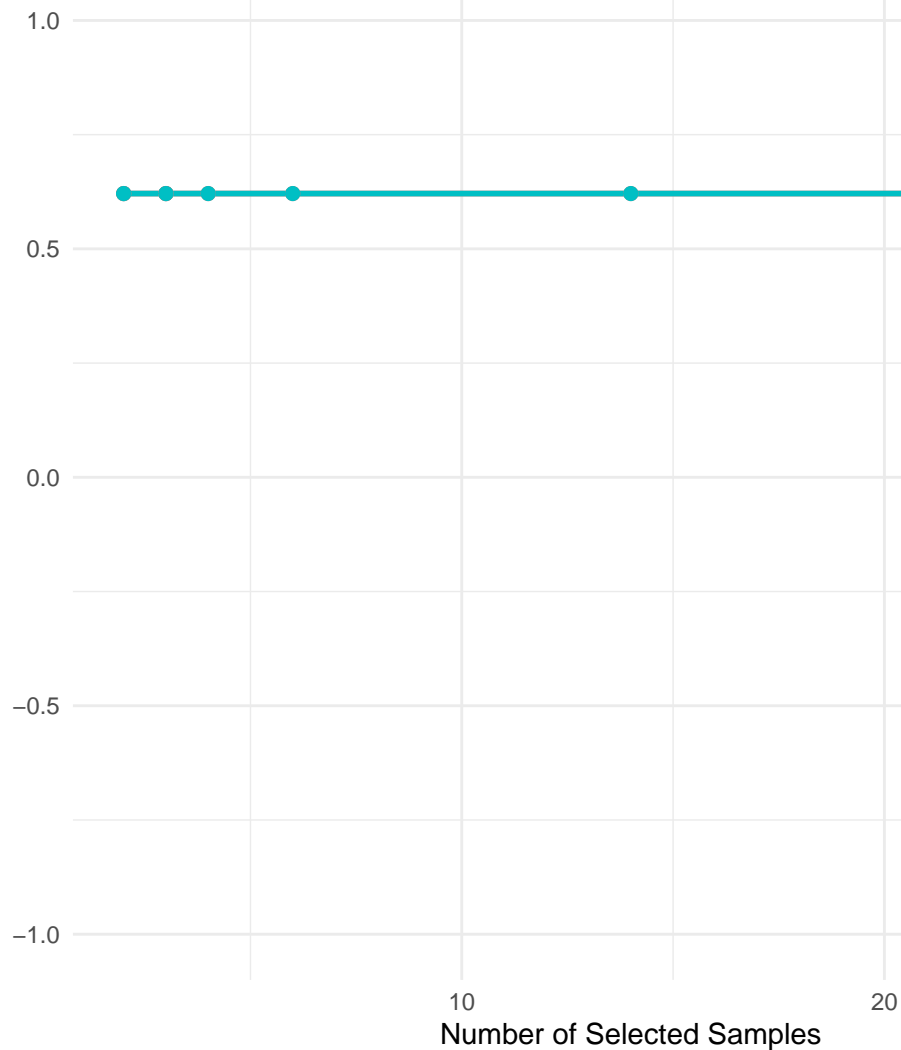


Score: CV\_t (Local, partial2 , Genotype: linear )

Kendall Correlation



Summary Statistic

- range
- variance

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

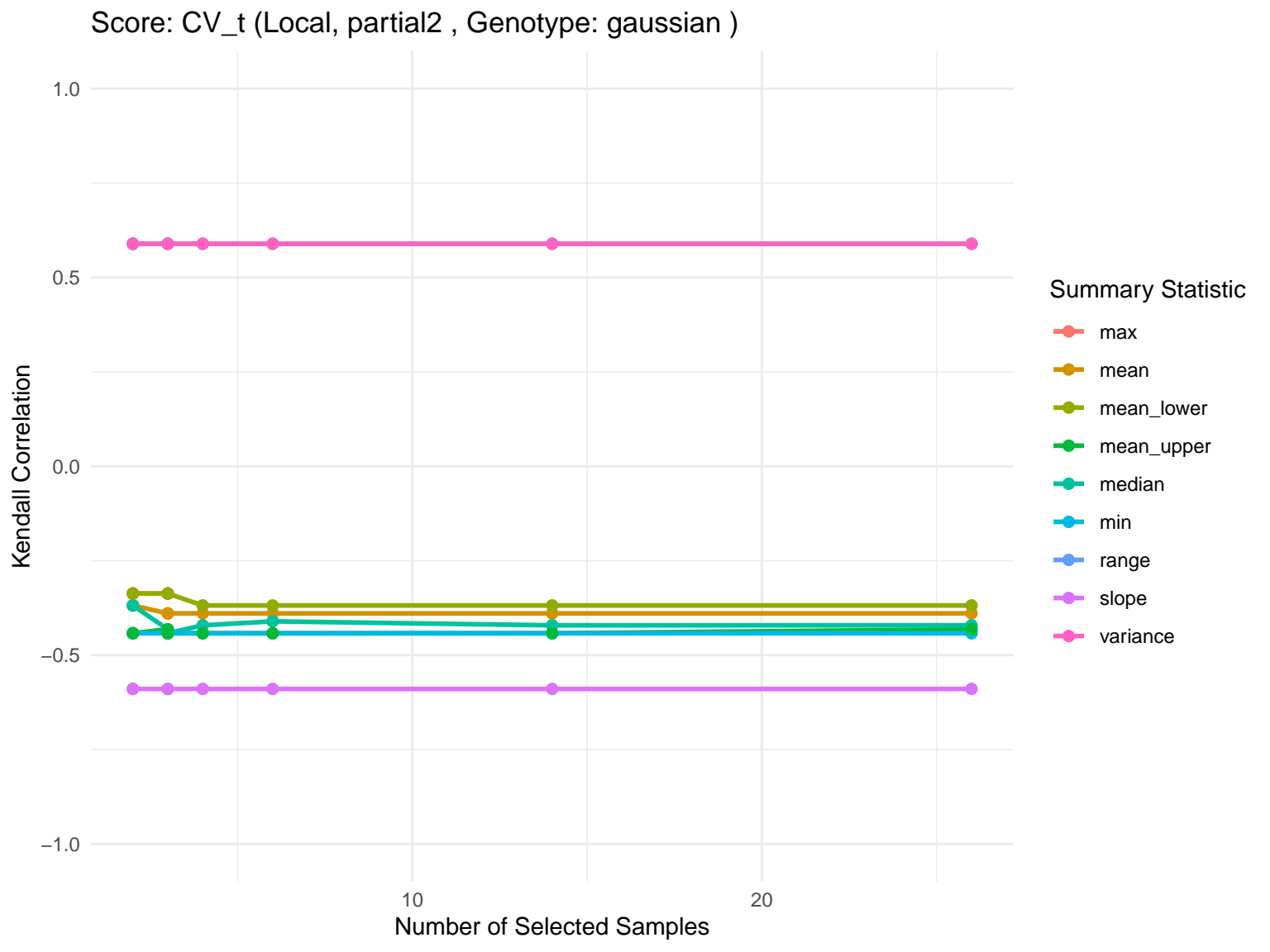
-0.5

-1.0

10

20

Number of Selected Samples

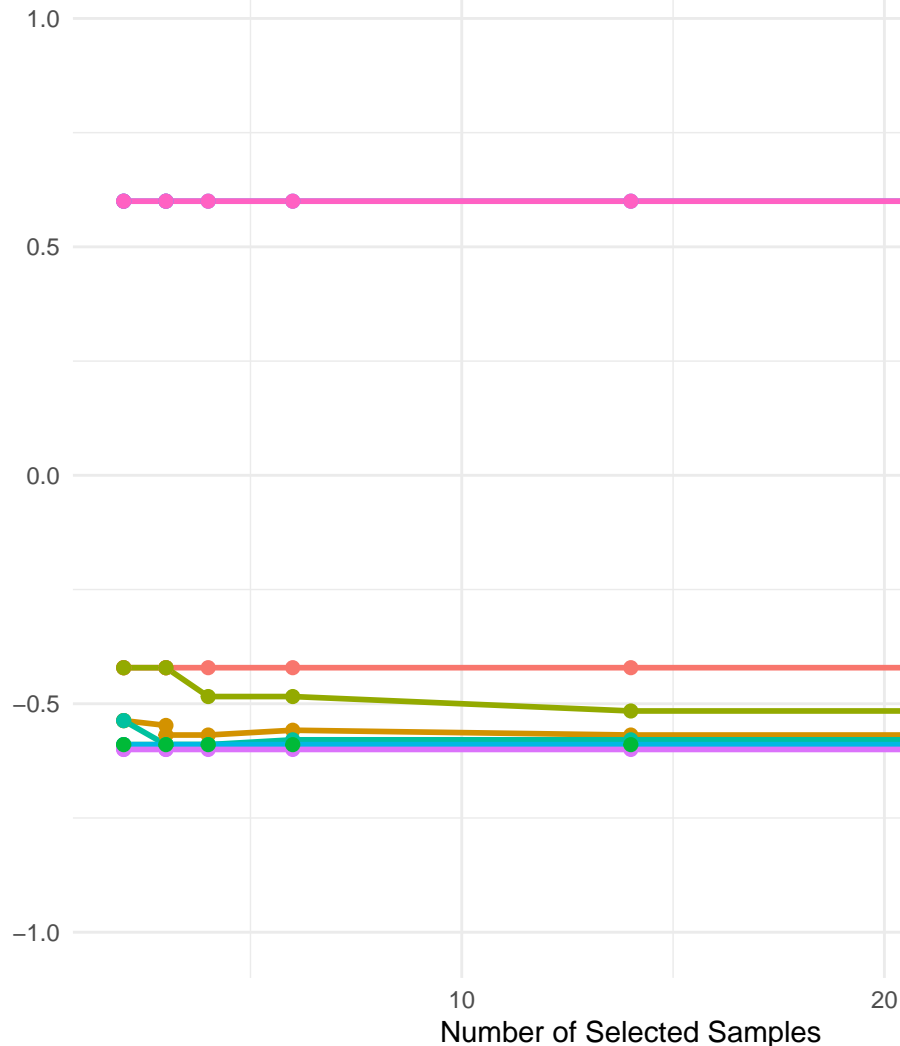


Score: CV\_t (Local, partial2 , Genotype: sinusoidal )

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

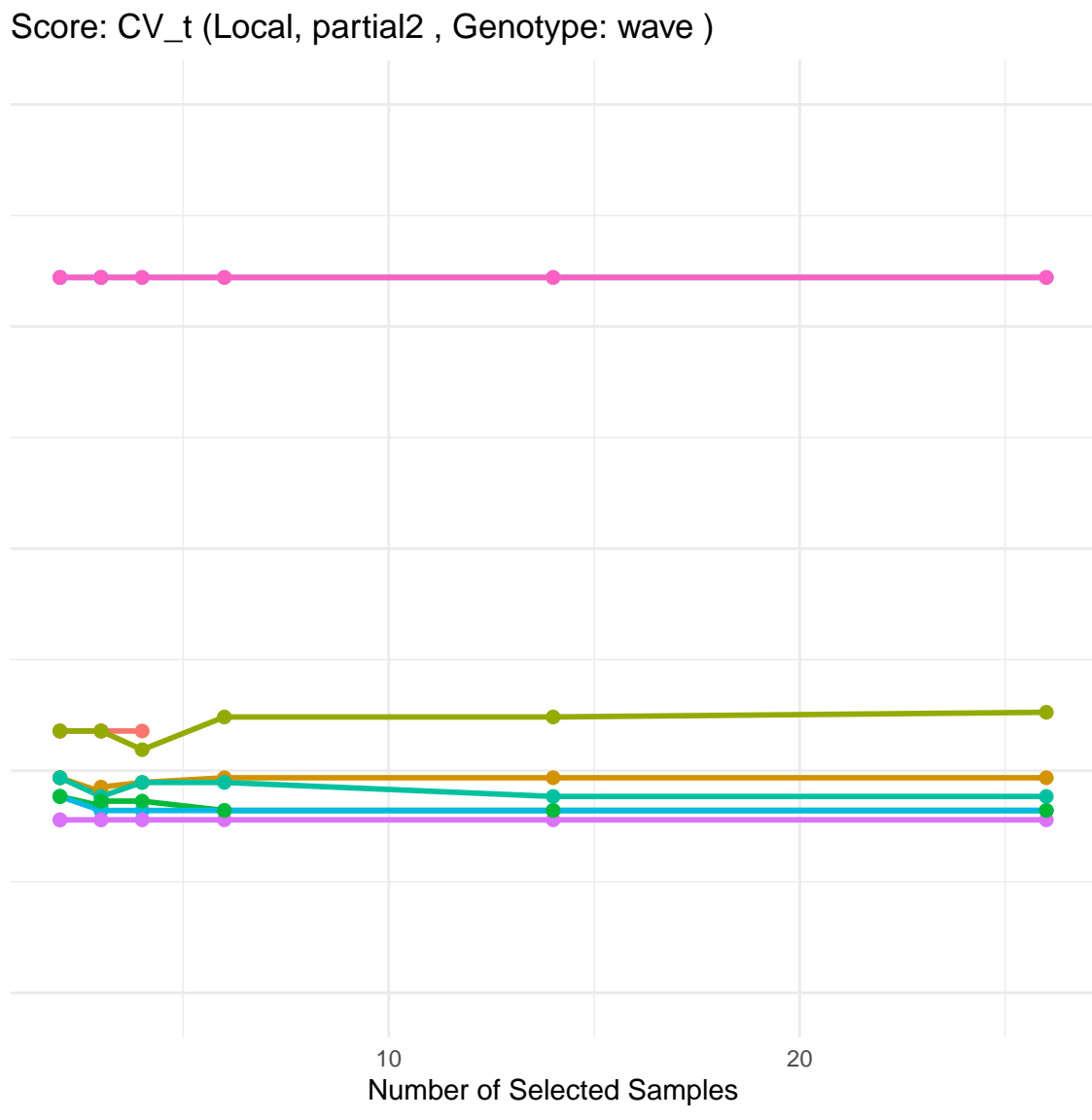
-0.5

-1.0

Number of Selected Samples

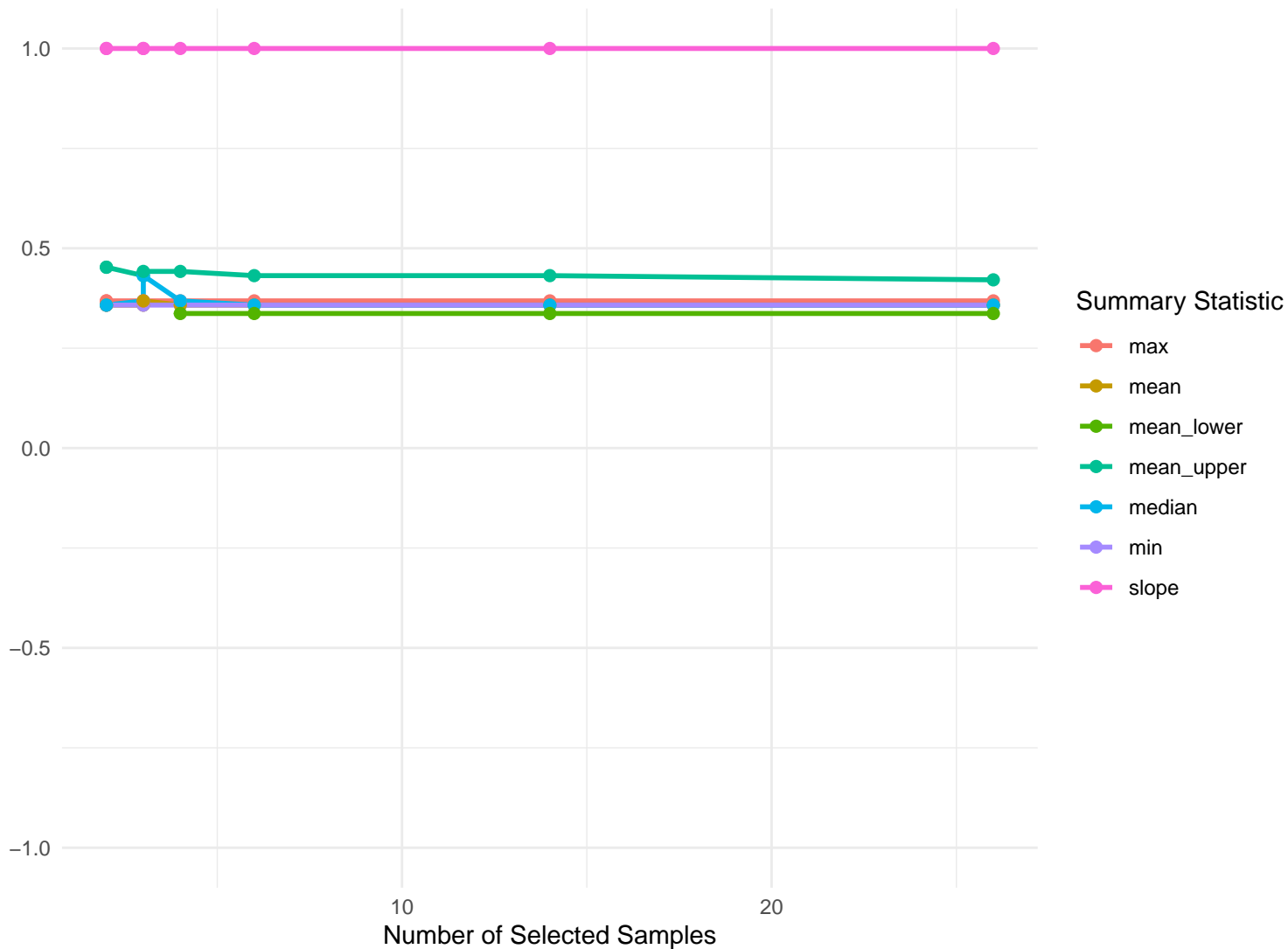
10

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

Number of Selected Samples

10

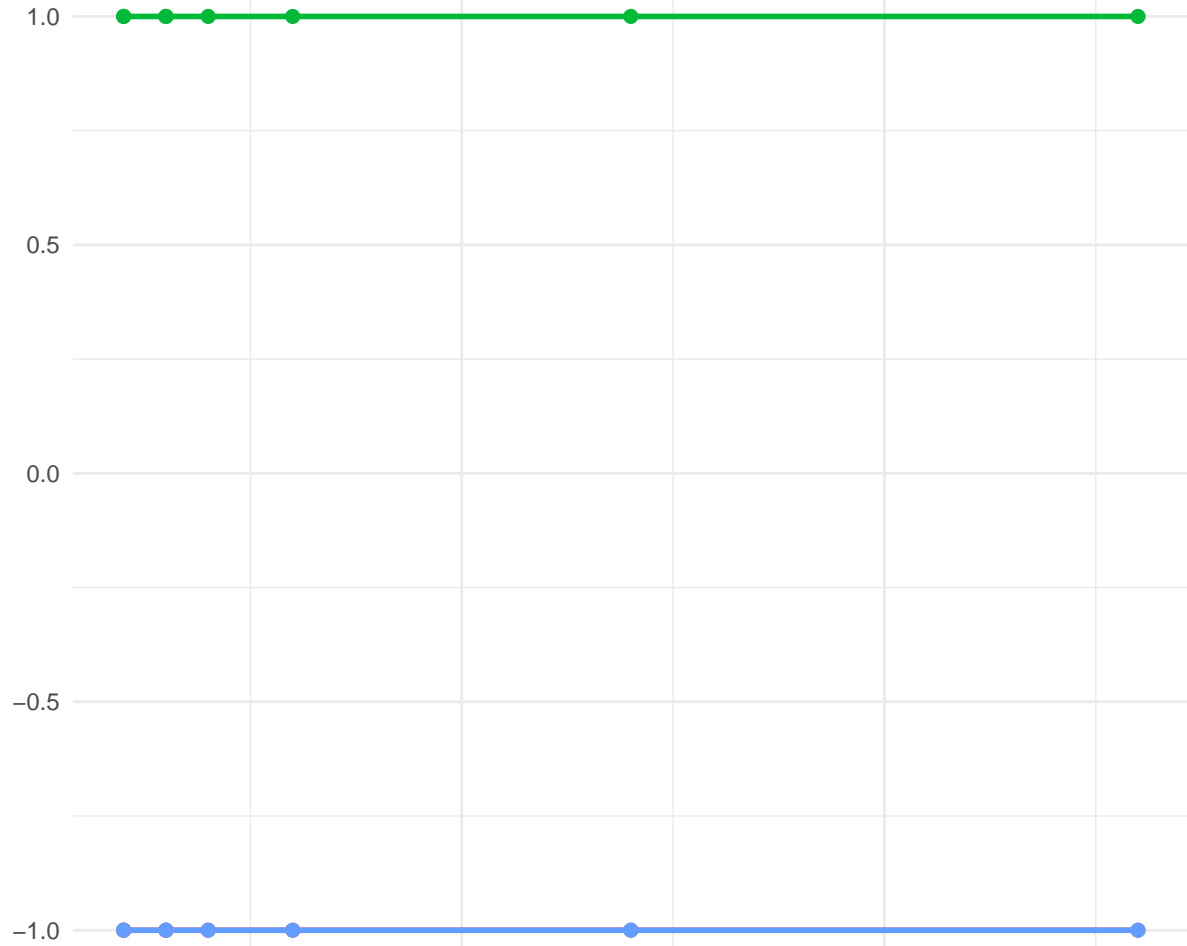
20

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

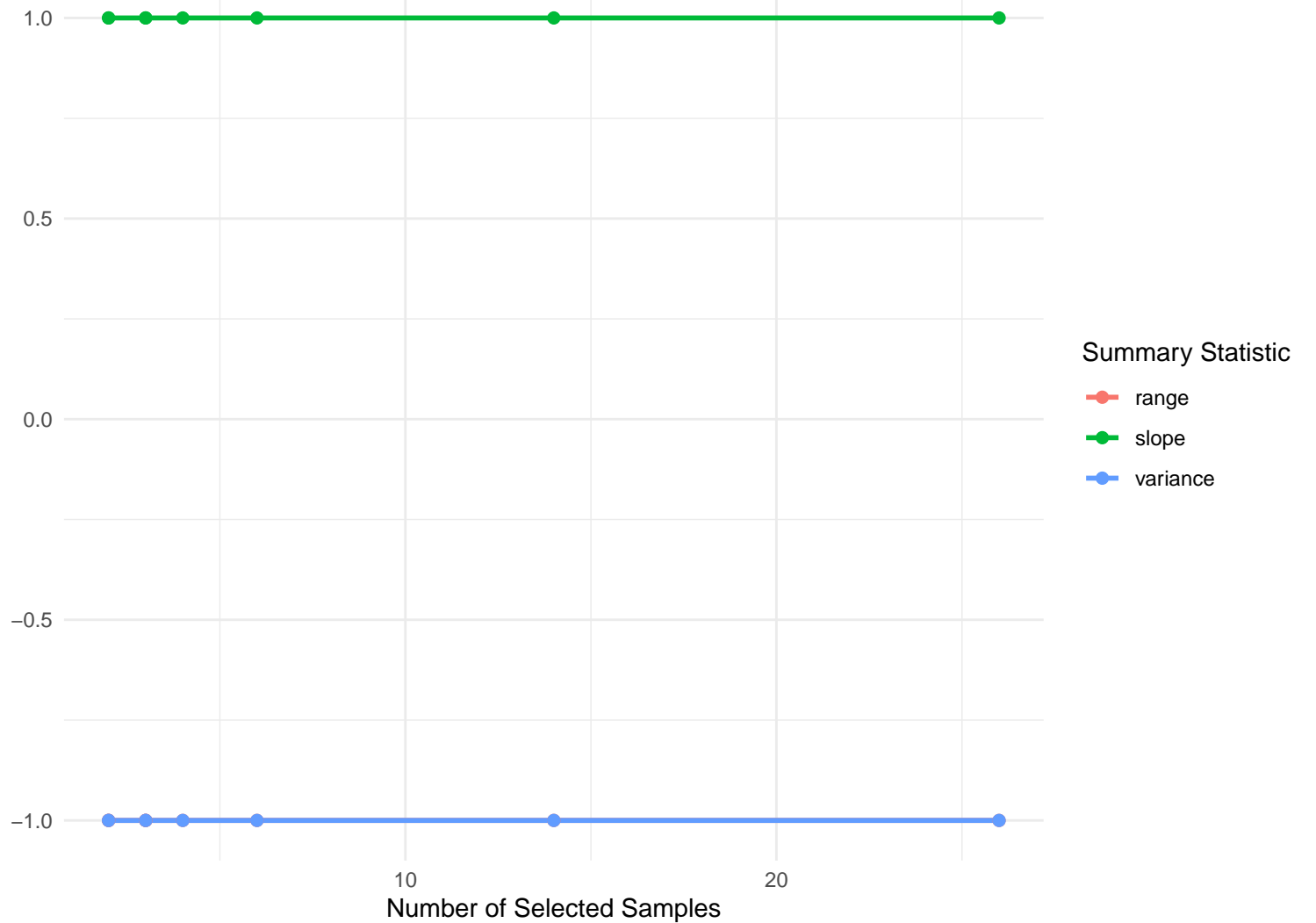
Number of Selected Samples

Summary Statistic

range

slope

variance

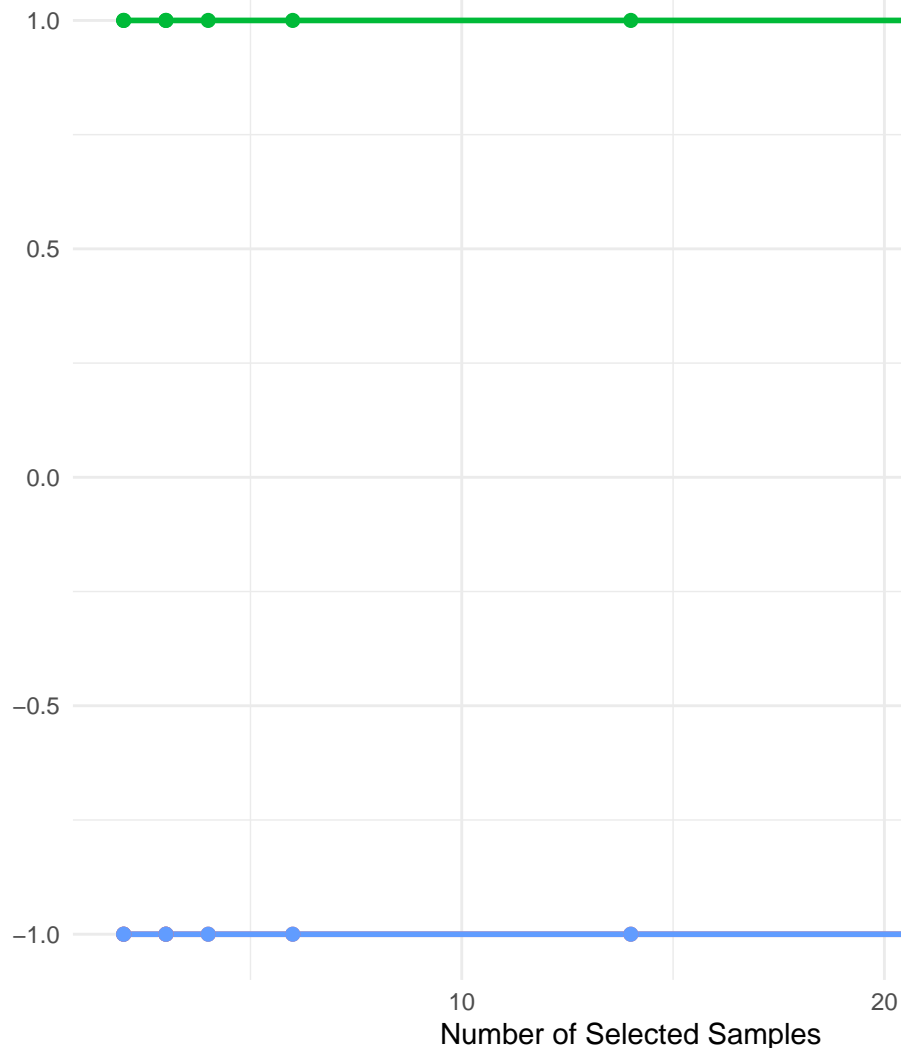


Score: RN (Local, partial2 , Genotype: wave )

Kendall Correlation

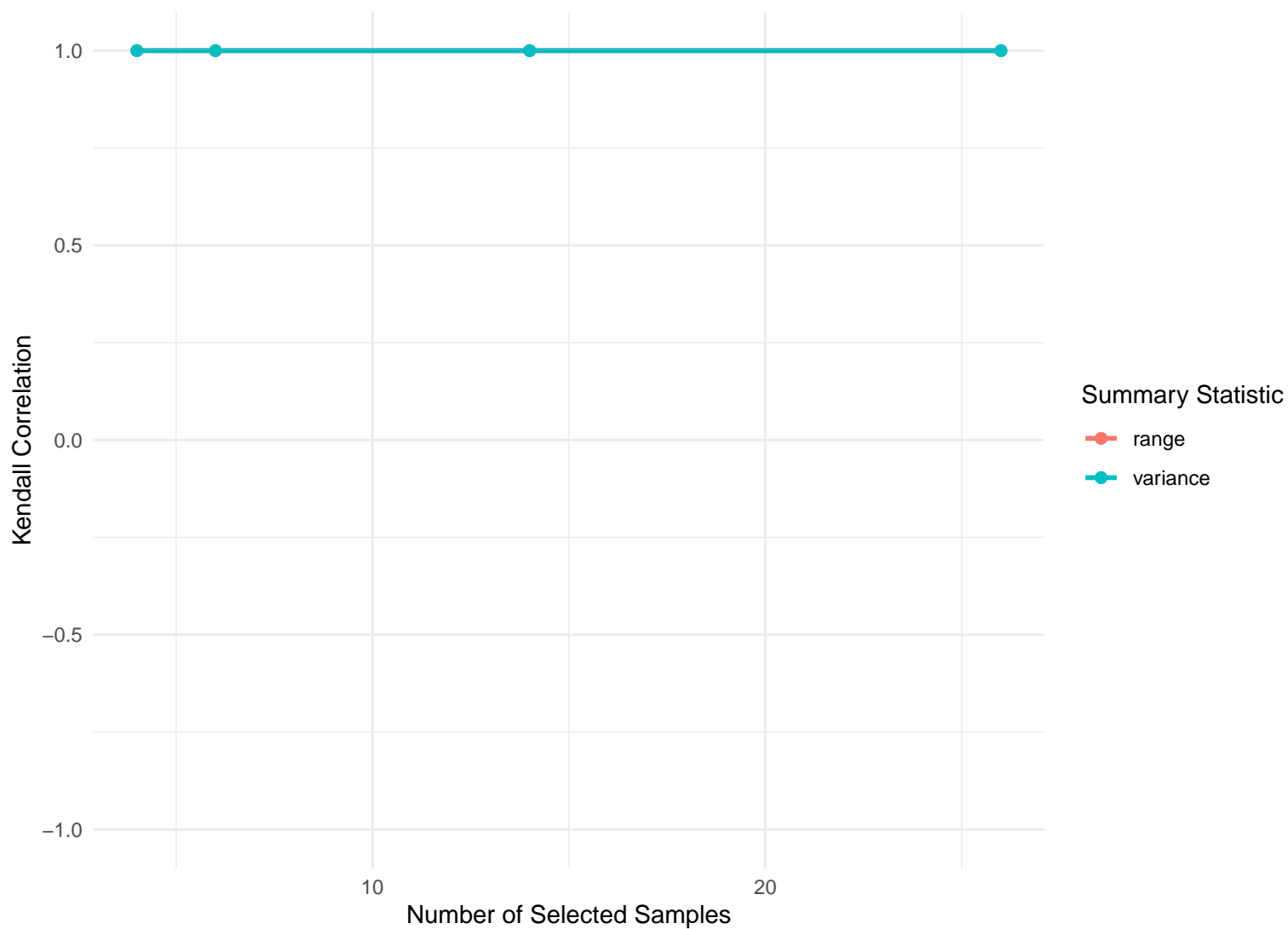
Summary Statistic

- 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

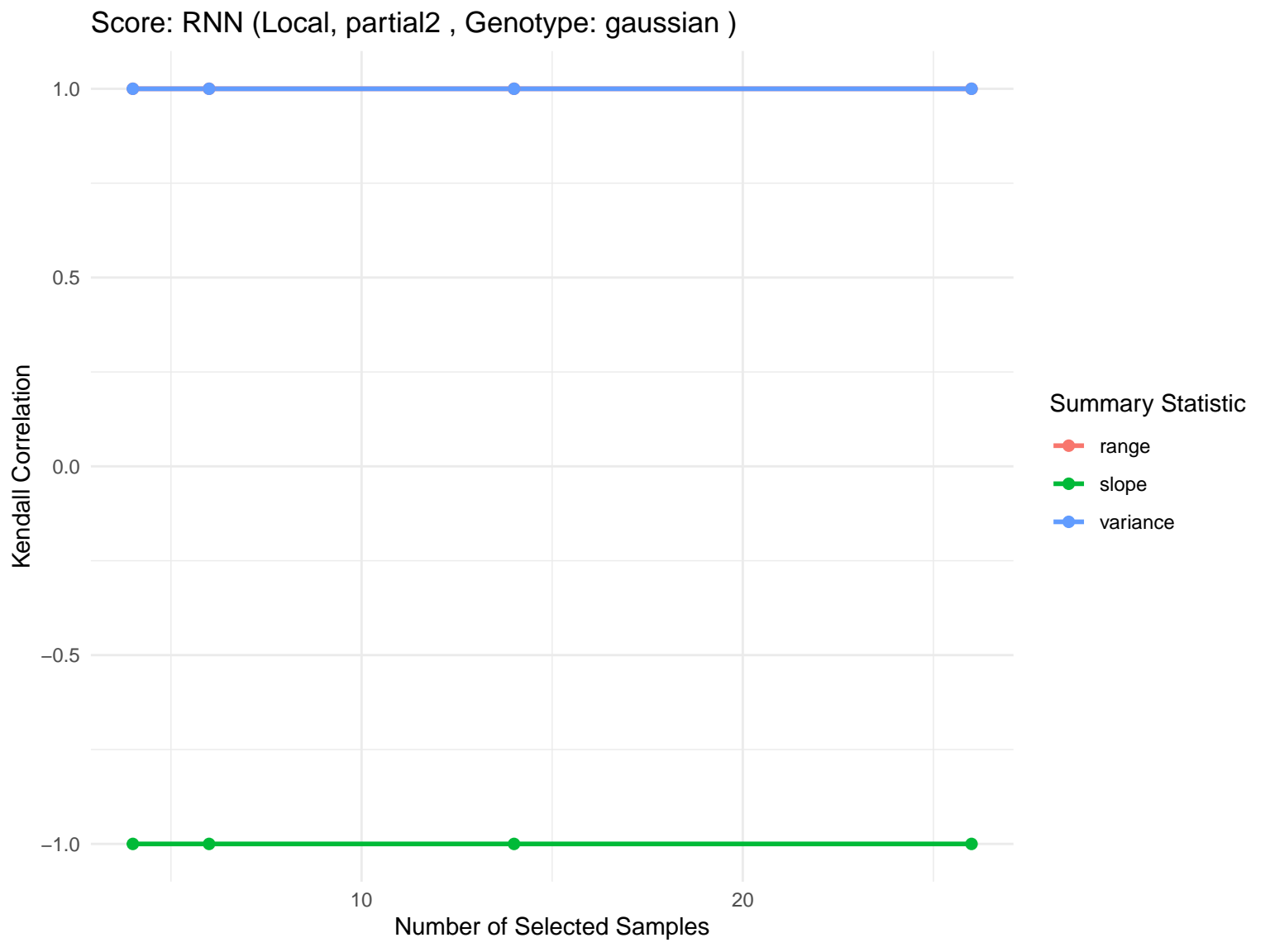
10

20

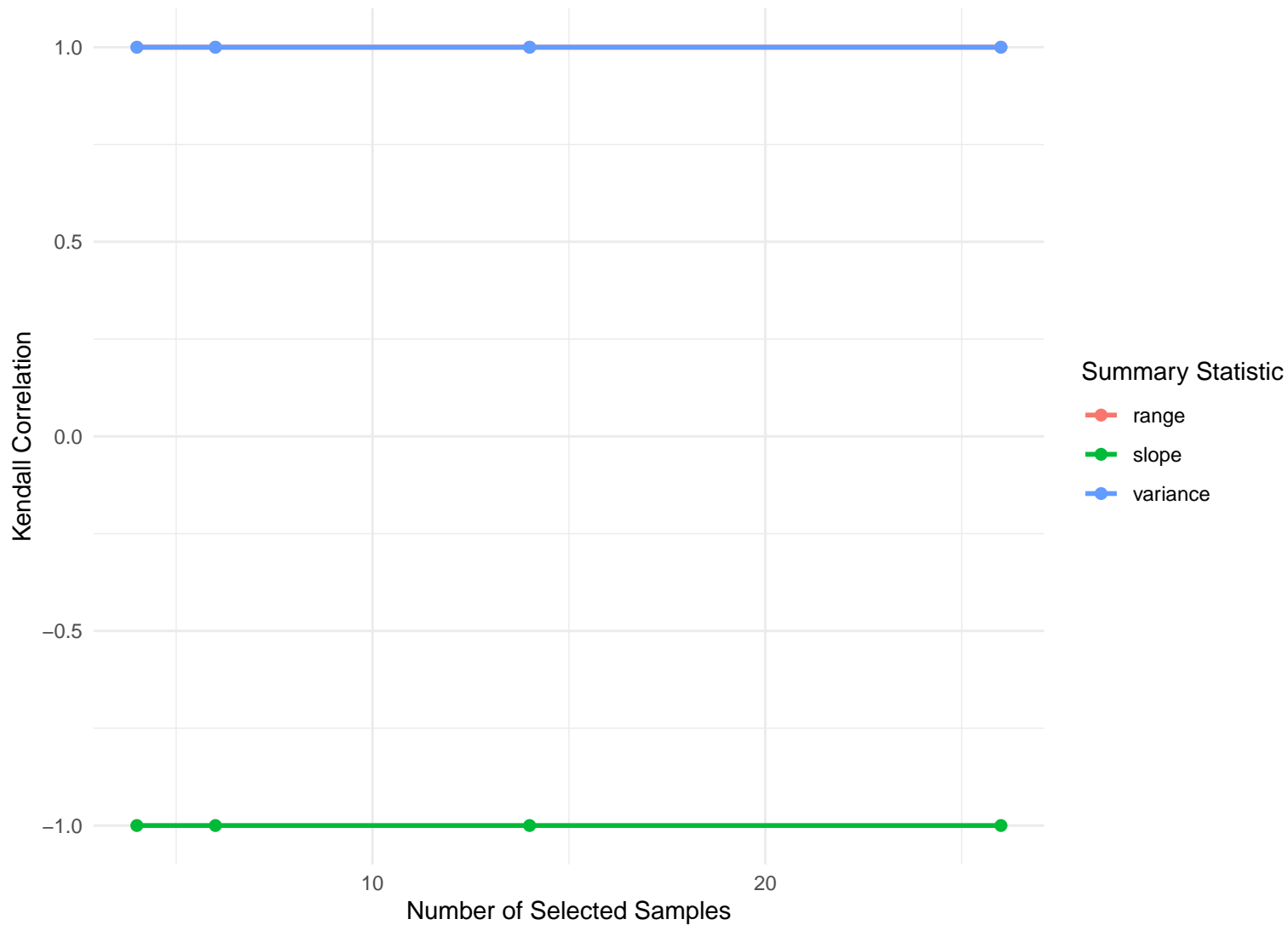
Number of Selected Samples

Summary Statistic

- range
- slope
- variance



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

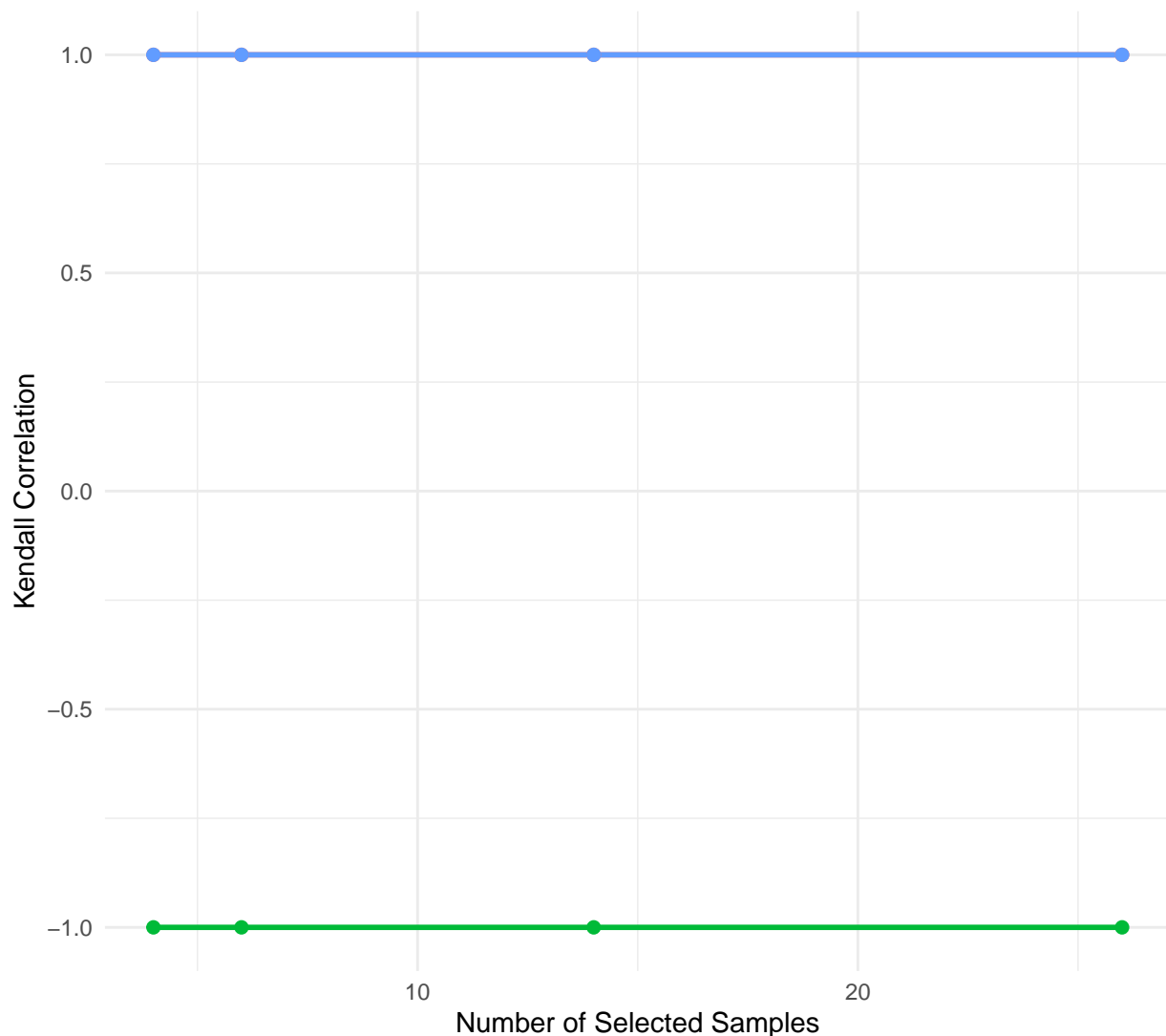
10

20

Number of Selected Samples

Summary Statistic

- range
- slope
- variance



Score: D\_slope (Local, partial2 , Genotype: linear )

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

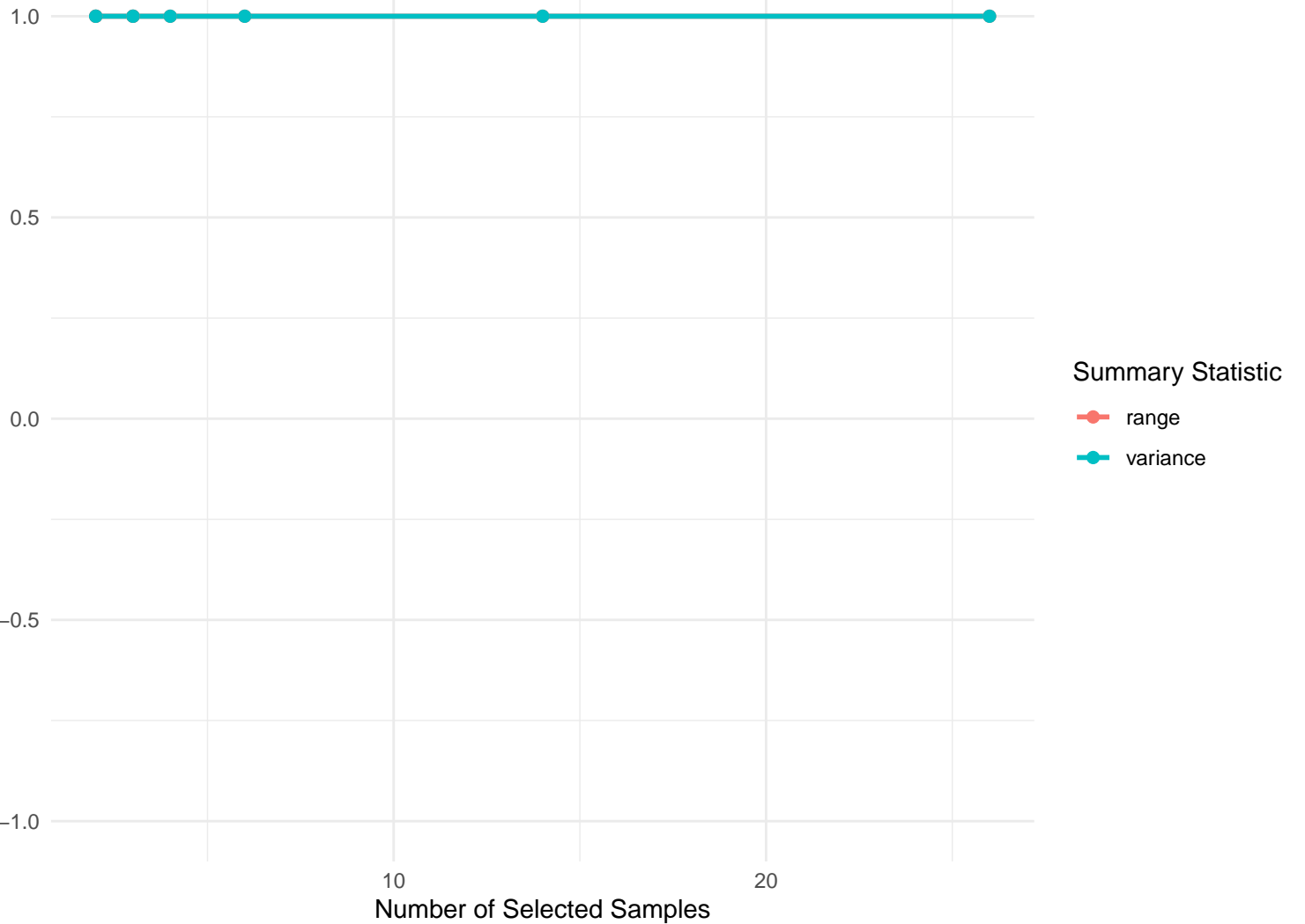
20

Number of Selected Samples

Summary Statistic

range

variance

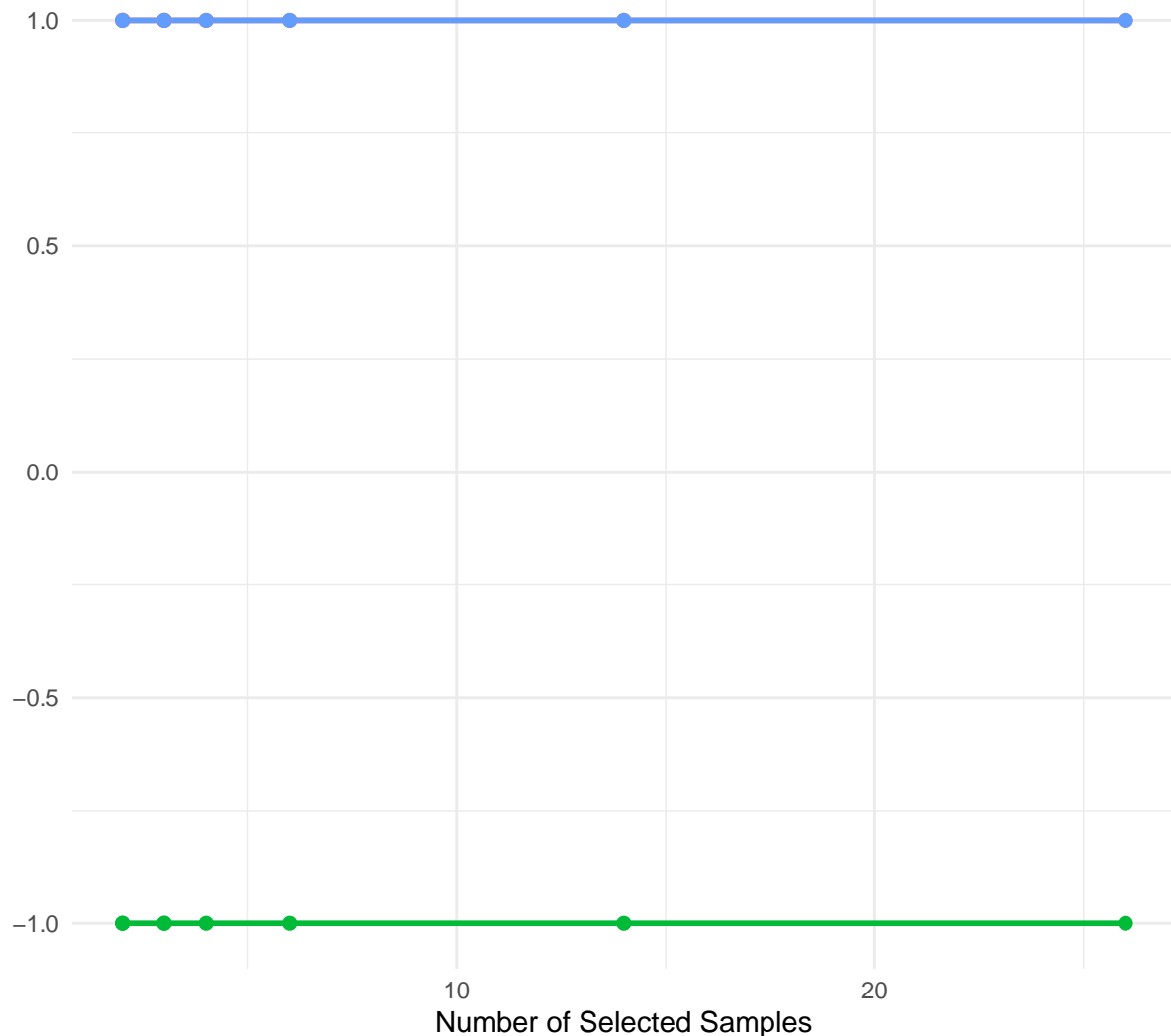


Score: D\_slope (Local, partial2 , Genotype: gaussian )

Kendall Correlation

Summary Statistic

- range
- slope
- variance



Score: D\_slope (Local, partial2 , Genotype: sinusoidal )

Kendall Correlation

1.0  
0.5  
0.0  
-0.5  
-1.0

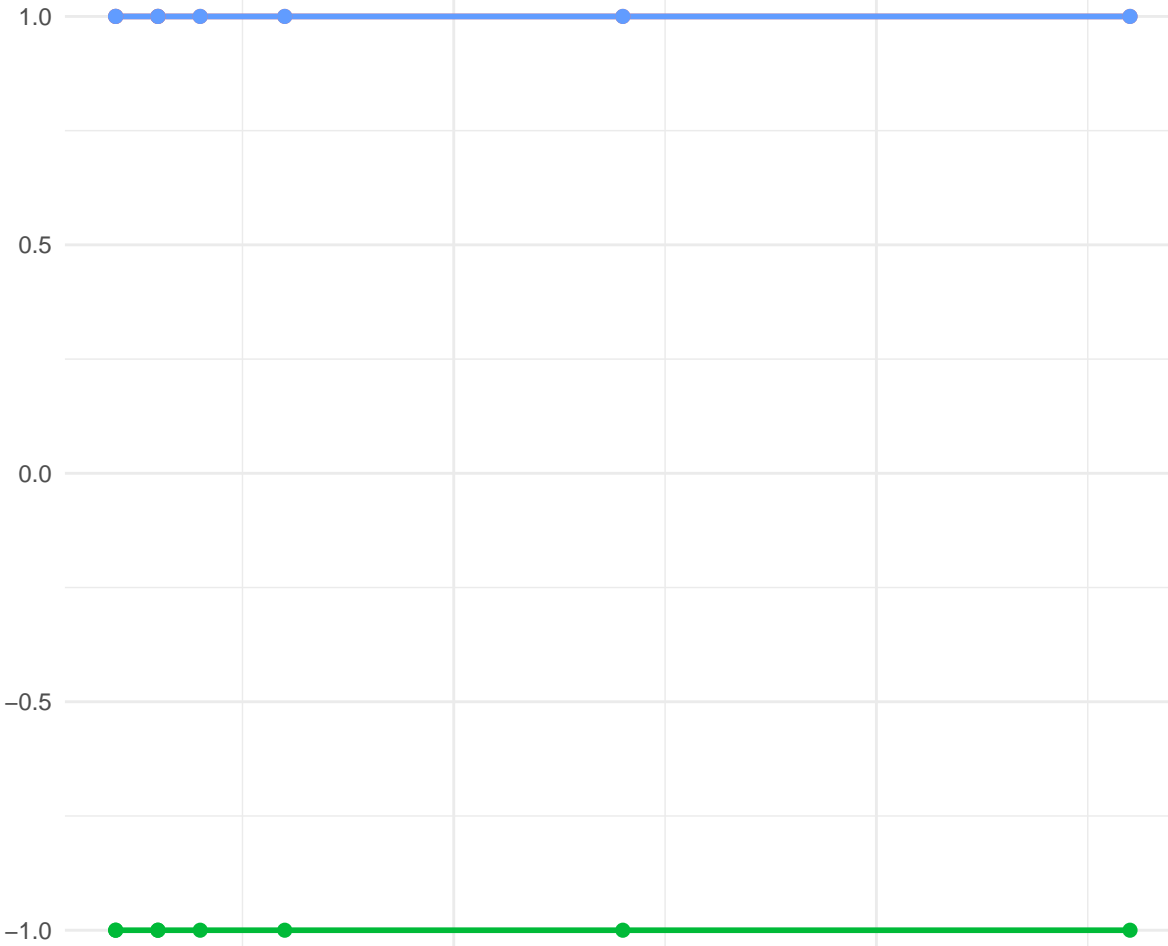
Number of Selected Samples

Summary Statistic

- range
- slope
- variance

10

20



Score: D\_slope (Local, partial2 , Genotype: wave )

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

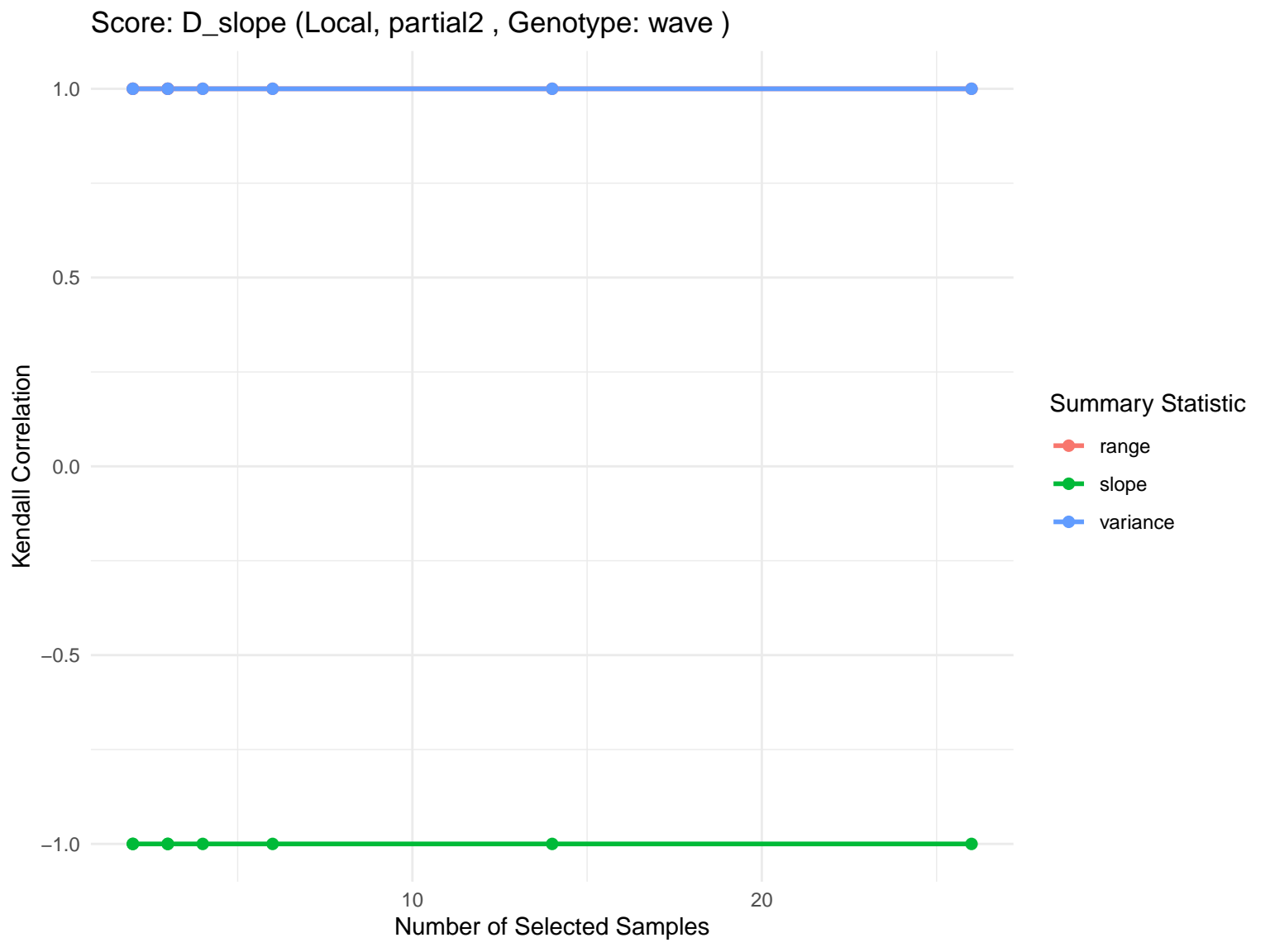
10

20

Number of Selected Samples

Summary Statistic

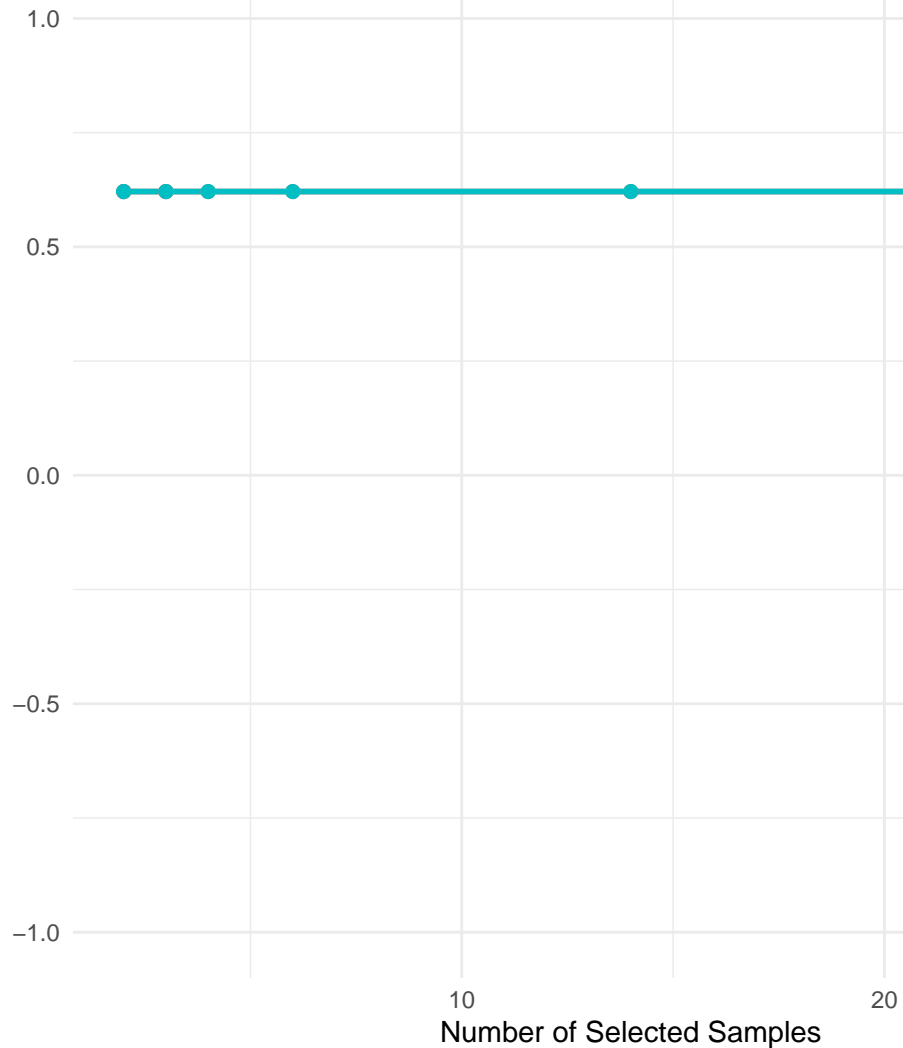
- range
- slope
- variance





Score: 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

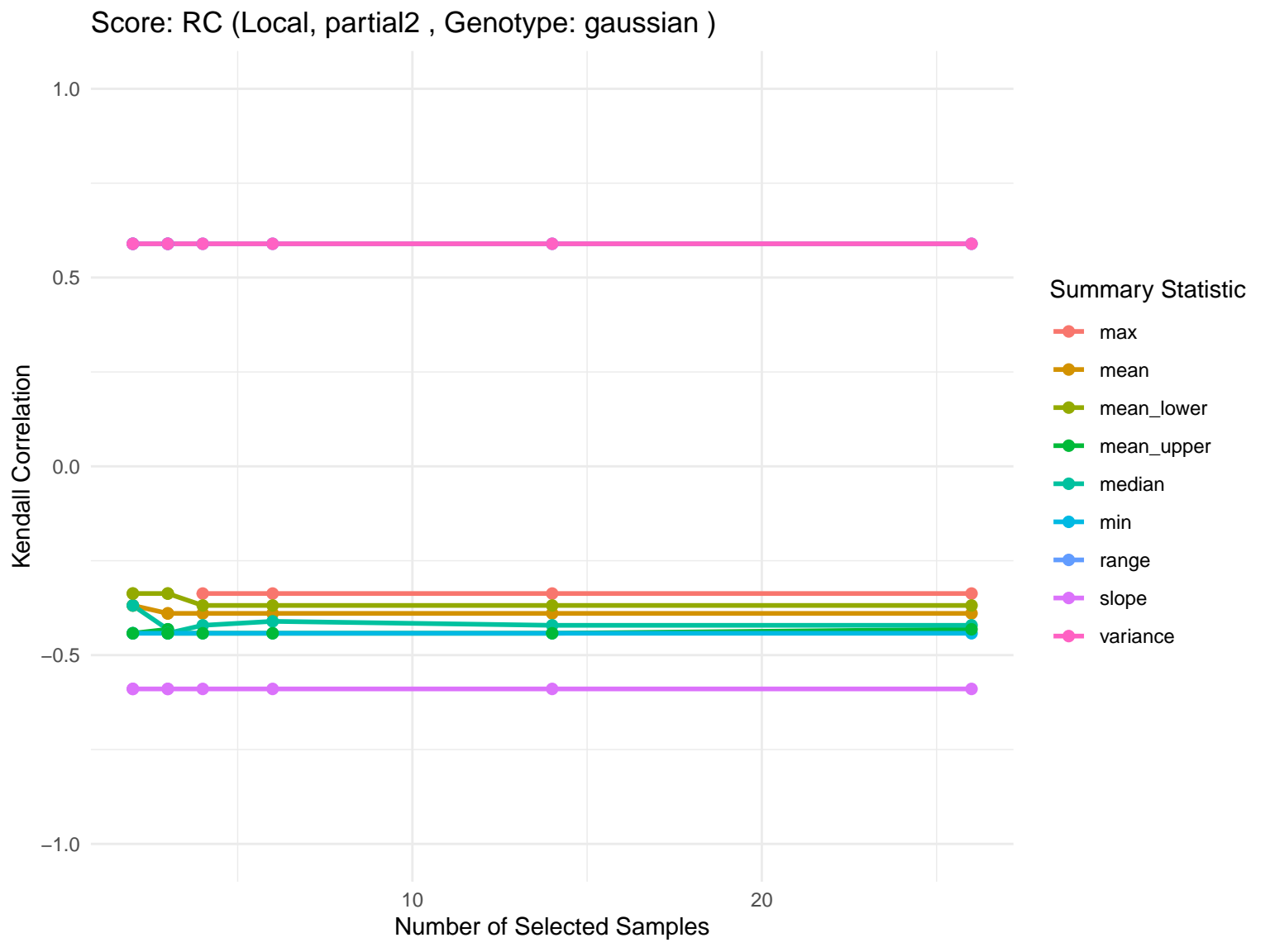
-0.5

-1.0

10

20

Number of Selected Samples

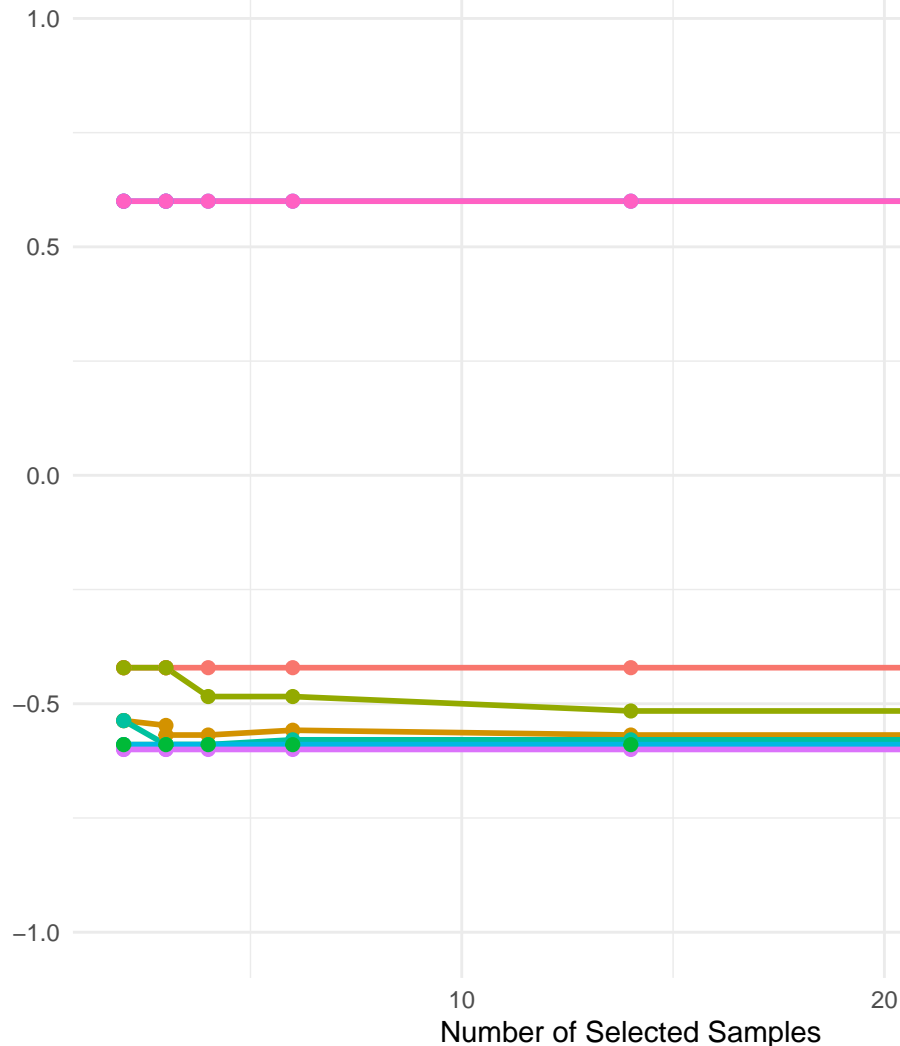


Score: RC (Local, partial2 , Genotype: sinusoidal )

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

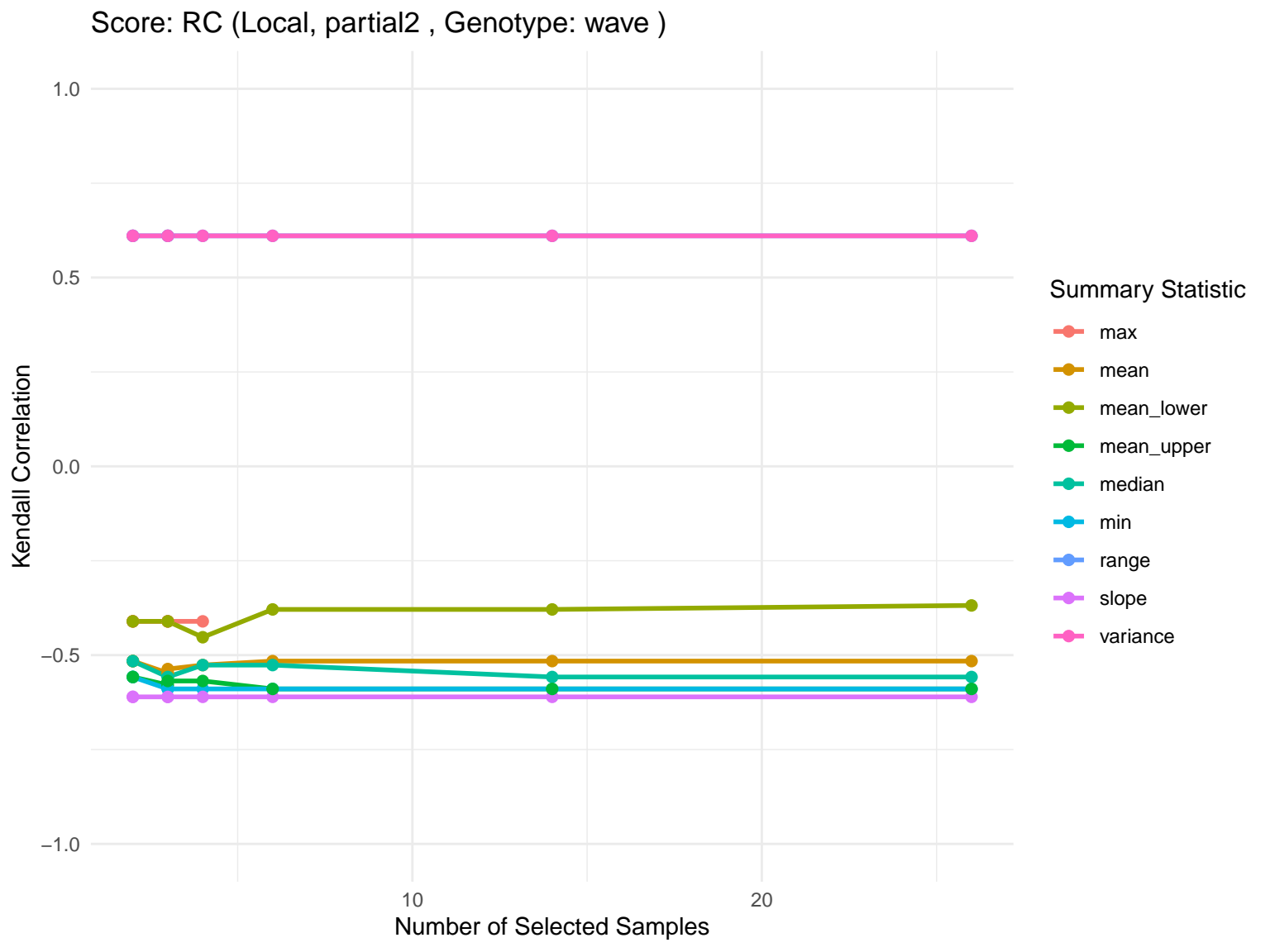
-0.5

-1.0

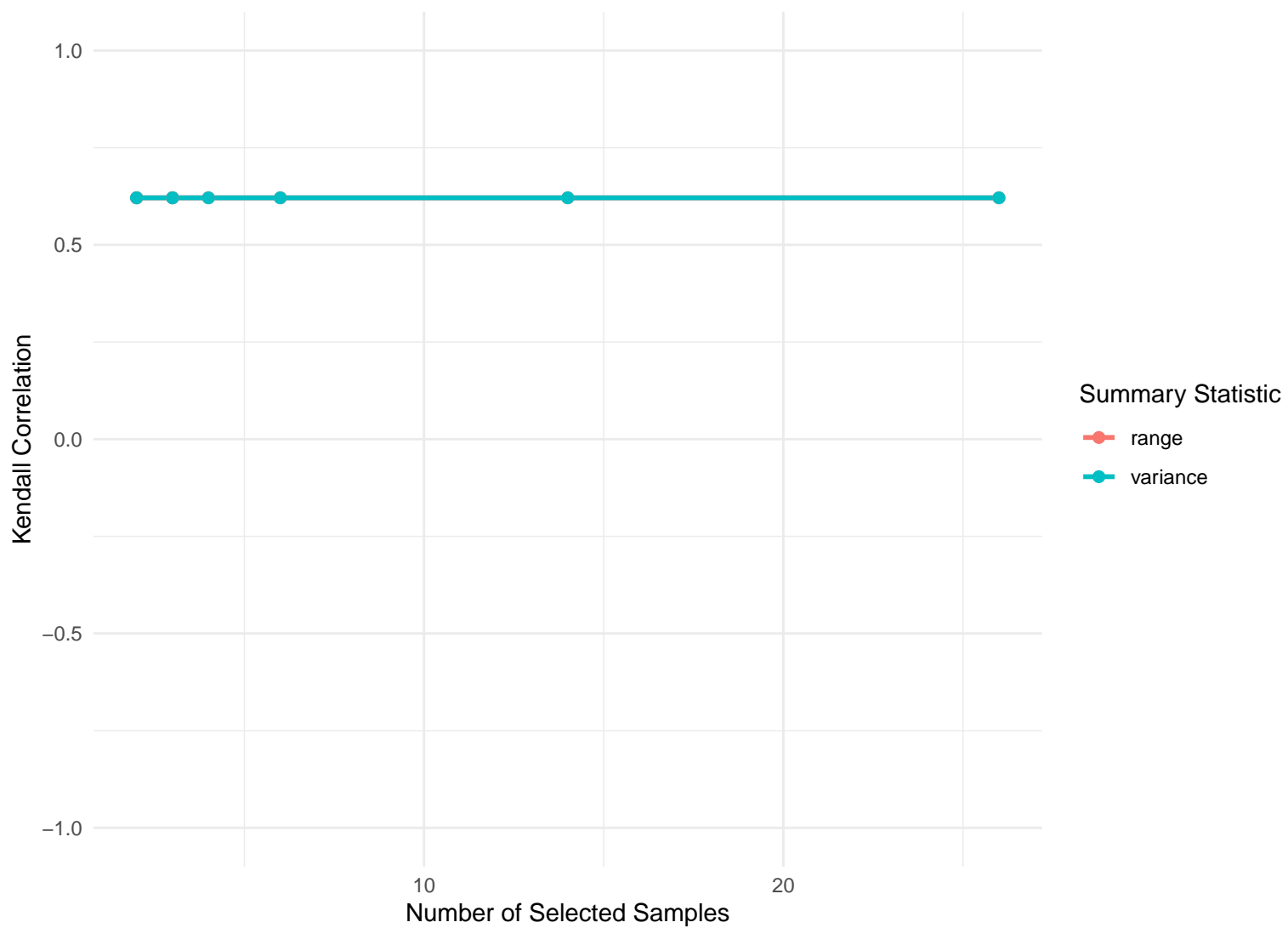
Number of Selected Samples

10

20



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

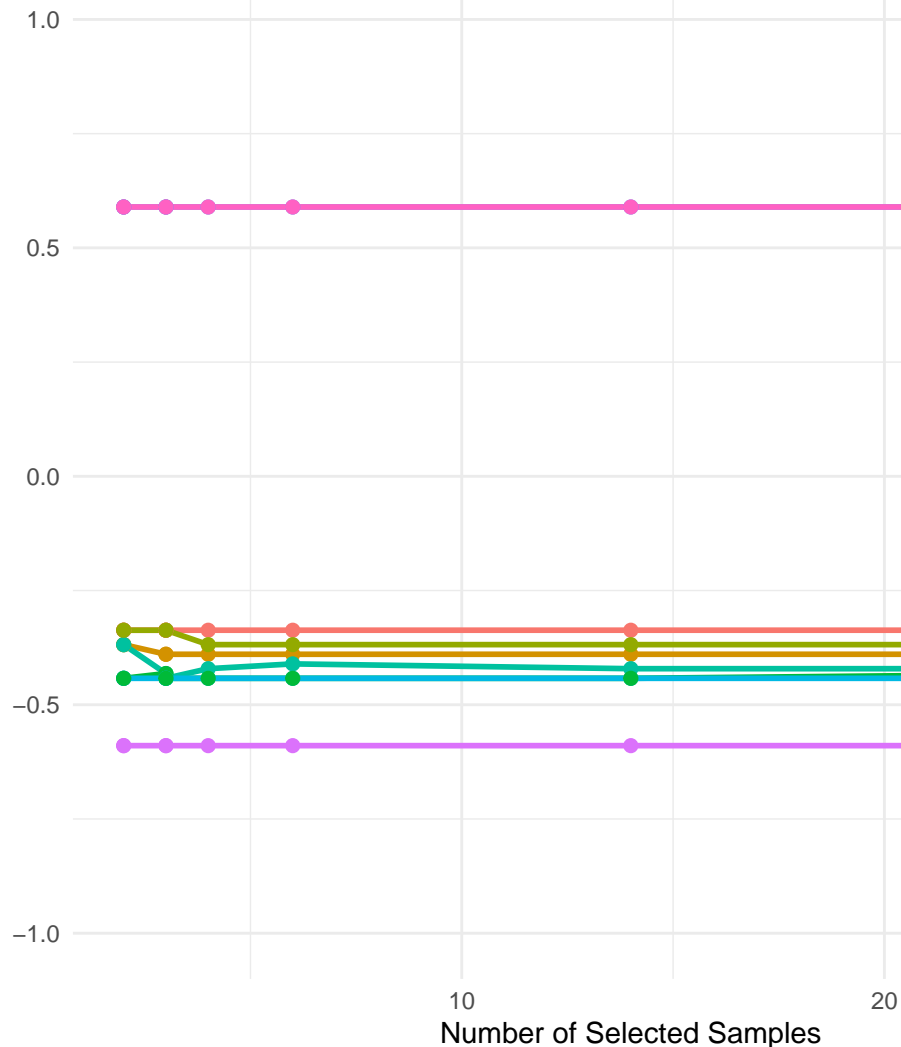


Score: gPi (Local, partial2 , Genotype: gaussian )

Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance



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

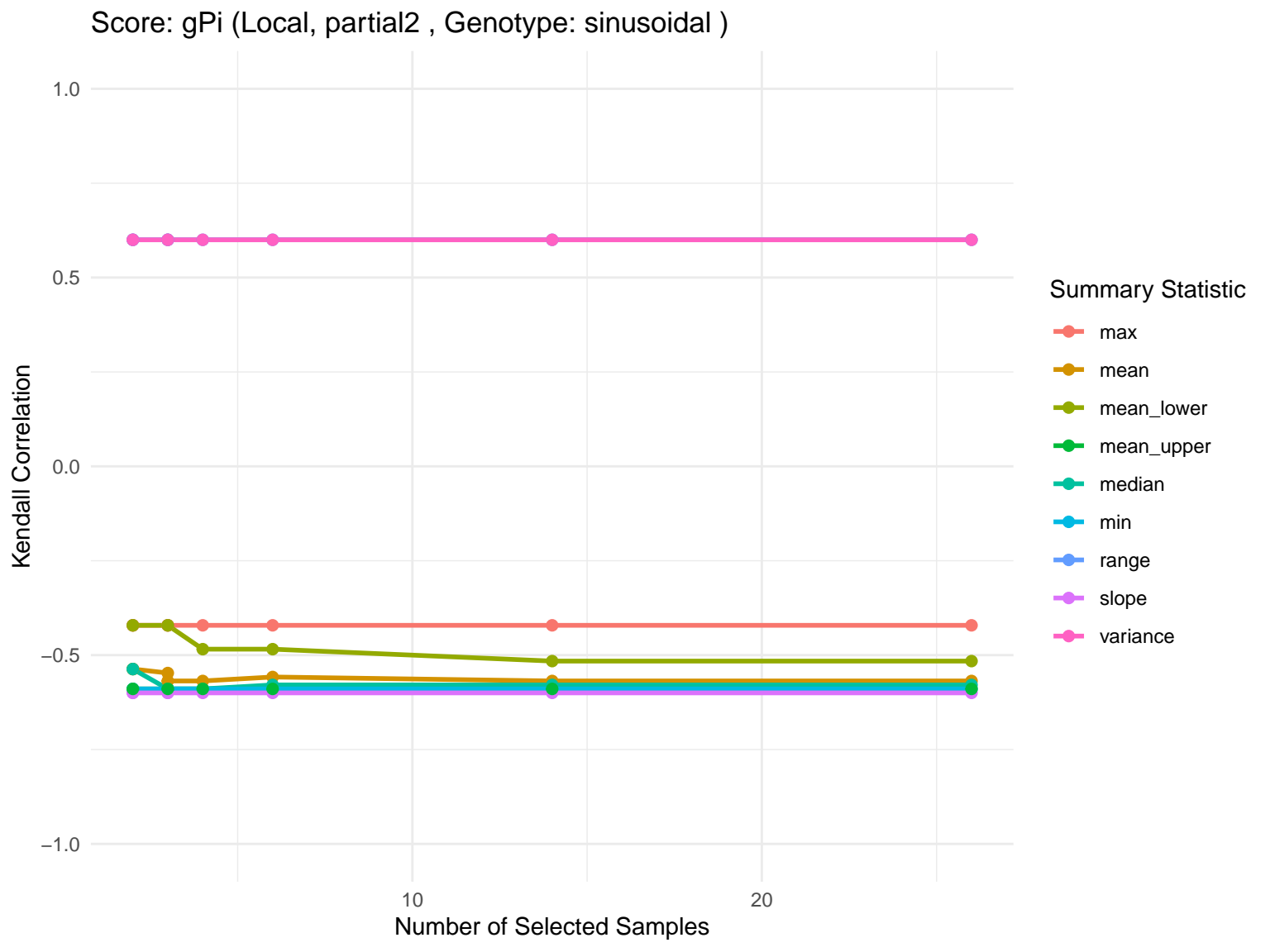
Kendall Correlation

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

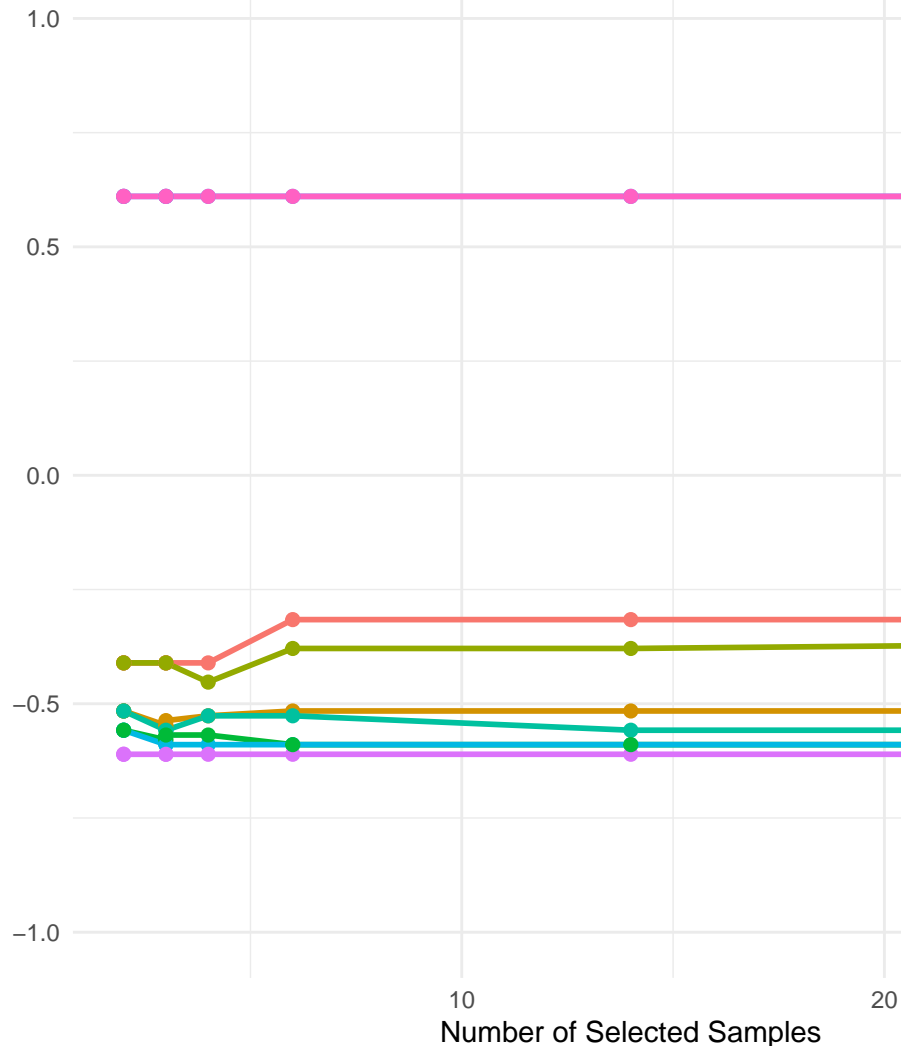


Score: gPi (Local, partial2 , Genotype: wave )

Kendall Correlation

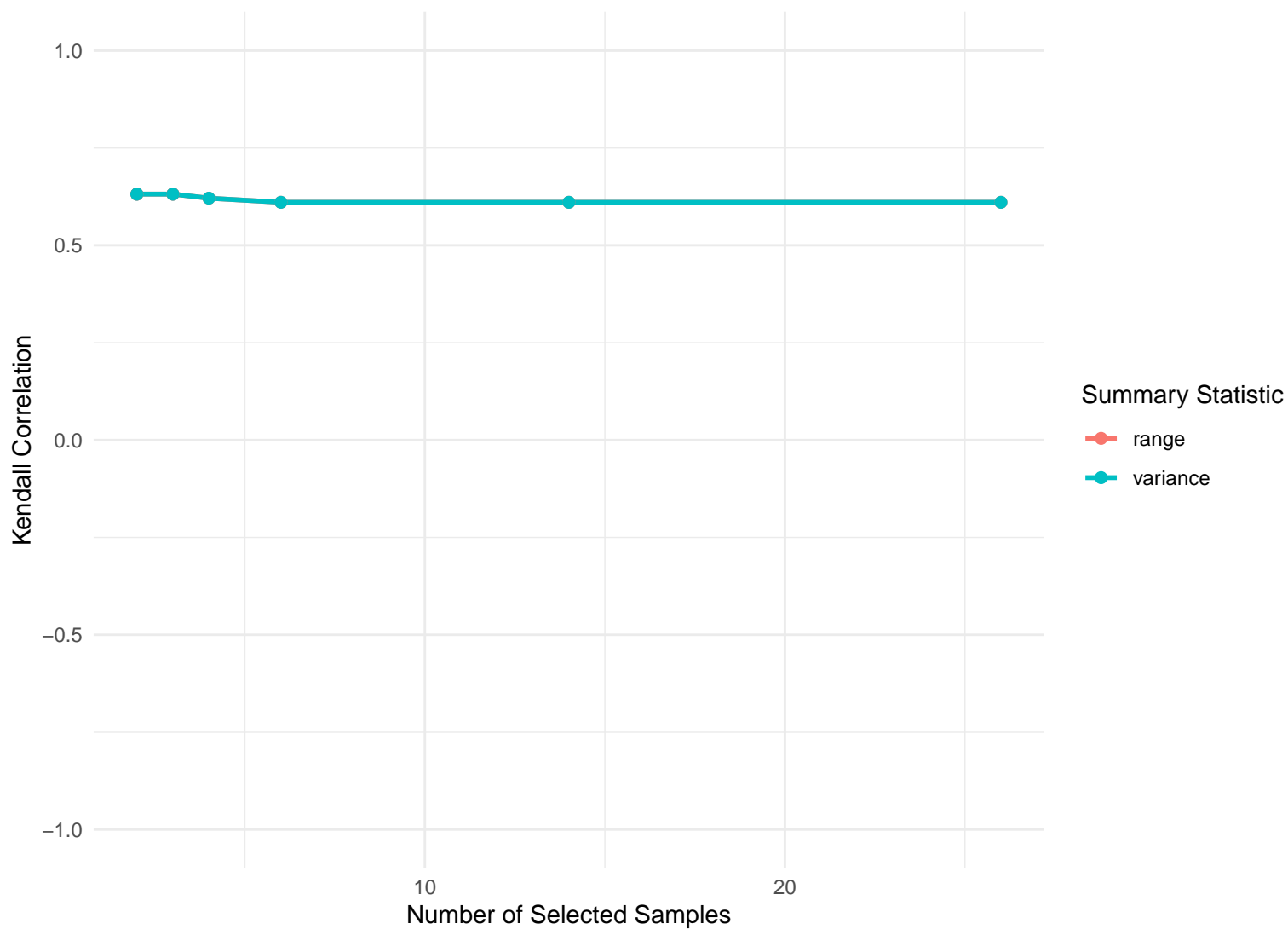
Summary Statistic

- 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

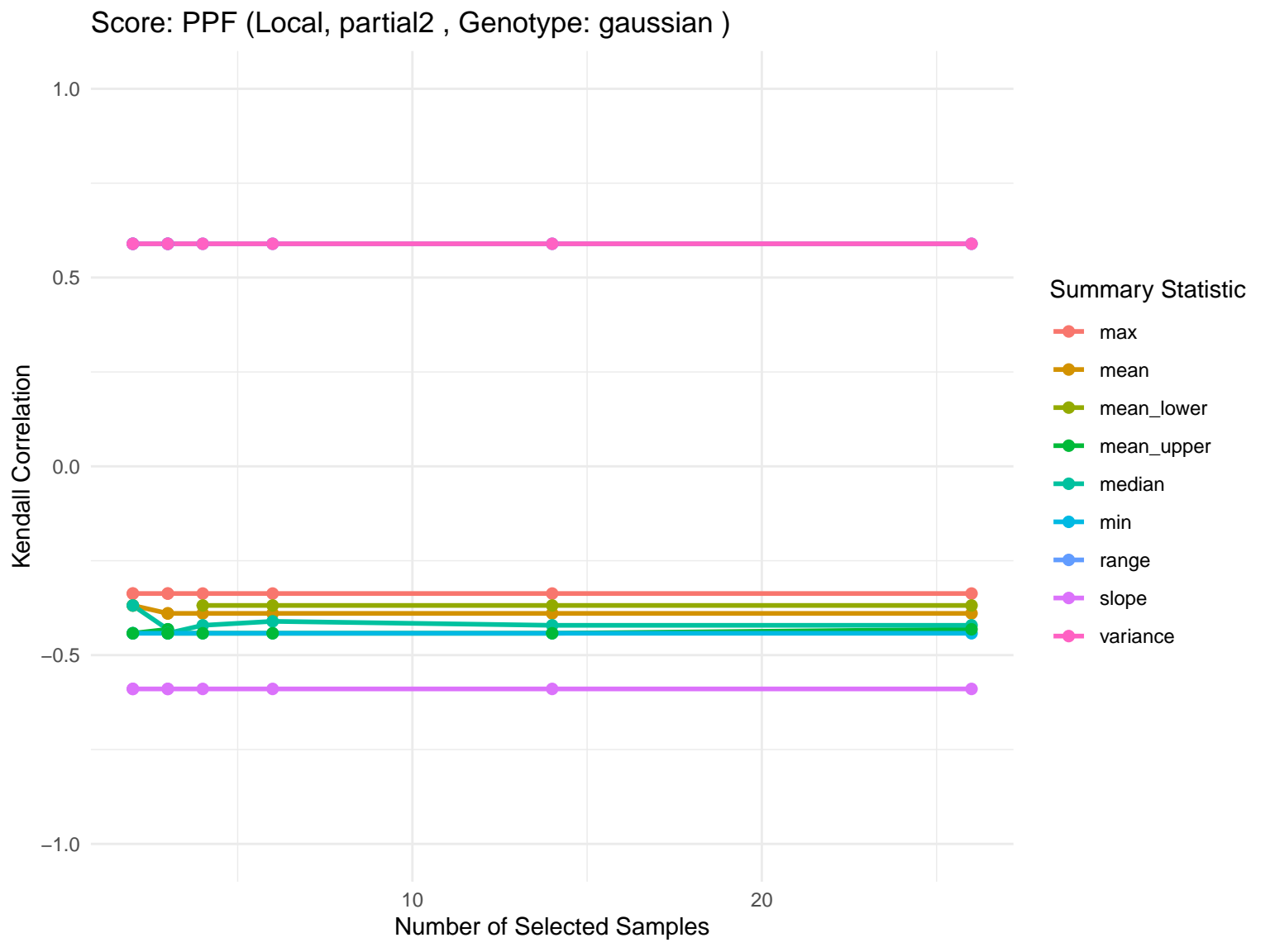
-0.5

-1.0

Number of Selected Samples

10

20

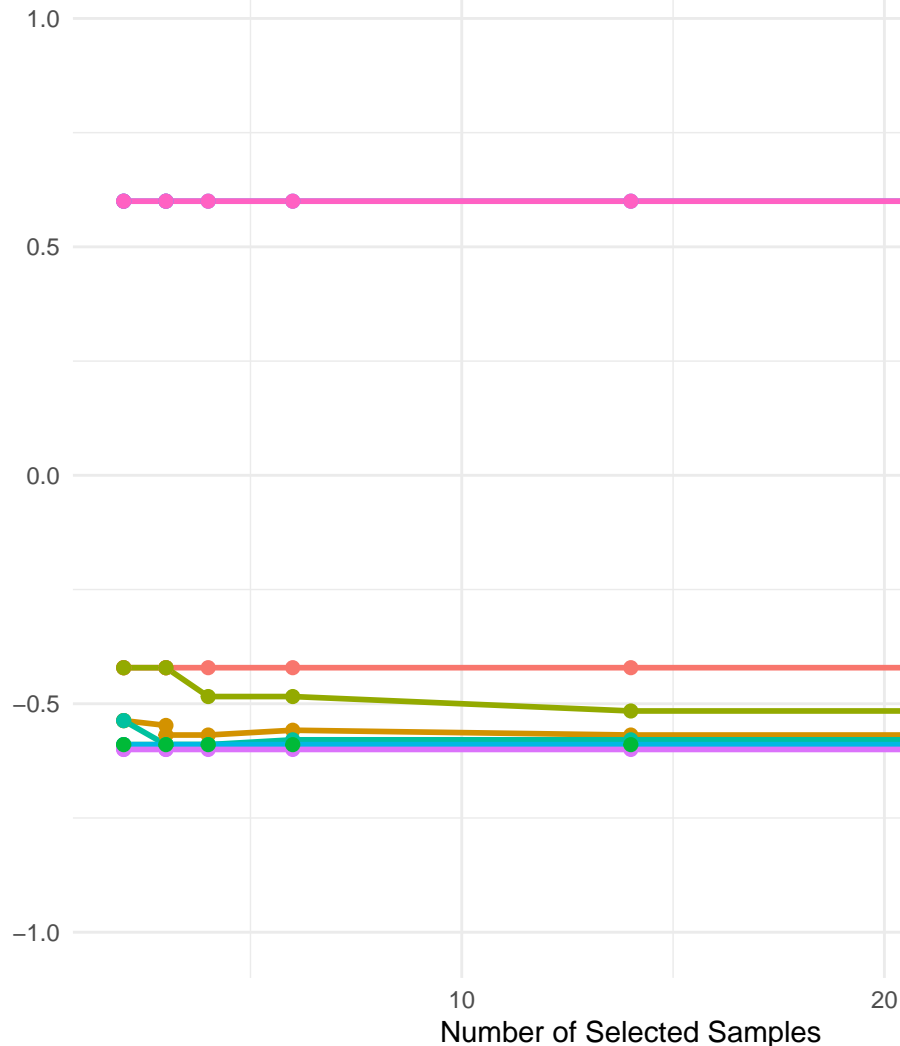


Score: PPF (Local, partial2 , Genotype: sinusoidal )

Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance



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

Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

1.0

0.5

0.0

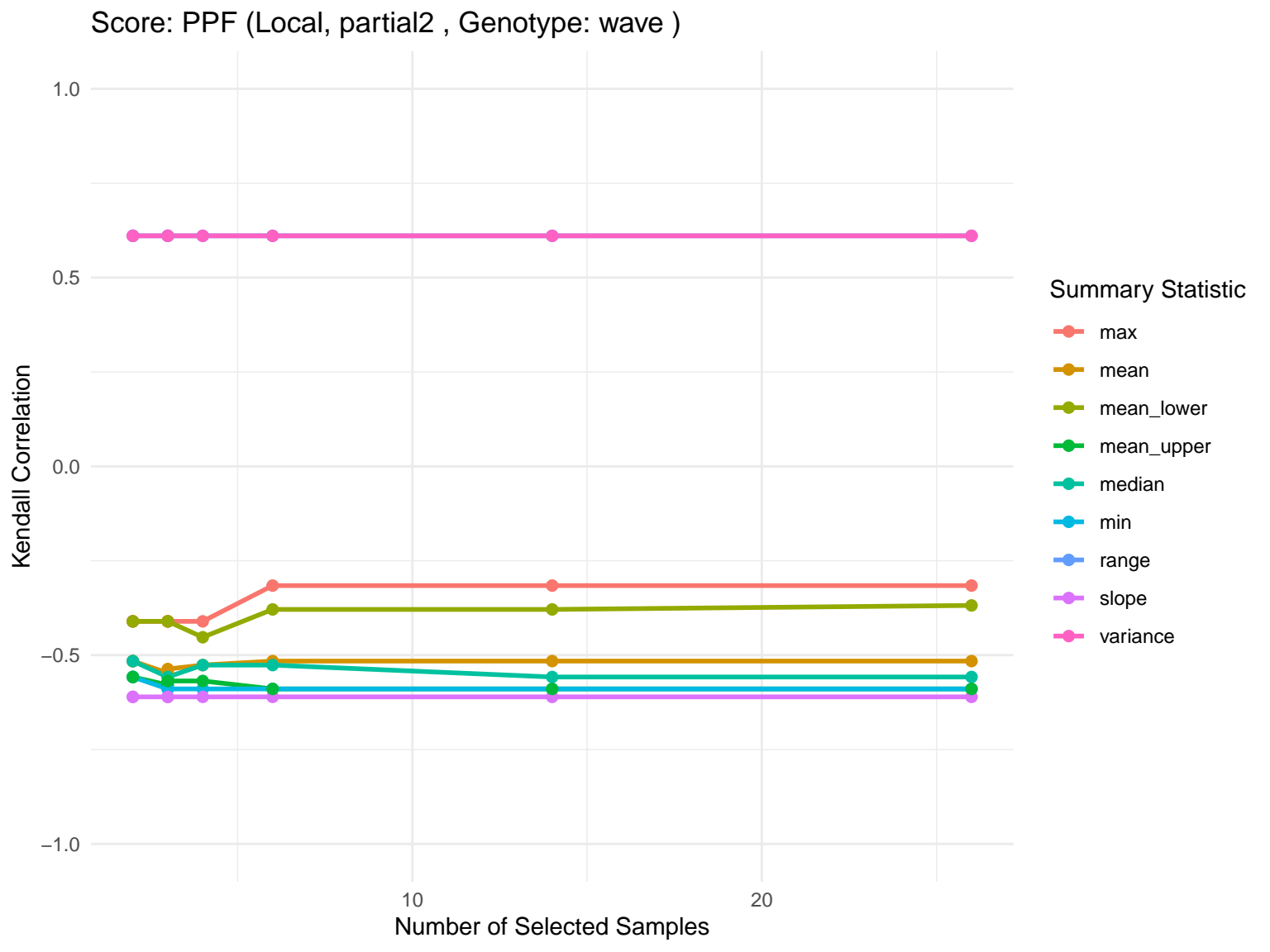
-0.5

-1.0

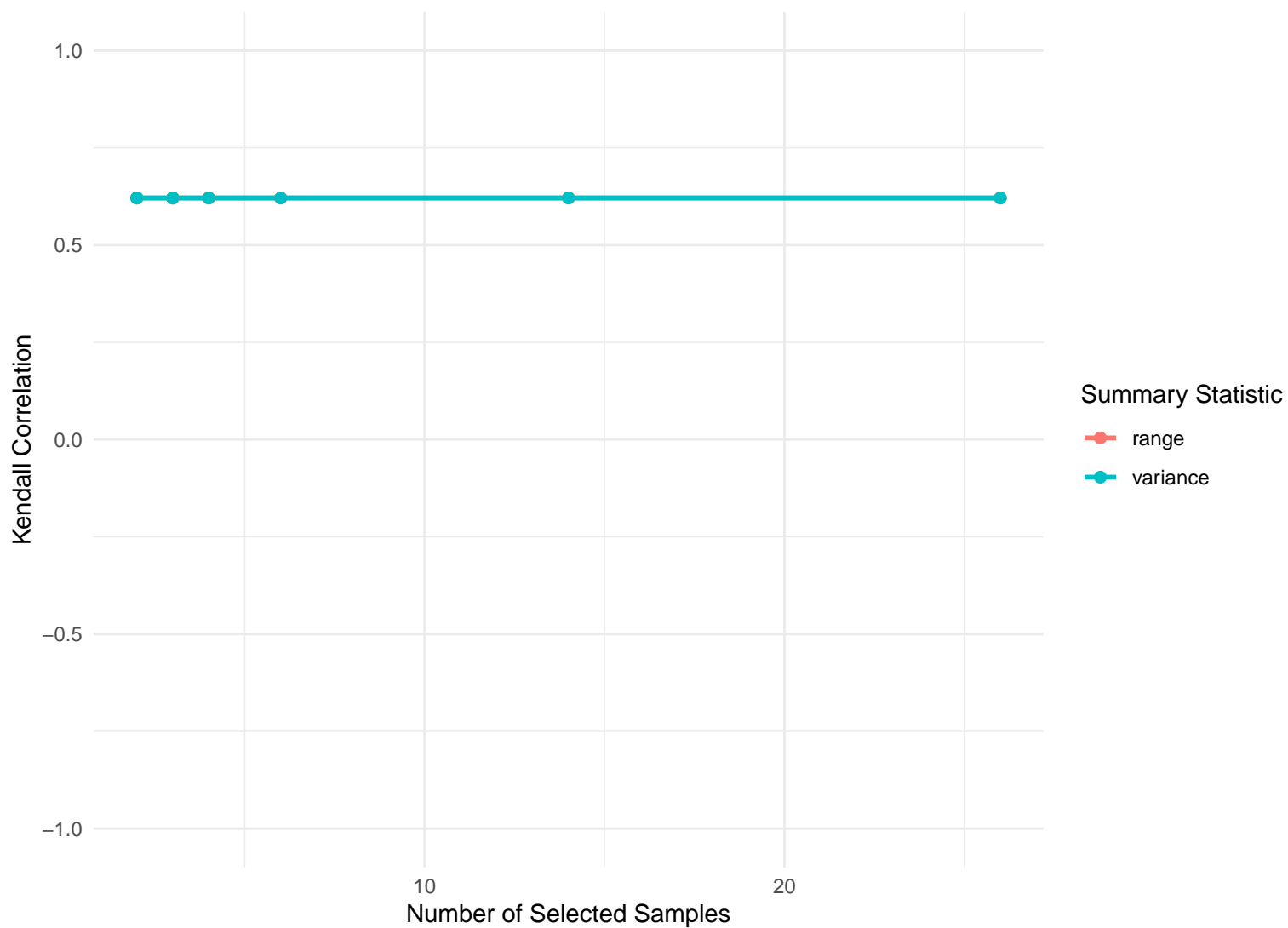
Number of Selected Samples

10

20



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



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

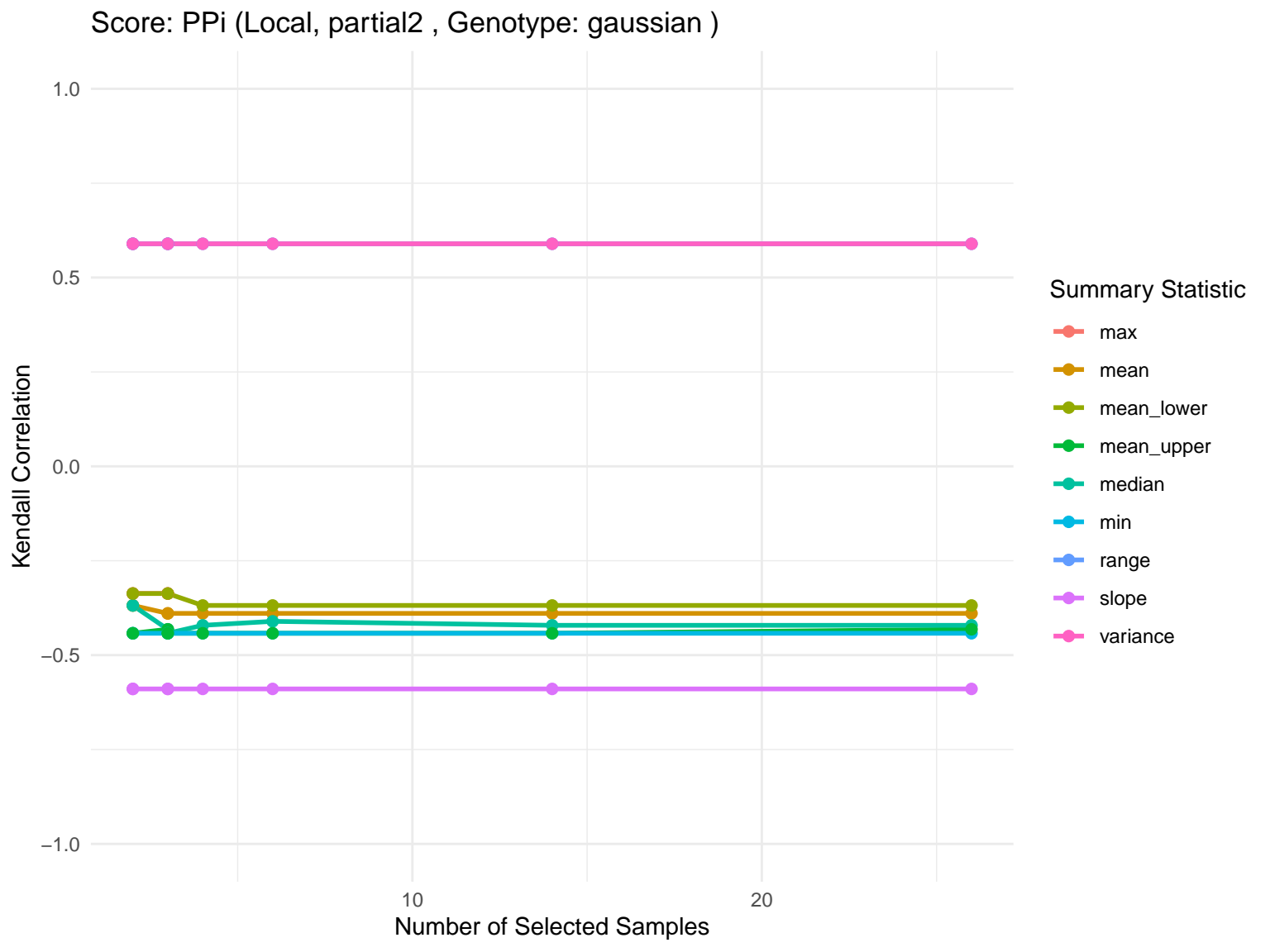
Kendall Correlation

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: PPI (Local, partial2 , Genotype: sinusoidal )



Number of Selected Samples

## Summary Statistic

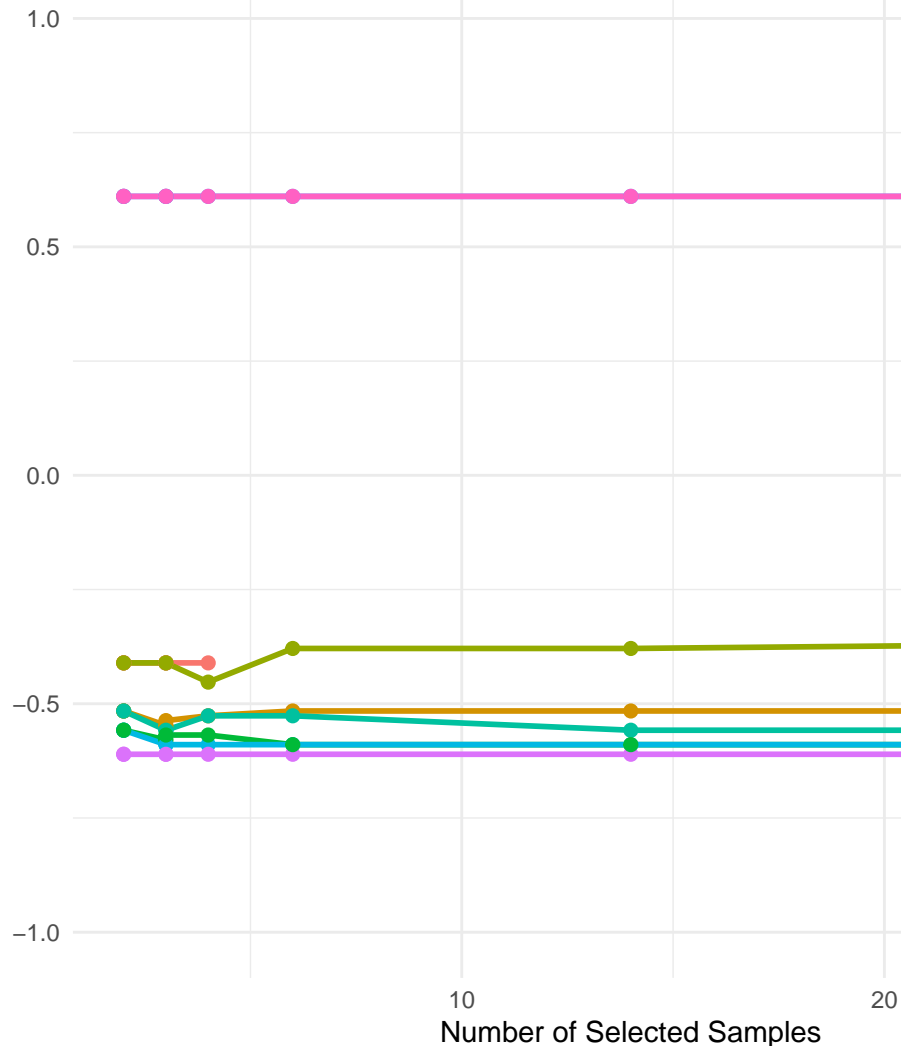
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

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

Kendall Correlation

Summary Statistic

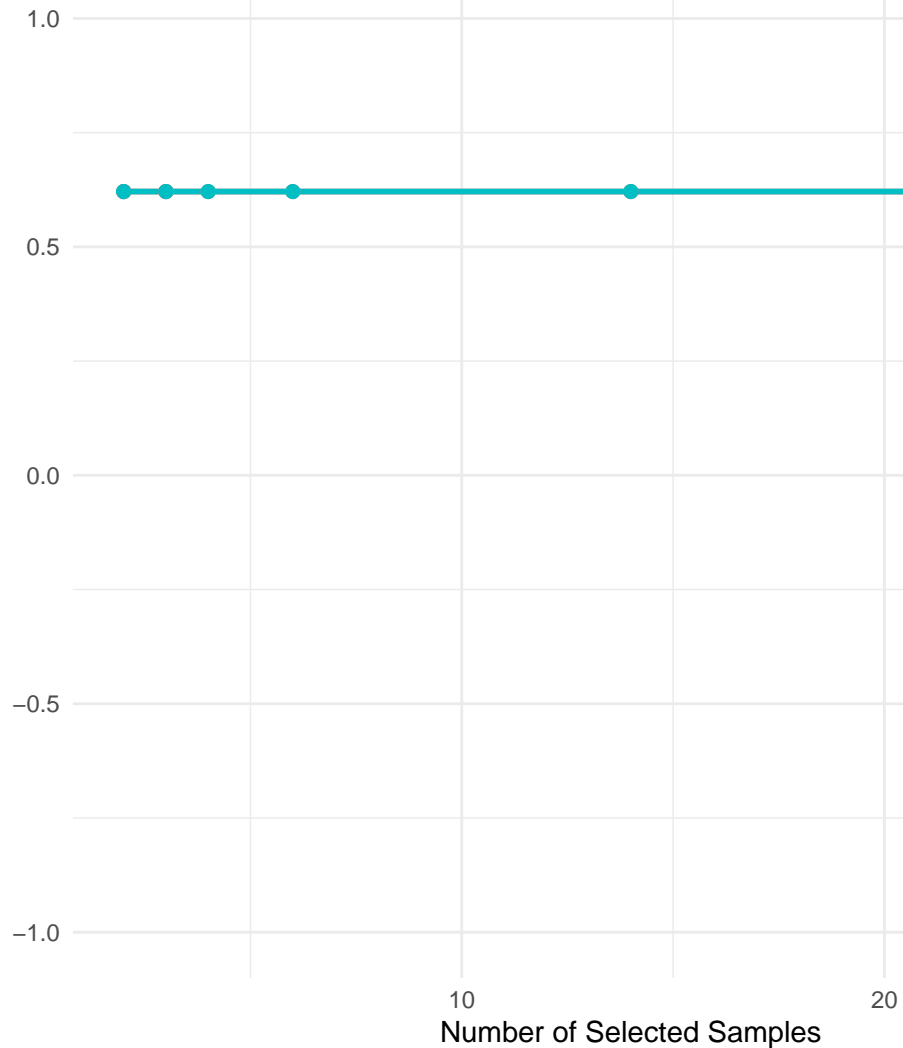
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance





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

Kendall Correlation



Summary Statistic

- range
- variance

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

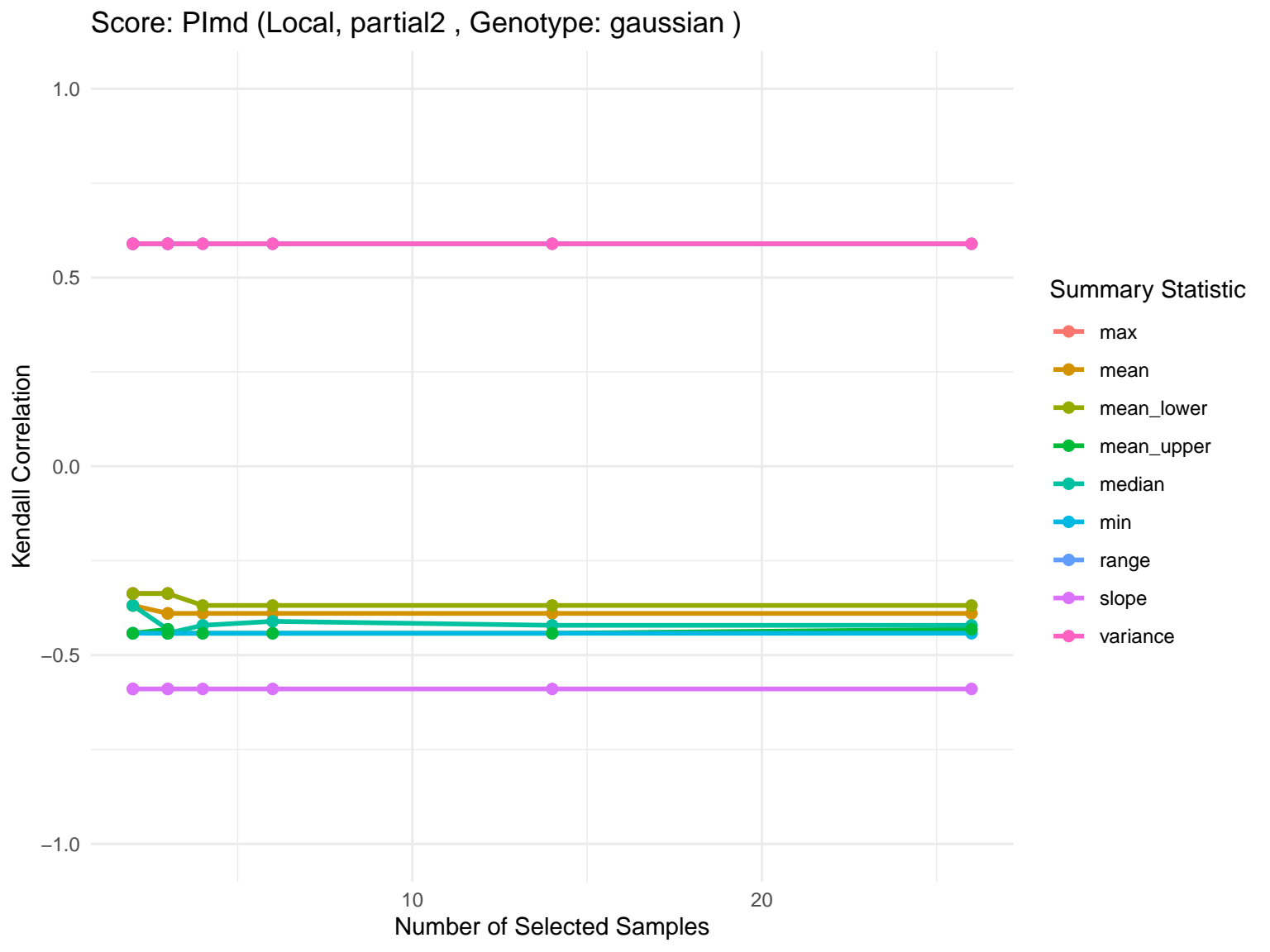
Kendall Correlation

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

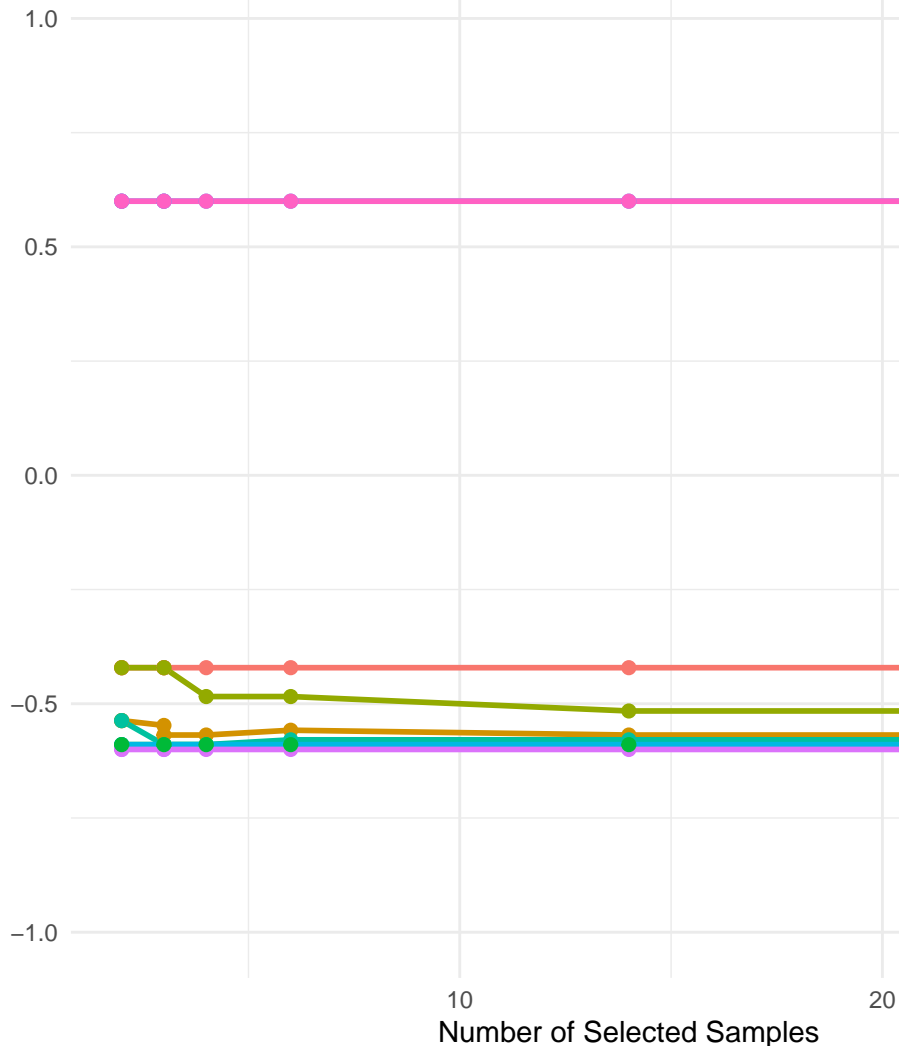


Score: Plmd (Local, partial2 , Genotype: sinusoidal )

Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance



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

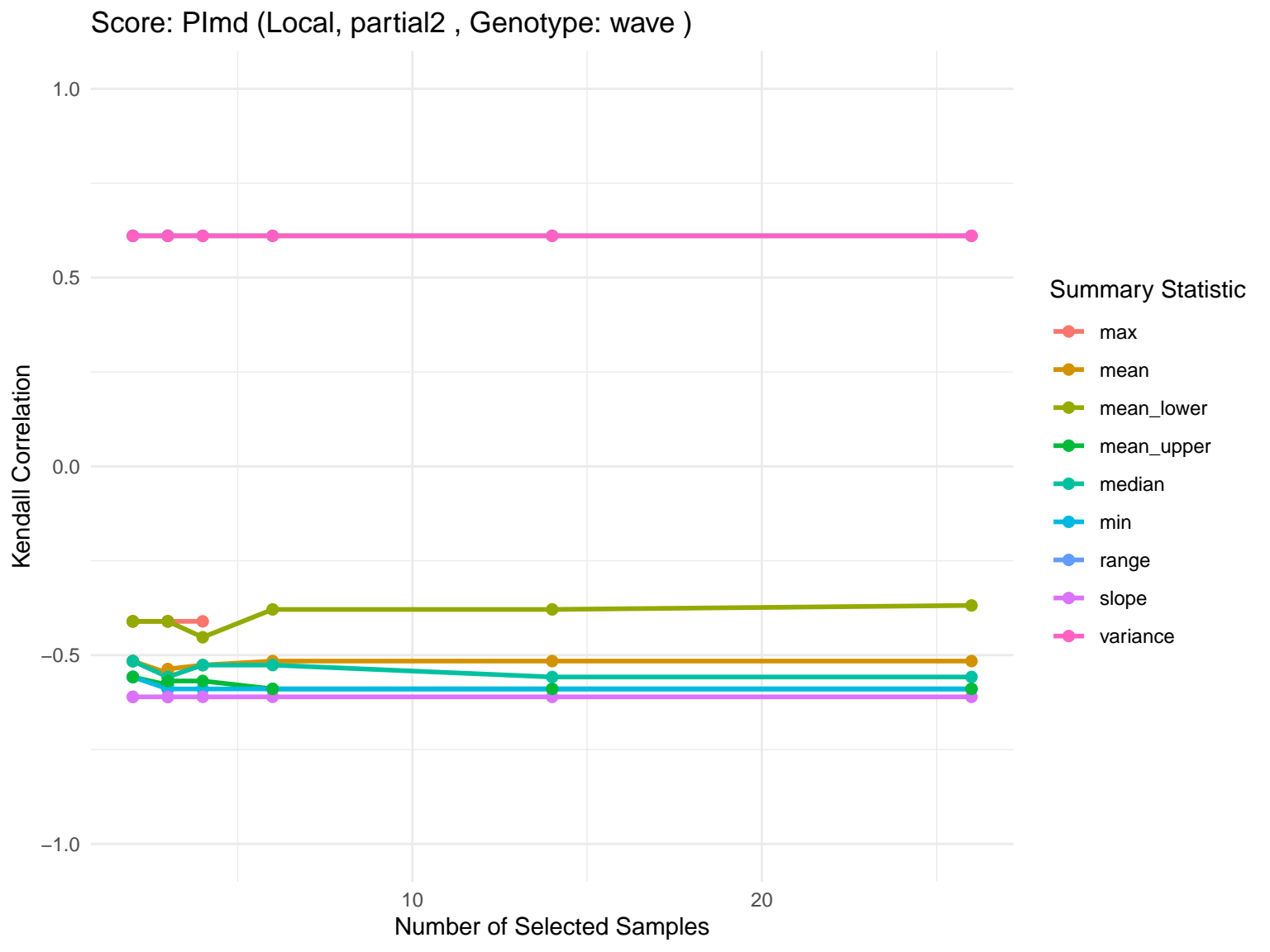
Kendall Correlation

Summary Statistic

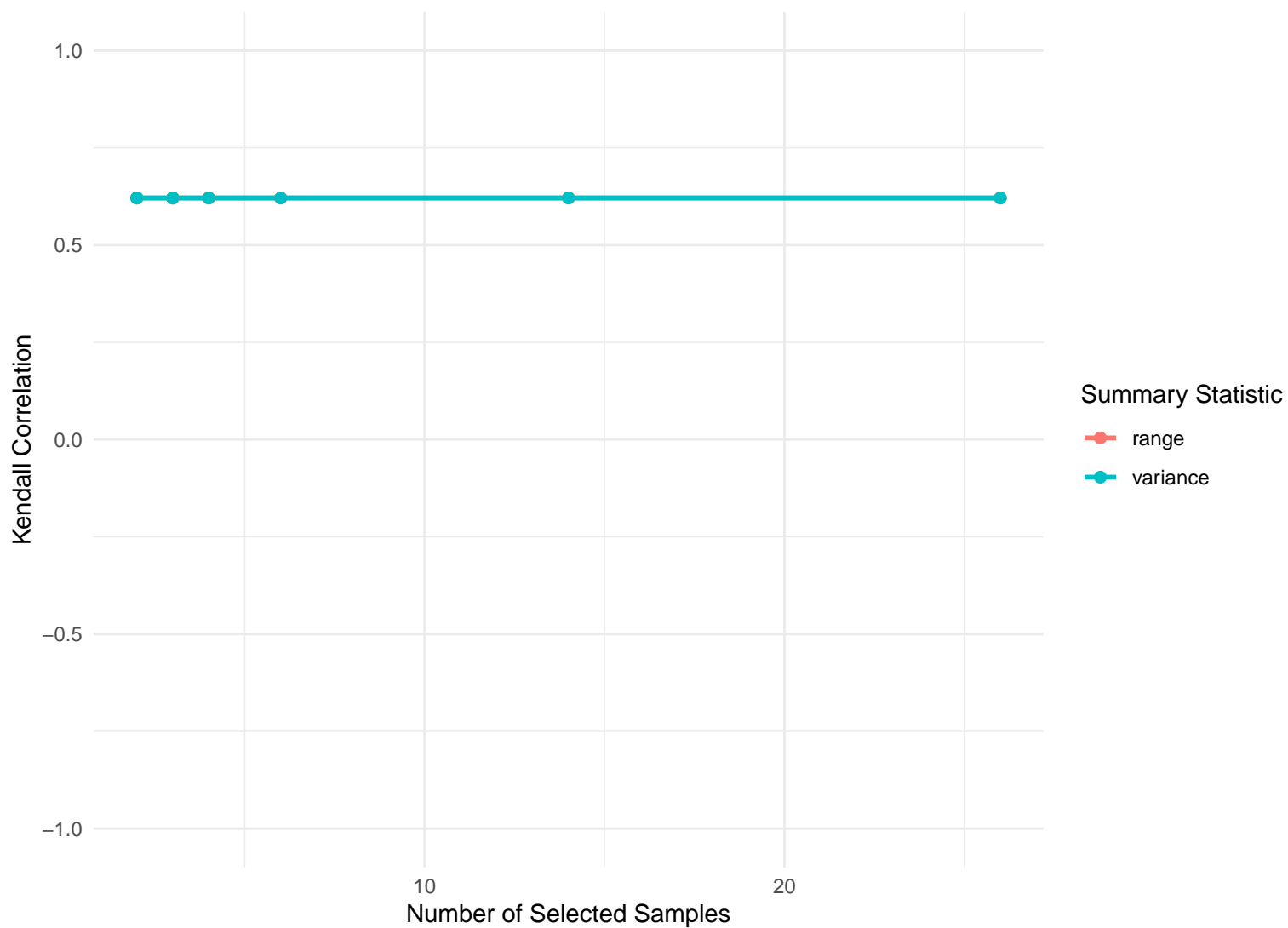
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

1.0  
0.5  
0.0  
-0.5  
-1.0

10 20  
Number of Selected Samples



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



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

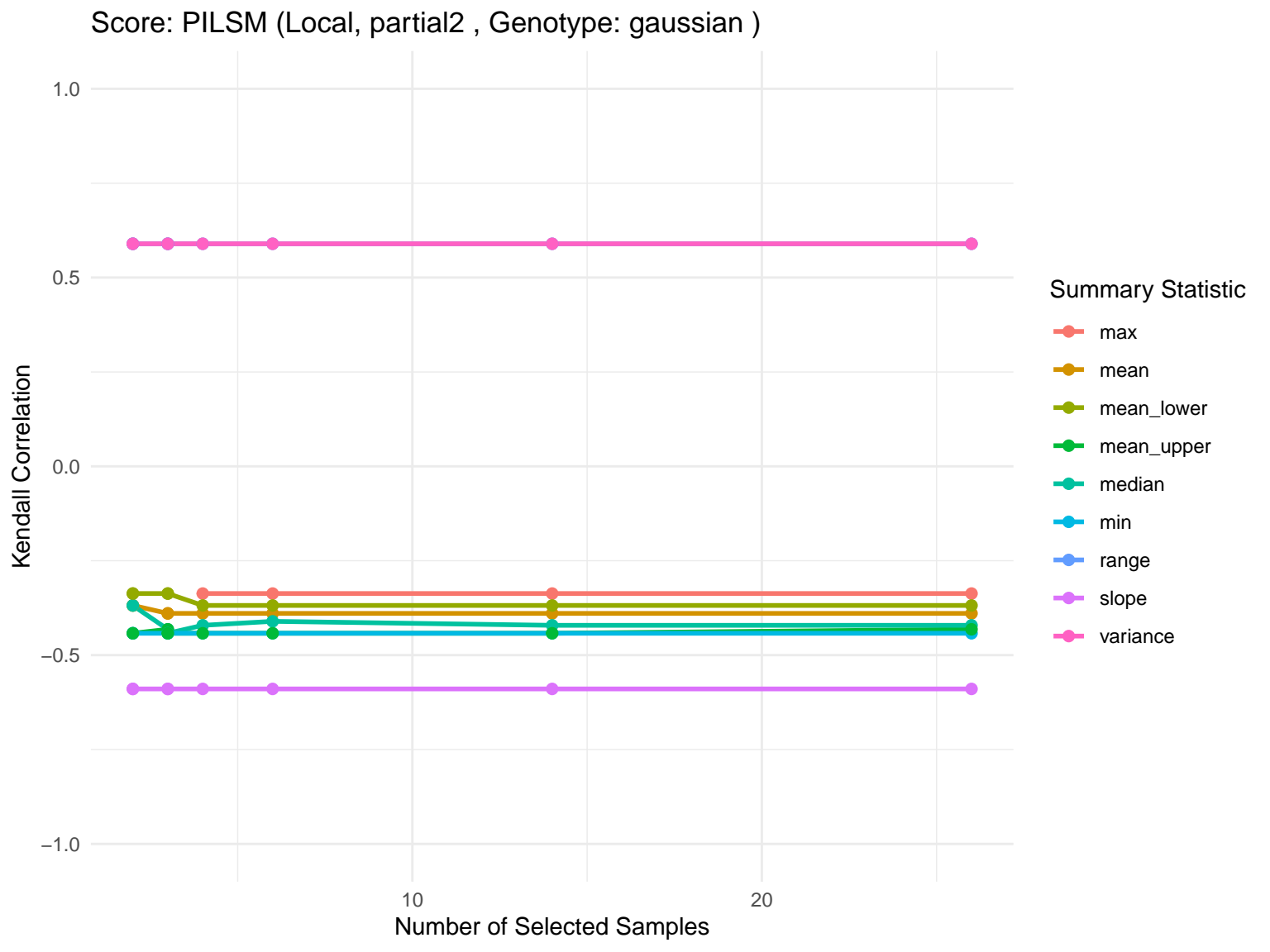
Kendall Correlation

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 )

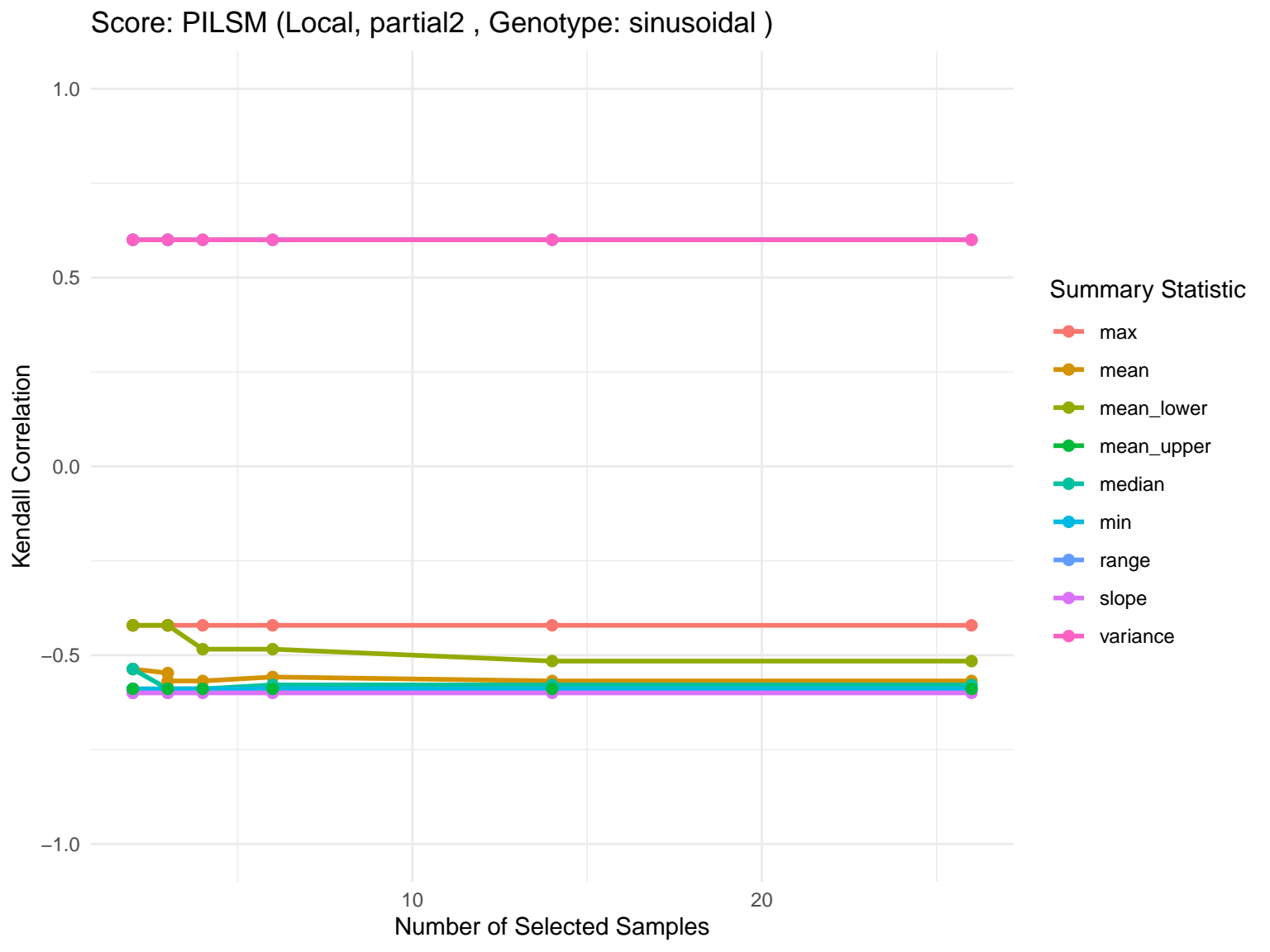
Kendall Correlation

Summary Statistic

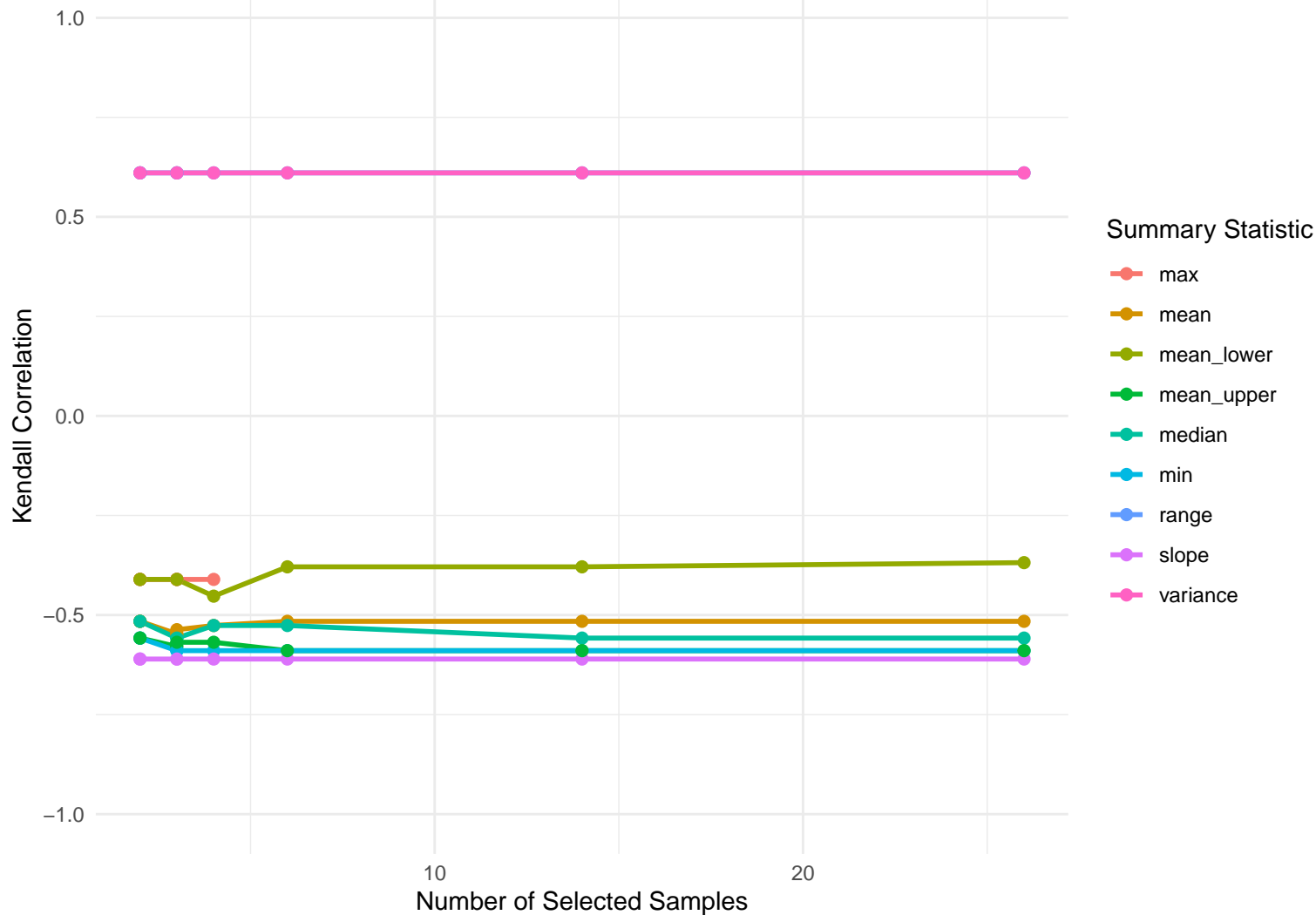
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

1.0  
0.5  
0.0  
-0.5  
-1.0

10 20  
Number of Selected Samples

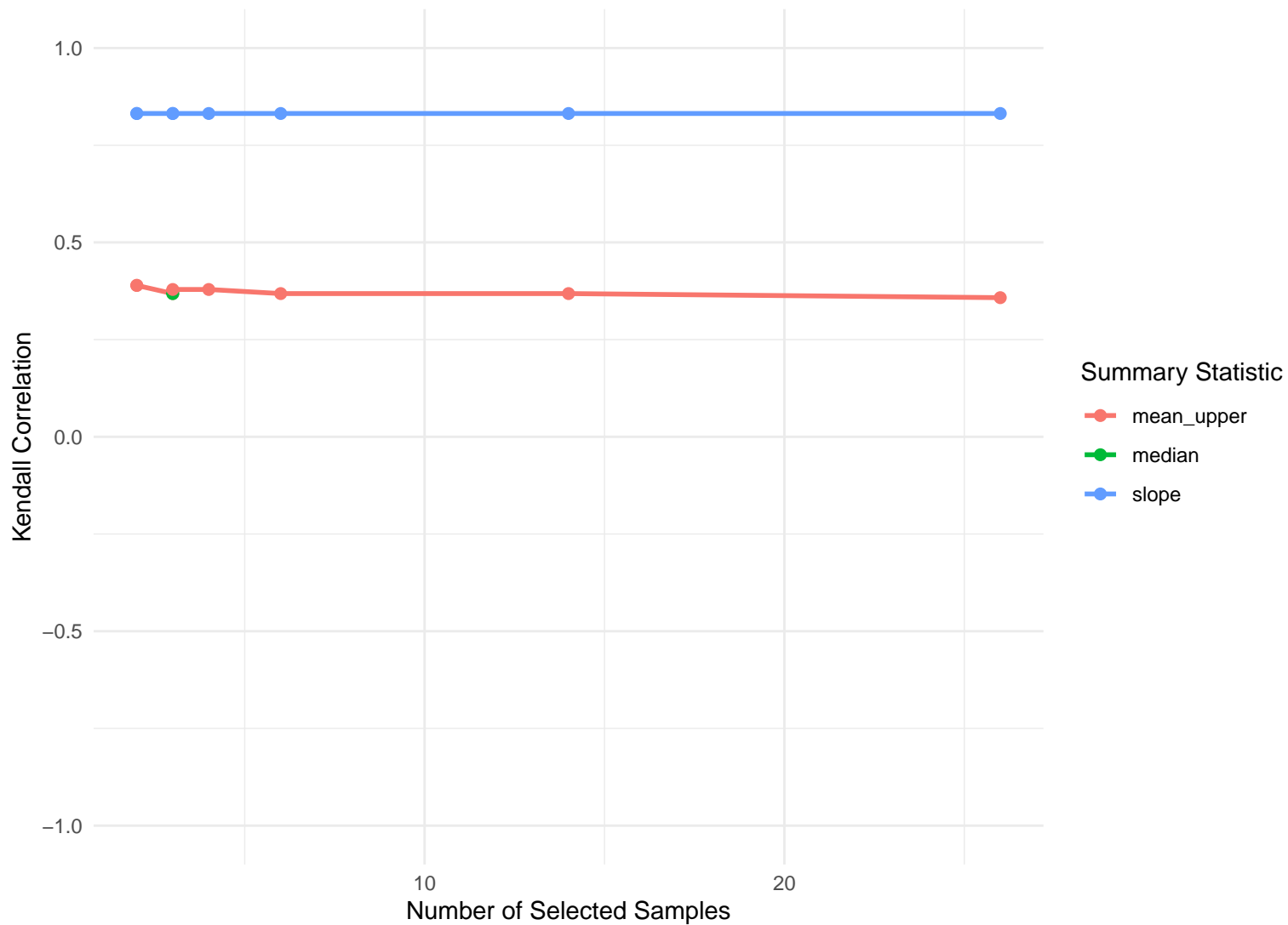


Score: PILSM (Local, partial2 , Genotype: wave )





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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

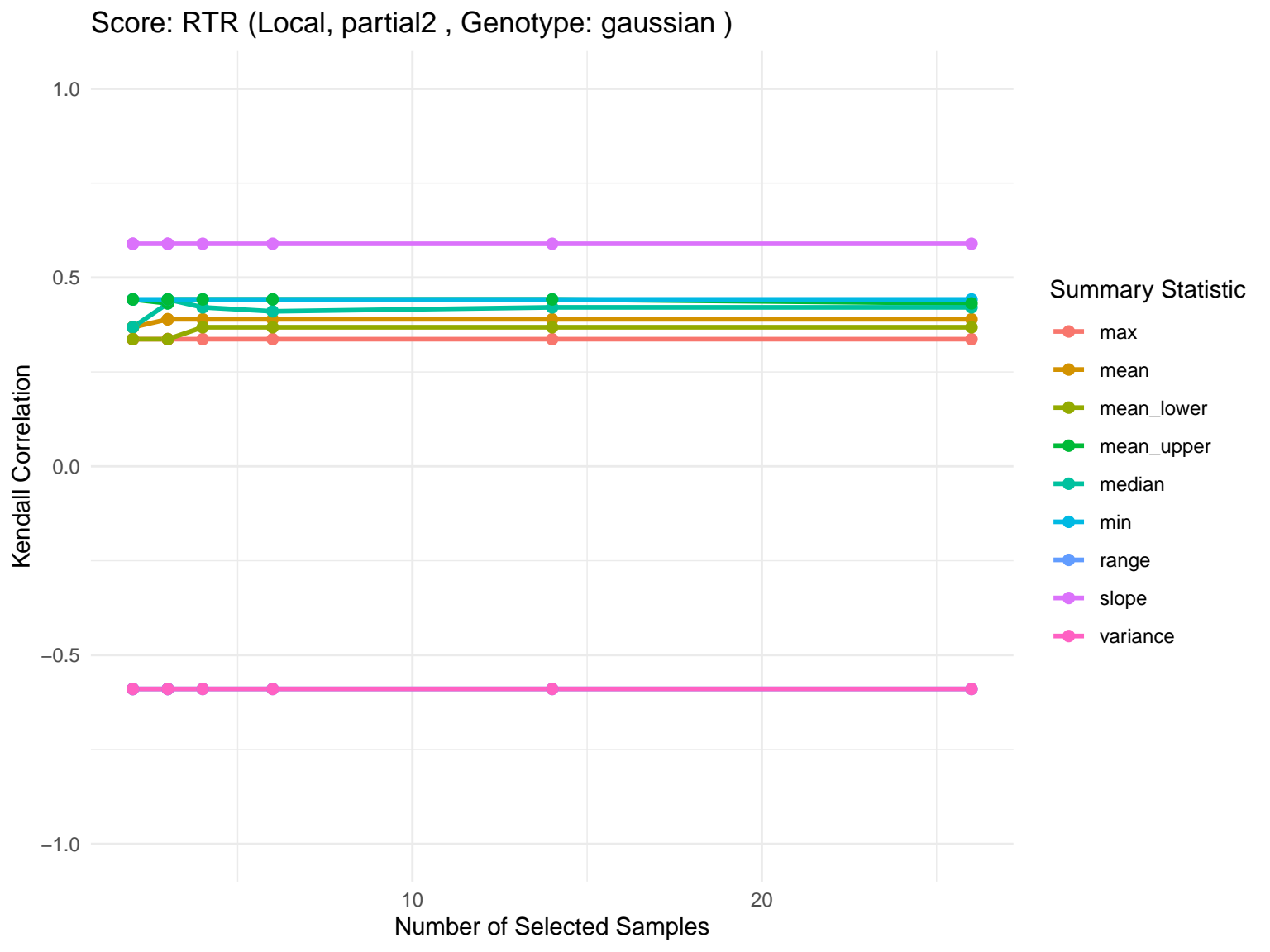
10

20

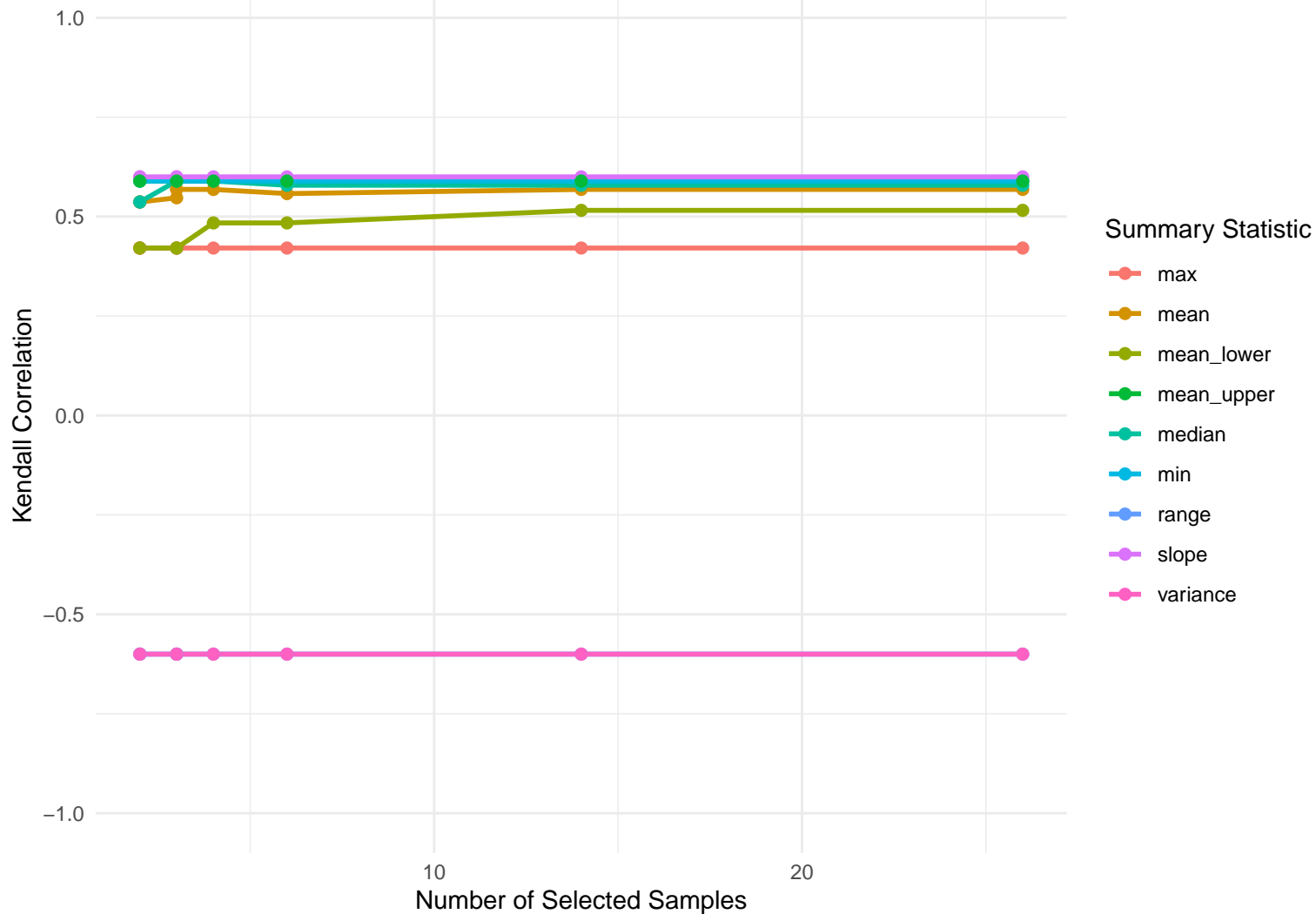
Number of Selected Samples

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 )

Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

1.0

0.5

0.0

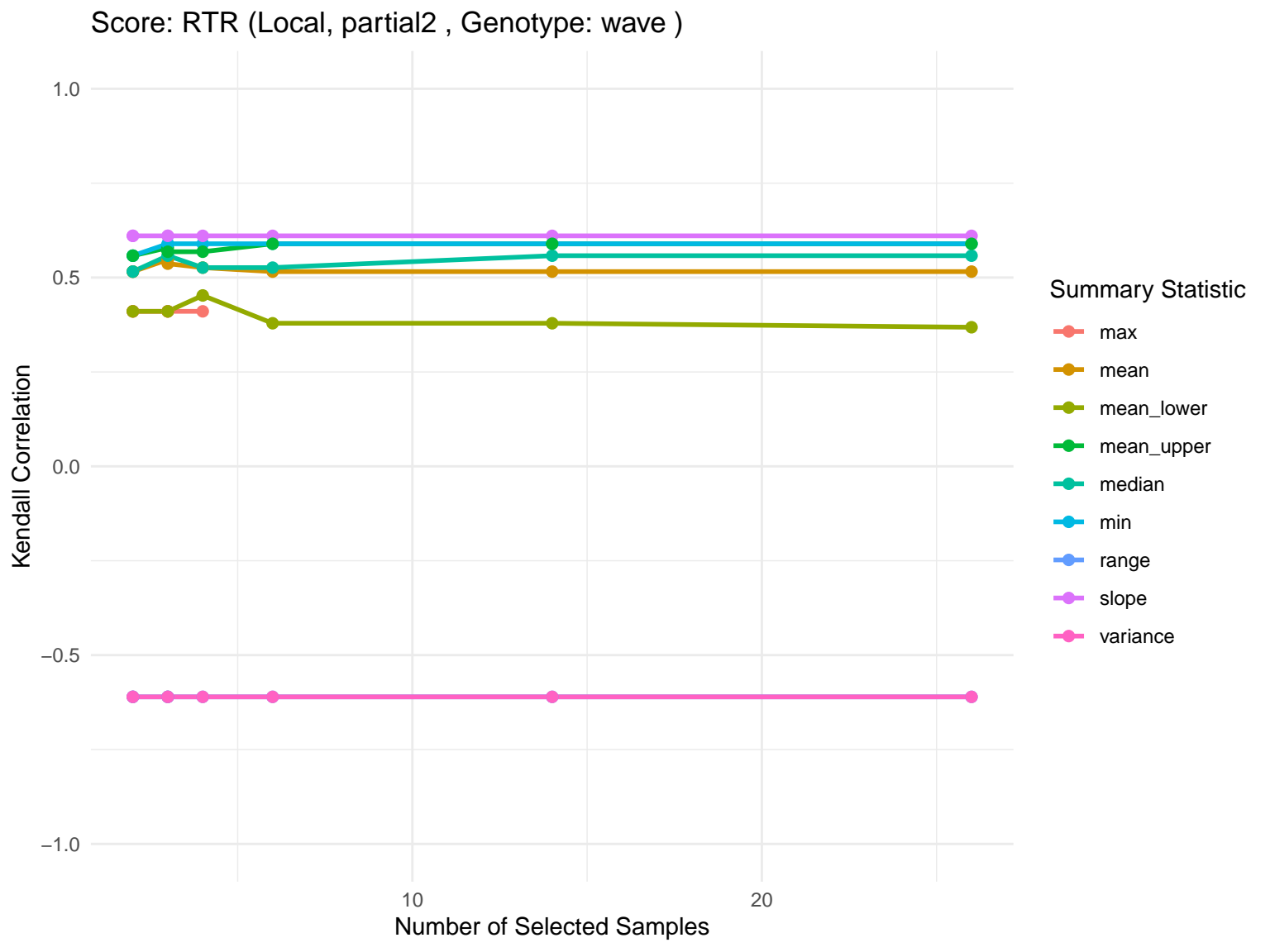
-0.5

-1.0

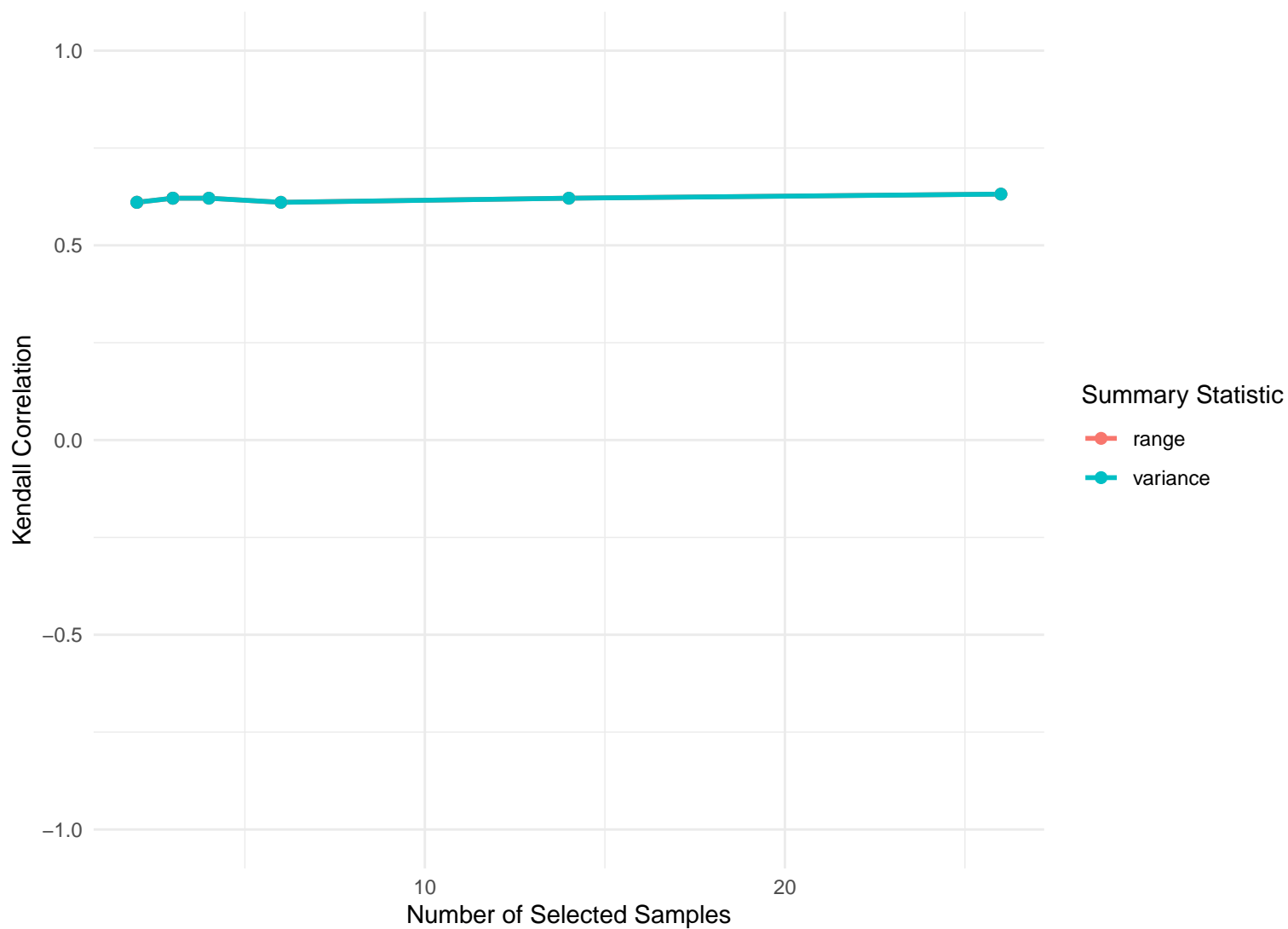
Number of Selected Samples

10

20



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



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

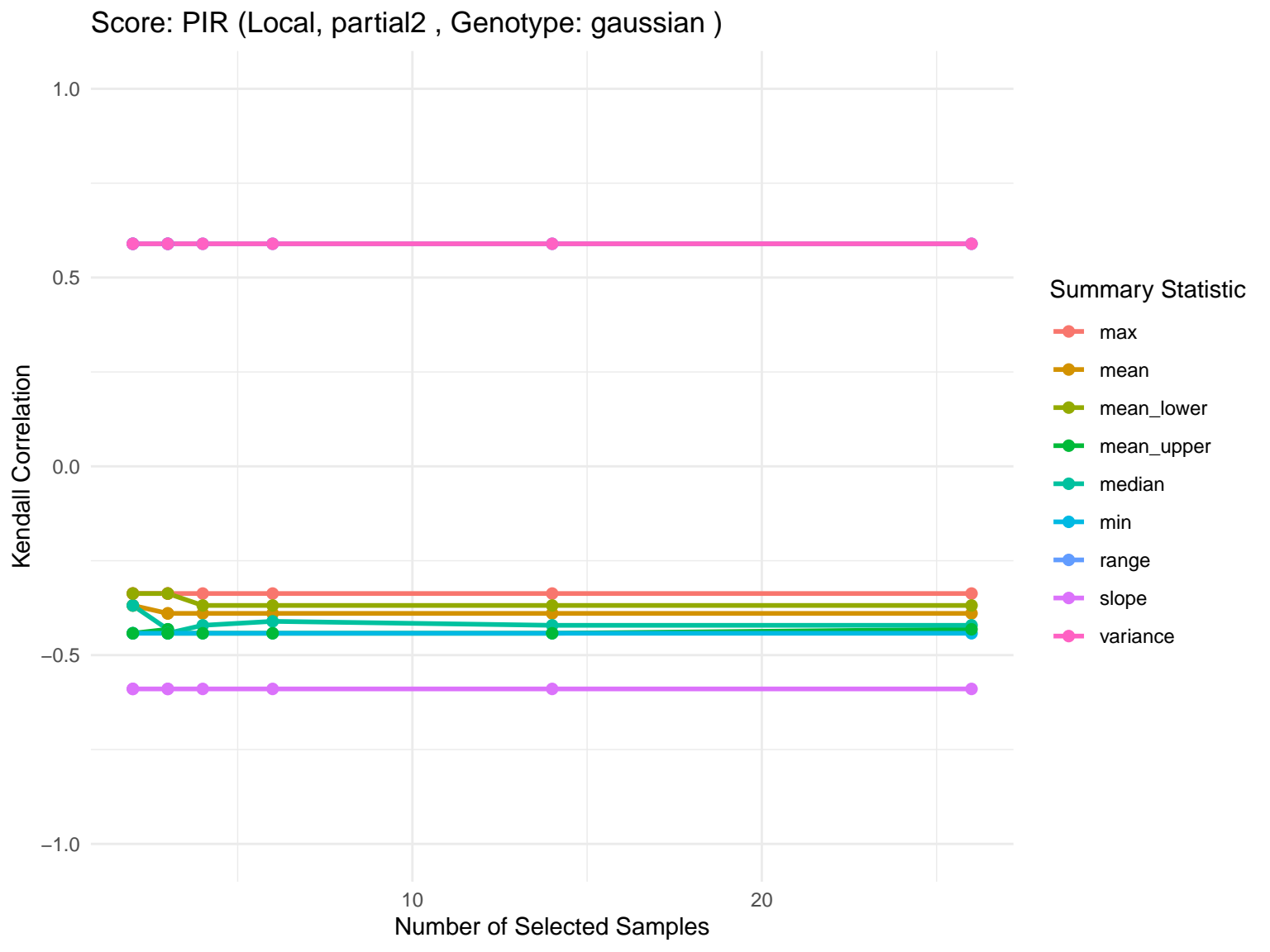
Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

1.0  
0.5  
0.0  
-0.5  
-1.0

10 20  
Number of Selected Samples

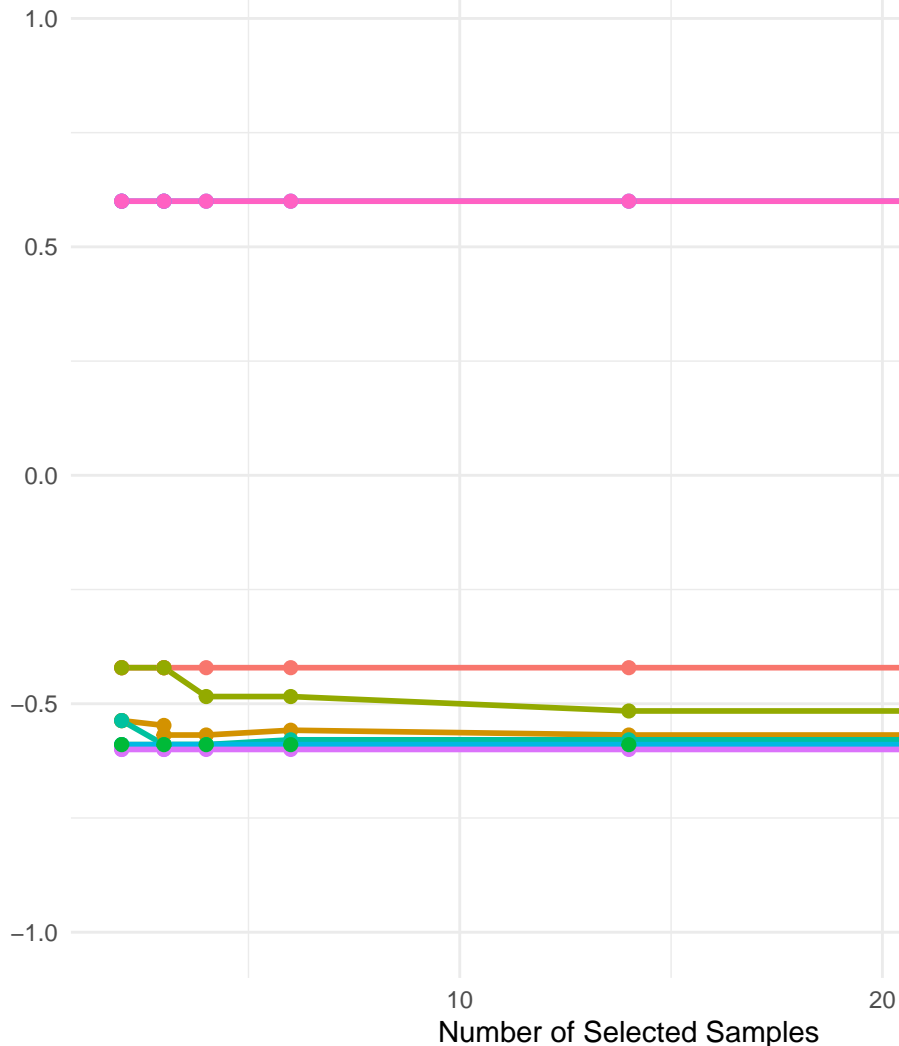


Score: PIR (Local, partial2 , Genotype: sinusoidal )

Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

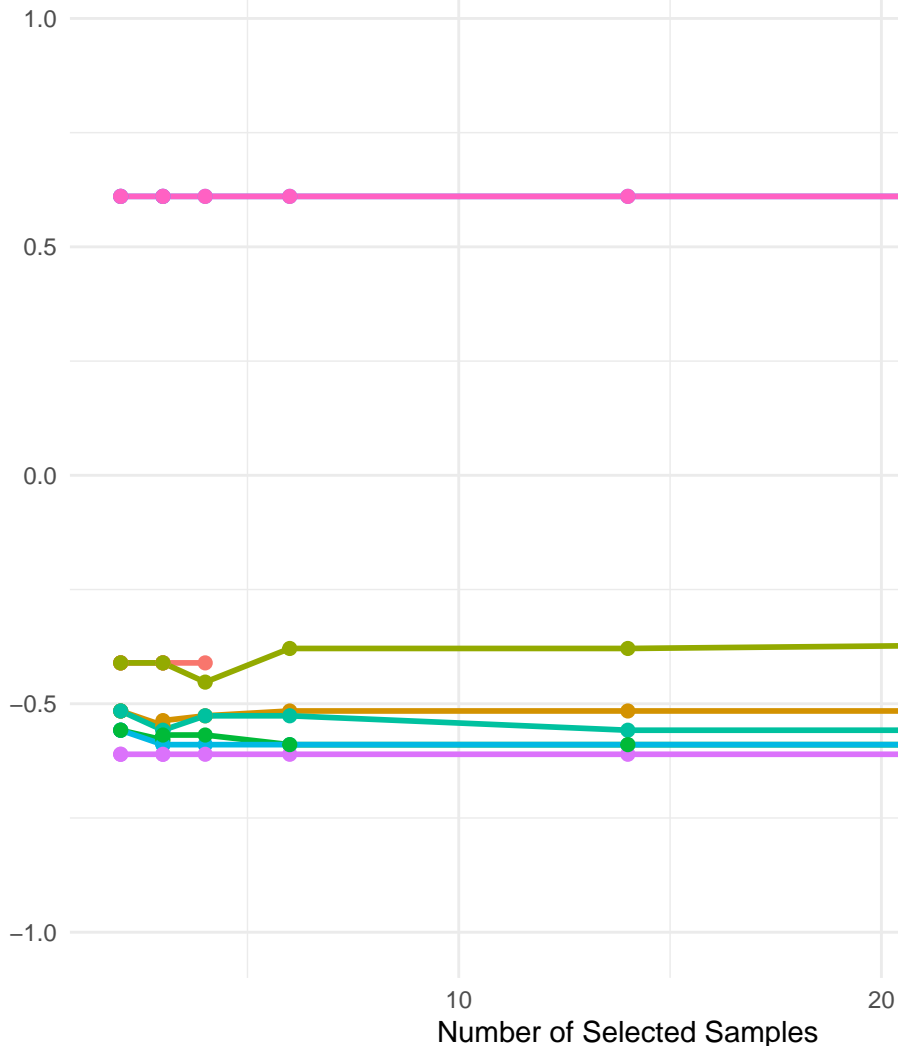


Score: PIR (Local, partial2 , Genotype: wave )

Kendall Correlation

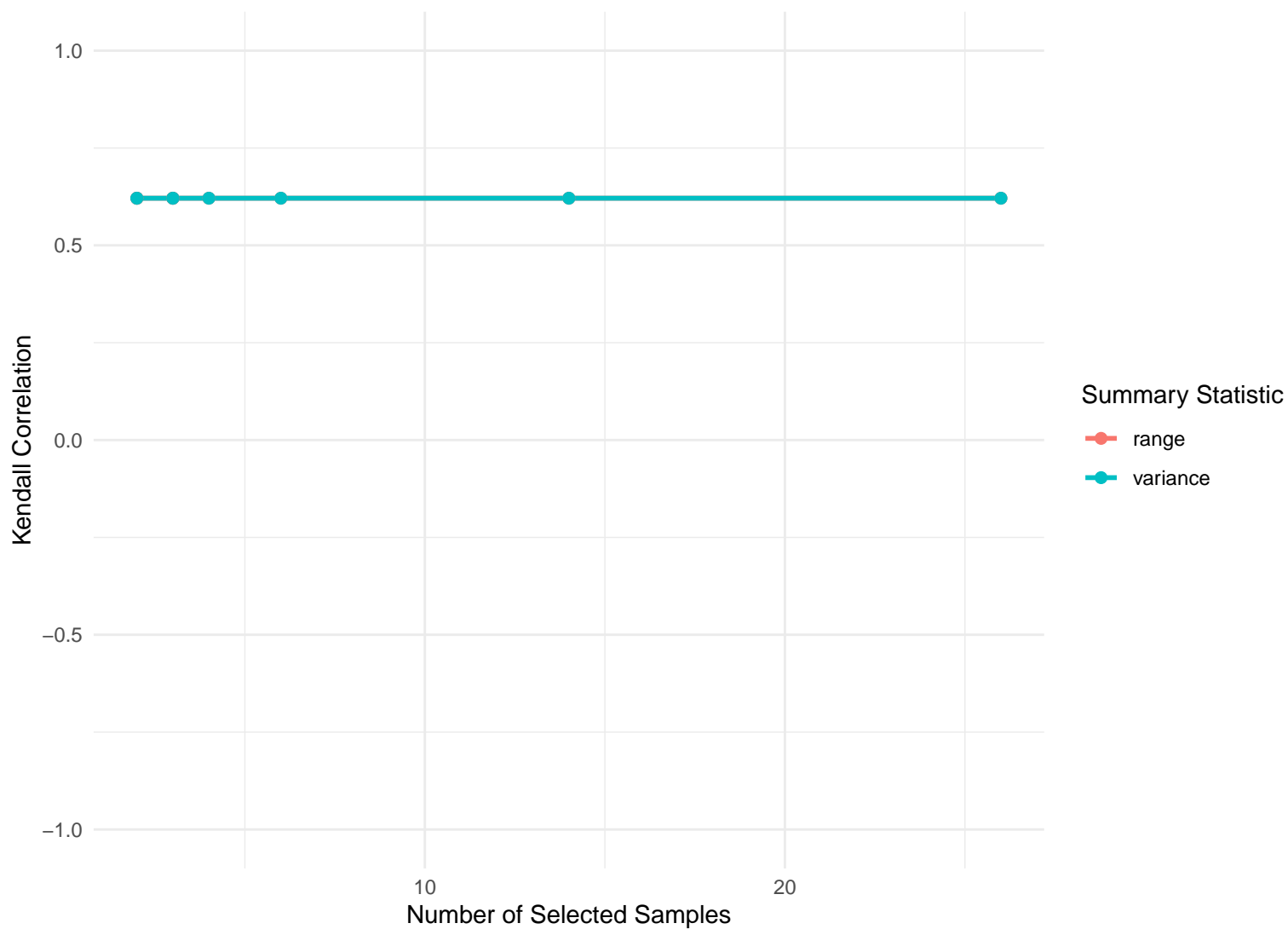
Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance





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

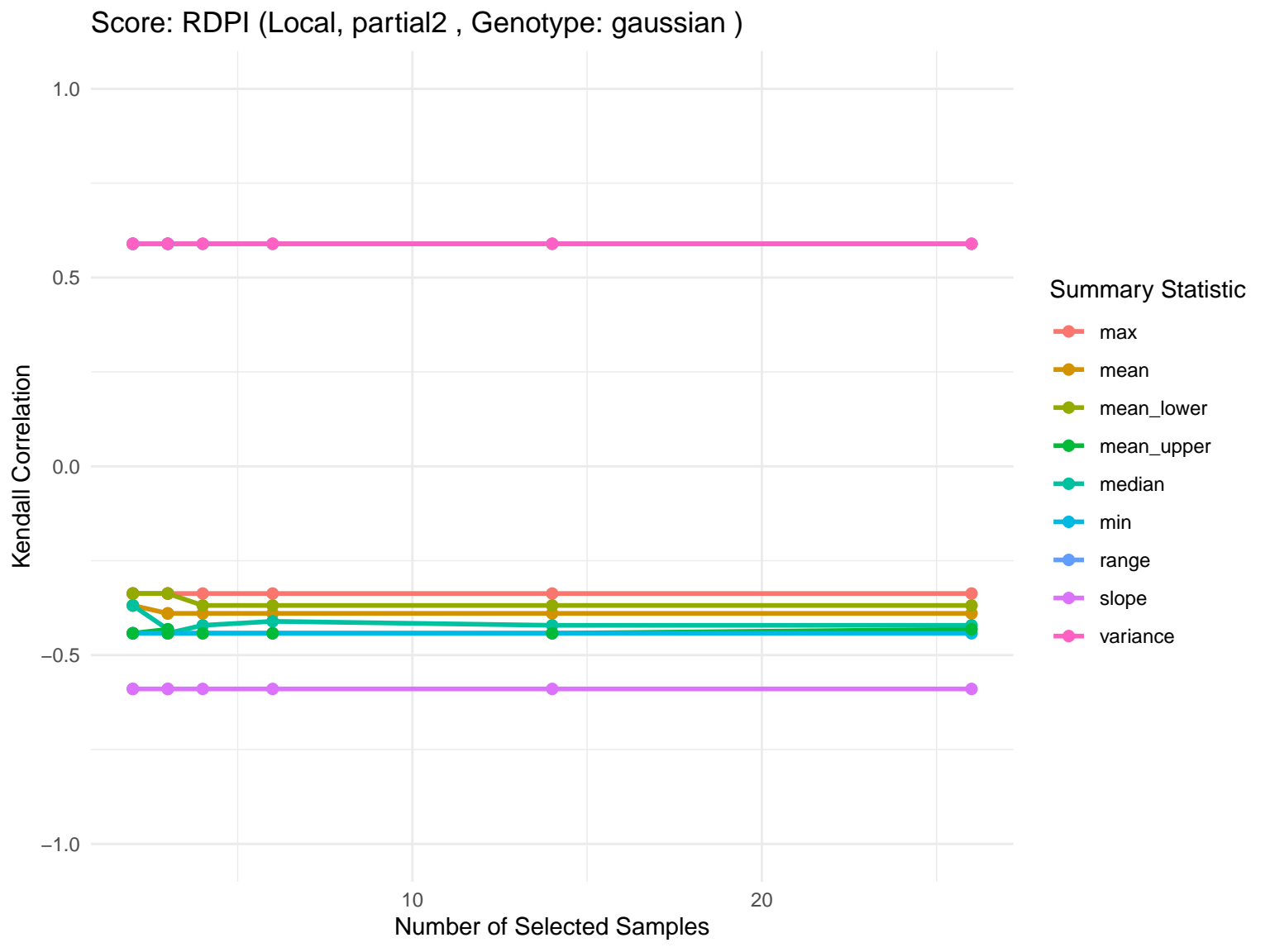
-0.5

-1.0

Number of Selected Samples

10

20



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



Number of Selected Samples

## Summary Statistic

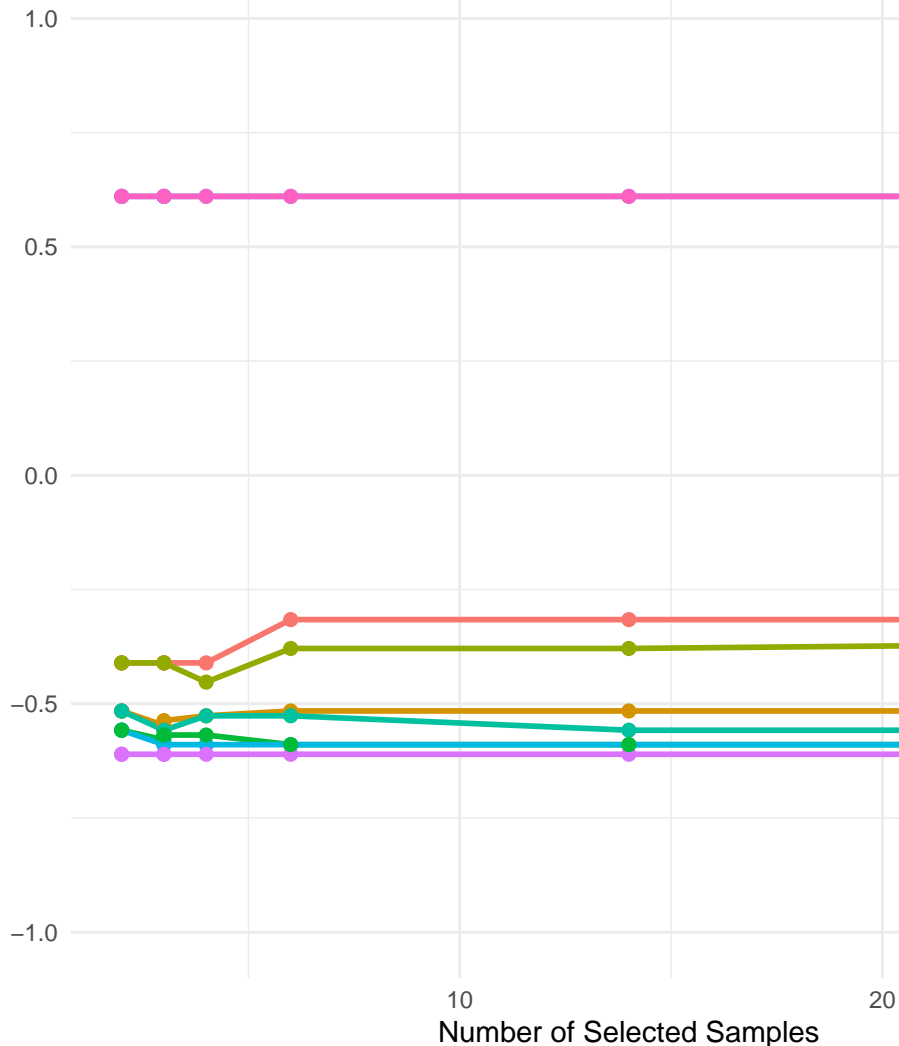
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

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 )

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

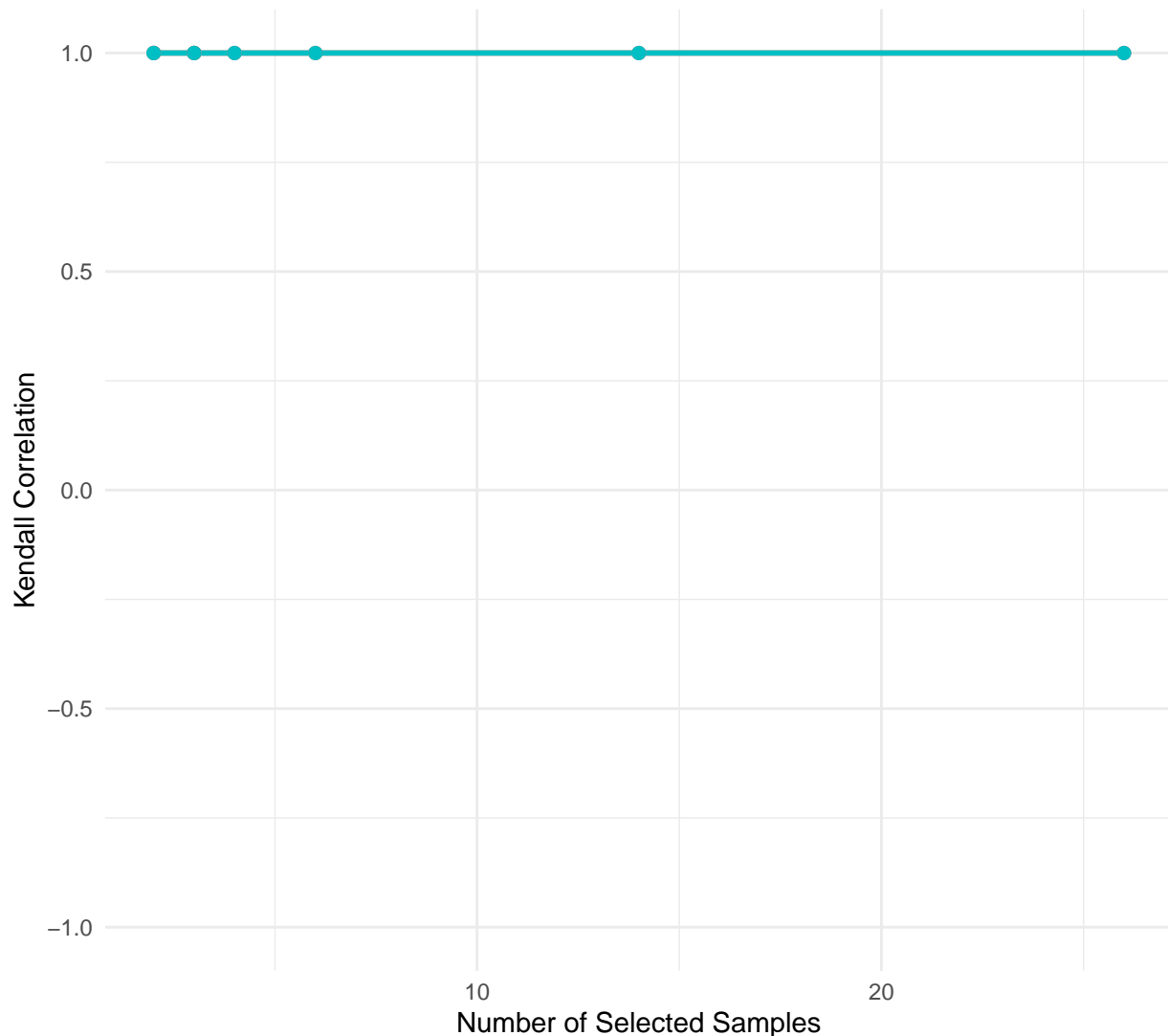
20

Number of Selected Samples

Summary Statistic

range

variance



Score: ESPI (Local, 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

1.0

0.5

0.0

-0.5

-1.0

1.0

0.5

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-0.5

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-1.0

1.0

0.5

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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

10

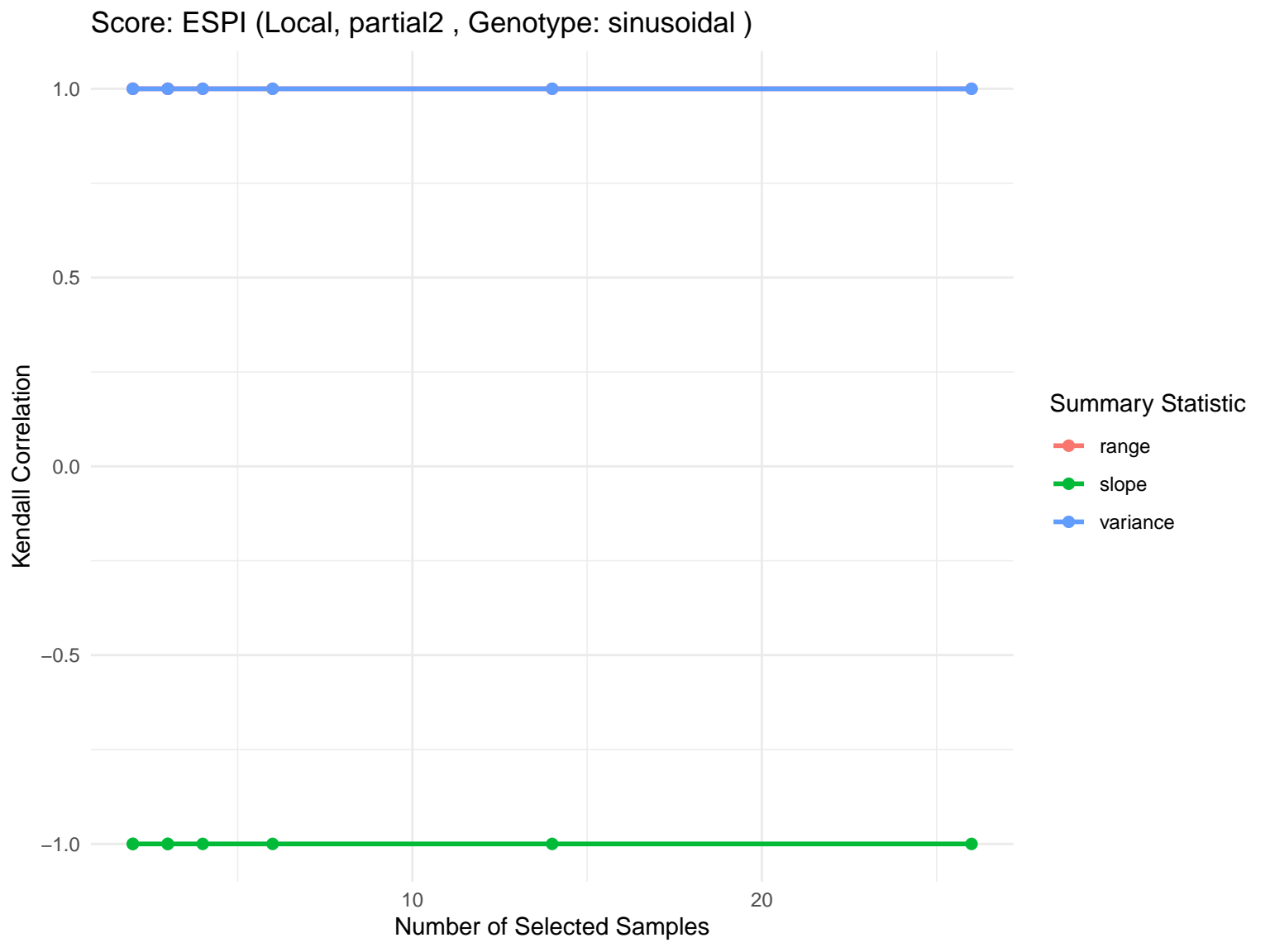
20

Summary Statistic

range

slope

variance

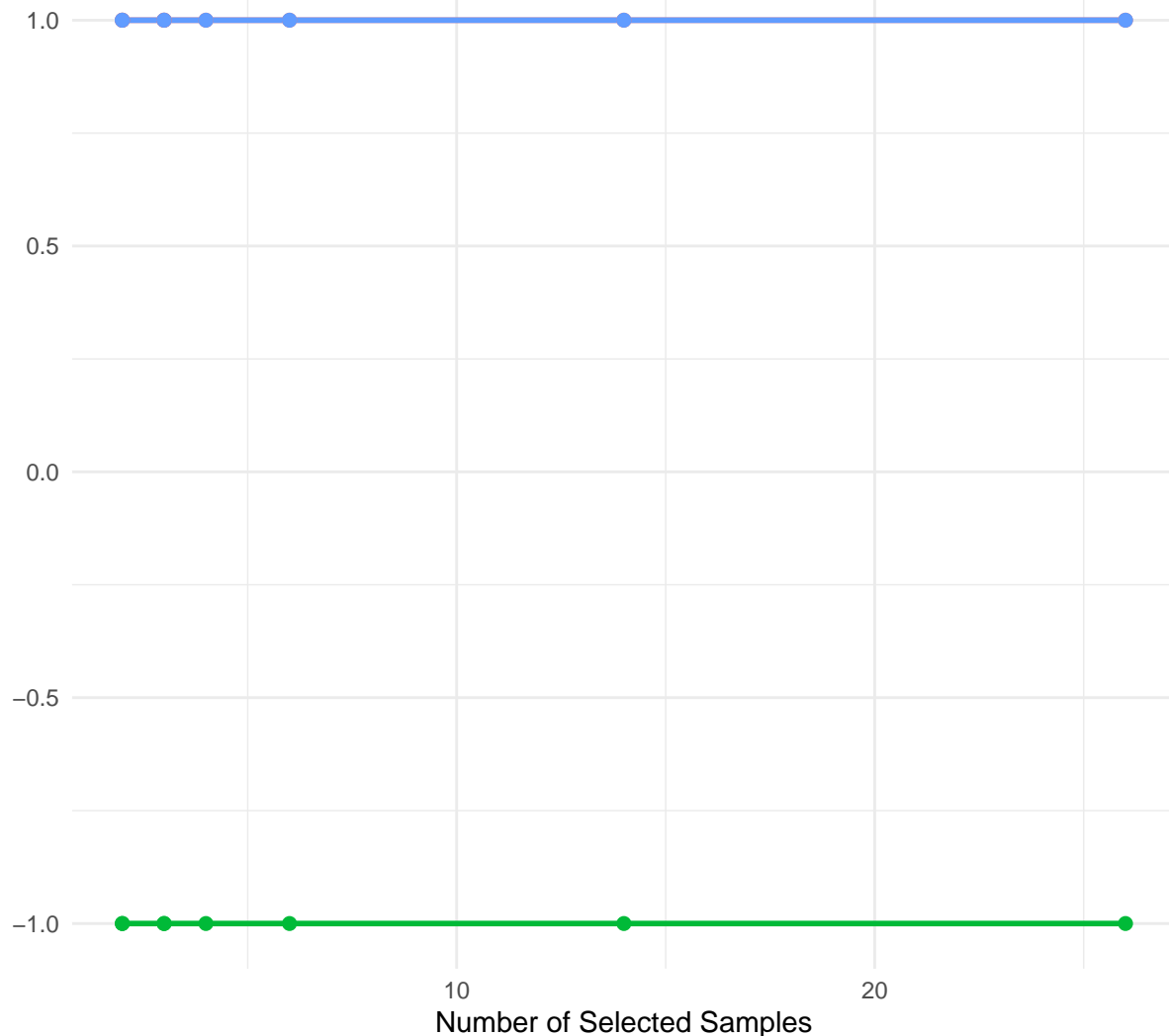


Score: ESPI (Local, partial2 , Genotype: wave )

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

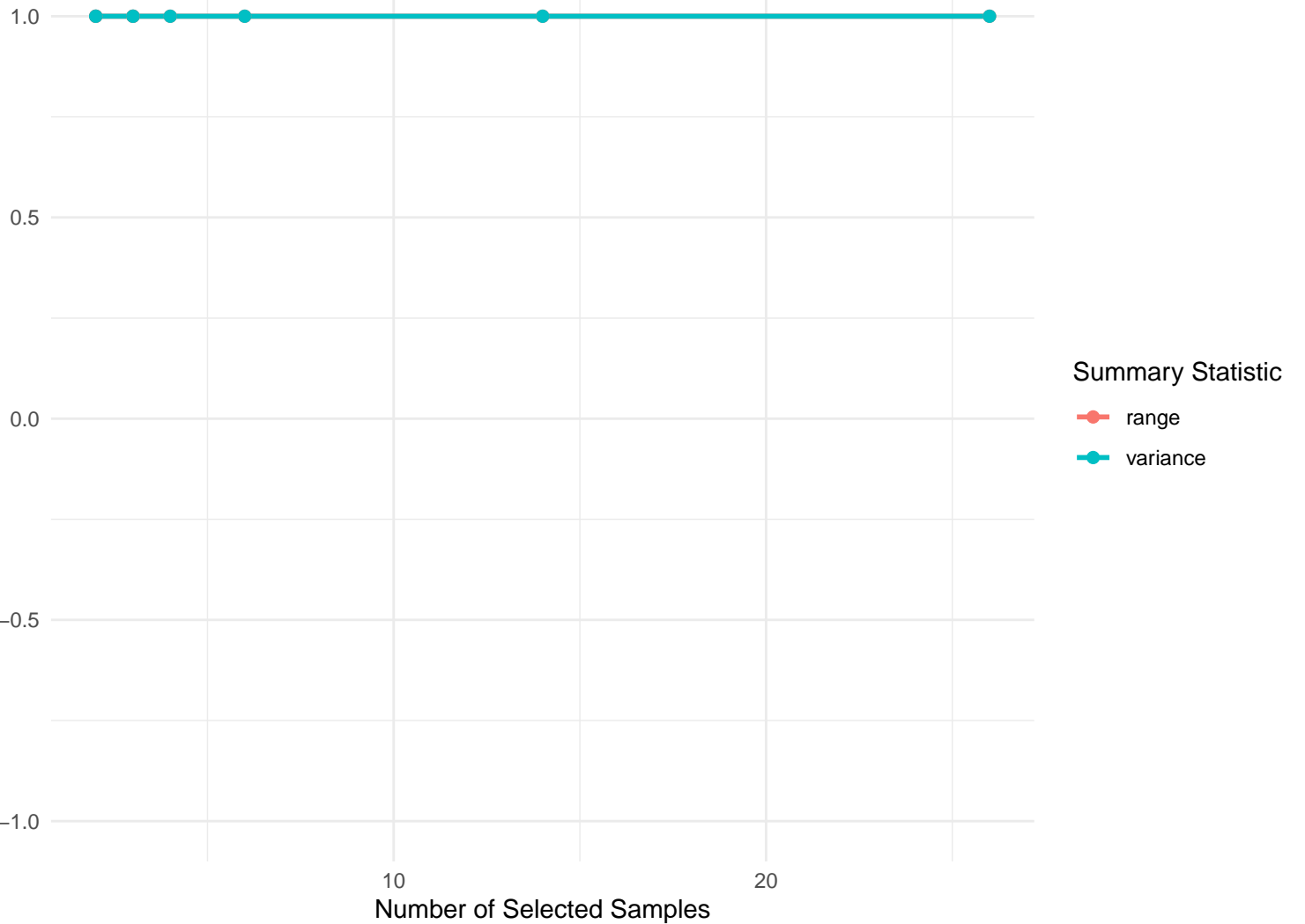
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Number of Selected Samples

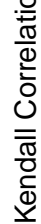
Summary Statistic

range

variance



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



### Summary Statistic

 range

 slope

 variance

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

10

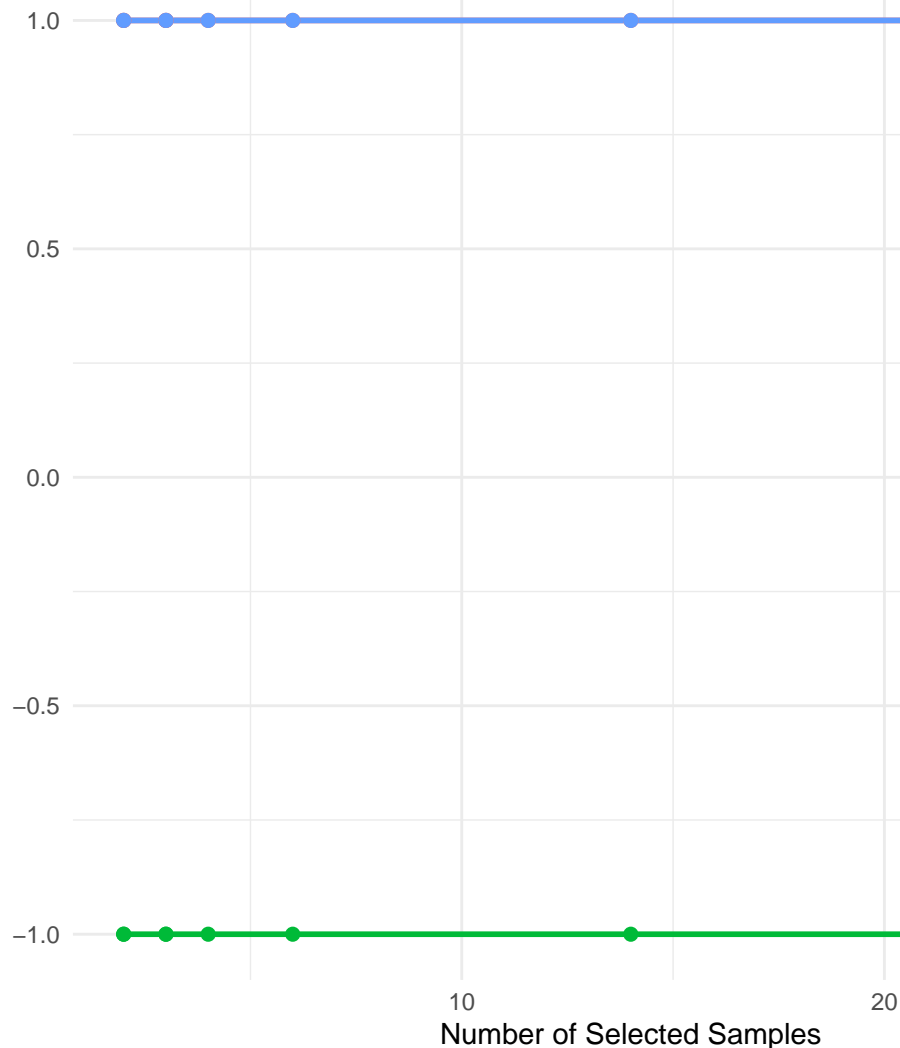
20

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

Kendall Correlation

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

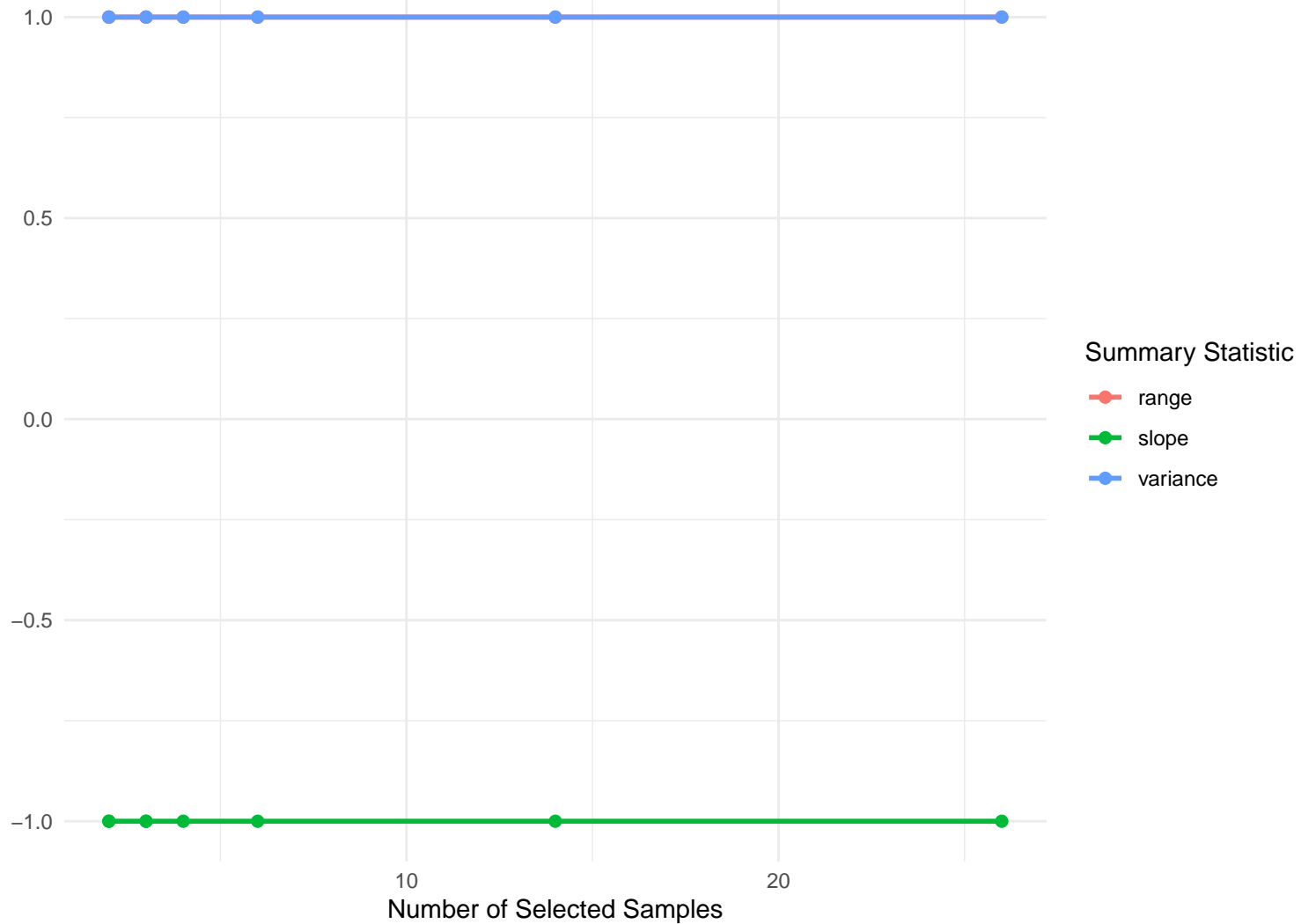
10

20

Number of Selected Samples

Summary Statistic

- range
- slope
- variance

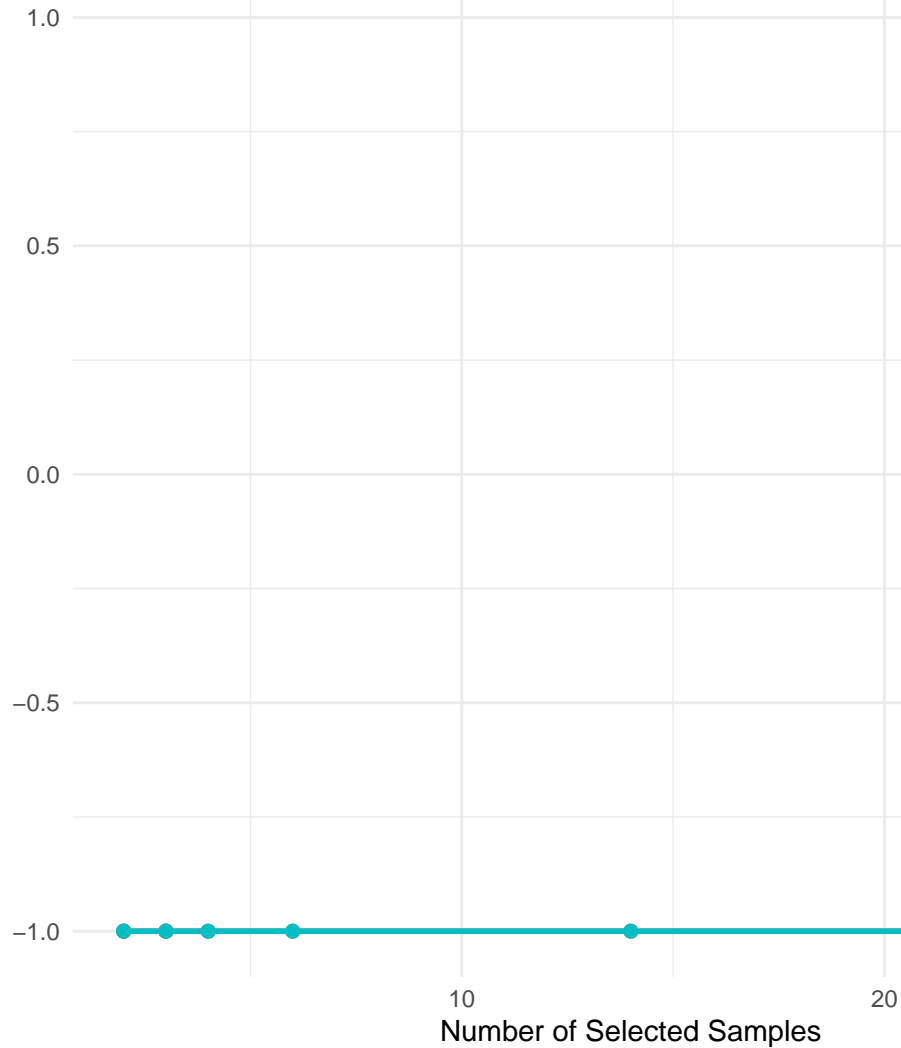


Score: PSI (Local, partial2 , Genotype: linear )

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

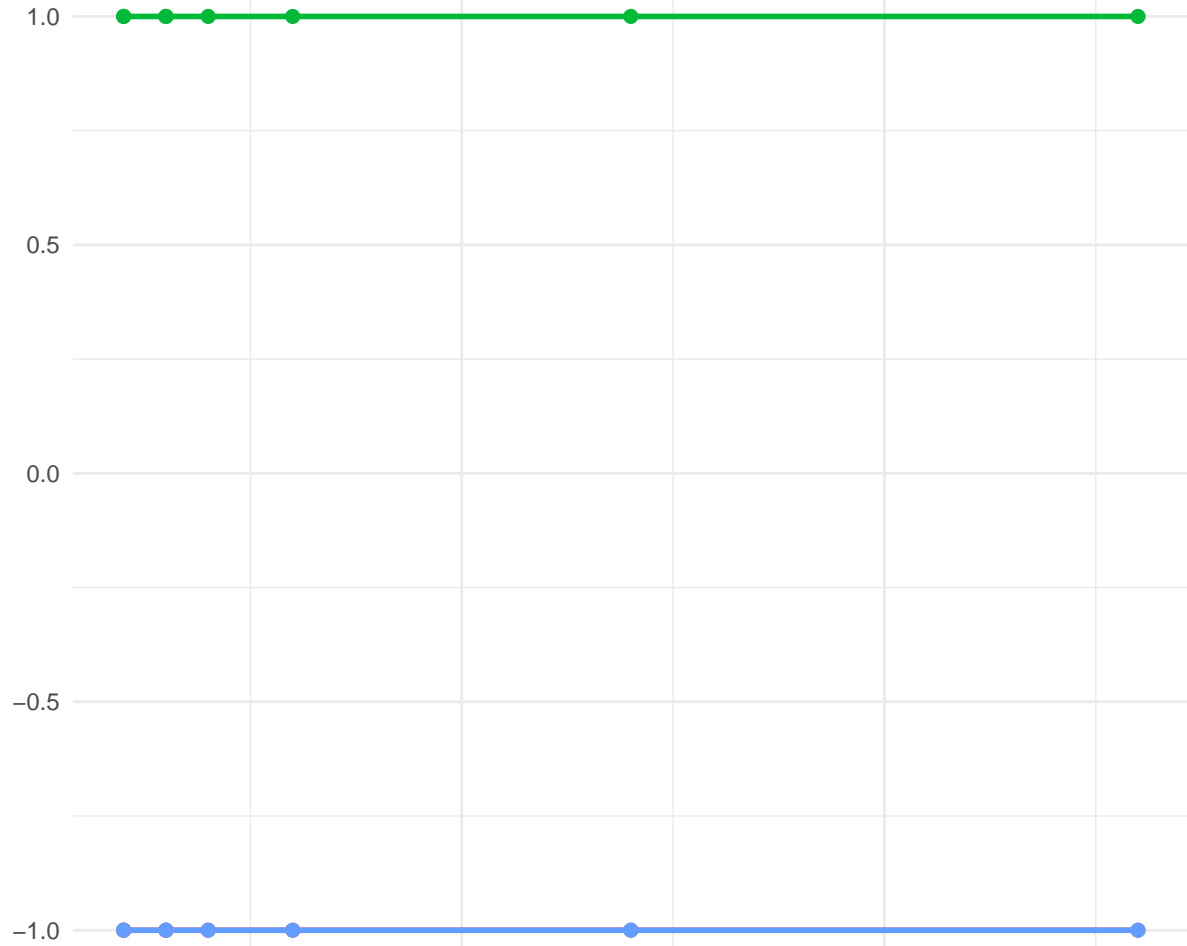
20

Summary Statistic

range

slope

variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

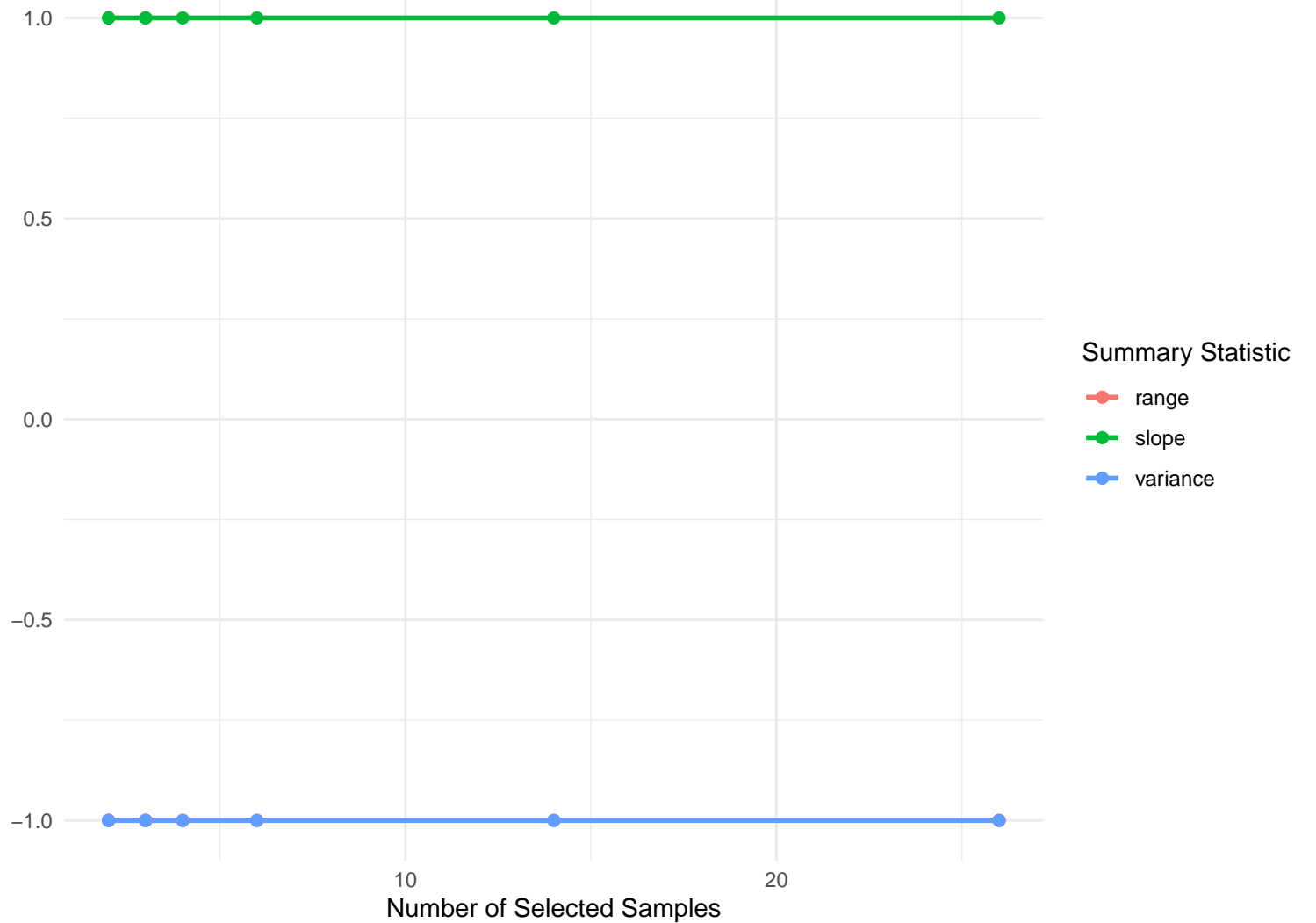
10

20

Number of Selected Samples

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

10

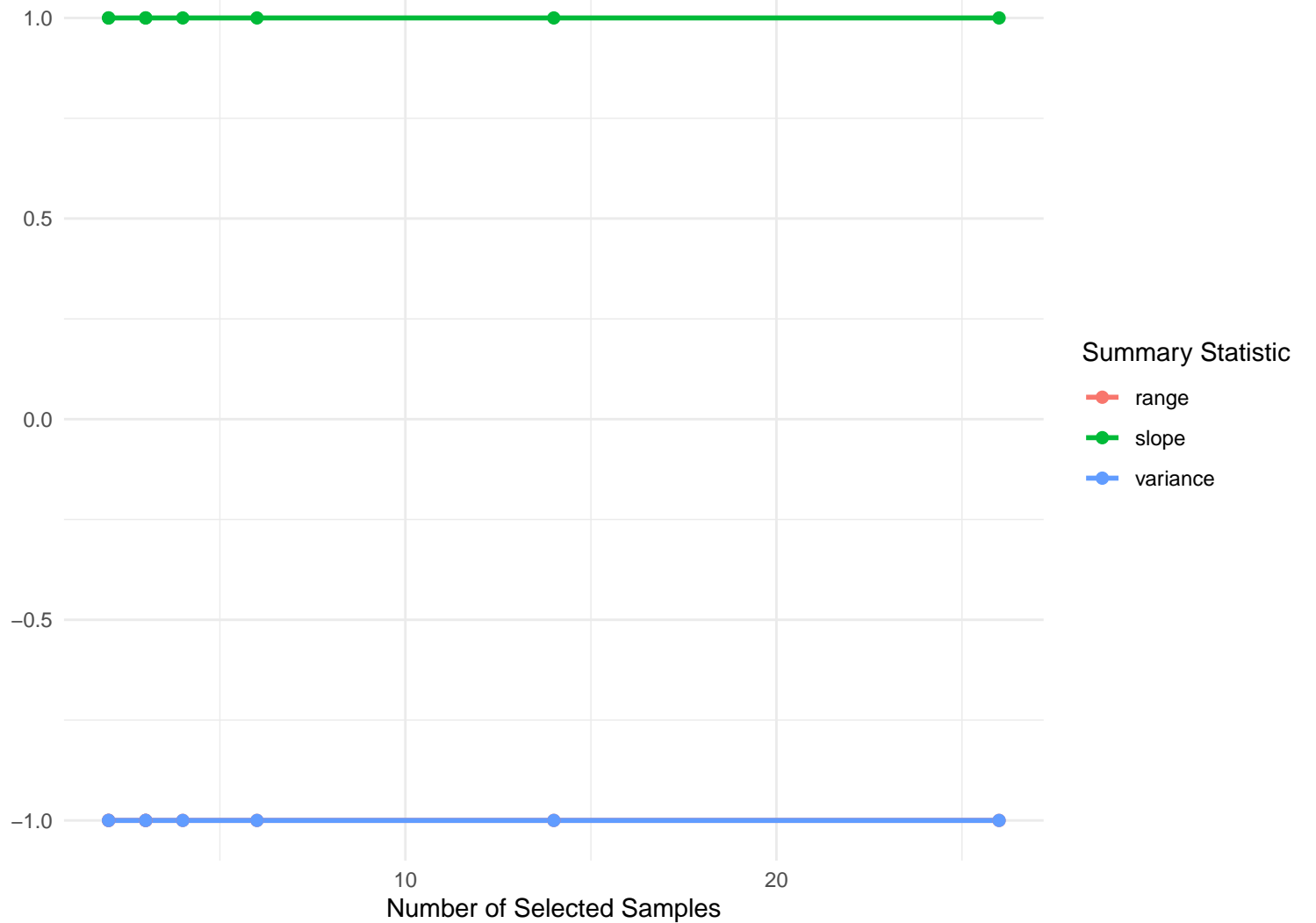
20

Summary Statistic

range

slope

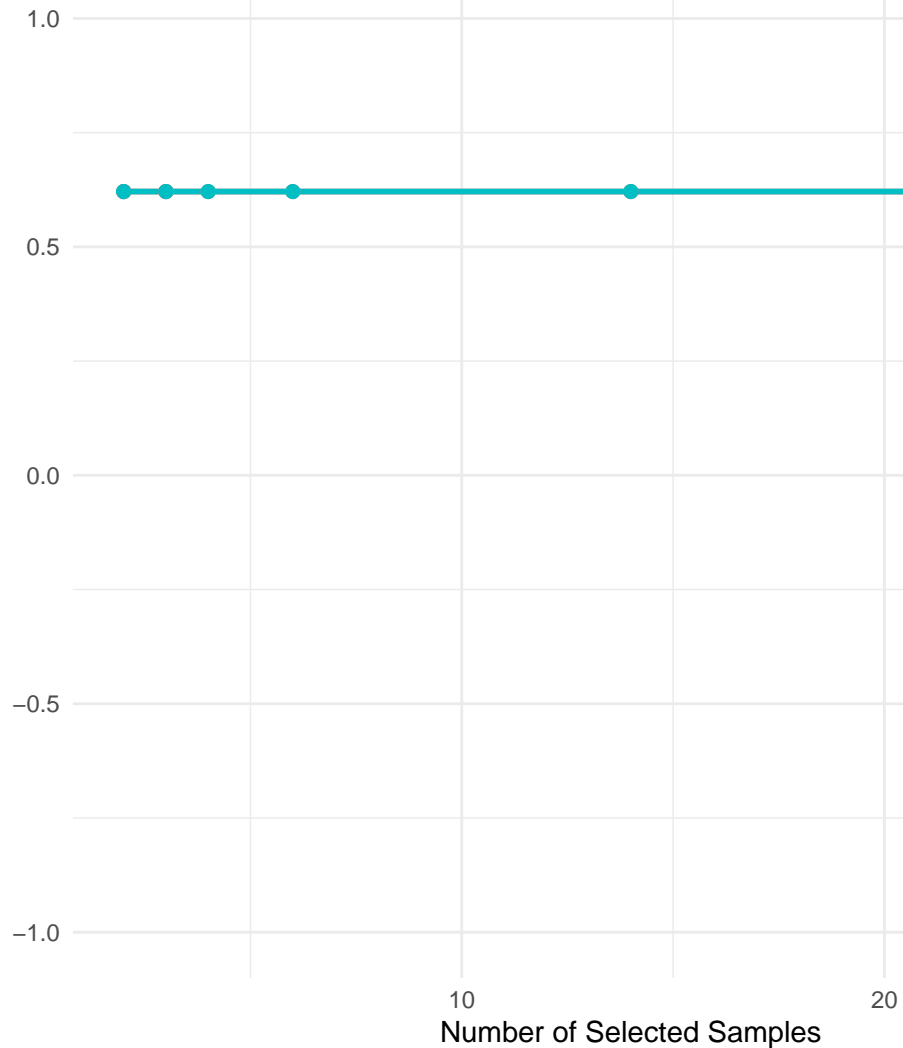
variance





Score: 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

1.0

0.5

0.0

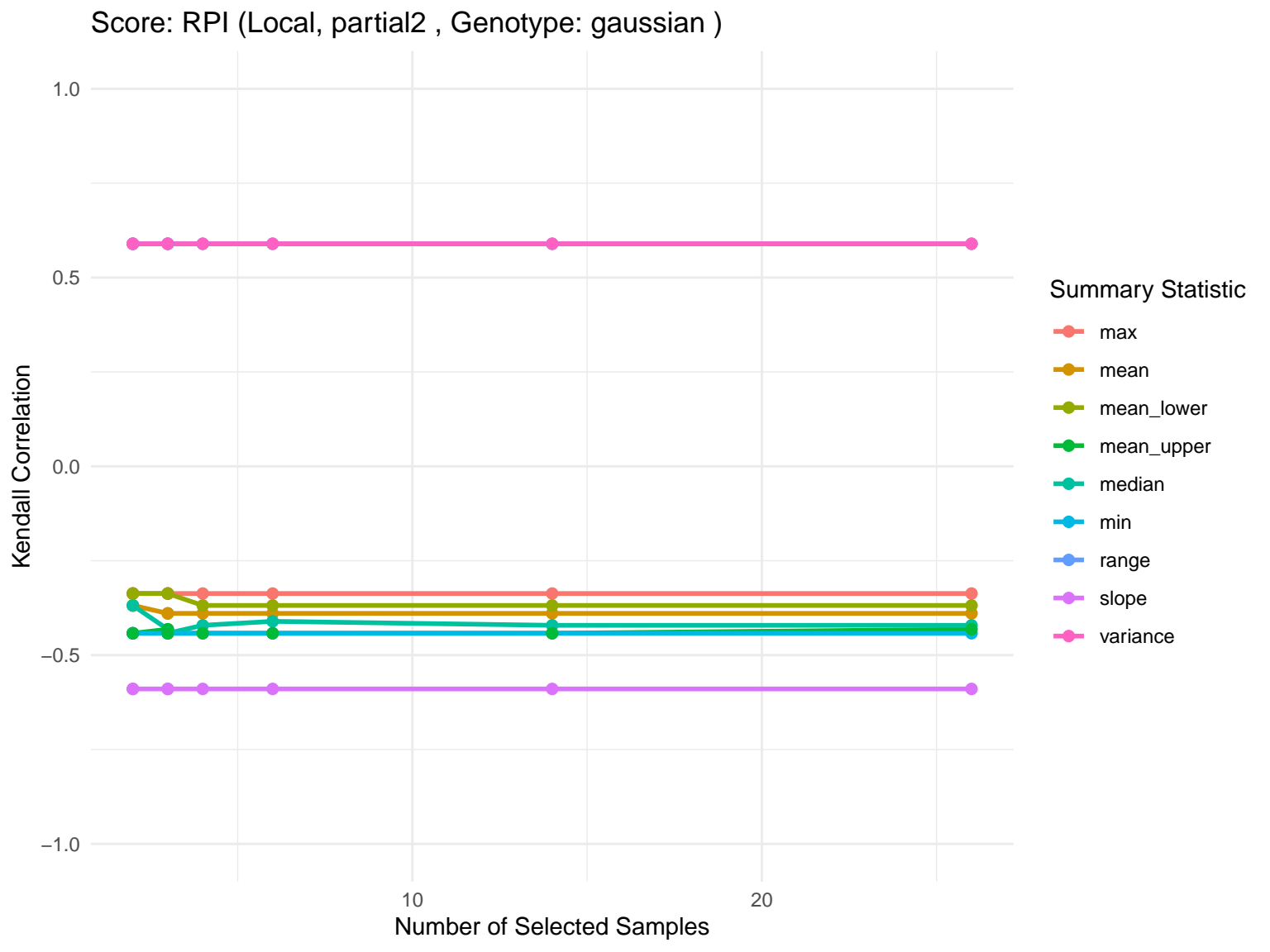
-0.5

-1.0

10

20

Number of Selected Samples

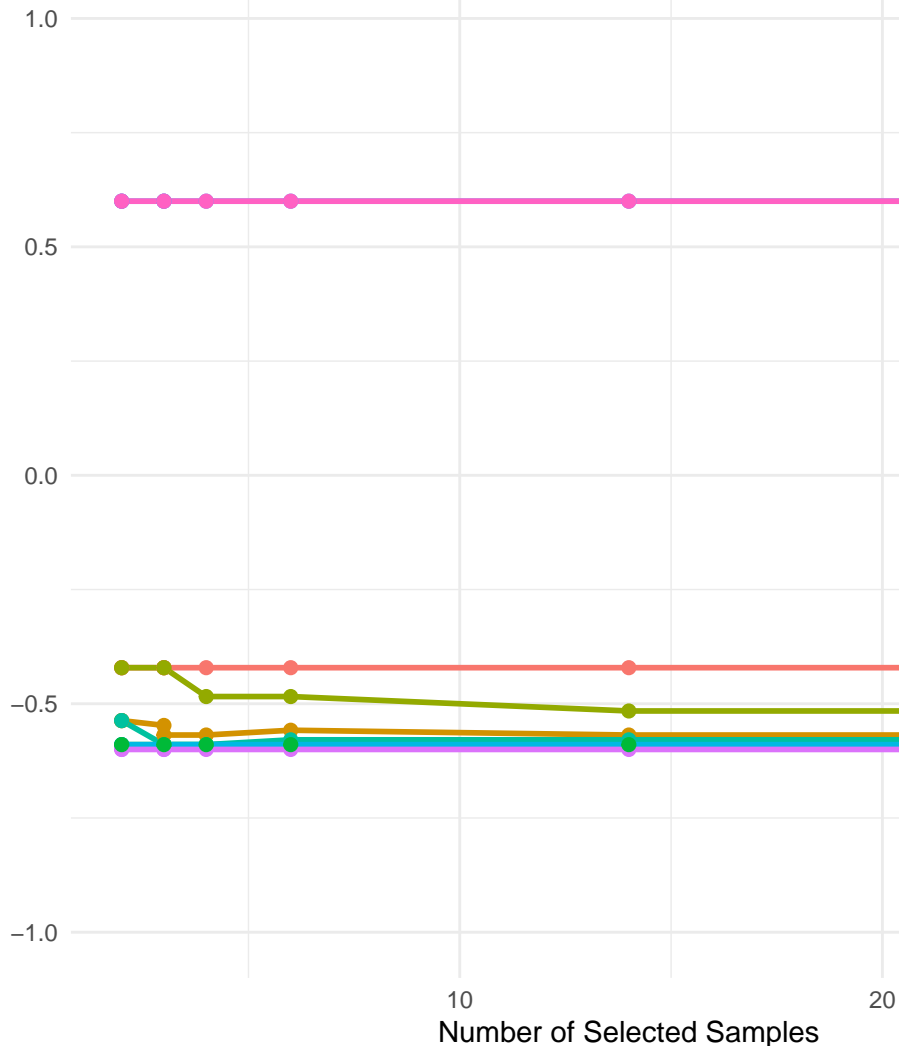


Score: RPI (Local, partial2 , Genotype: sinusoidal )

Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

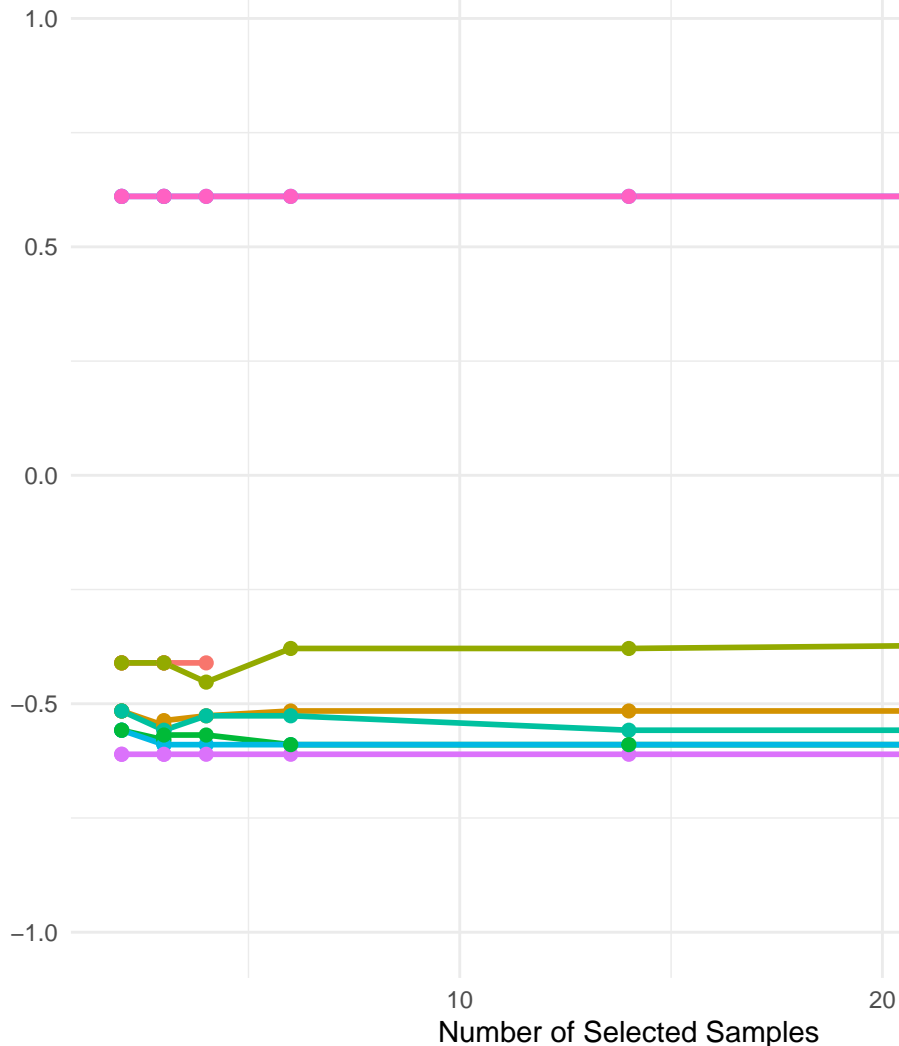


Score: RPI (Local, partial2 , Genotype: wave )

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

10

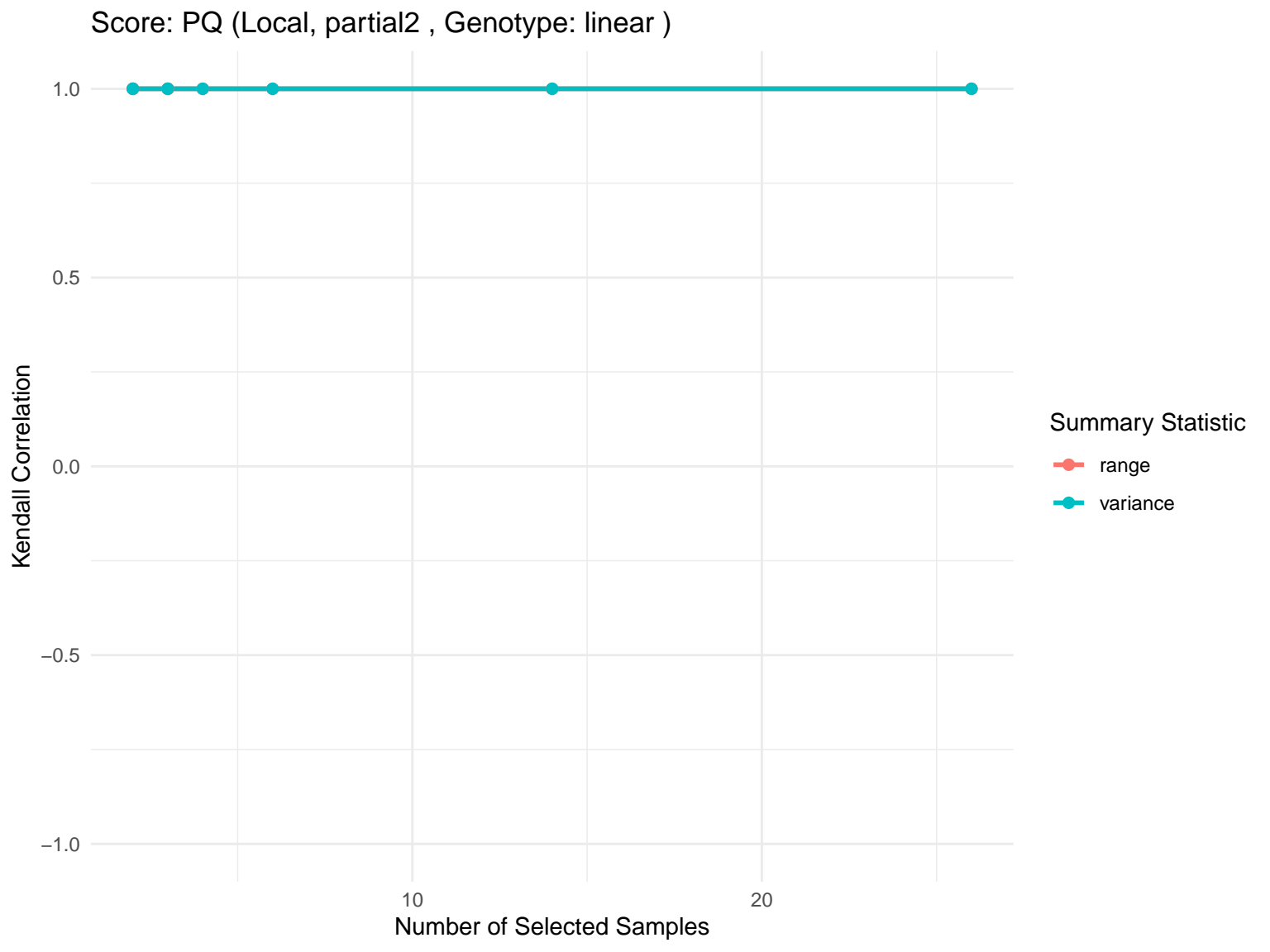
20

Number of Selected Samples

Summary Statistic

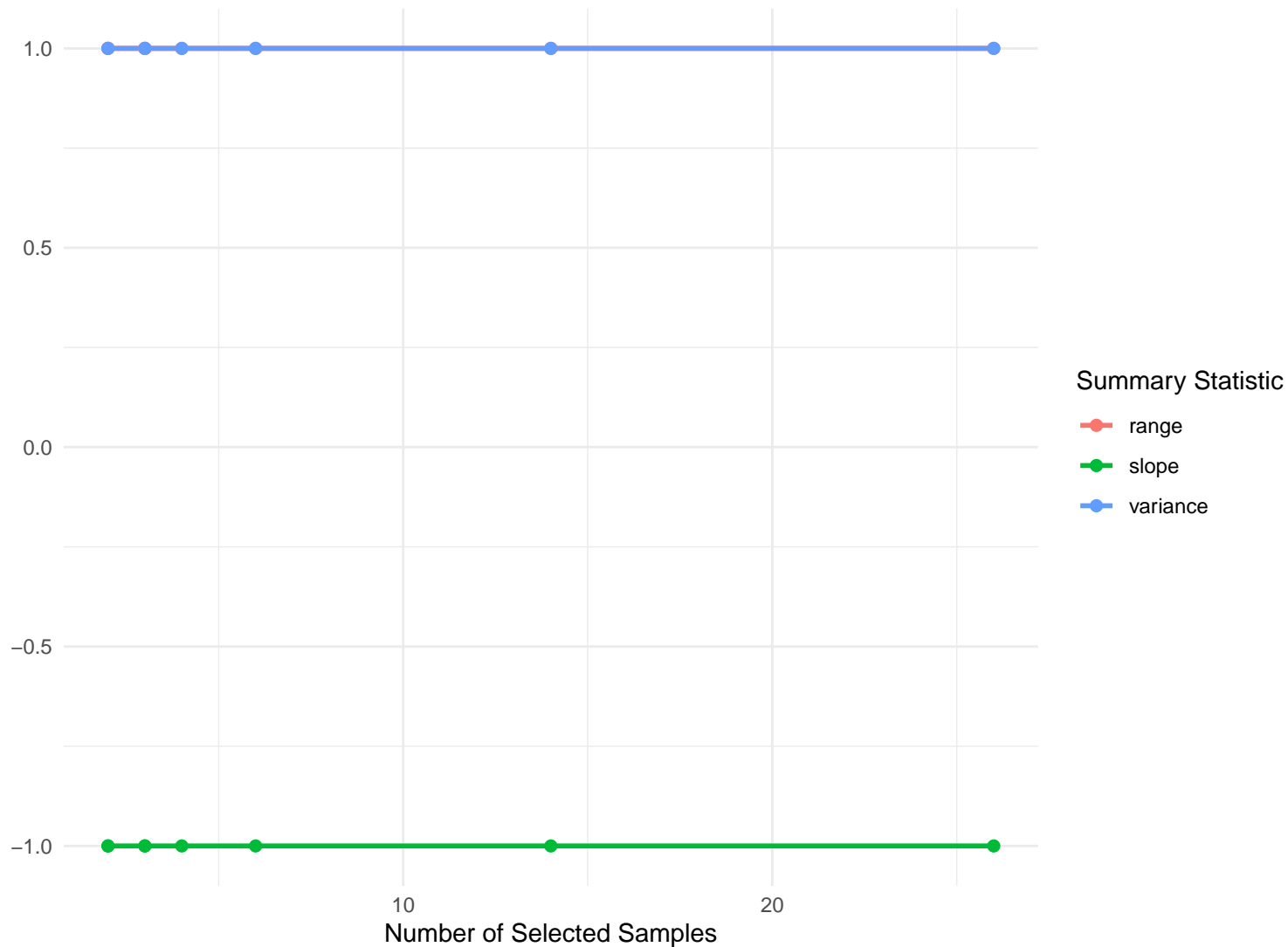
range

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

10

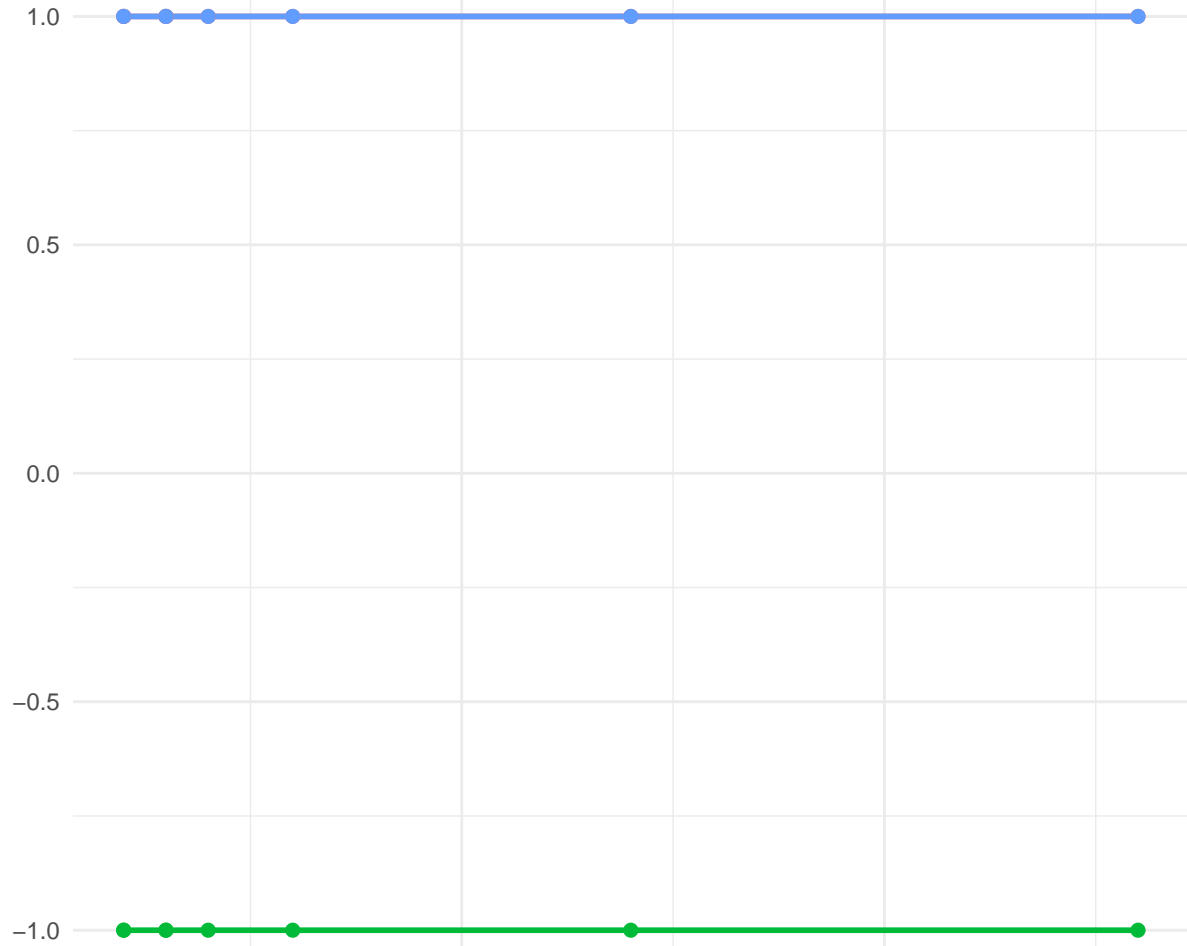
20

Summary Statistic

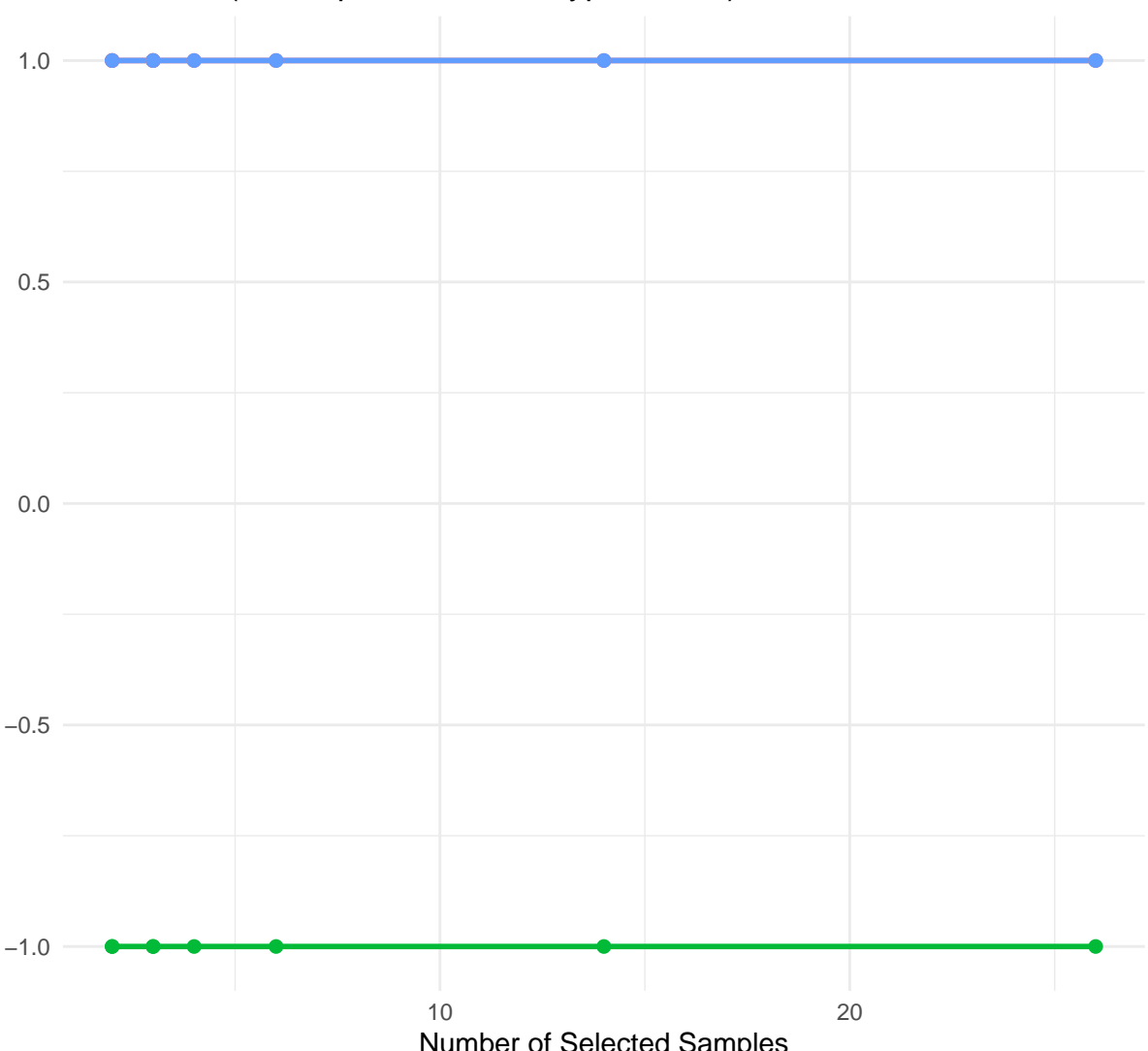
range

slope

variance



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



## Summary Statistic

 range

 slope

 variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

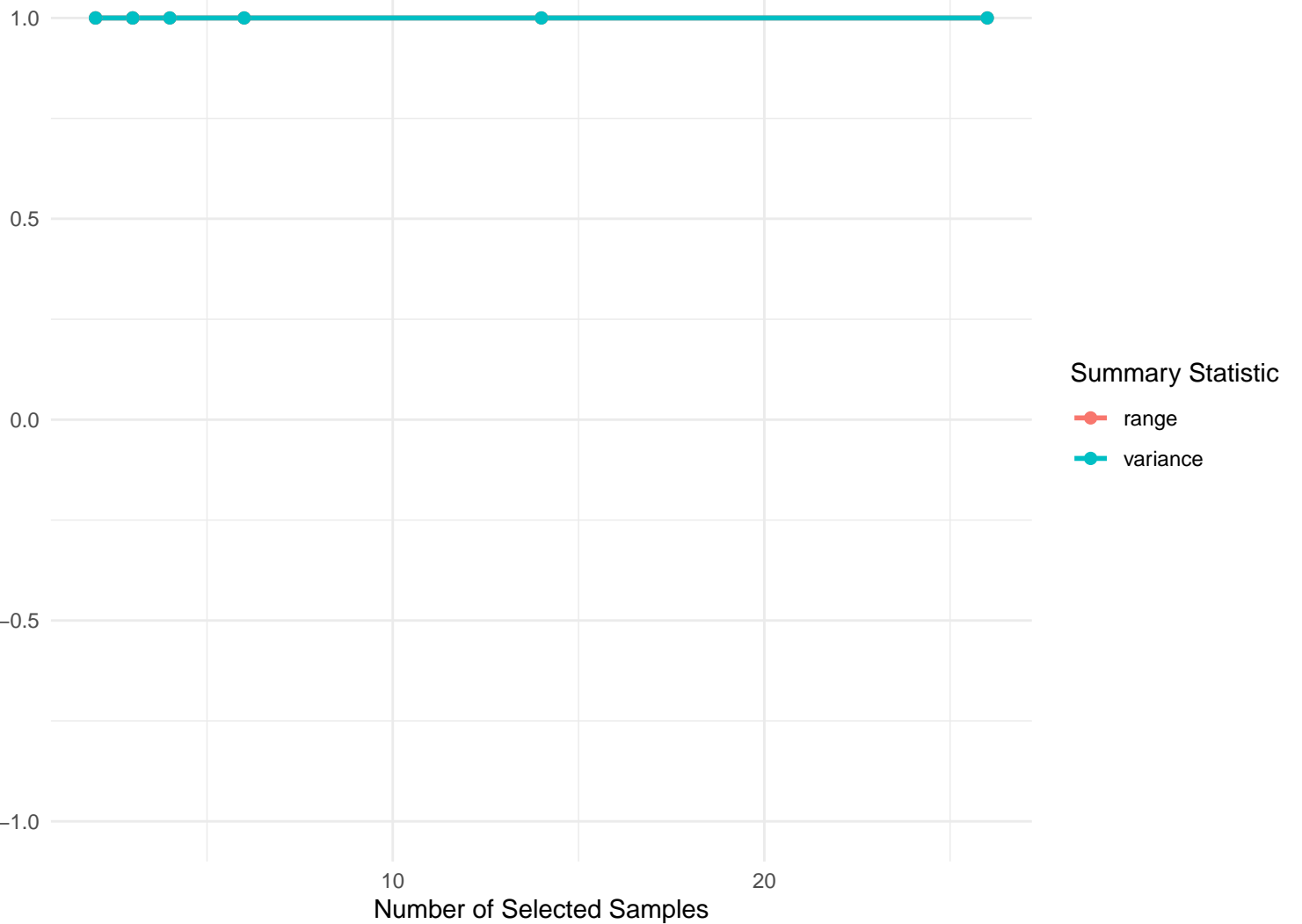
20

Number of Selected Samples

Summary Statistic

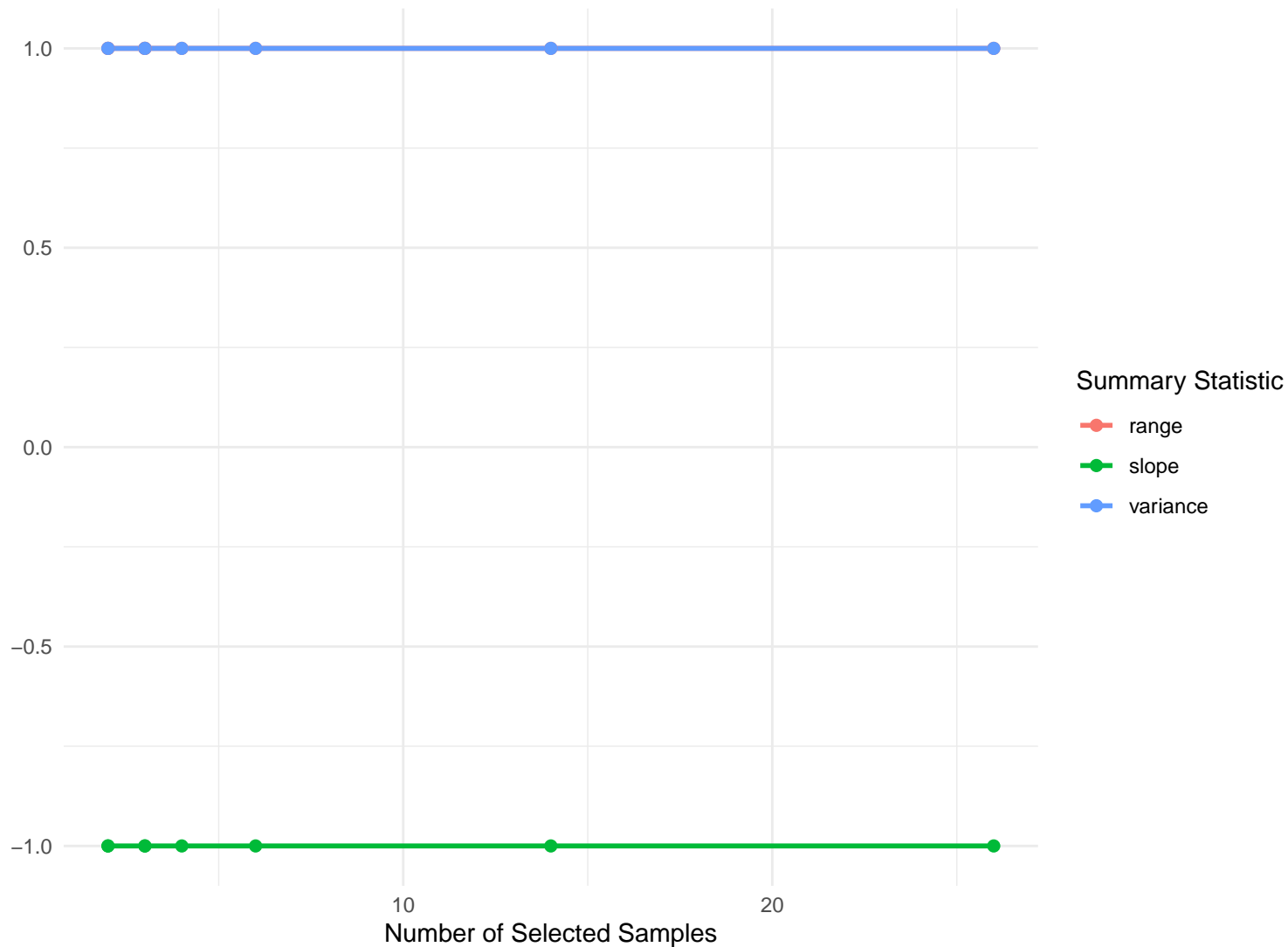
range

variance



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

Kendall Correlation



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

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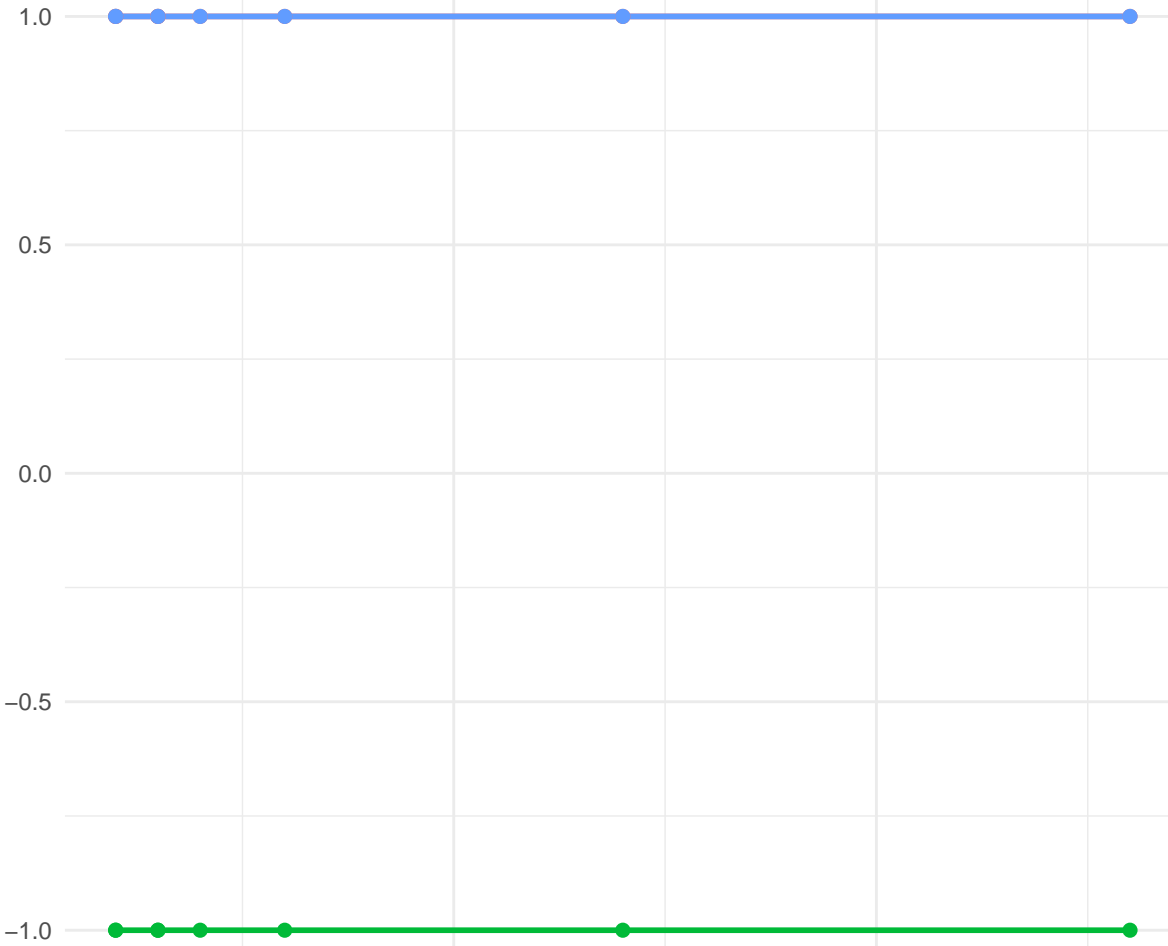
20

Summary Statistic

range

slope

variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

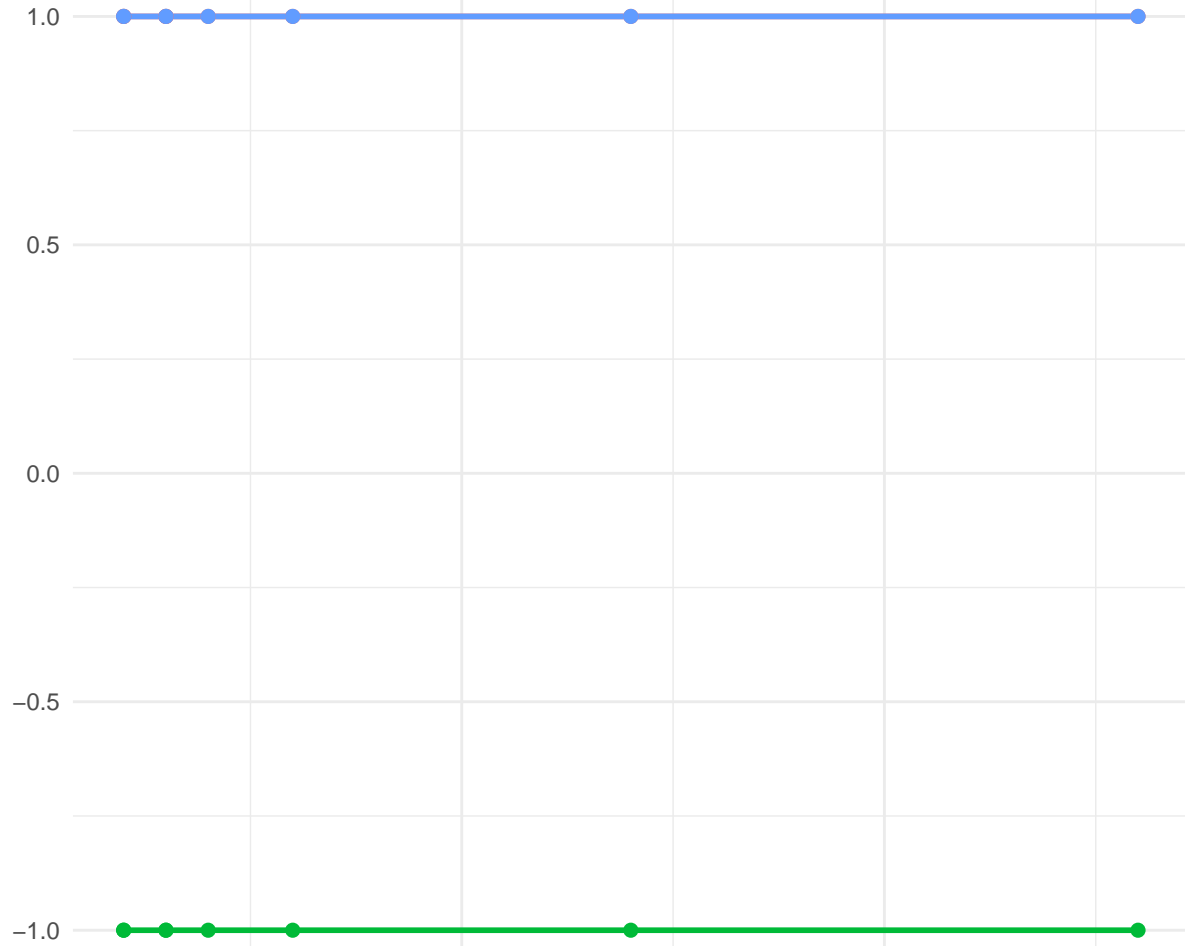
Number of Selected Samples

10

20

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

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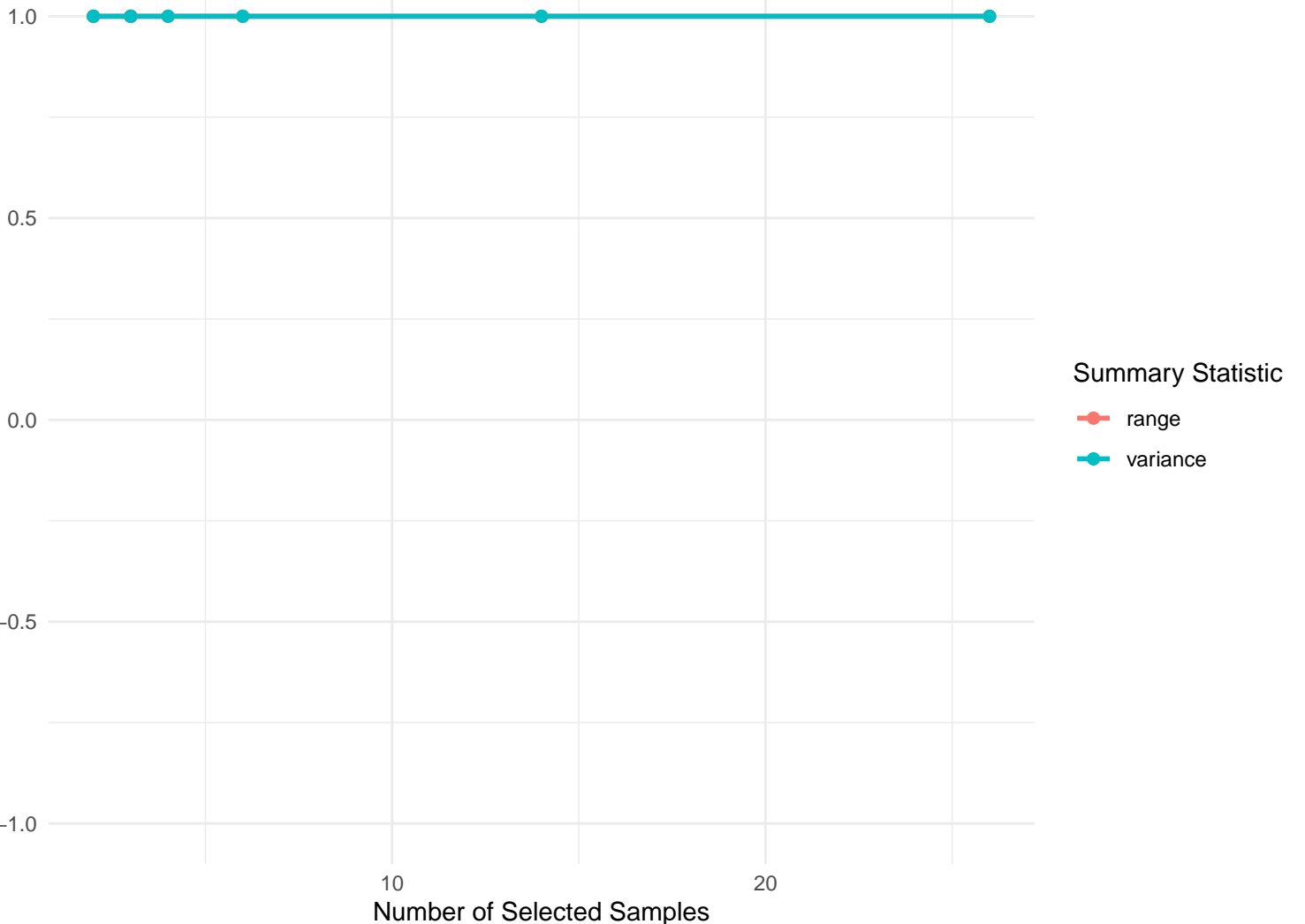
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Number of Selected Samples

Summary Statistic

range

variance

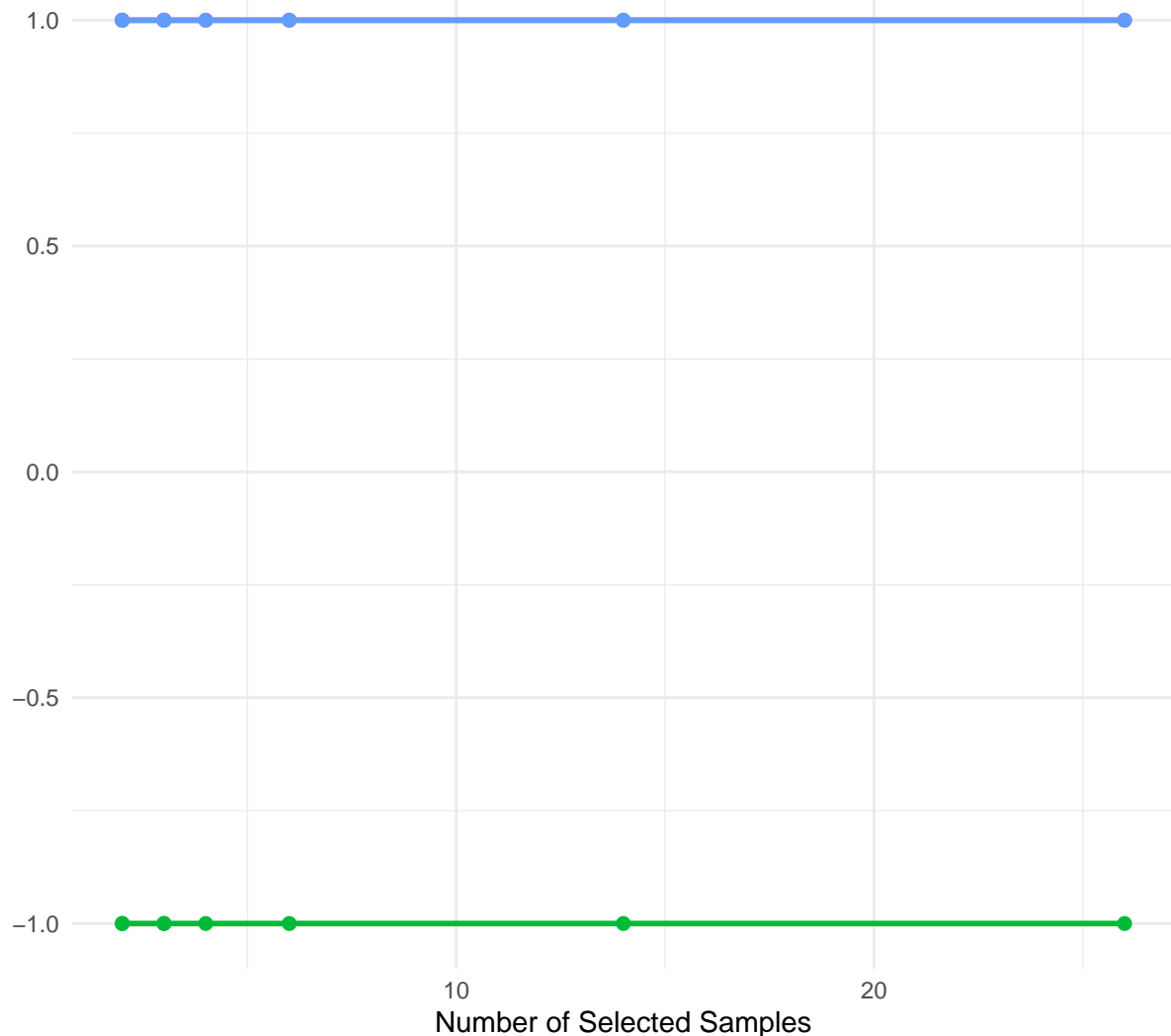


Score: NRW (Local, partial2 , Genotype: gaussian )

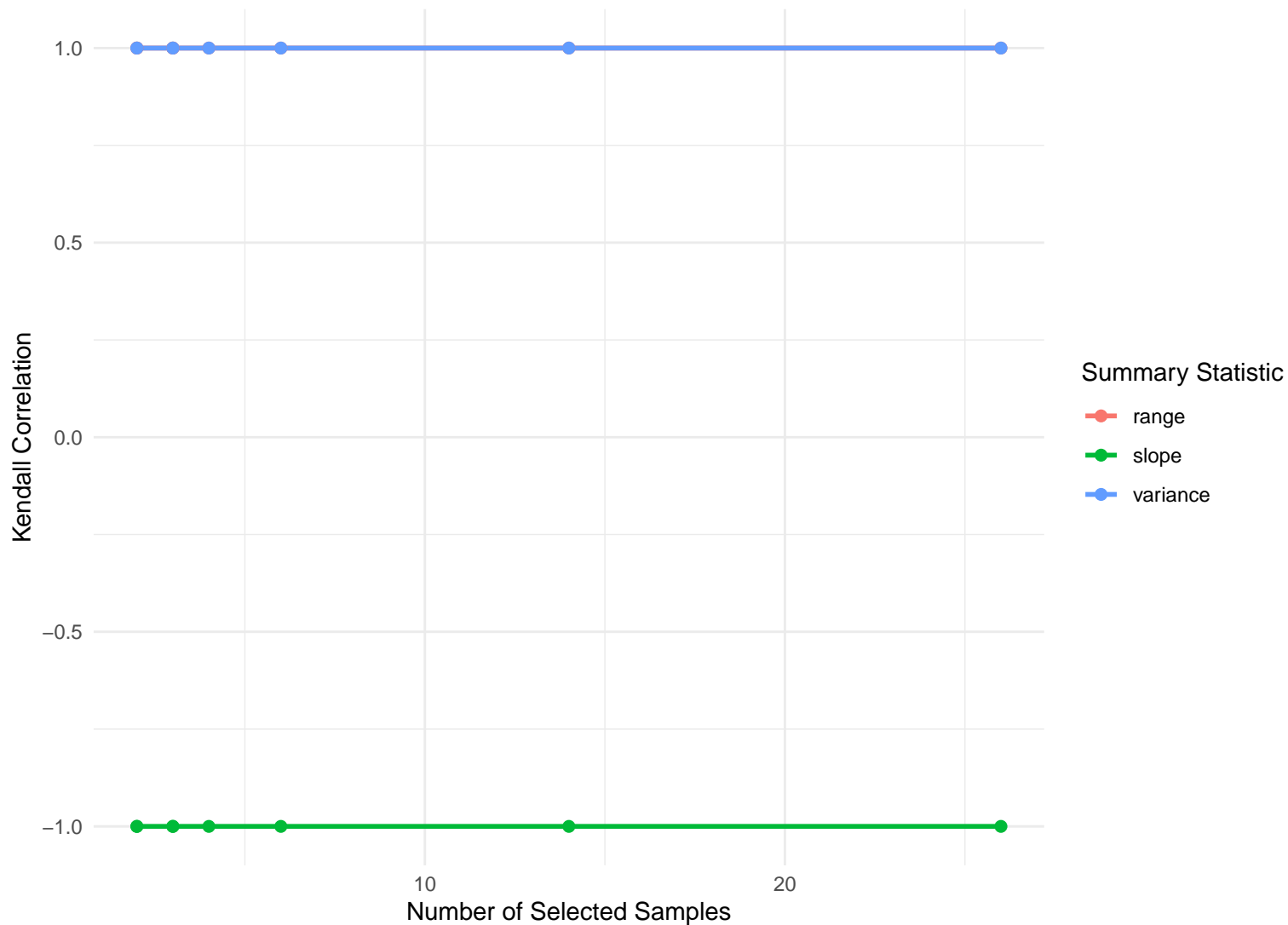
Kendall Correlation

Summary Statistic

- range
- slope
- variance



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

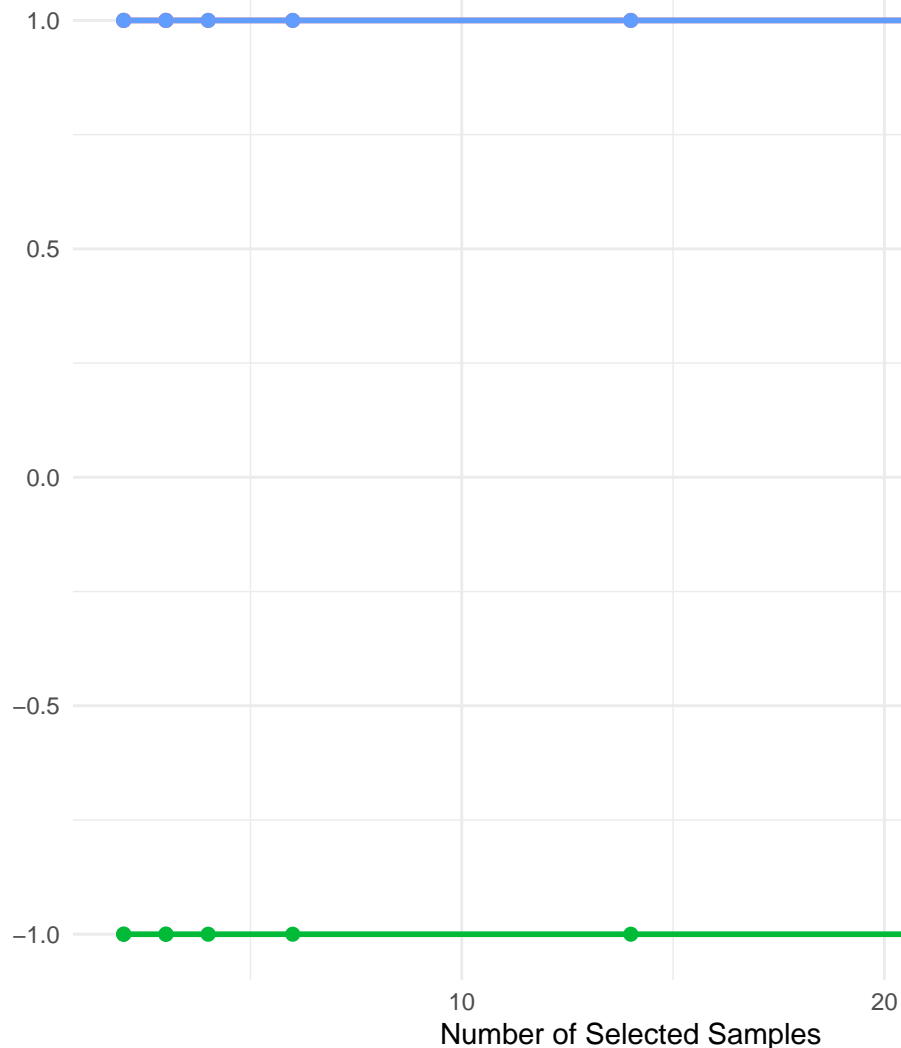


Score: NRW (Local, partial2 , Genotype: wave )

Kendall Correlation

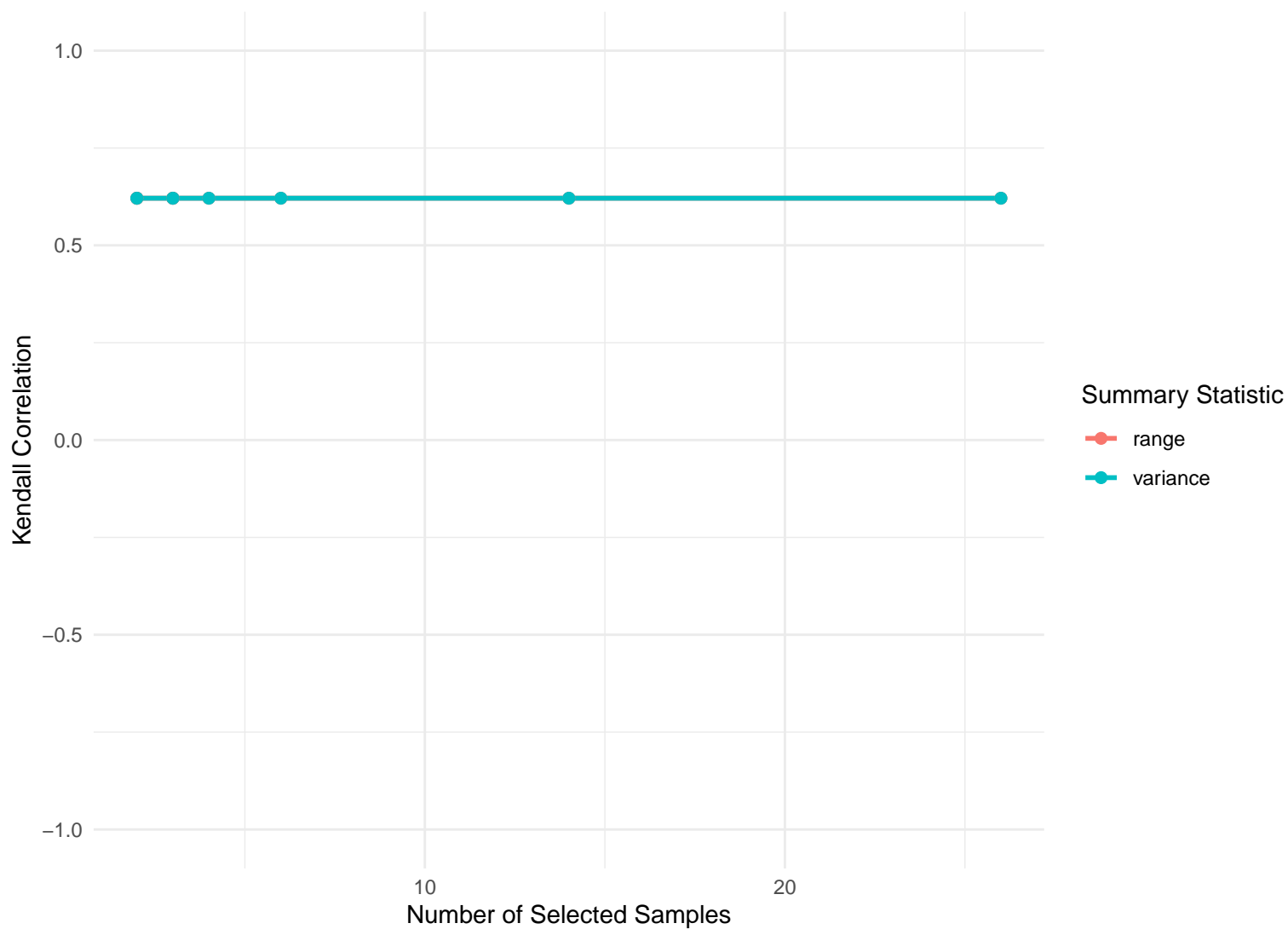
Summary Statistic

- range
- slope
- variance





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

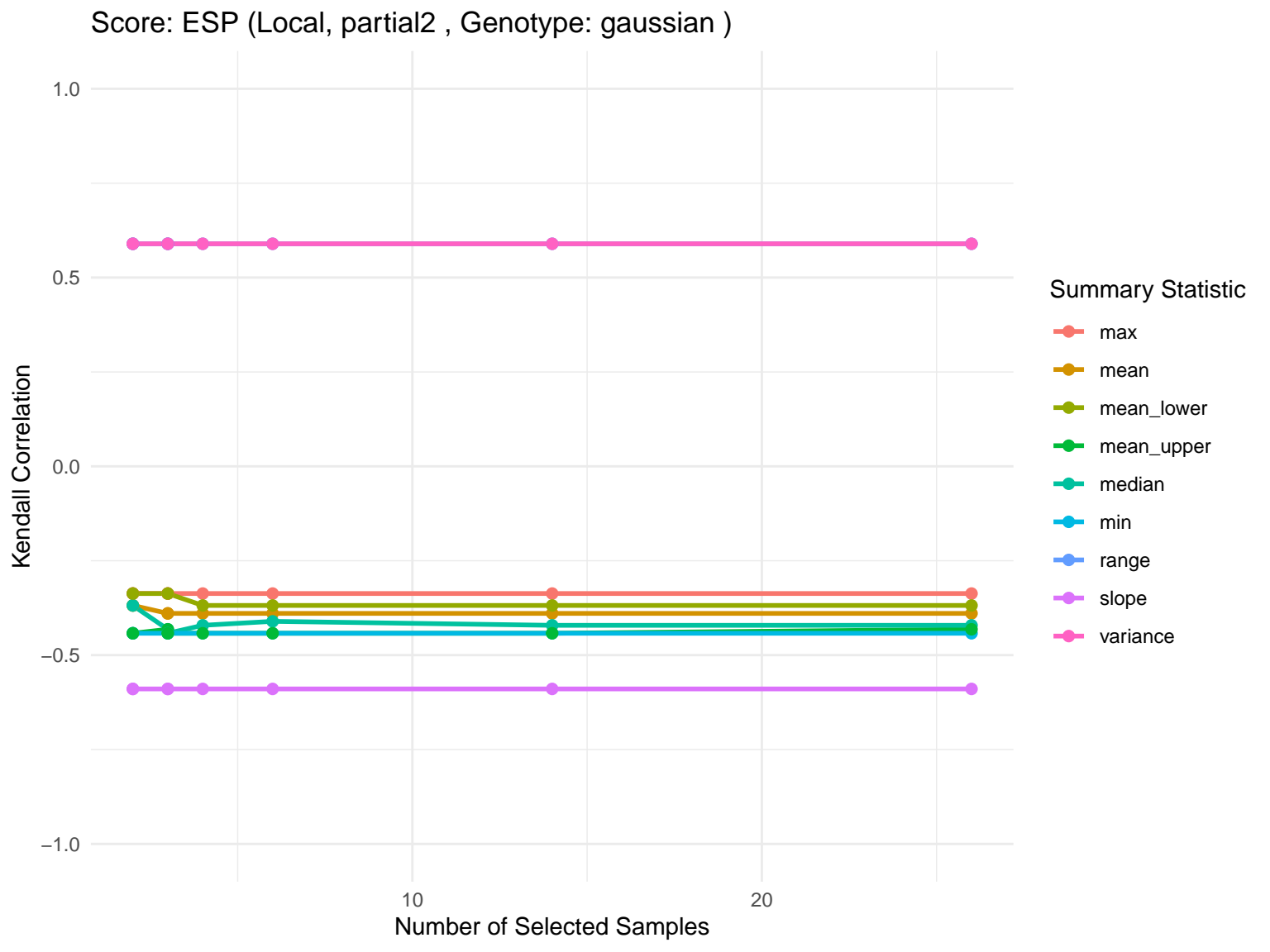
-0.5

-1.0

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Number of Selected Samples

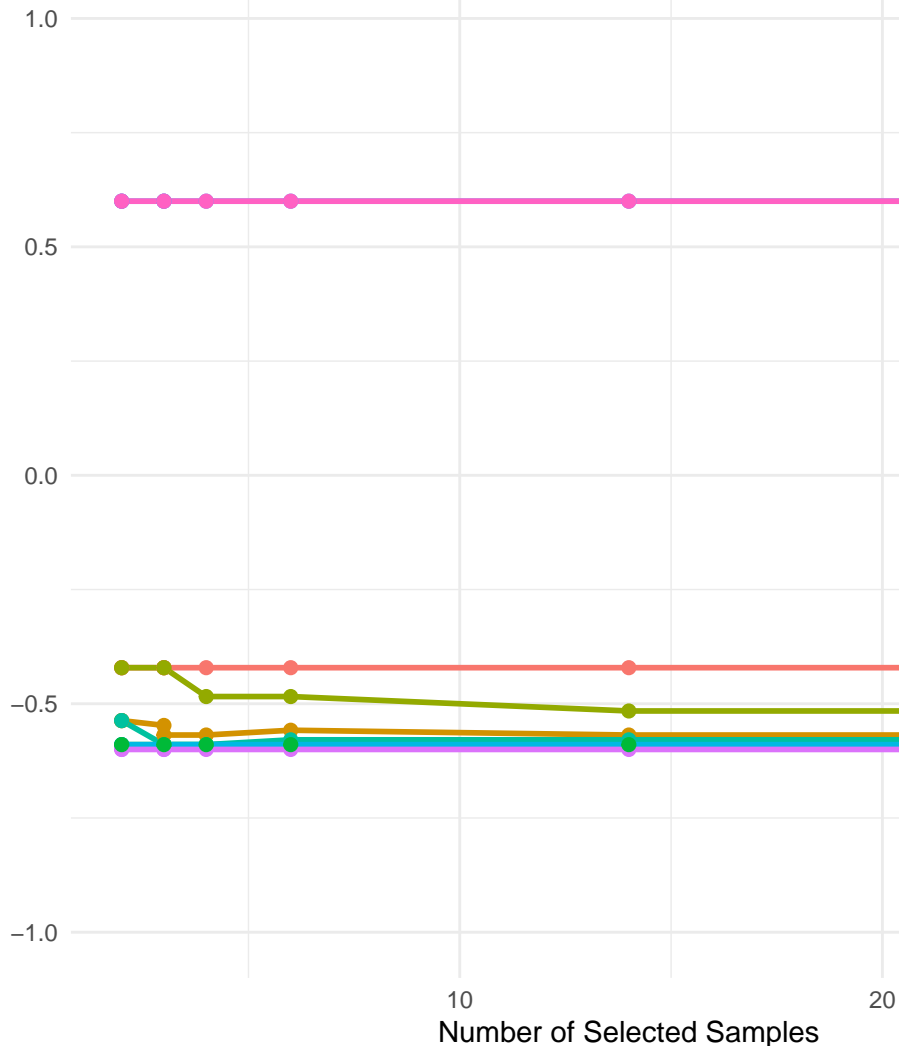


Score: ESP (Local, partial2 , Genotype: sinusoidal )

Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance



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

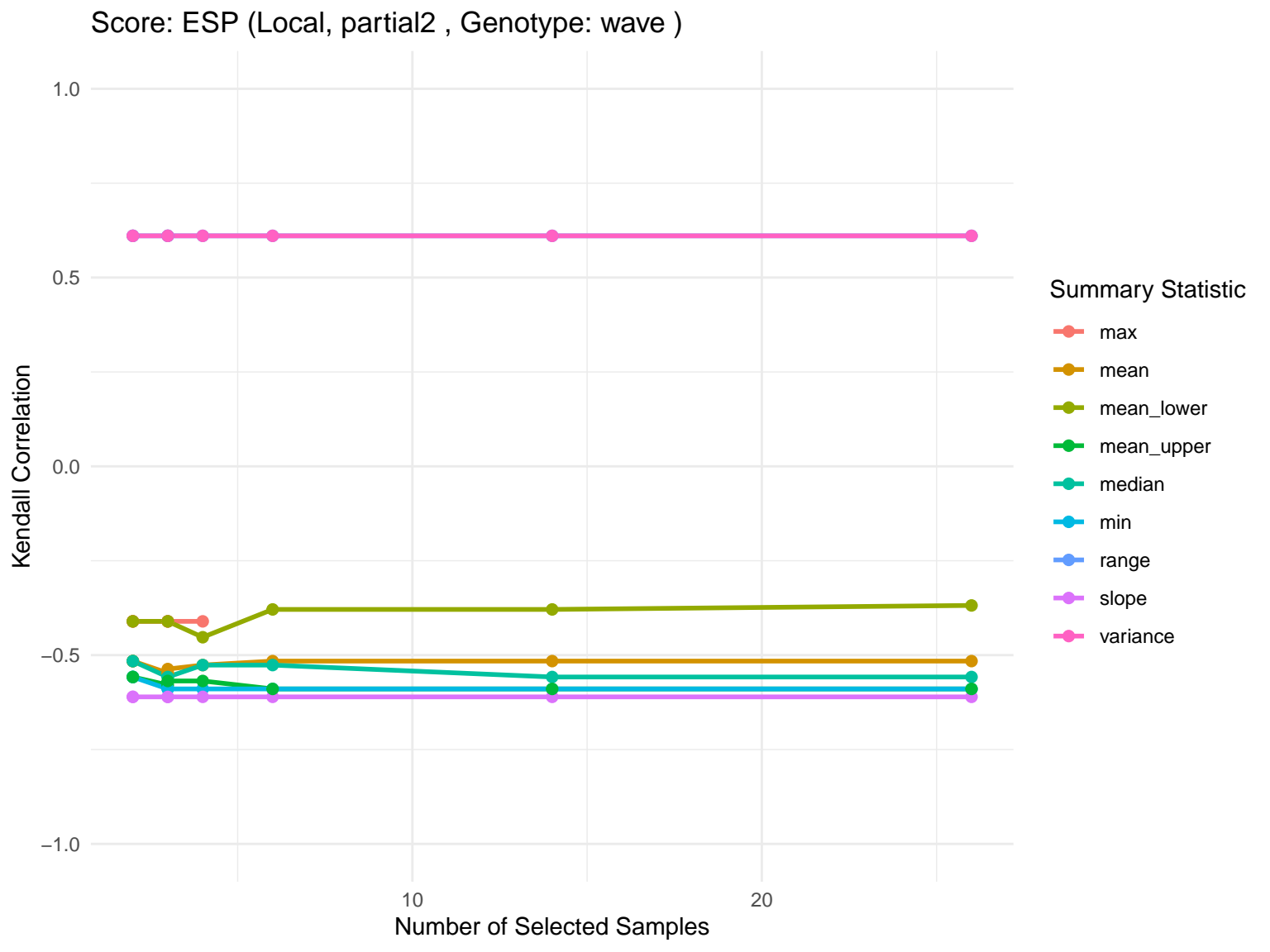
Kendall Correlation

Summary Statistic

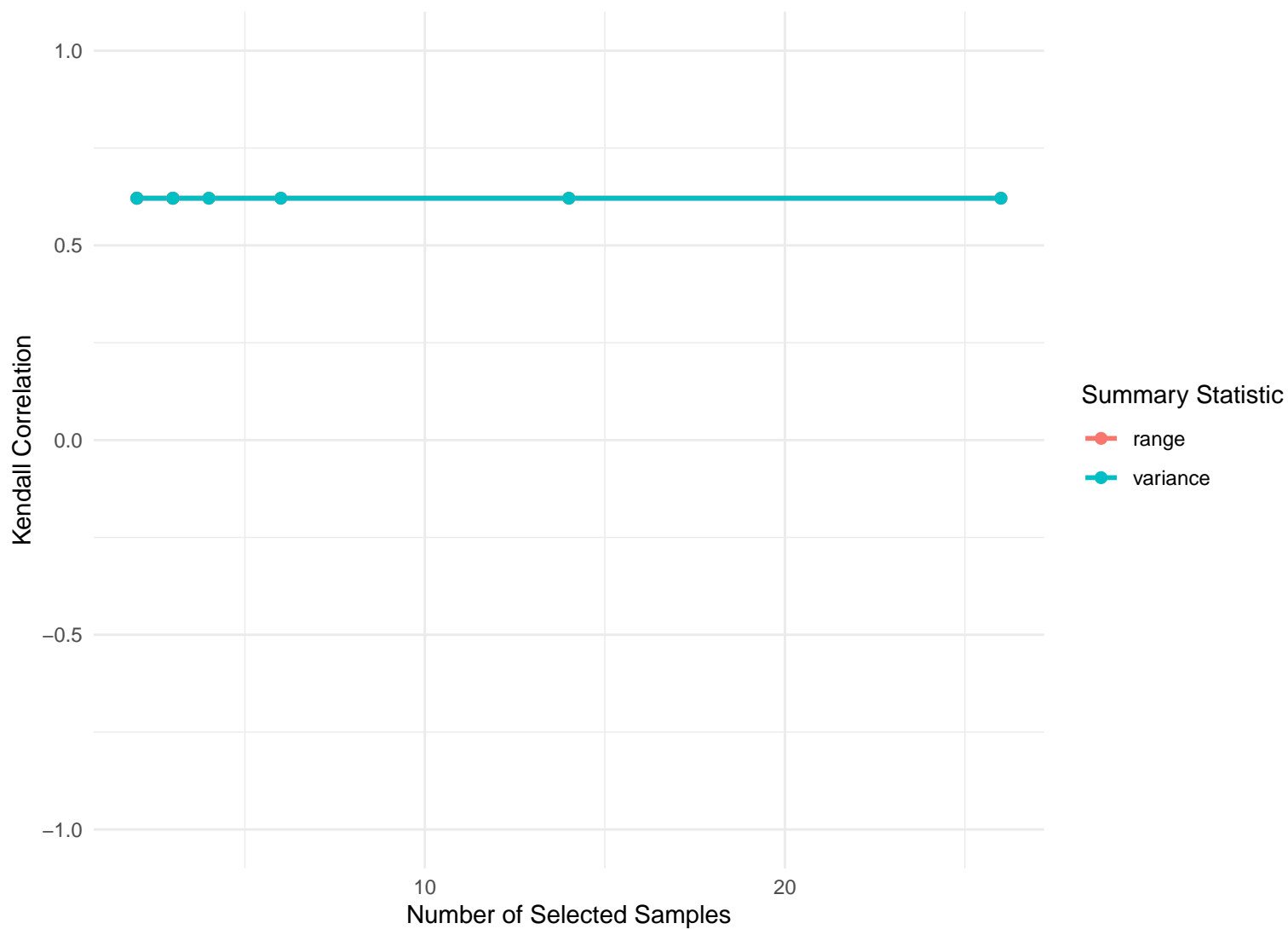
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

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

10 20  
Number of Selected Samples



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

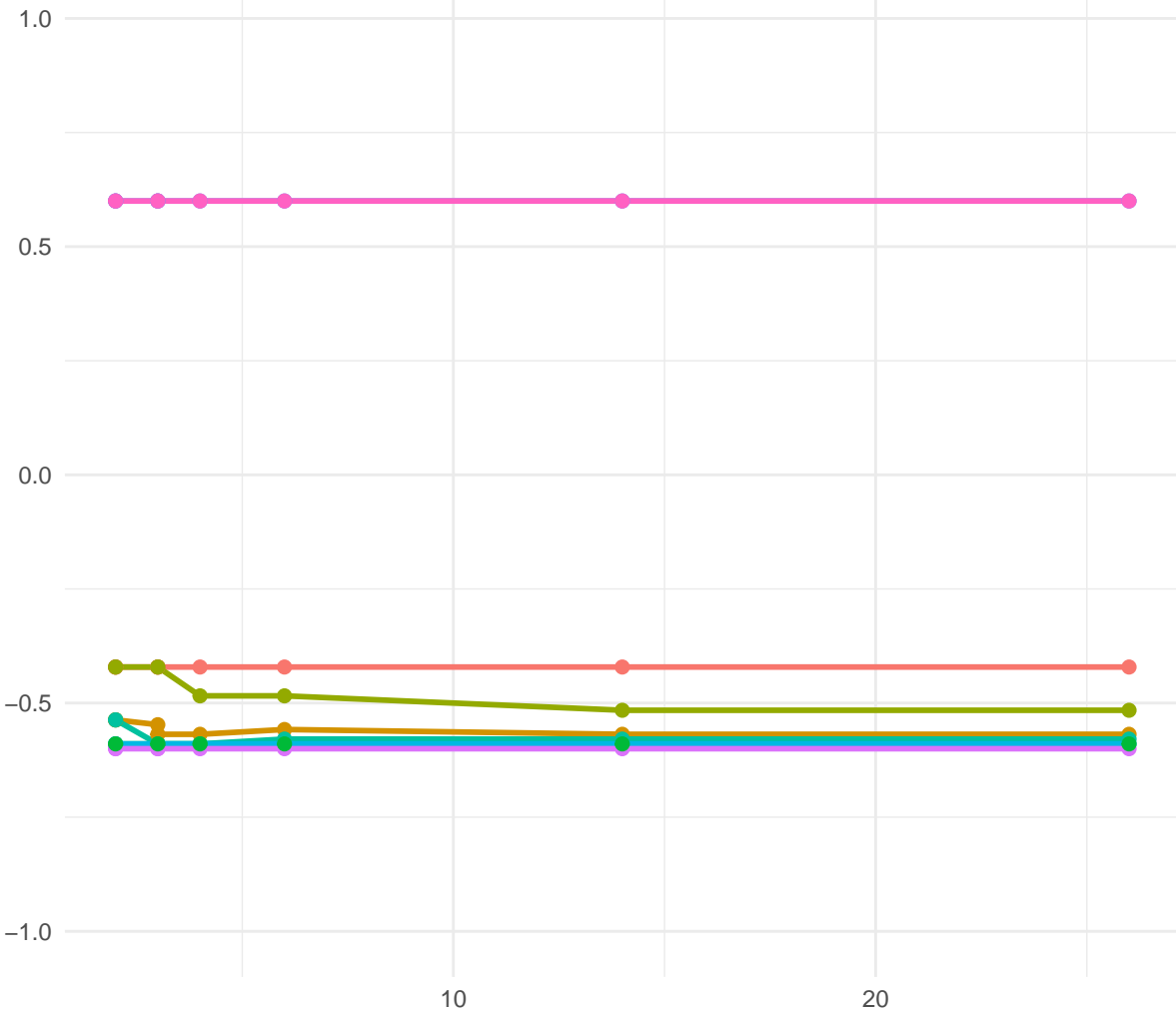
Kendall Correlation

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

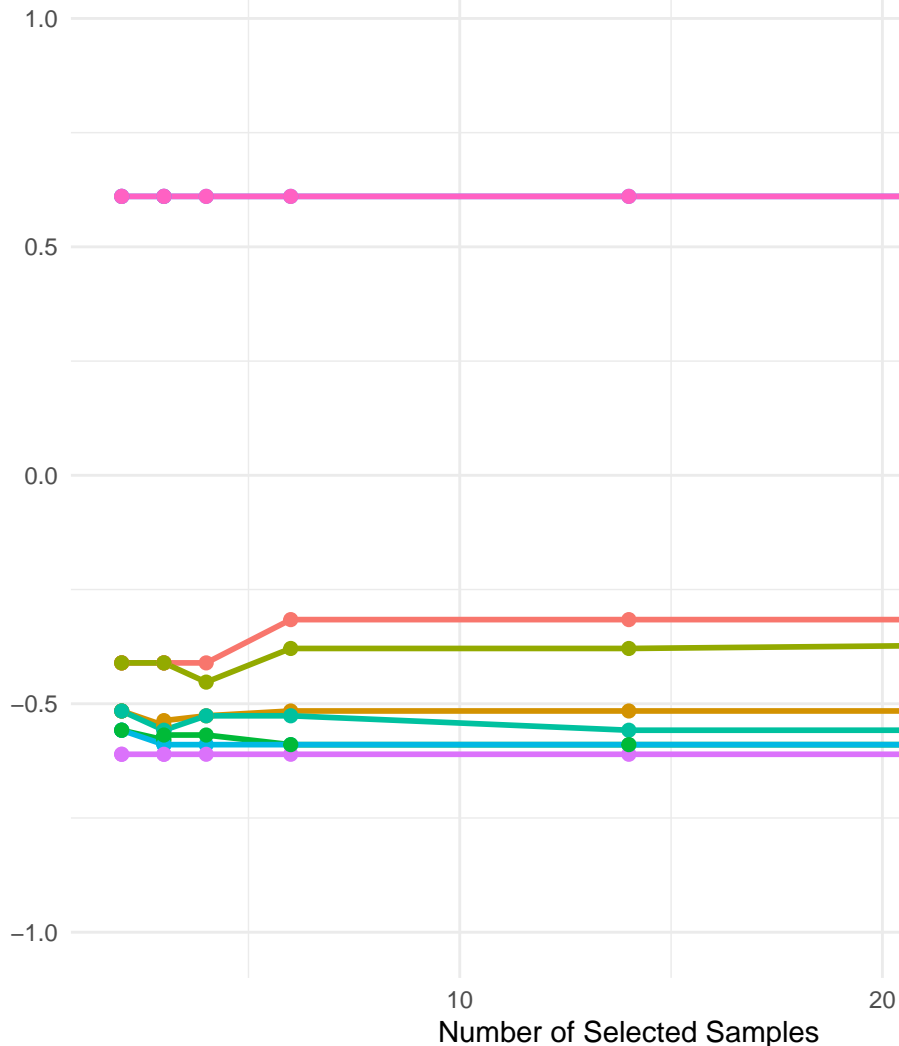


Score: CEV (Local, partial2 , Genotype: wave )

Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance





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

Kendall Correlation

1.0  
0.5  
0.0  
-0.5  
-1.0

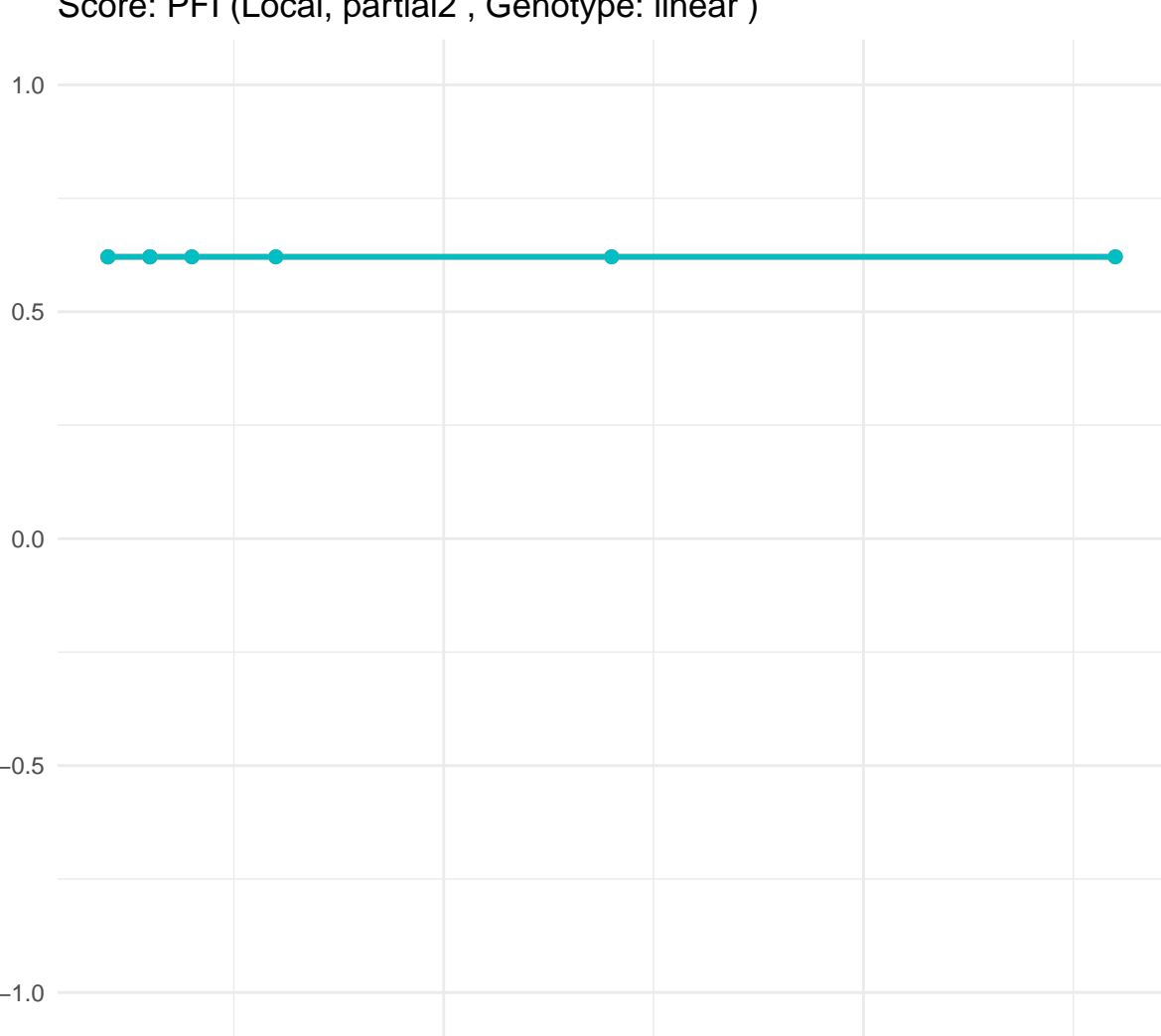
Number of Selected Samples

10

20

Summary Statistic

range  
variance



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

Kendall Correlation

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

1.0

0.5

0.0

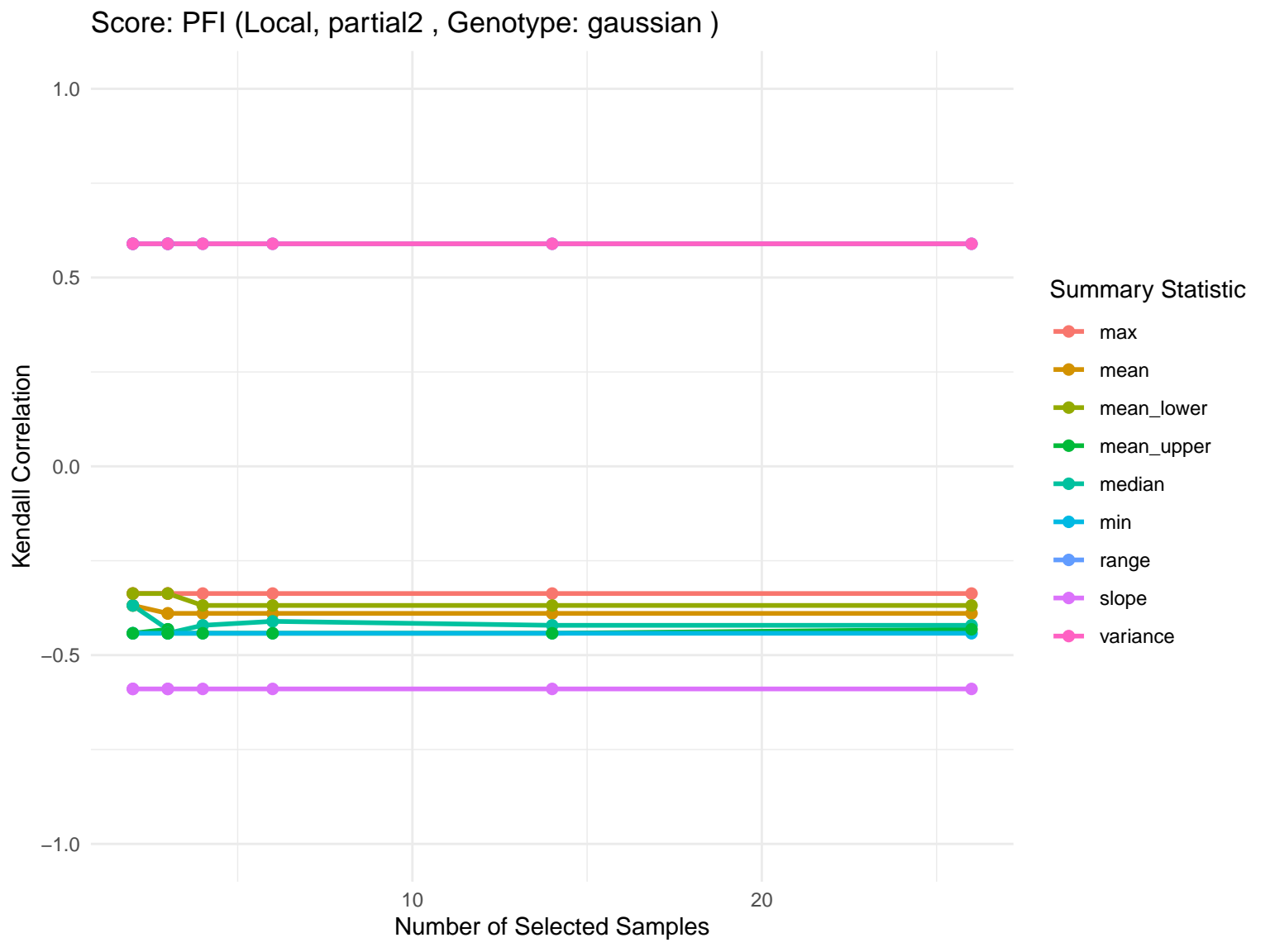
-0.5

-1.0

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Number of Selected Samples

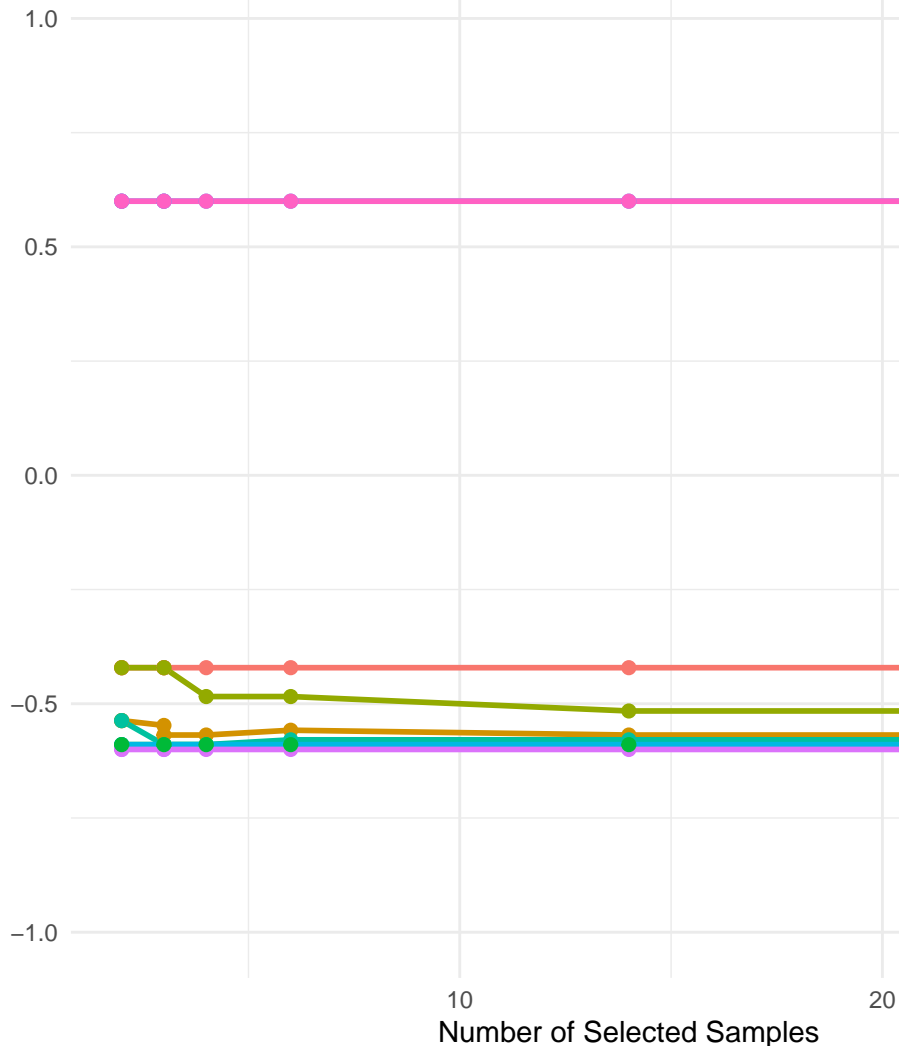


Score: PFI (Local, partial2 , Genotype: sinusoidal )

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

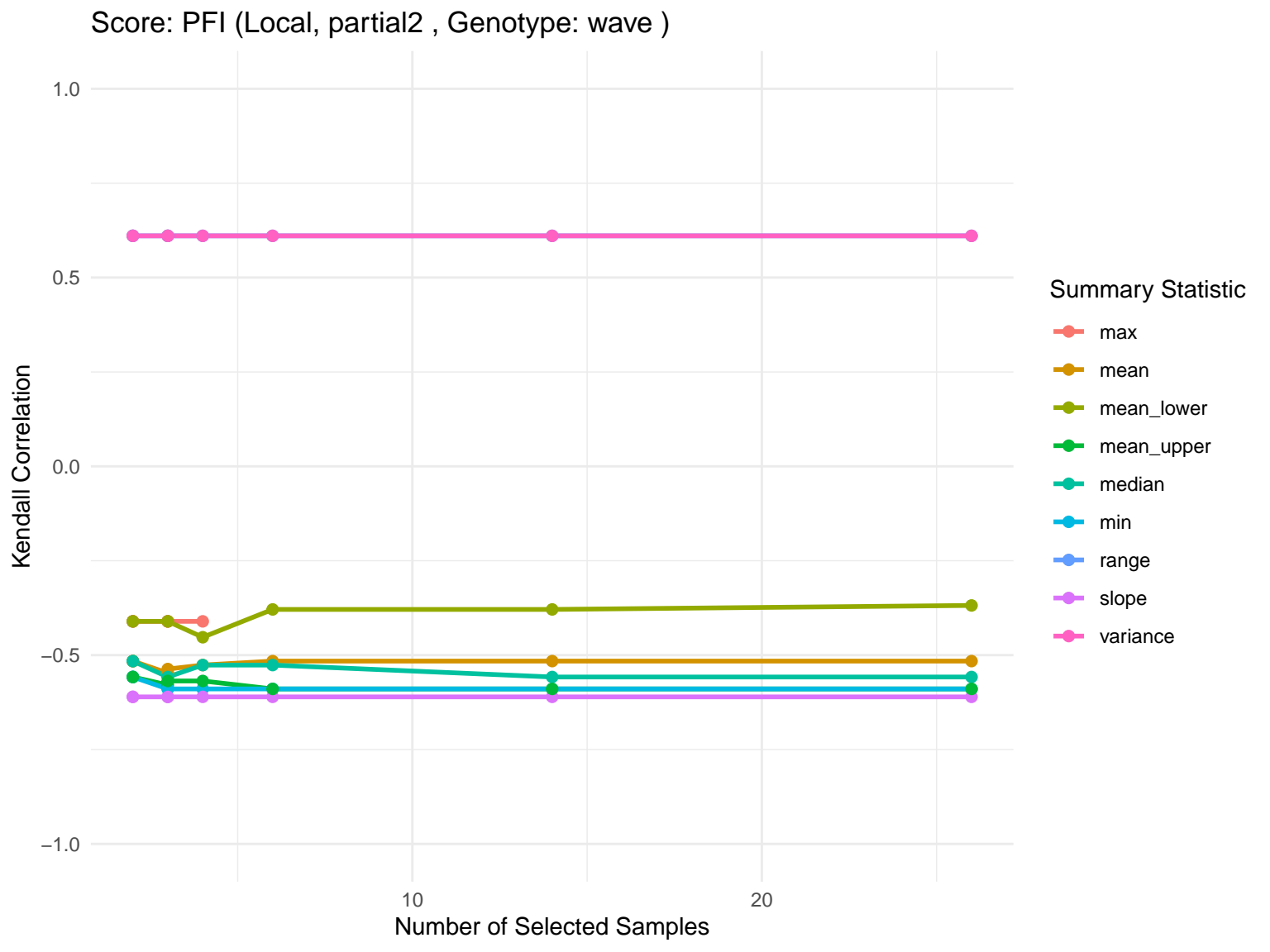
0.5

0.0

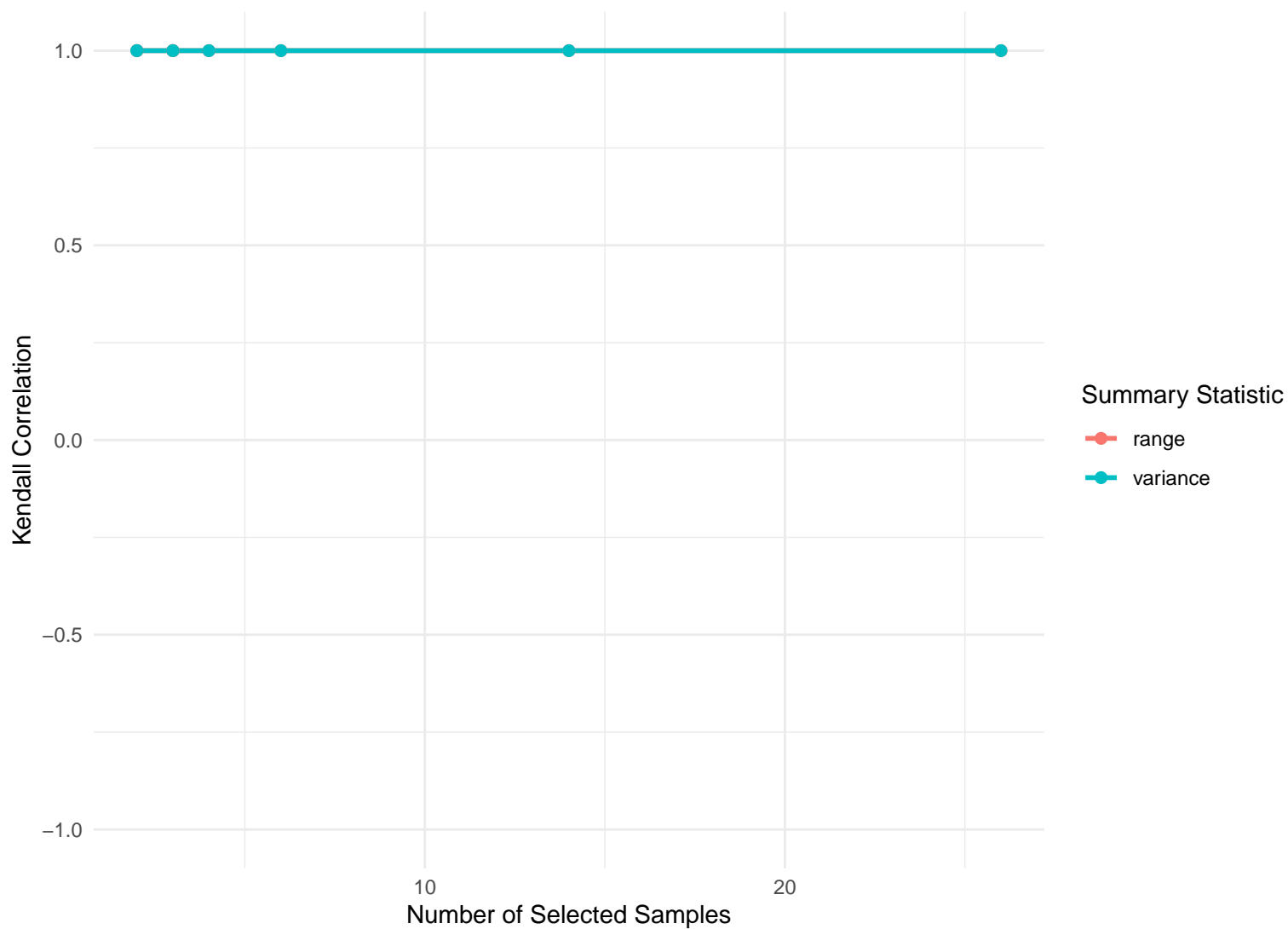
-0.5

-1.0

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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

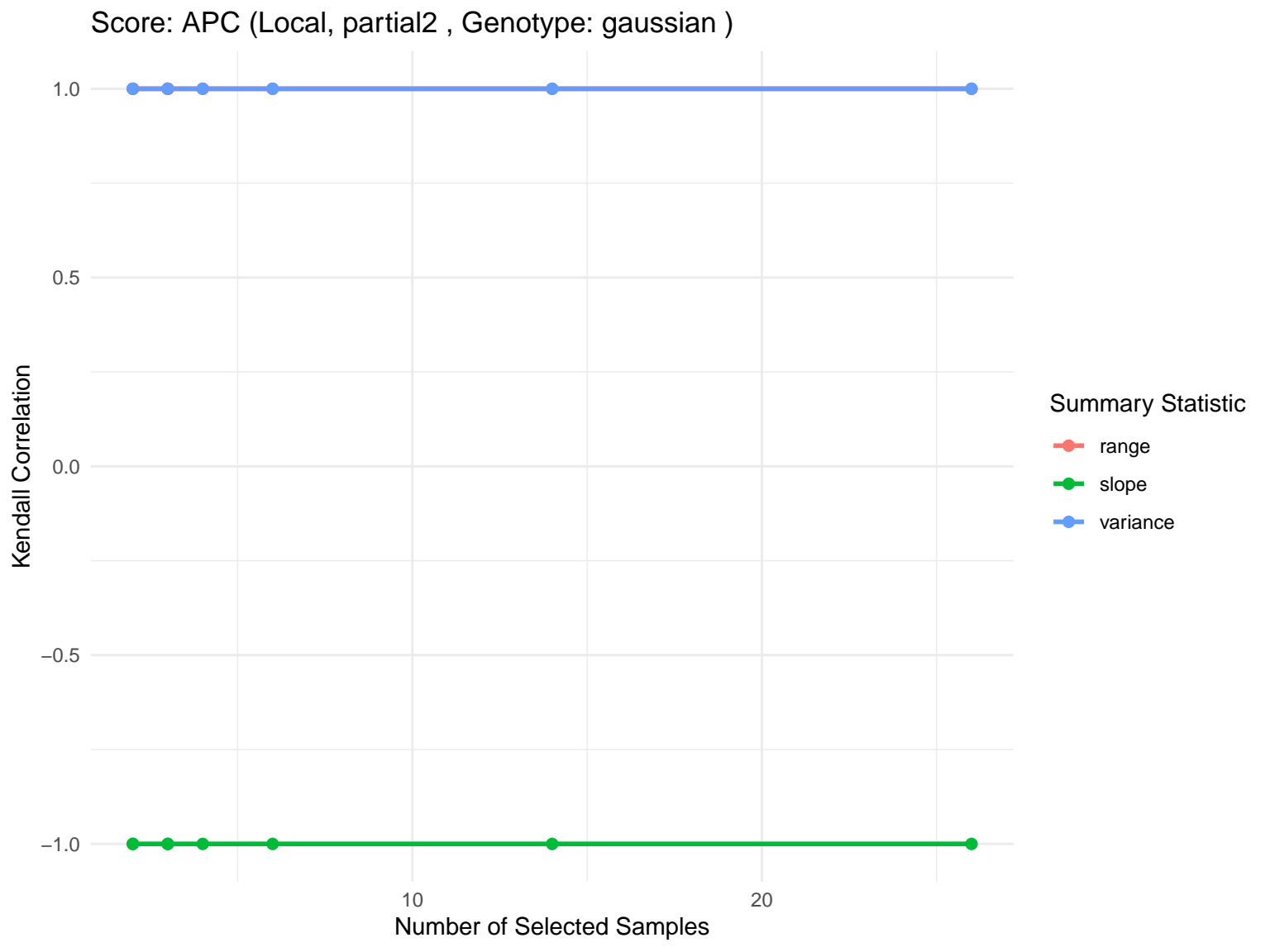
10

20

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

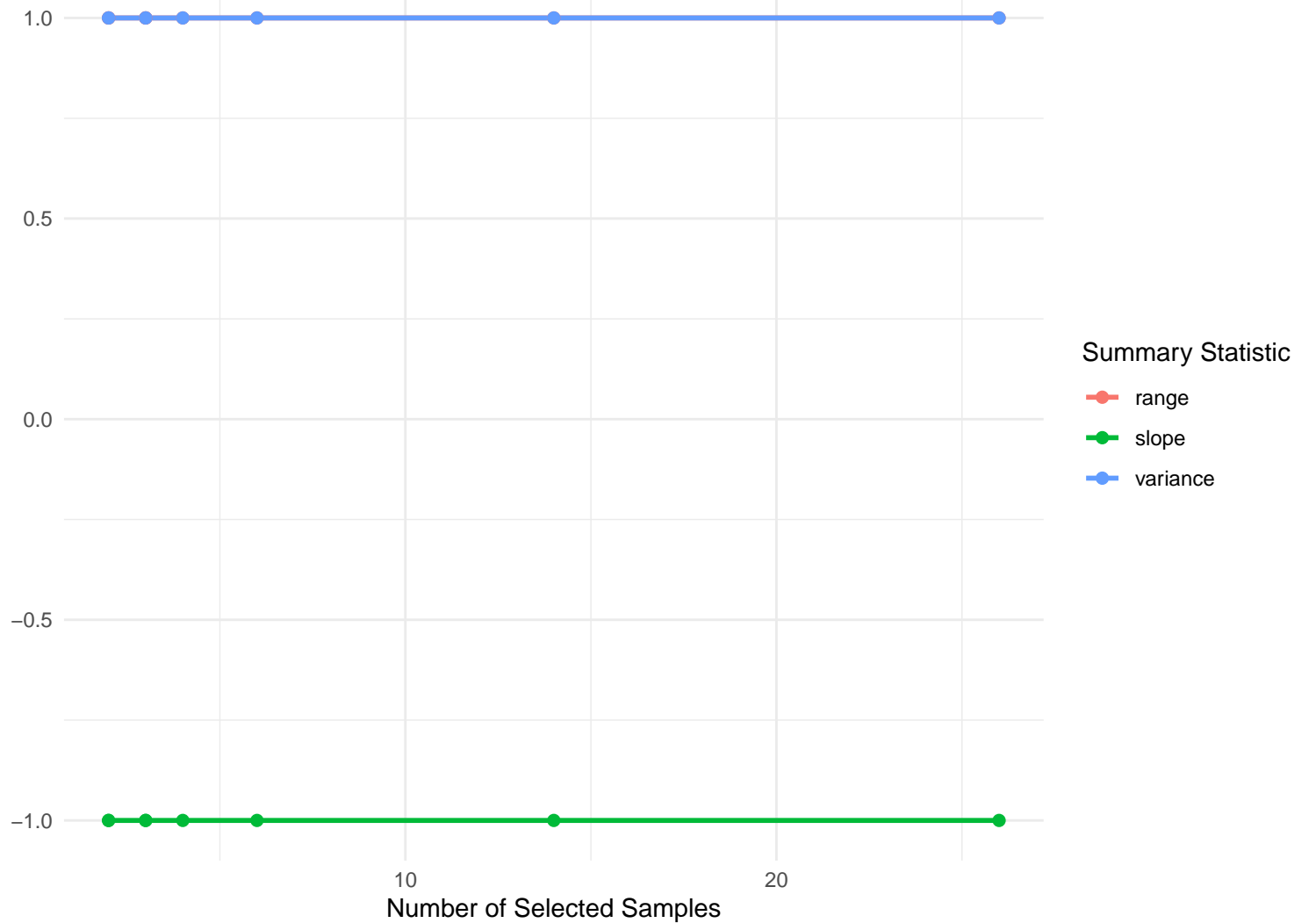
Number of Selected Samples

10

20

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

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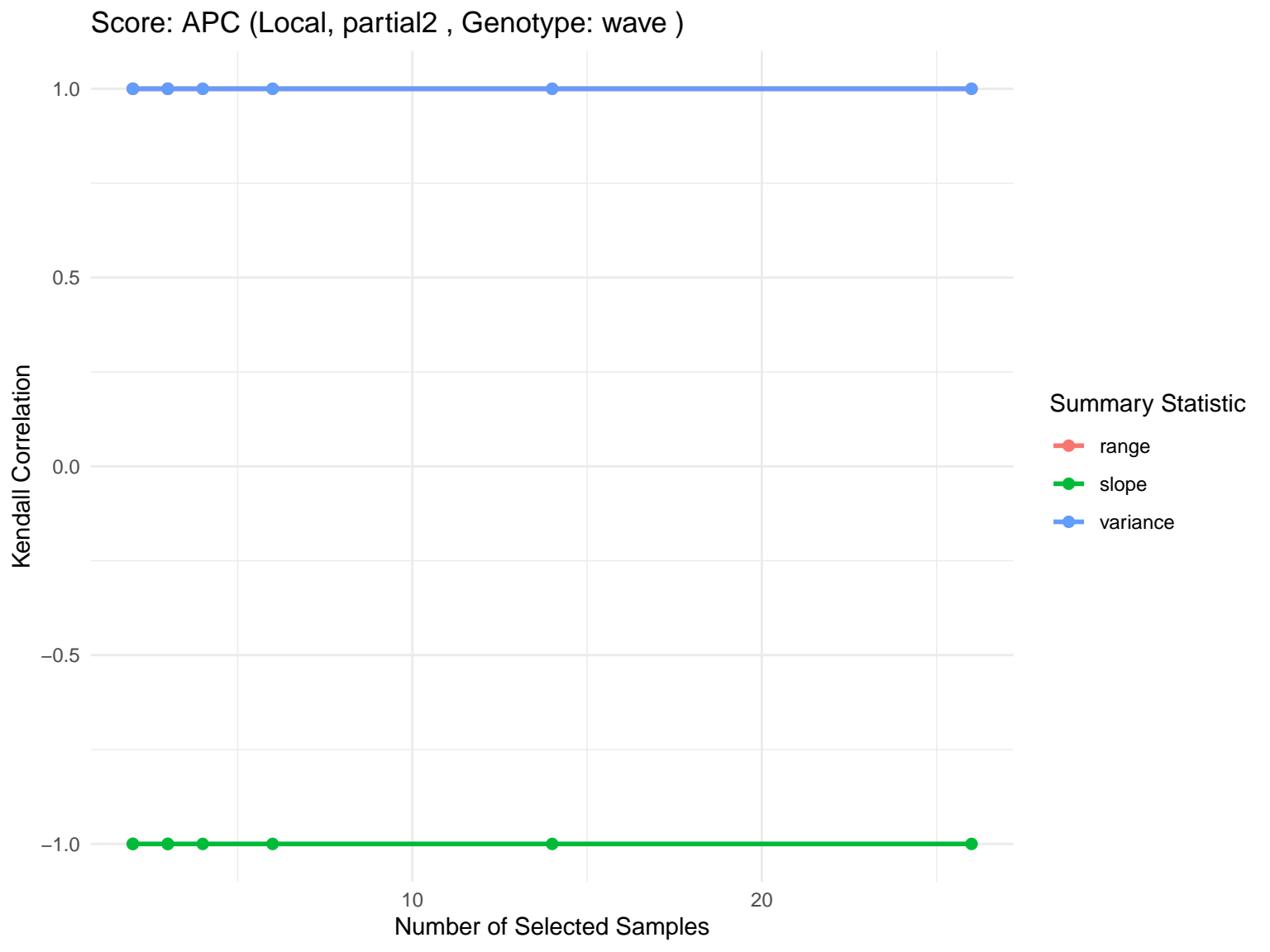
Number of Selected Samples

Summary Statistic

range

slope

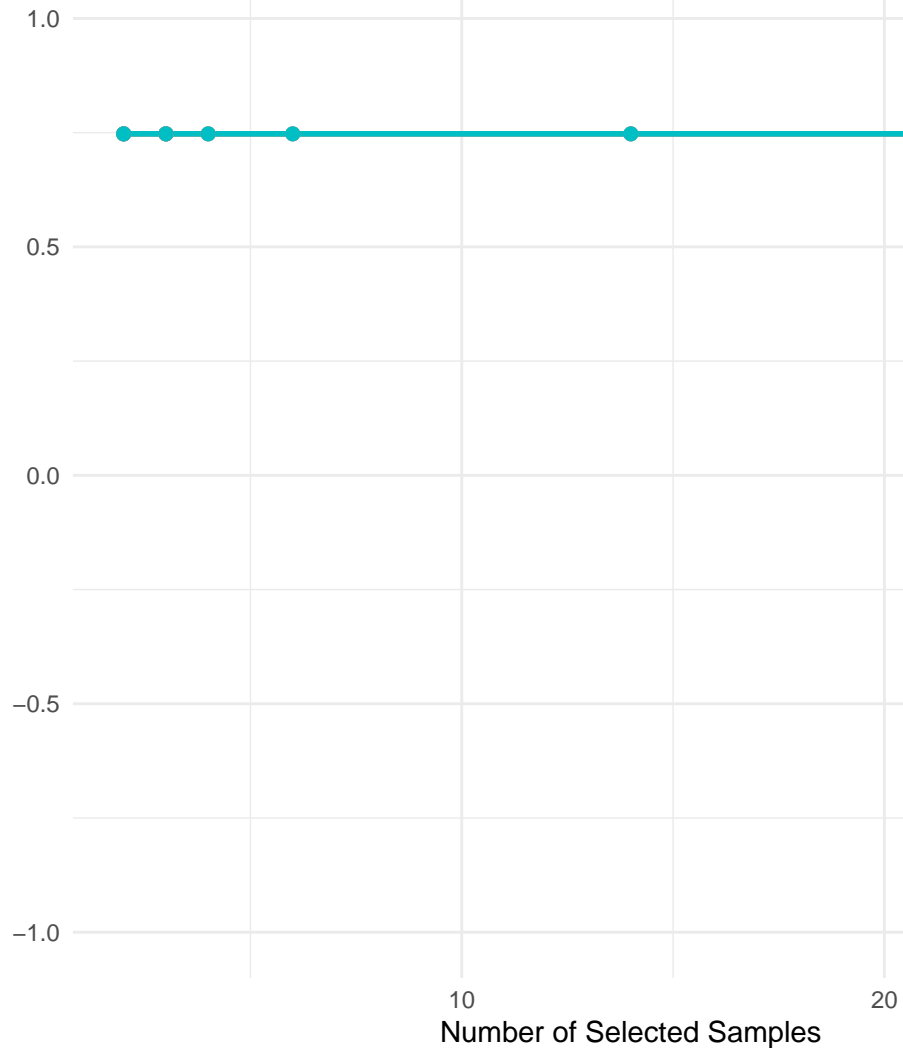
variance





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

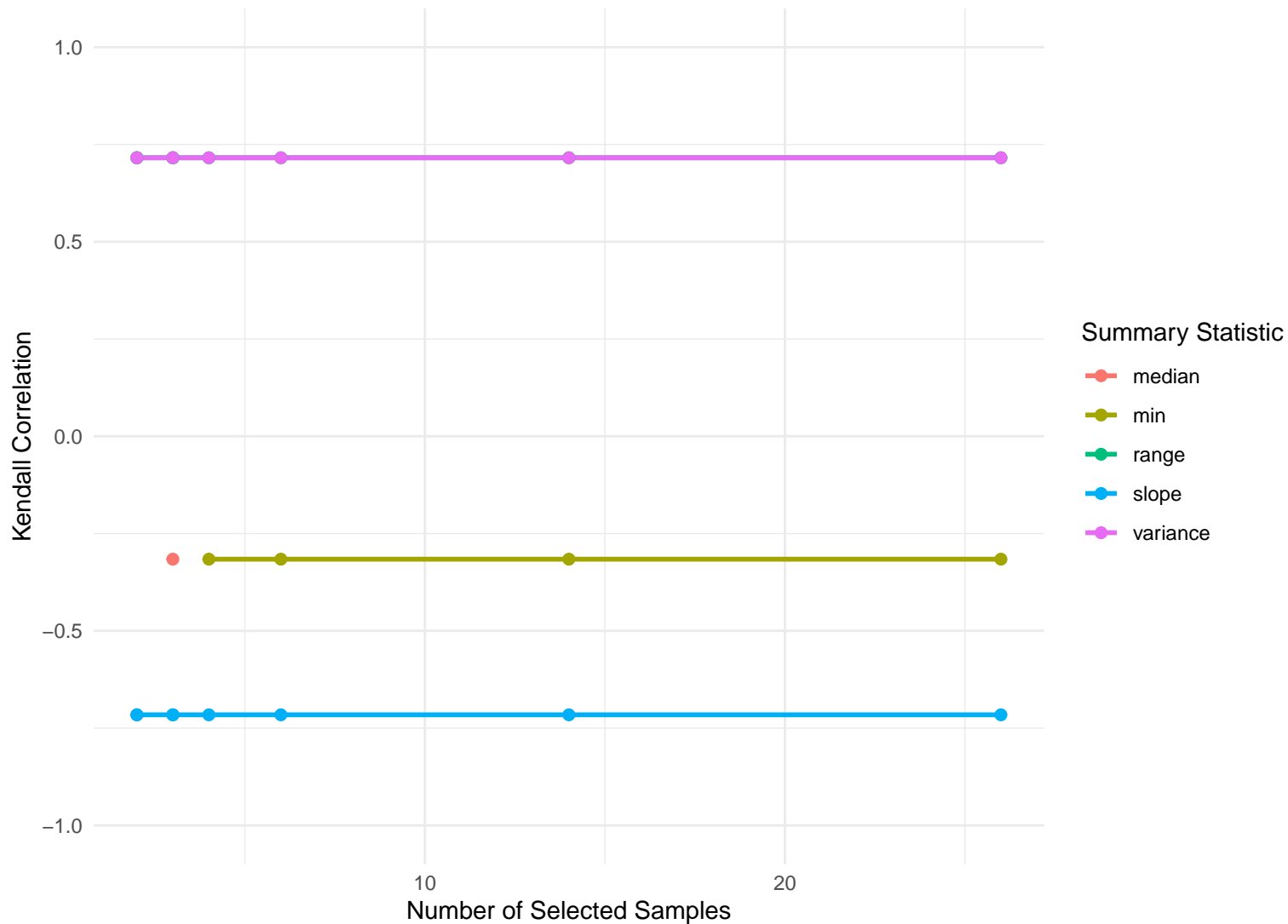
Kendall Correlation



Summary Statistic

- range
- variance

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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

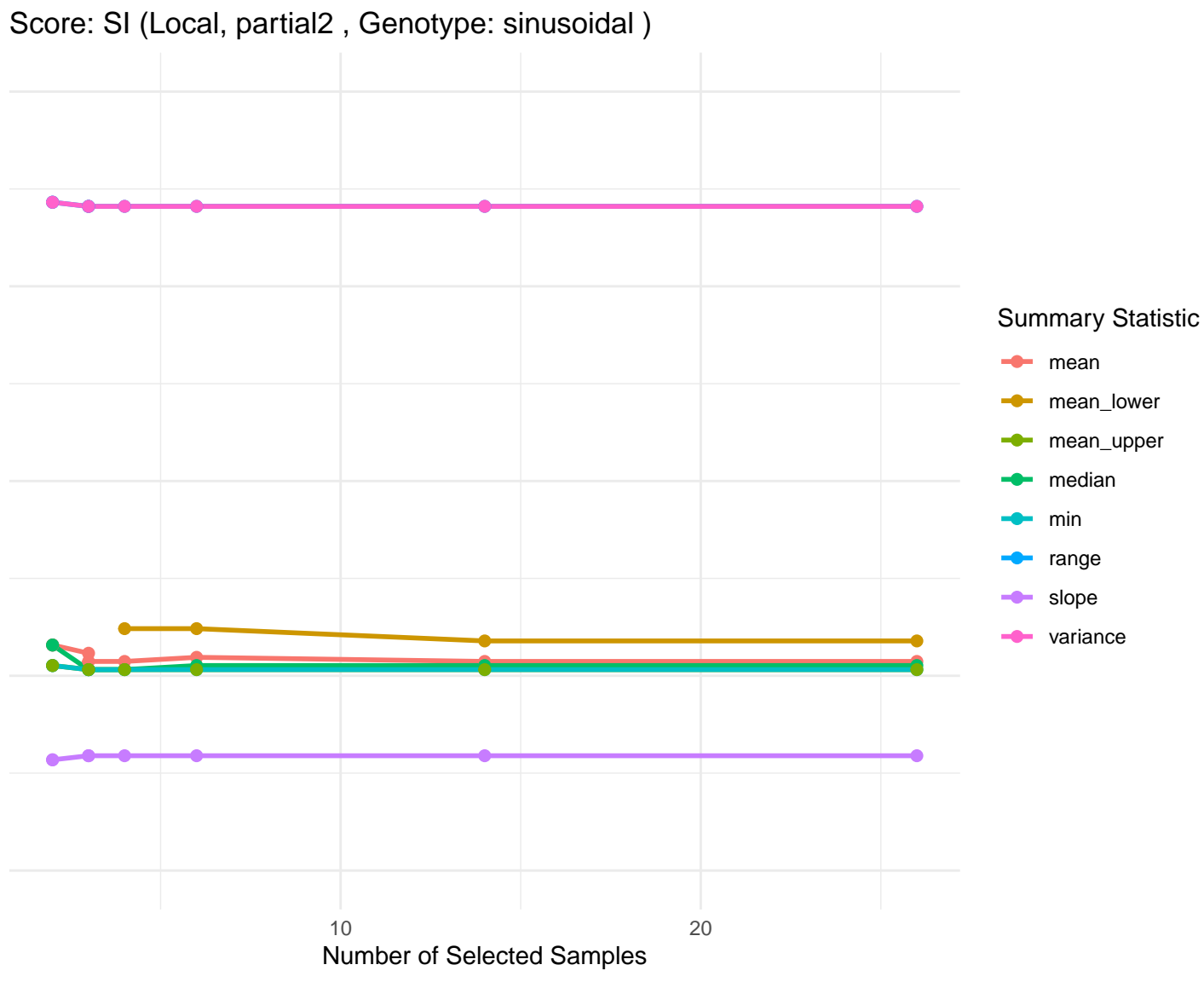
Number of Selected Samples

10

20

Summary Statistic

- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

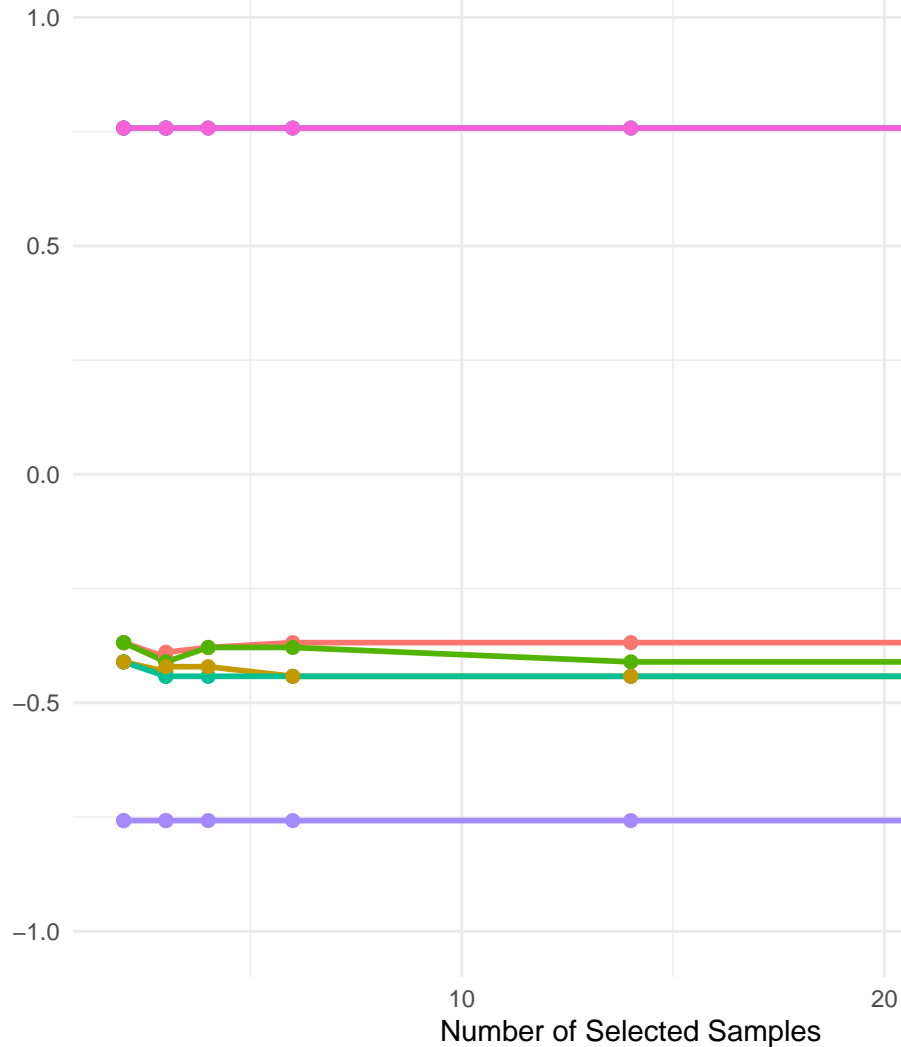


Score: SI (Local, partial2 , Genotype: wave )

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

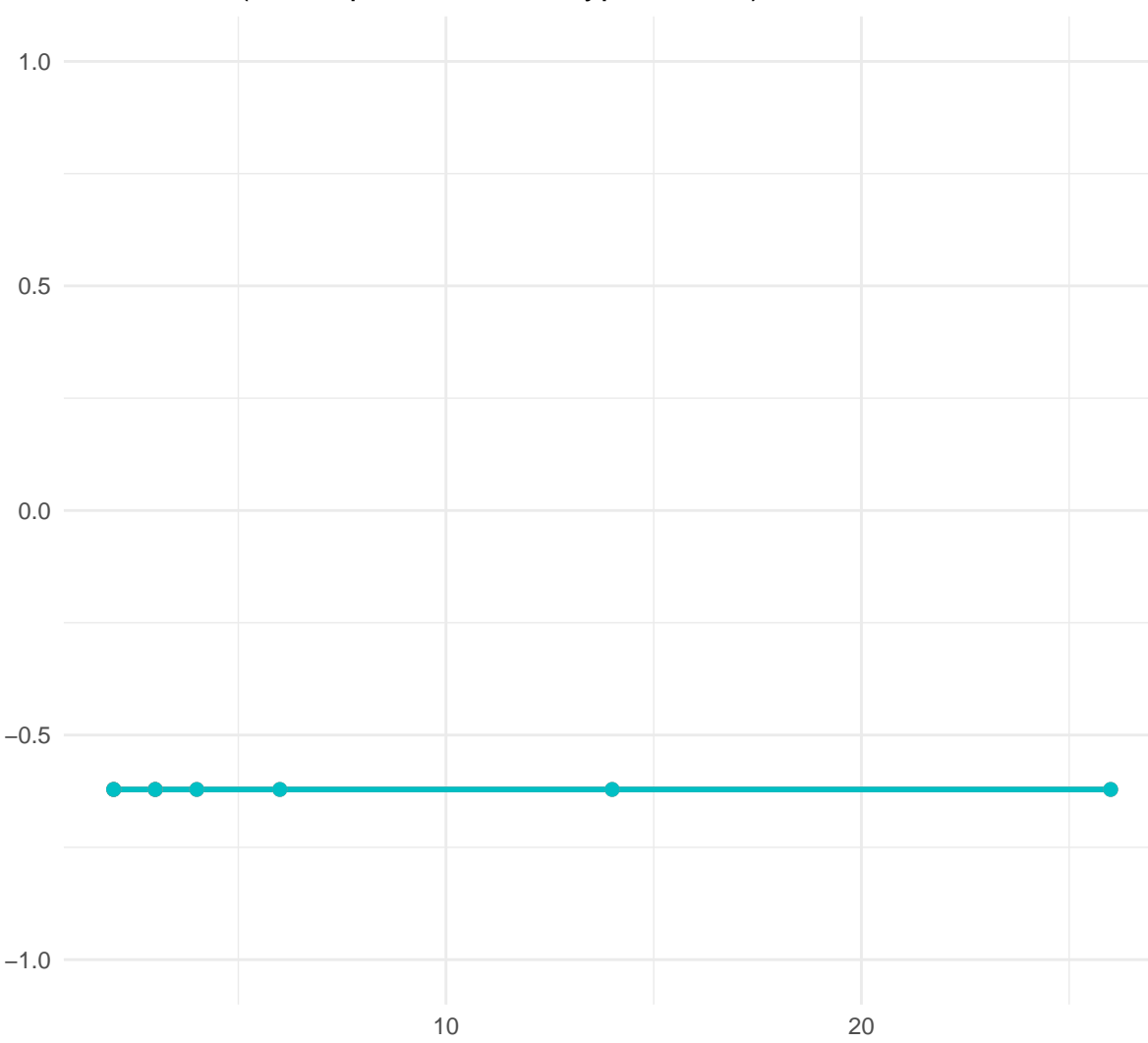
Summary Statistic

range  
variance

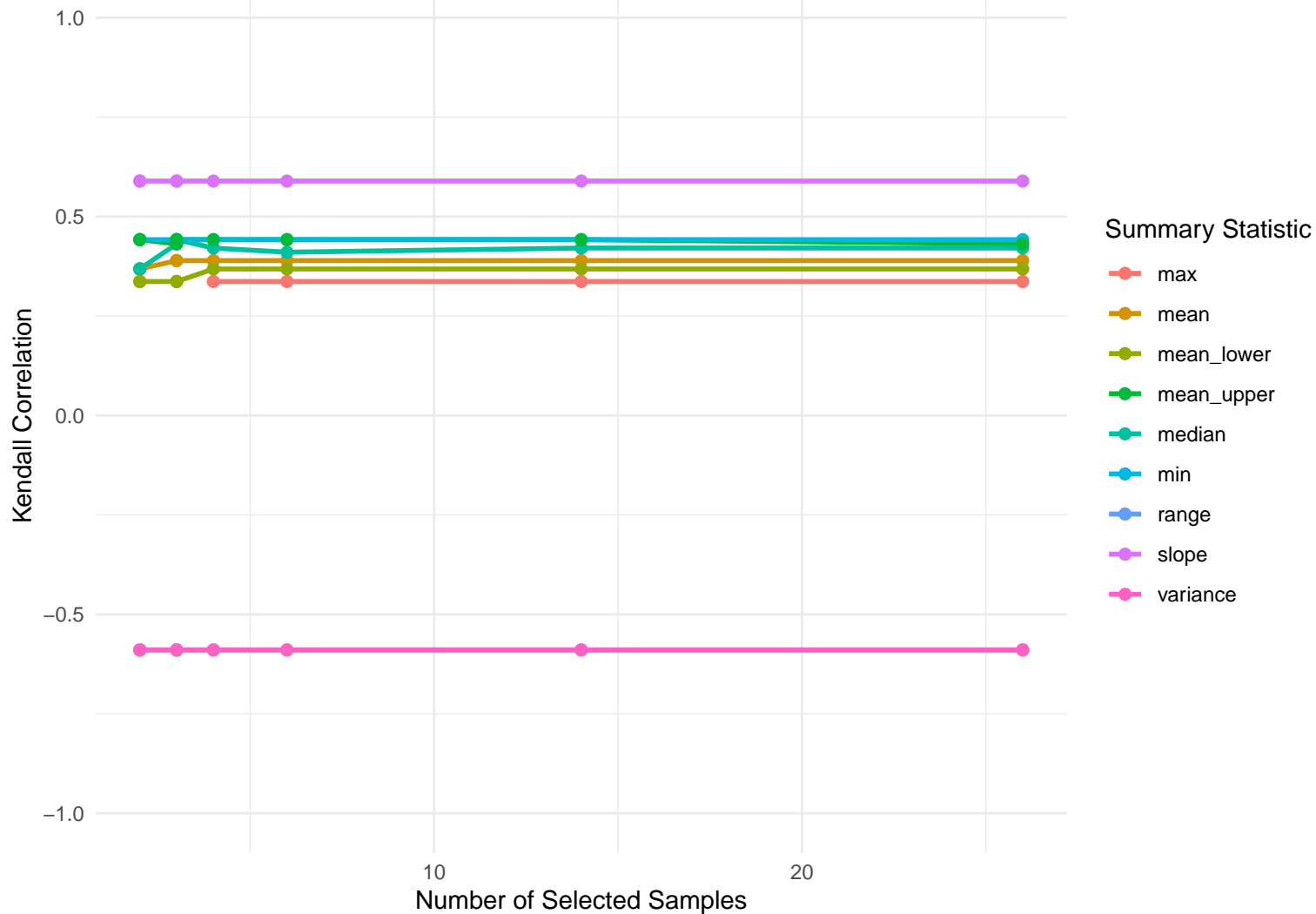
Number of Selected Samples

10

20



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

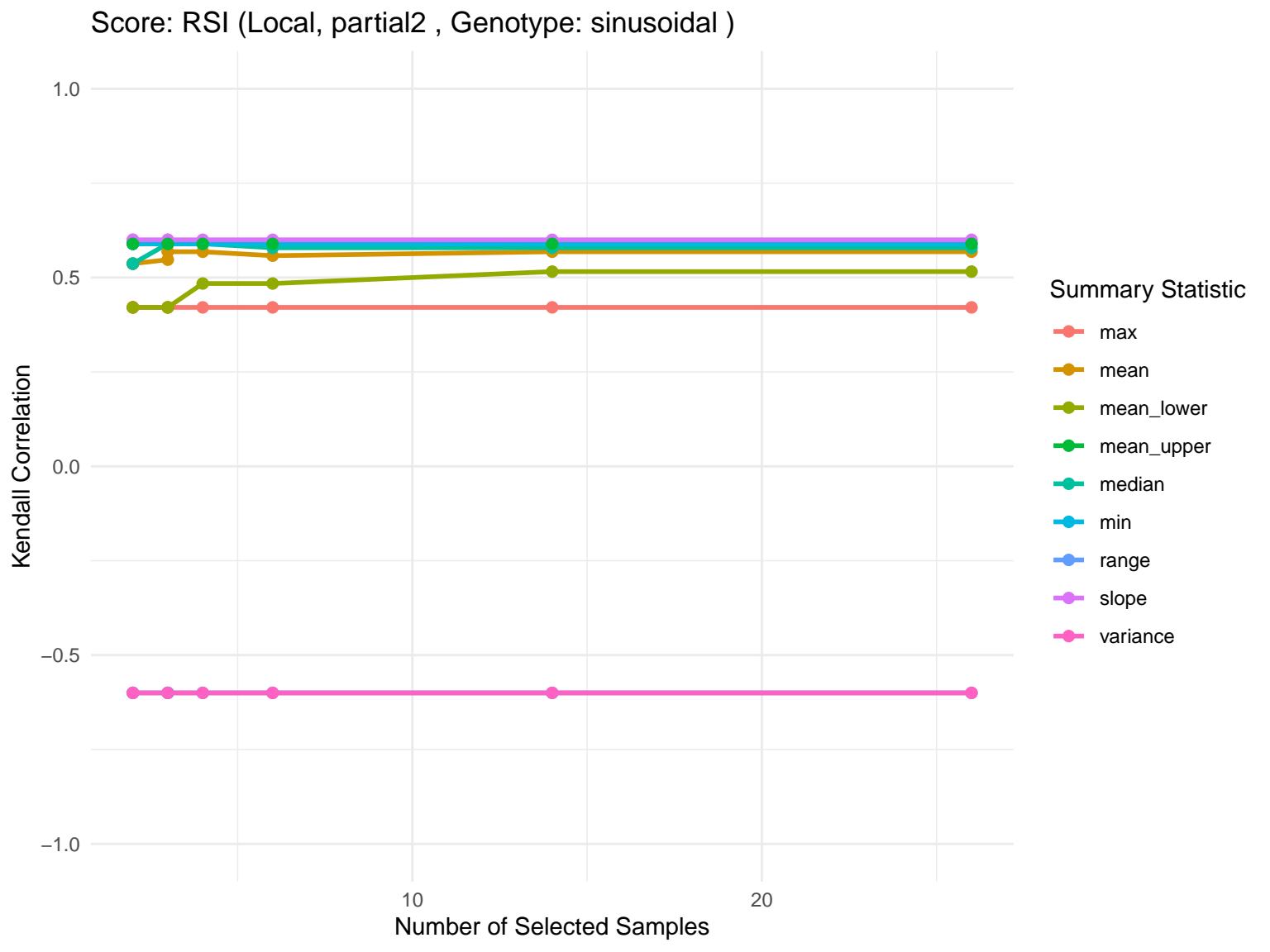
10

20

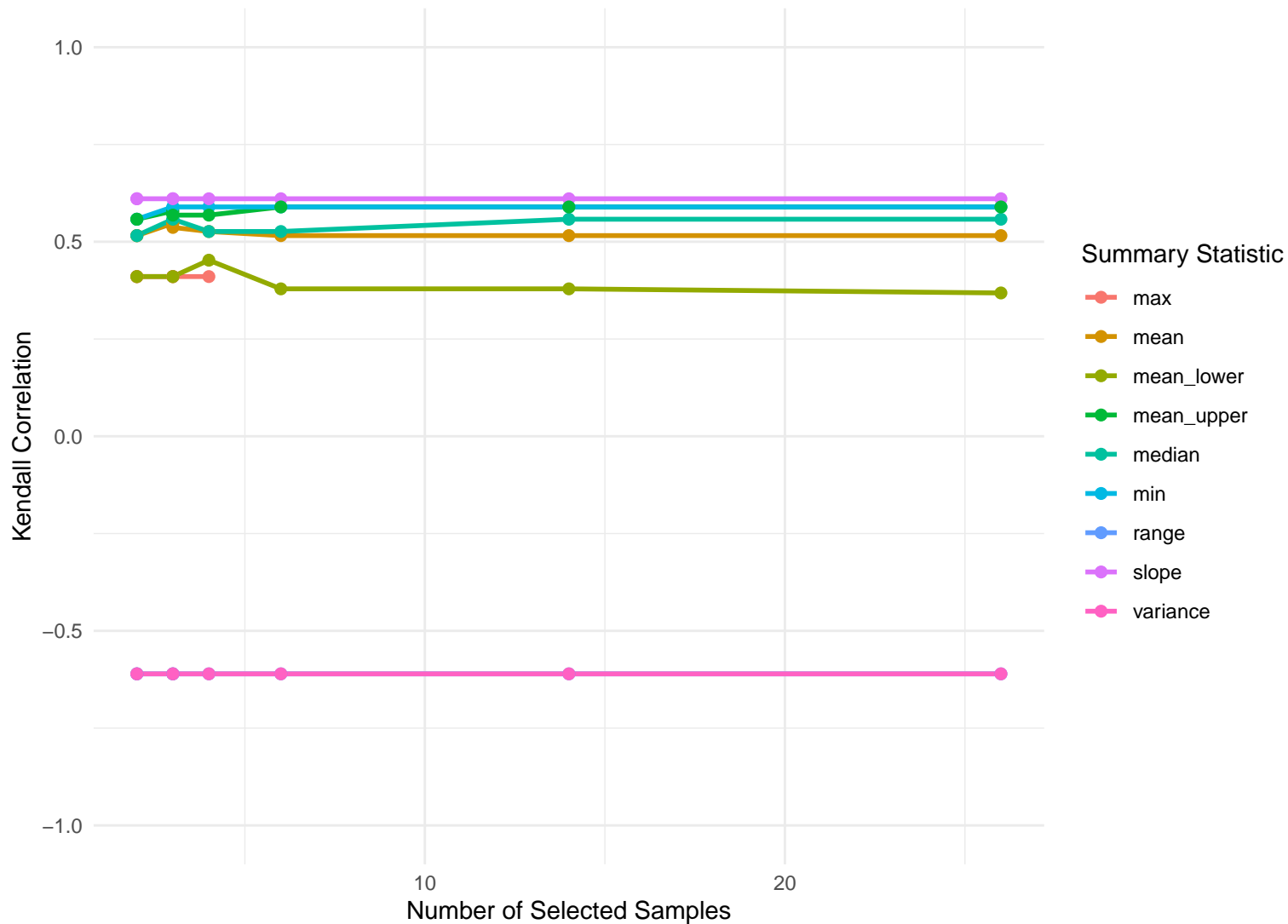
Number of Selected Samples

Summary Statistic

- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance



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





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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

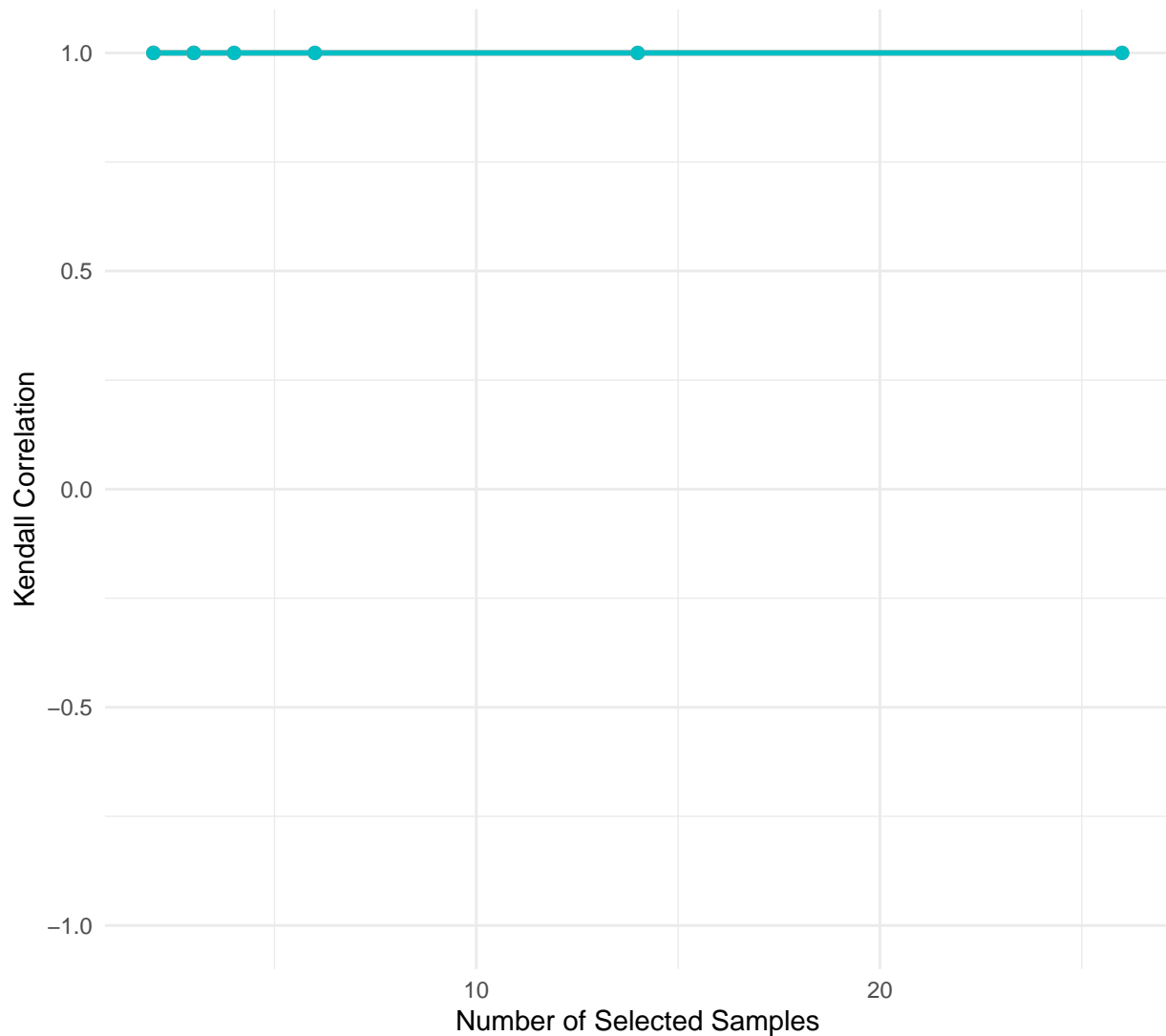
10

20

Number of Selected Samples

Summary Statistic

range  
variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

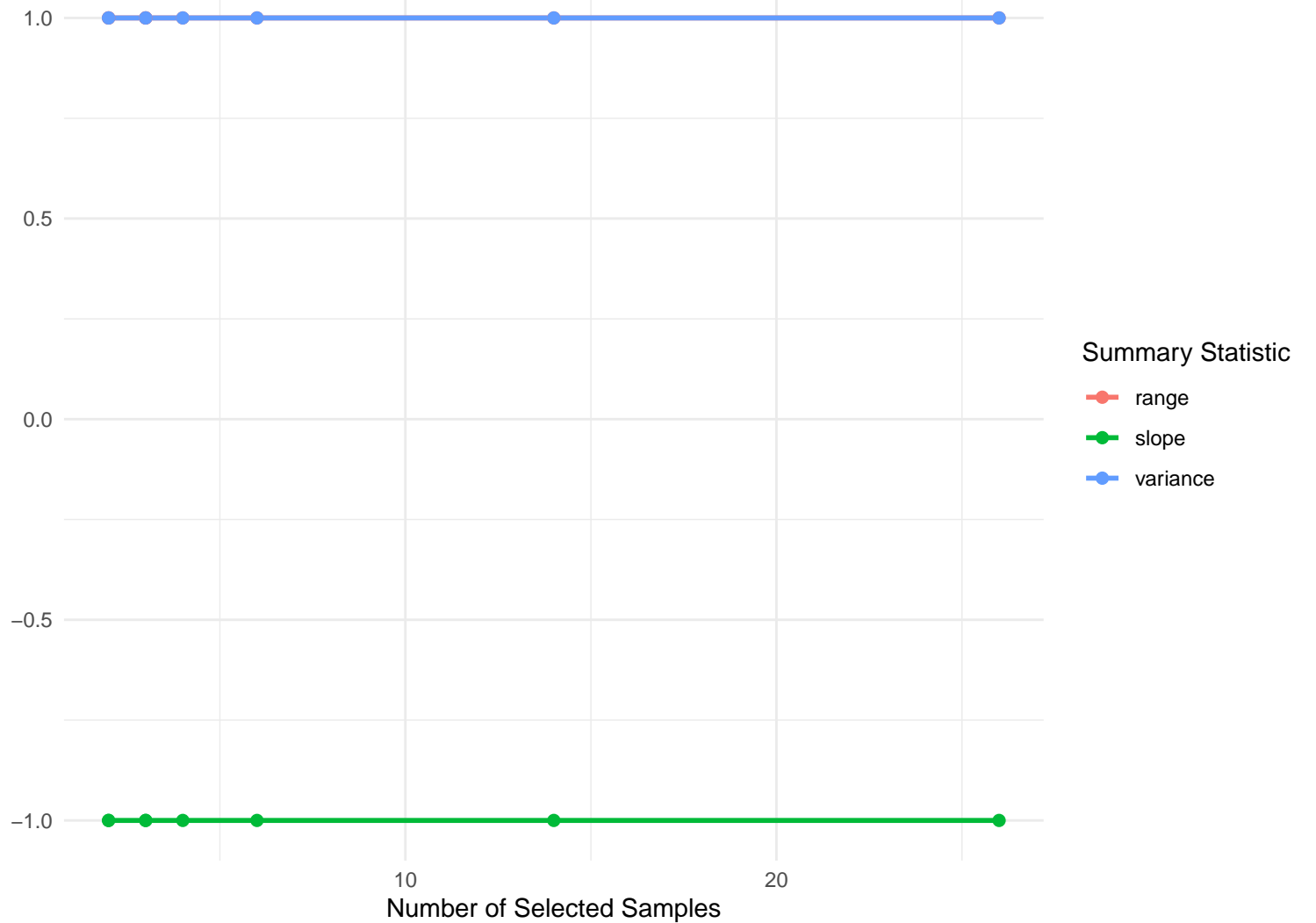
10

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

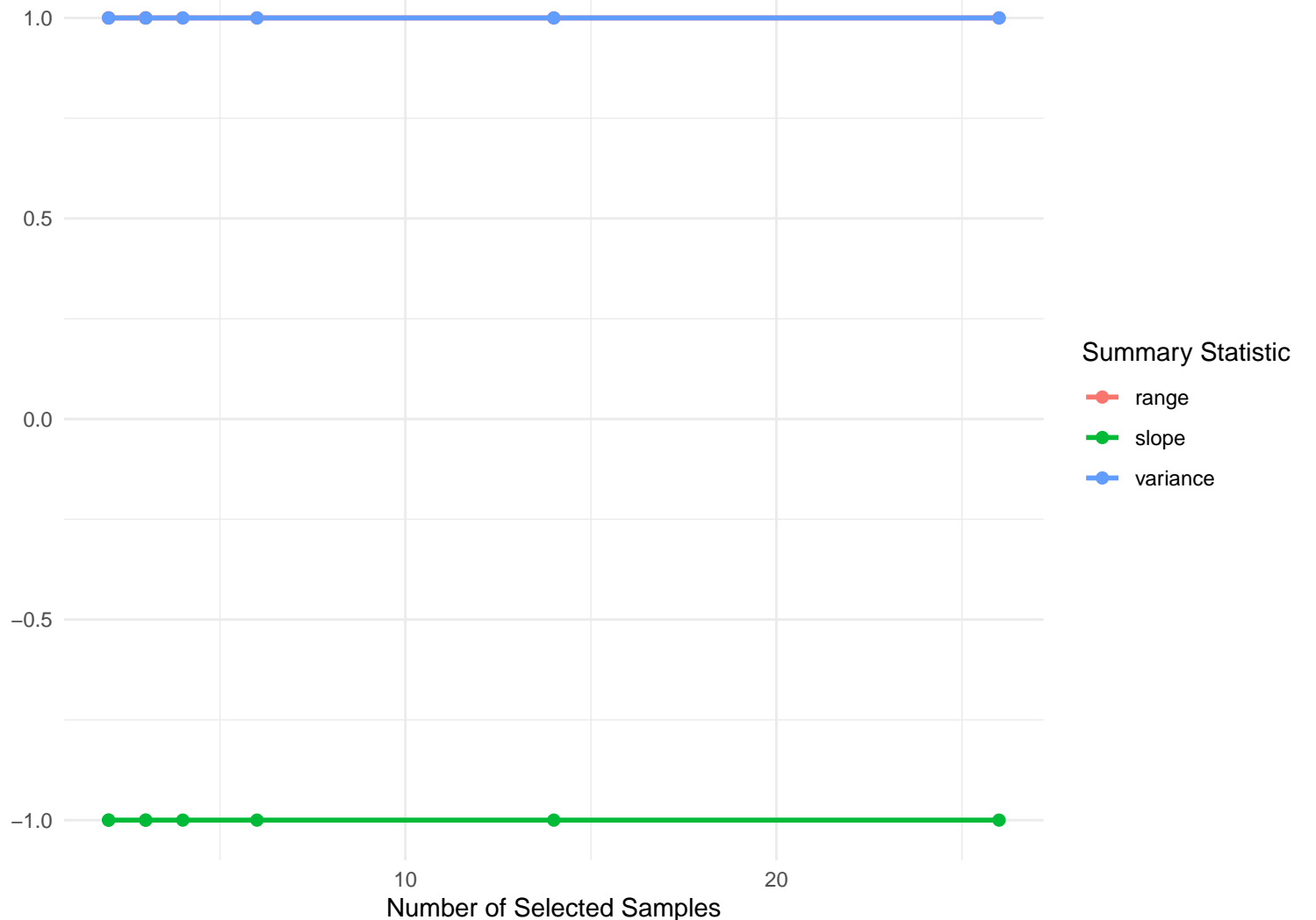
10

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

