

Score: CV\_t (Local, partial1 , Genotype: linear )

Kendall Correlation

1.0  
0.5  
0.0  
-0.5  
-1.0

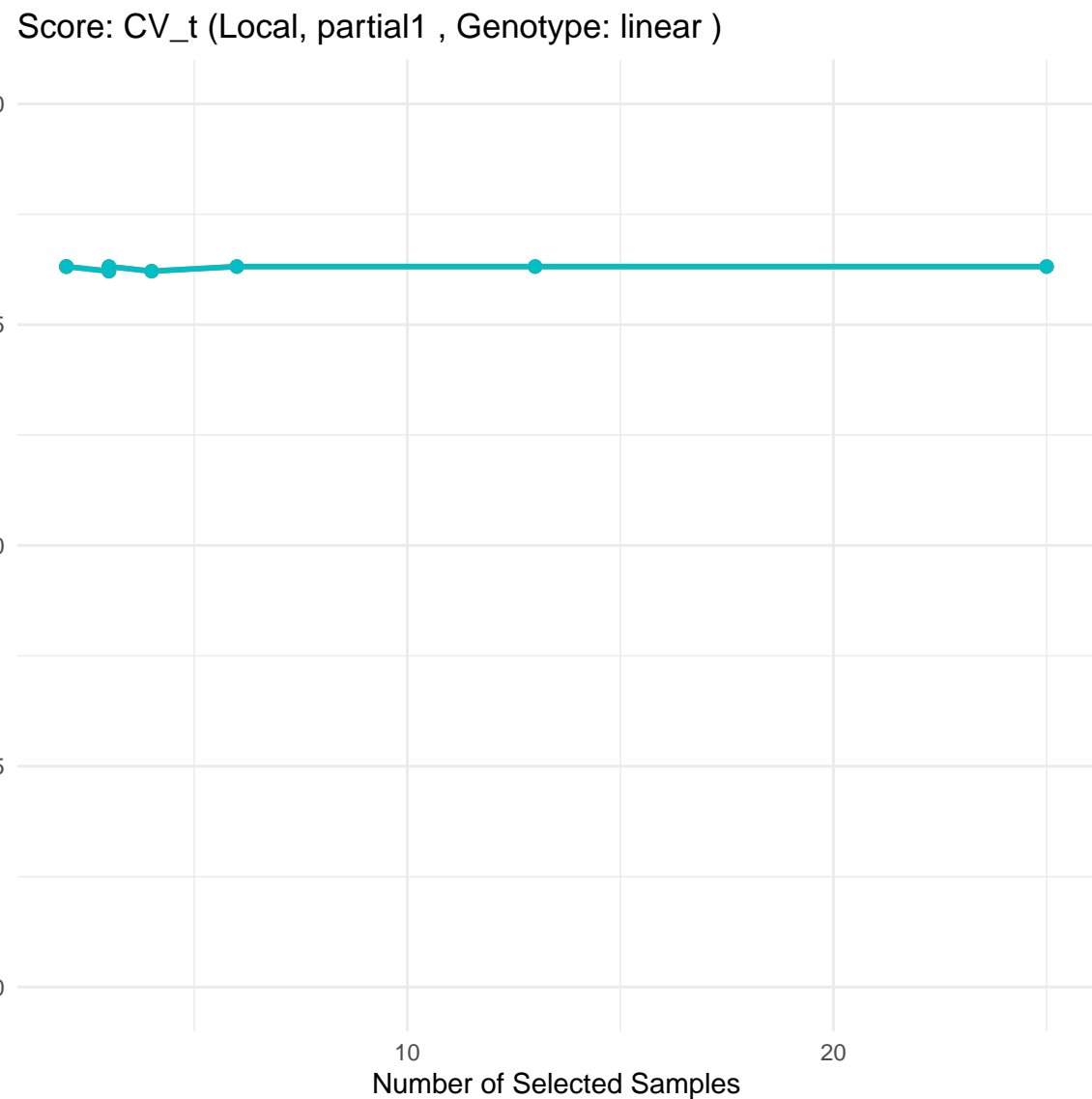
Number of Selected Samples

Summary Statistic

range  
variance

10

20



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

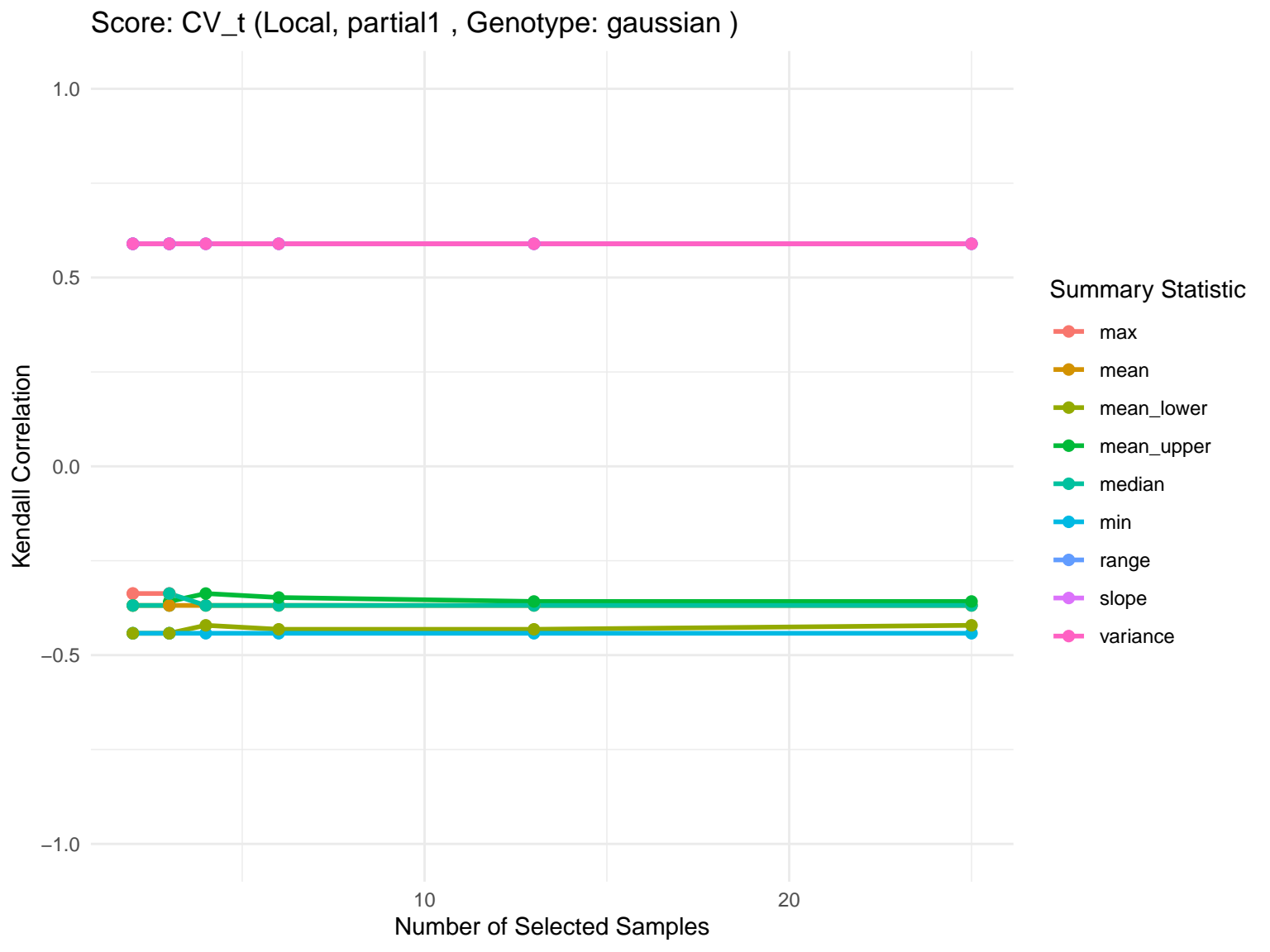
-0.5

-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

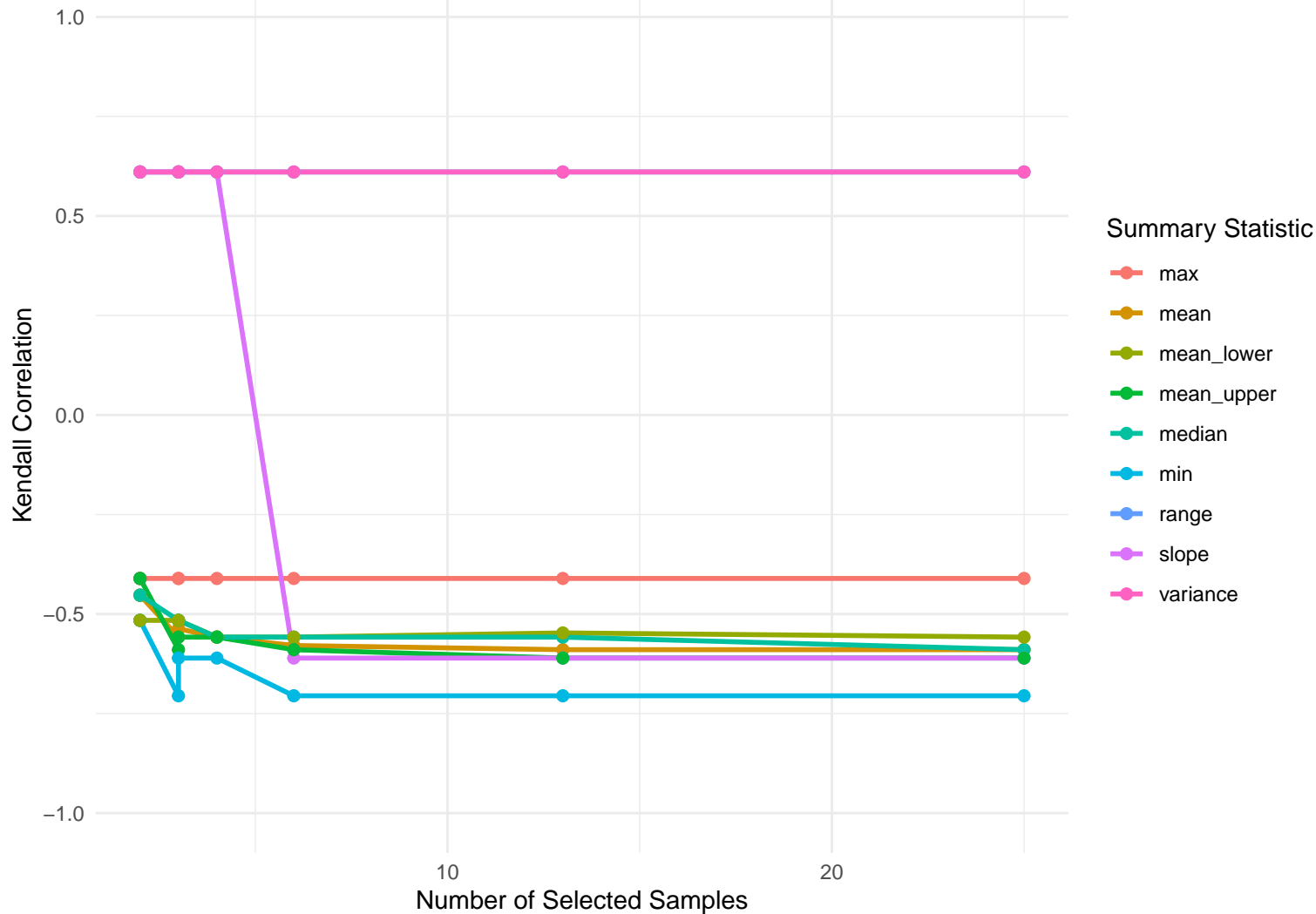
range

slope

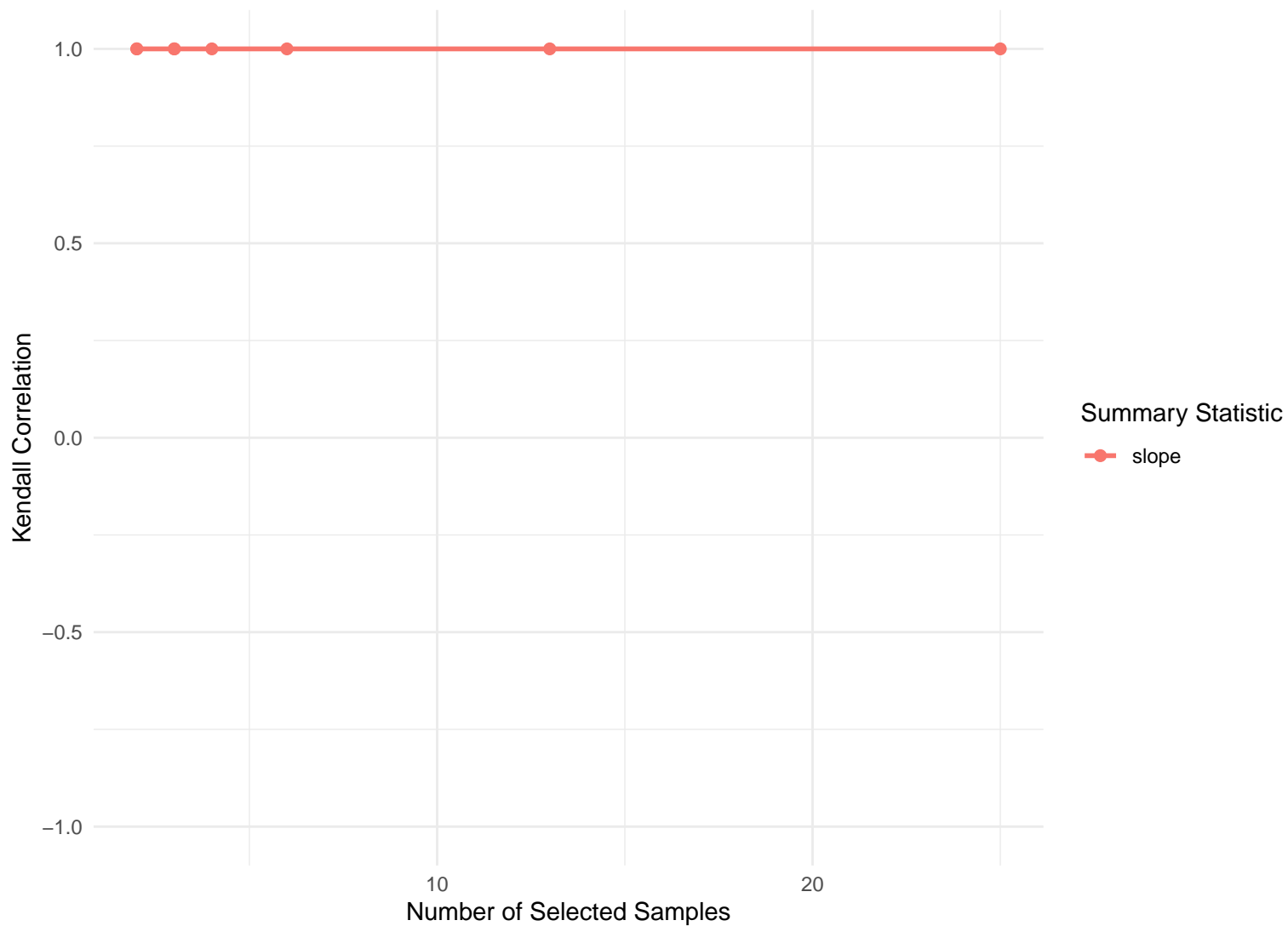
variance

Number of Selected Samples

Score: CV\_t (Local, partial1 , Genotype: wave )



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

20

Number of Selected Samples

Summary Statistic

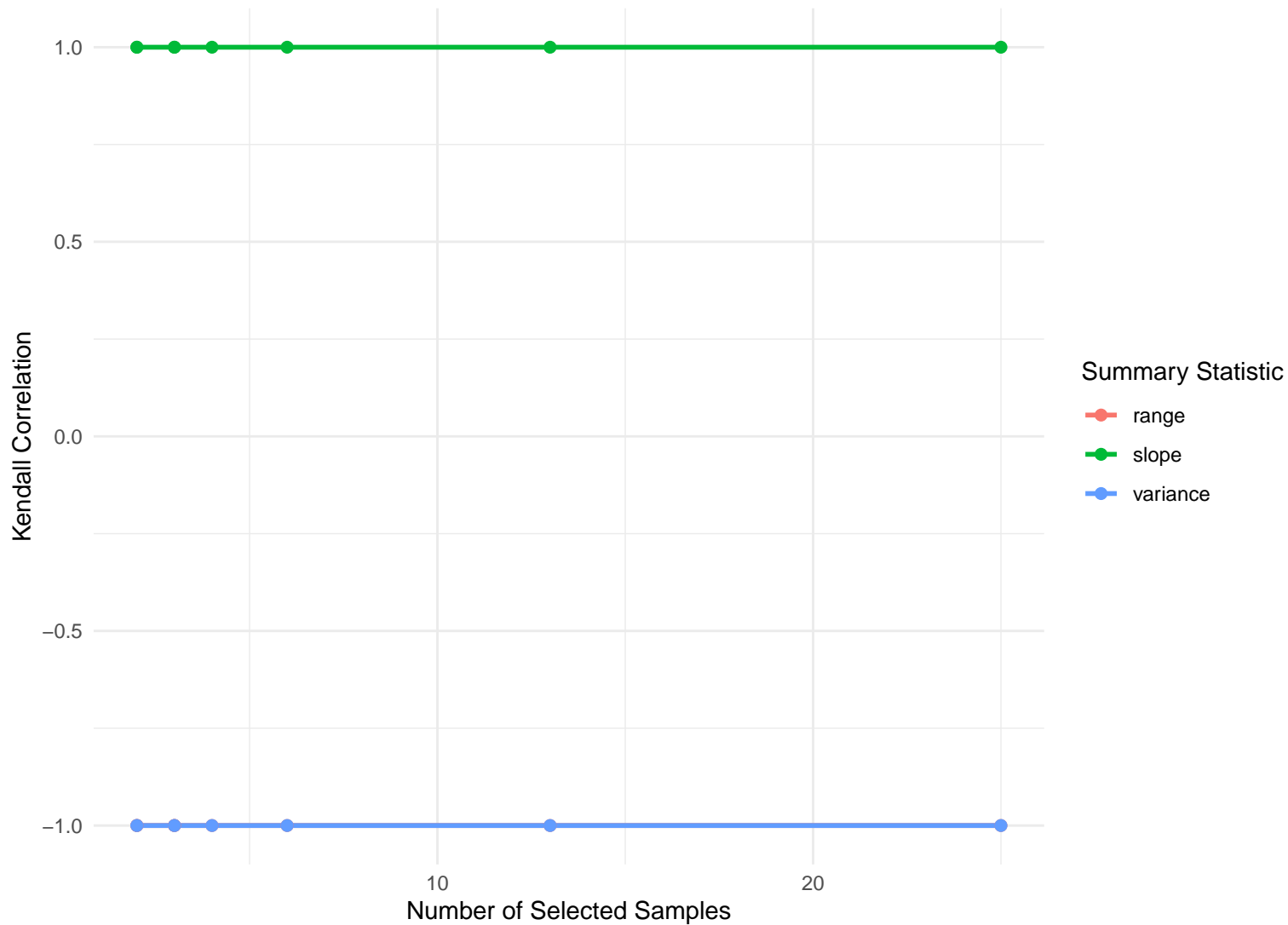
range

slope

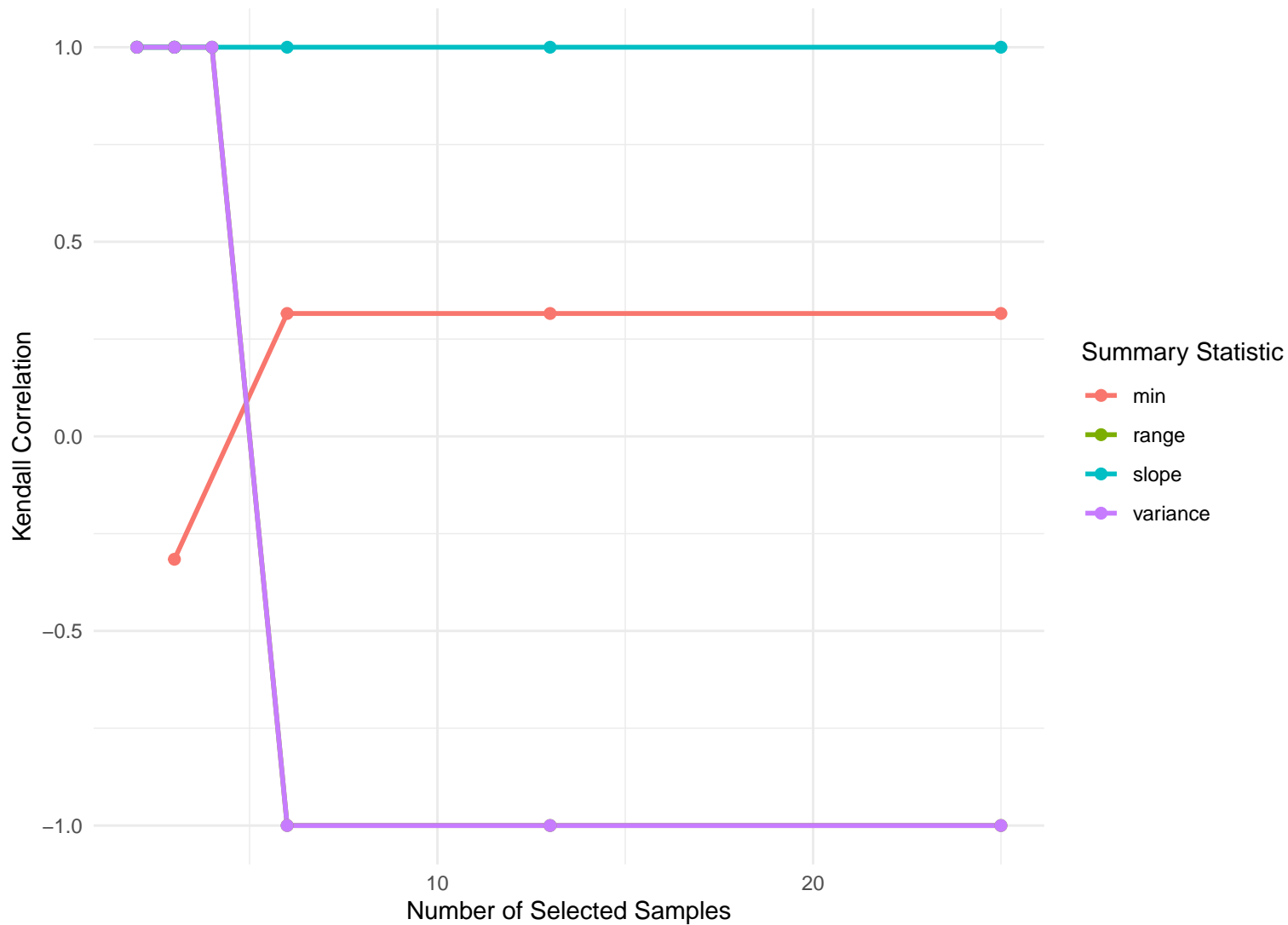
variance



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

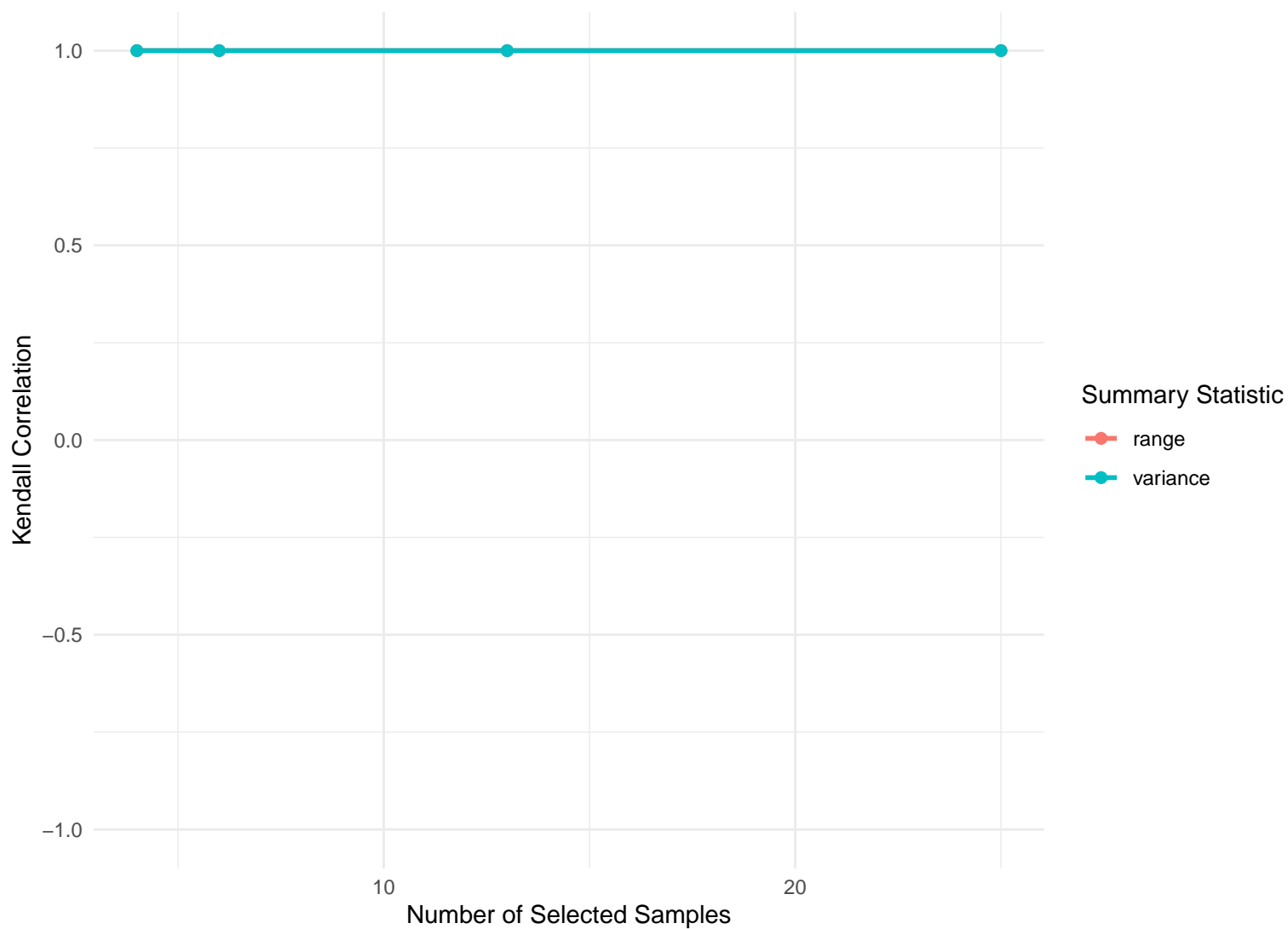


Score: RN (Local, partial1 , Genotype: wave )





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

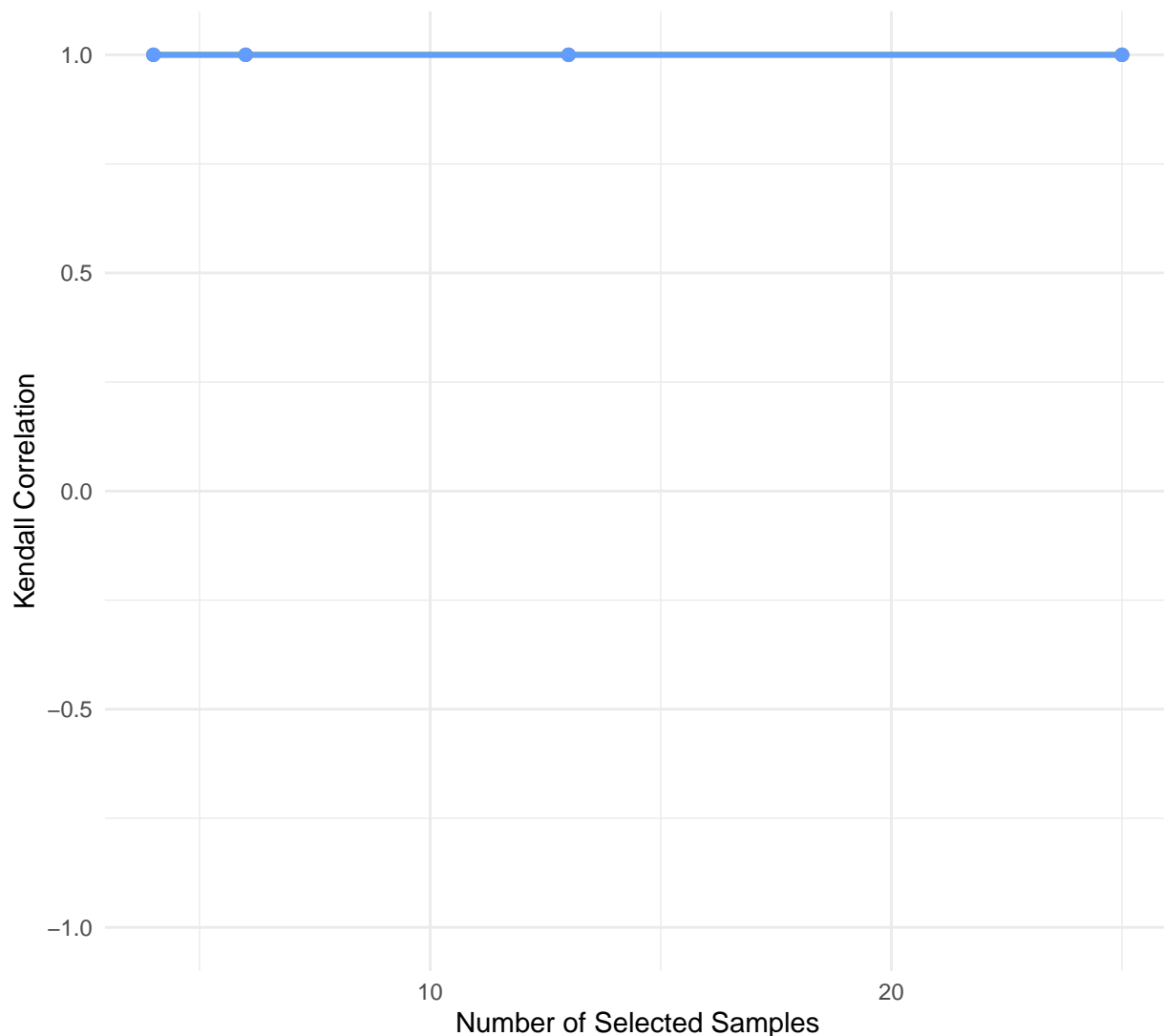
Number of Selected Samples

Summary Statistic

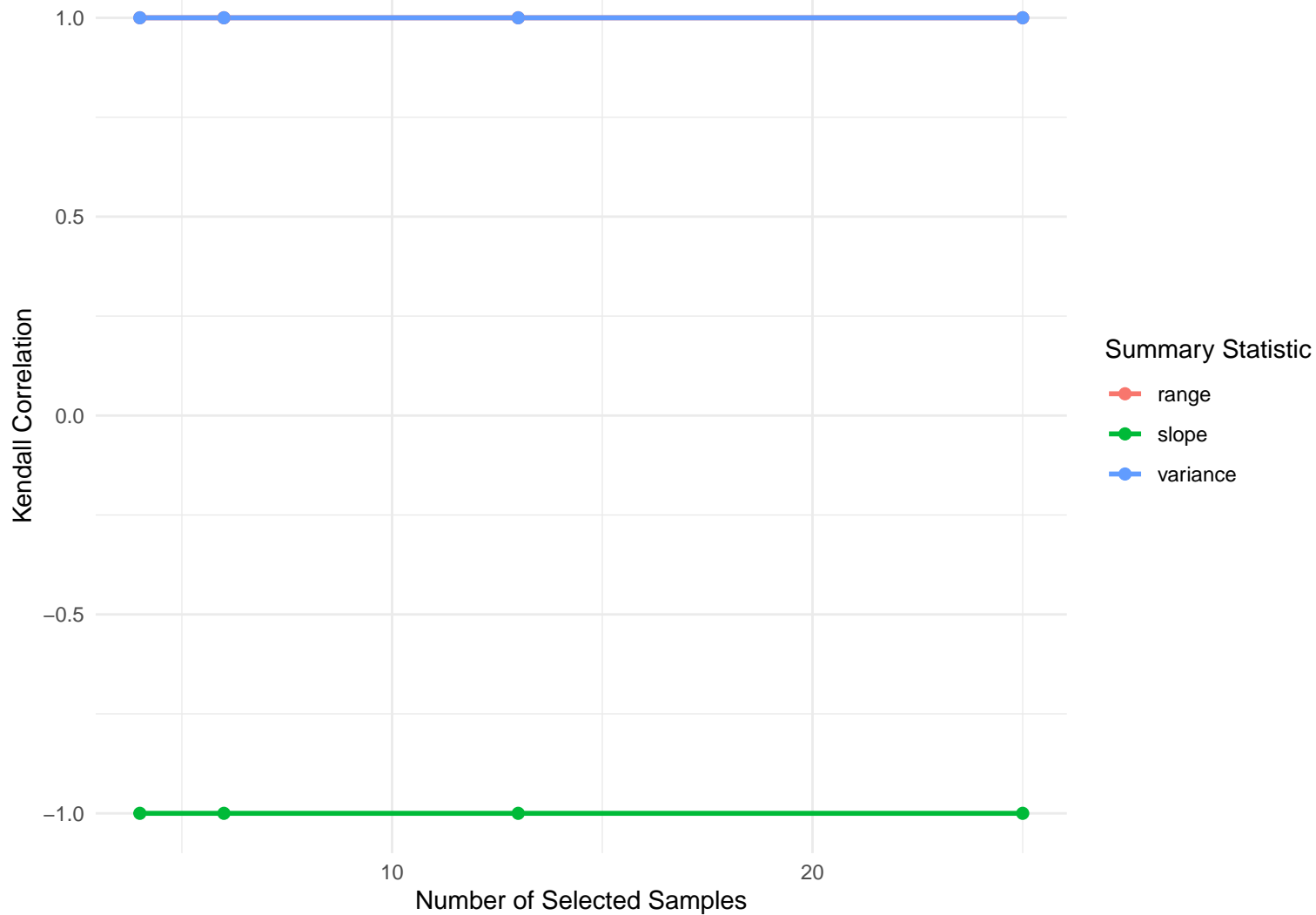
range

slope

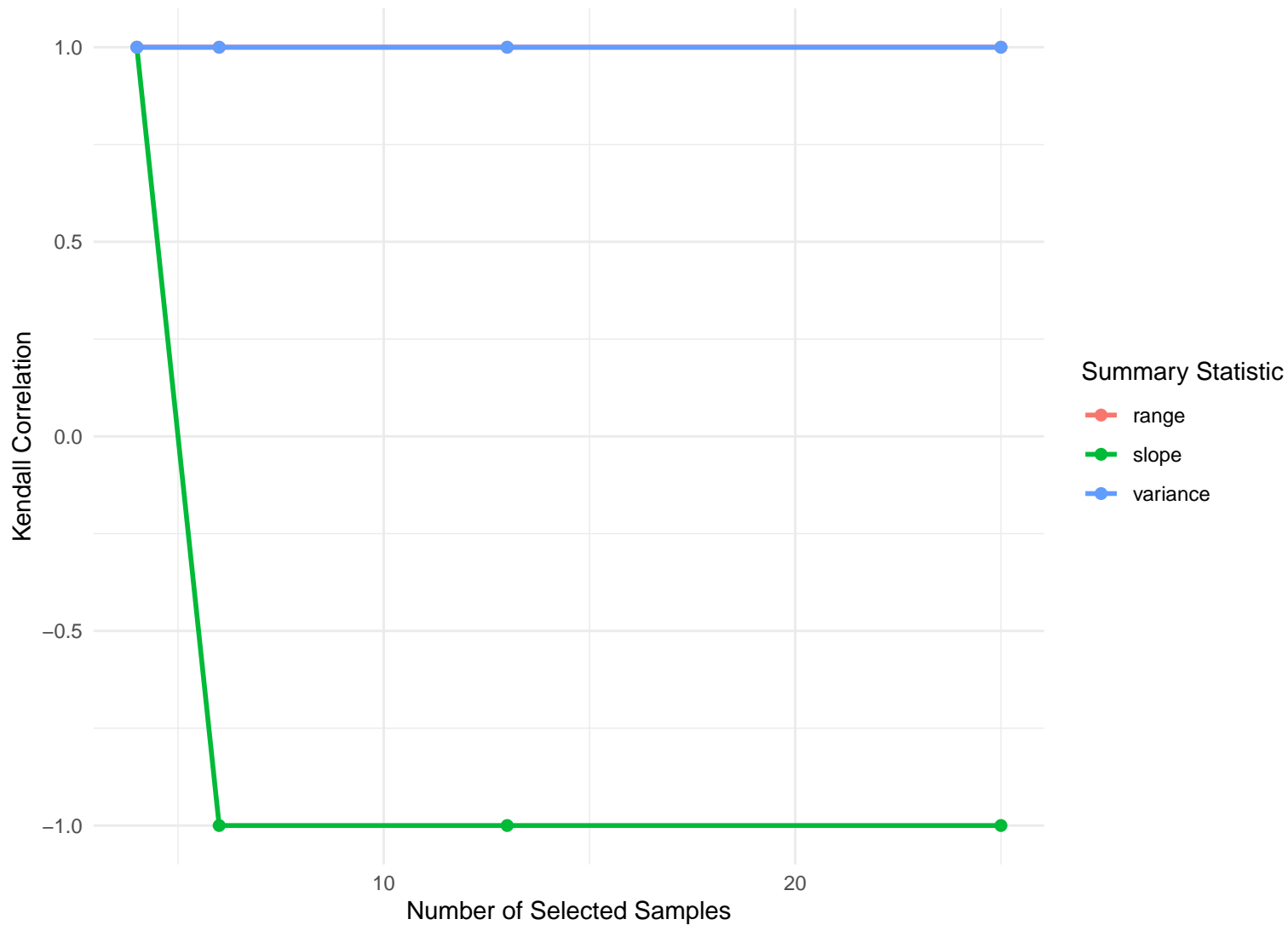
variance



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

10

20

Summary Statistic

range

variance

1.0

0.5

0.0

-0.5

-1.0

10

20

Summary Statistic

range

variance

Score: D\_slope (Local, partial1 , Genotype: gaussian )

Kendall Correlation

1.0  
0.5  
0.0  
-0.5  
-1.0

Number of Selected Samples

Summary Statistic

- range
- slope
- variance



Score: D\_slope (Local, partial1 , Genotype: sinusoidal )

Kendall Correlation

Summary Statistic

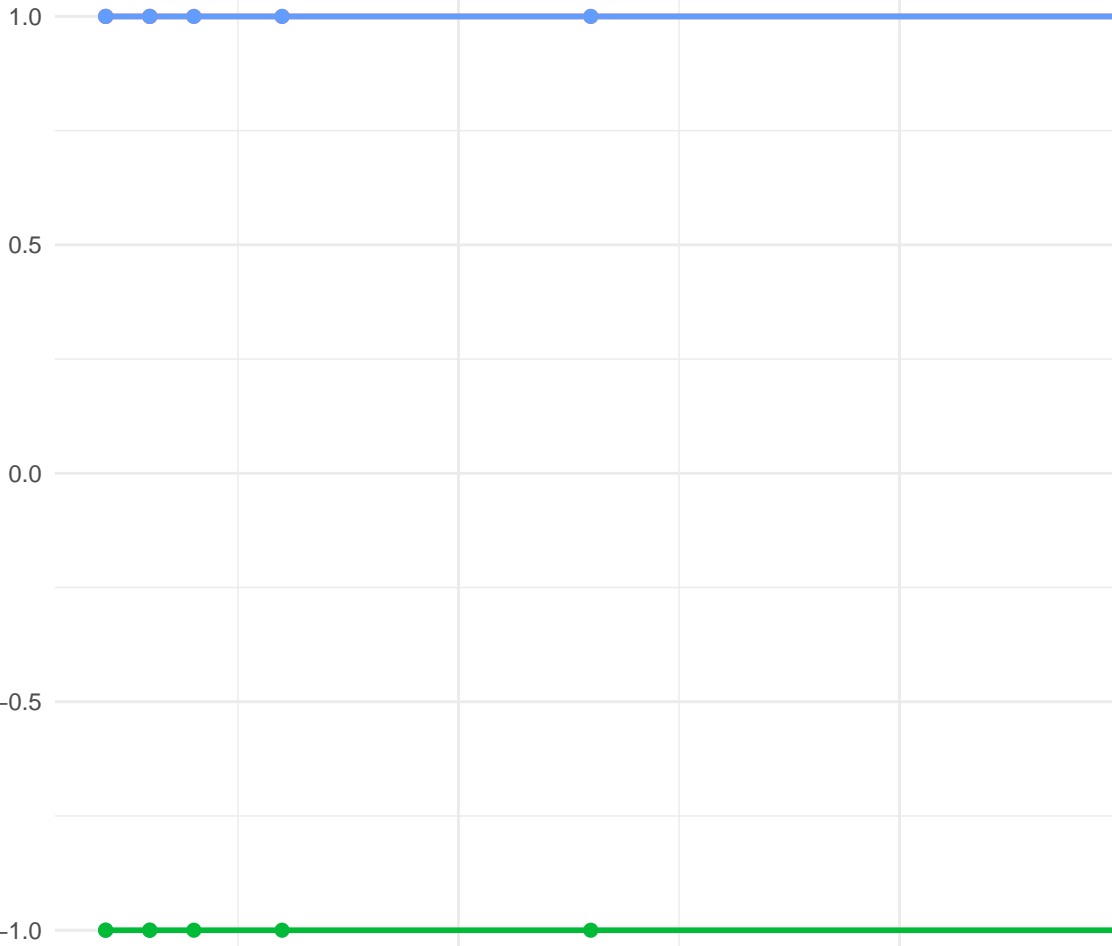
- range
- slope
- variance

1.0  
0.5  
0.0  
-0.5  
-1.0

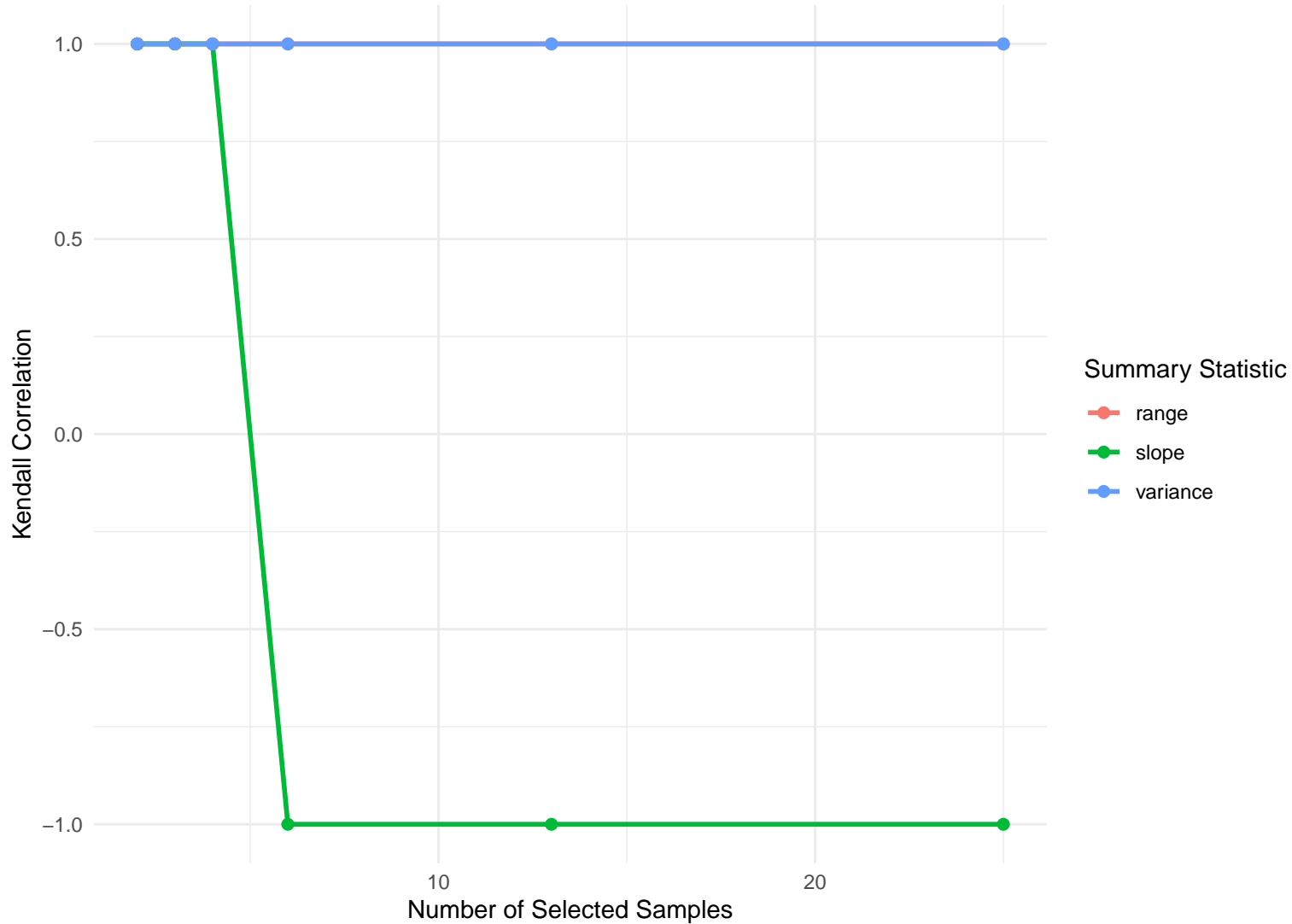
Number of Selected Samples

10

20

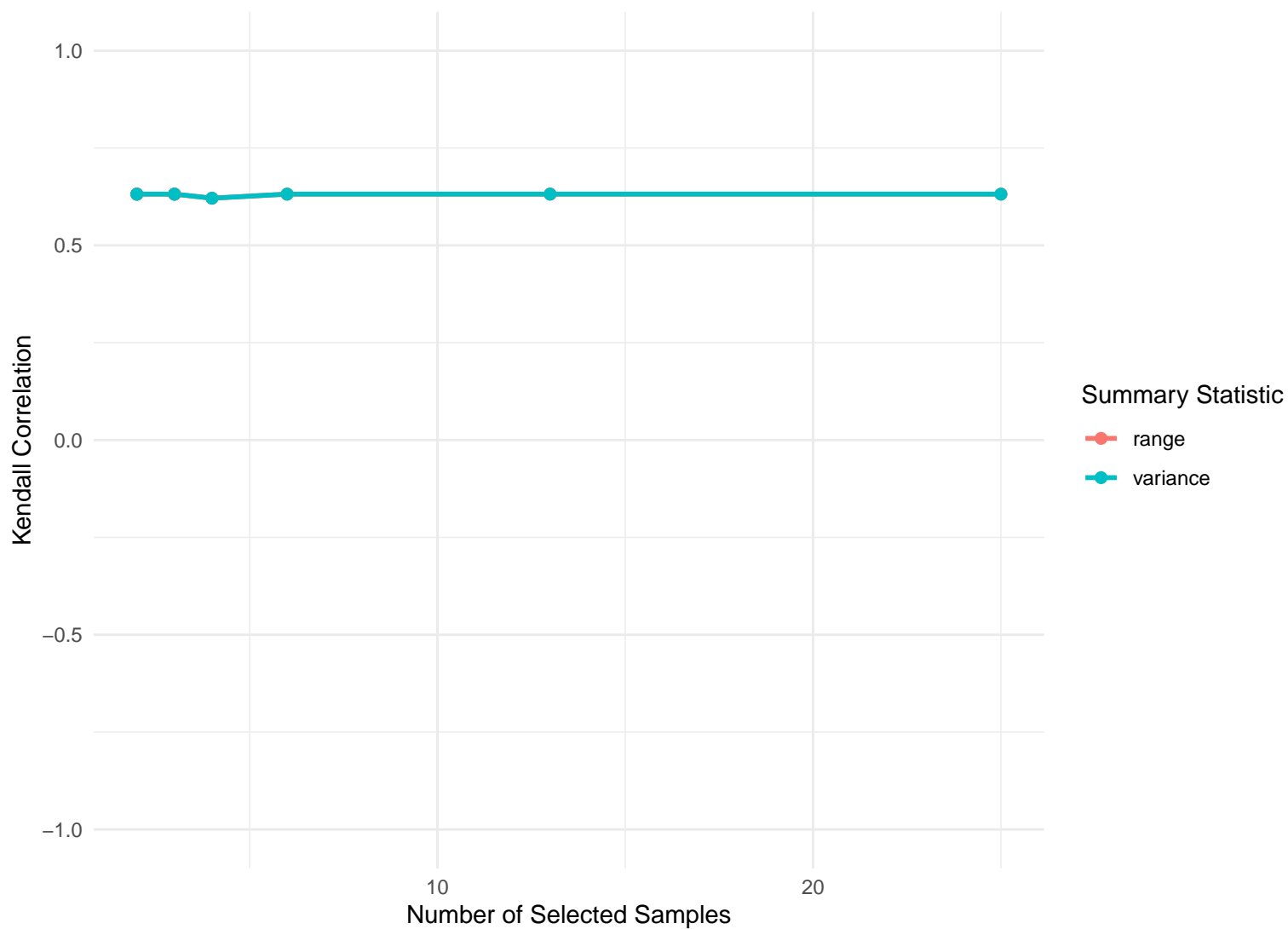


Score: D\_slope (Local, partial1 , Genotype: wave )





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

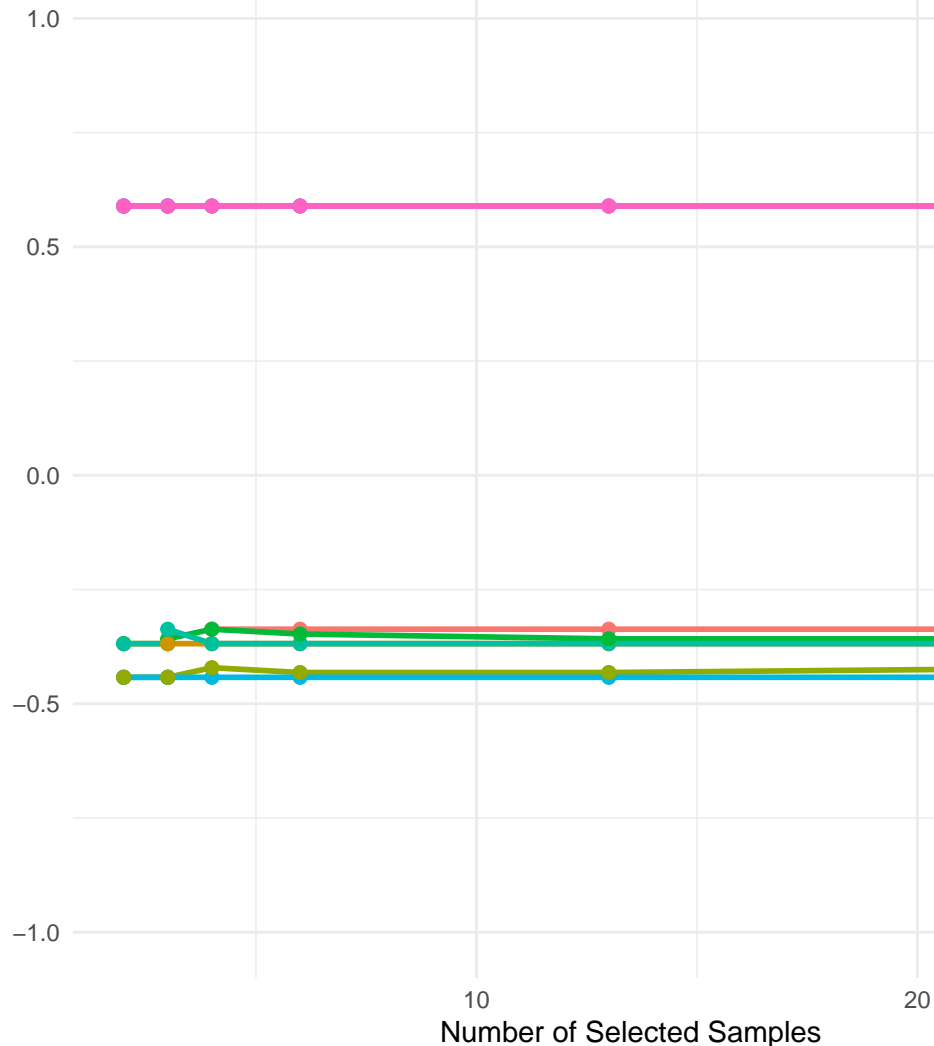


Score: RC (Local, partial1 , Genotype: gaussian )

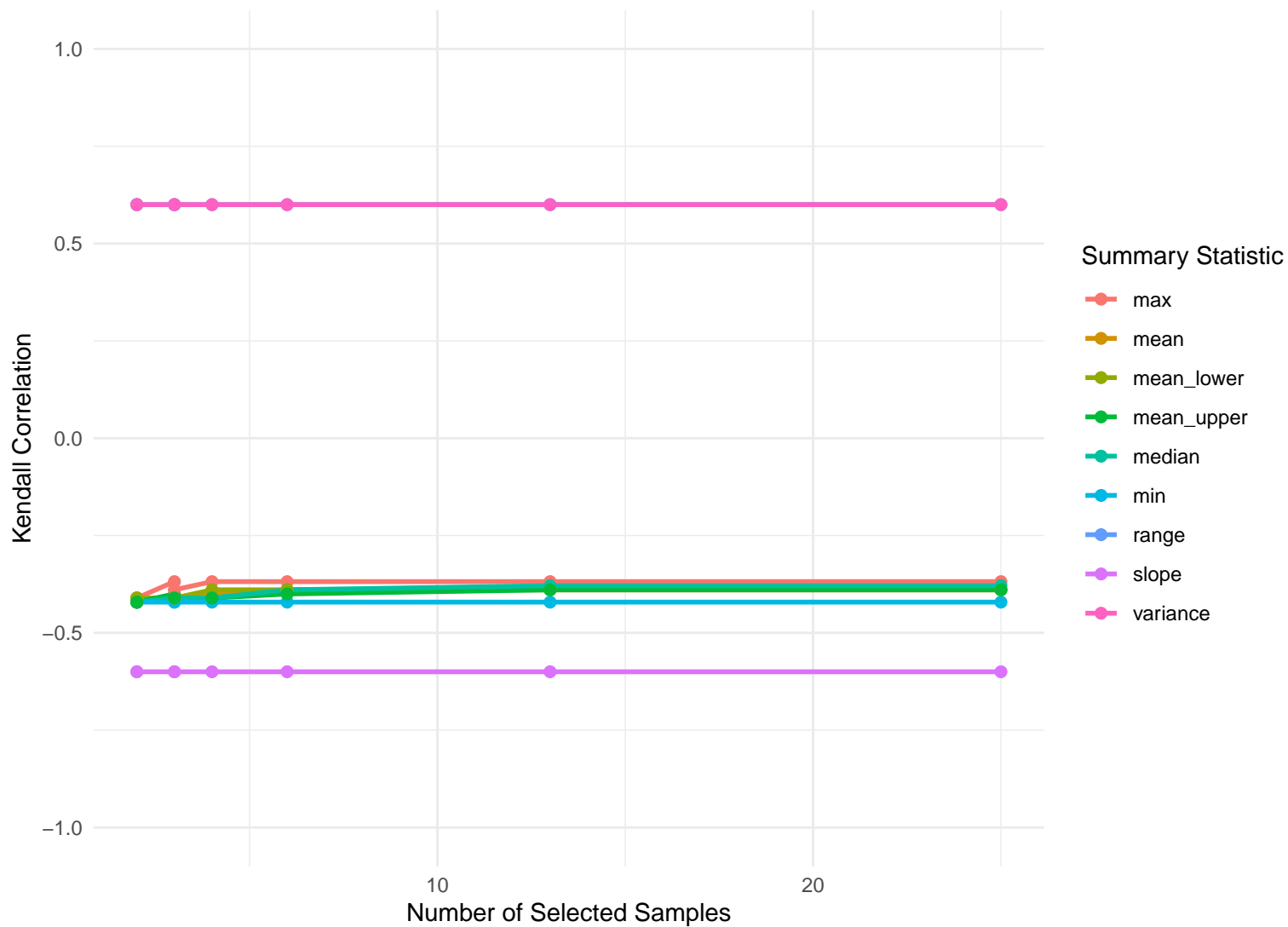
Kendall Correlation

Summary Statistic

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



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

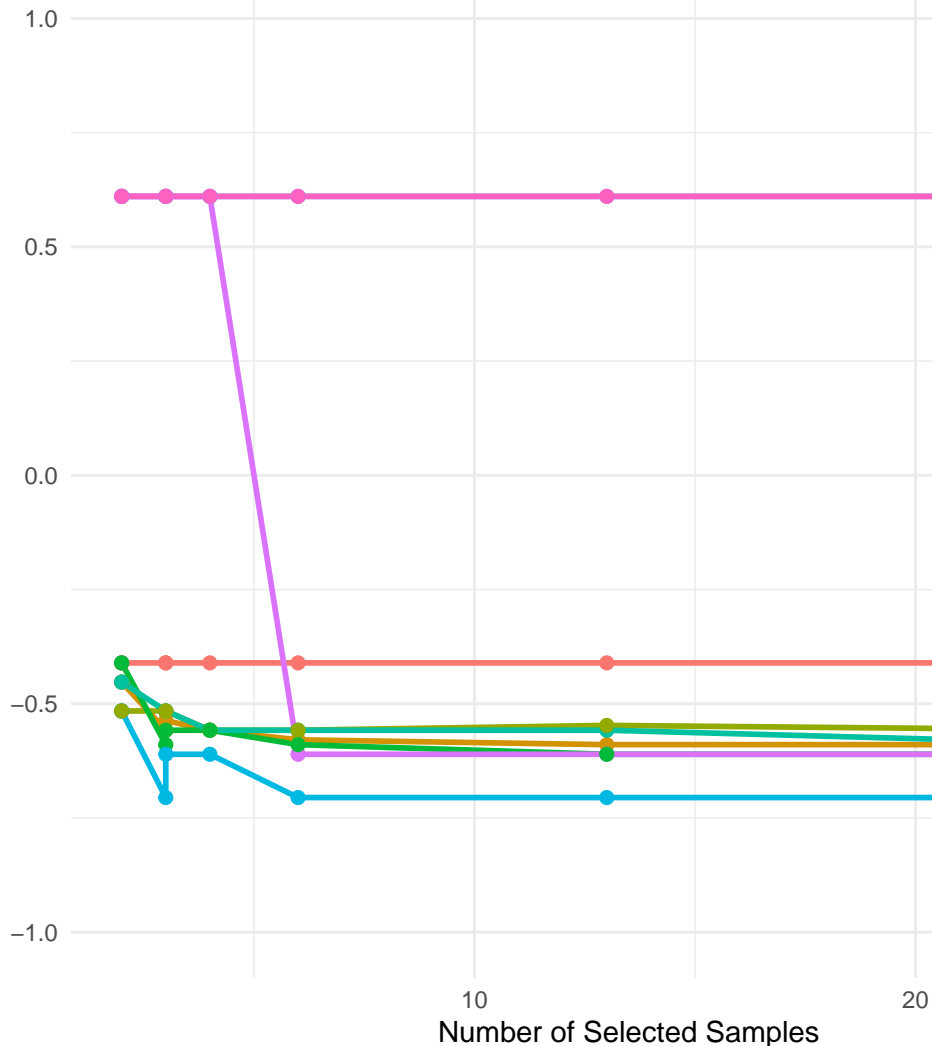


Score: RC (Local, partial1 , Genotype: wave )

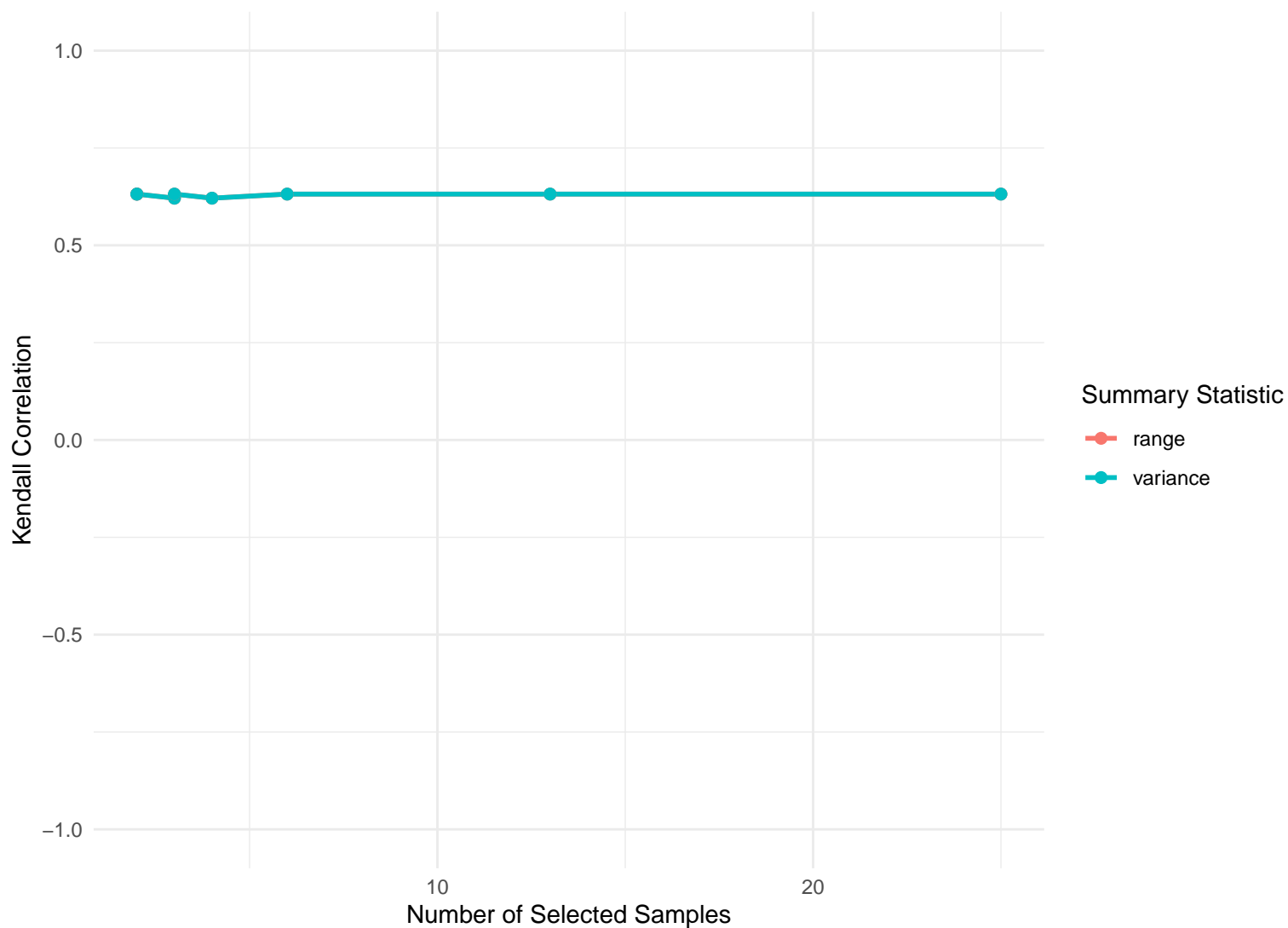
Kendall Correlation

Summary Statistic

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



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

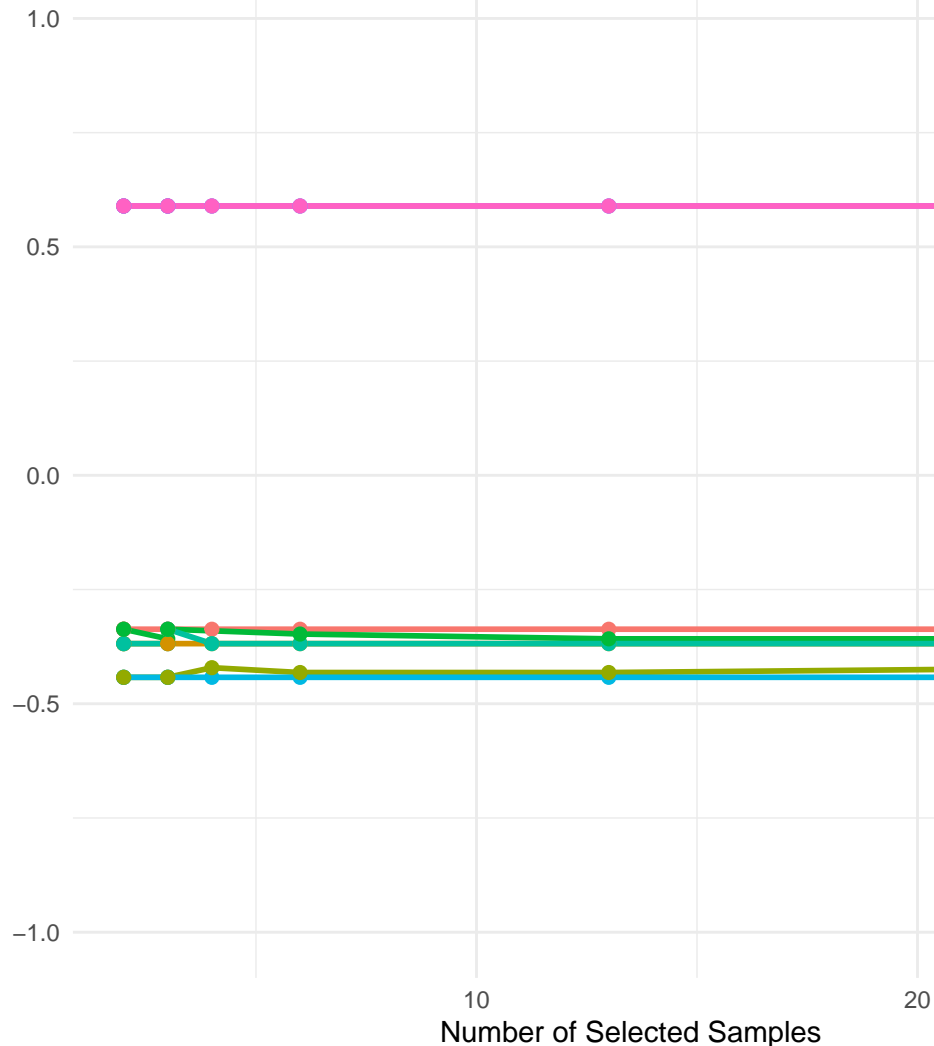


Score: gPi (Local, partial1 , Genotype: gaussian )

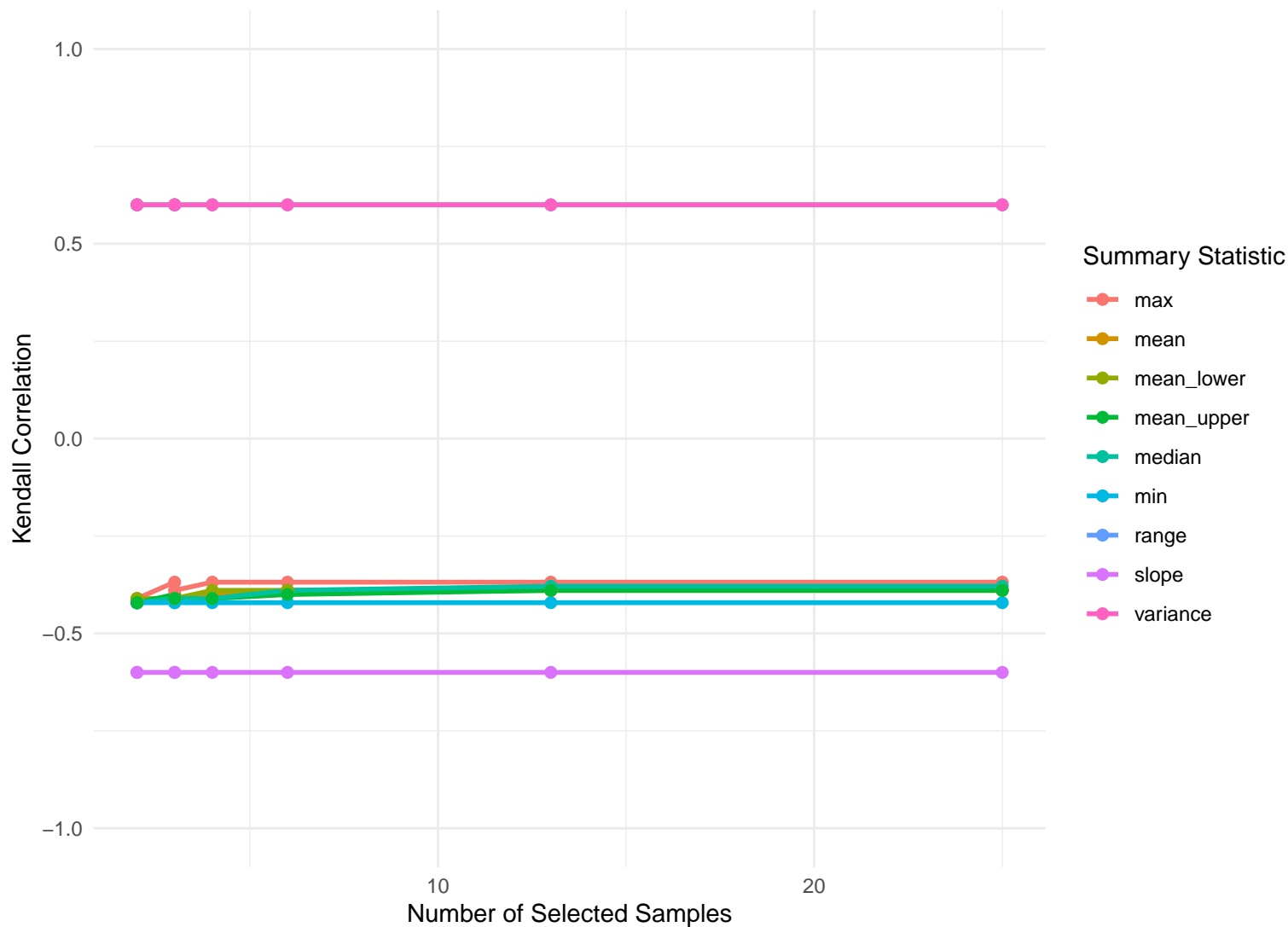
Kendall Correlation

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 )

Kendall Correlation

Summary Statistic

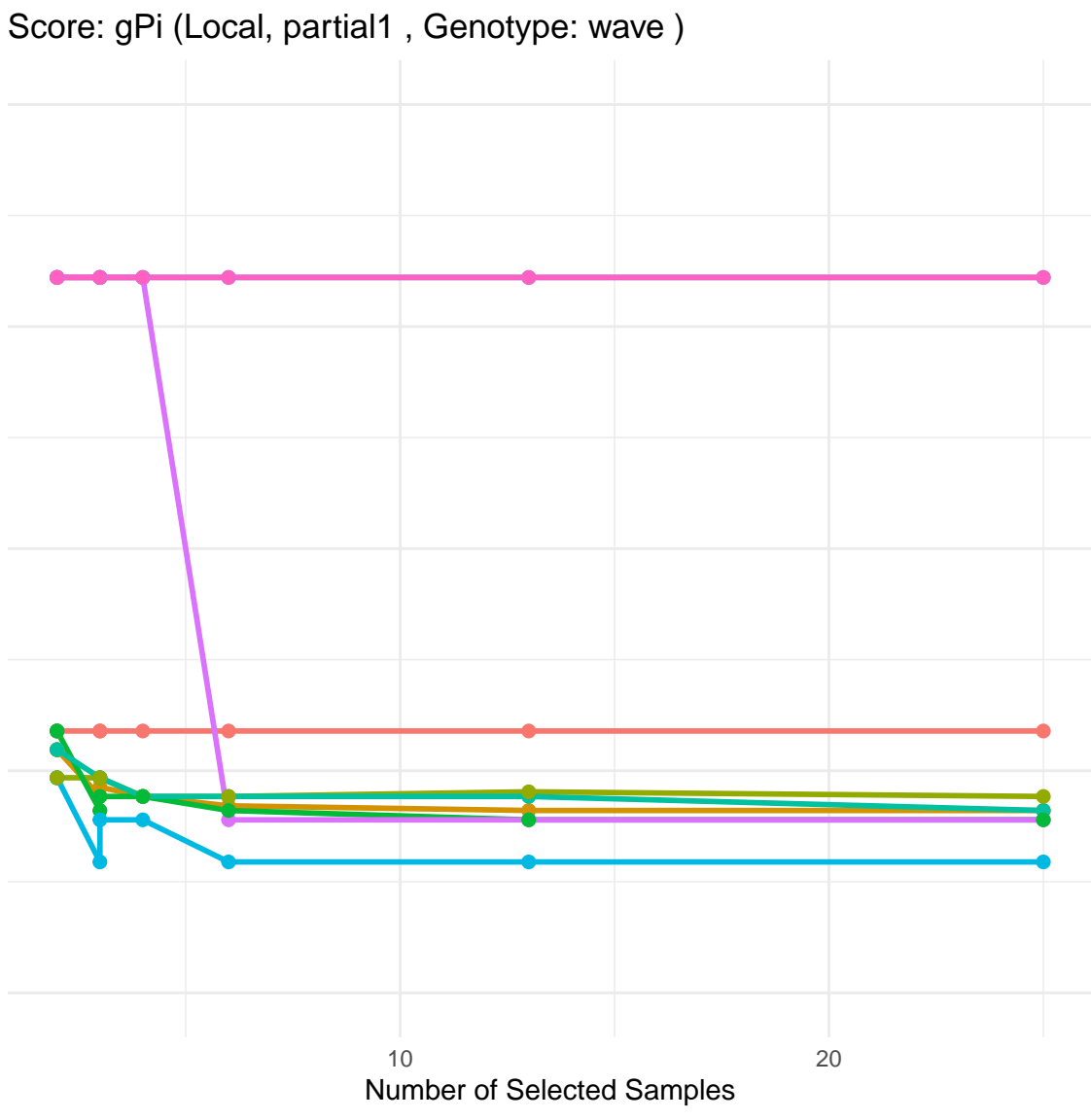
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

1.0  
0.5  
0.0  
-0.5  
-1.0

Number of Selected Samples

10

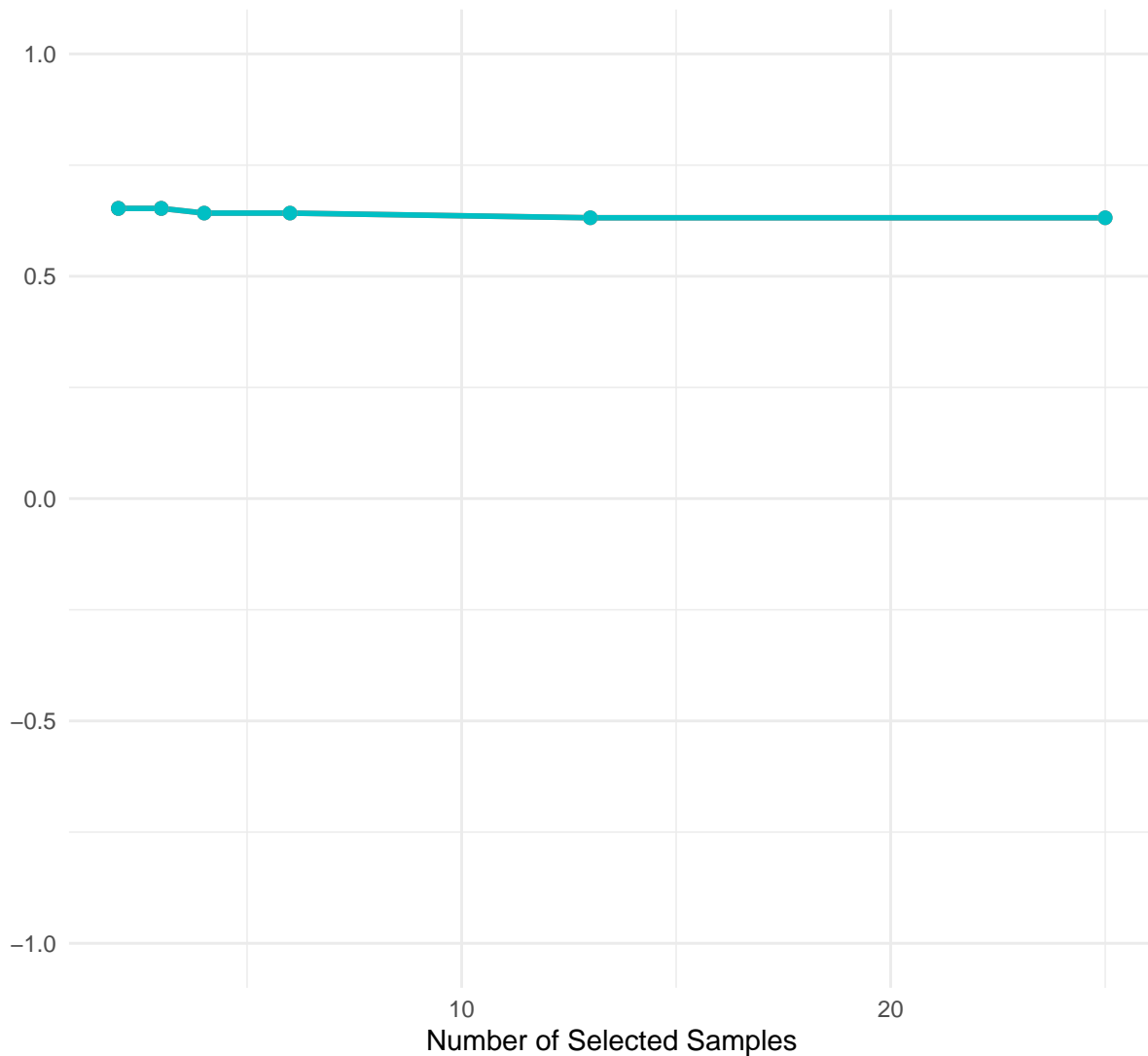
20





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

Kendall Correlation



Summary Statistic

- range
- variance

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

Kendall Correlation

Summary Statistic

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

1.0

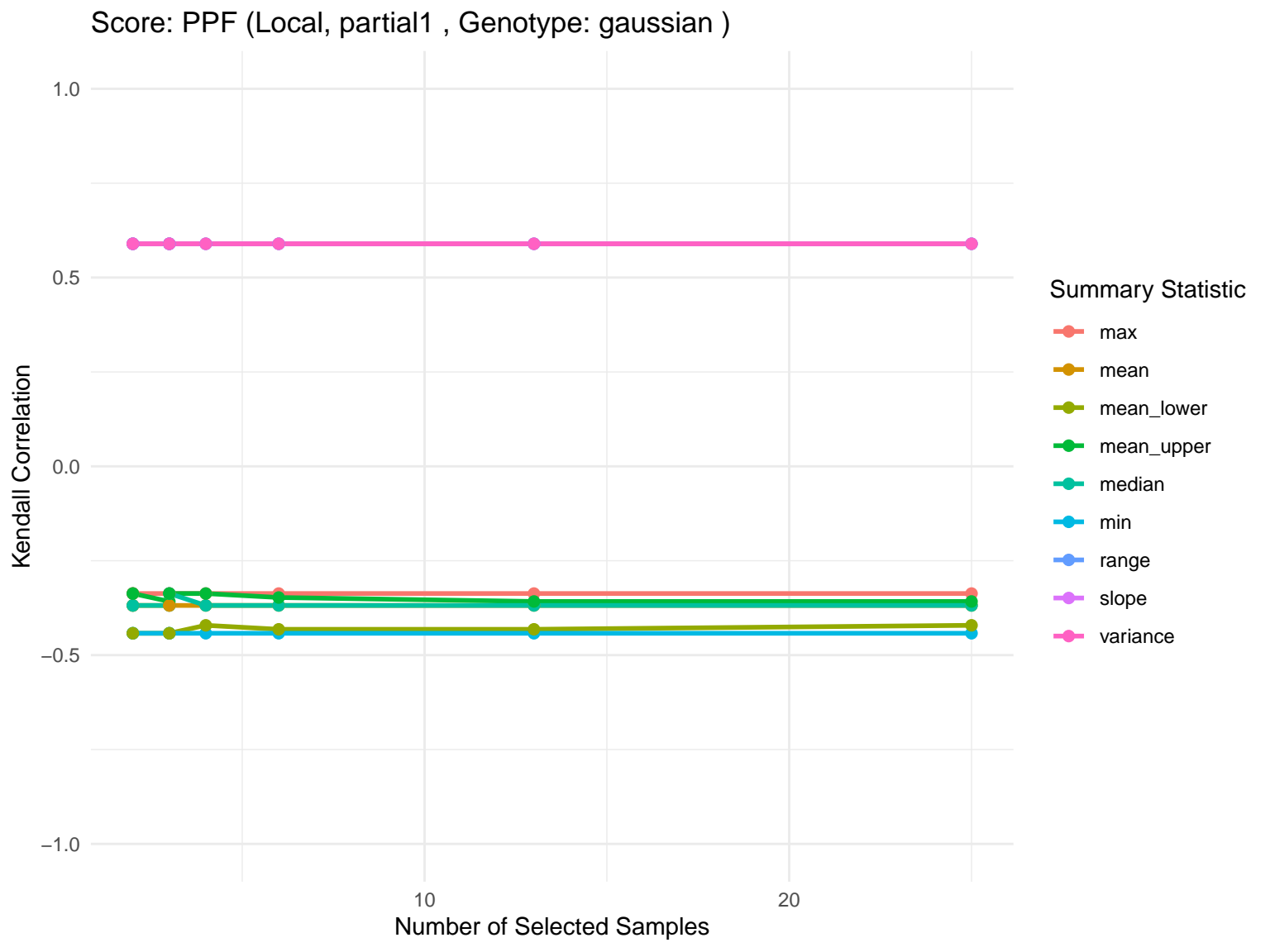
0.5

0.0

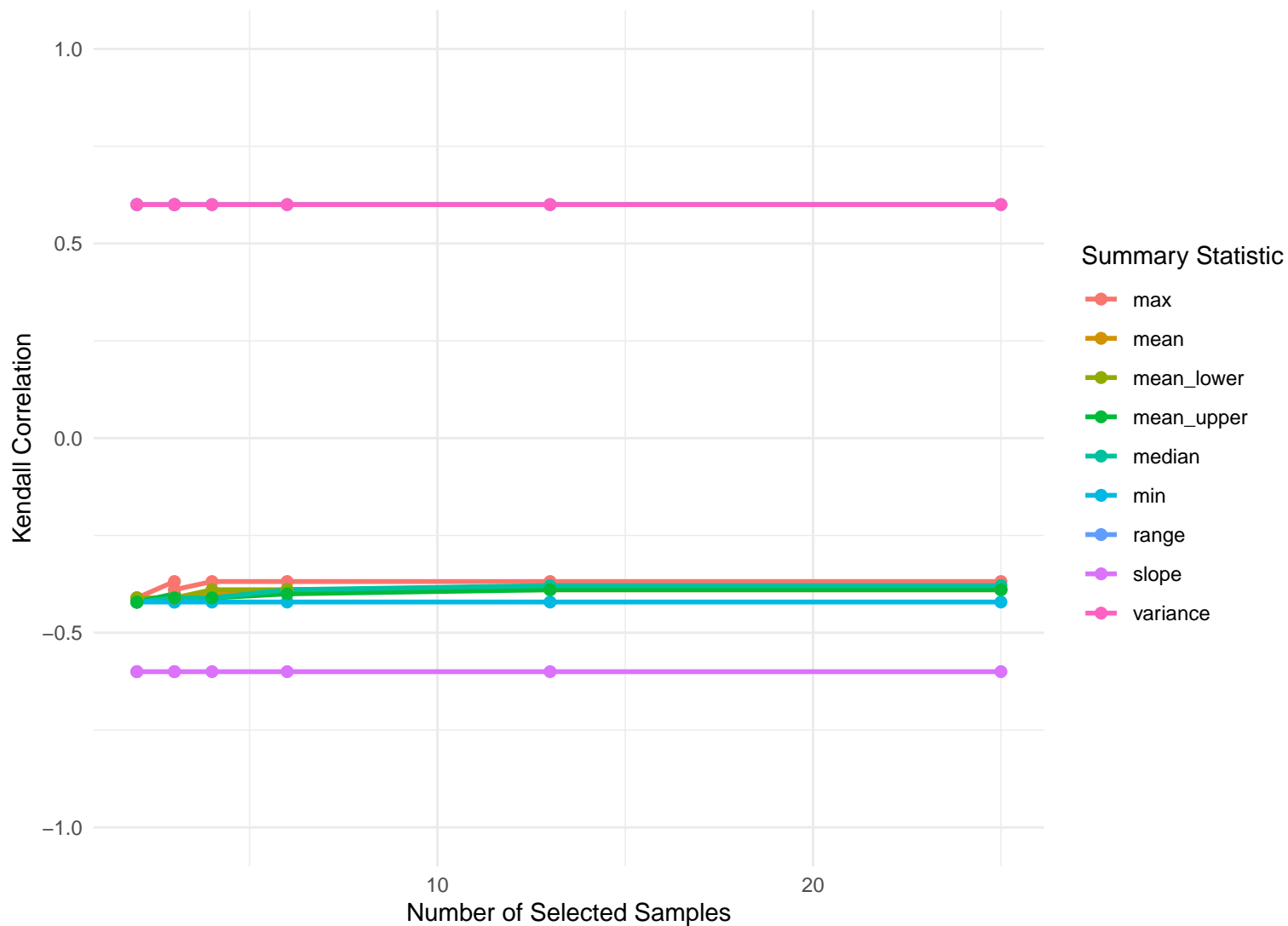
-0.5

-1.0

10  
20  
Number of Selected Samples



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



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

Kendall Correlation

Summary Statistic

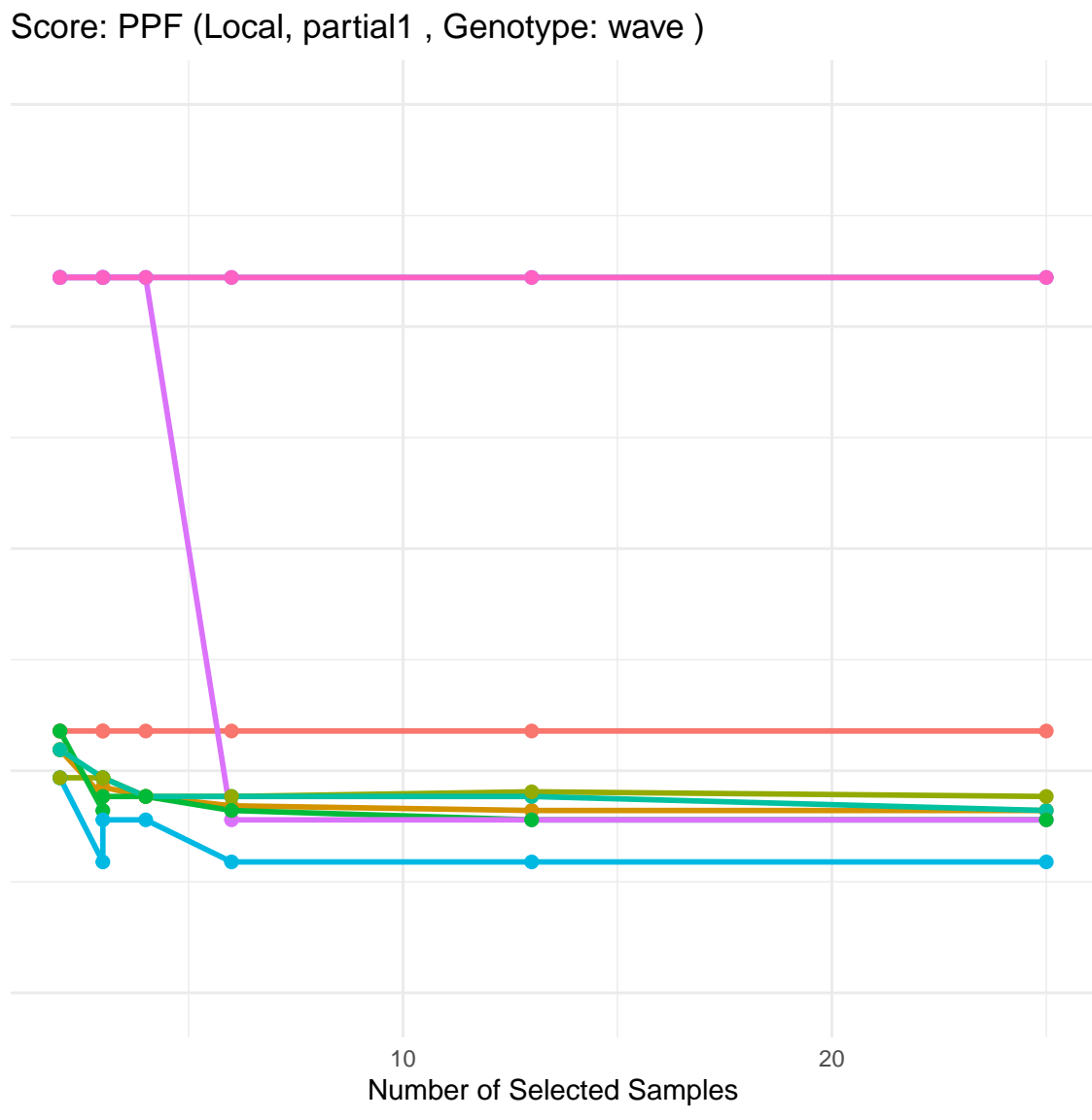
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

1.0  
0.5  
0.0  
-0.5  
-1.0

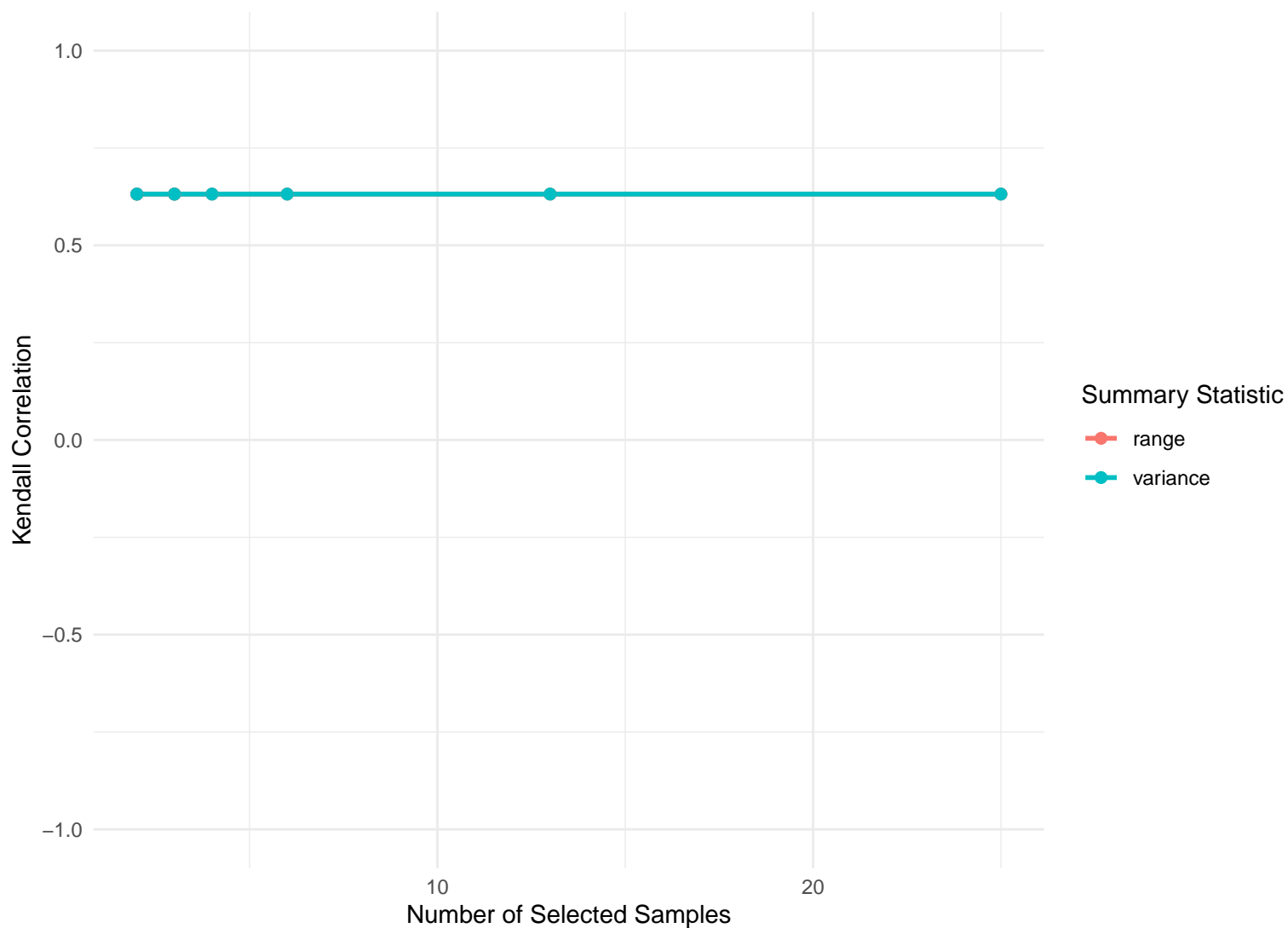
Number of Selected Samples

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

1.0

0.5

0.0

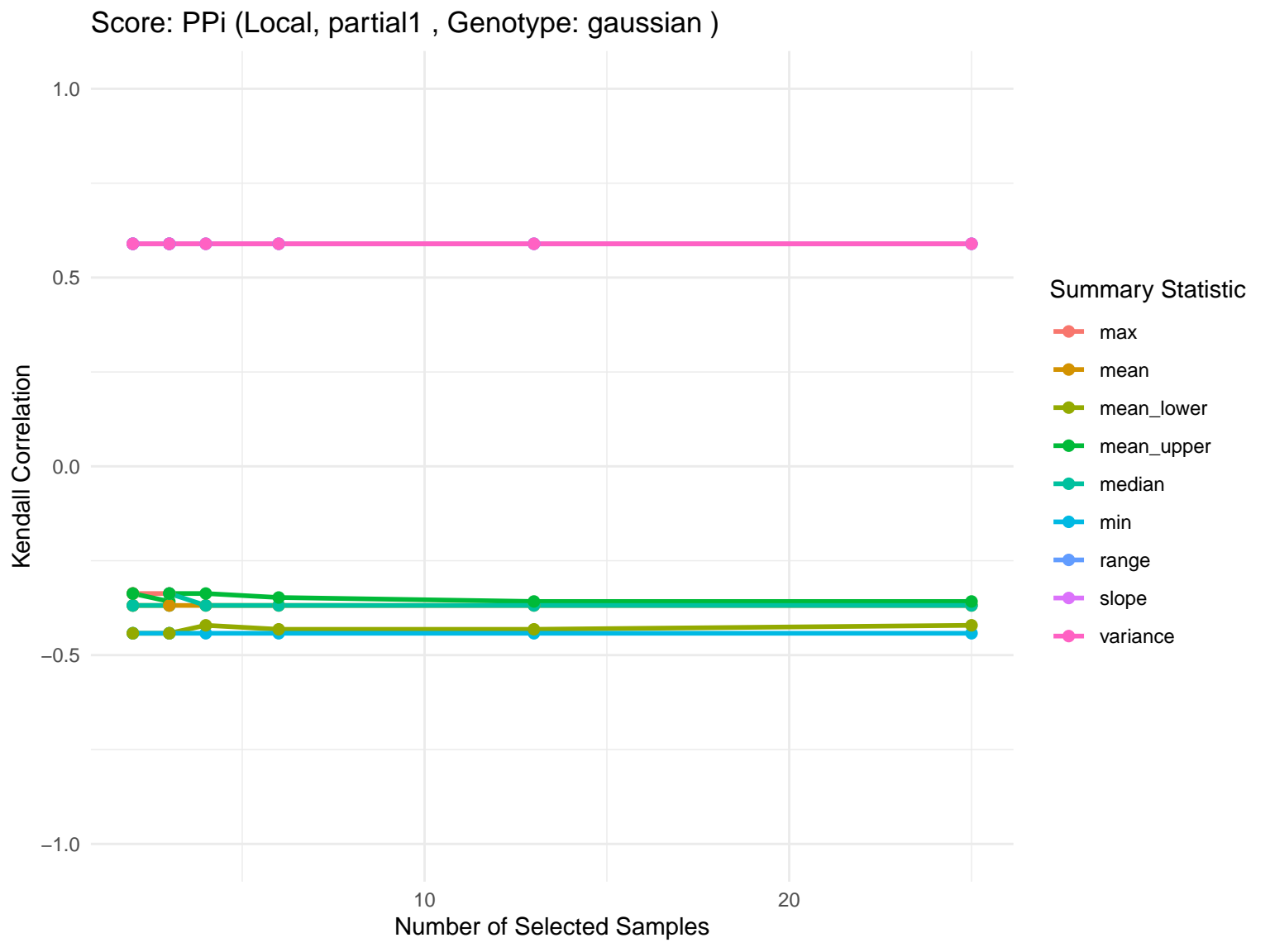
-0.5

-1.0

Number of Selected Samples

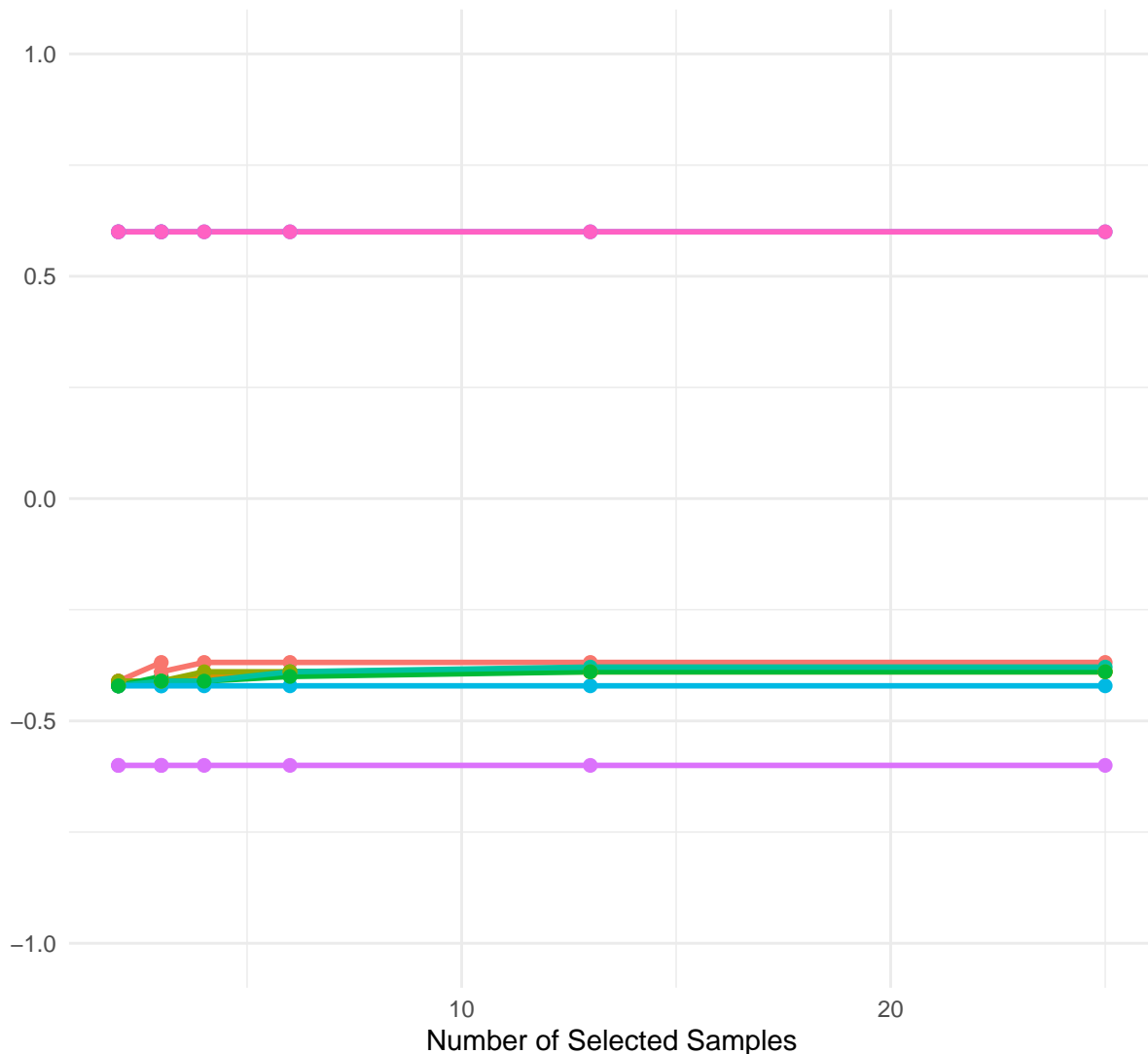
10

20



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

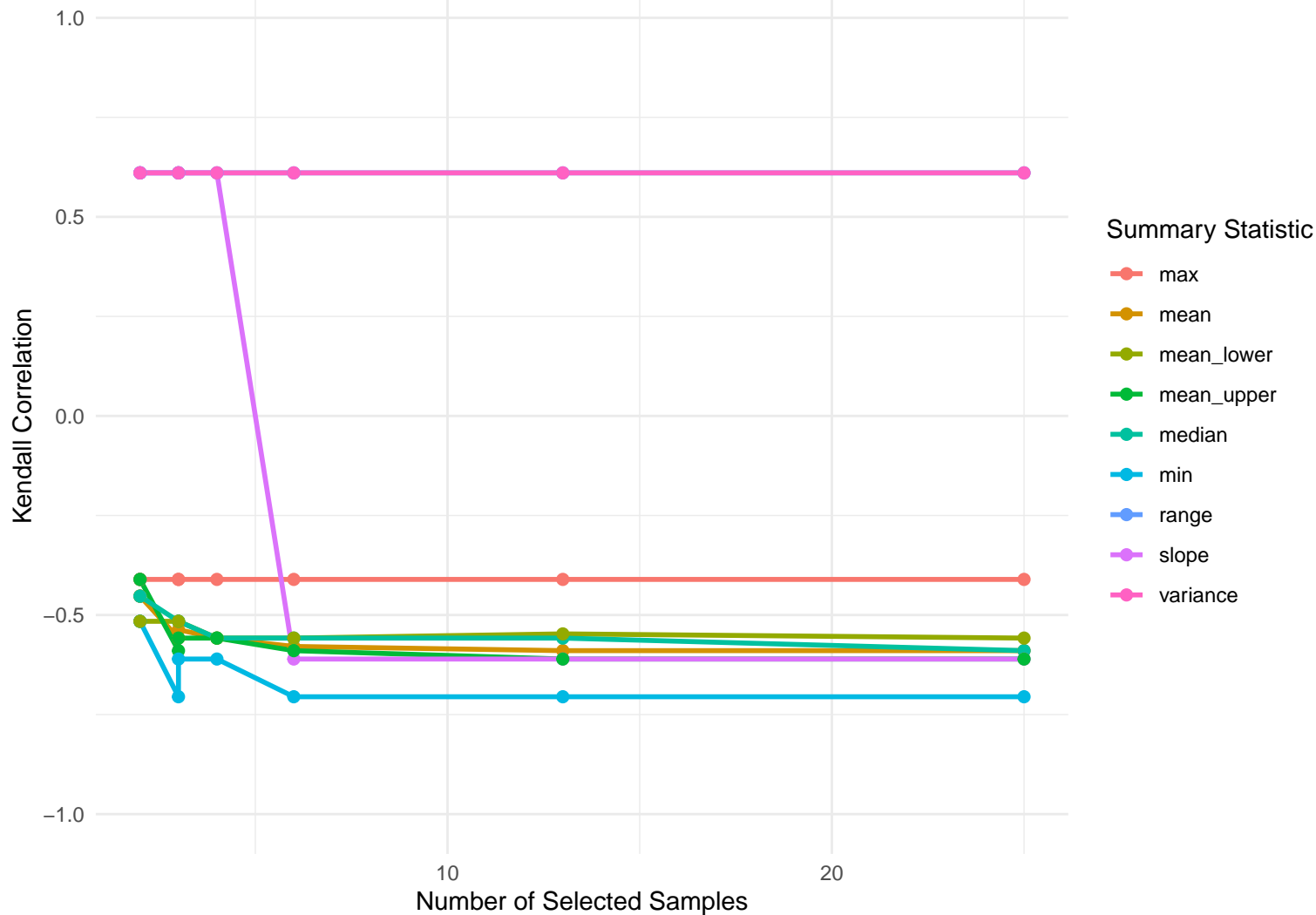
Kendall Correlation



Summary Statistic

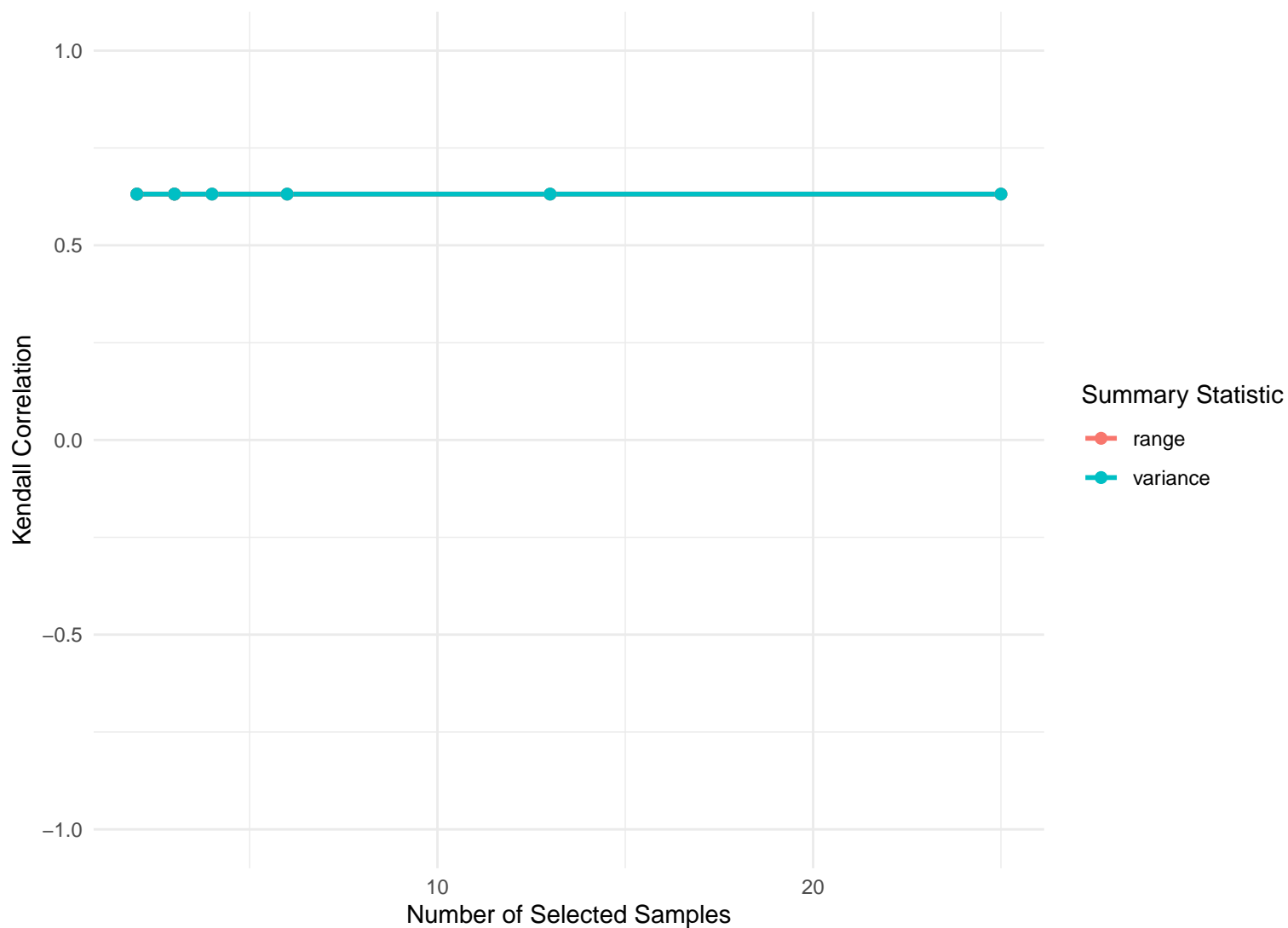
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance

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

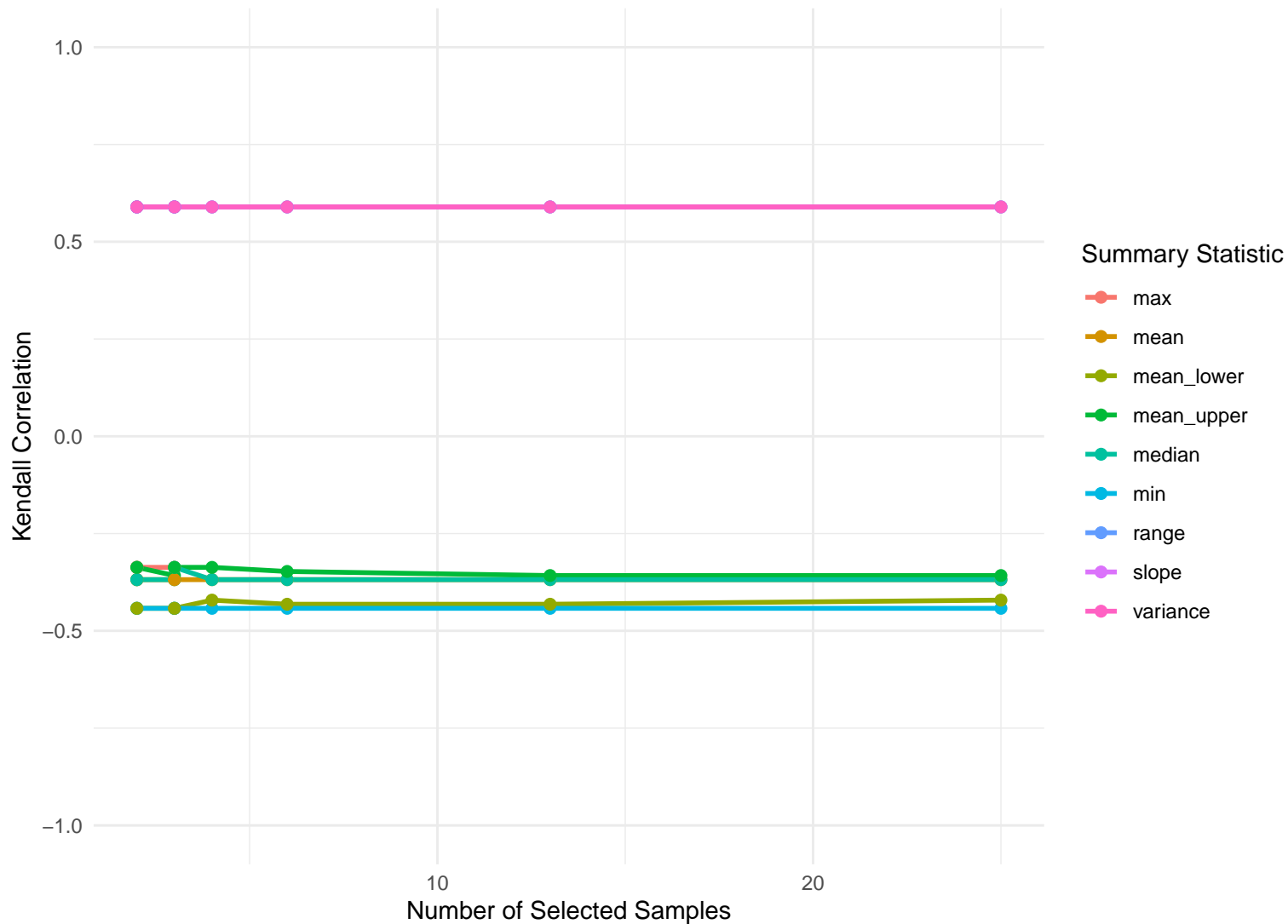




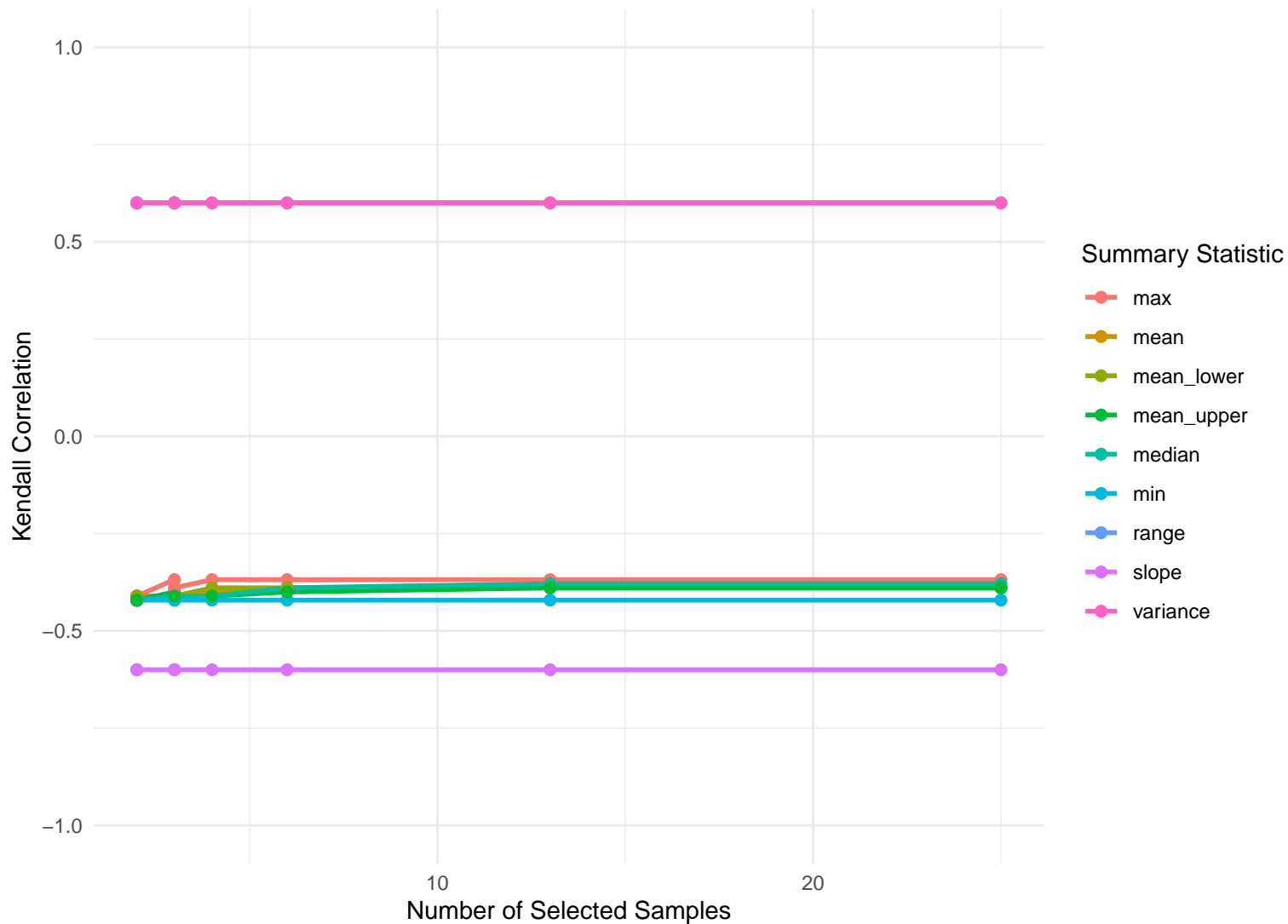
Score: Plmd (Local, partial1 , Genotype: linear )



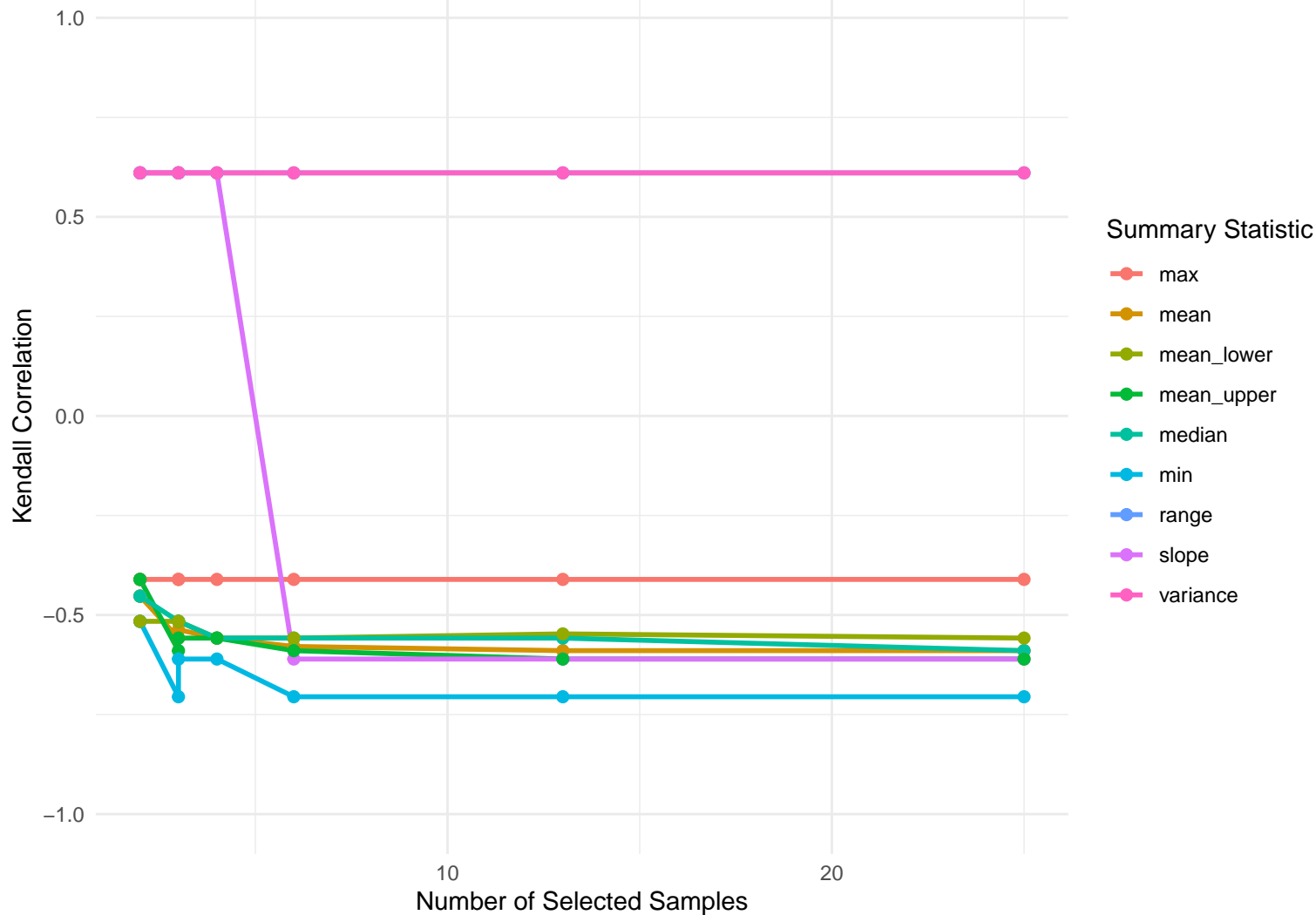
Score: PImd (Local, partial1 , Genotype: gaussian )



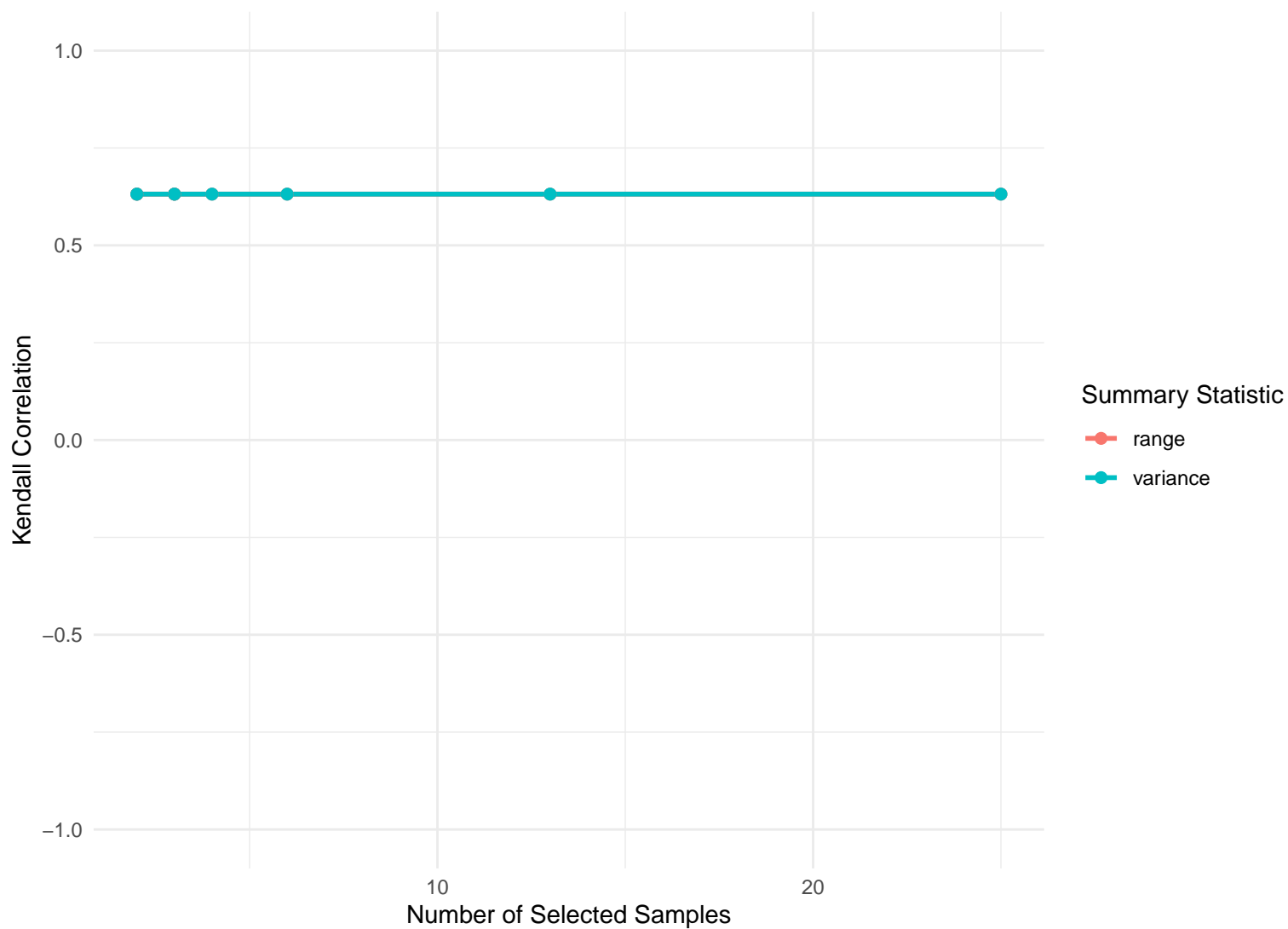
Score: Plmd (Local, partial1 , Genotype: sinusoidal )



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

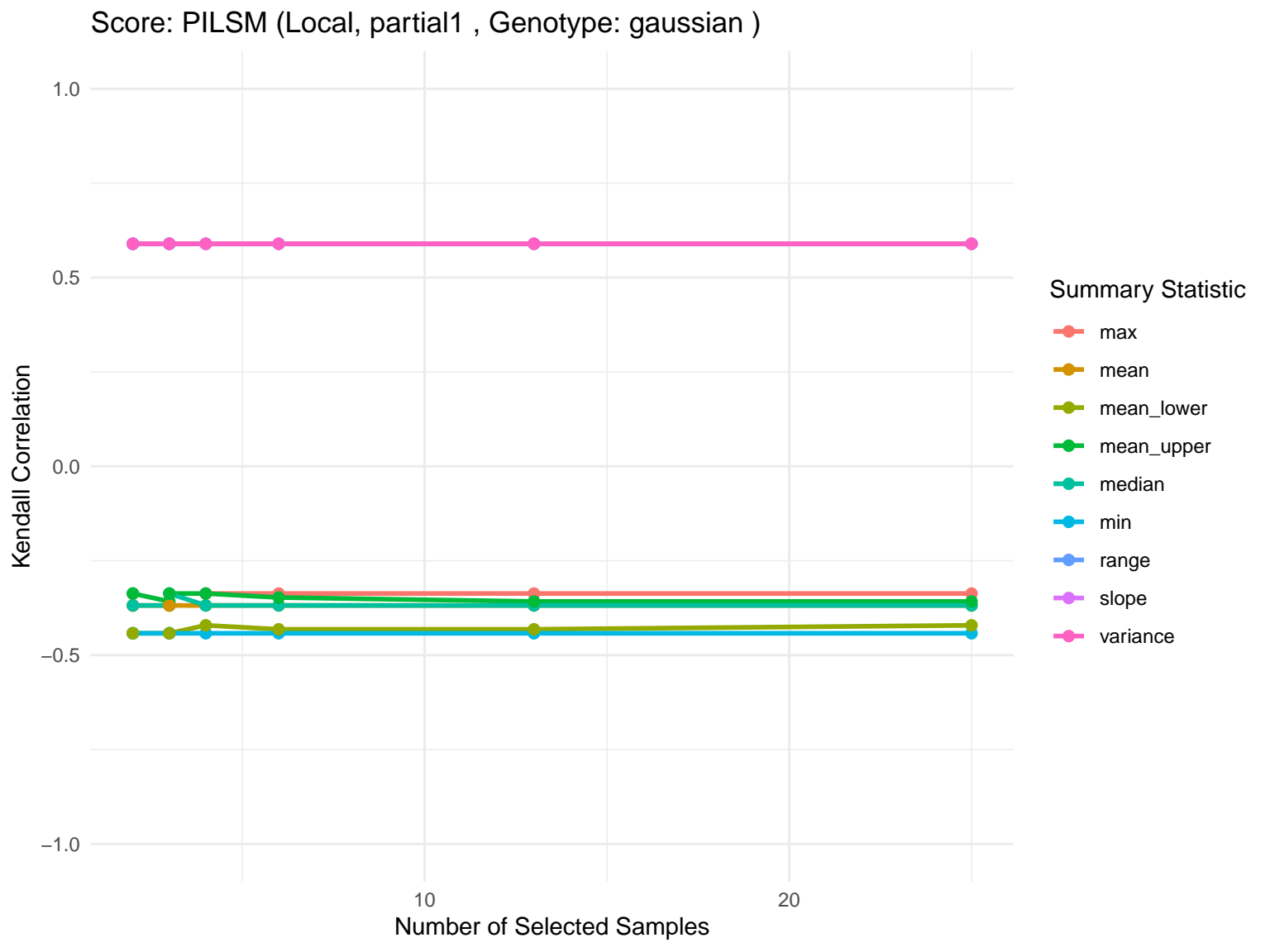
-0.5

-1.0

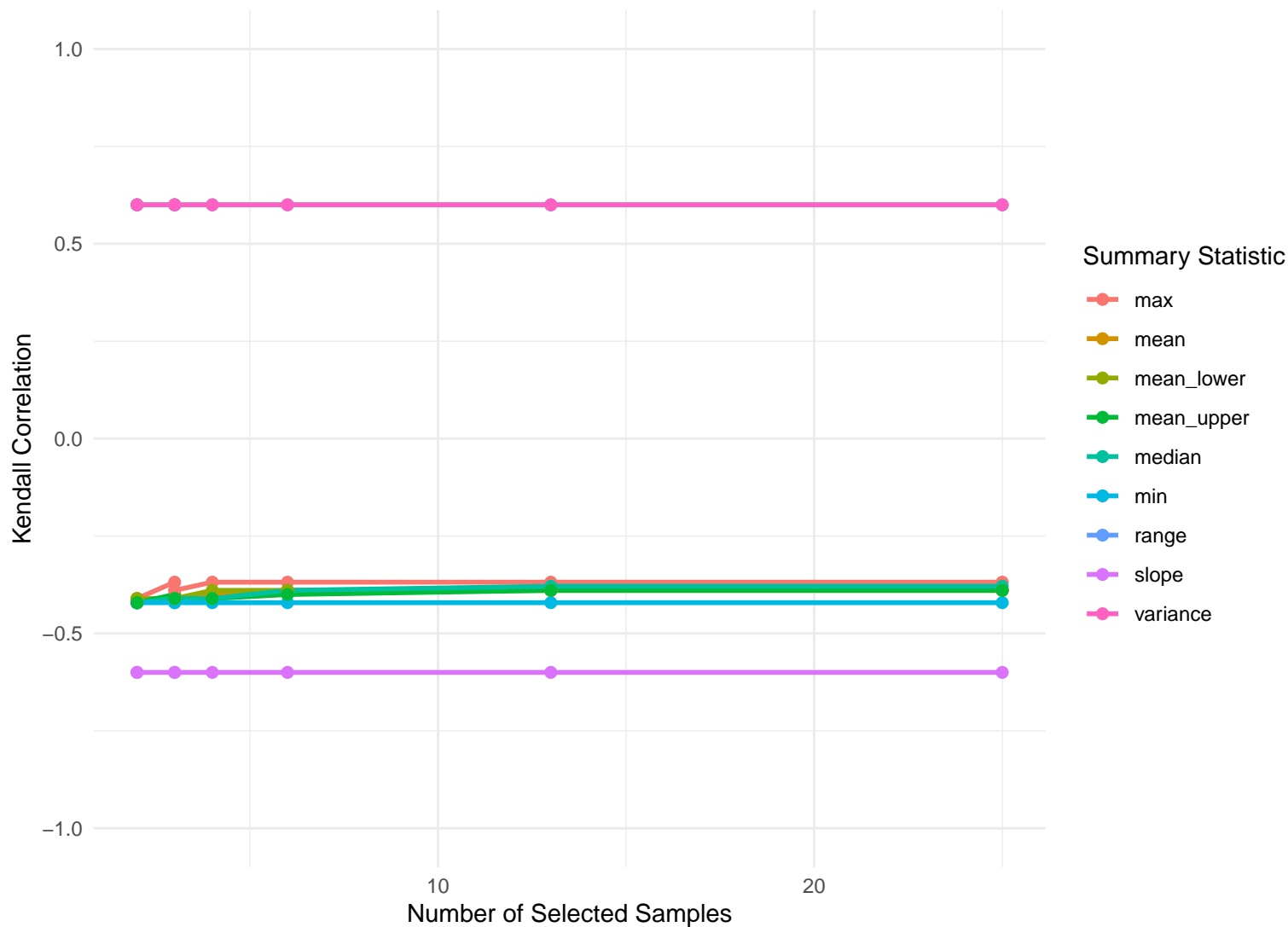
Number of Selected Samples

10

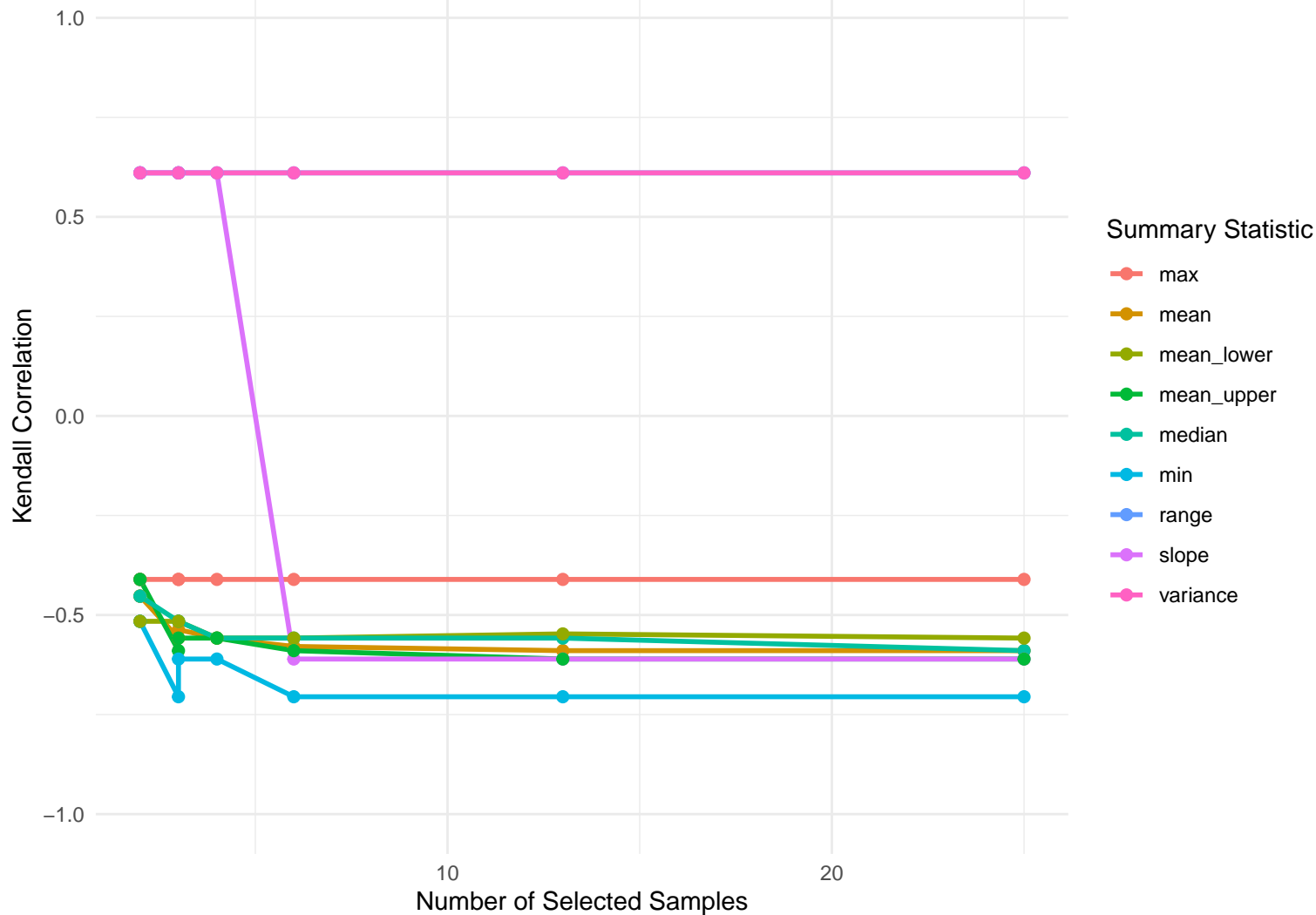
20



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

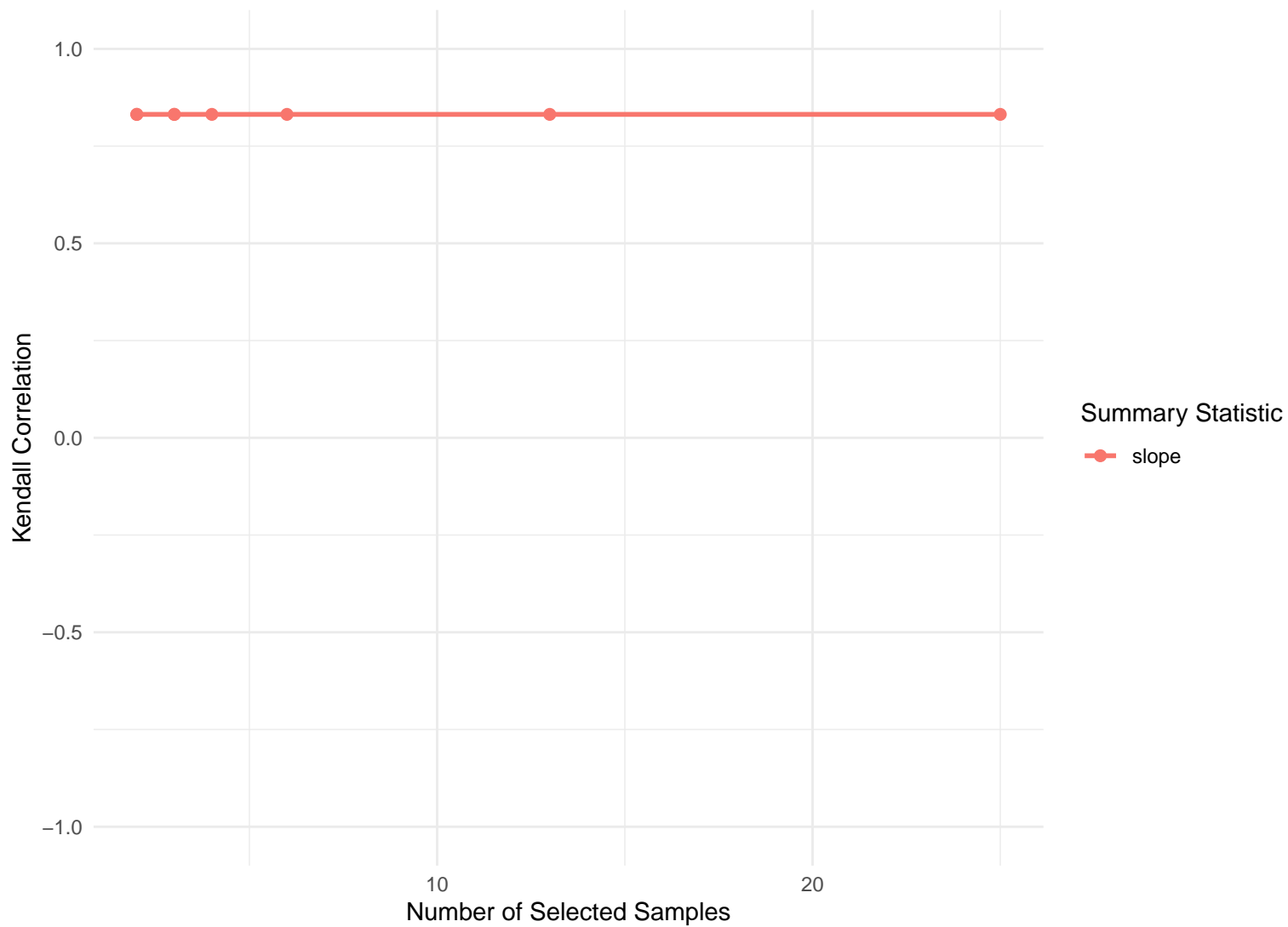


Score: PILSM (Local, partial1 , Genotype: wave )





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

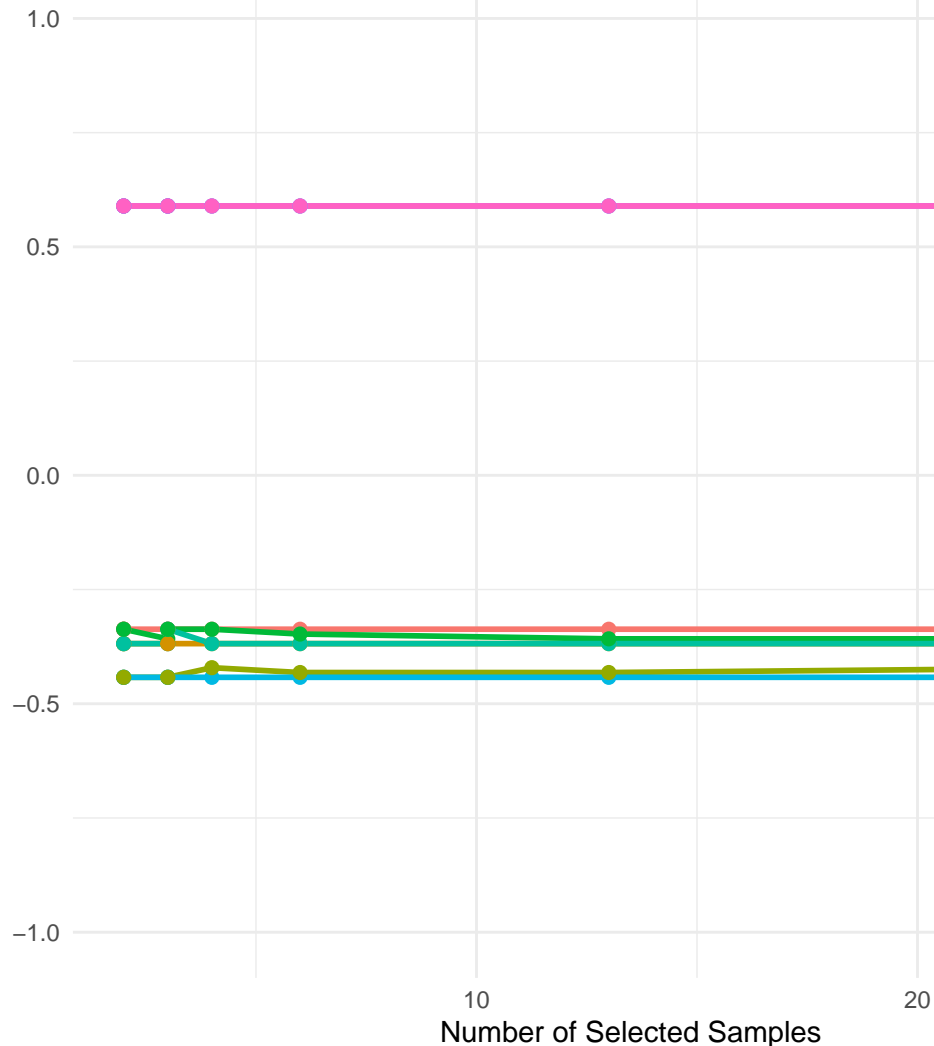


Score: RTR (Local, partial1 , Genotype: gaussian )

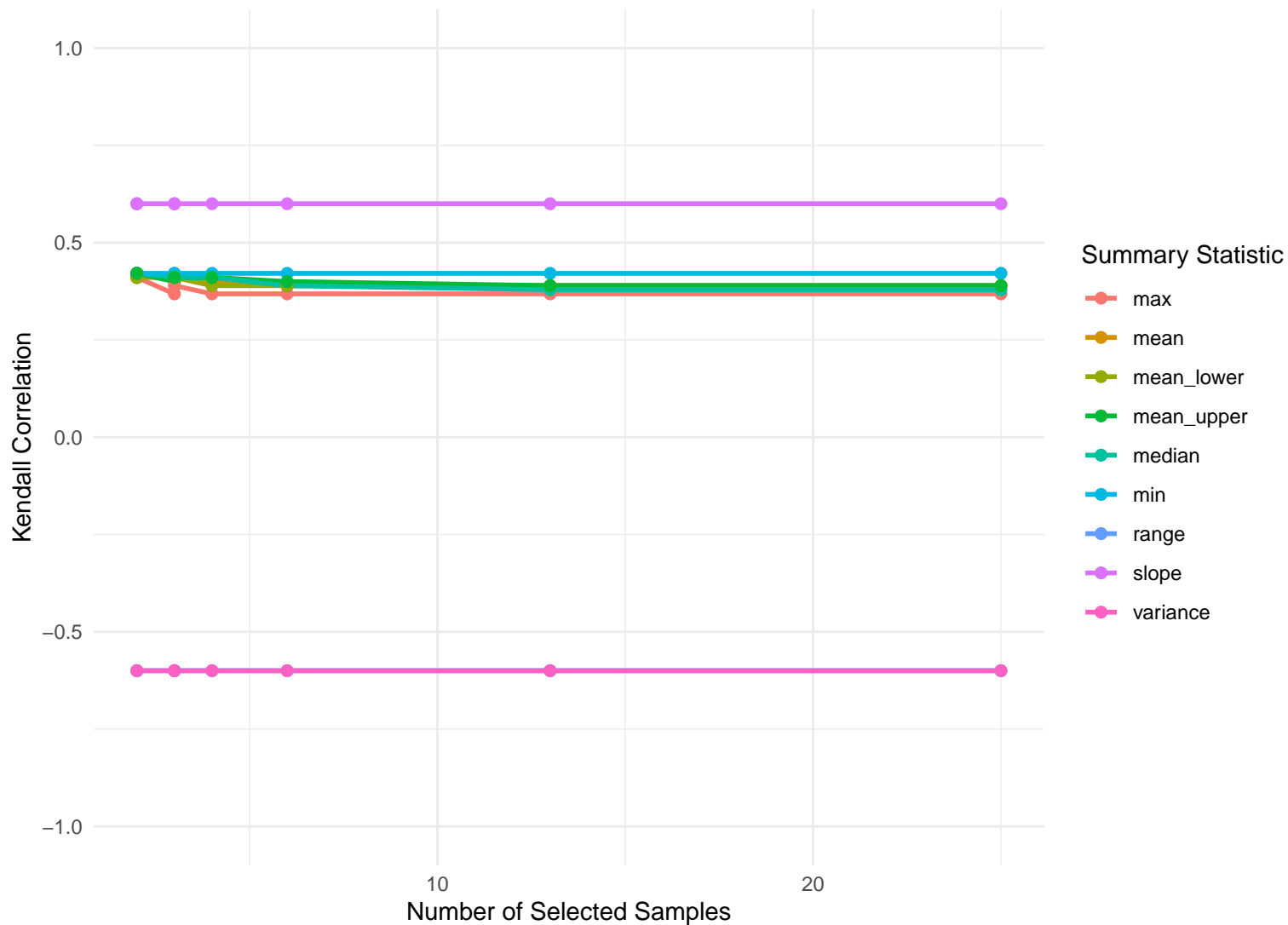
Kendall Correlation

Summary Statistic

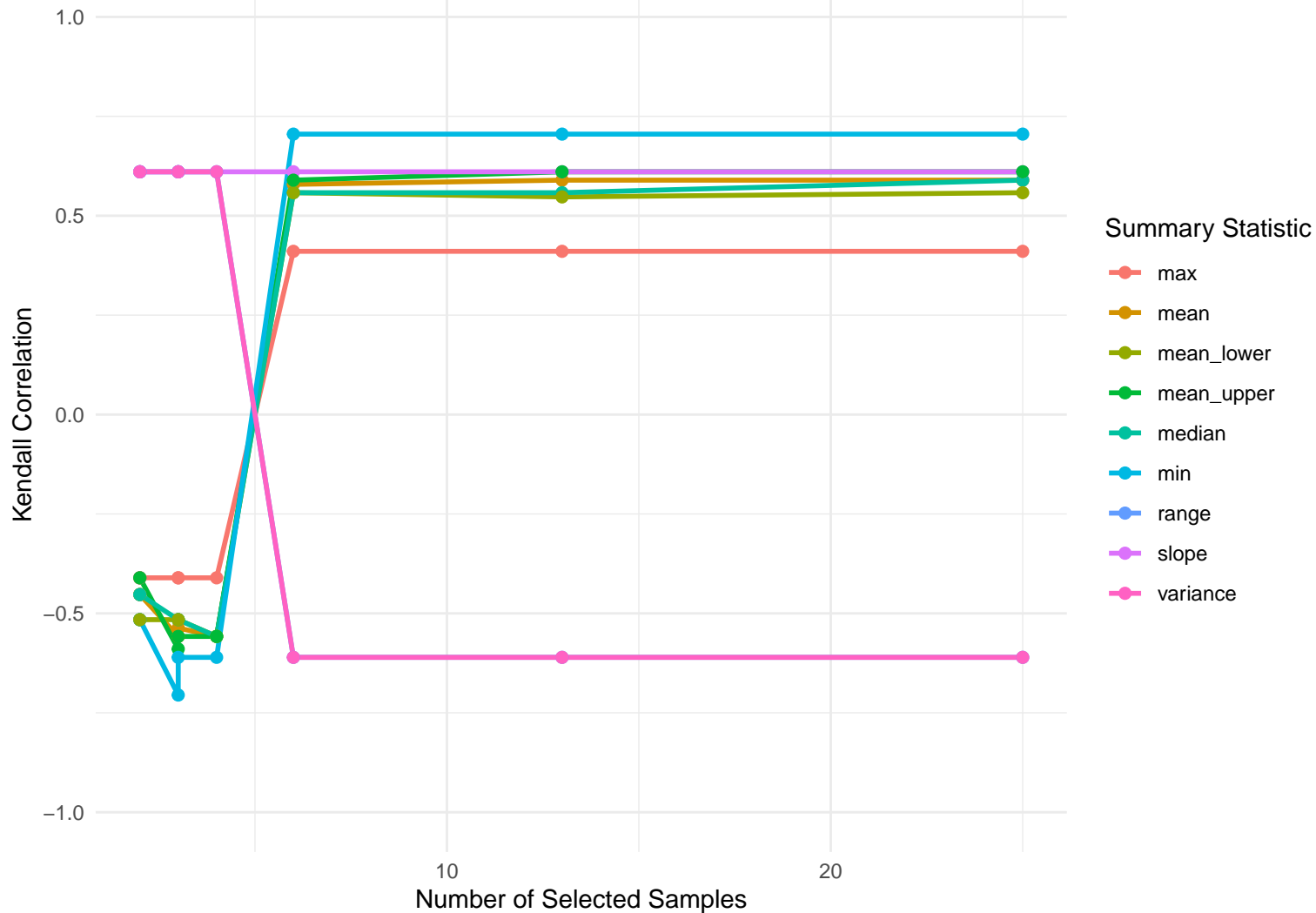
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance



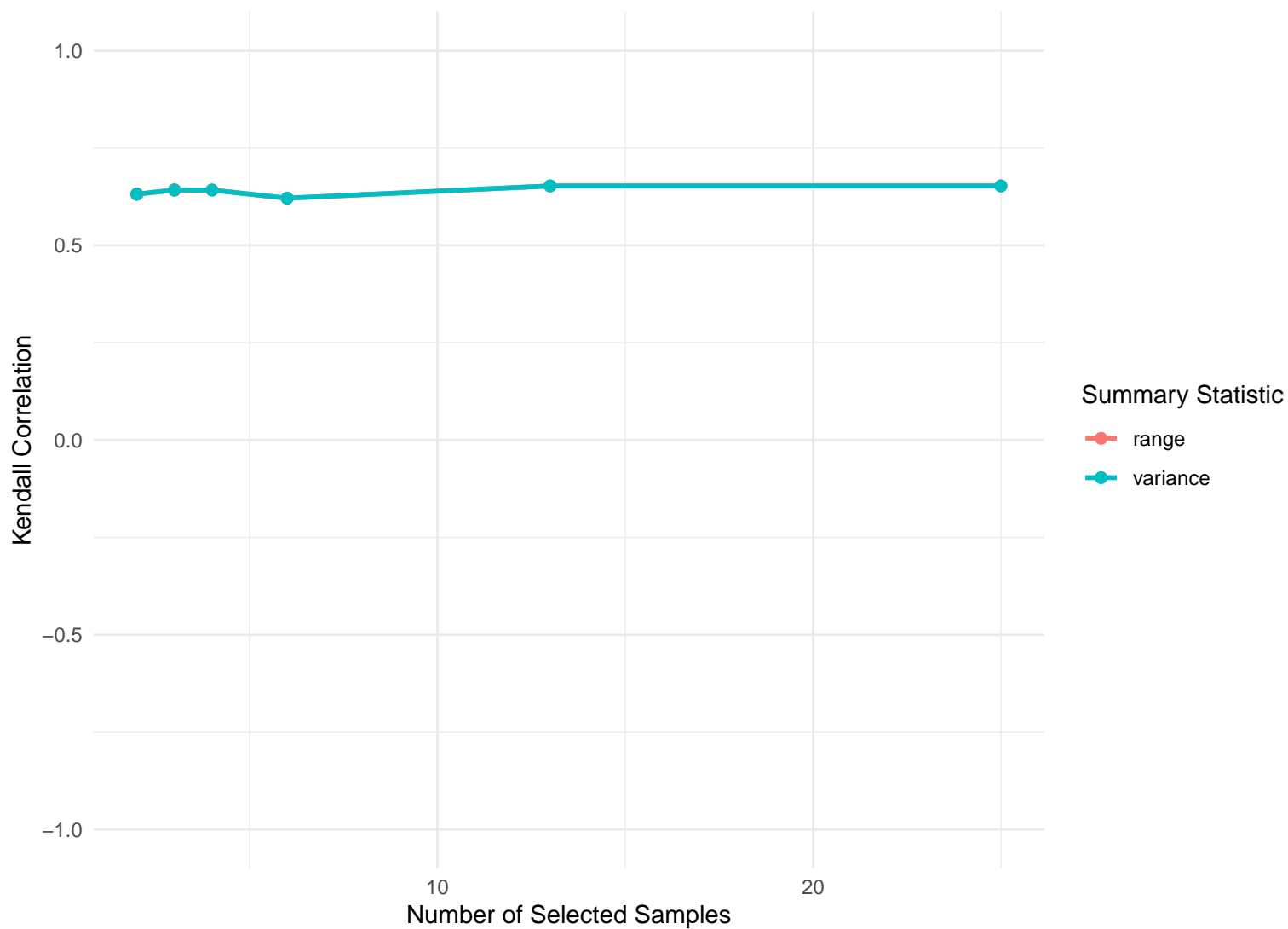
Score: RTR (Local, partial1 , Genotype: sinusoidal )



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



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

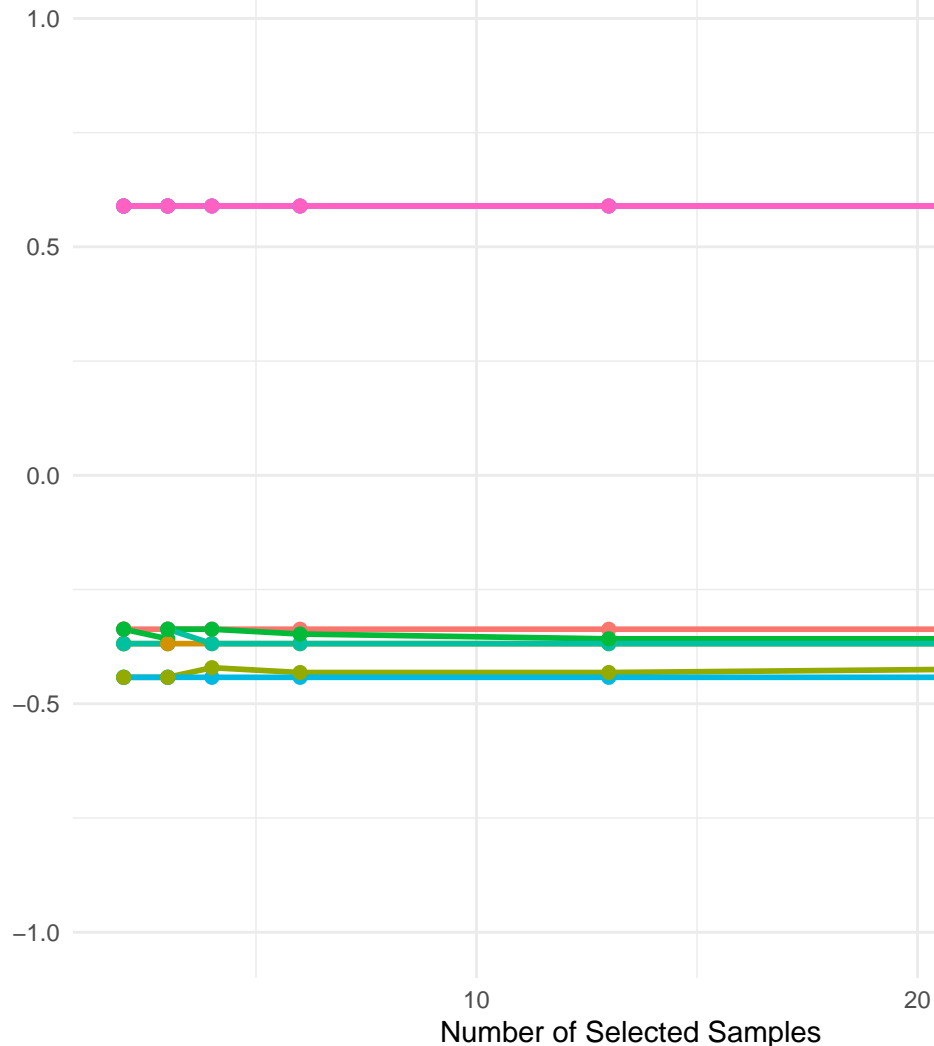


Score: PIR (Local, partial1 , Genotype: gaussian )

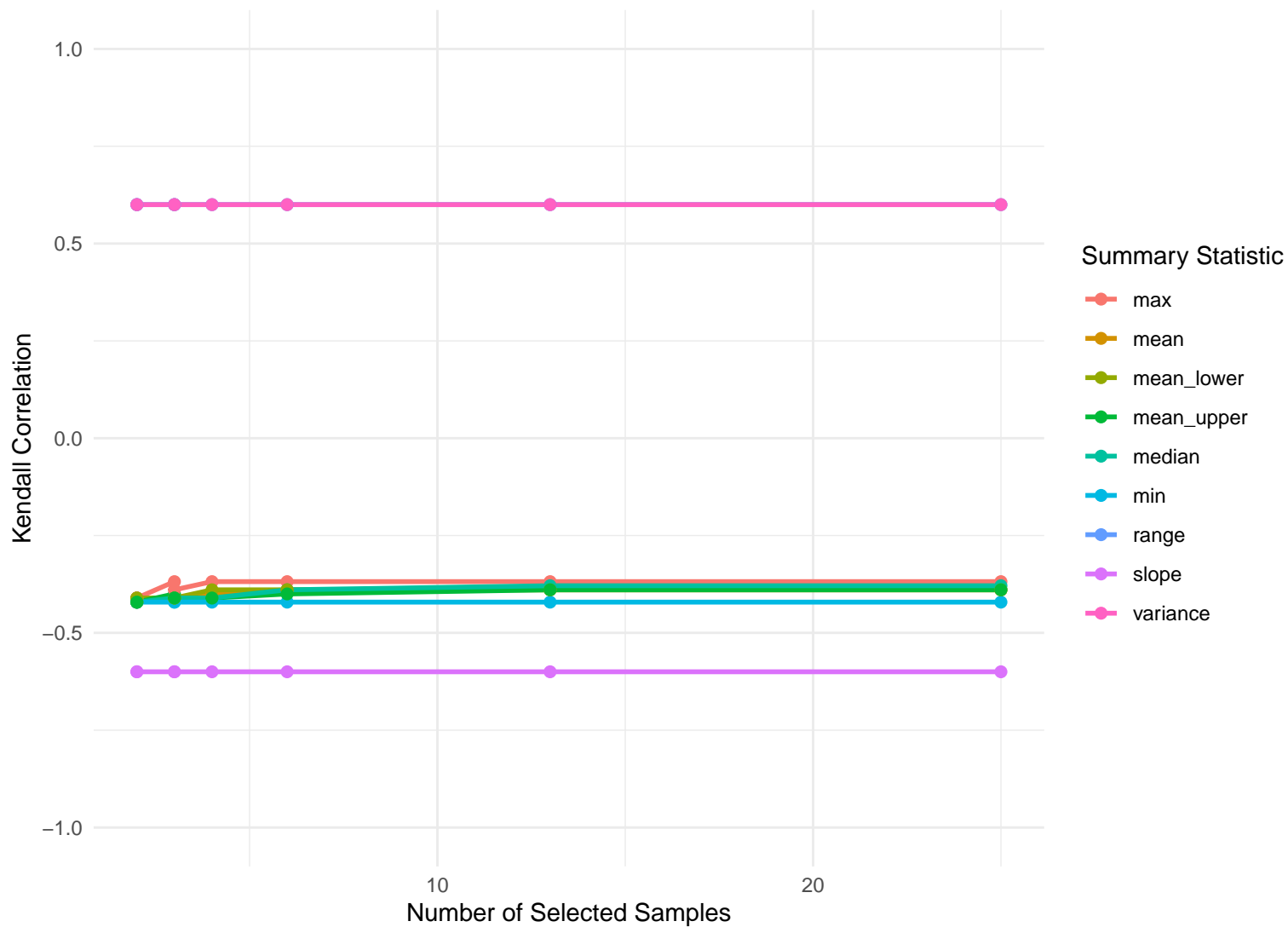
Kendall Correlation

Summary Statistic

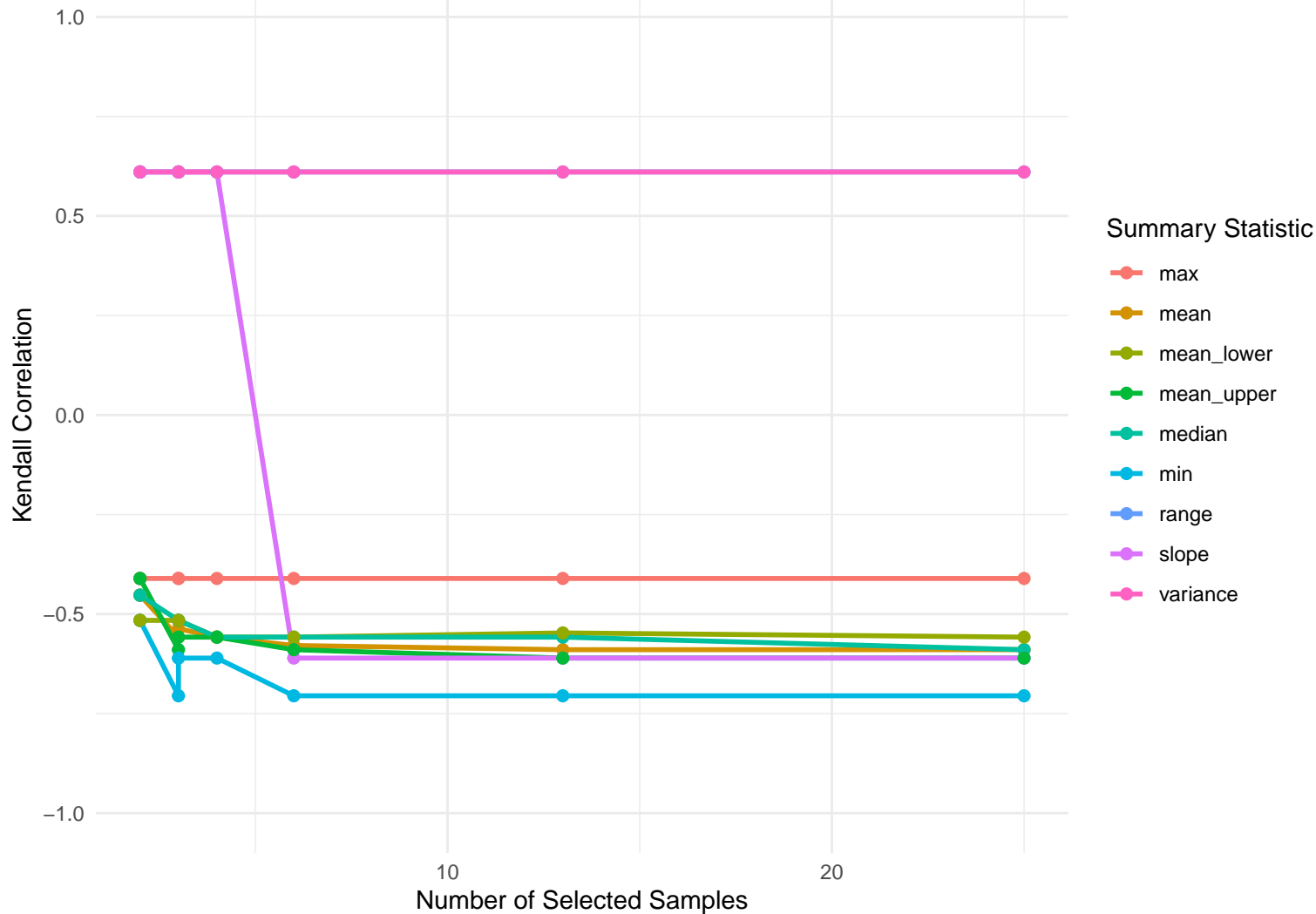
- max
- mean
- mean\_lower
- mean\_upper
- median
- min
- range
- slope
- variance



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

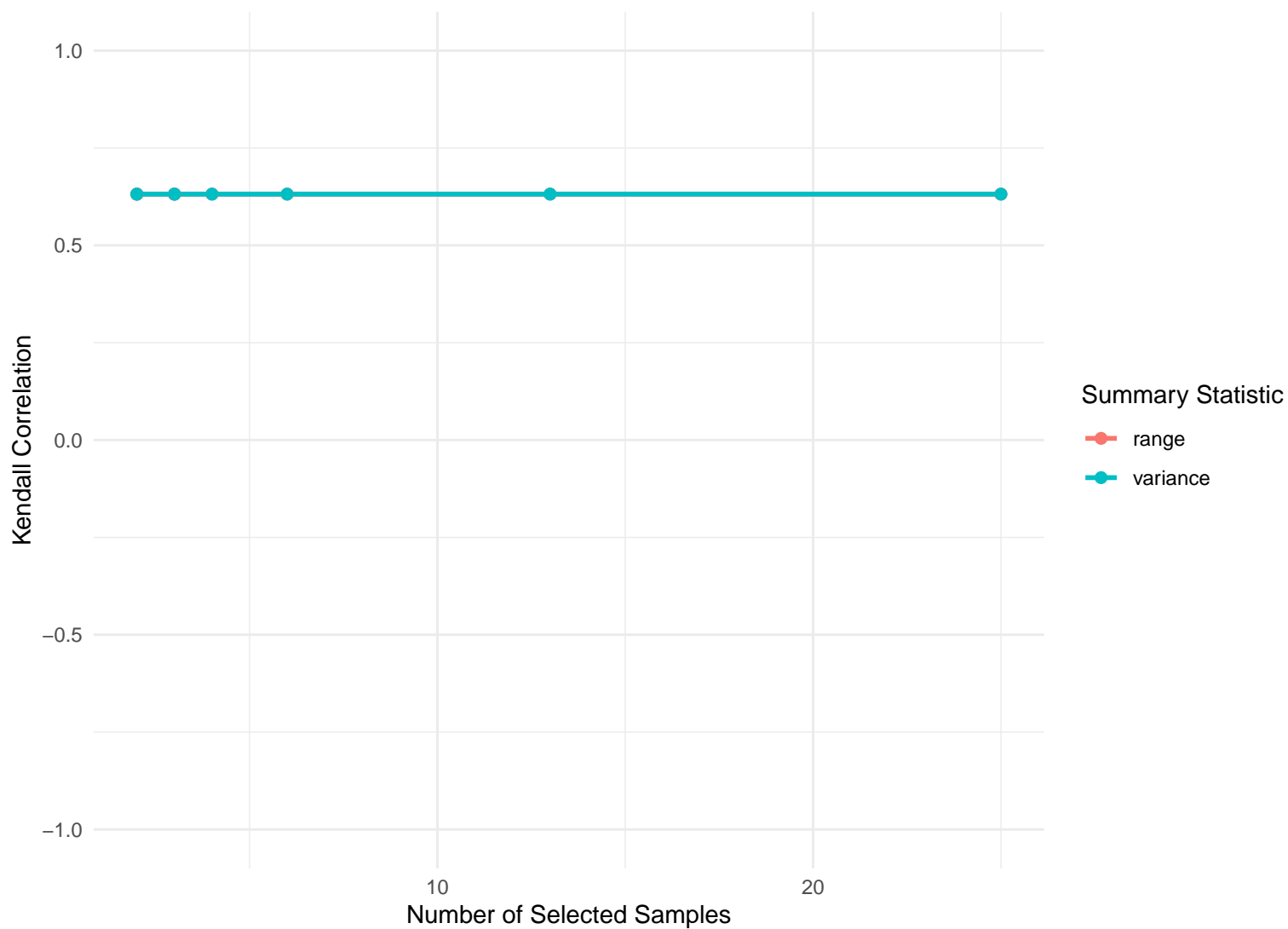


Score: PIR (Local, partial1 , Genotype: wave )

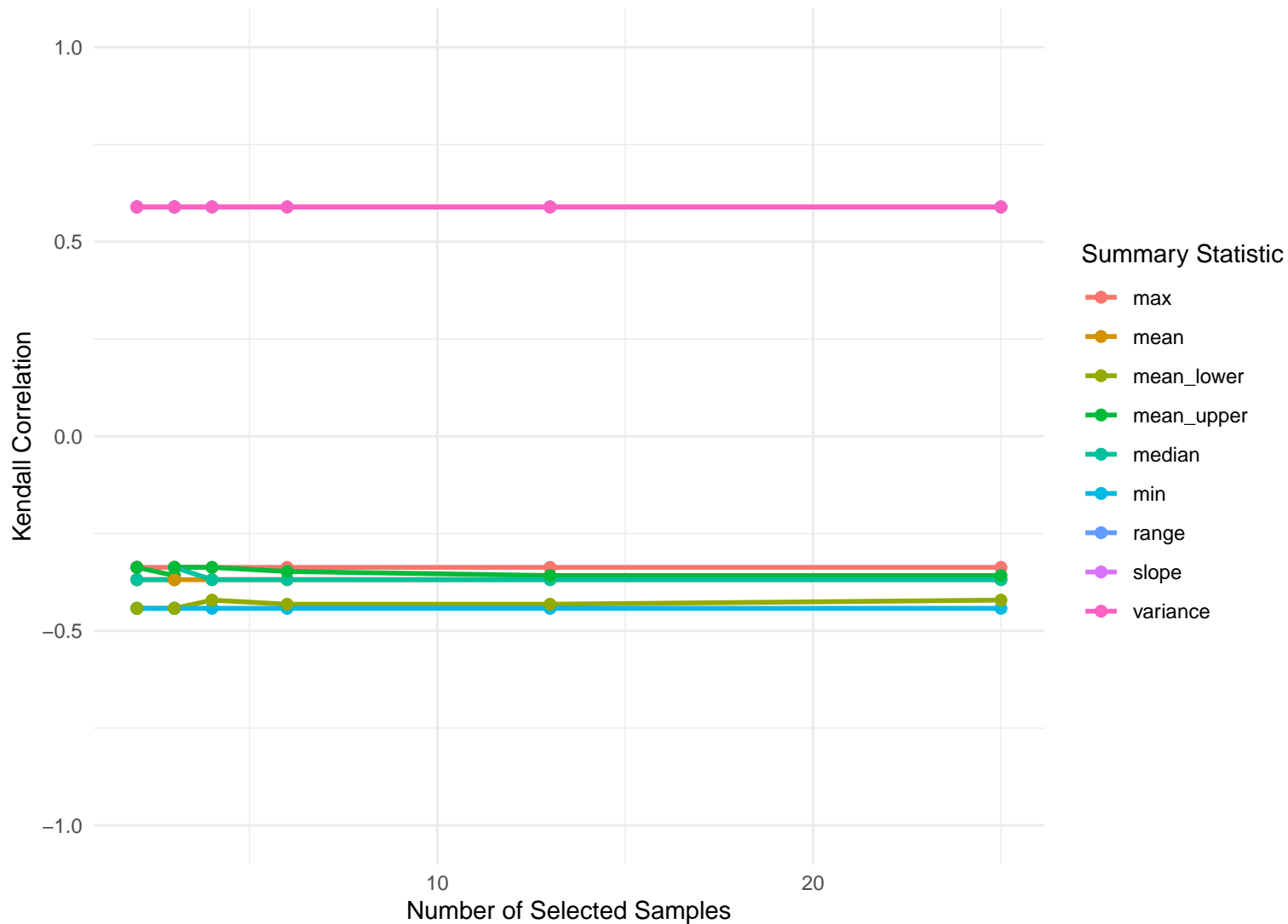




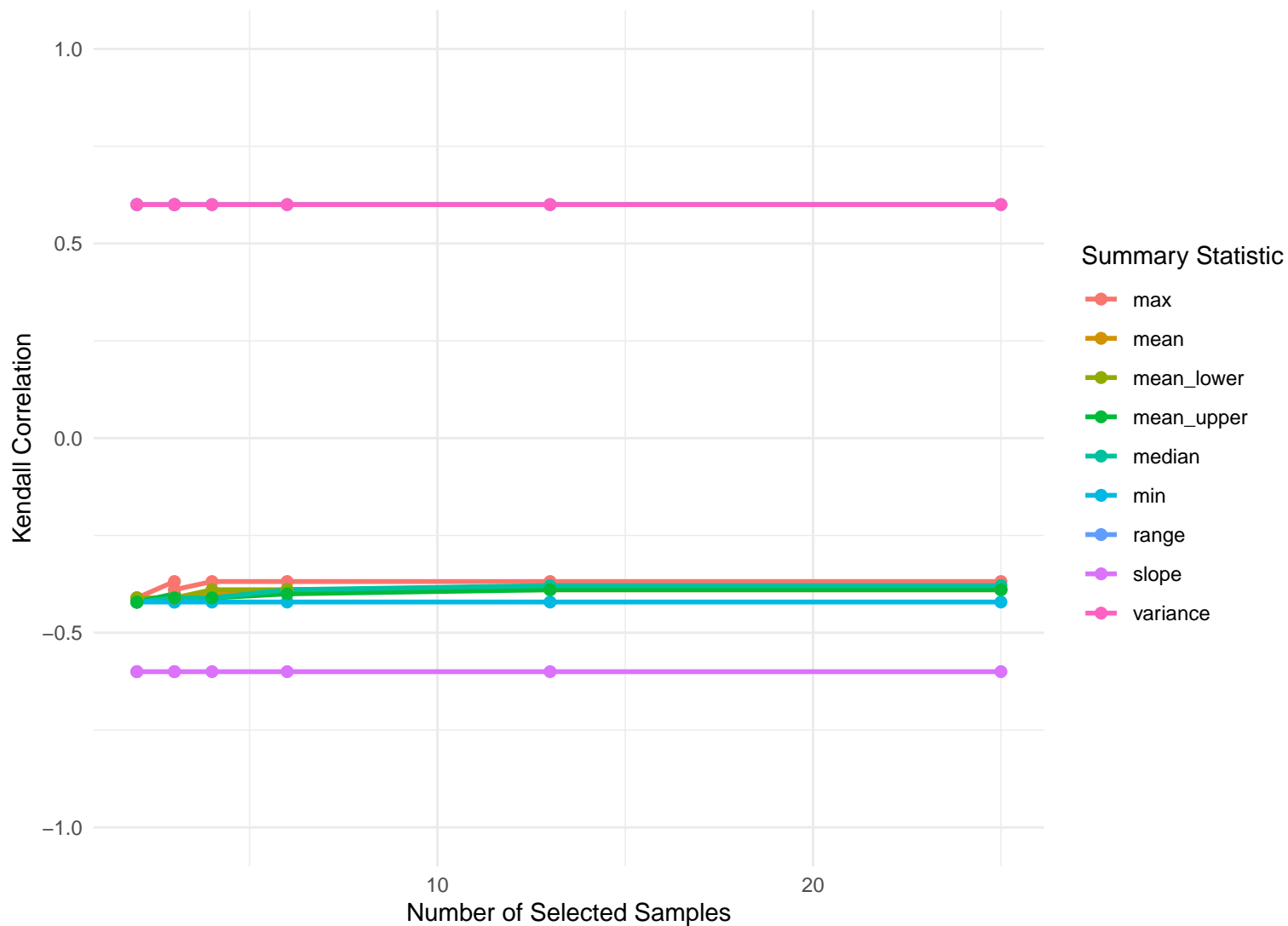
Score: RDPI (Local, partial1 , Genotype: linear )



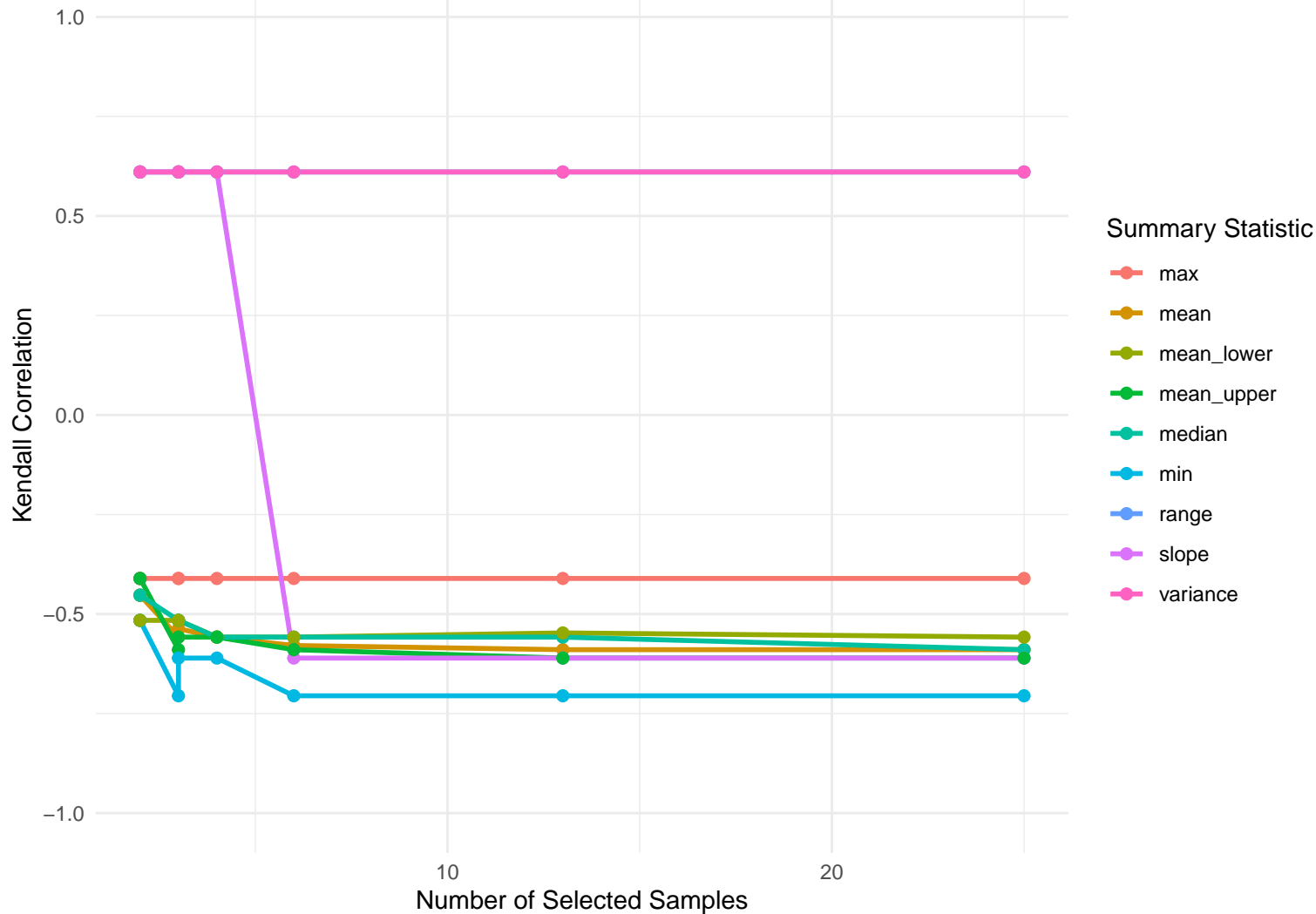
Score: RDPI (Local, partial1 , Genotype: gaussian )



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

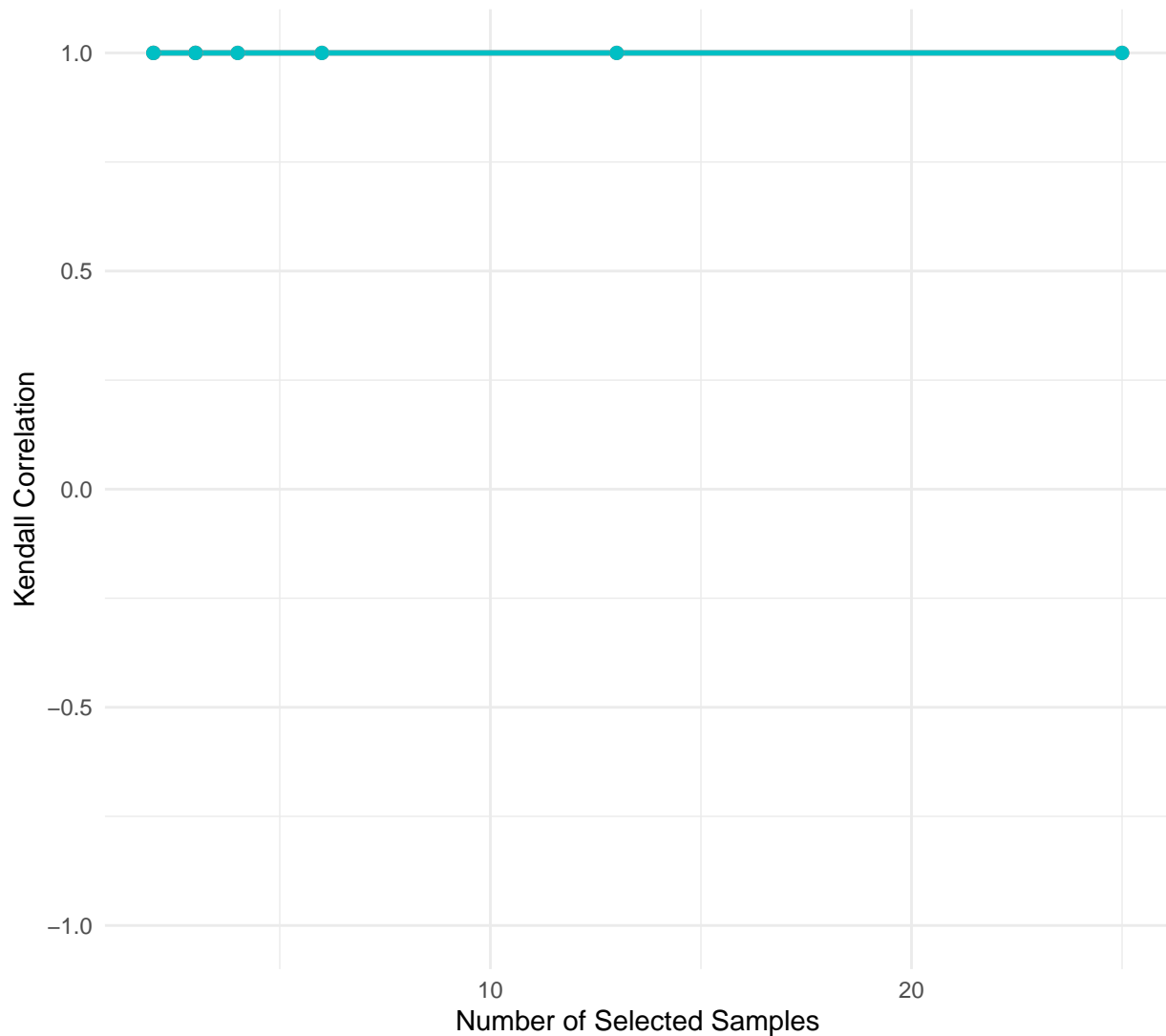
20

Number of Selected Samples

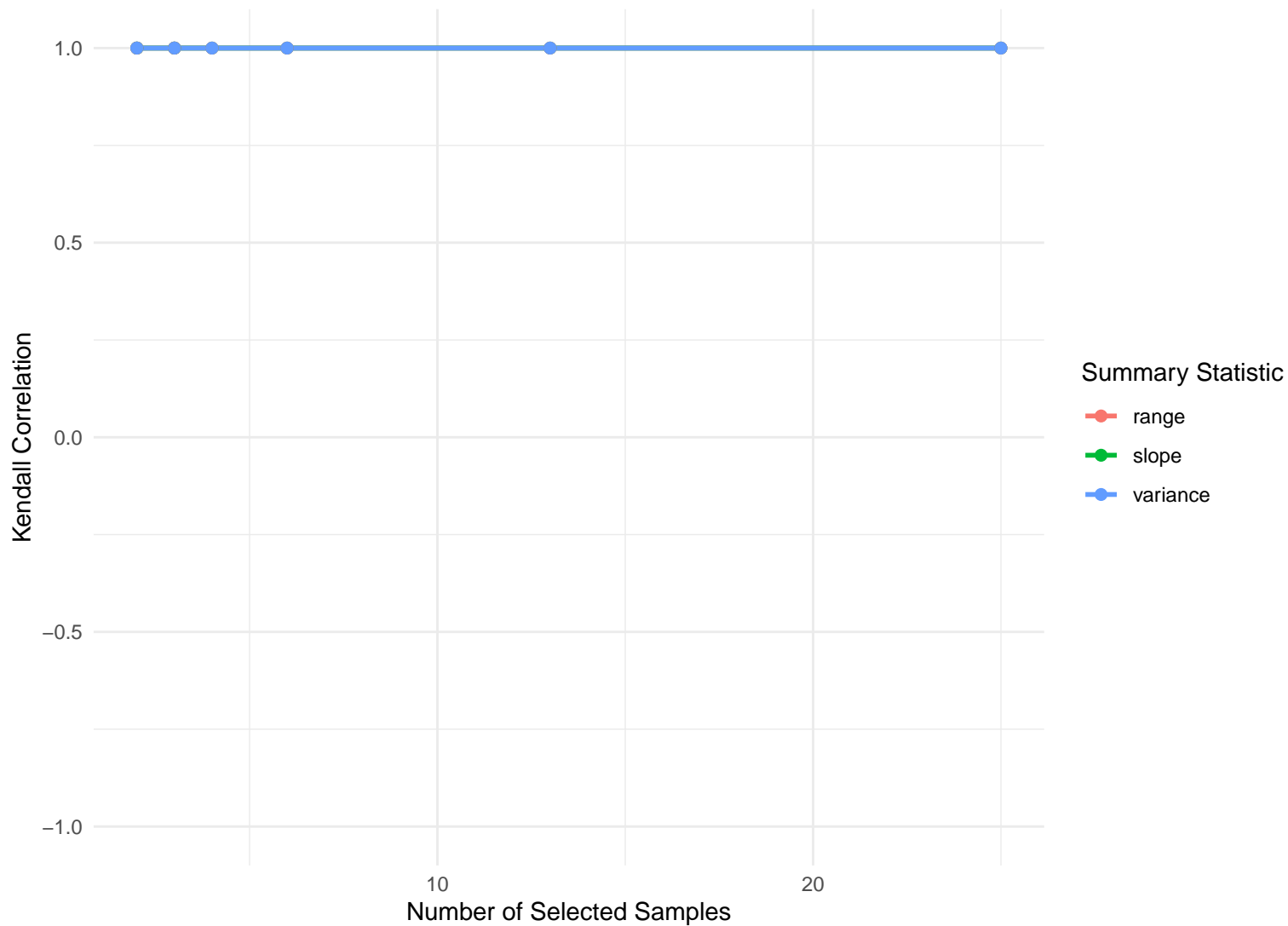
Summary Statistic

range

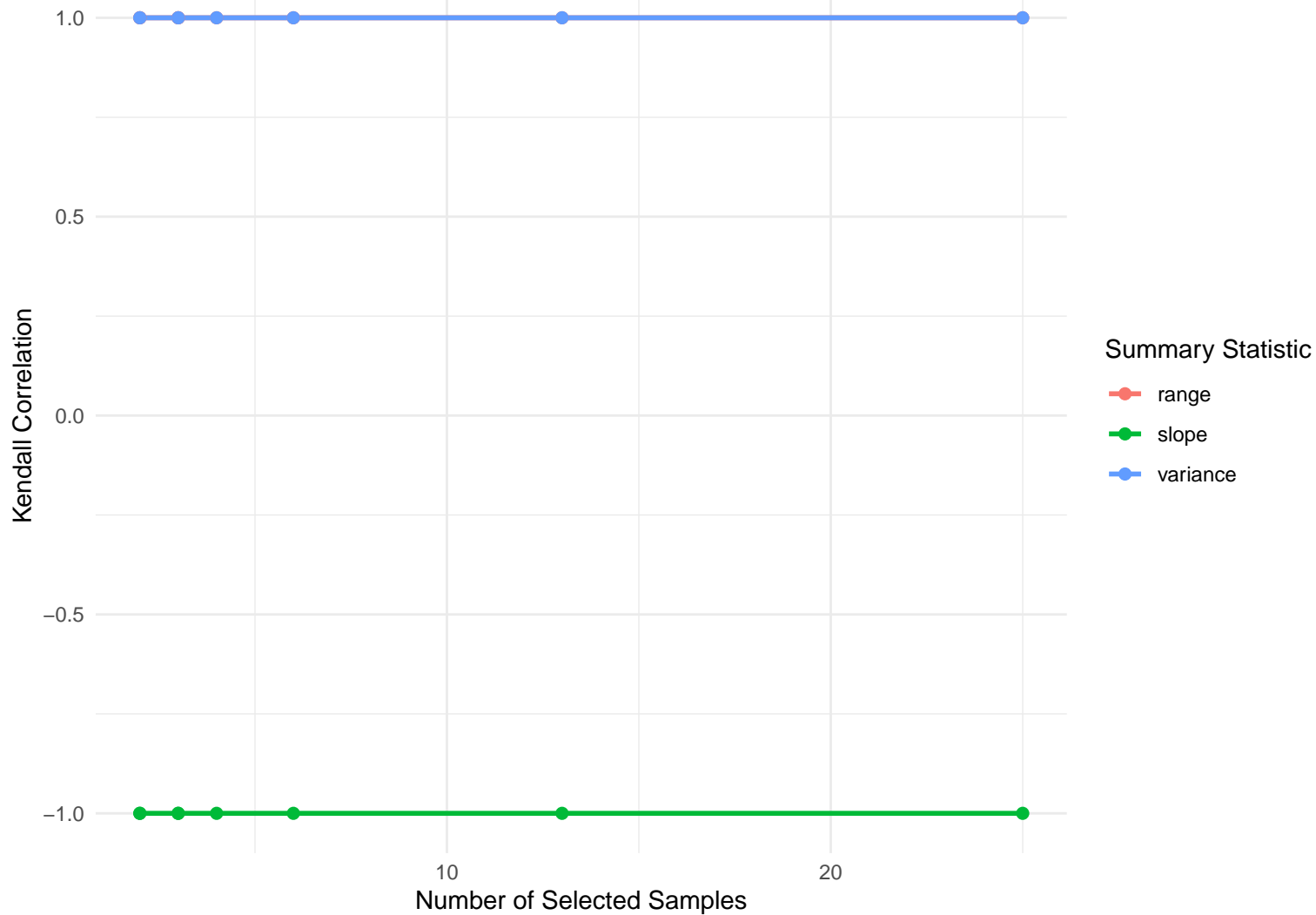
variance



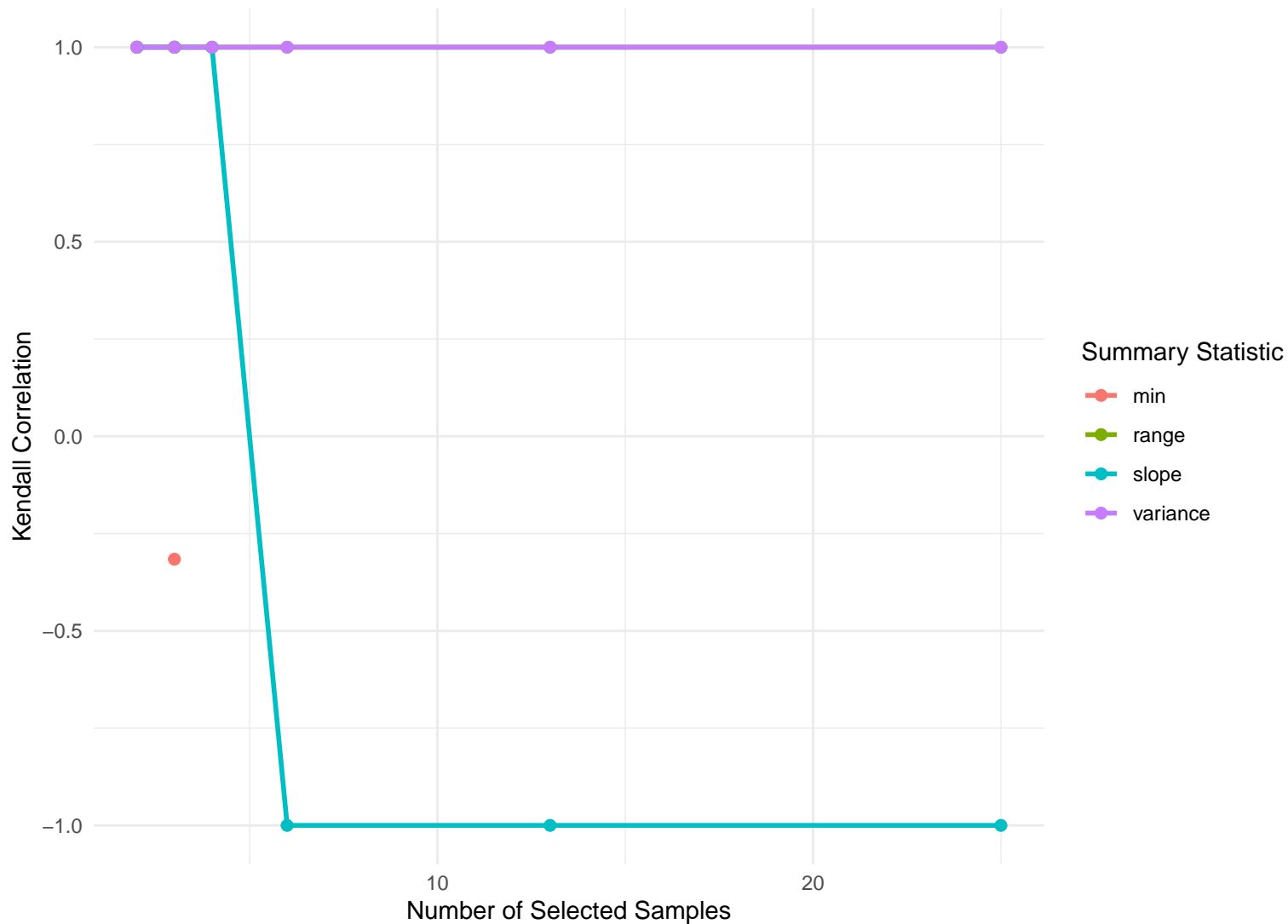
Score: ESPI (Local, partial1 , Genotype: gaussian )



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

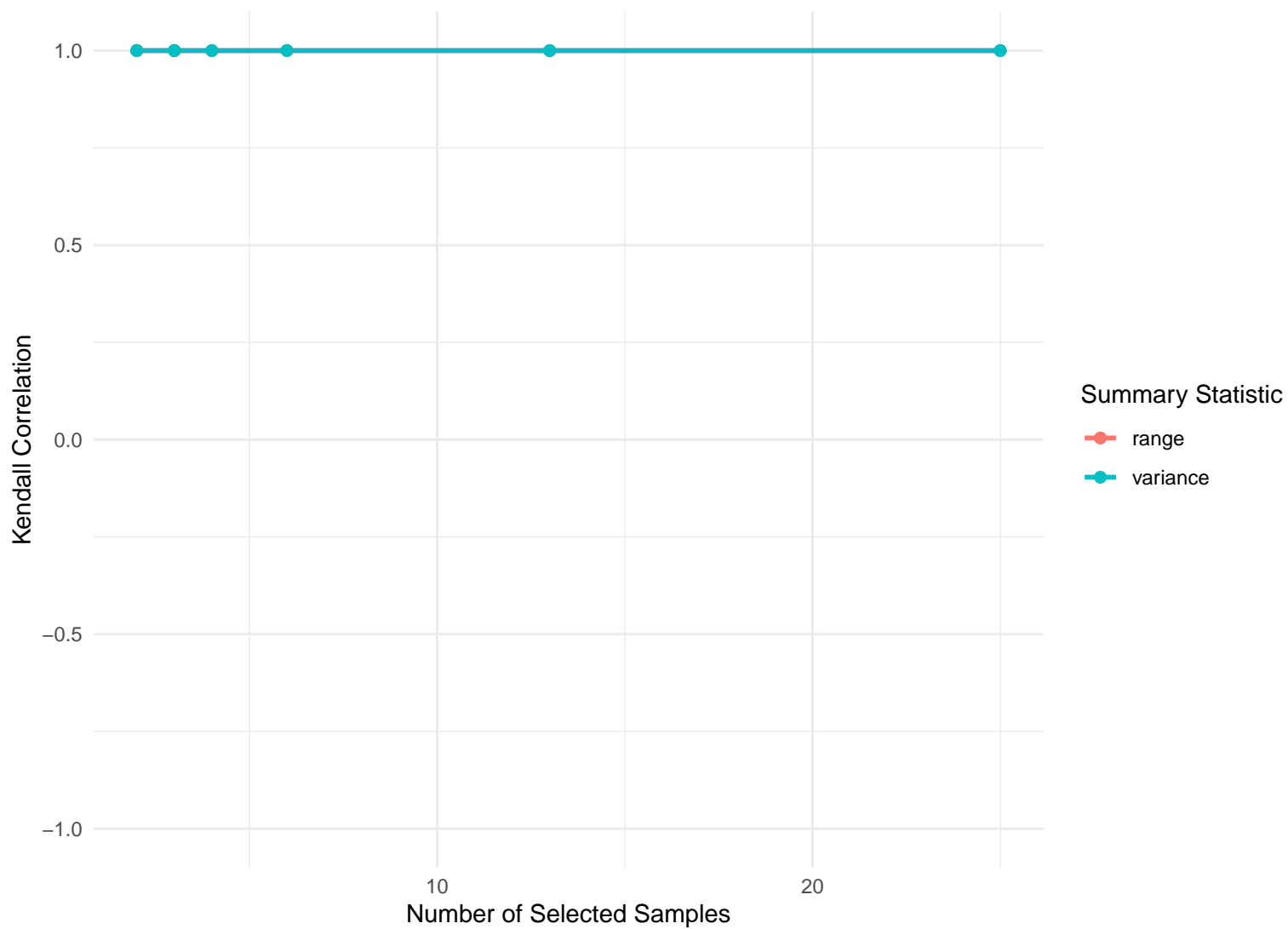


Score: ESPI (Local, partial1 , Genotype: wave )





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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

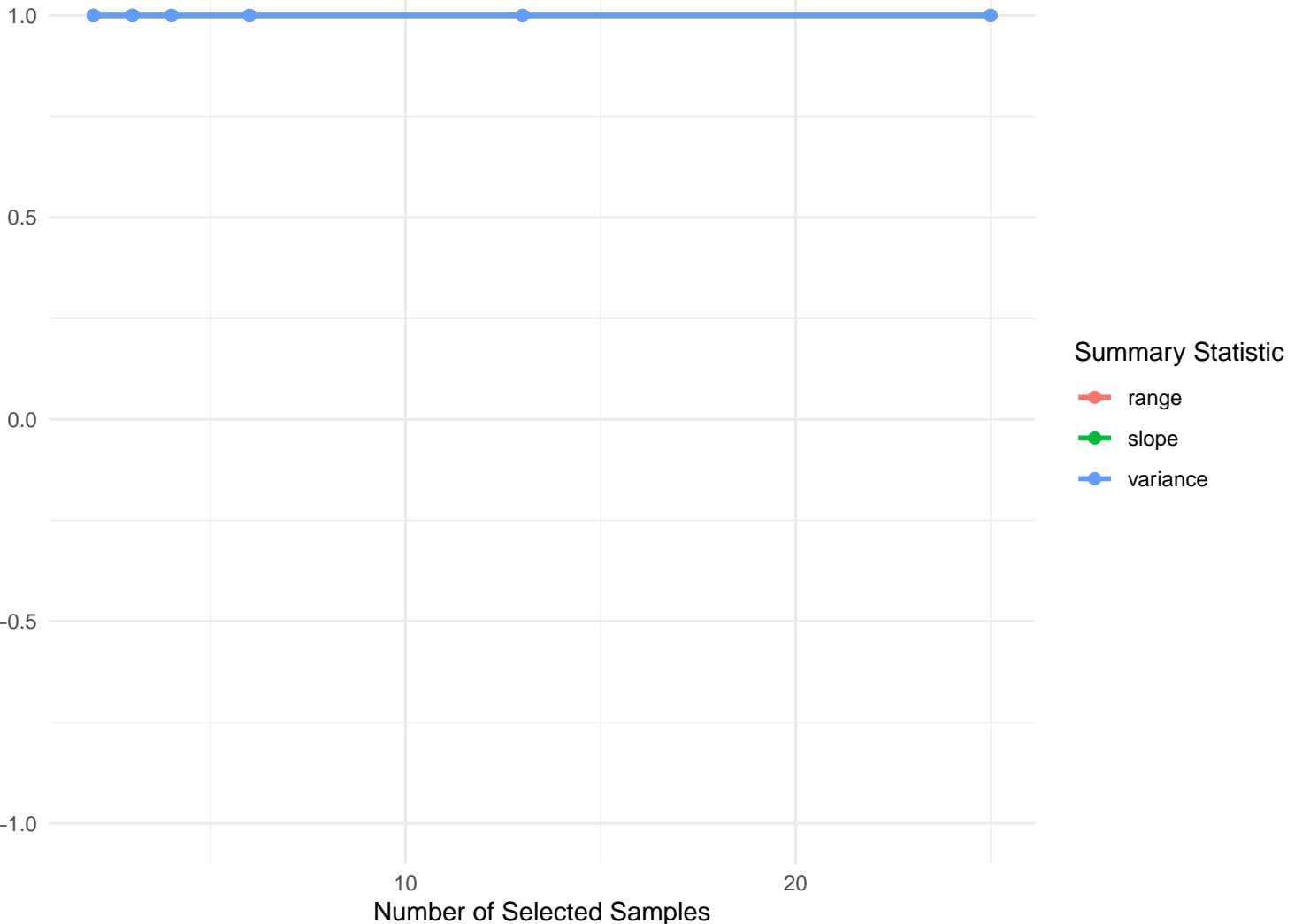
10

20

Number of Selected Samples

Summary Statistic

- range
- slope
- variance



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

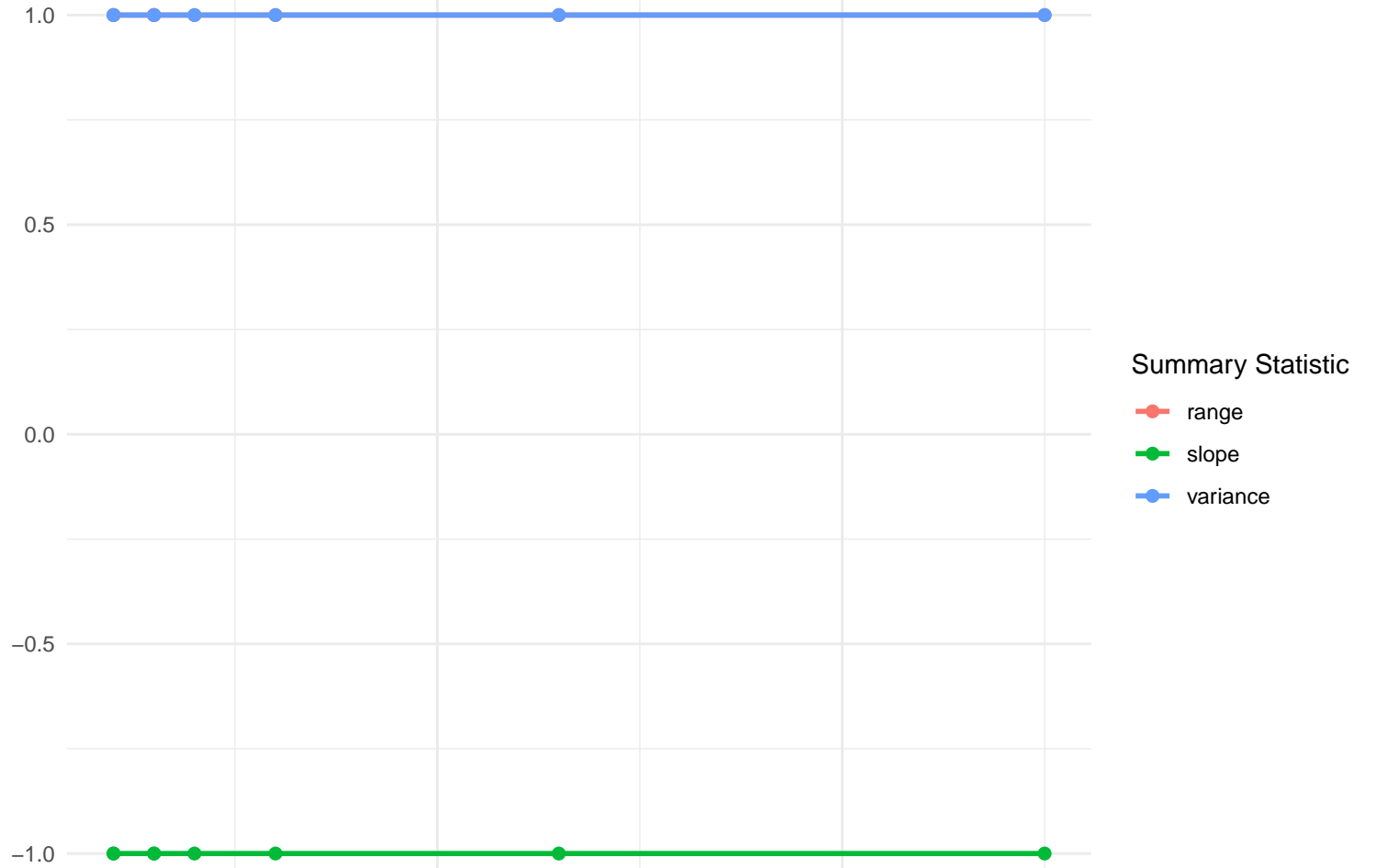
Number of Selected Samples

10

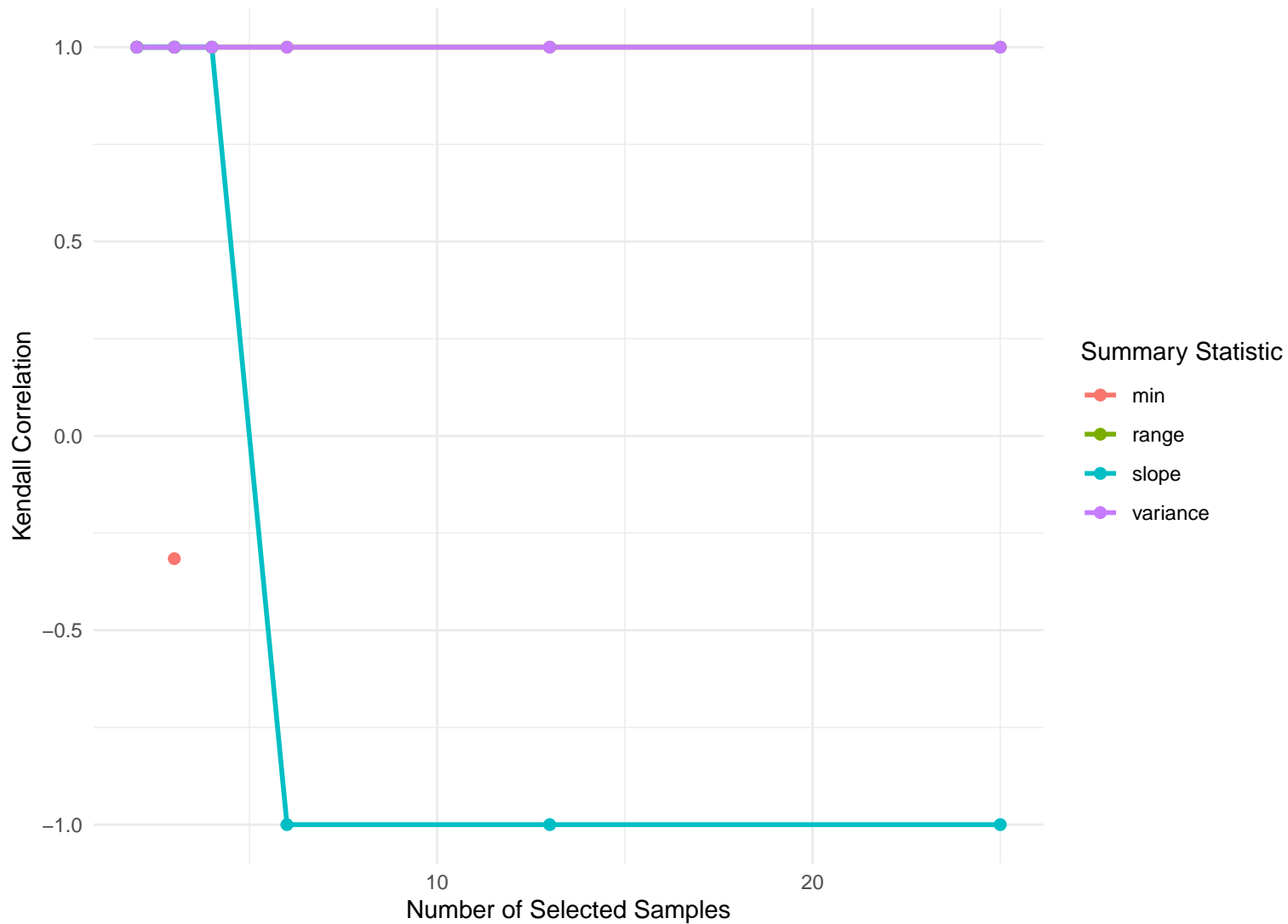
20

Summary Statistic

- range
- slope
- variance



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

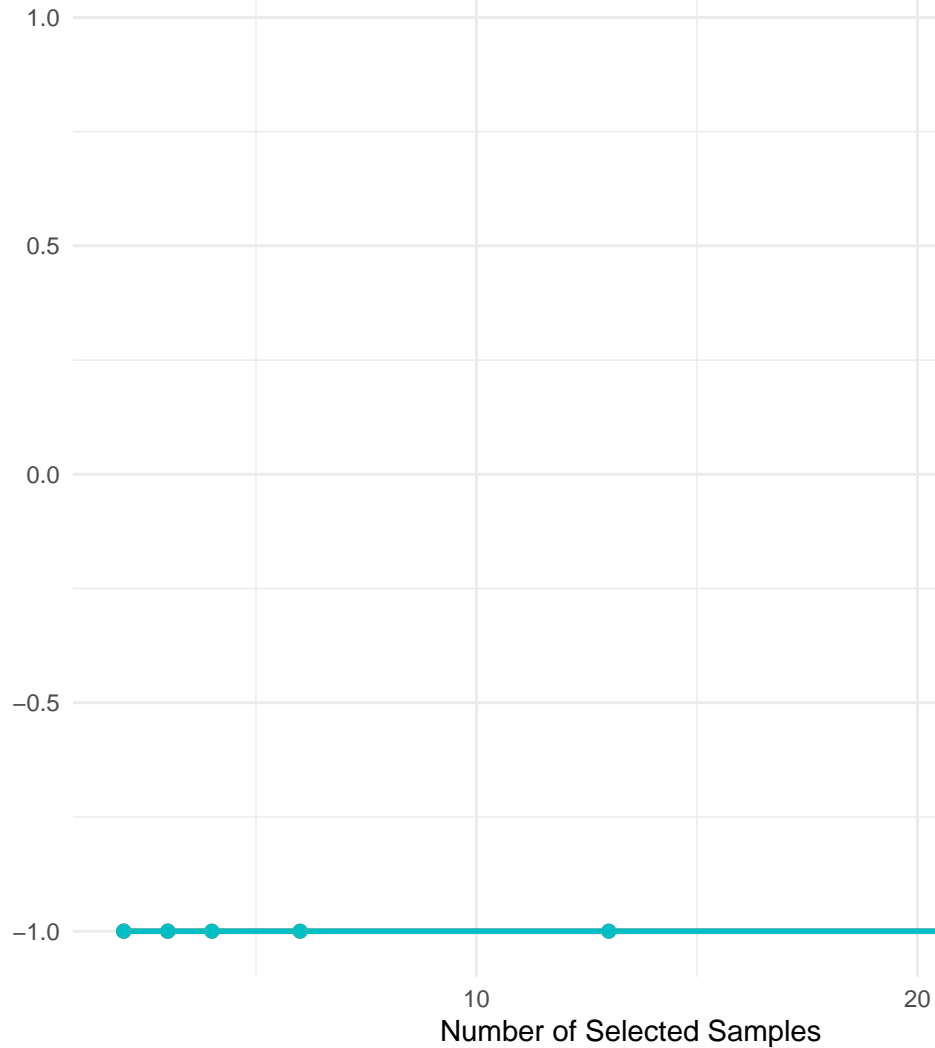


Score: PSI (Local, partial1 , Genotype: linear )

Kendall Correlation

Summary Statistic

- range
- variance

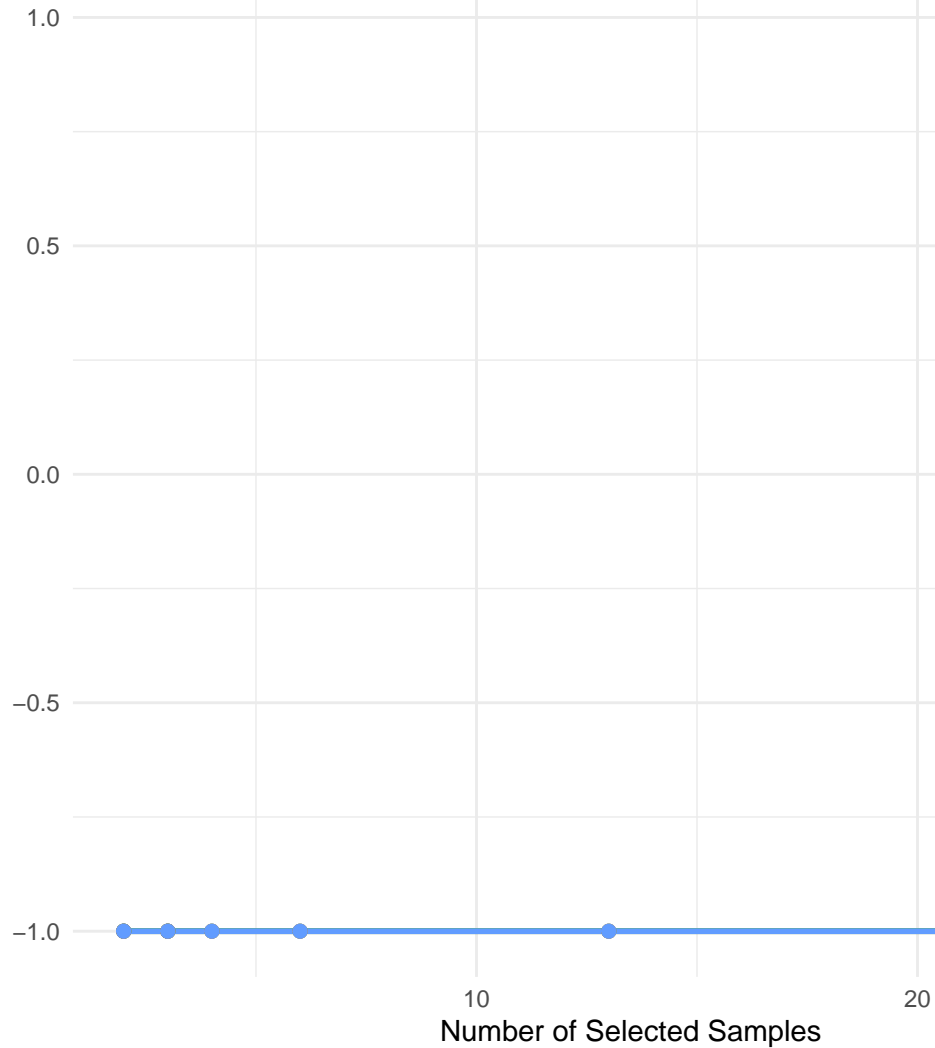


Score: PSI (Local, partial1 , Genotype: gaussian )

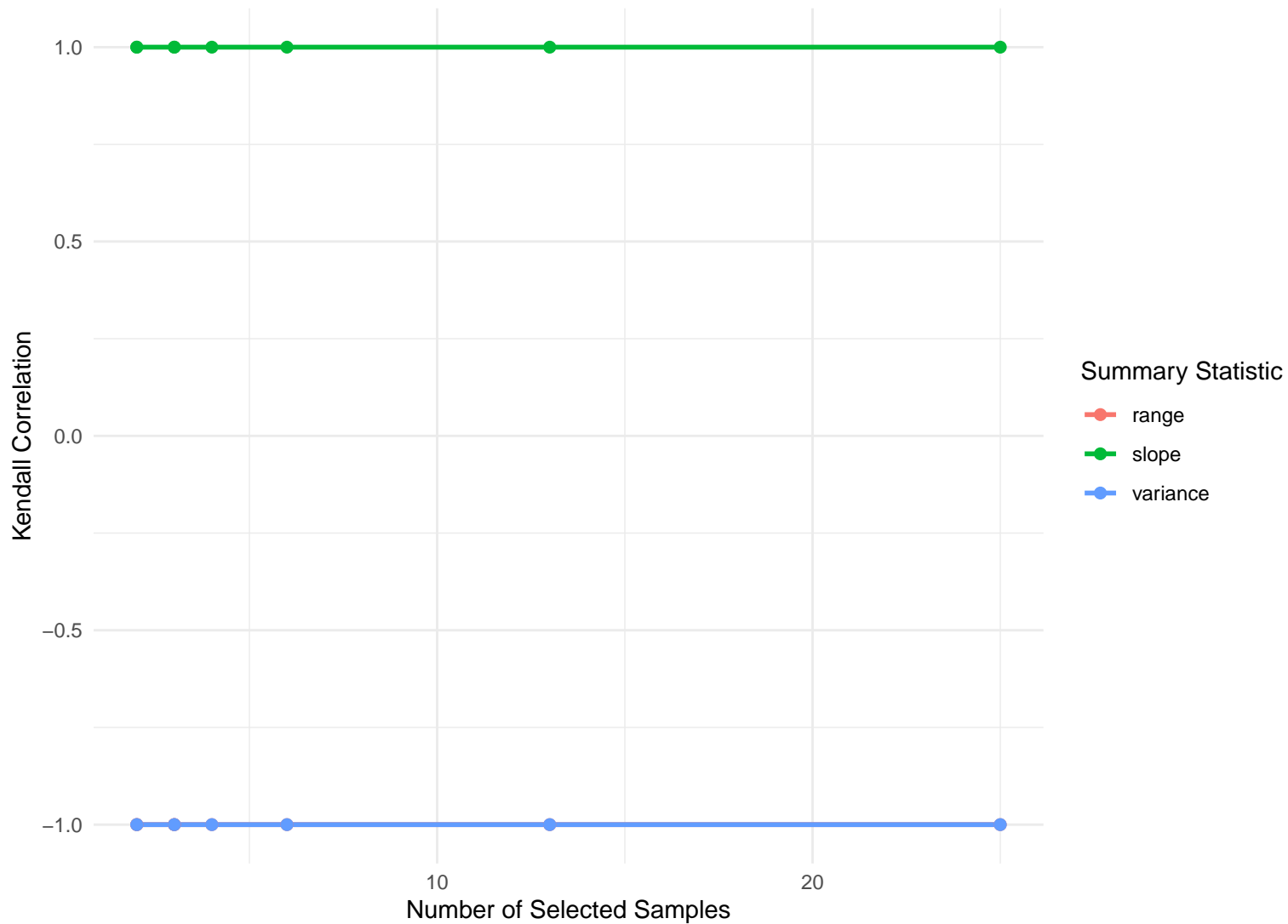
Kendall Correlation

Summary Statistic

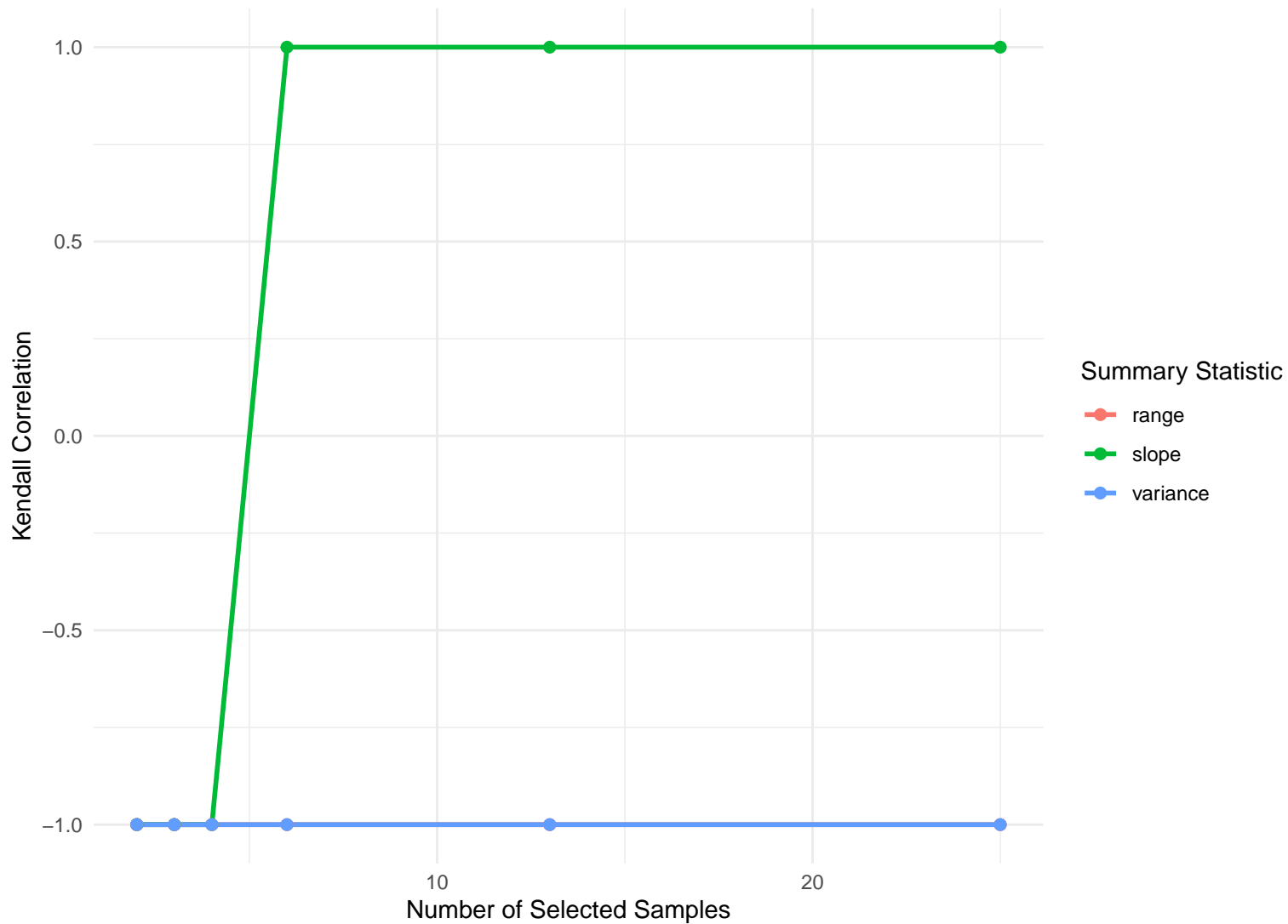
- range
- slope
- variance



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

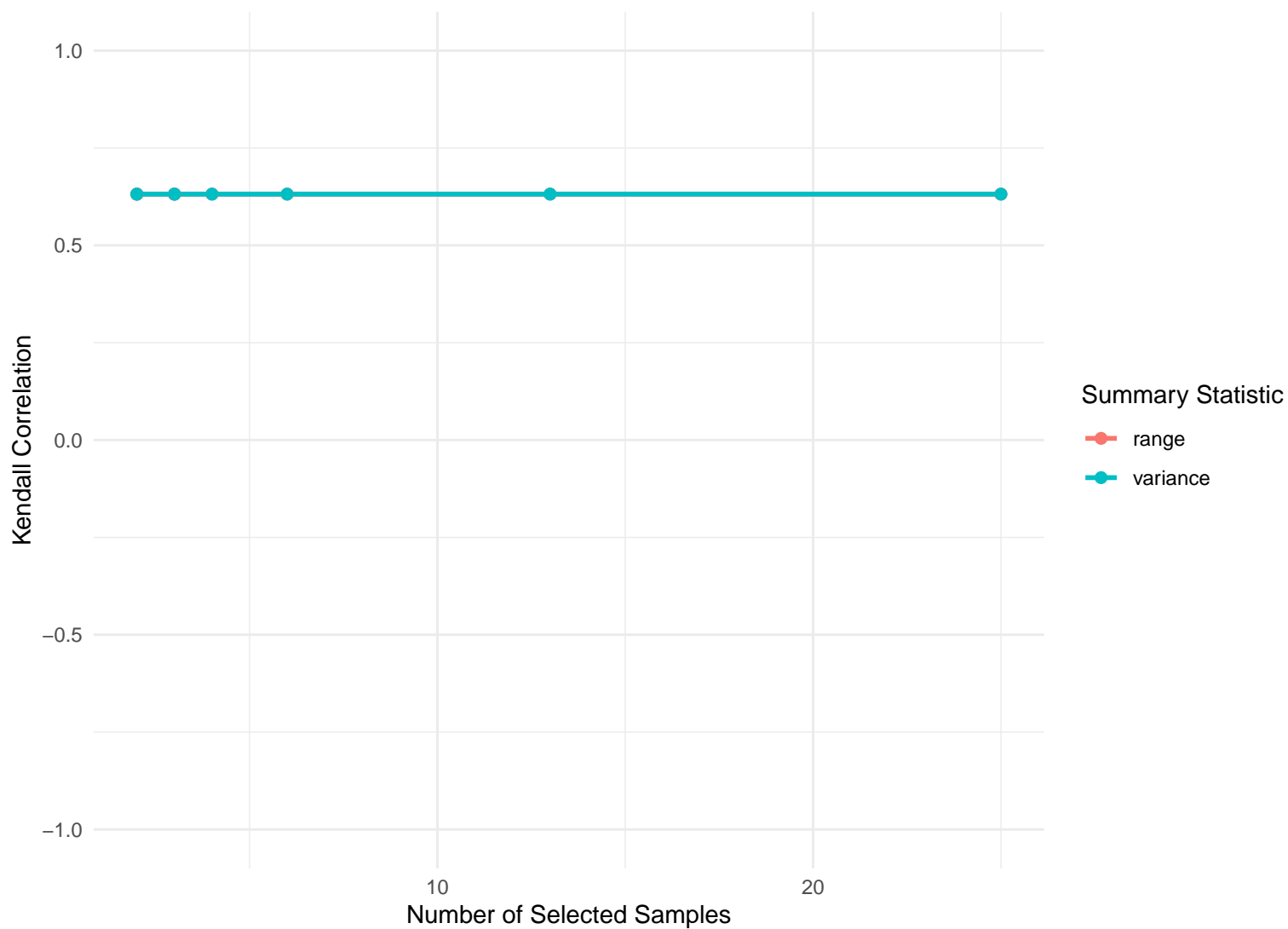


Score: PSI (Local, partial1 , Genotype: wave )





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

1.0

0.5

0.0

-0.5

-1.0

Number of Selected Samples

10

20



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



Number of Selected Samples

## Summary Statistic

max

mean

mean\_lower

mean\_upper

median

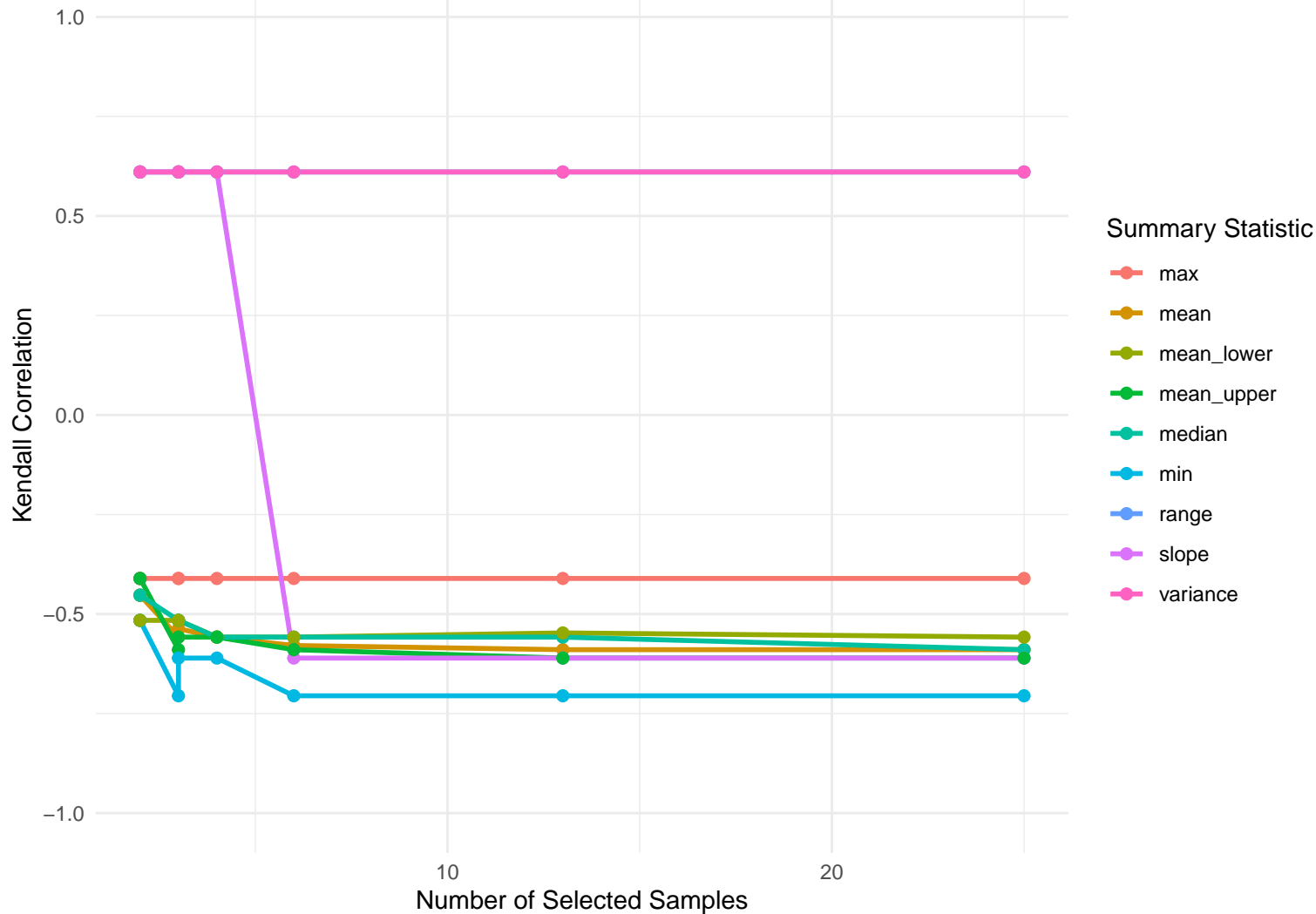
min

range

slope

variance

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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

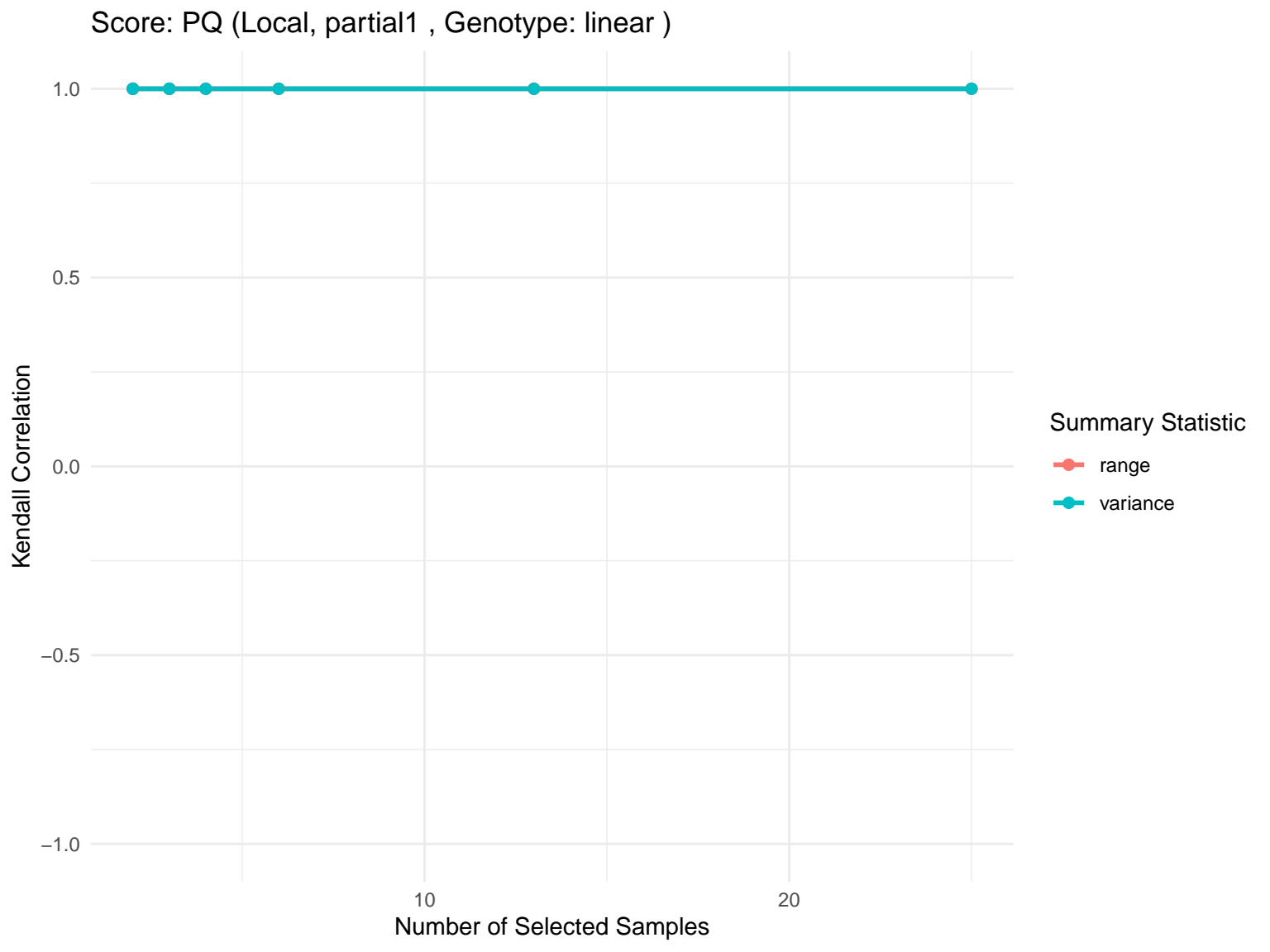
10

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

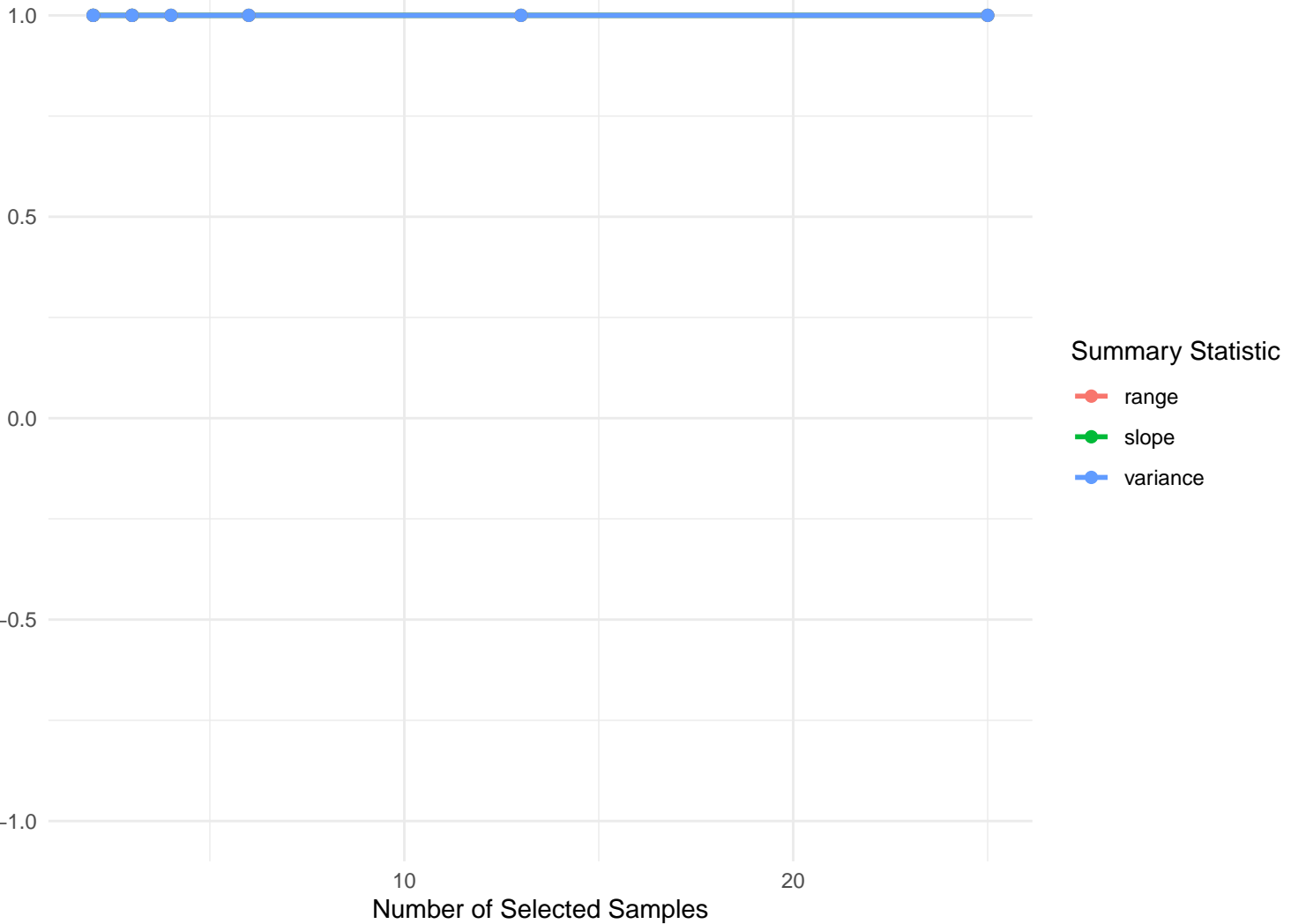
10

20

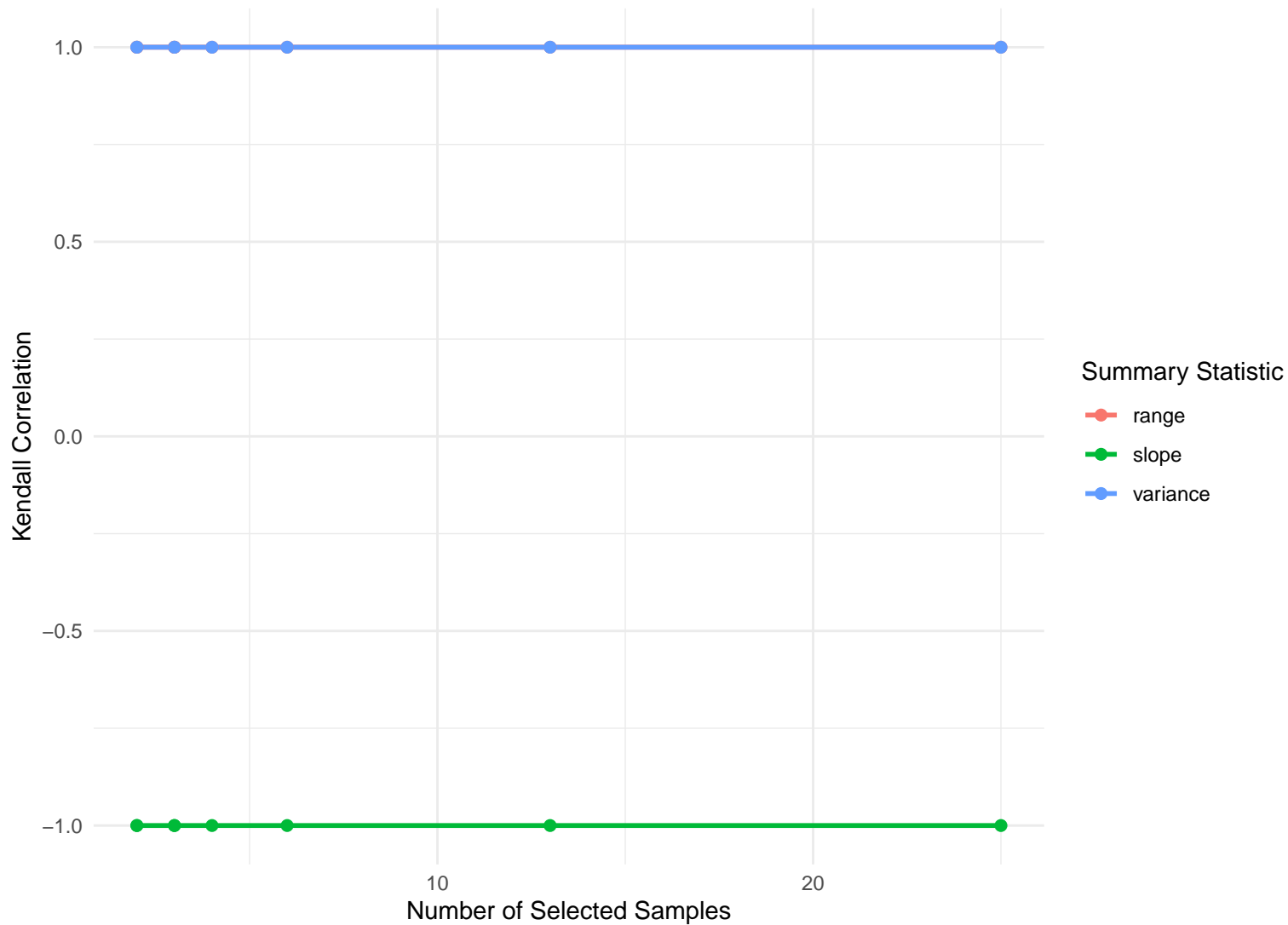
Number of Selected Samples

Summary Statistic

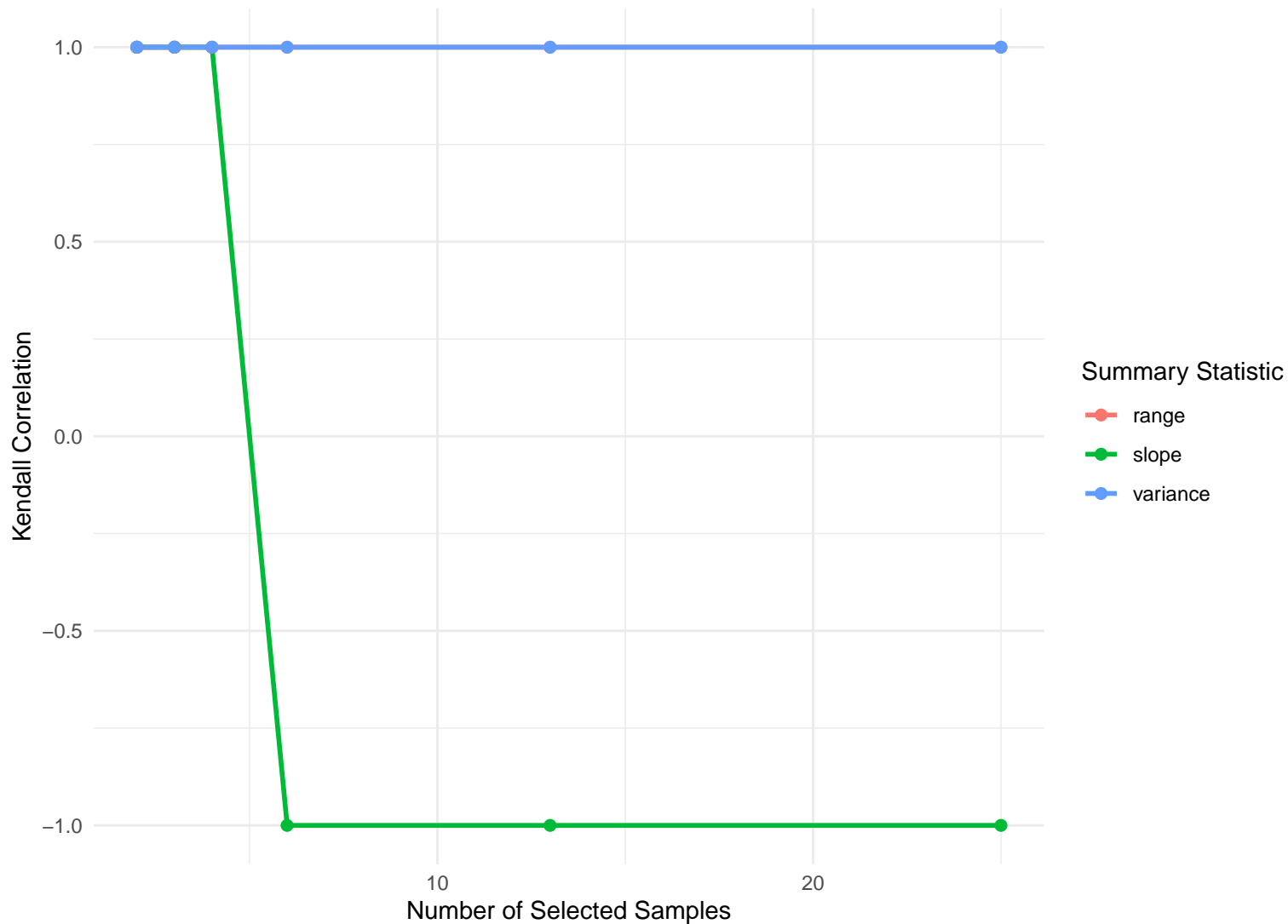
- range
- slope
- variance



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

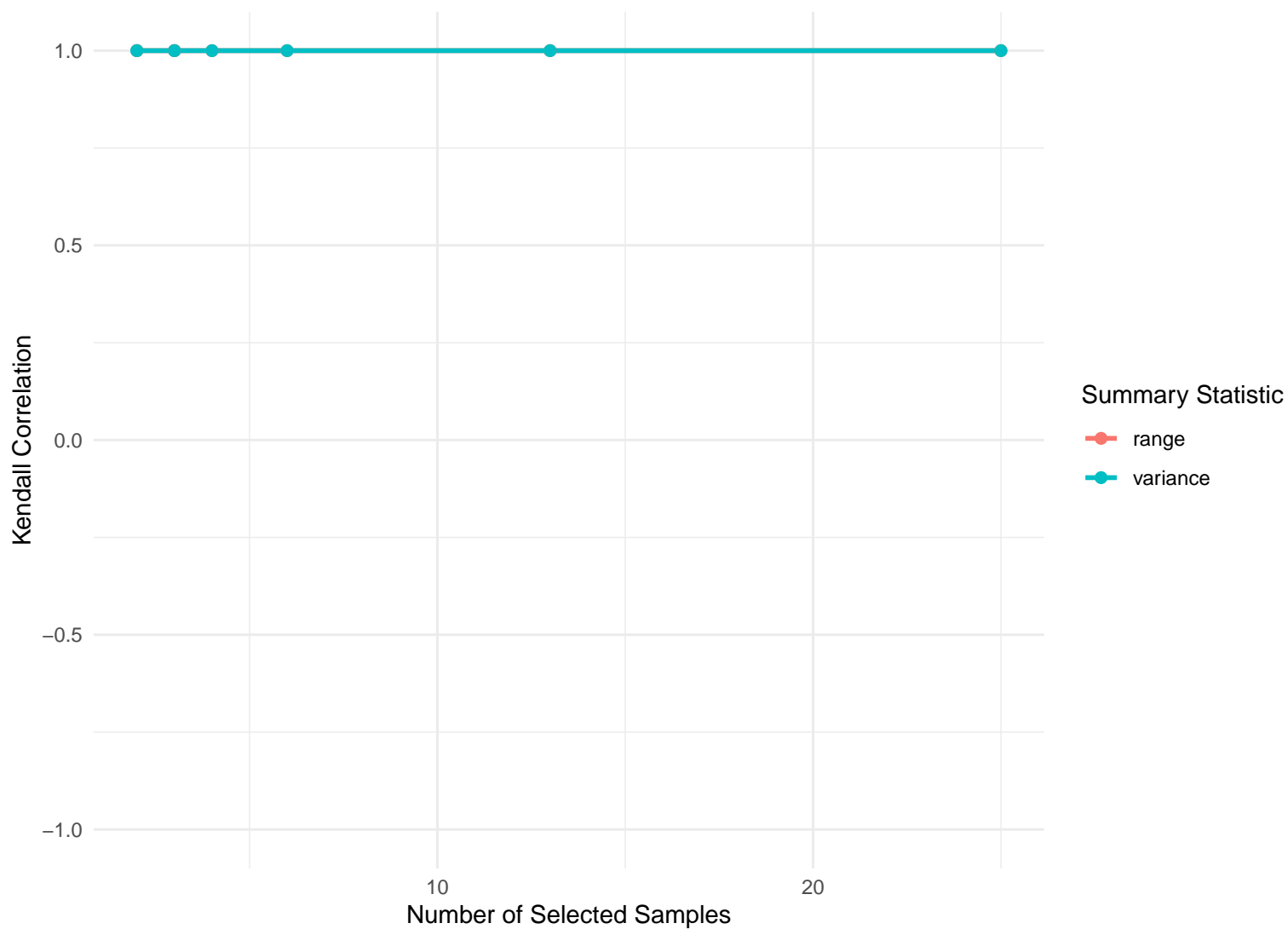


Score: PQ (Local, partial1 , Genotype: wave )





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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

20

Number of Selected Samples

Summary Statistic

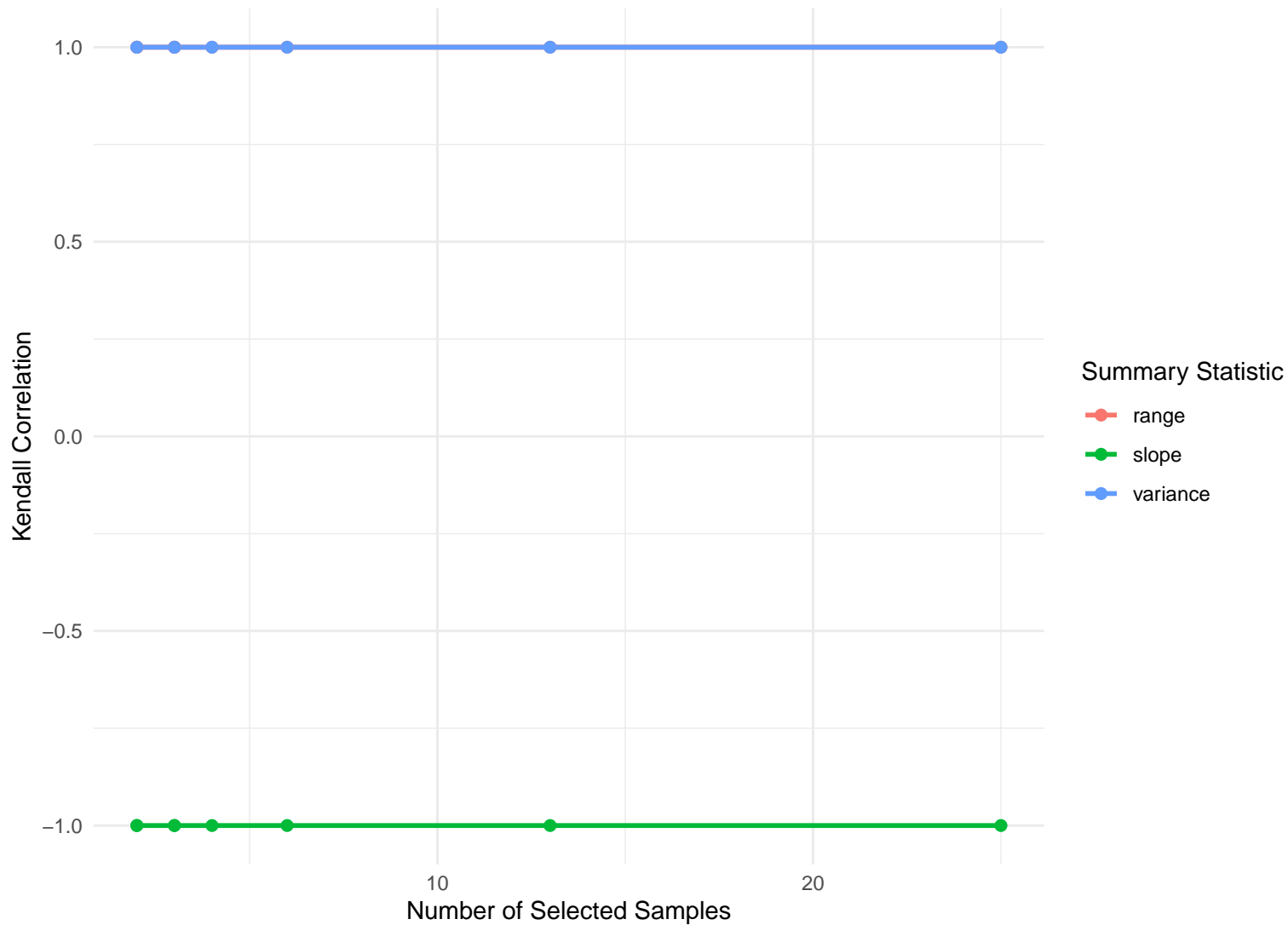
range

slope

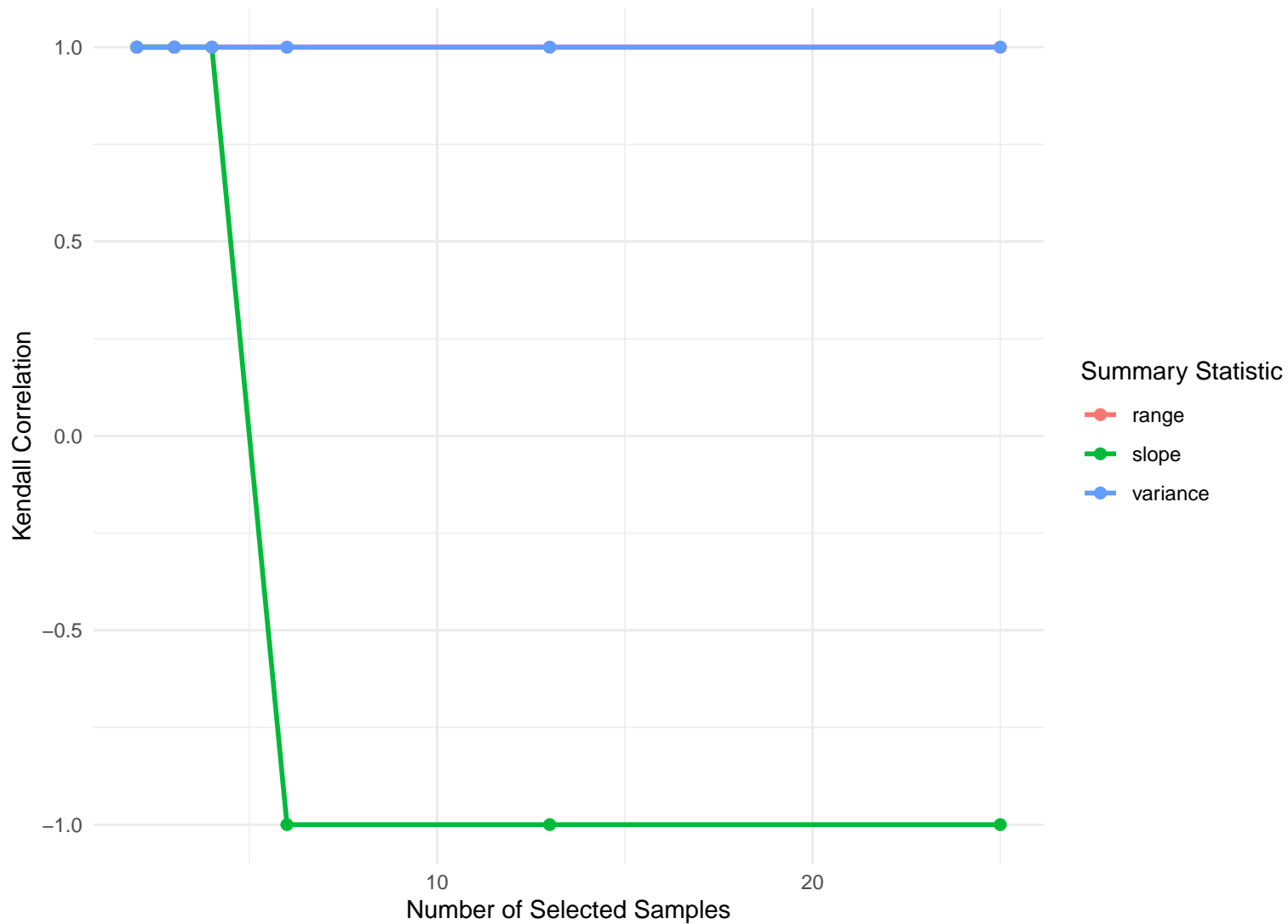
variance



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

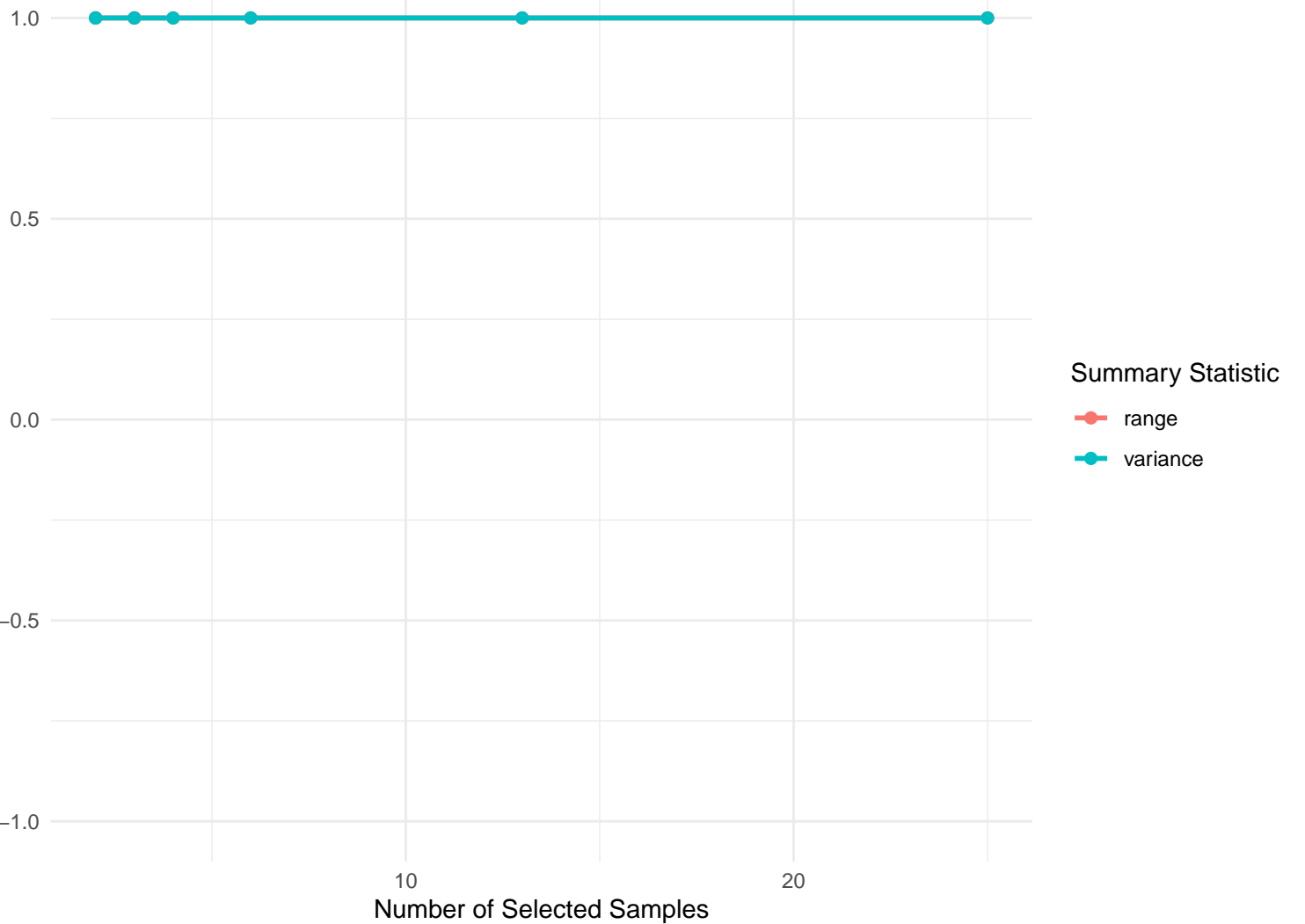
20

Number of Selected Samples

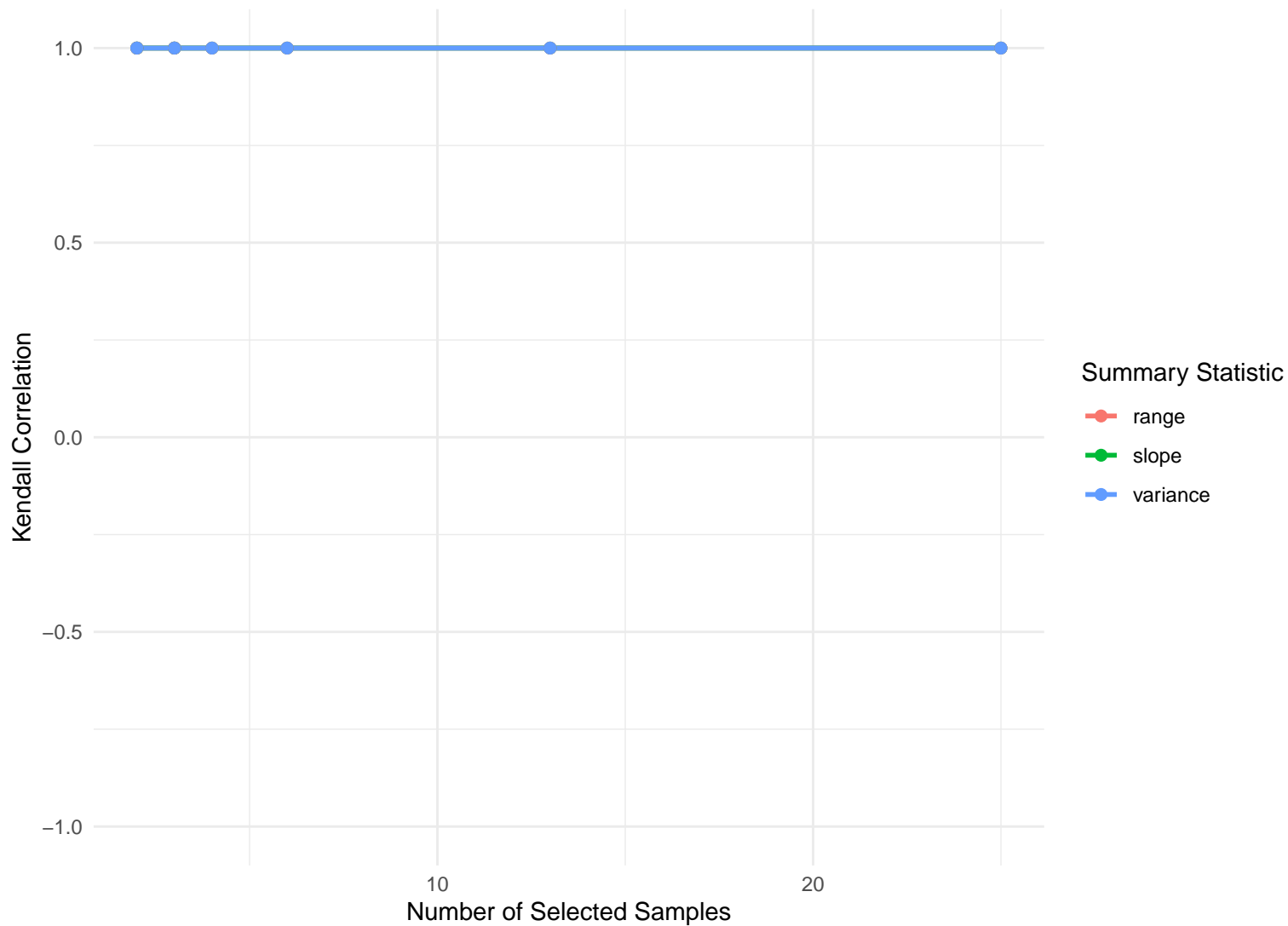
Summary Statistic

range

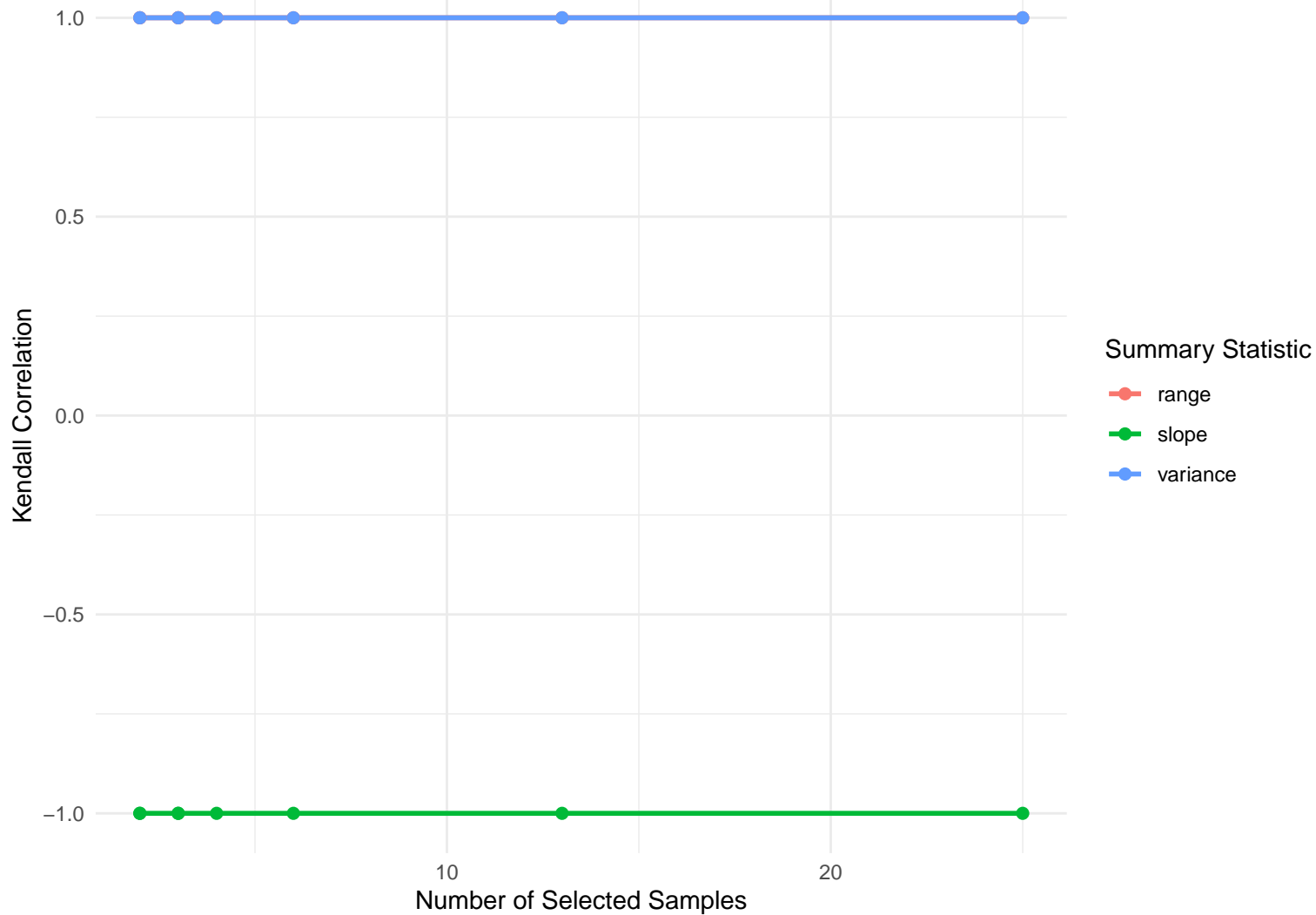
variance



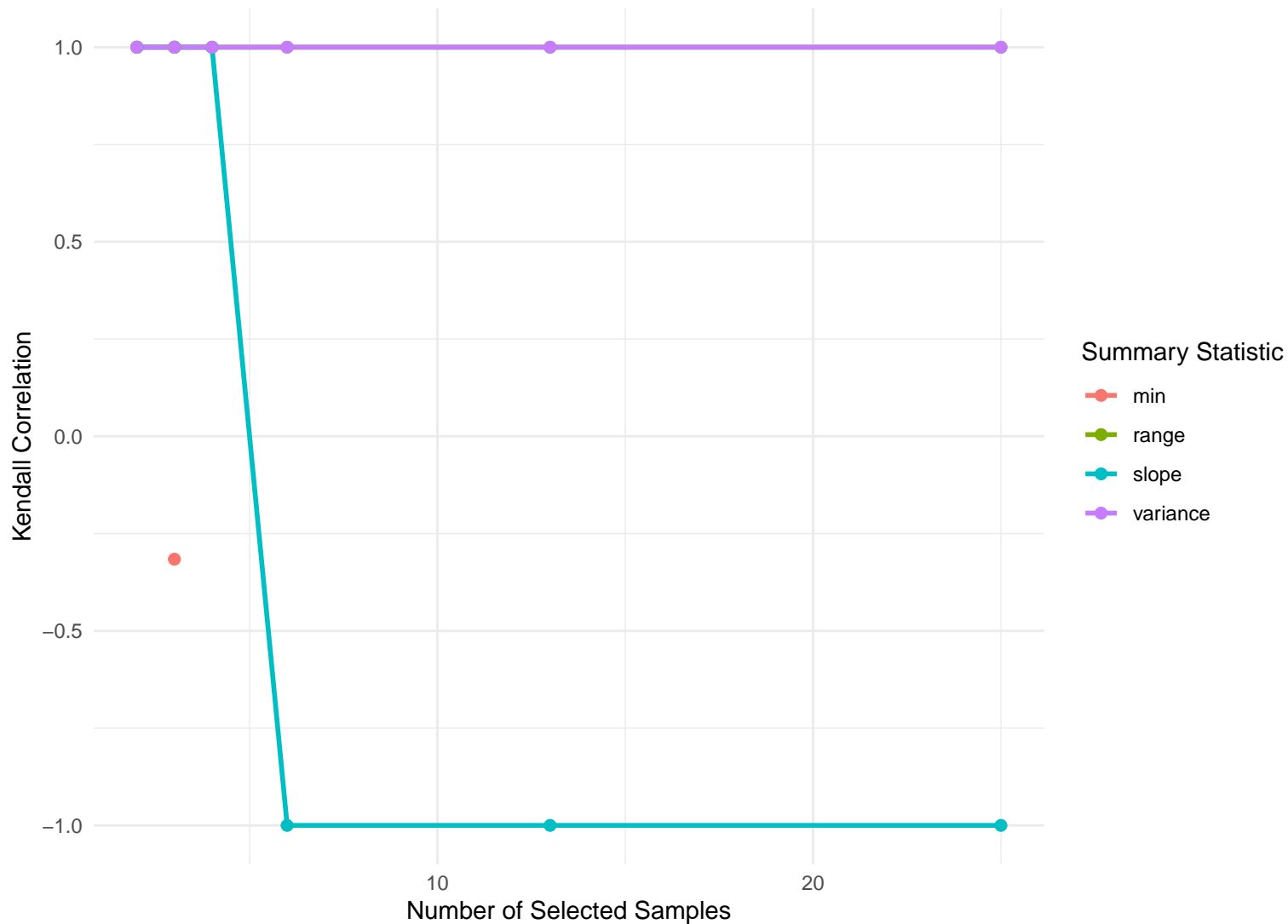
Score: NRW (Local, partial1 , Genotype: gaussian )



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

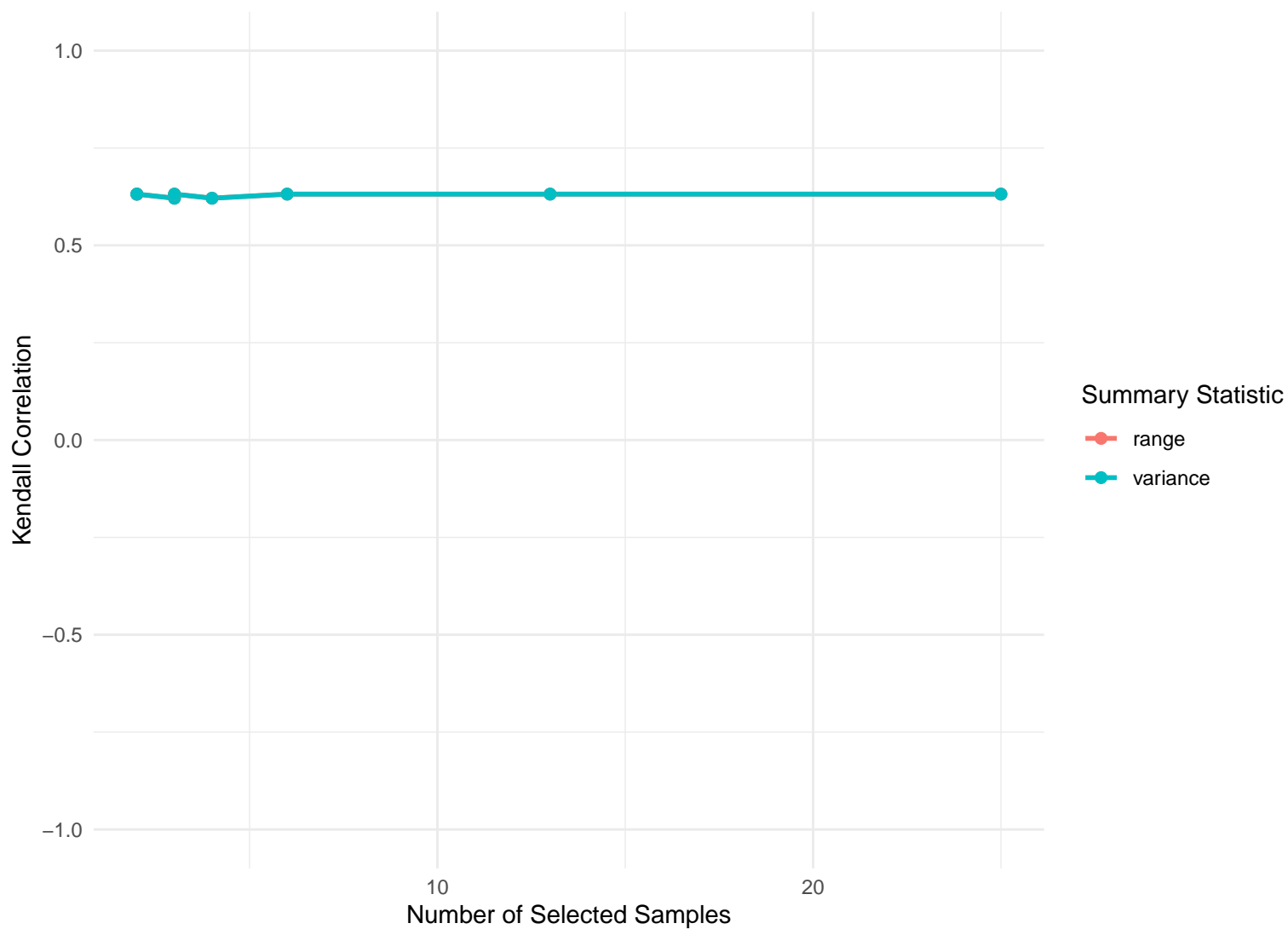


Score: NRW (Local, partial1 , Genotype: wave )





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



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

Kendall Correlation

Summary Statistic

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

1.0

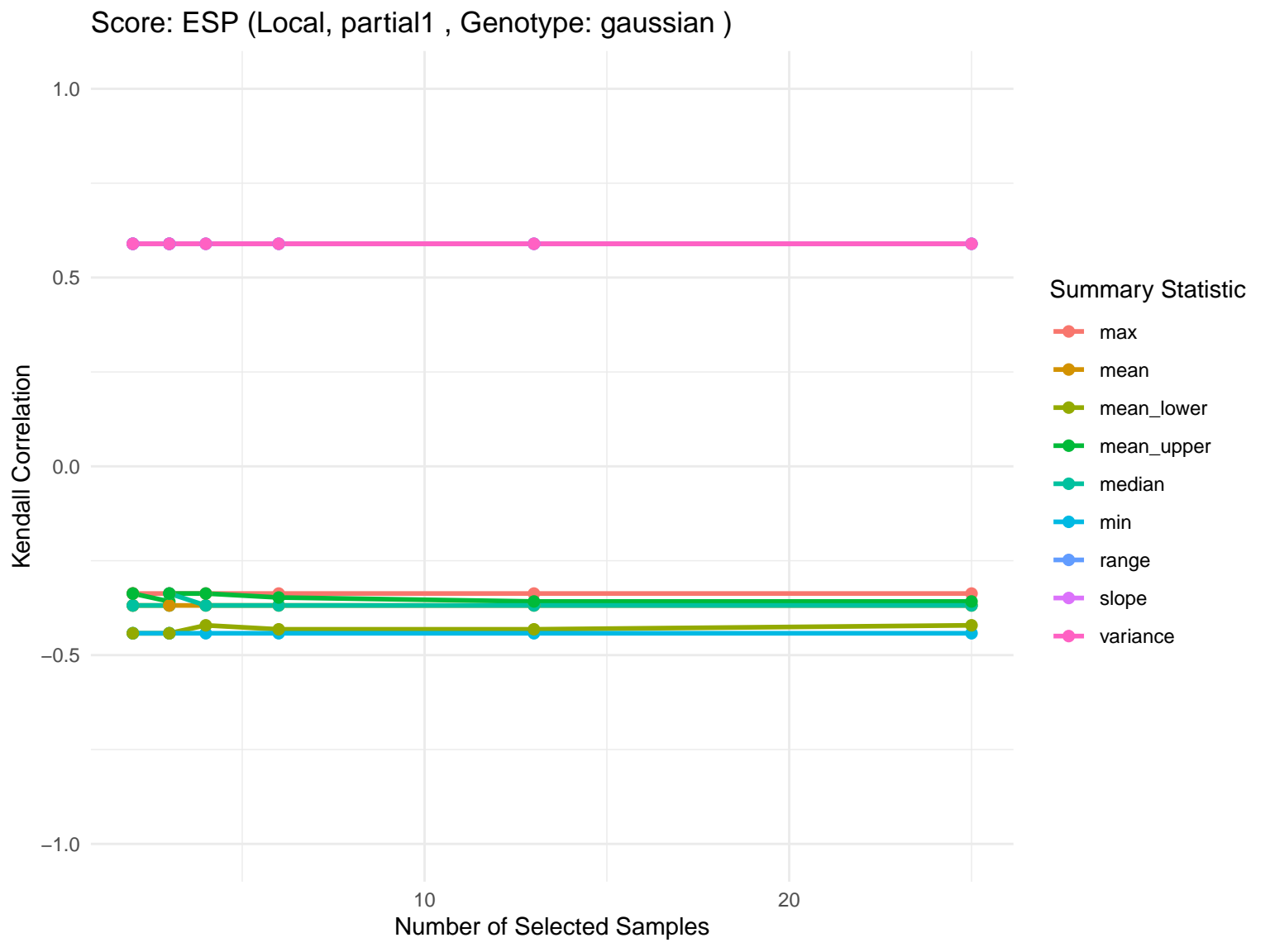
0.5

0.0

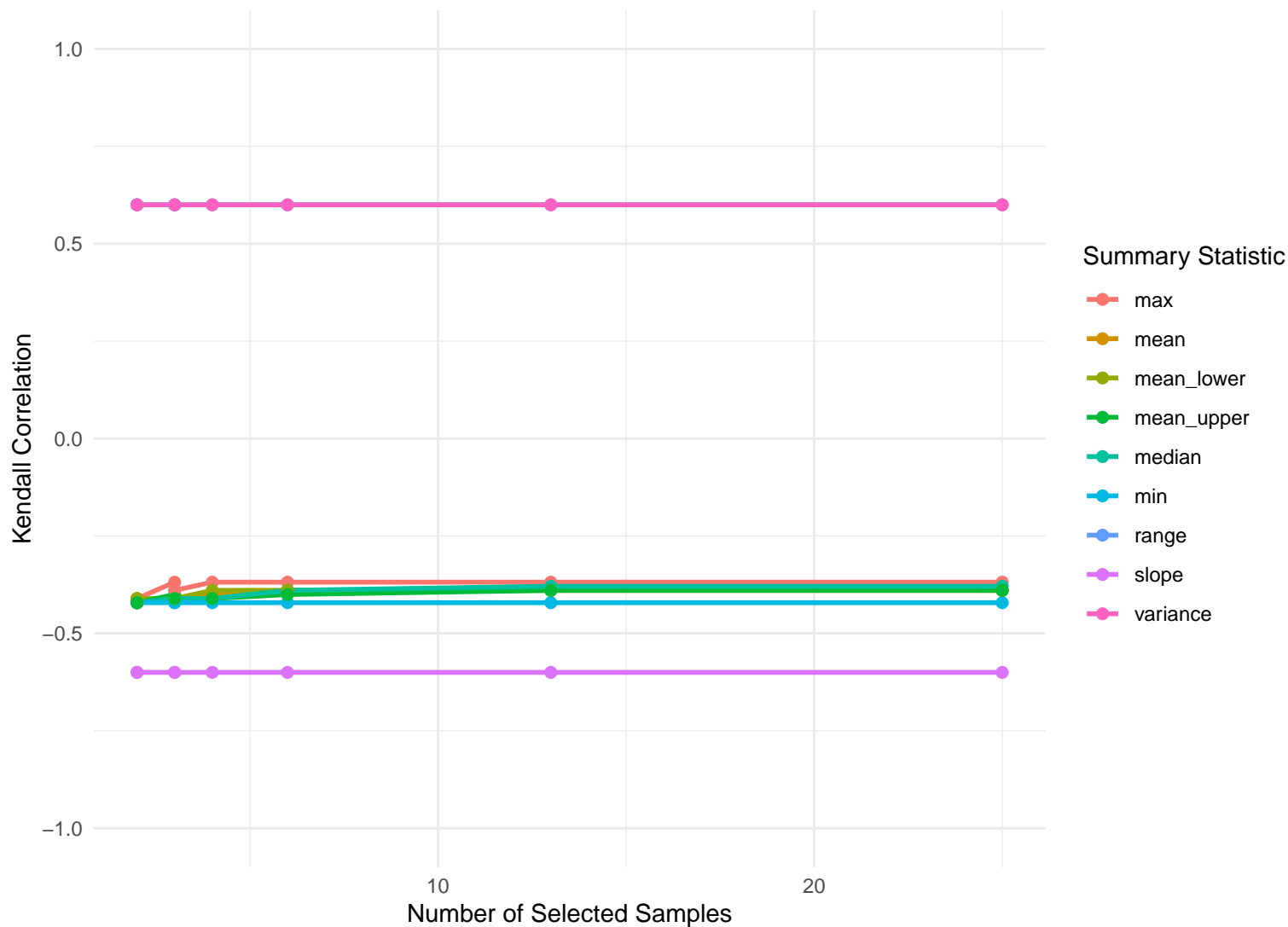
-0.5

-1.0

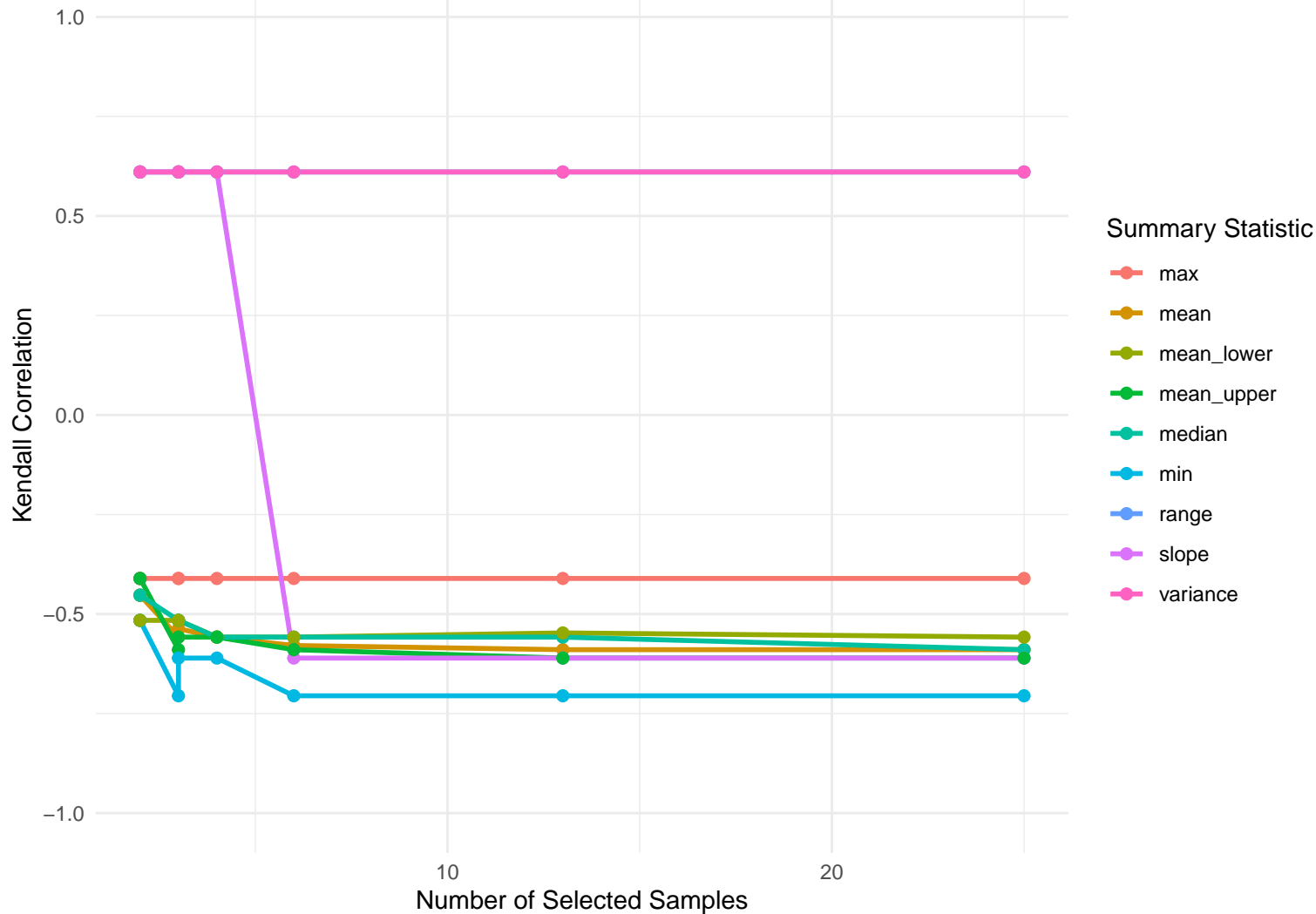
10  
20  
Number of Selected Samples



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

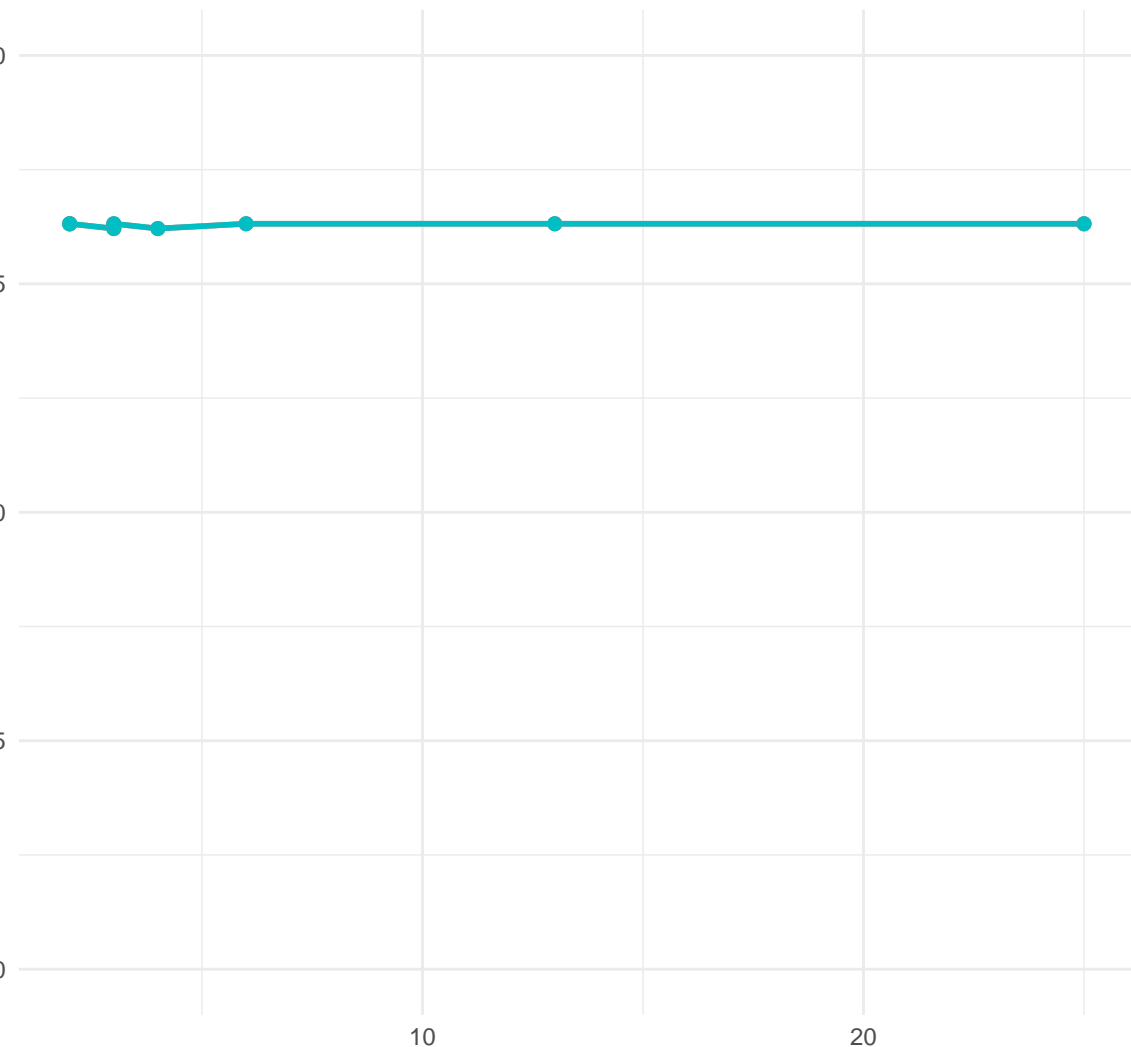
20

Number of Selected Samples

Summary Statistic

range

variance



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

Kendall Correlation

Summary Statistic

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

1.0

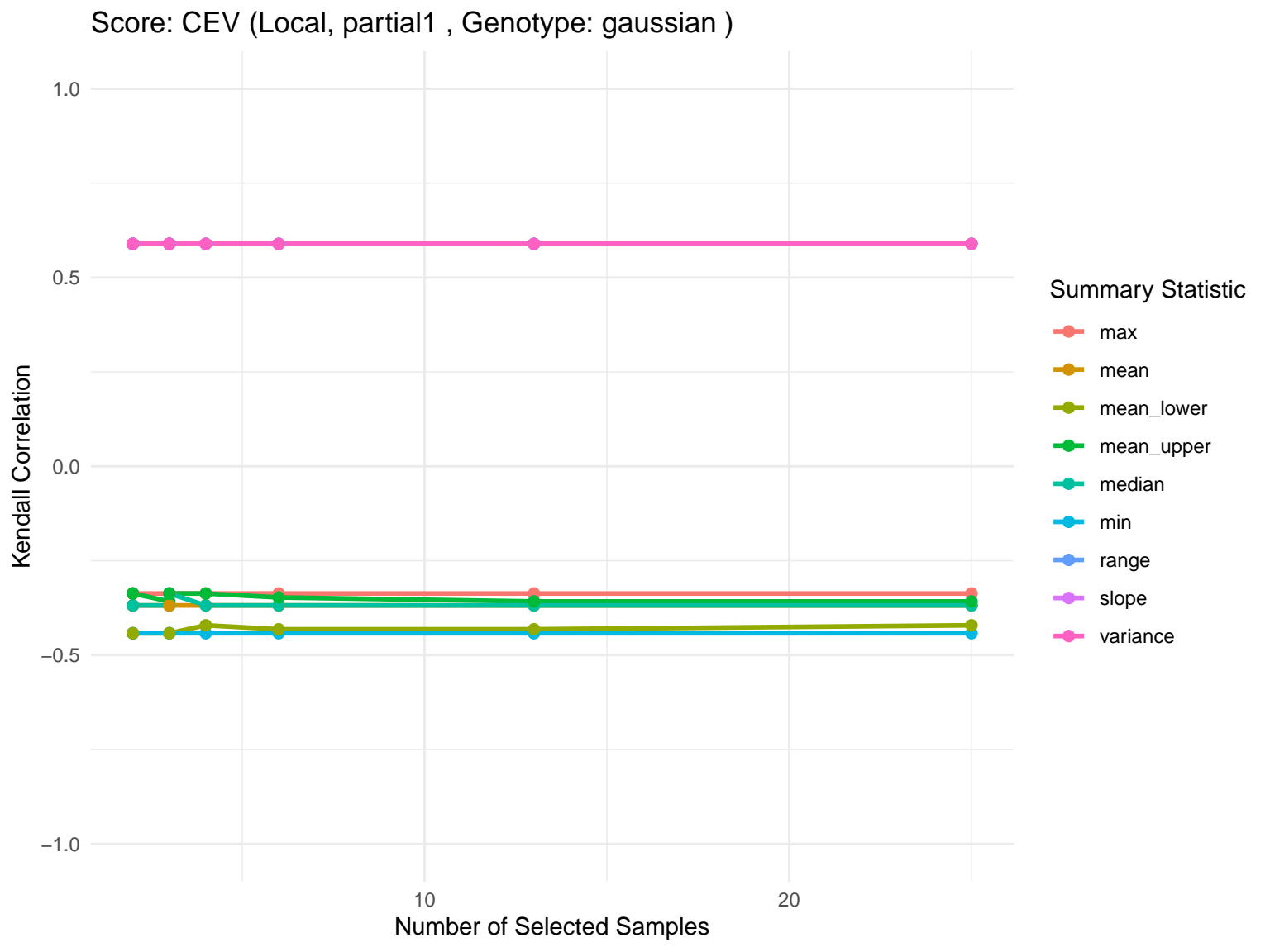
0.5

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

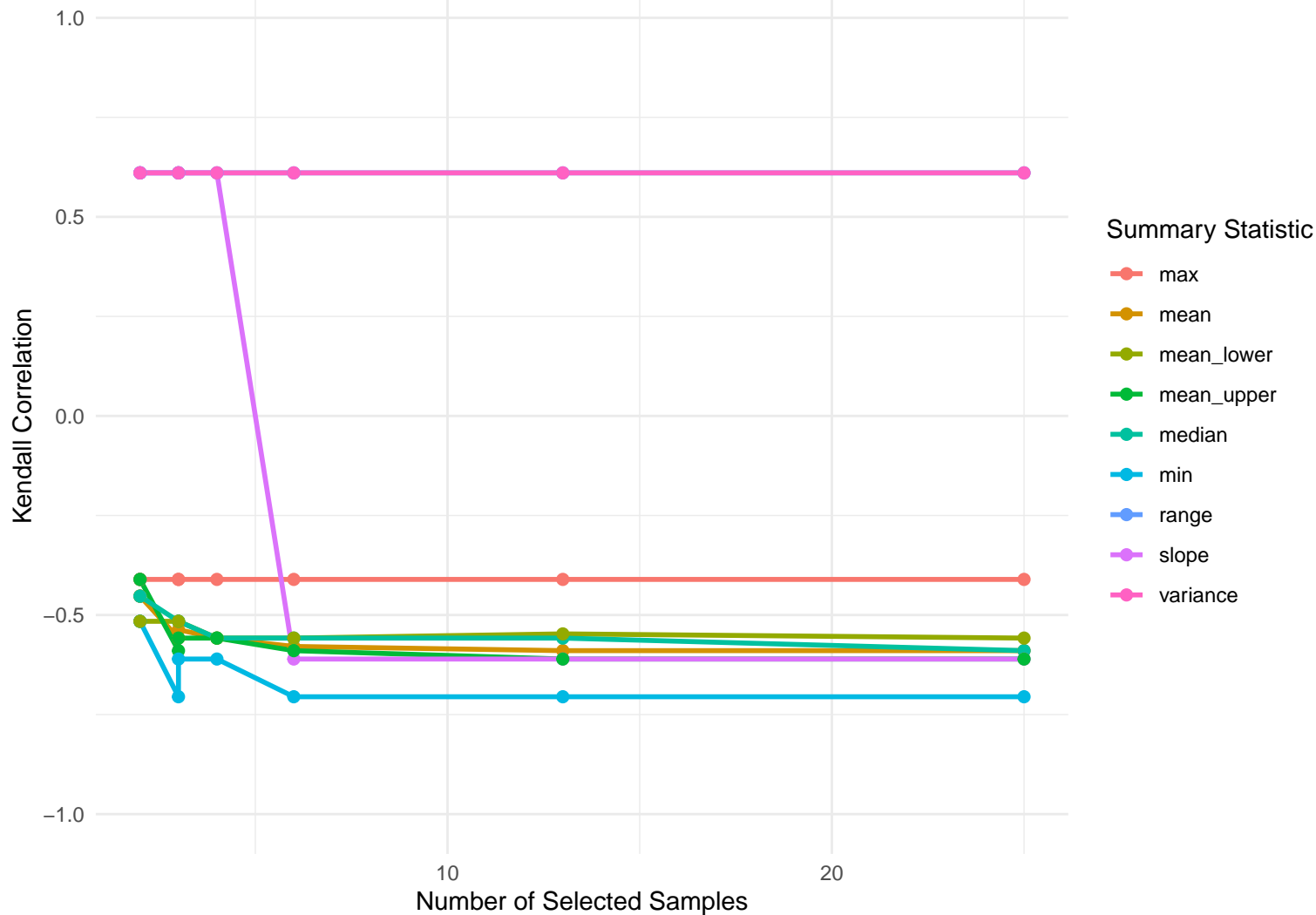
min

range

slope

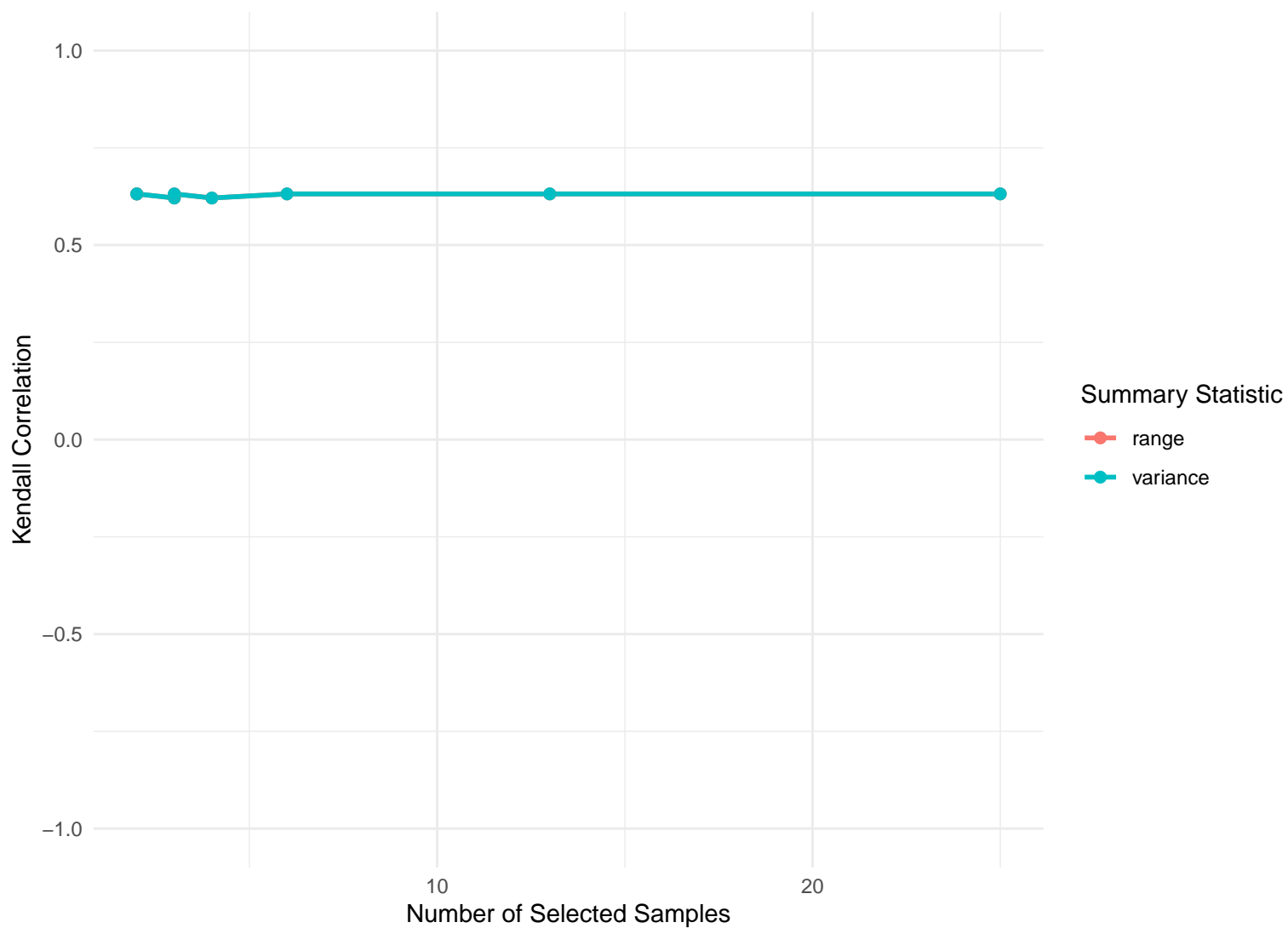
variance

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

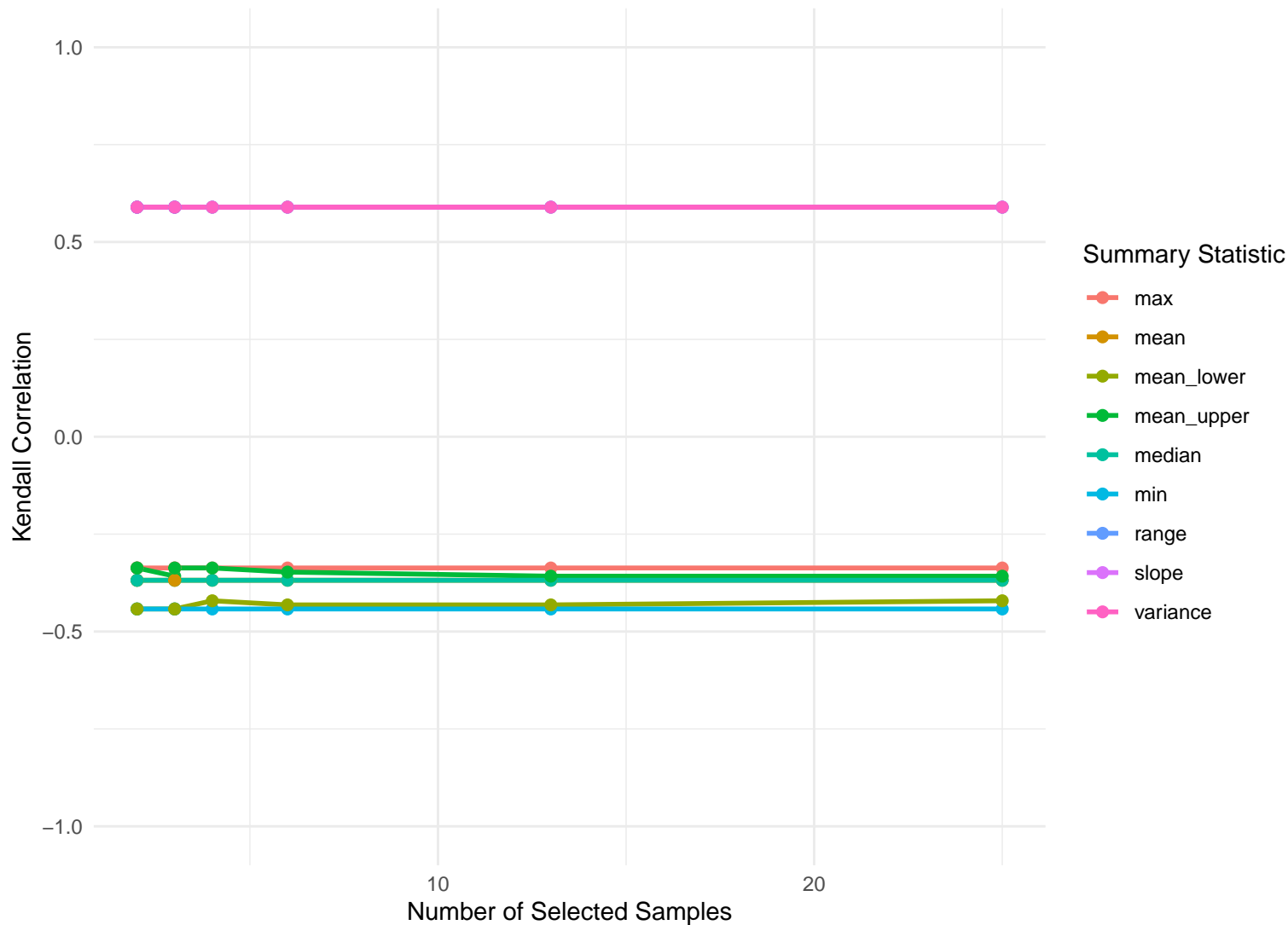




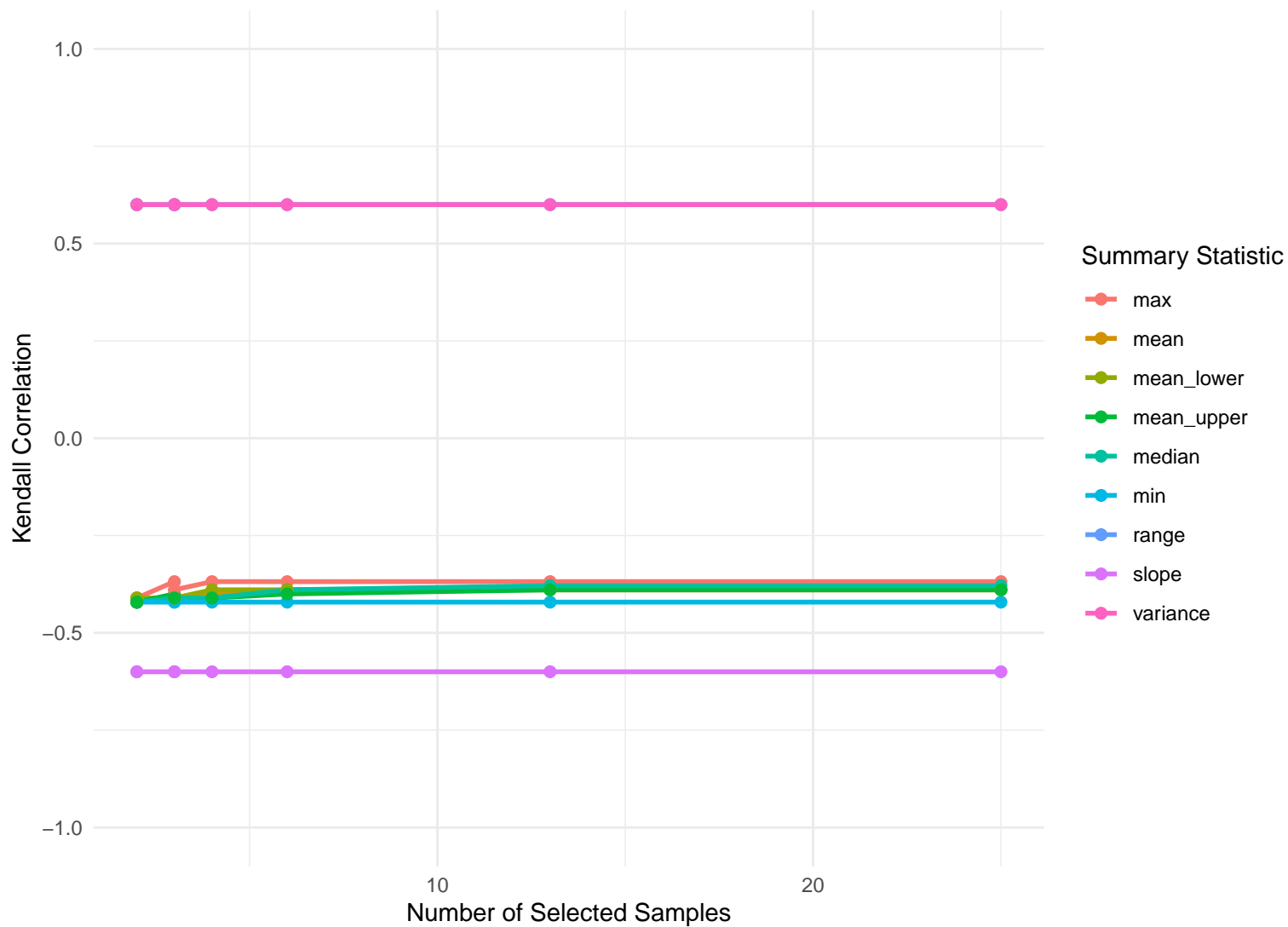
Score: PFI (Local, partial1 , Genotype: linear )



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



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



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

Kendall Correlation

1.0

0.5

0.0

-0.5

-1.0

10

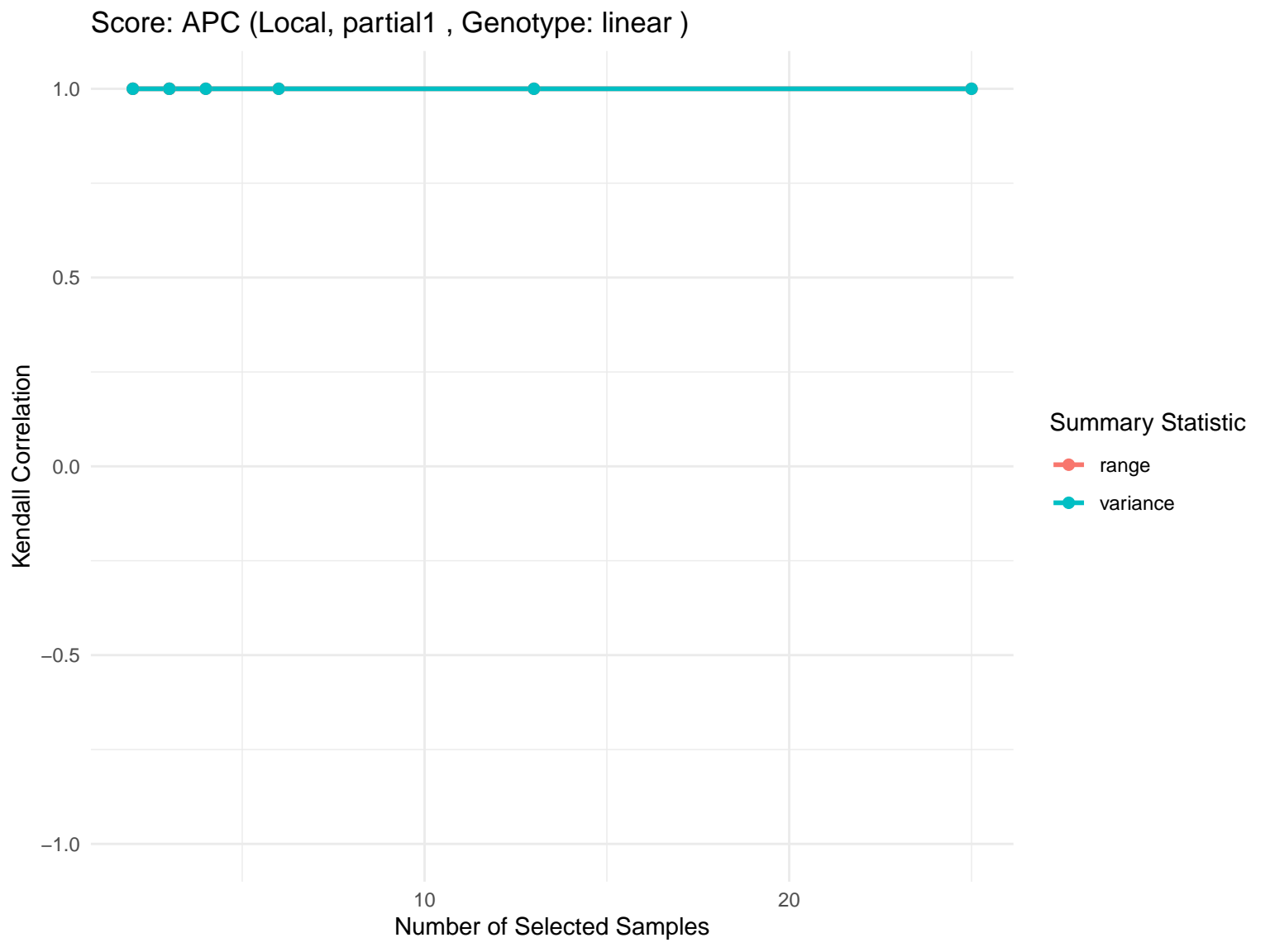
20

Number of Selected Samples

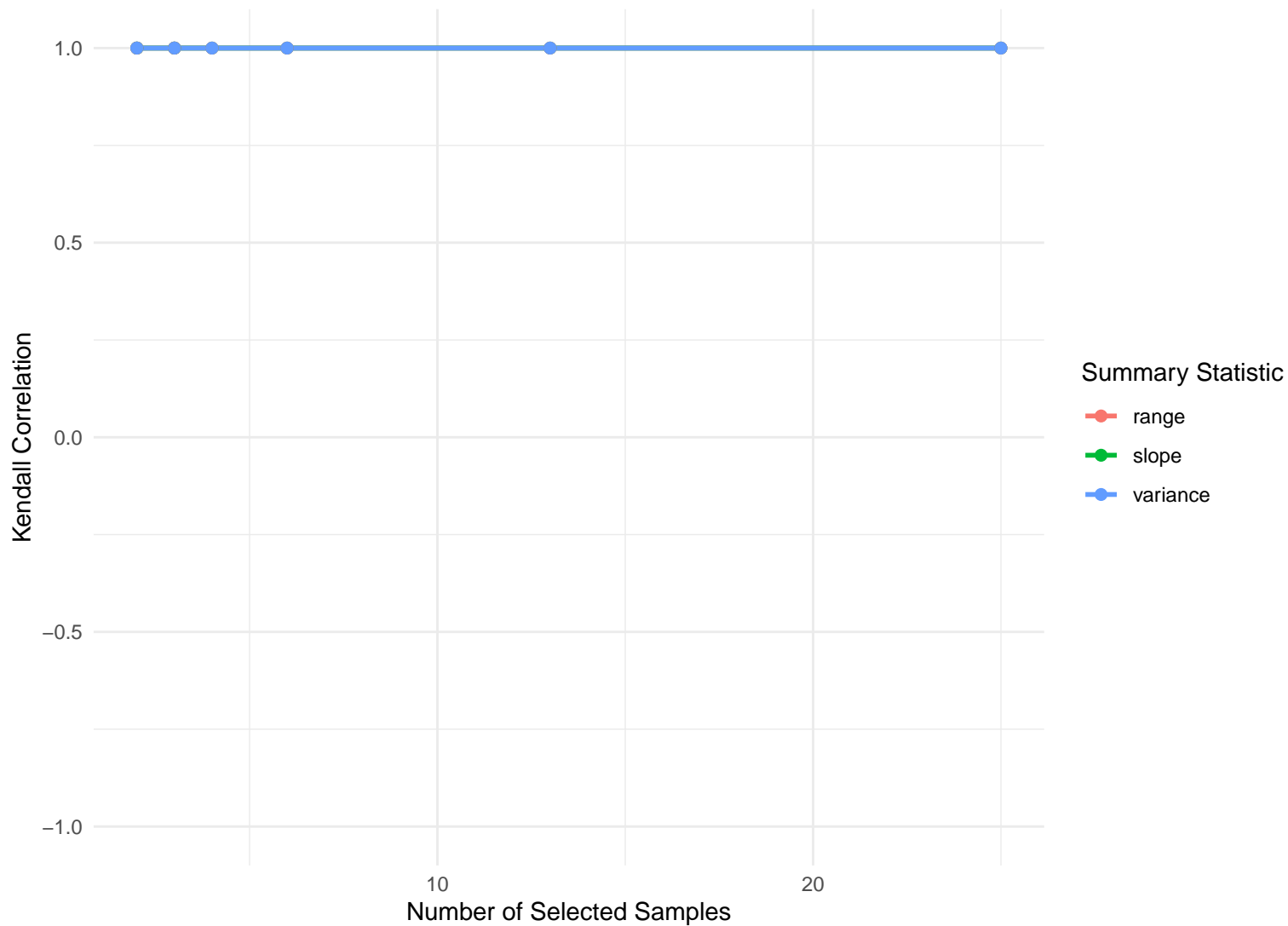
Summary Statistic

range

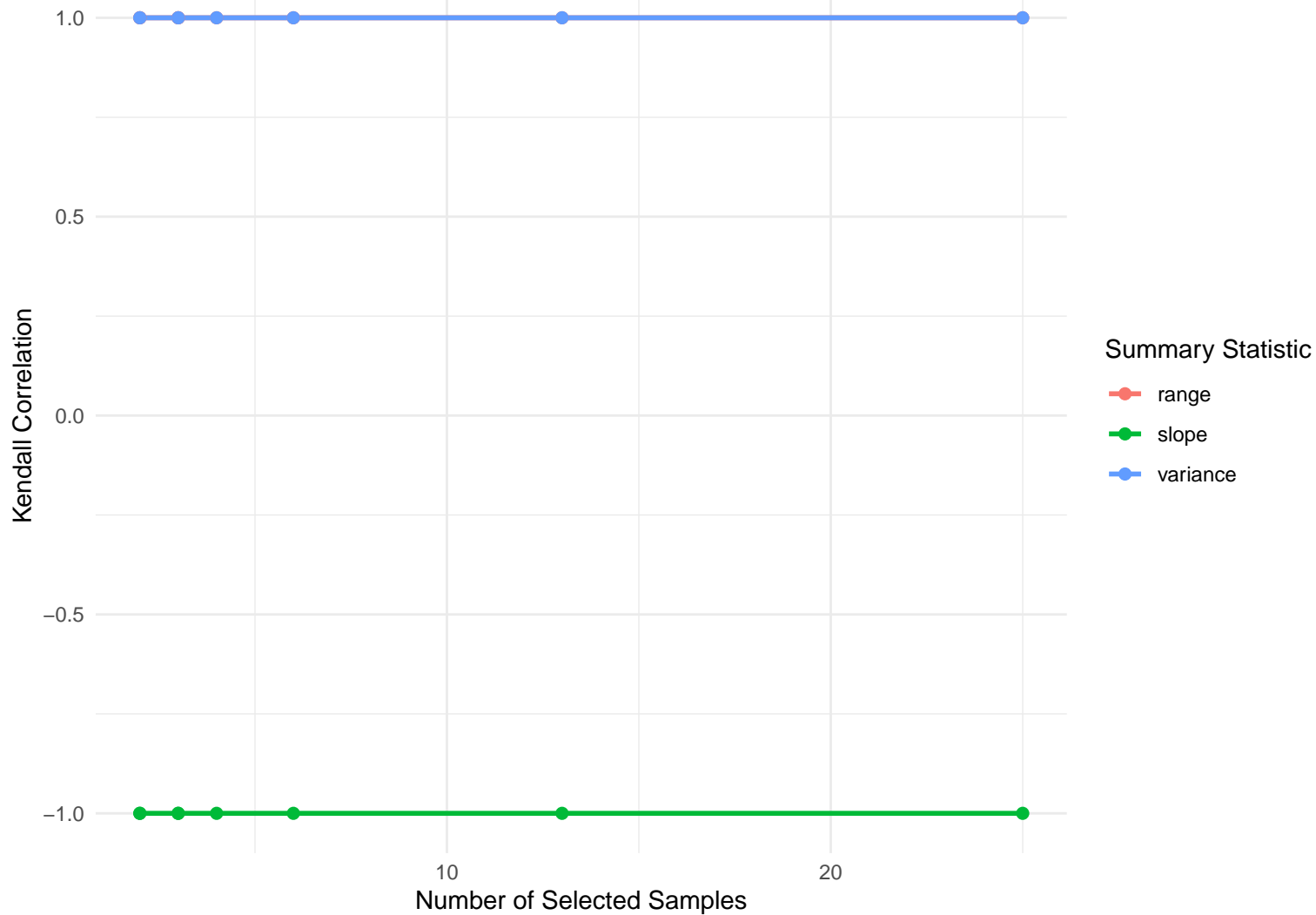
variance



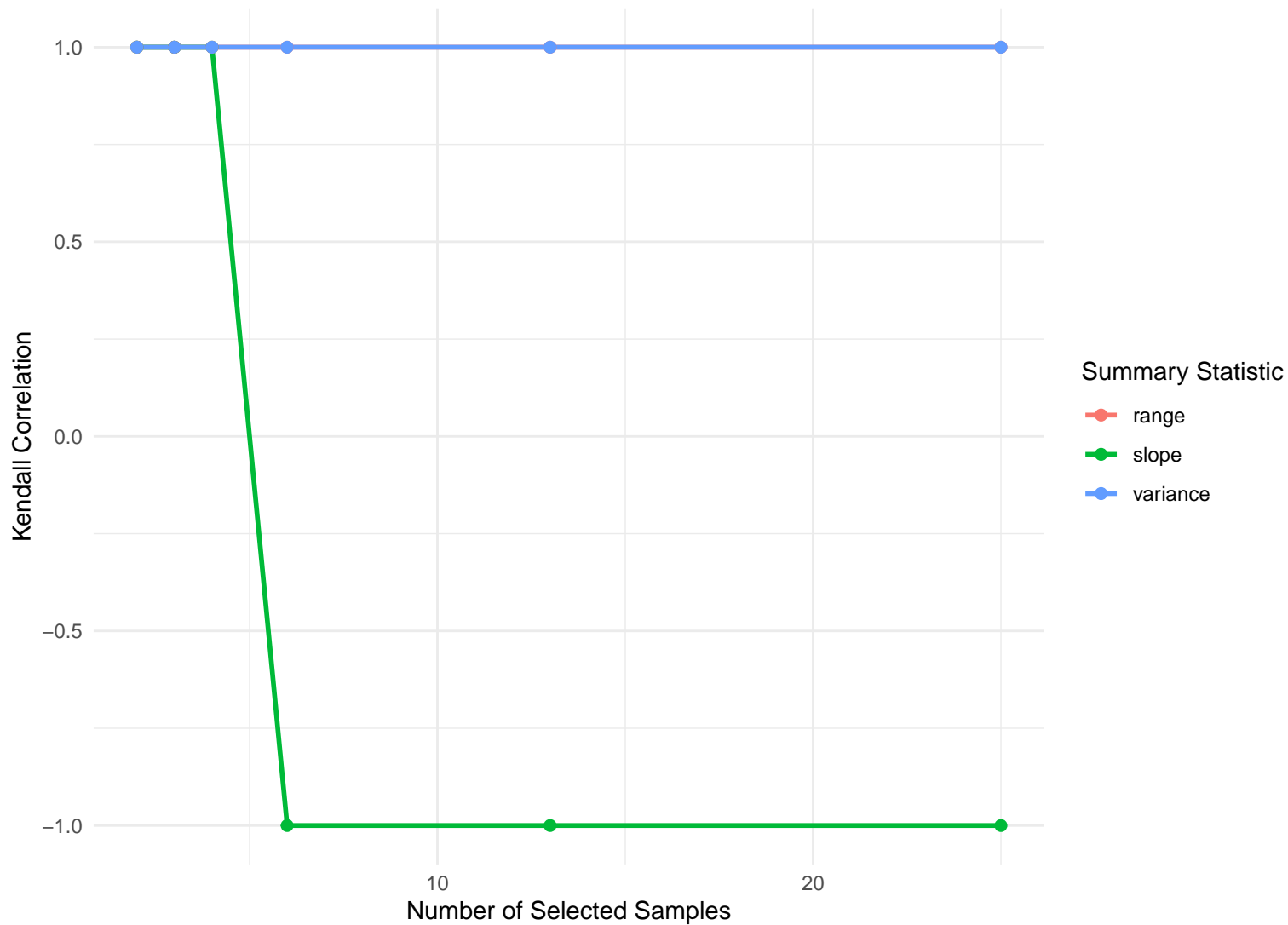
Score: APC (Local, partial1 , Genotype: gaussian )



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

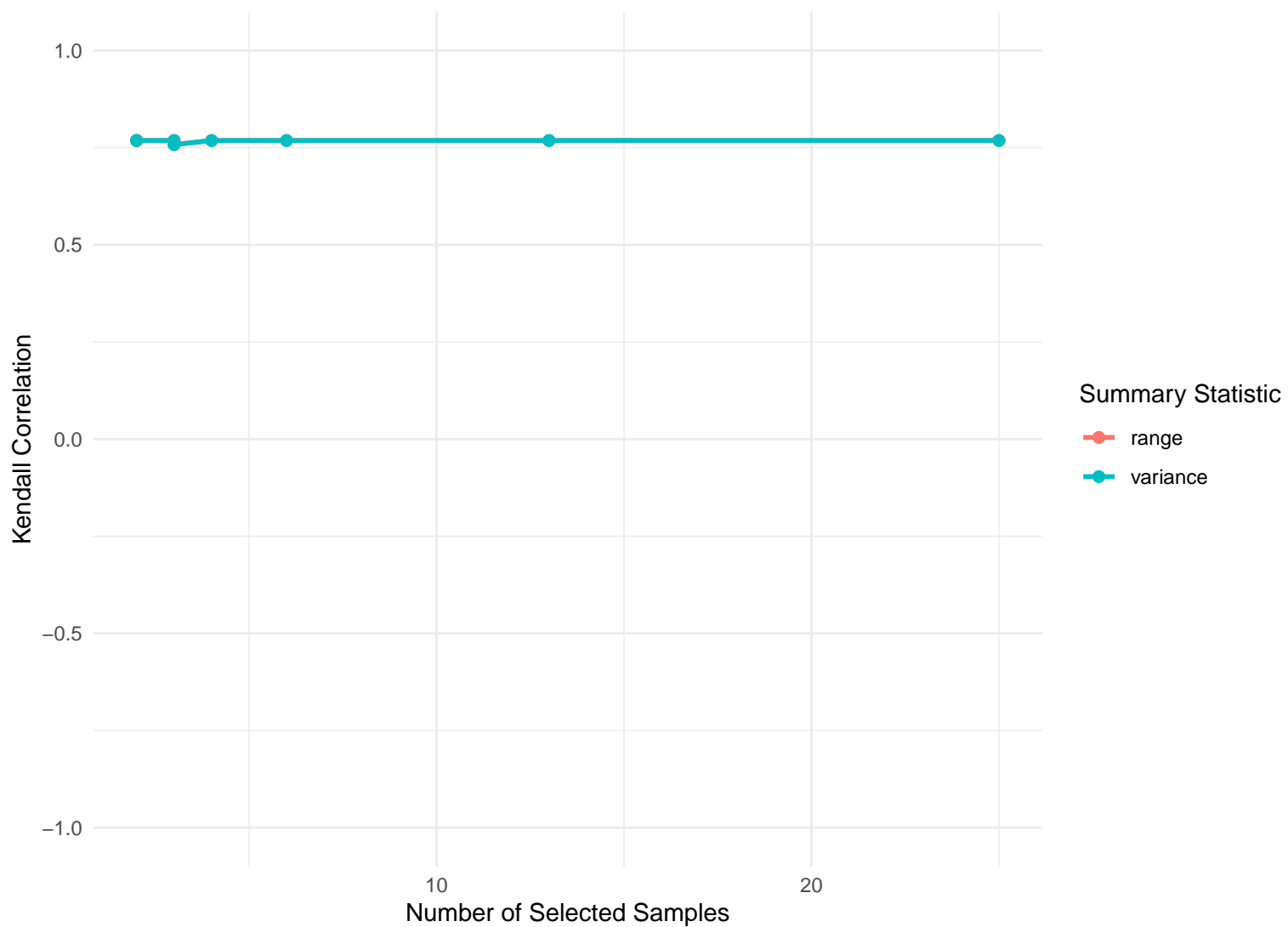


Score: APC (Local, partial1 , Genotype: wave )

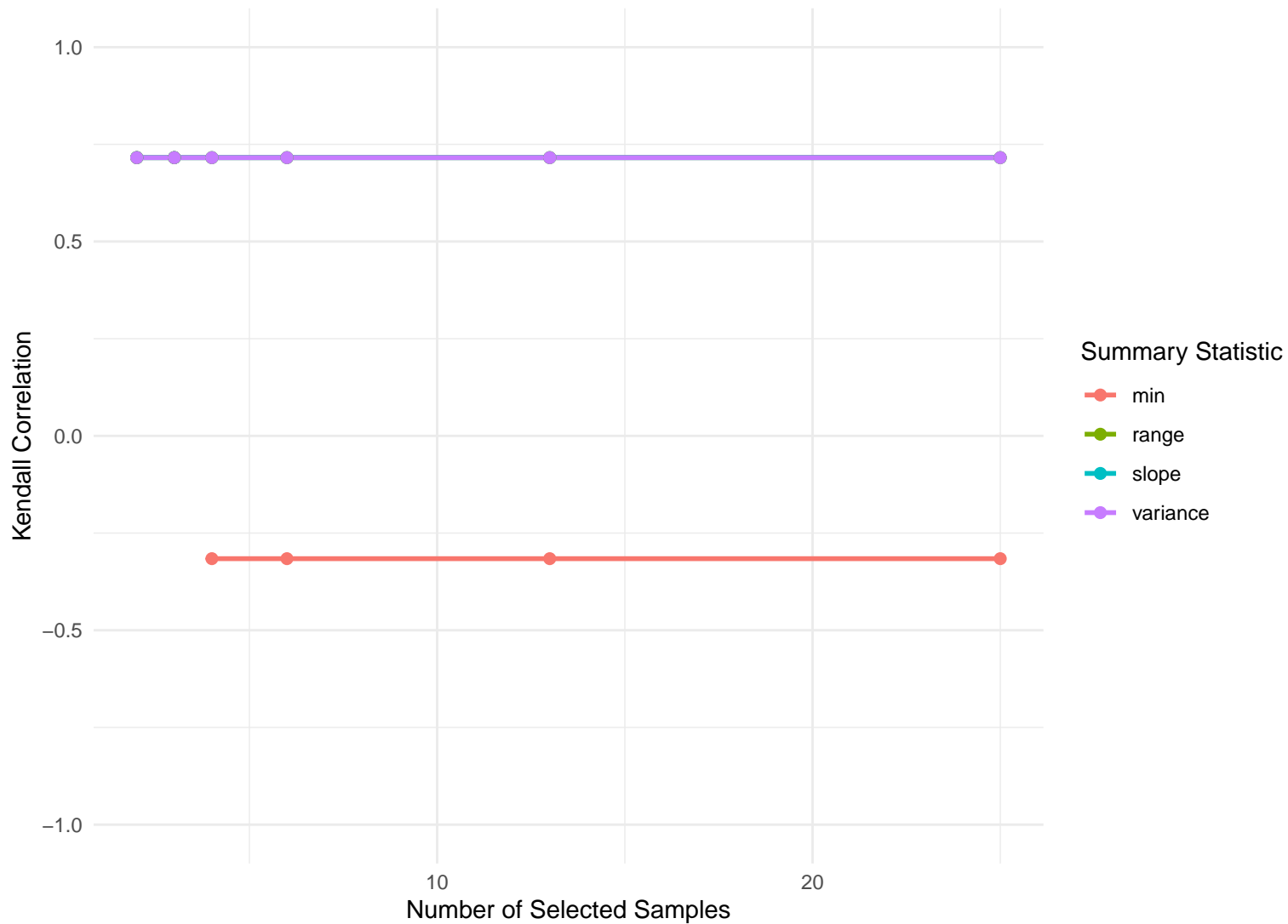




Score: SI (Local, partial1 , Genotype: linear )

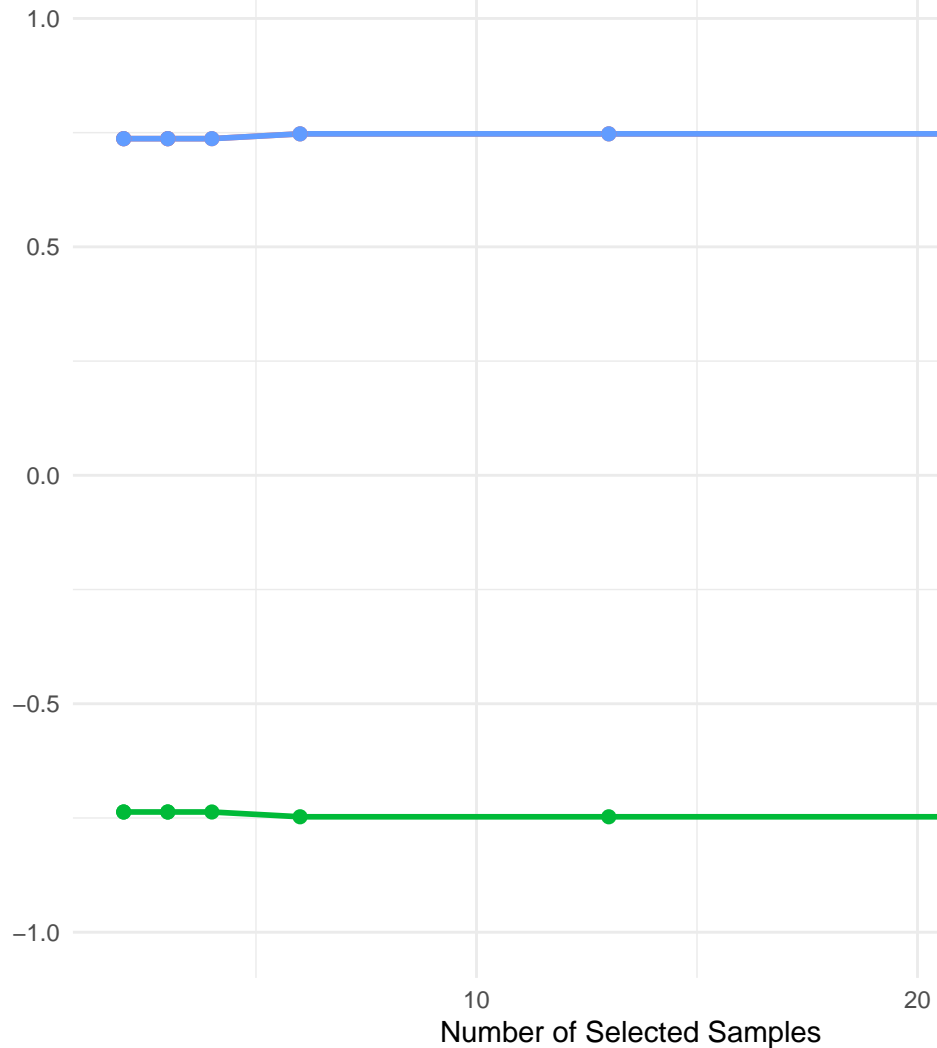


Score: SI (Local, partial1 , Genotype: gaussian )



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

Kendall Correlation



Summary Statistic

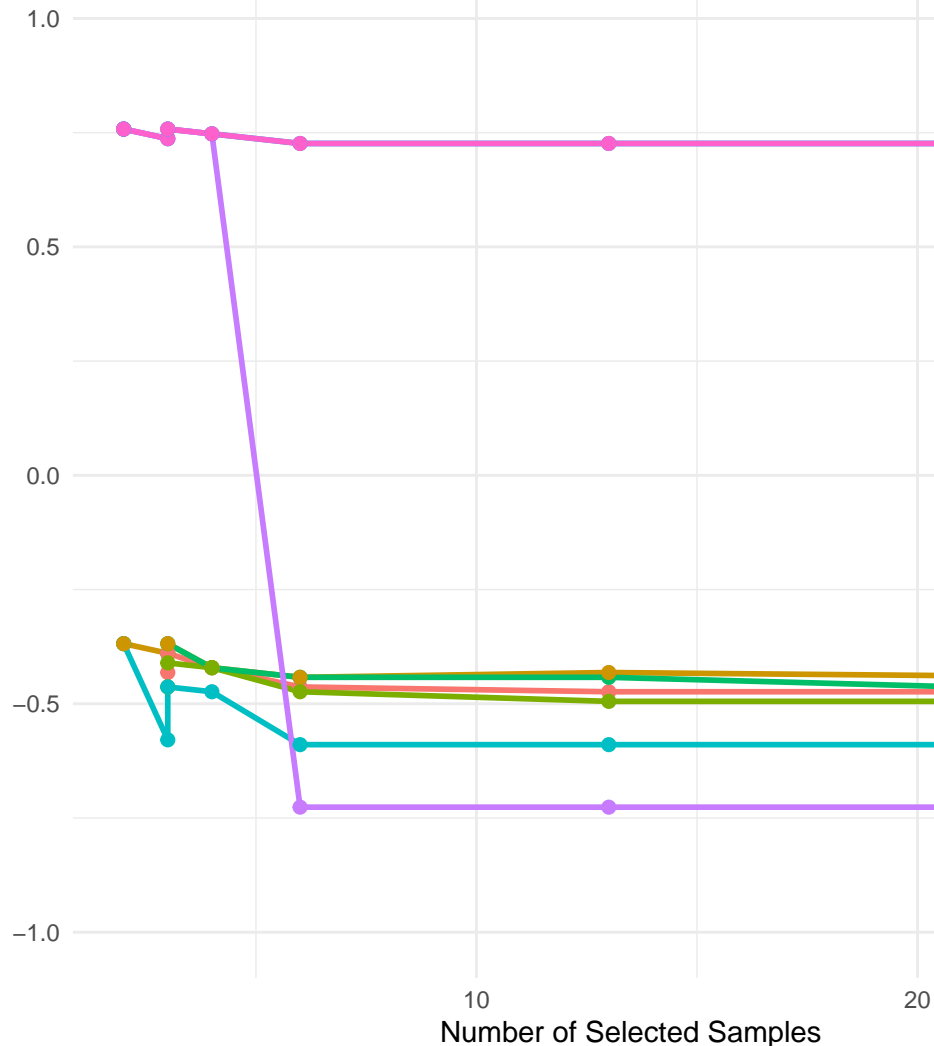
- range
- slope
- variance

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

Kendall Correlation

Summary Statistic

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

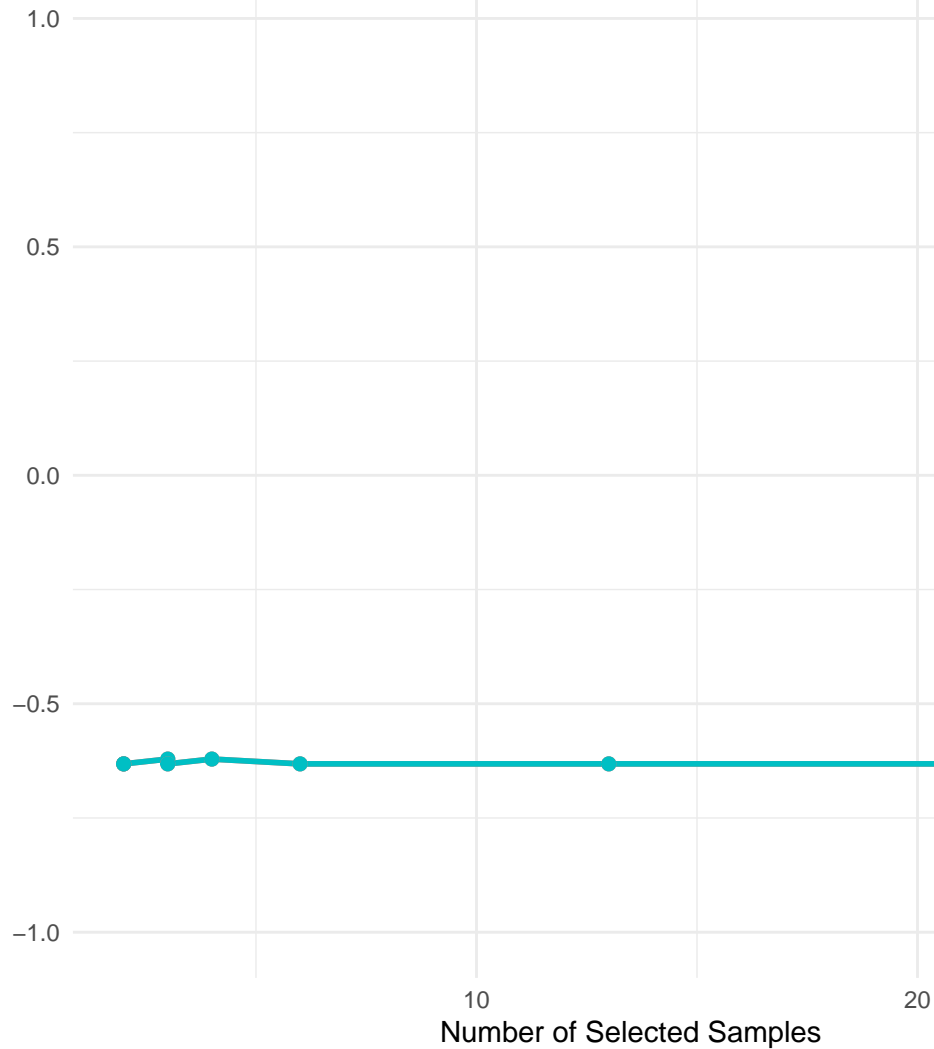


Score: RSI (Local, partial1 , Genotype: linear )

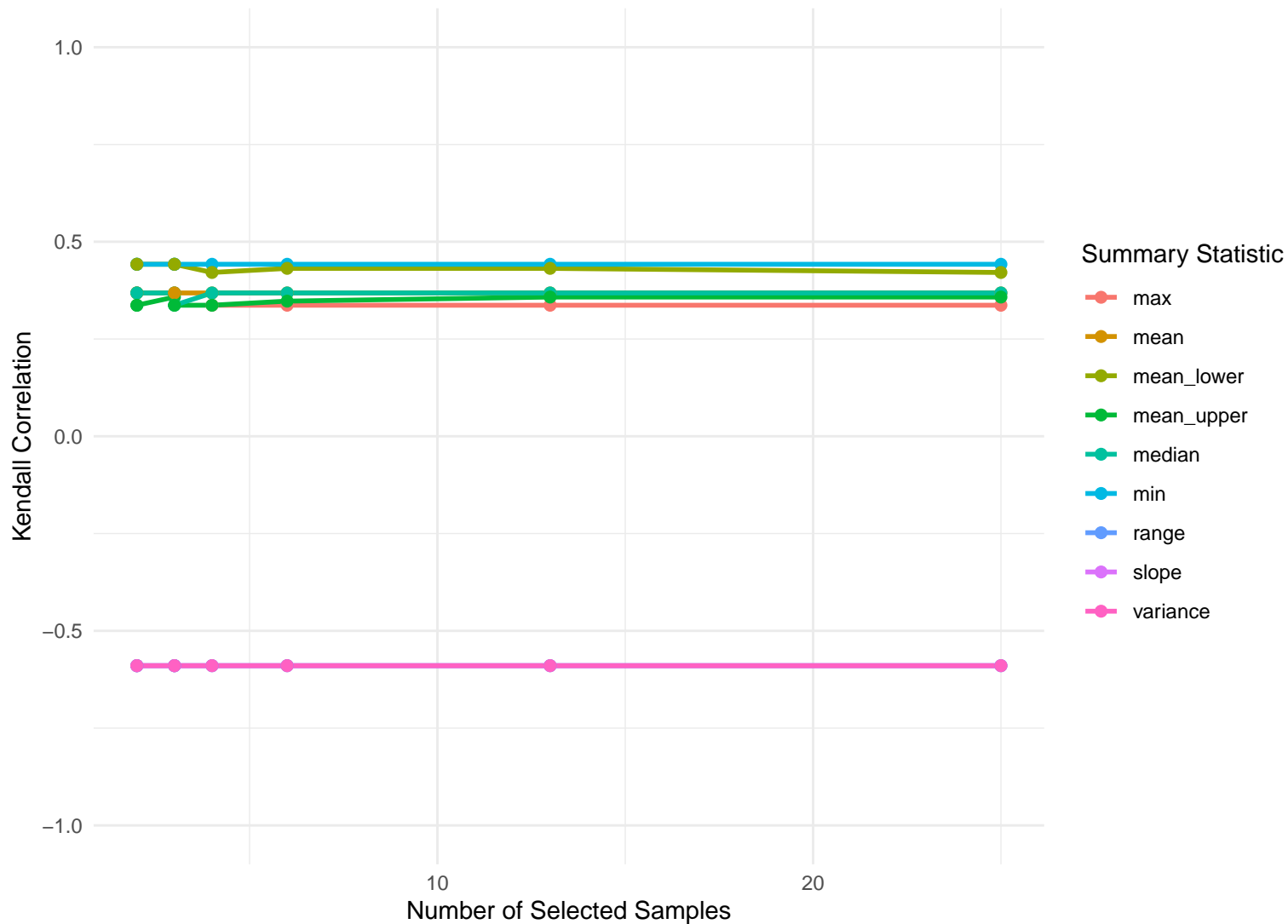
Kendall Correlation

Summary Statistic

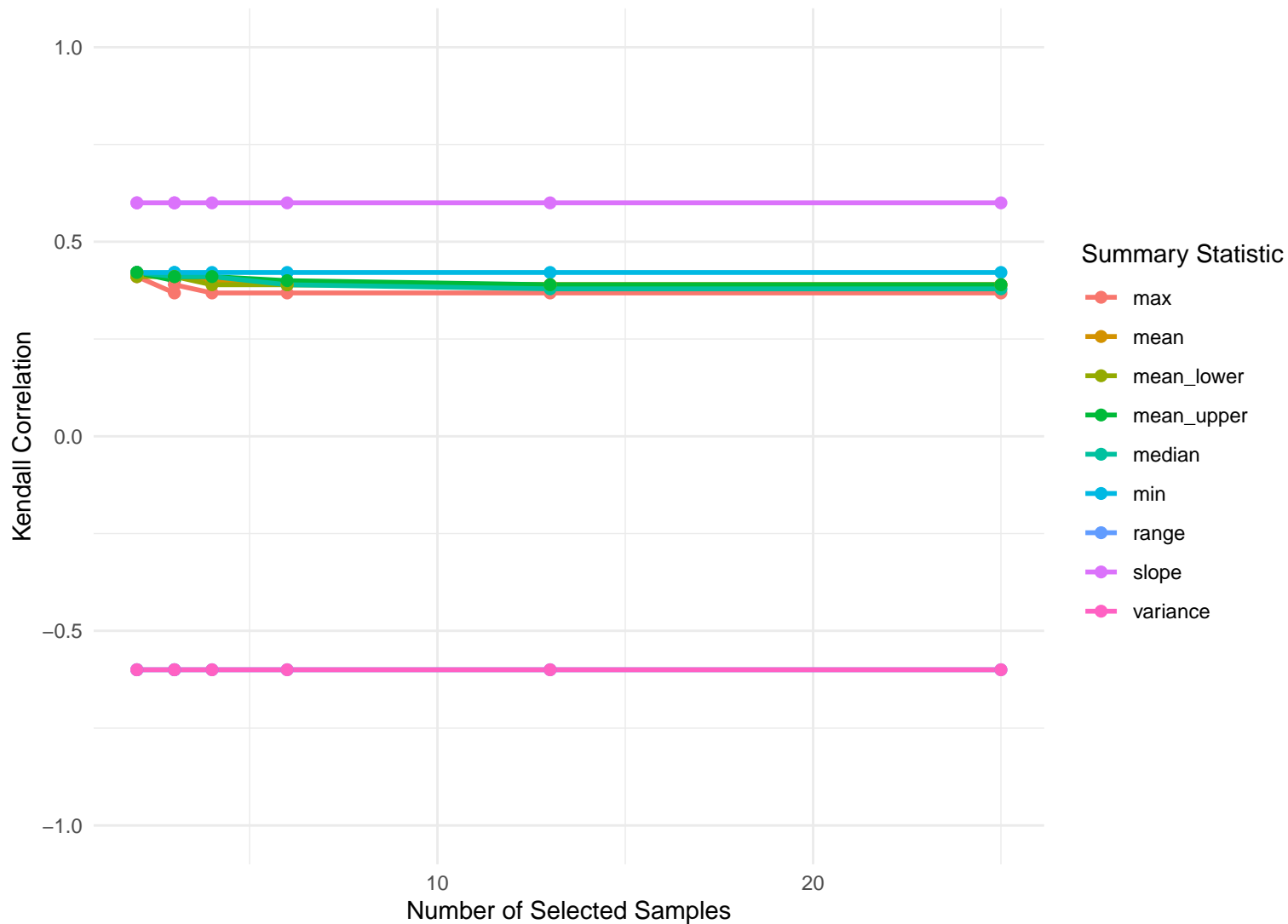
range  
variance



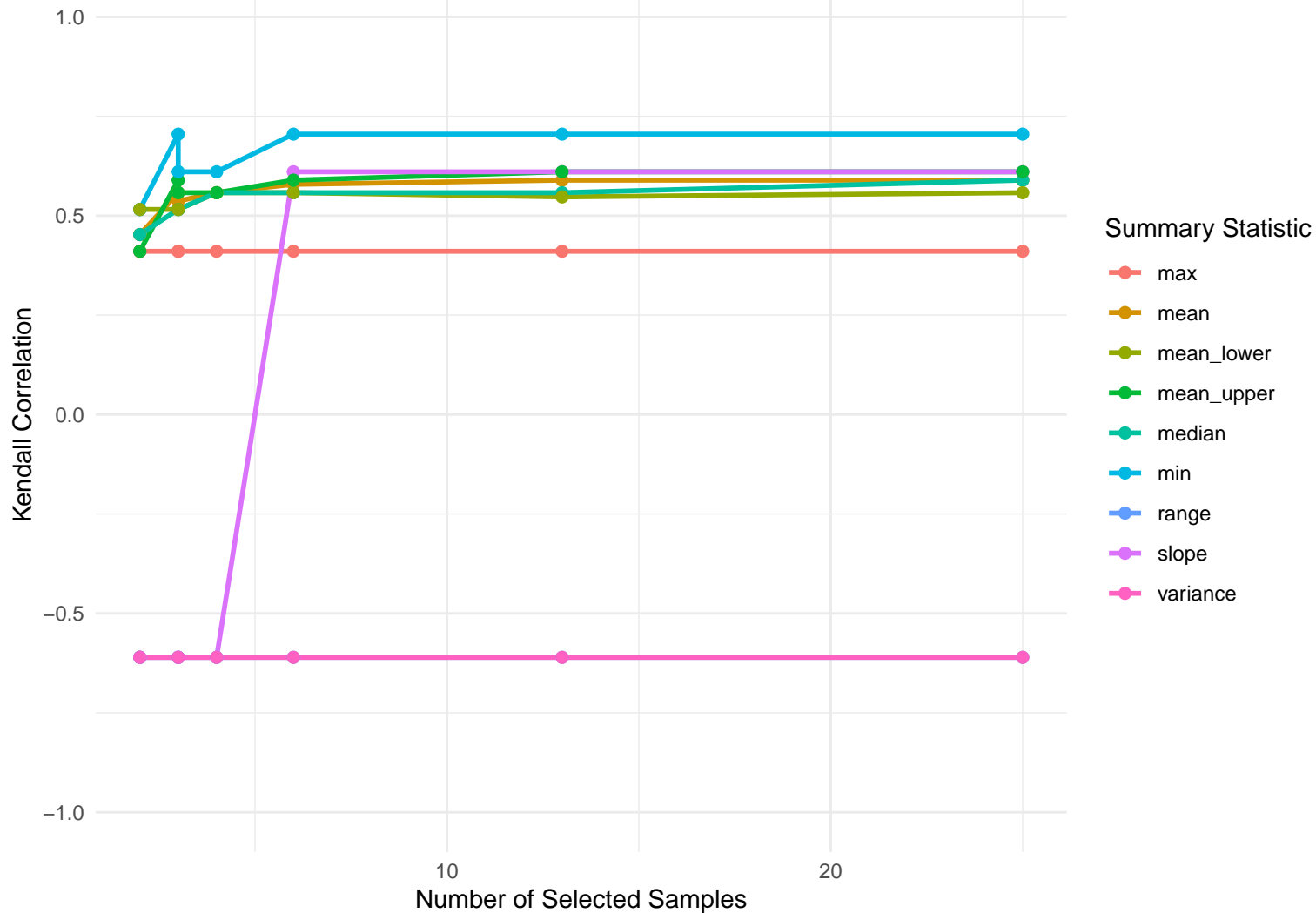
Score: RSI (Local, partial1 , Genotype: gaussian )



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

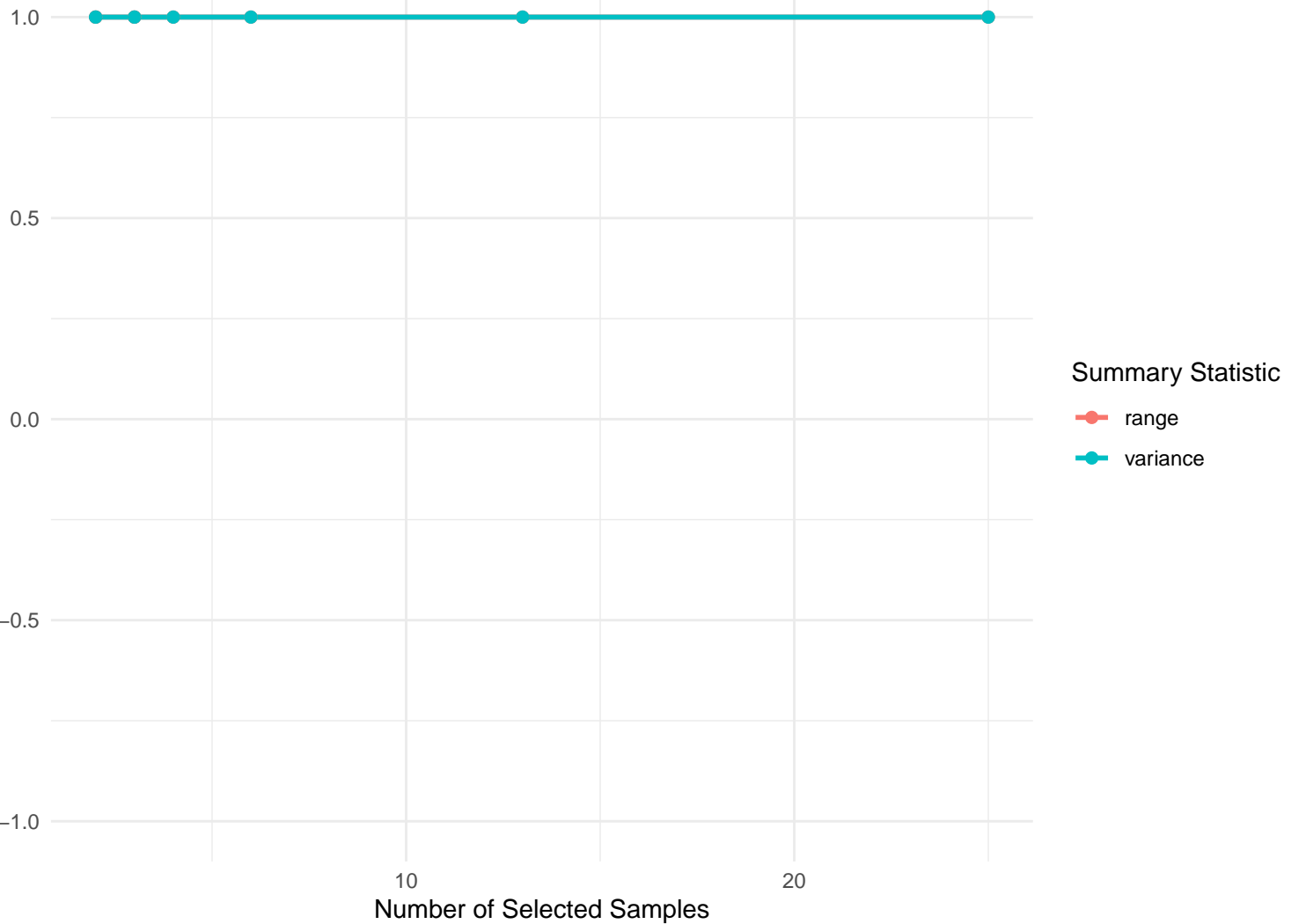
20

Number of Selected Samples

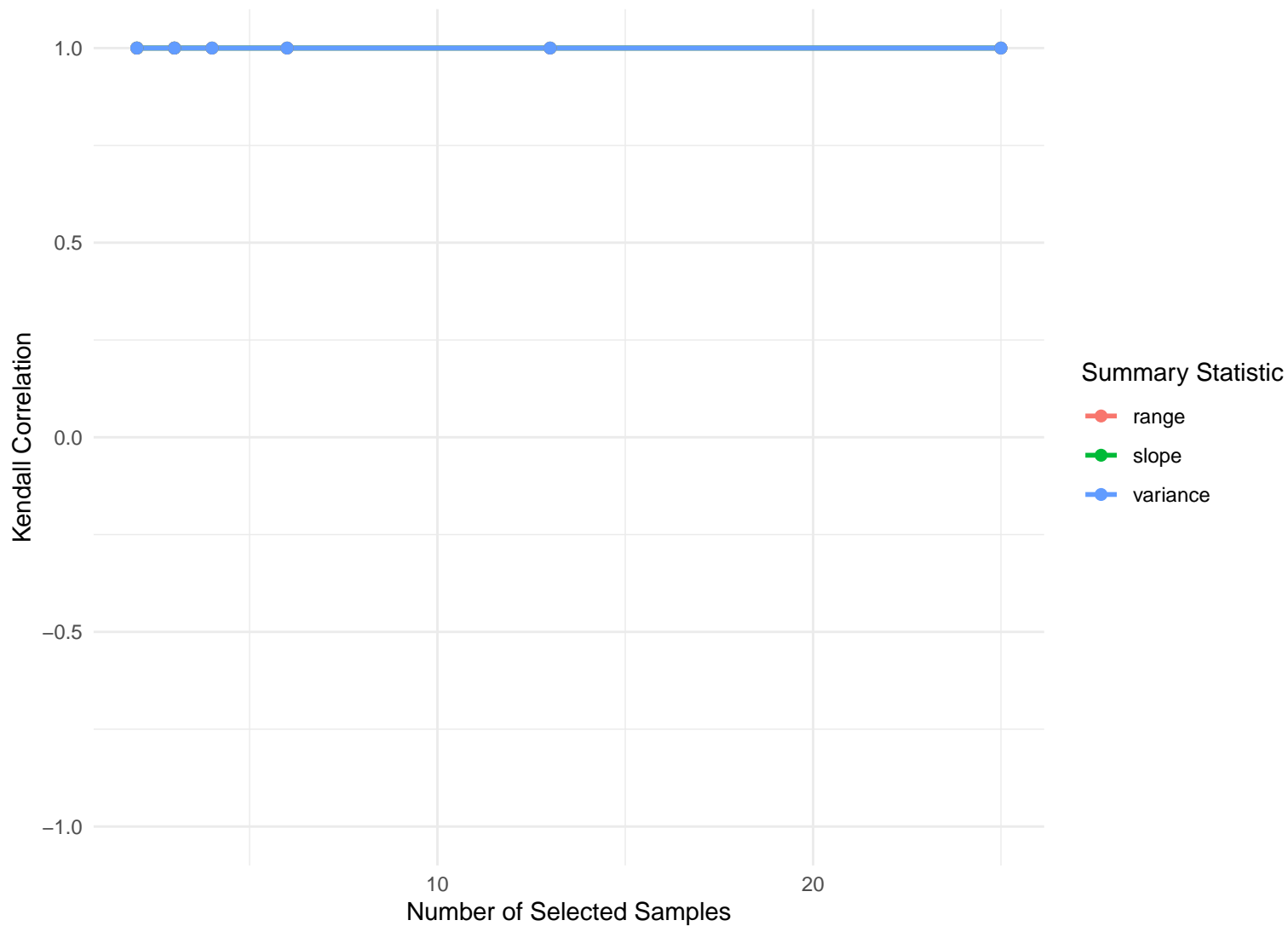
Summary Statistic

range

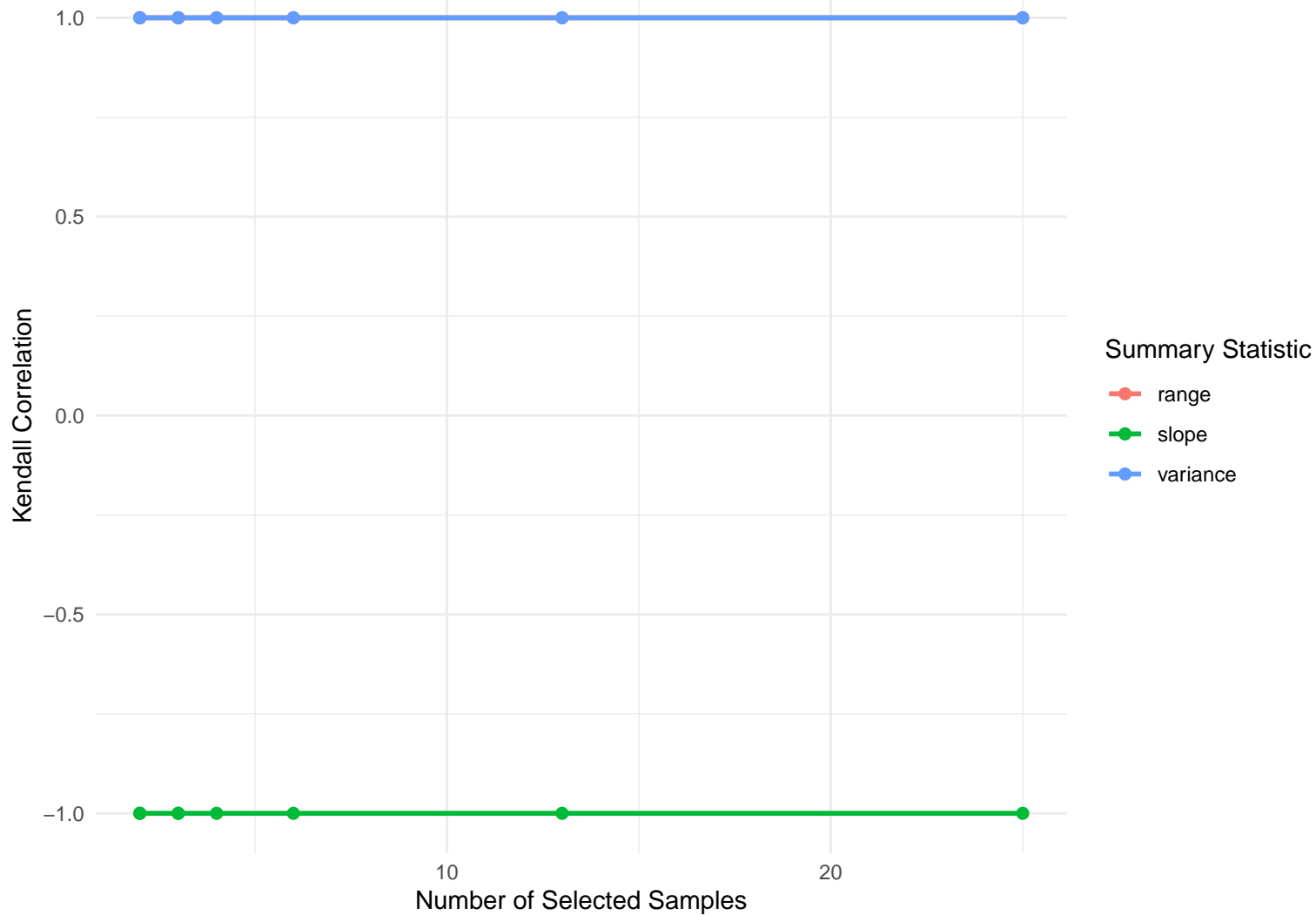
variance



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



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



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

Kendall Correlation

Summary Statistic

- range
- slope
- variance

