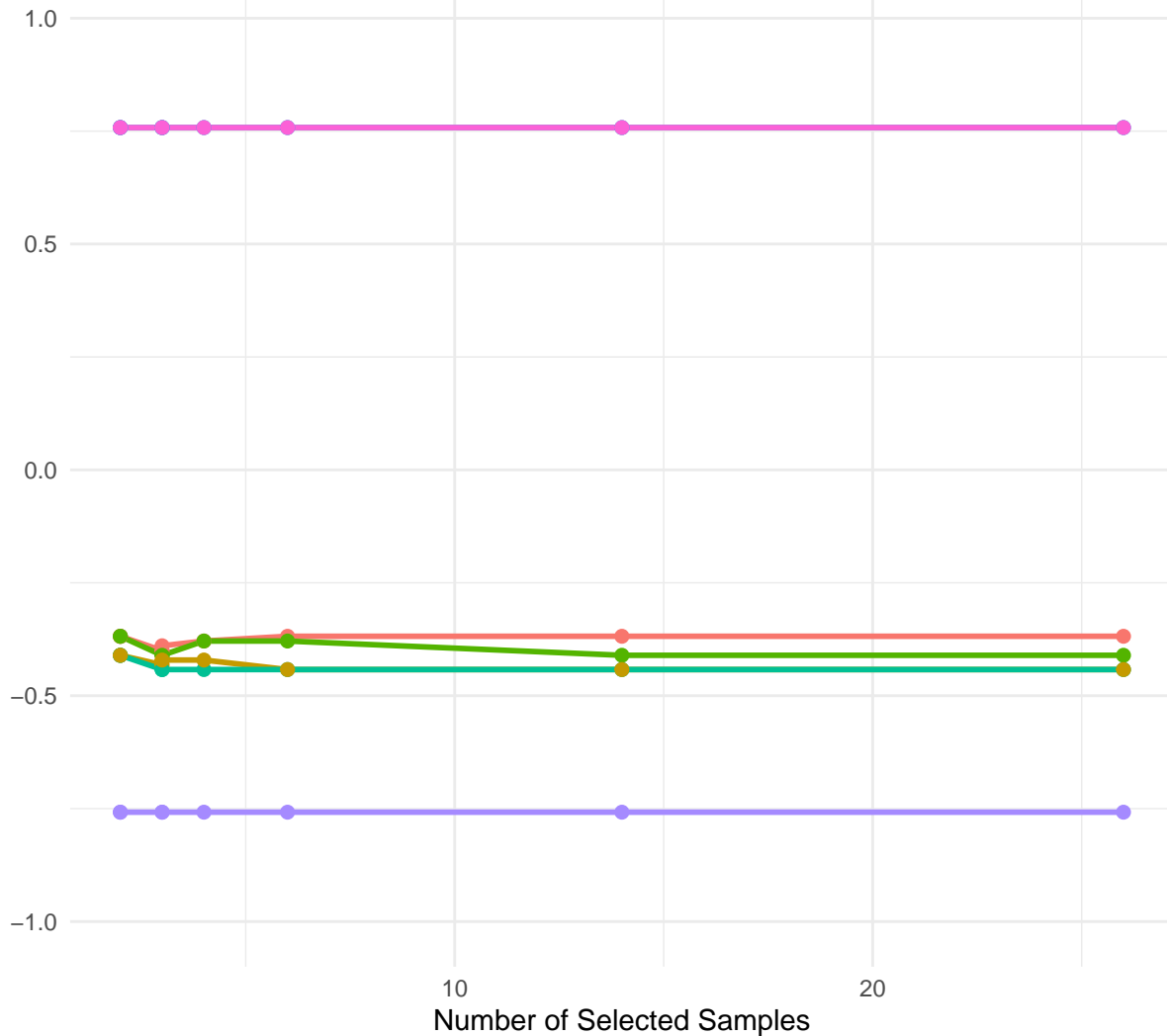
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance

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

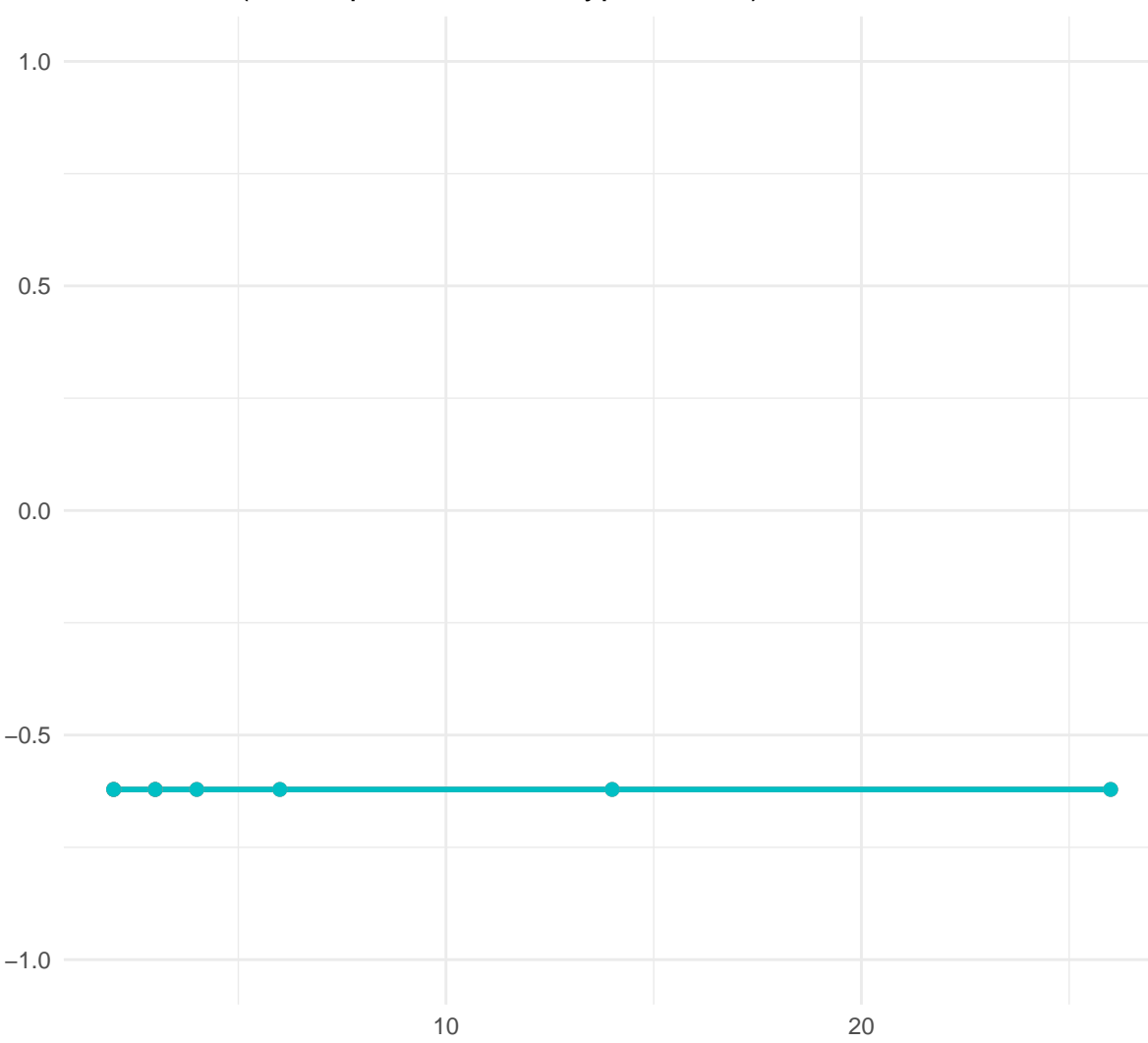
Summary Statistic

range
variance

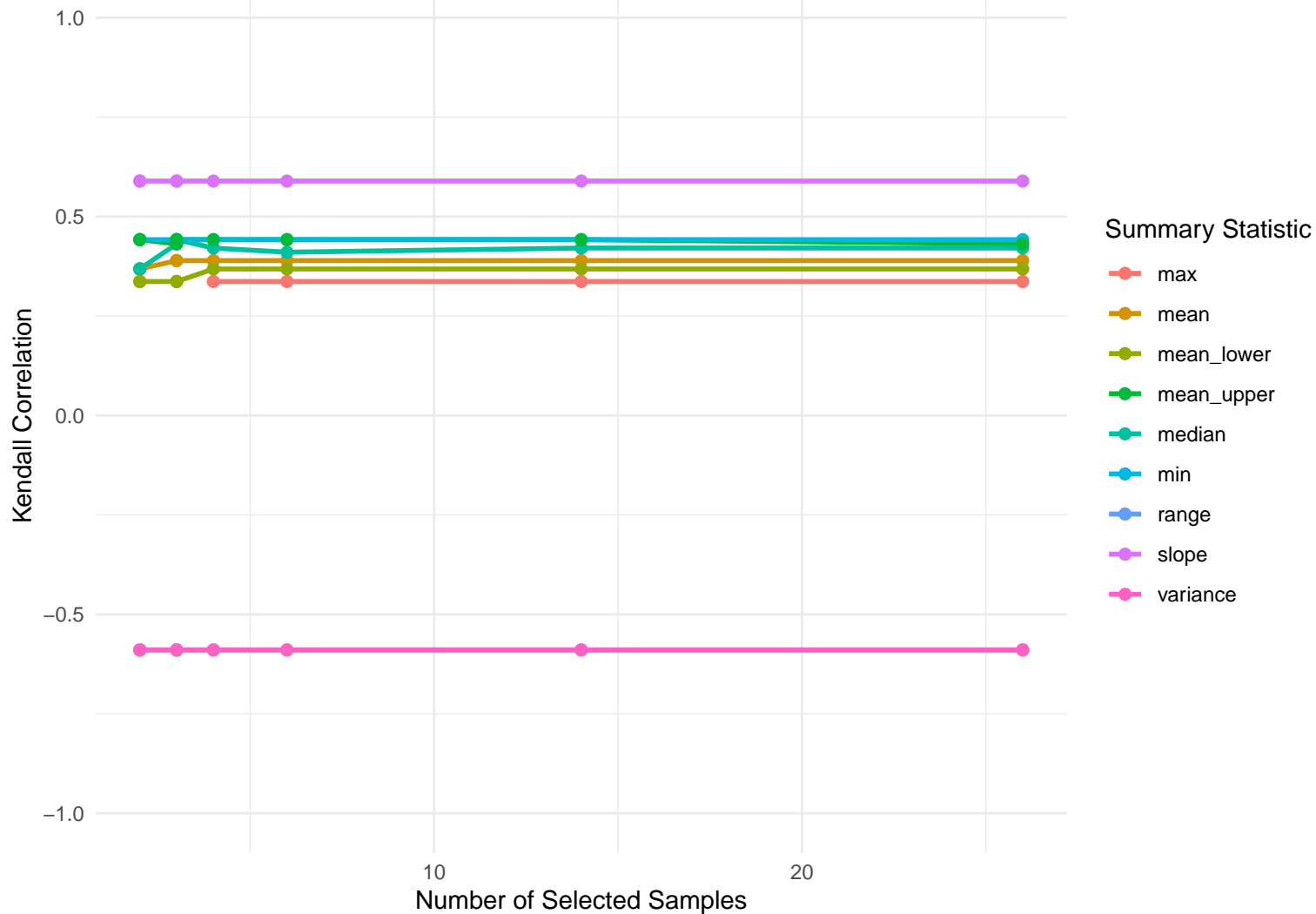
Number of Selected Samples

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



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

Kendall Correlation

1.0

0.5

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-0.5

-1.0

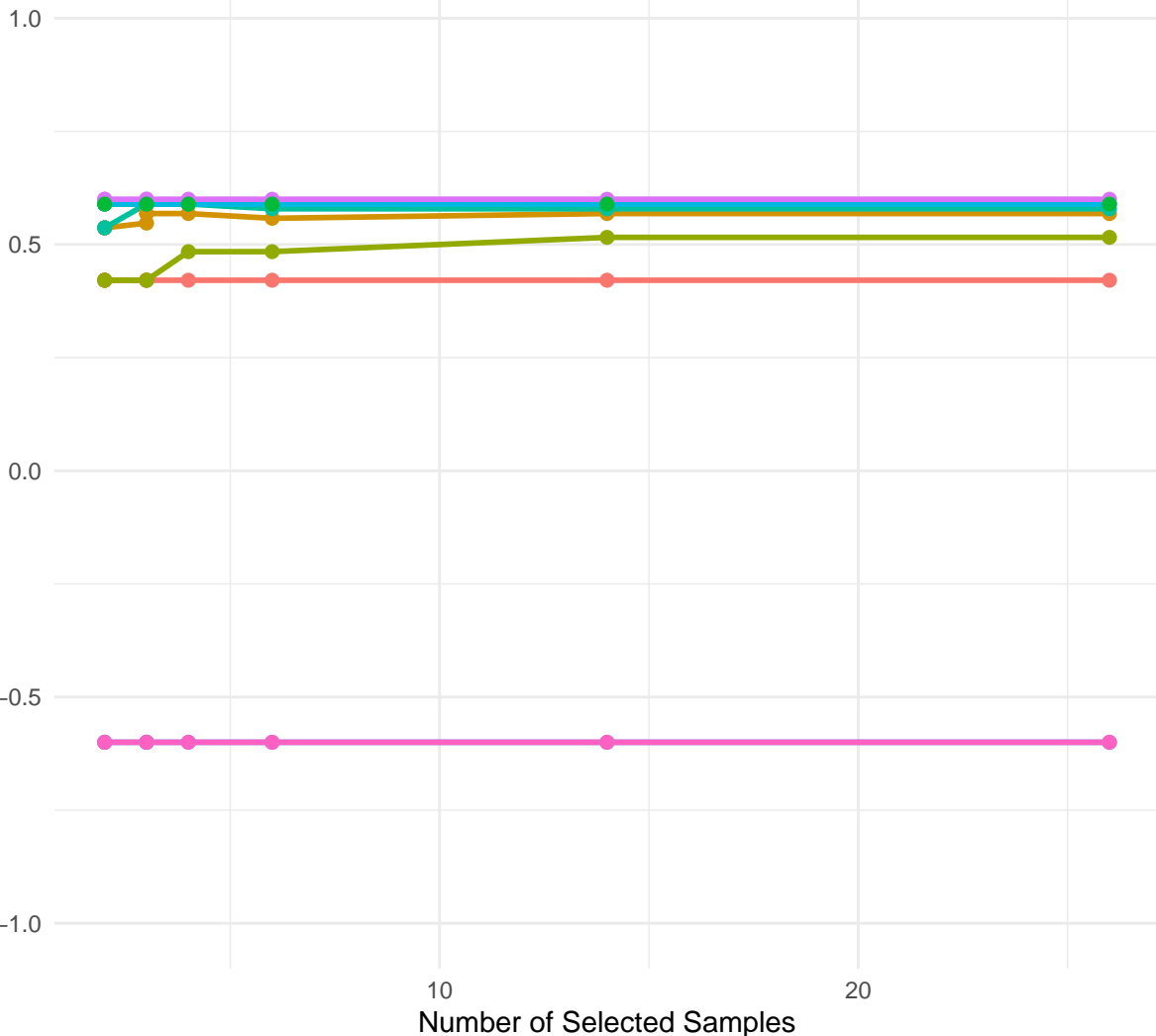
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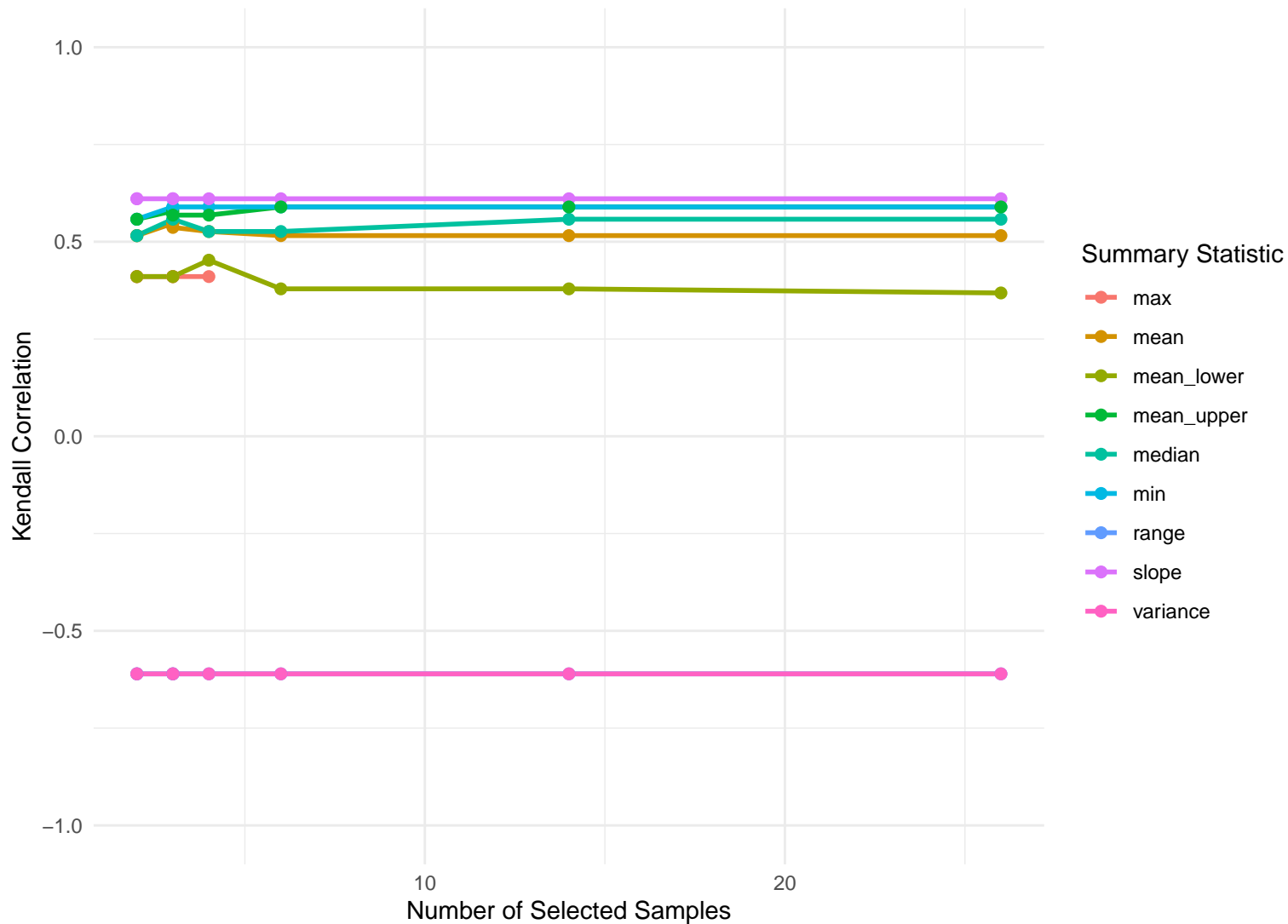
Number of Selected Samples

Summary Statistic

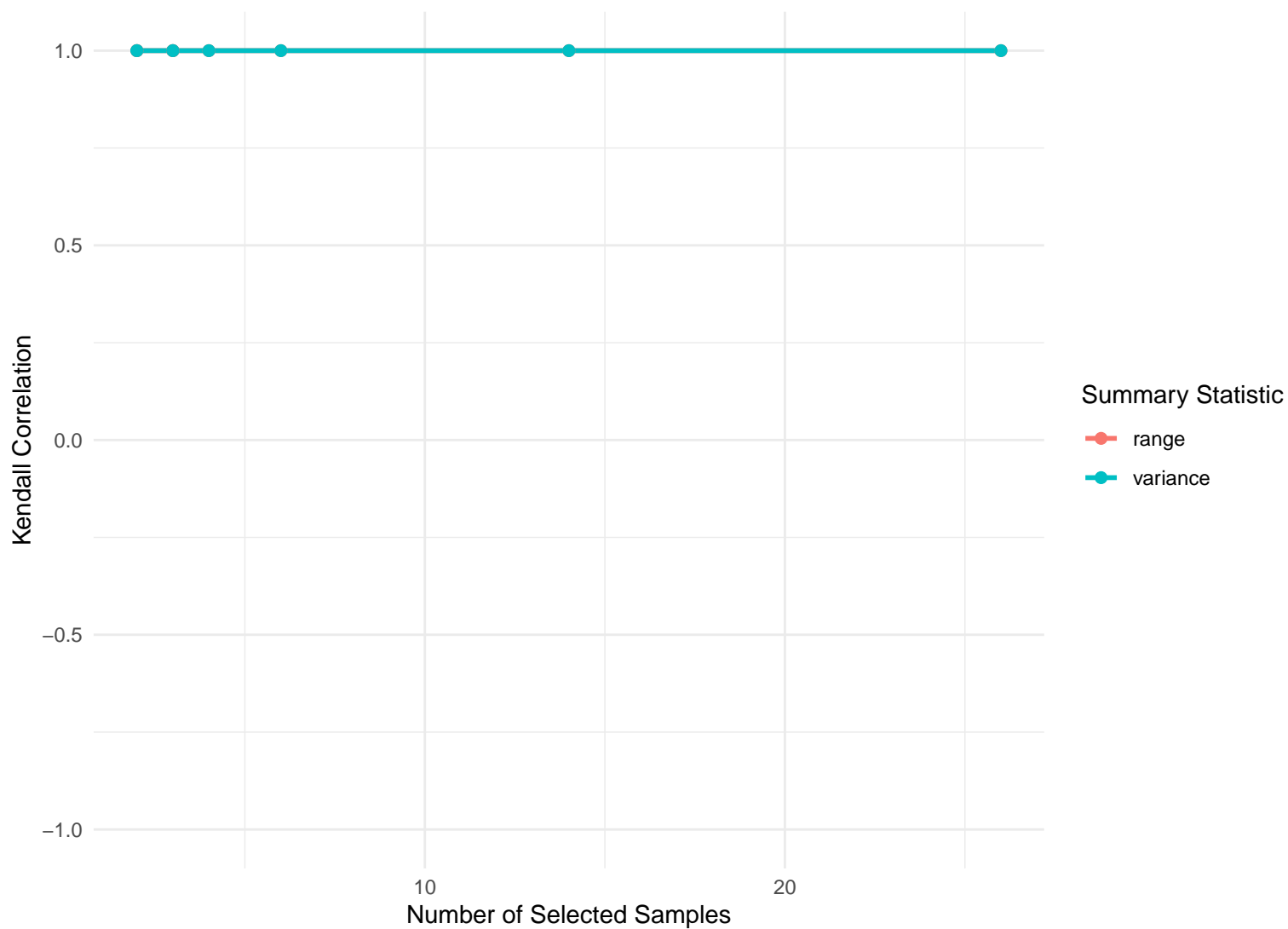
- max
- mean
- mean_lower
- mean_upper
- median
- min
- range
- slope
- variance



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

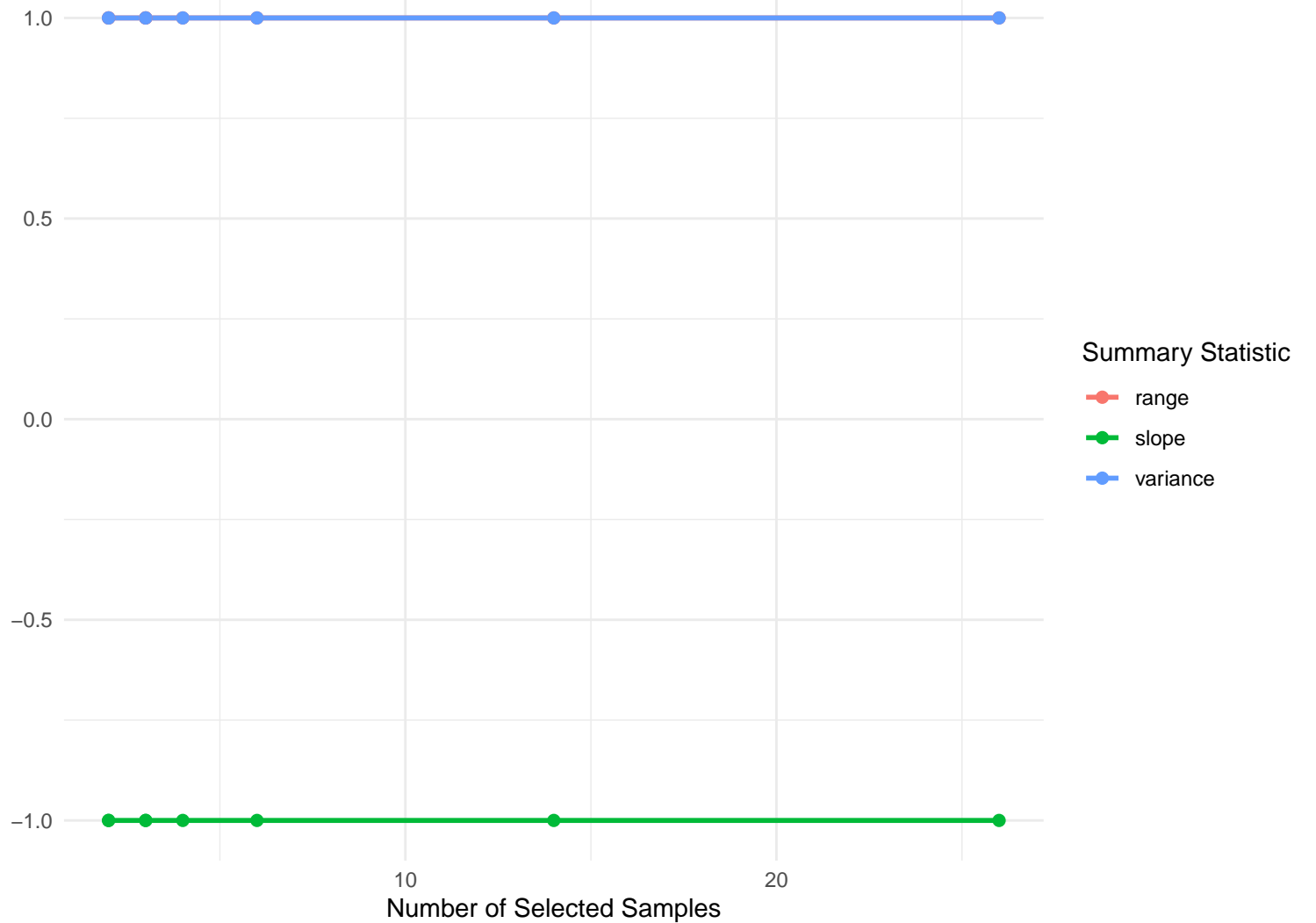
10

20

Number of Selected Samples

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

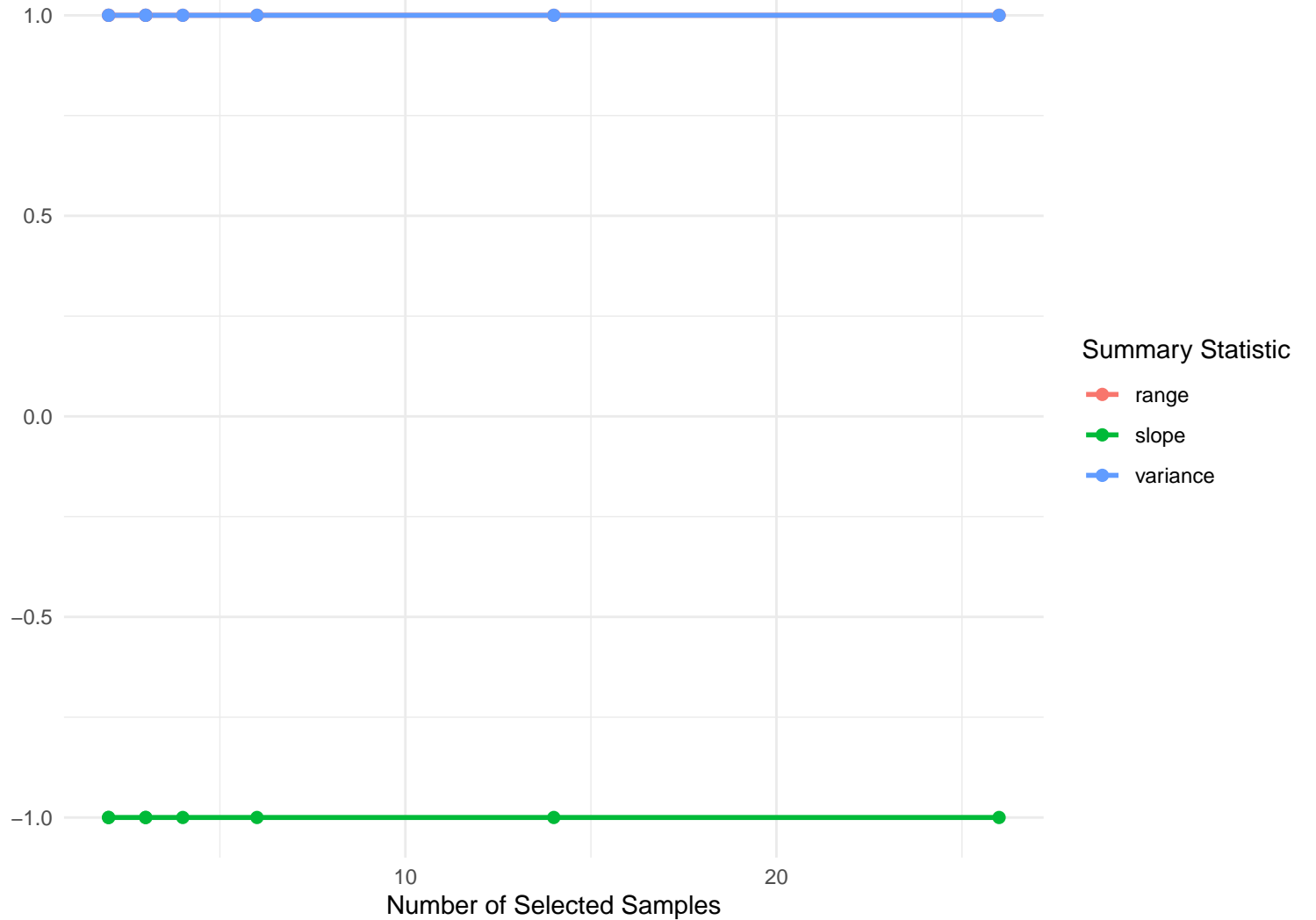
10

20

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

10

20

Number of Selected Samples

Summary Statistic

- range
- slope
- variance

