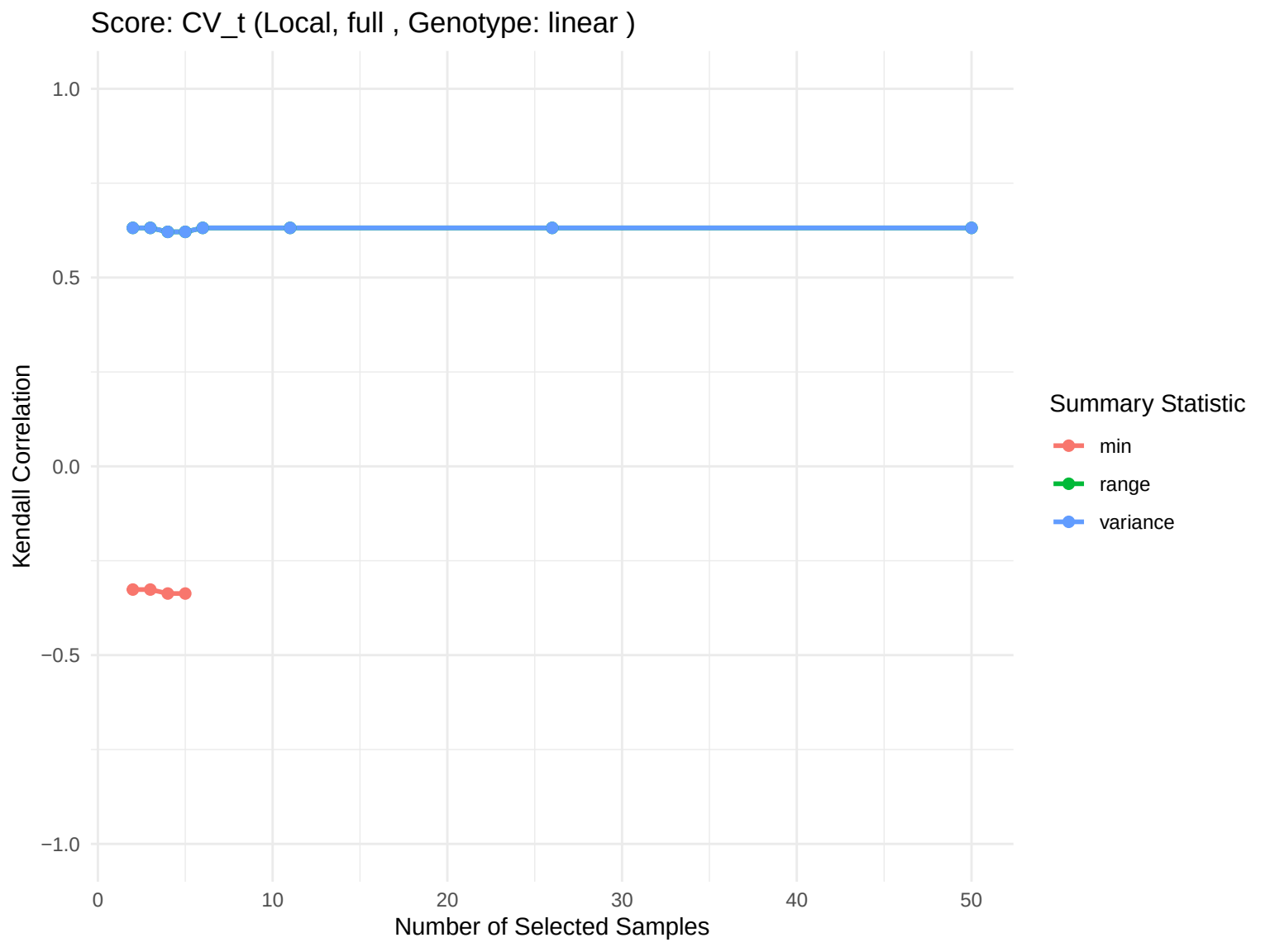
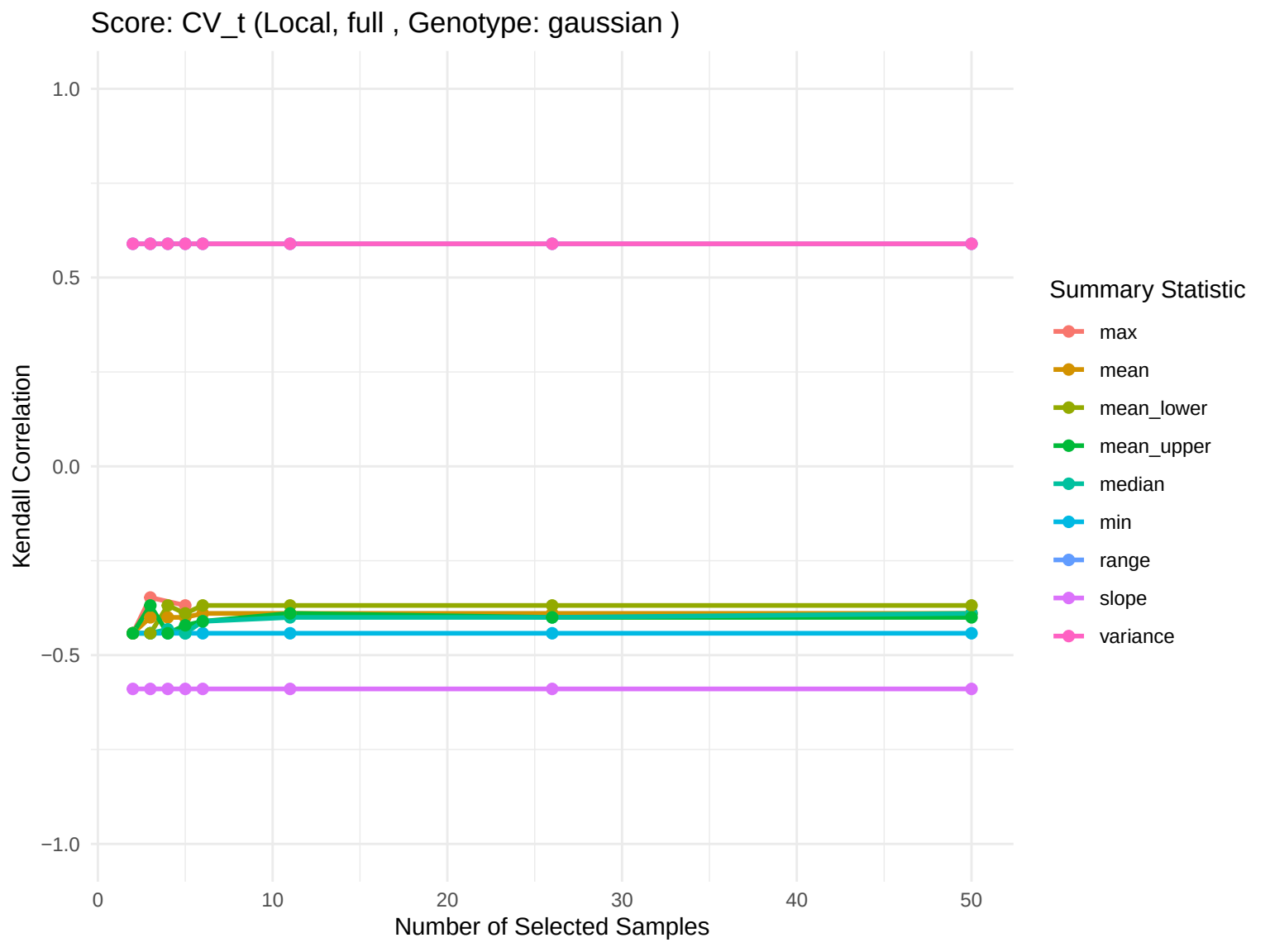

Score: CV_t (Local, full , Genotype: linear)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
min
range
variance


Score: CV_t (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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20
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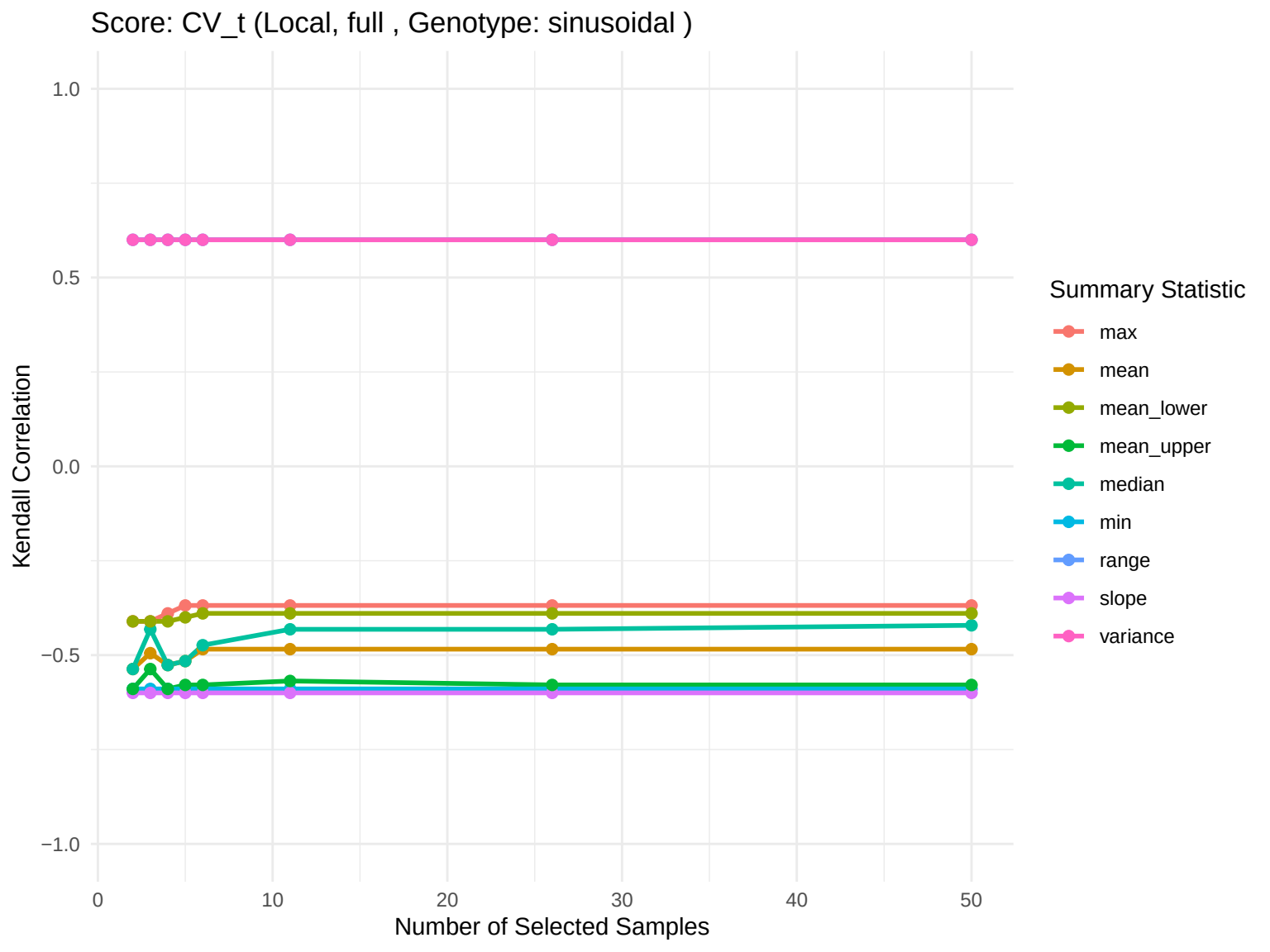
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: CV_t (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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20
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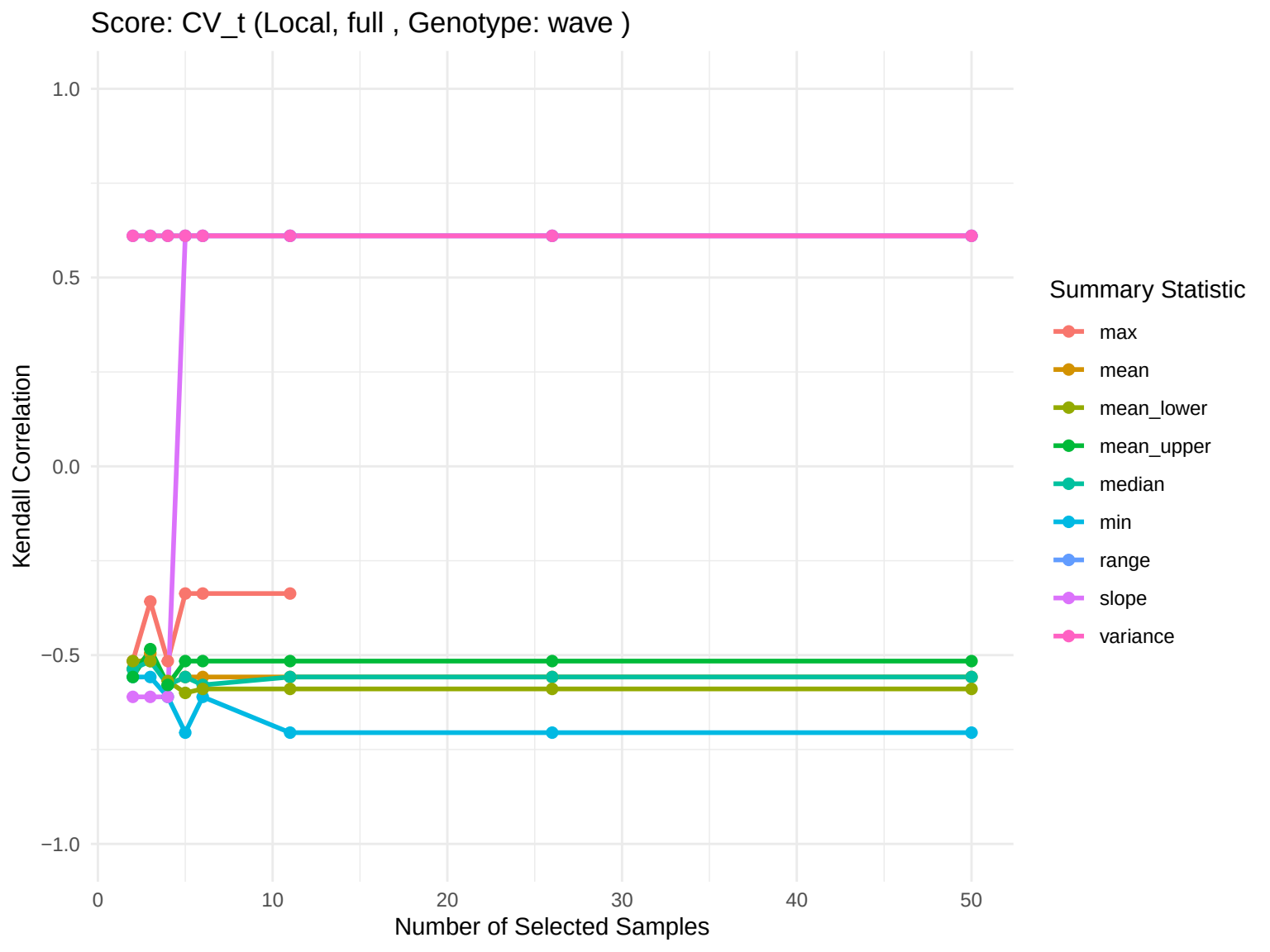
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: CV_t (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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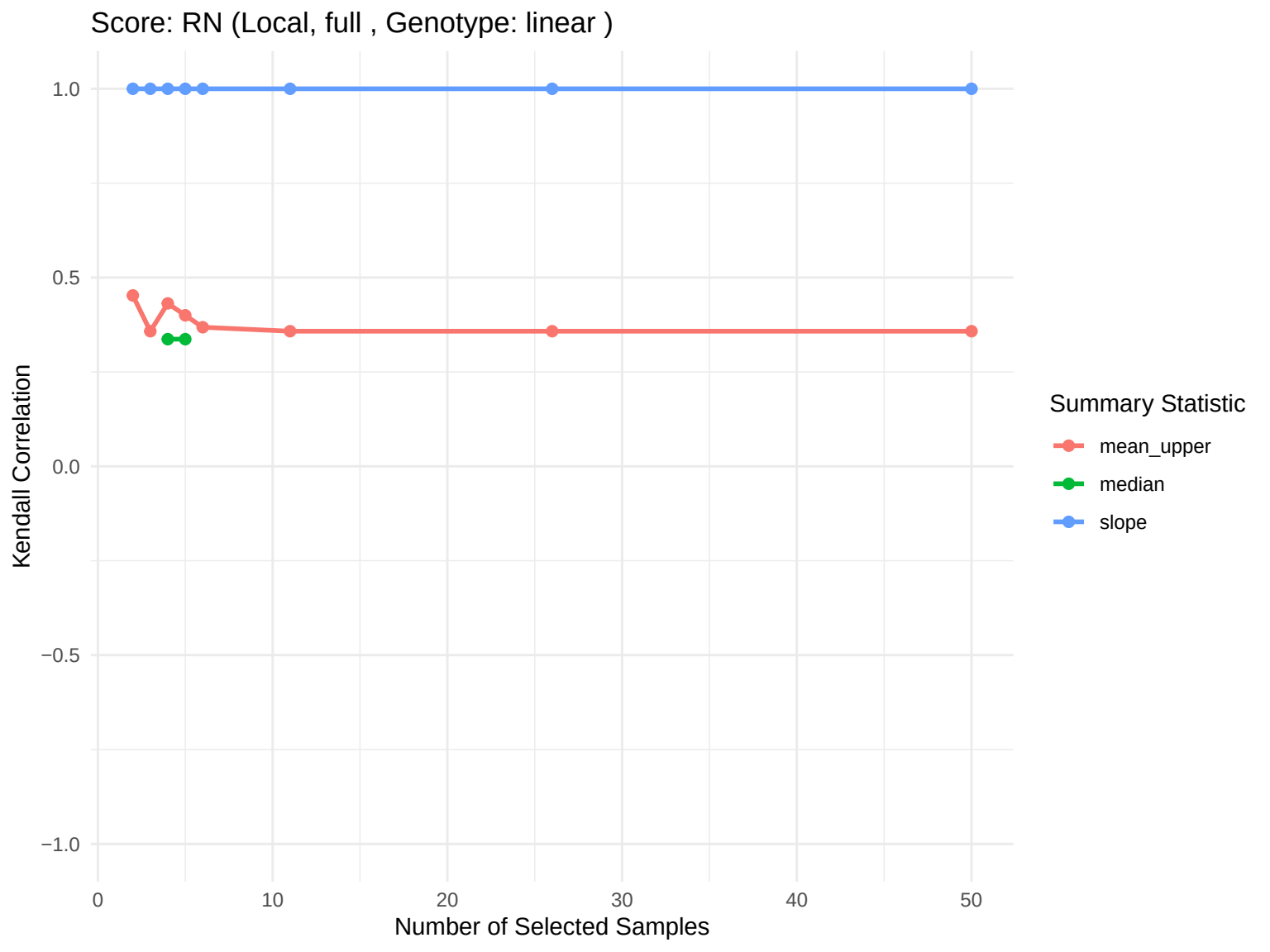
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RN (Local, full , Genotype: linear)
Kendall Correlation
1.0
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−0.5
−1.0
0
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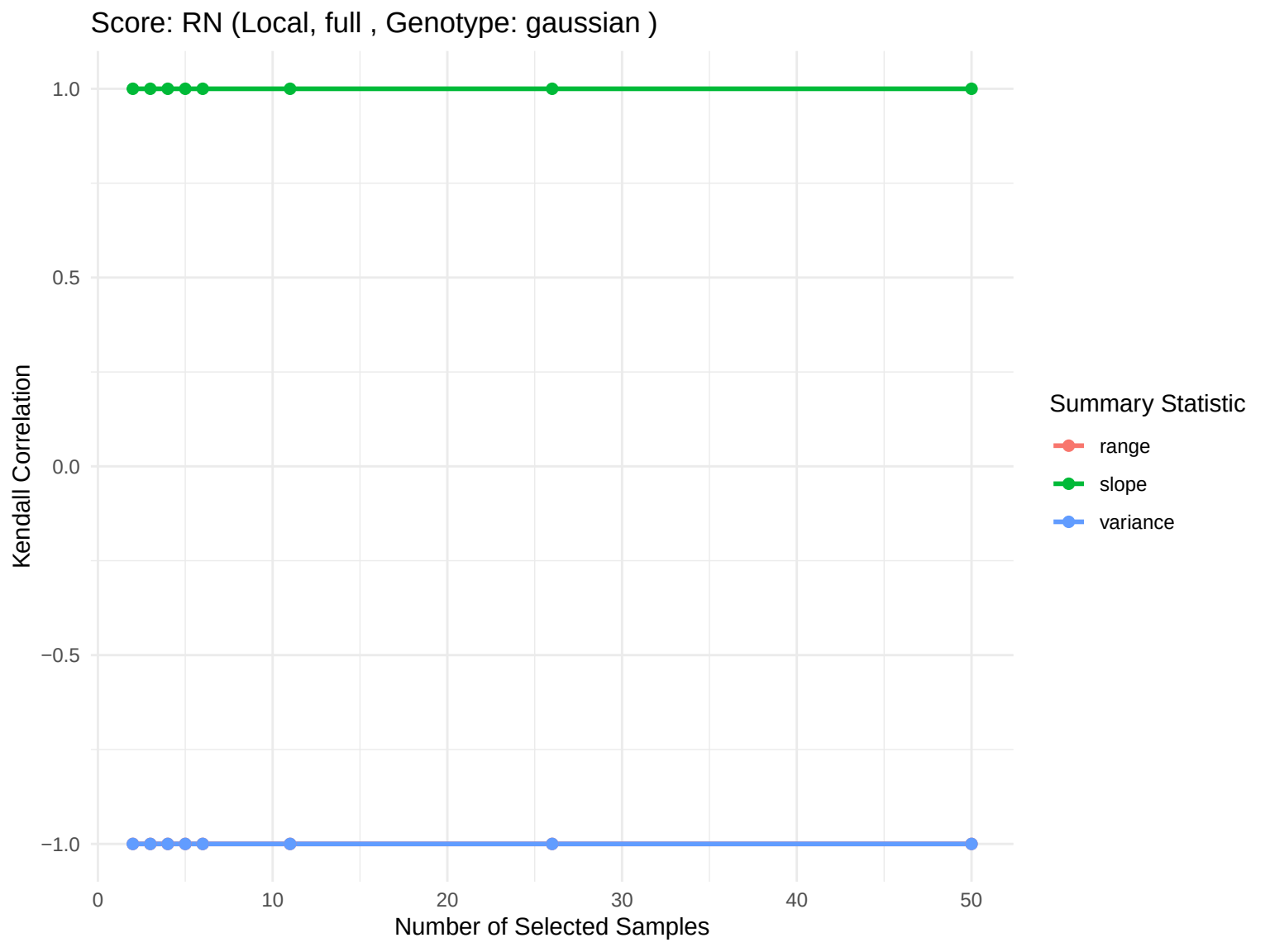
20
30
40
50
Number of Selected Samples
Summary Statistic
mean_upper
median
slope


Score: RN (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
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0.0
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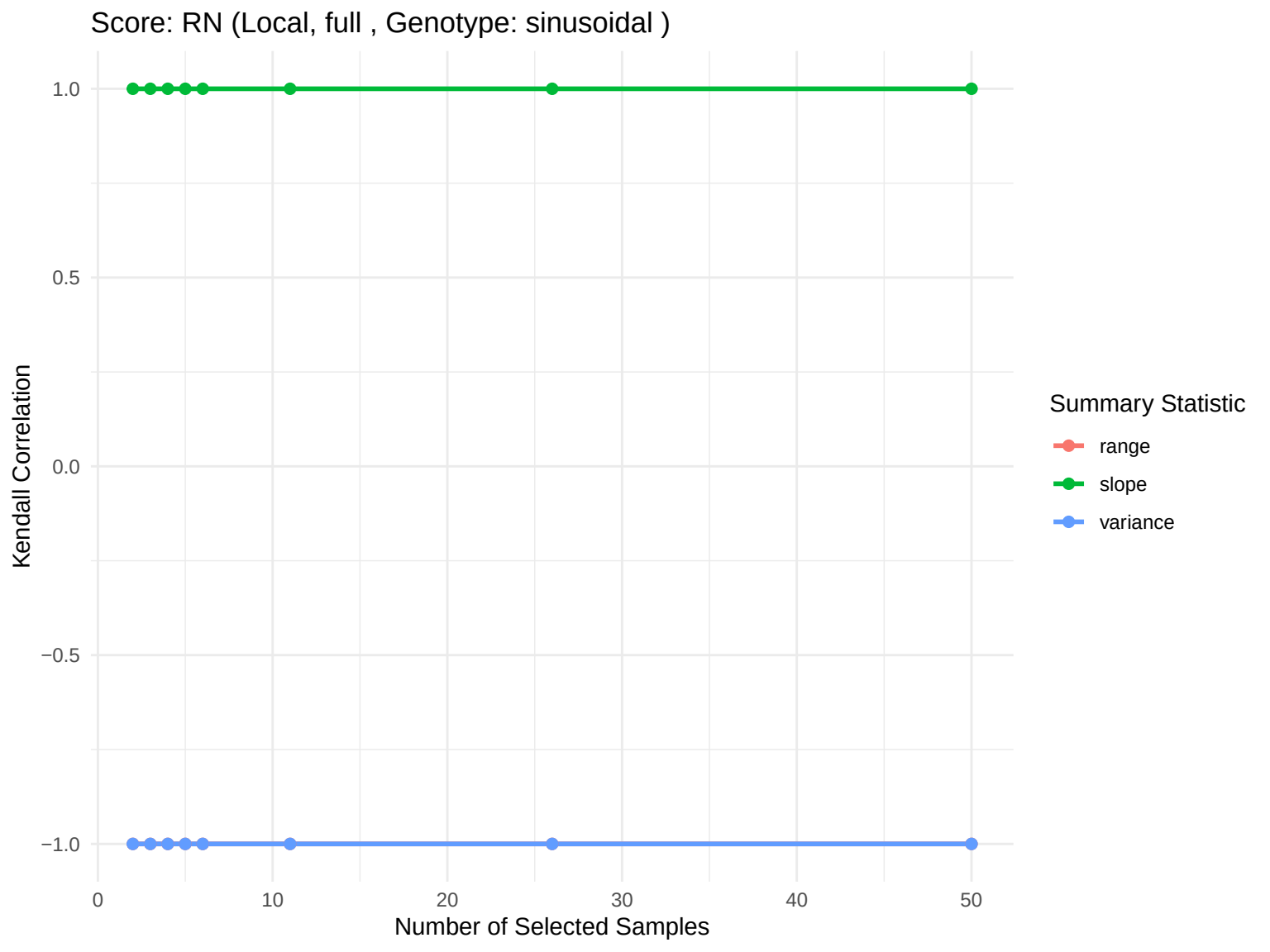
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: RN (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
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−1.0
0
10
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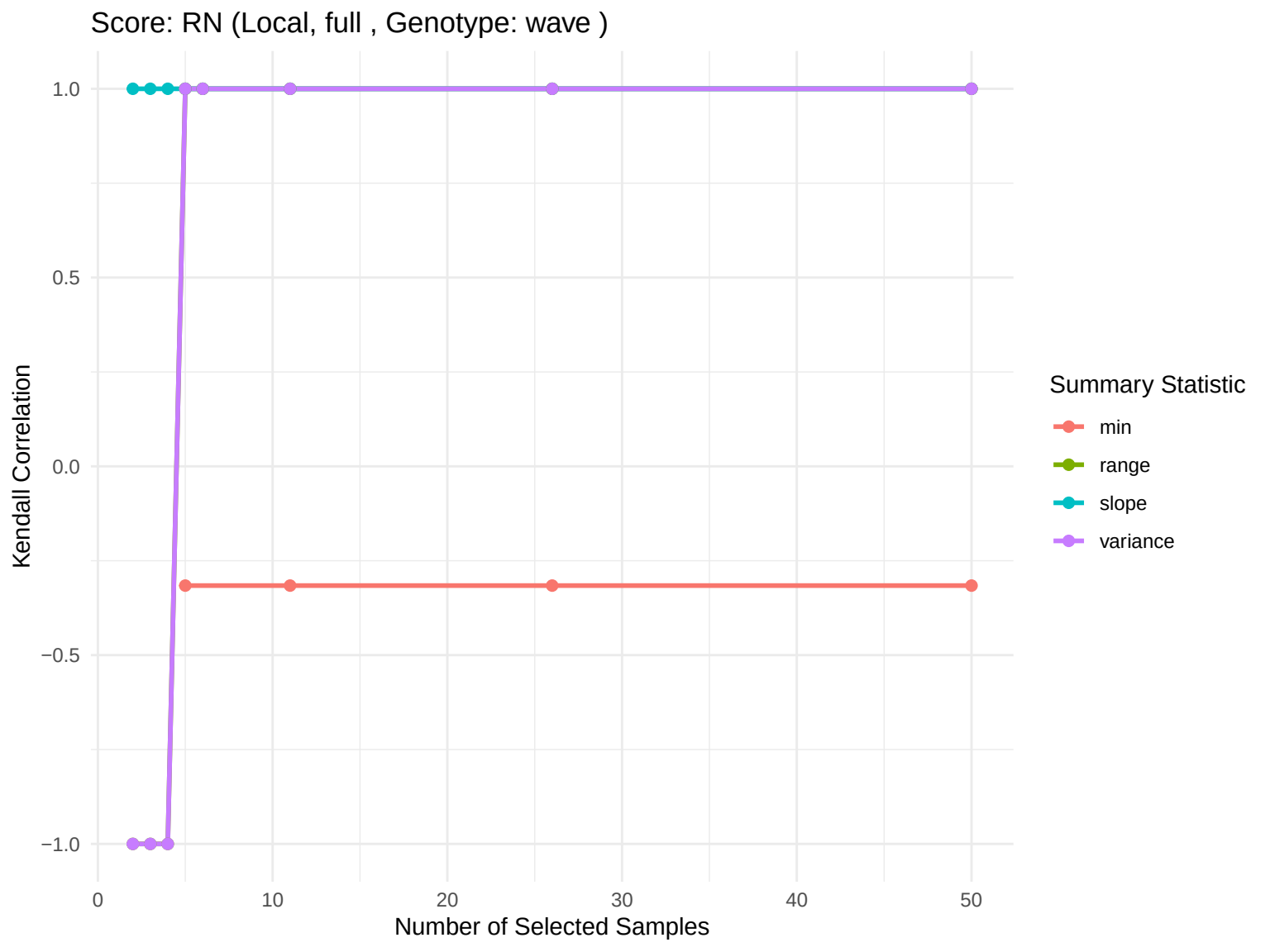
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: RN (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
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−1.0
0
10
20
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40
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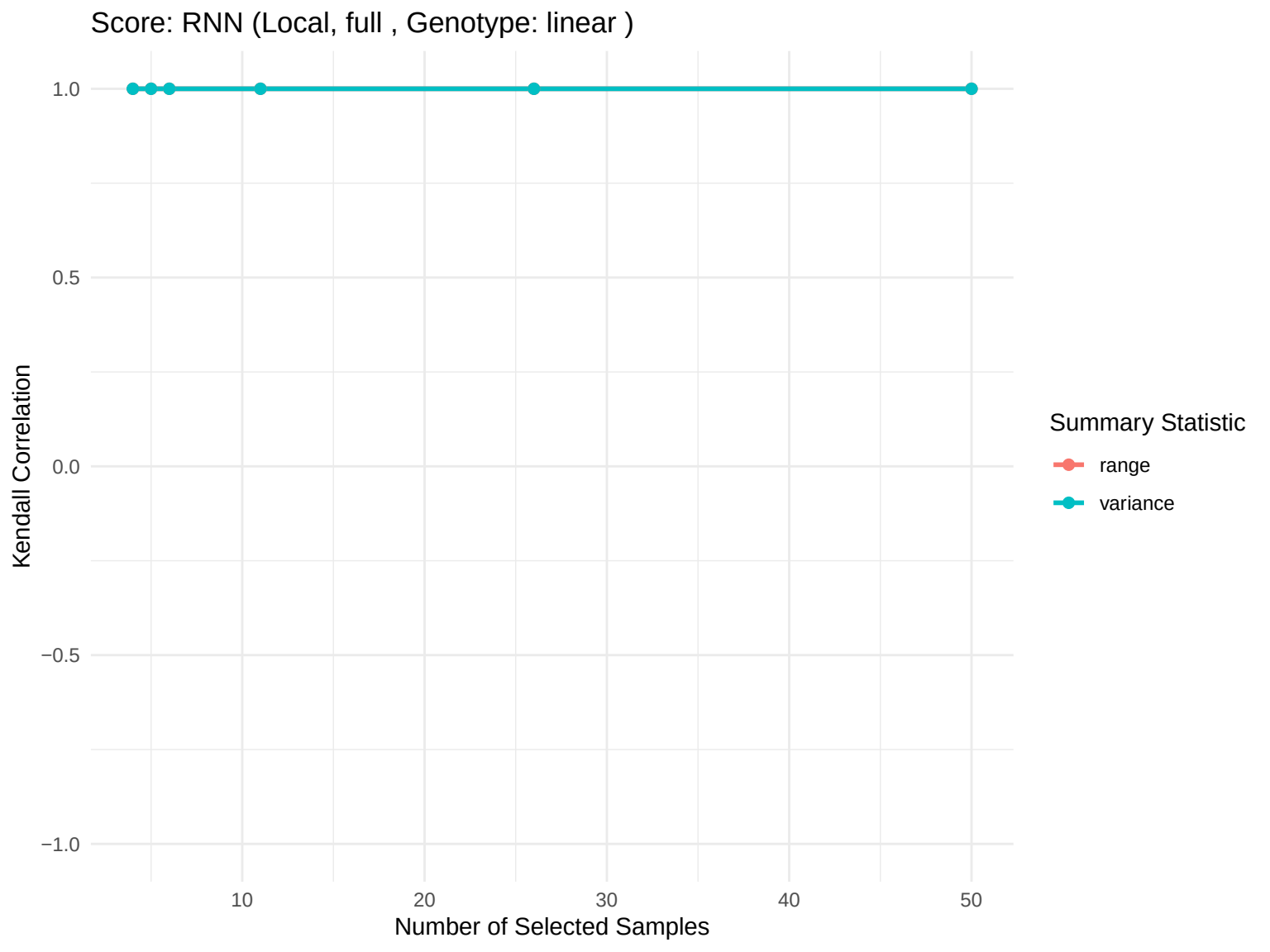
Number of Selected Samples
Summary Statistic
min
range
slope
variance


Score: RNN (Local, full , Genotype: linear)
1.0
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0.0
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−1.0
Kendall Correlation
10
20
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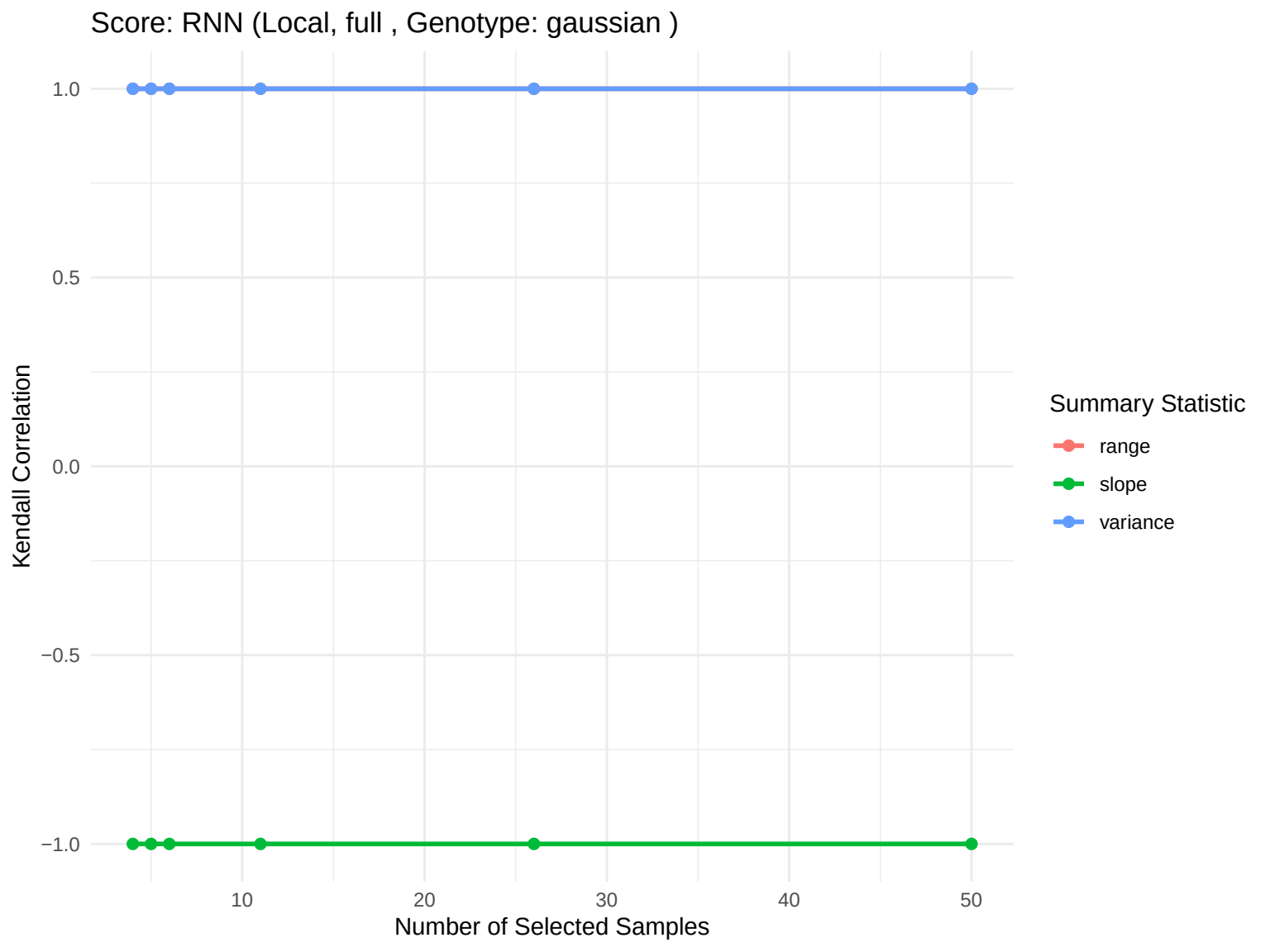
40
50
Number of Selected Samples
Summary Statistic
range
variance


Score: RNN (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
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−1.0
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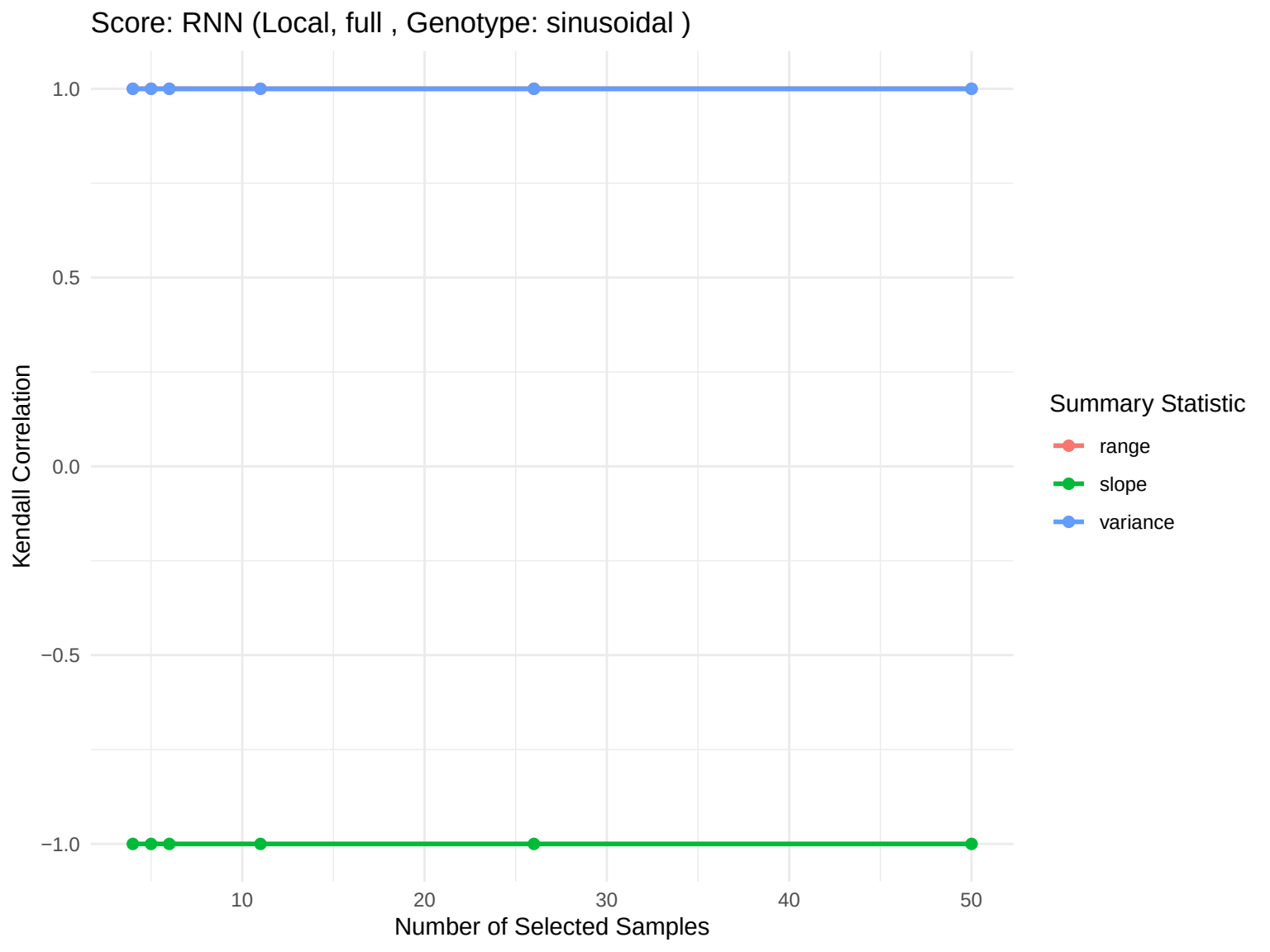
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: RNN (Local, full , Genotype: sinusoidal)
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0.5
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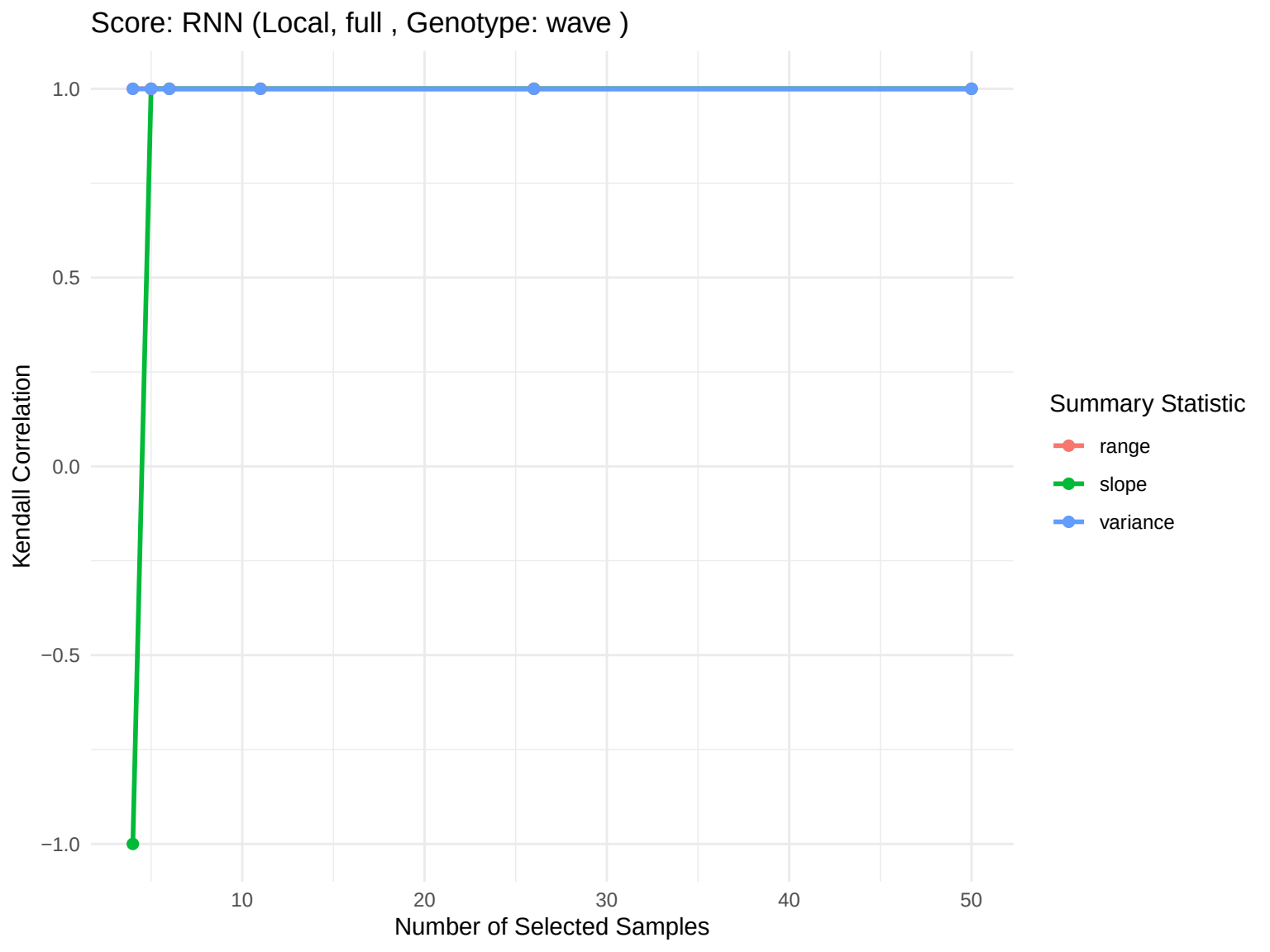
−0.5
−1.0
Kendall Correlation
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: RNN (Local, full , Genotype: wave)
Kendall Correlation
1.0
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−0.5
−1.0
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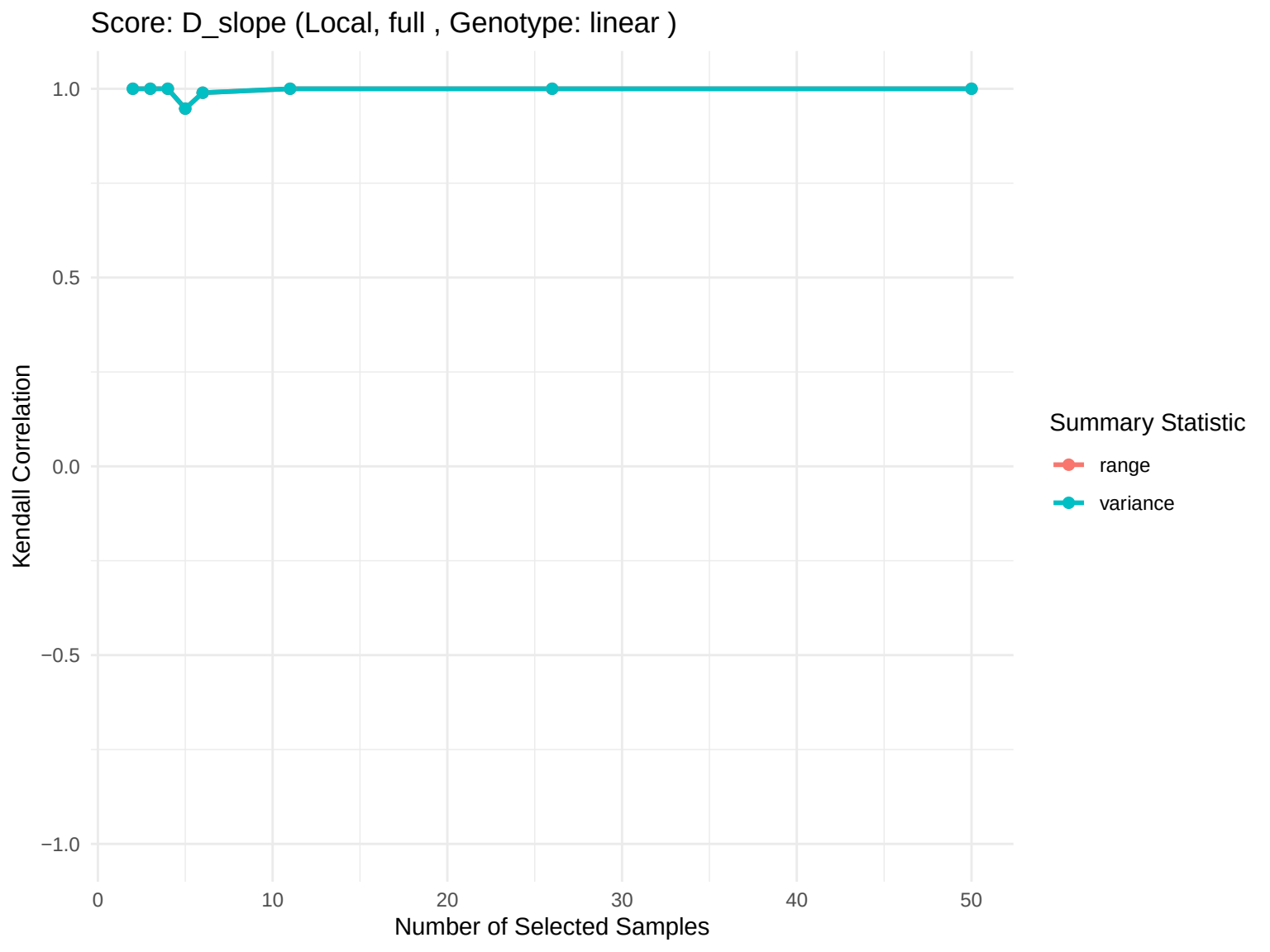
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: D_slope (Local, full , Genotype: linear)
Kendall Correlation
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−0.5
−1.0
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40
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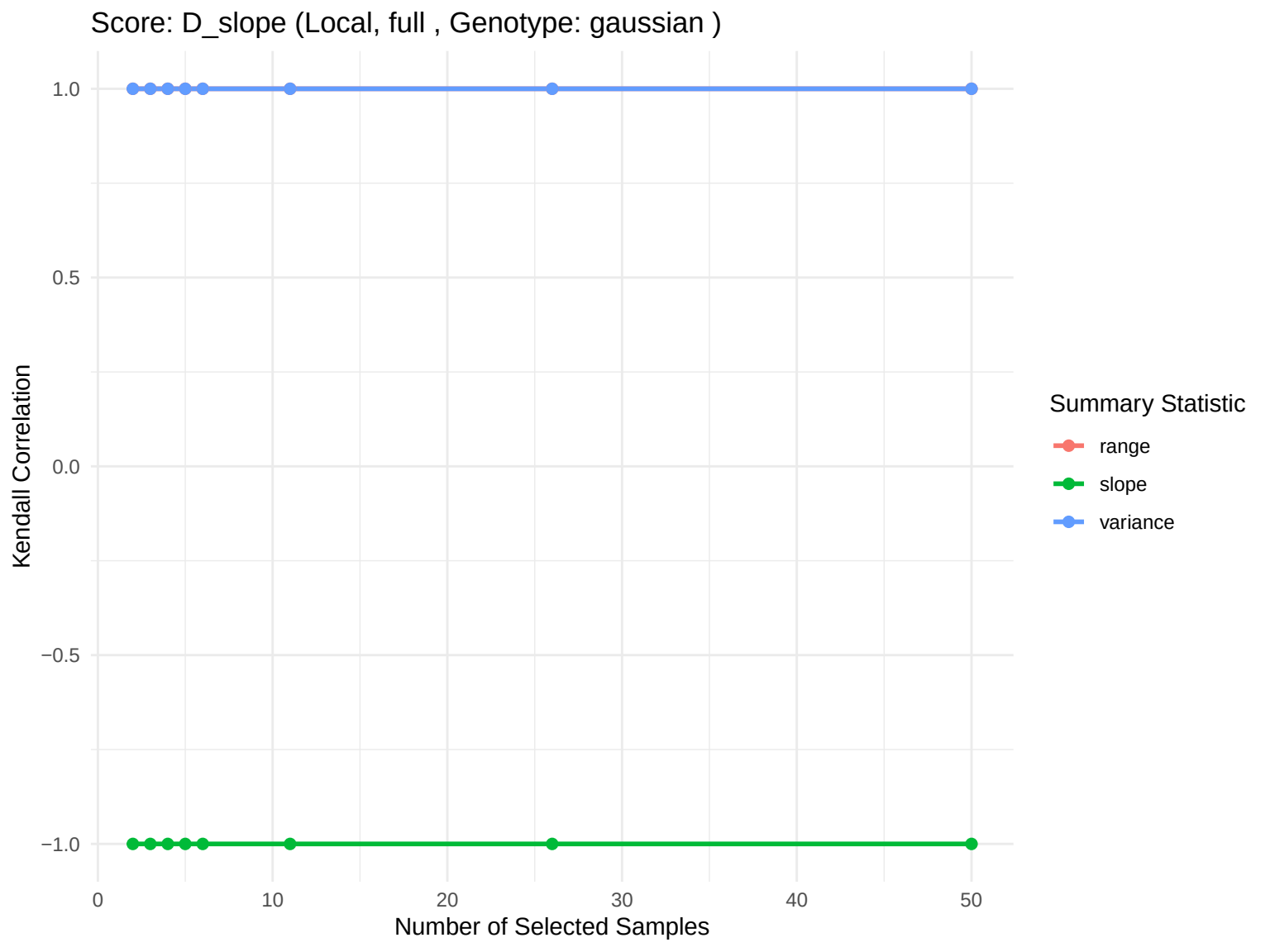
Number of Selected Samples
Summary Statistic
range
variance


Score: D_slope (Local, full , Genotype: gaussian)
Kendall Correlation
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−1.0
0
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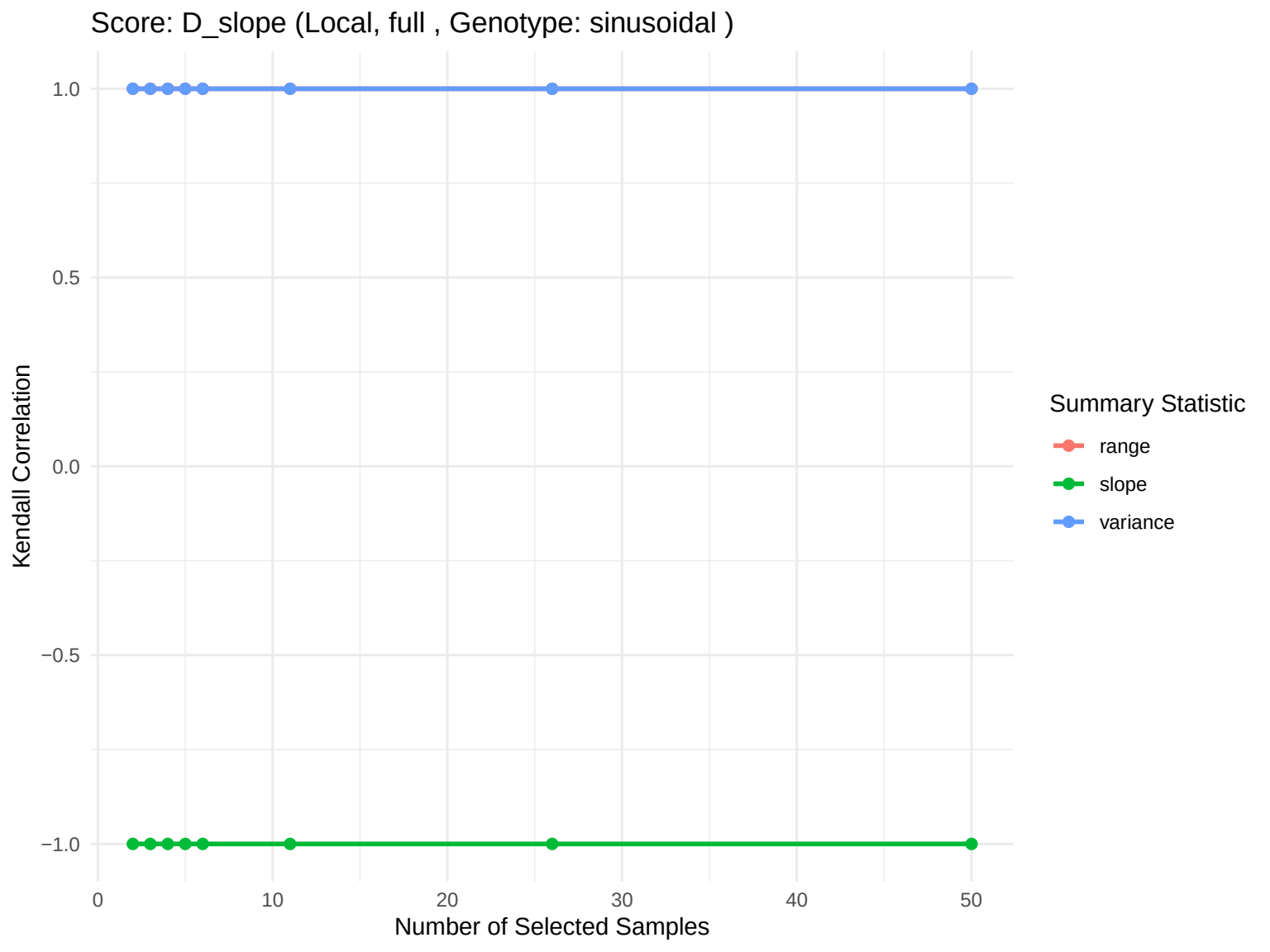
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: D_slope (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
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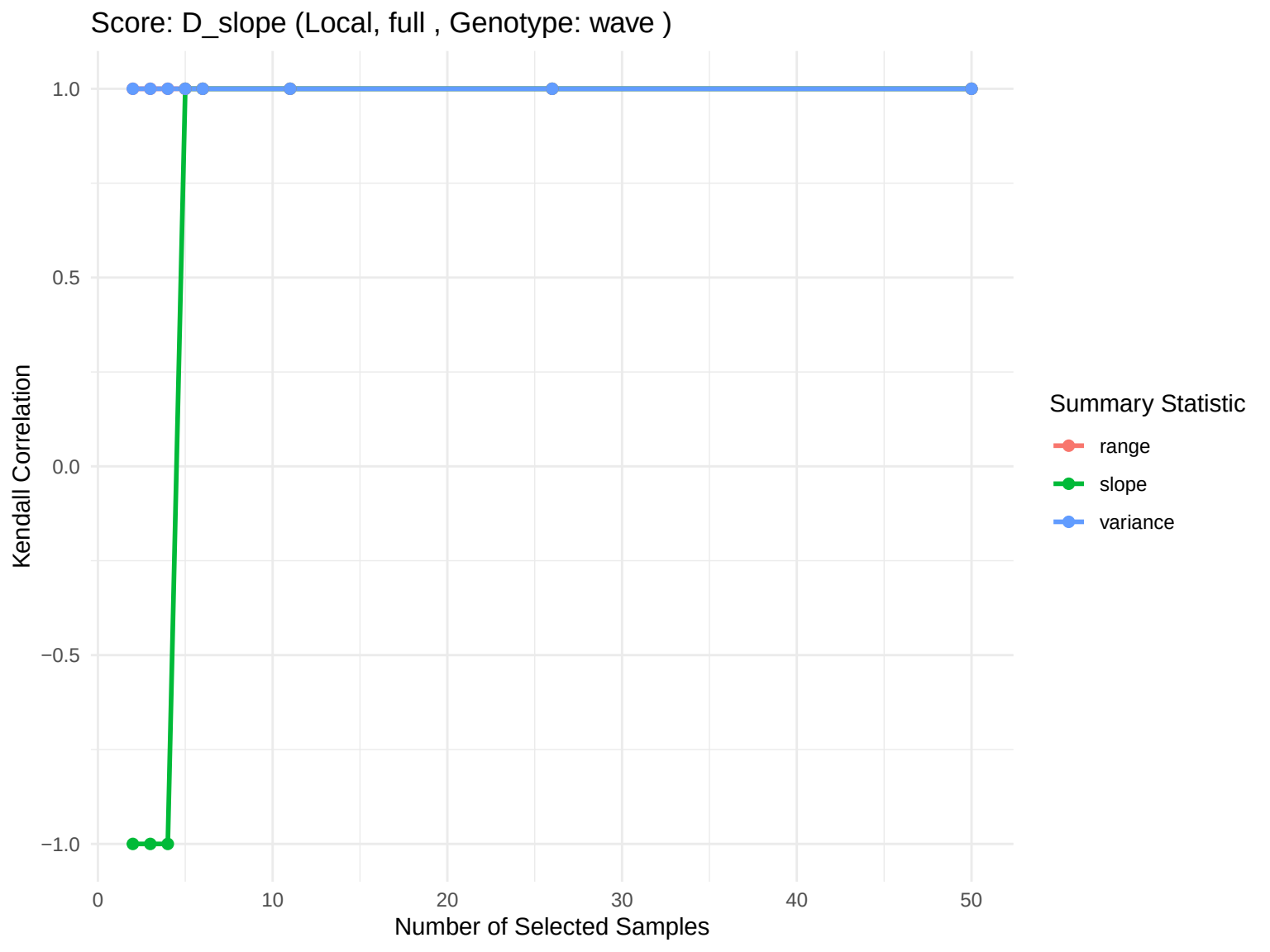
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: D_slope (Local, full , Genotype: wave)
Kendall Correlation
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−1.0
0
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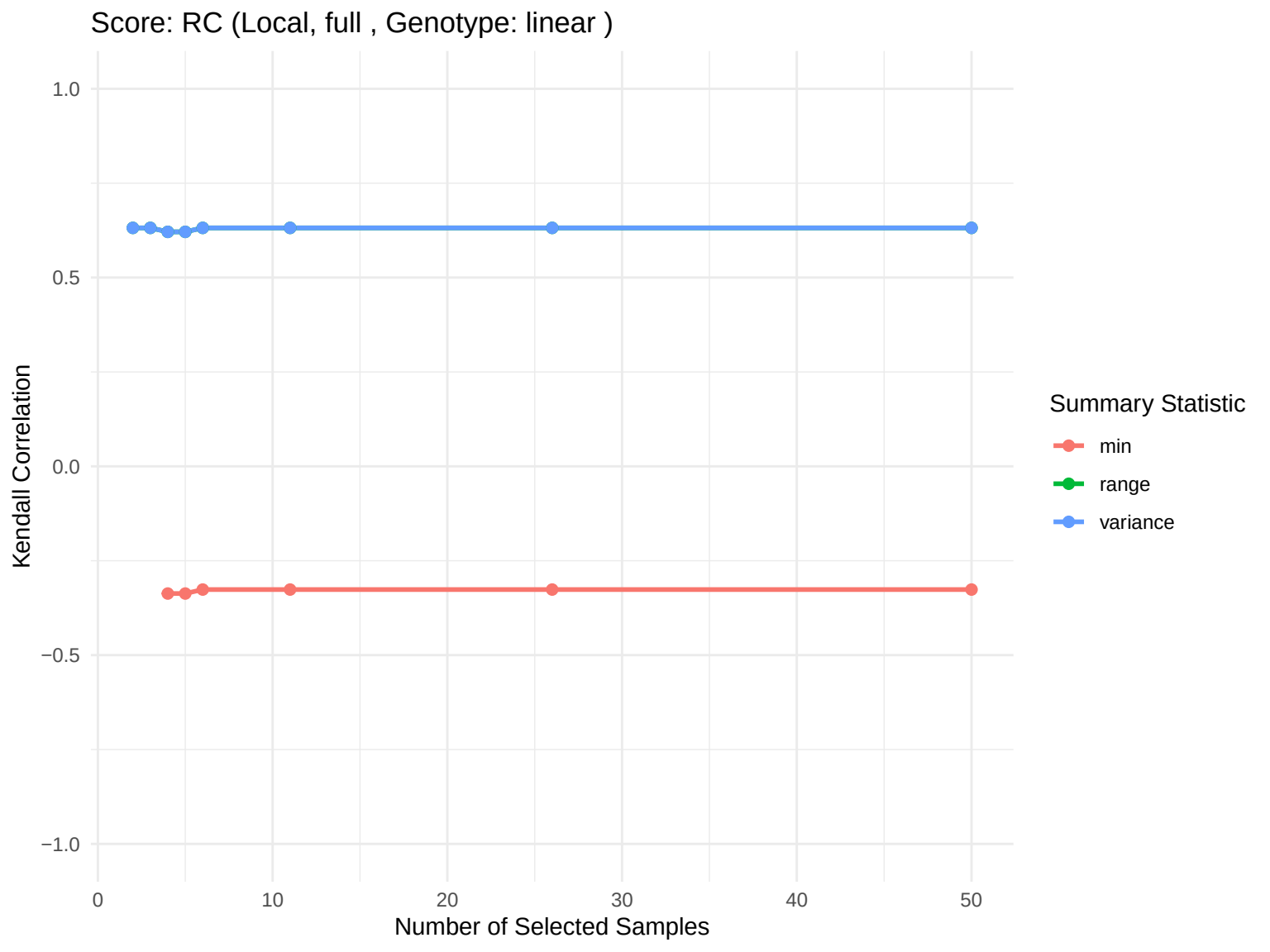
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: RC (Local, full , Genotype: linear)
Kendall Correlation
1.0
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−1.0
0
10
20
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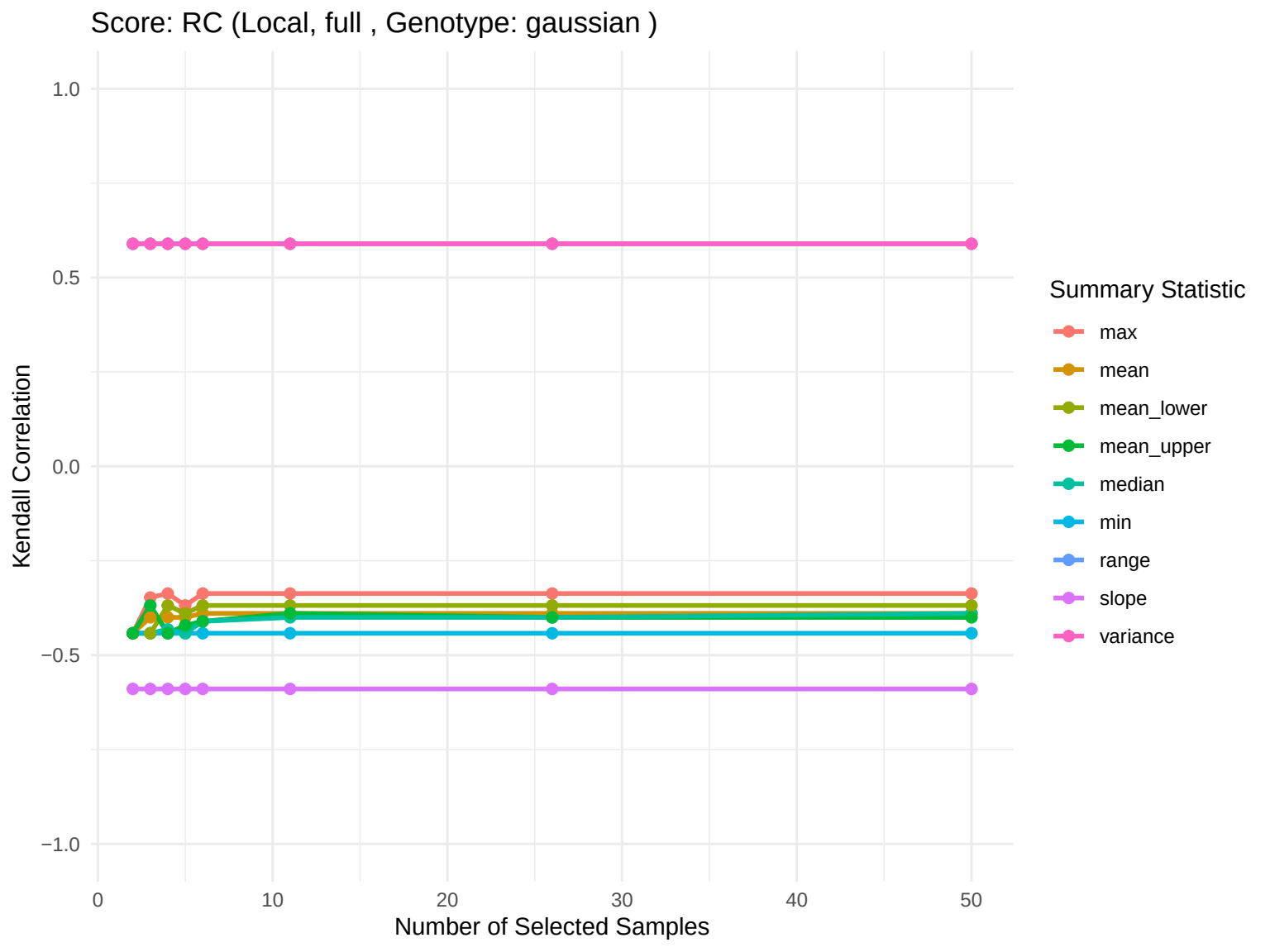
50
Number of Selected Samples
Summary Statistic
min
range
variance


Score: RC (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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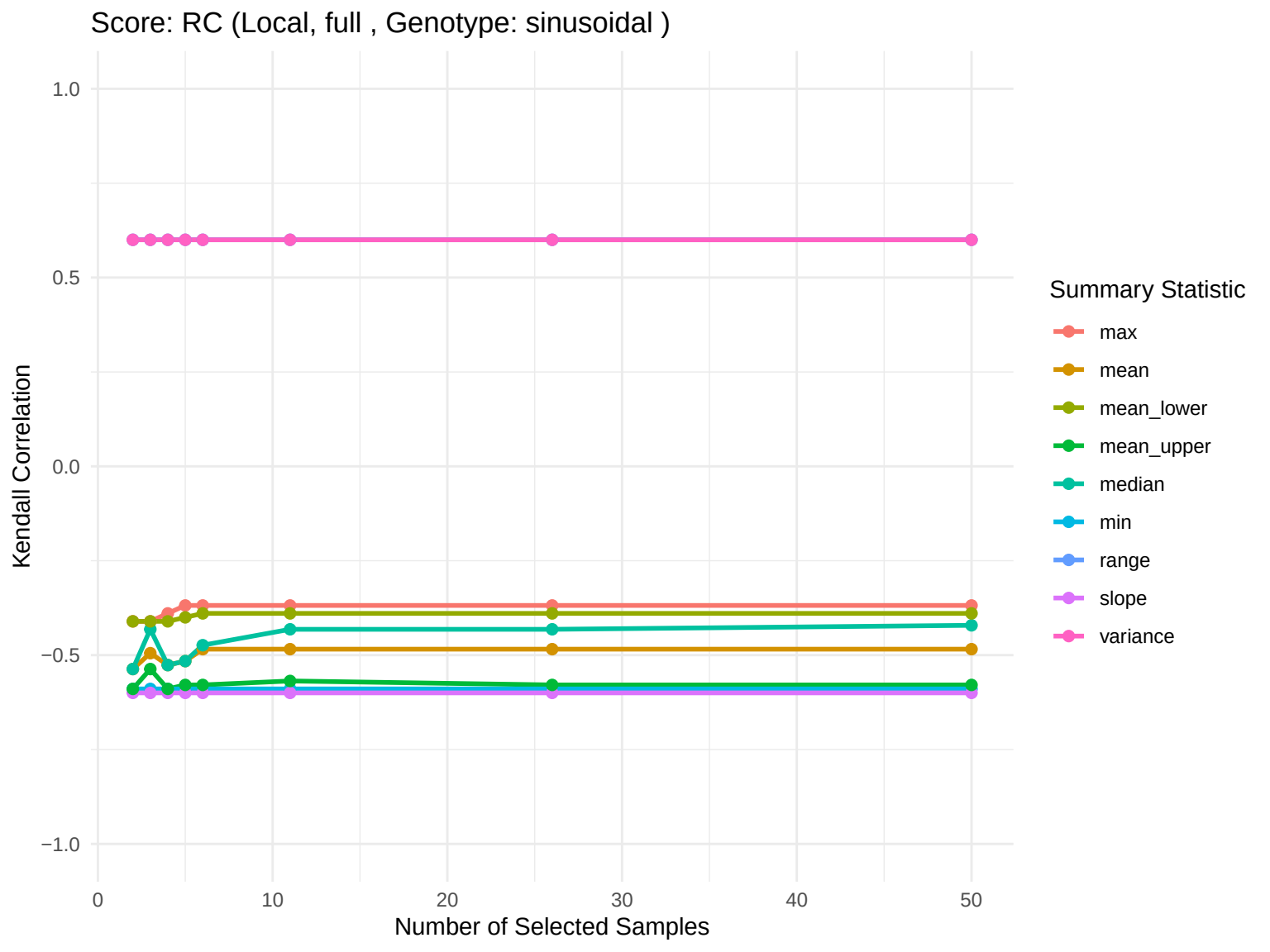
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RC (Local, full , Genotype: sinusoidal)
Kendall Correlation
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0.5
0.0
−0.5
−1.0
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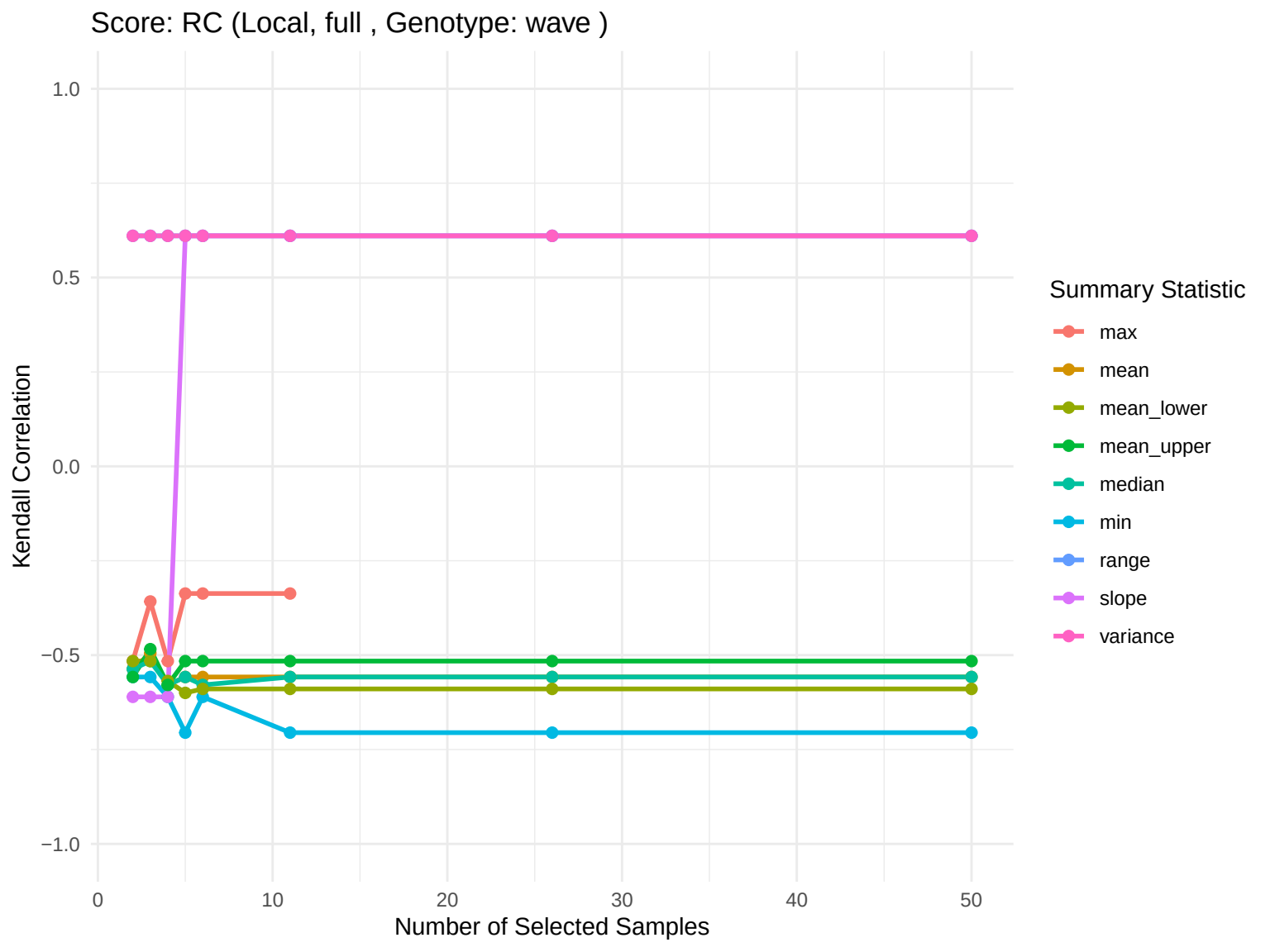
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RC (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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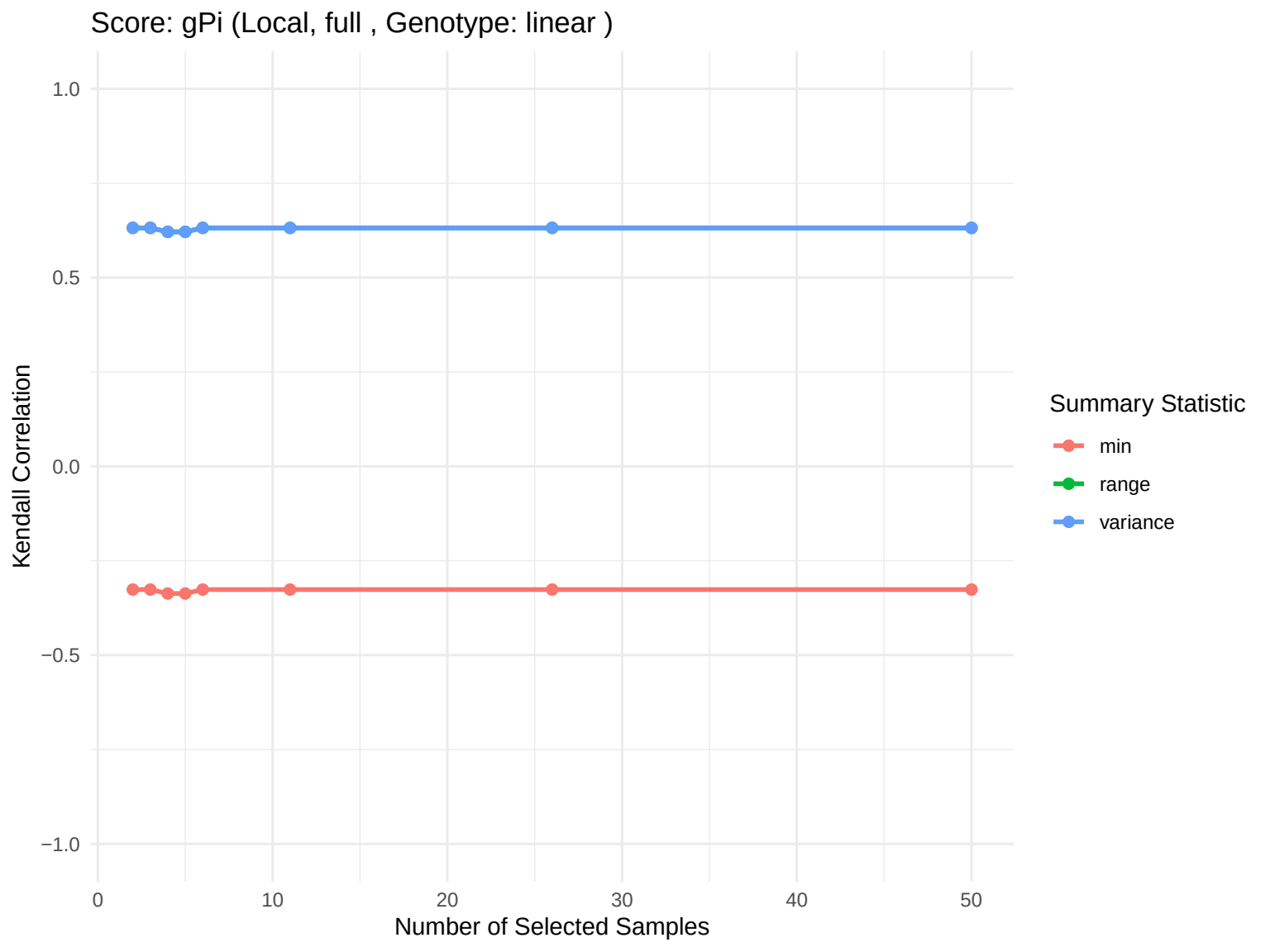
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: gPi (Local, full , Genotype: linear)
Kendall Correlation
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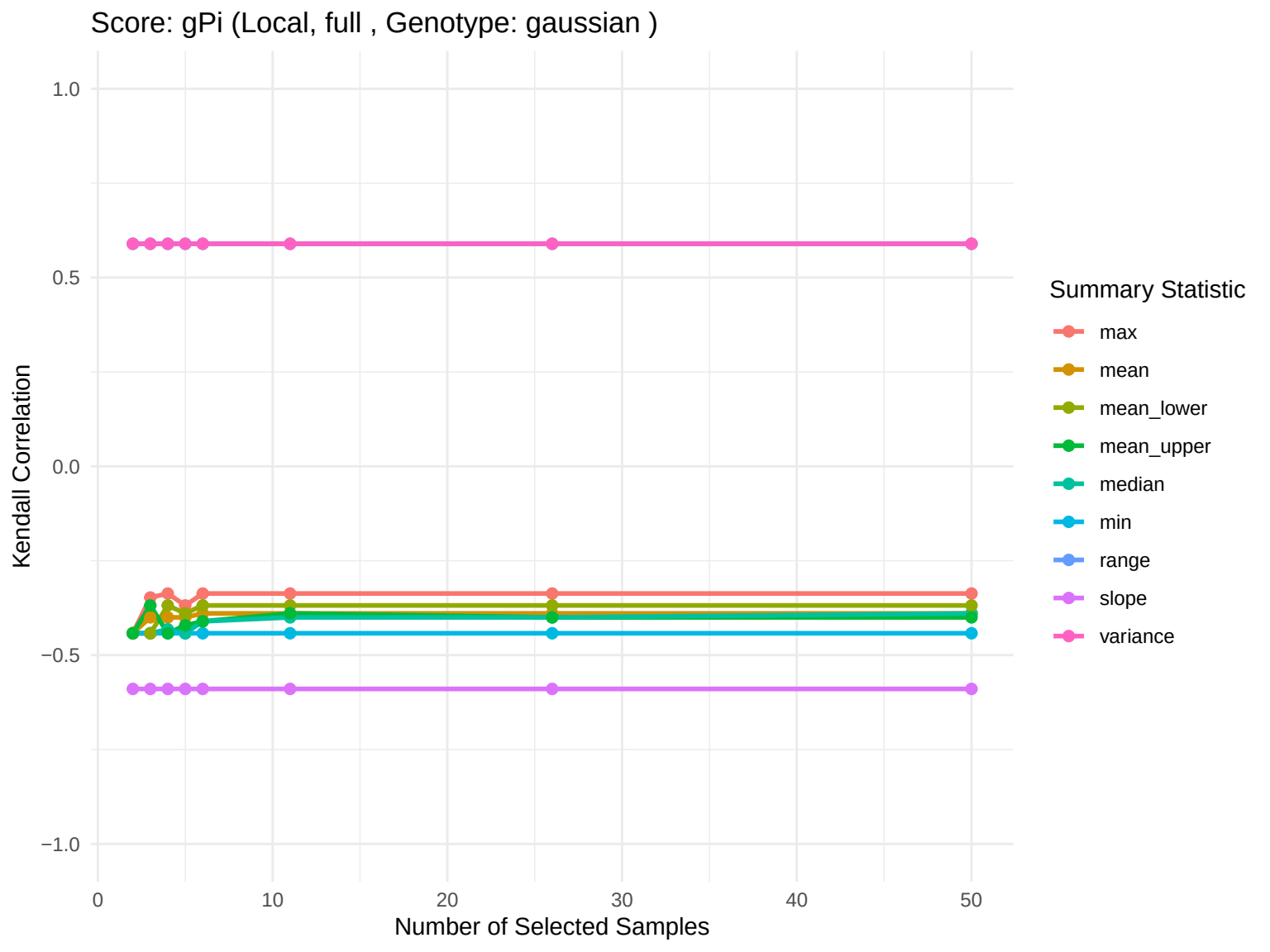
Number of Selected Samples
Summary Statistic
min
range
variance


Score: gPi (Local, full , Genotype: gaussian)
Kendall Correlation
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Number of Selected Samples
Summary Statistic
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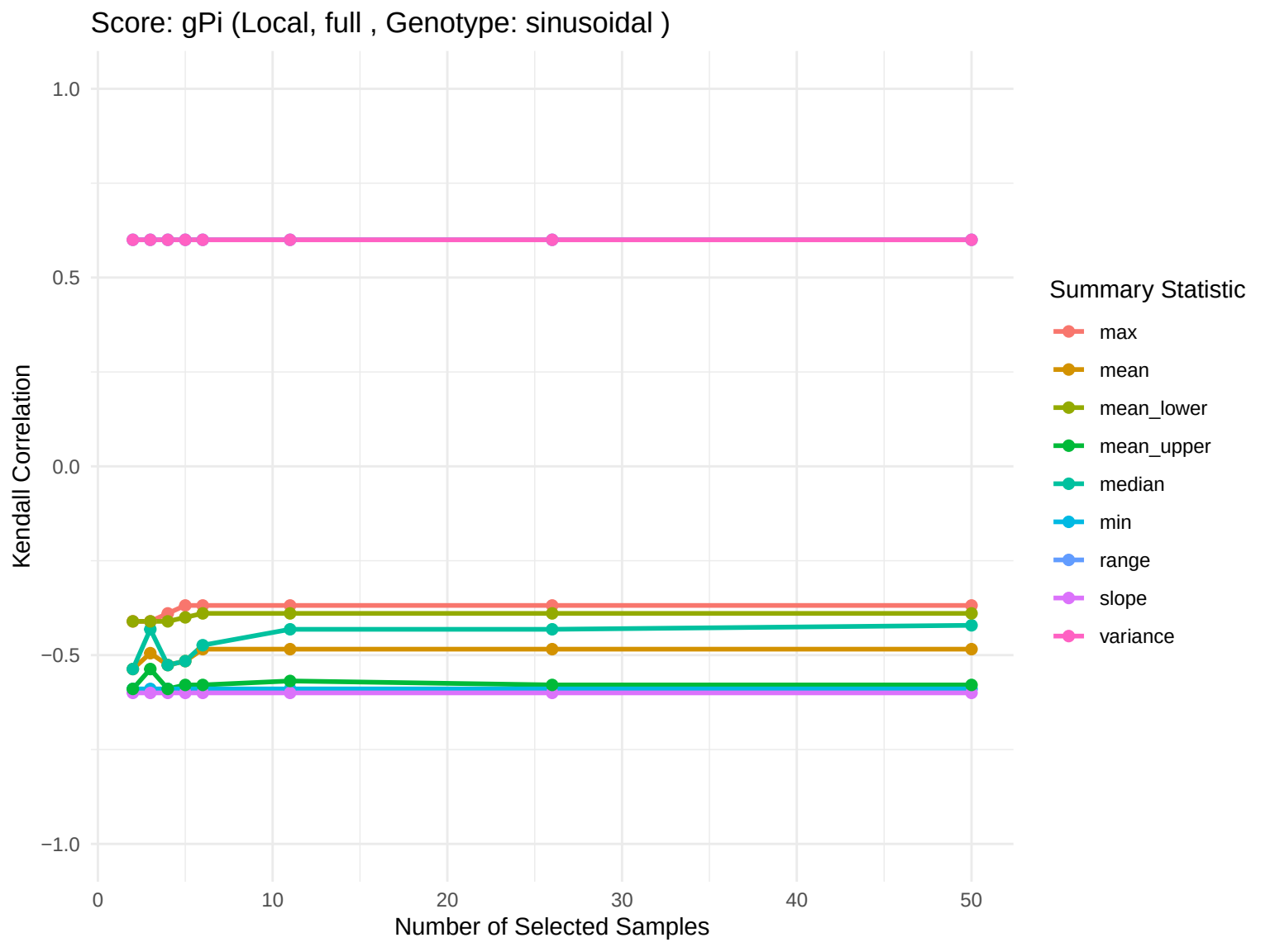
Score: gPi (Local, full , Genotype: sinusoidal)
Kendall Correlation
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0.0
−0.5
−1.0
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Number of Selected Samples
Summary Statistic
max
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mean_upper
median
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range
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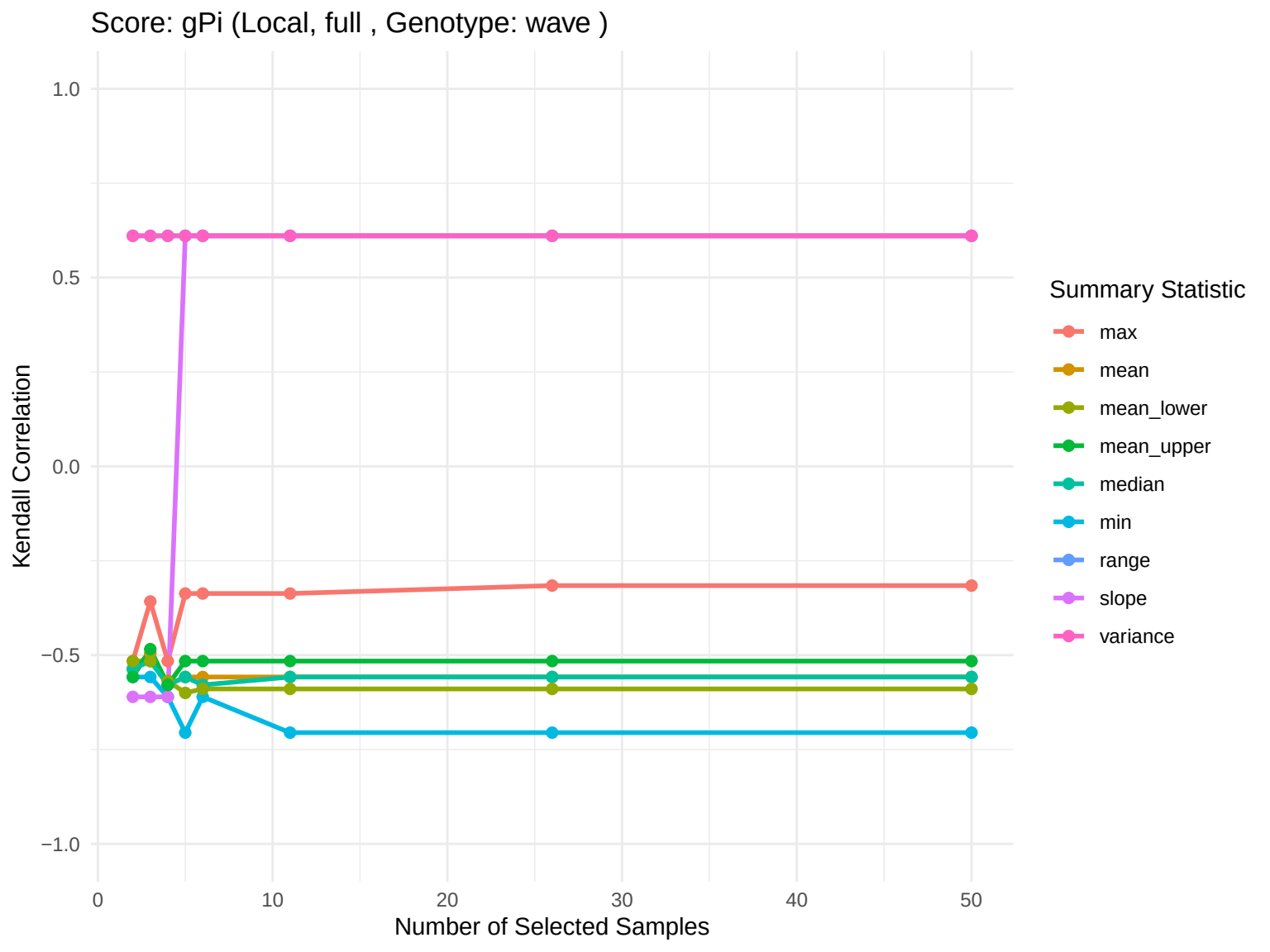
Score: gPi (Local, full , Genotype: wave)
Kendall Correlation
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0.0
−0.5
−1.0
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Number of Selected Samples
Summary Statistic
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mean_upper
median
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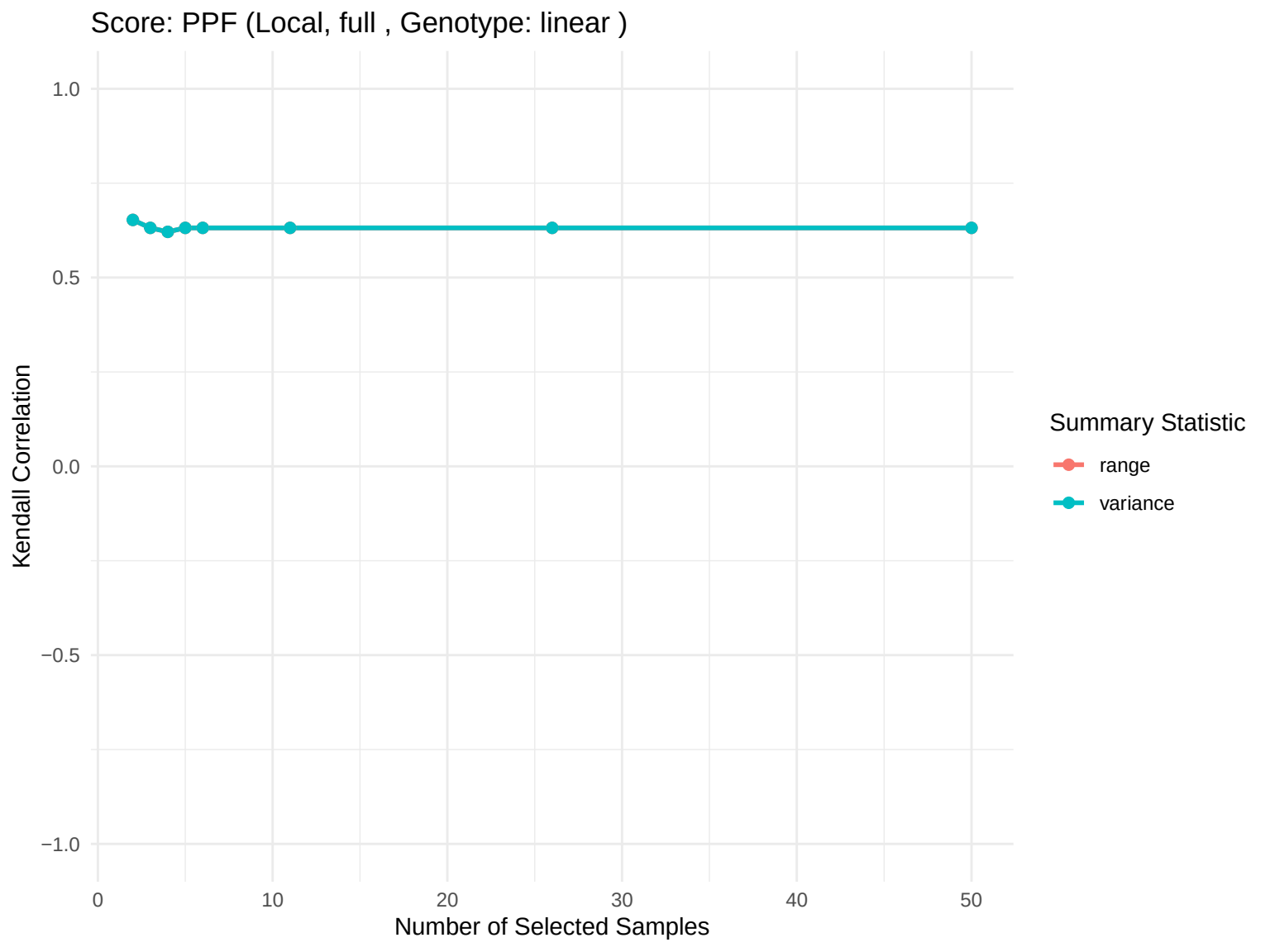
slope
variance


Score: PPF (Local, full , Genotype: linear)
Kendall Correlation
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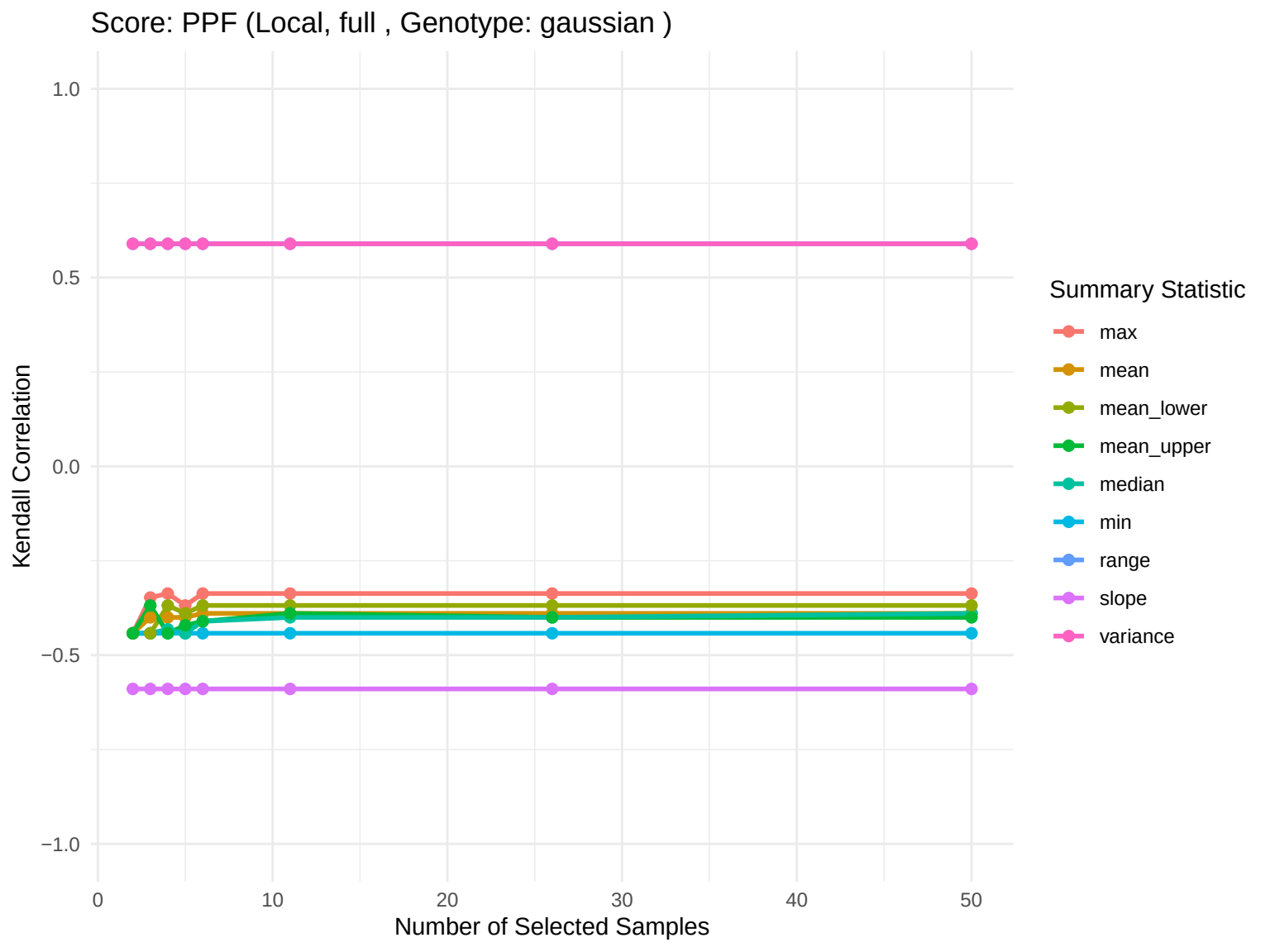
Number of Selected Samples
Summary Statistic
range
variance


Score: PPF (Local, full , Genotype: gaussian)
Kendall Correlation
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0.0
−0.5
−1.0
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Number of Selected Samples
Summary Statistic
max
mean
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mean_upper
median
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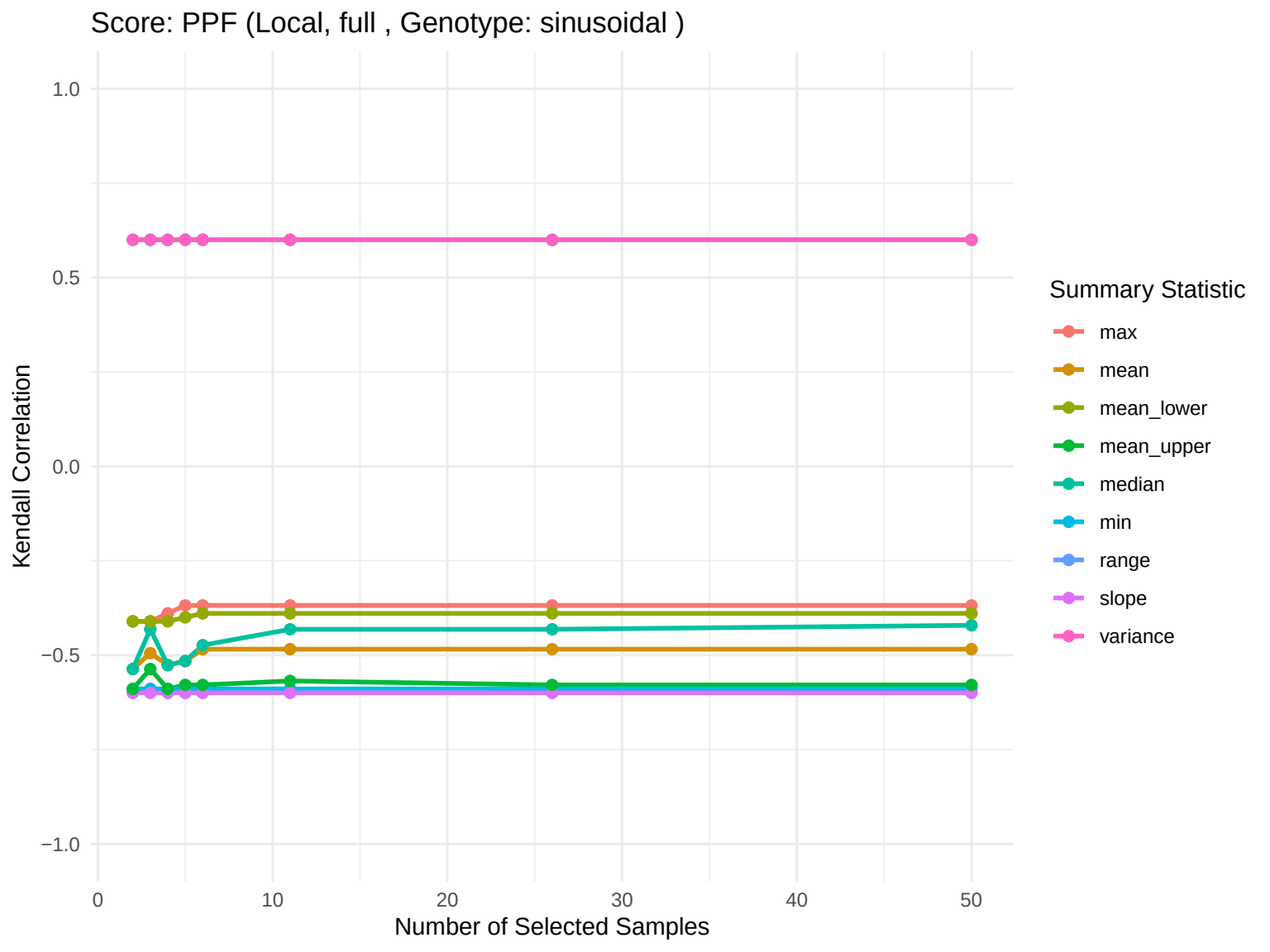
Score: PPF (Local, full , Genotype: sinusoidal)
Kendall Correlation
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0.0
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−1.0
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Number of Selected Samples
Summary Statistic
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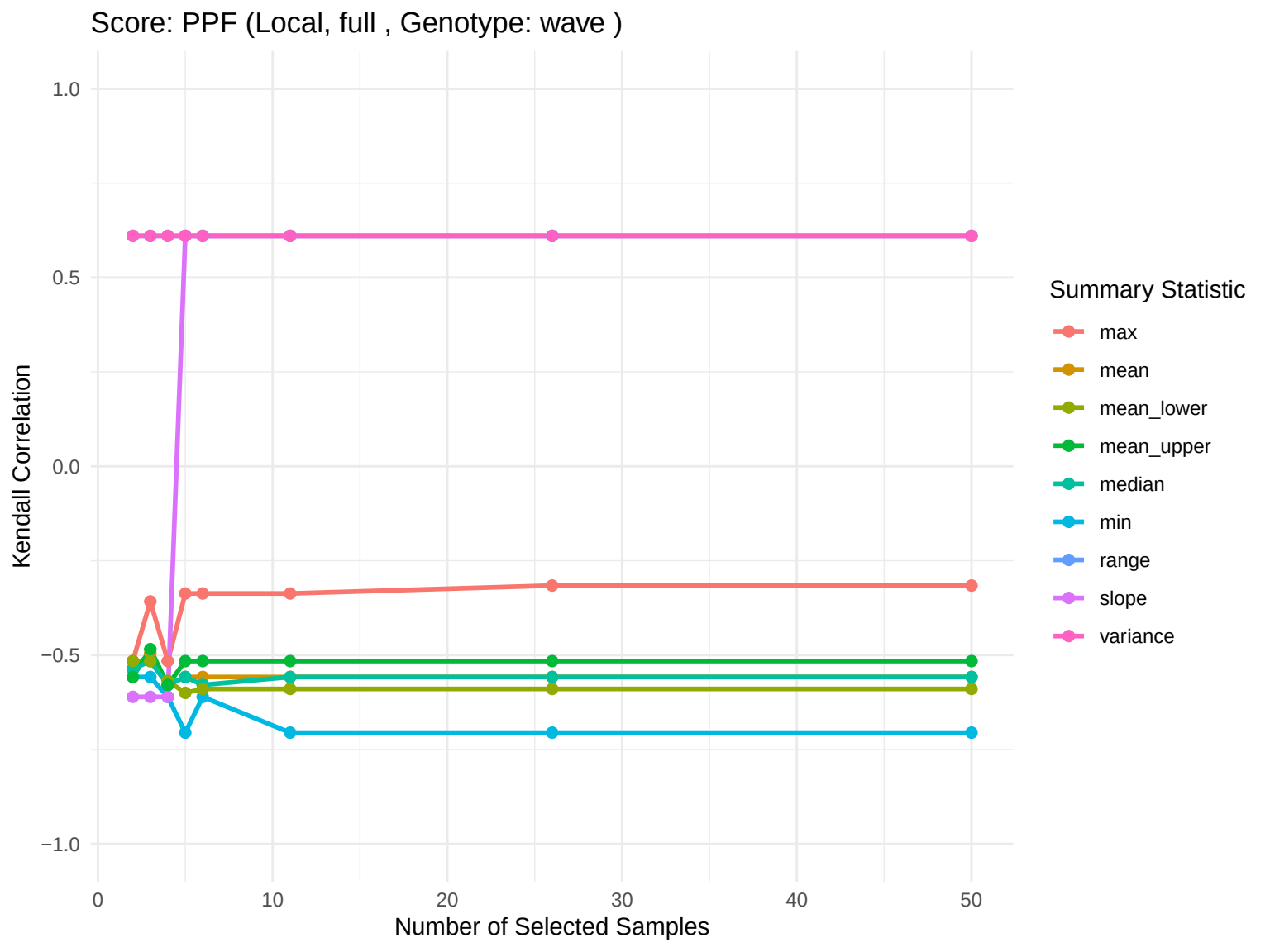
Score: PPF (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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Number of Selected Samples
Summary Statistic
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mean
mean_lower
mean_upper
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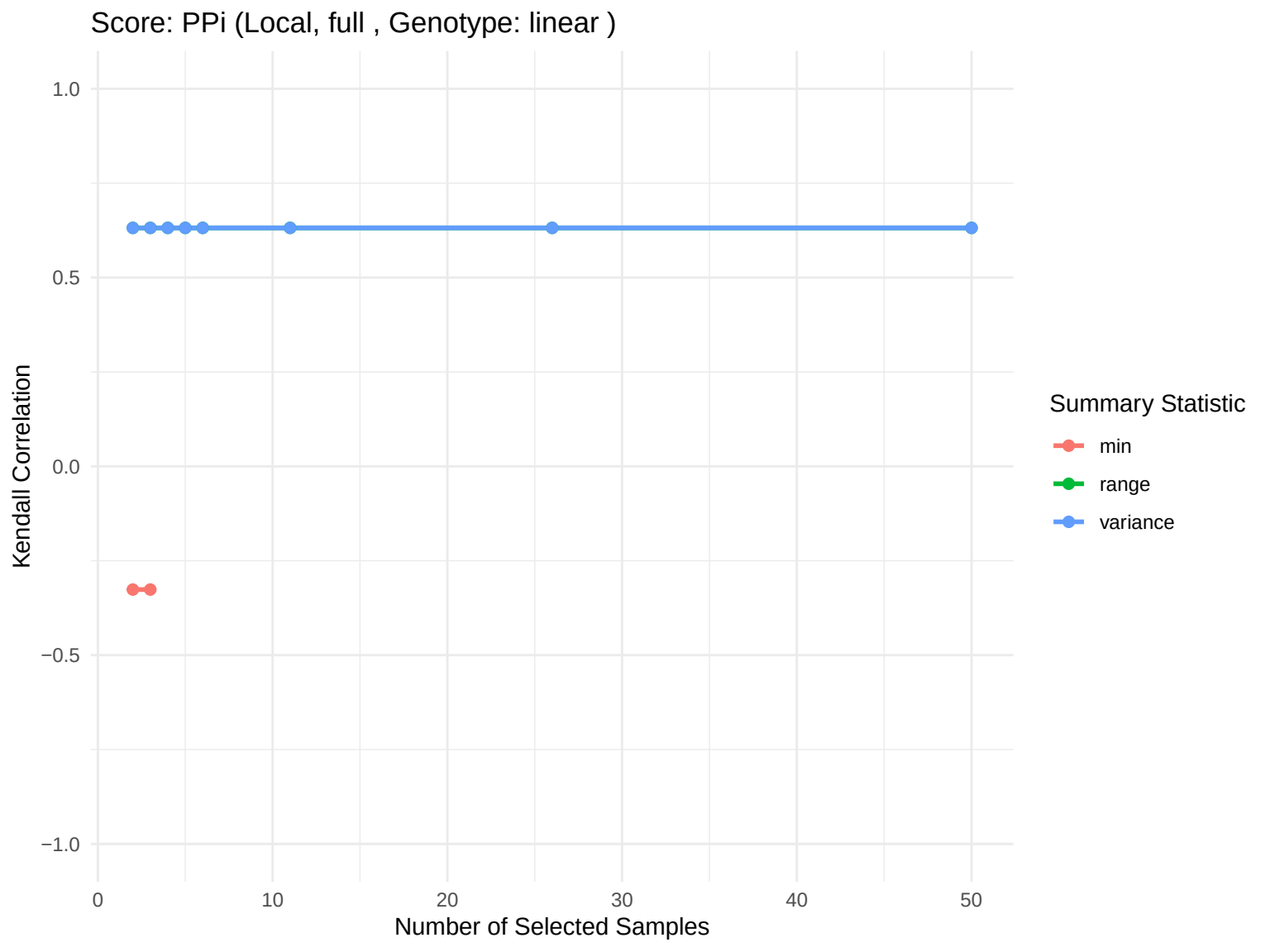
variance


Score: PPi (Local, full , Genotype: linear)
Kendall Correlation
1.0
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−1.0
0
10
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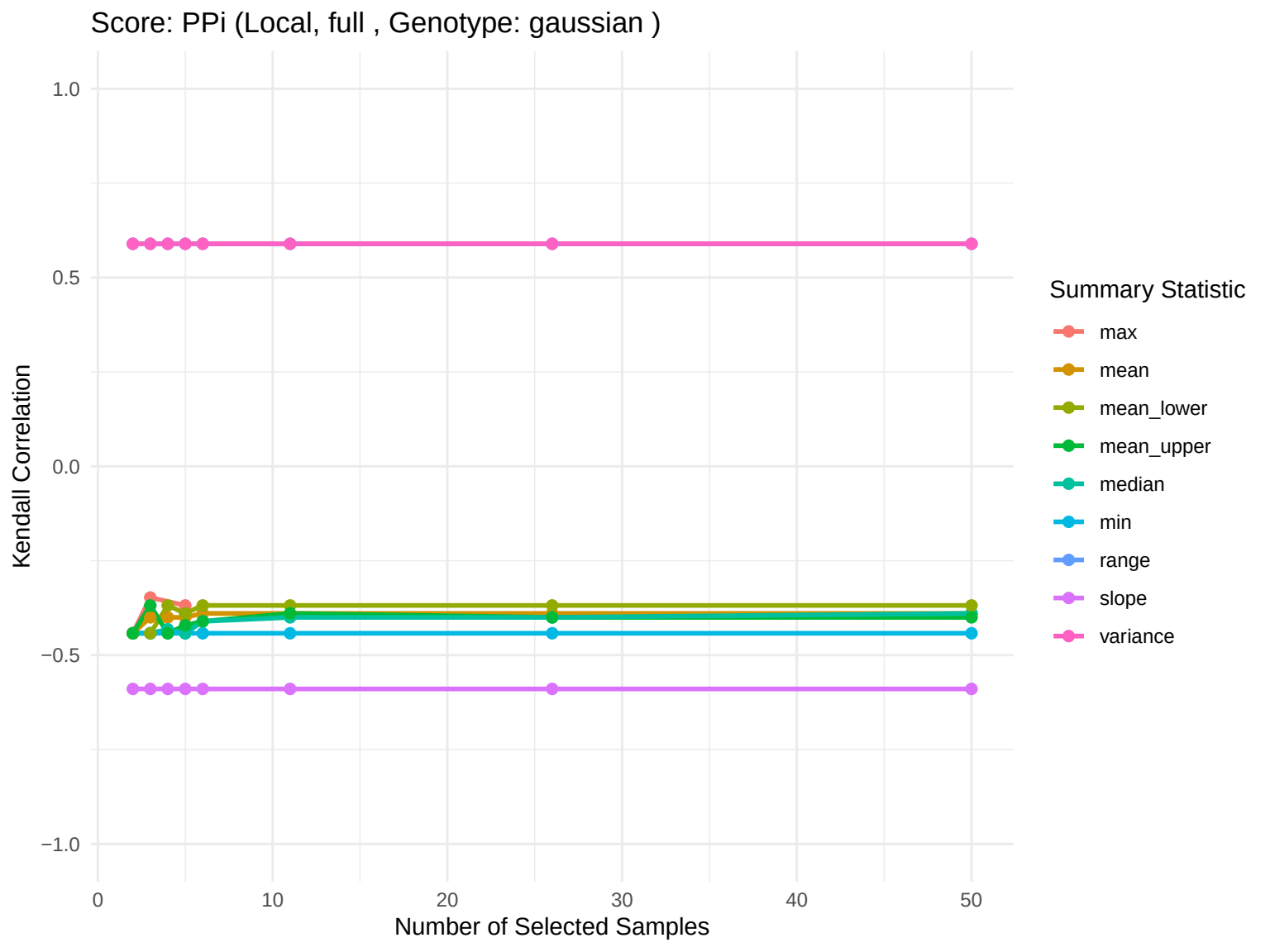
Number of Selected Samples
Summary Statistic
min
range
variance


Score: PPi (Local, full , Genotype: gaussian)
Kendall Correlation
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0.0
−0.5
−1.0
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Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
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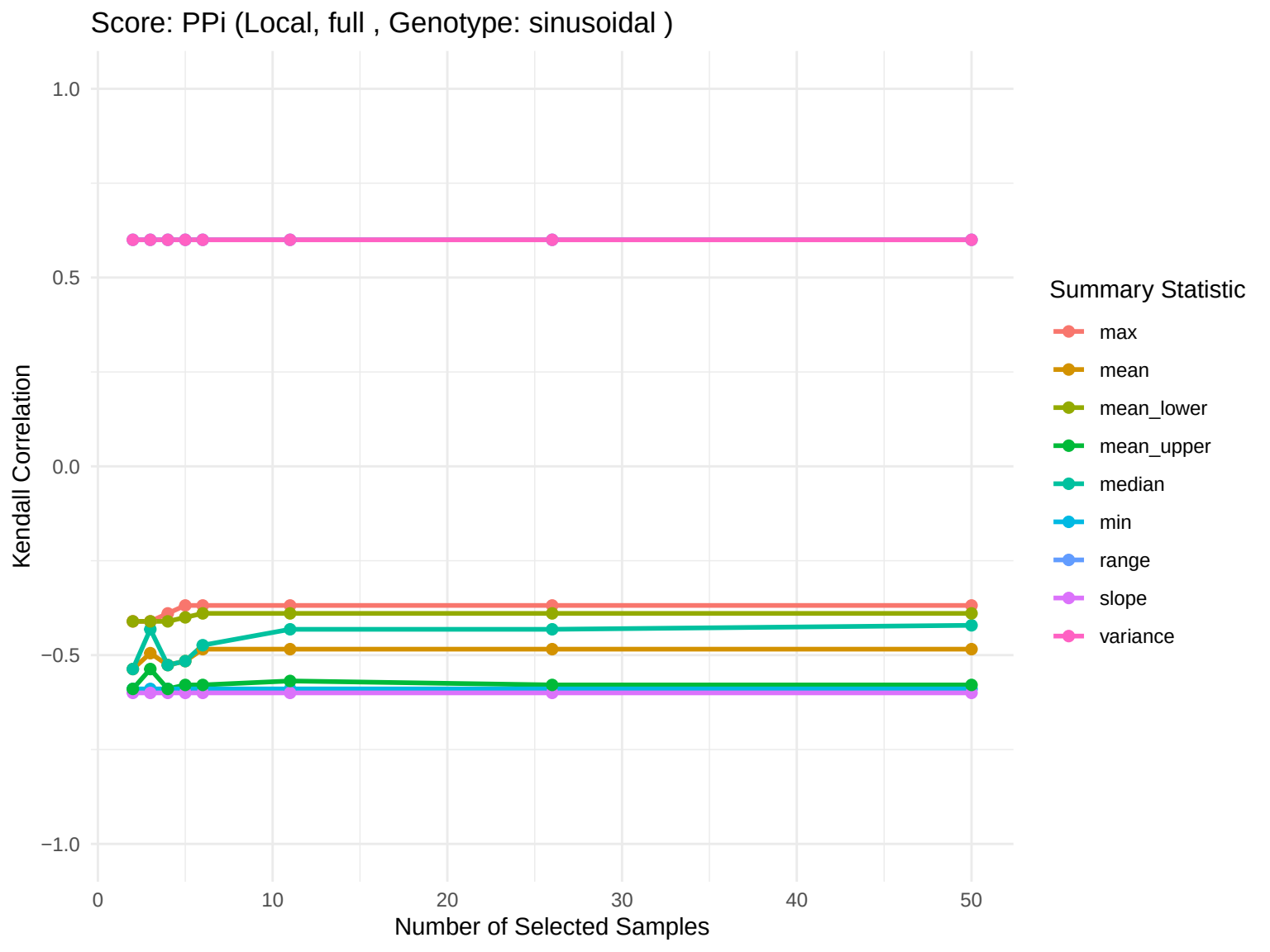
slope
variance


Score: PPi (Local, full , Genotype: sinusoidal)
Kendall Correlation
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0.5
0.0
−0.5
−1.0
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Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
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slope
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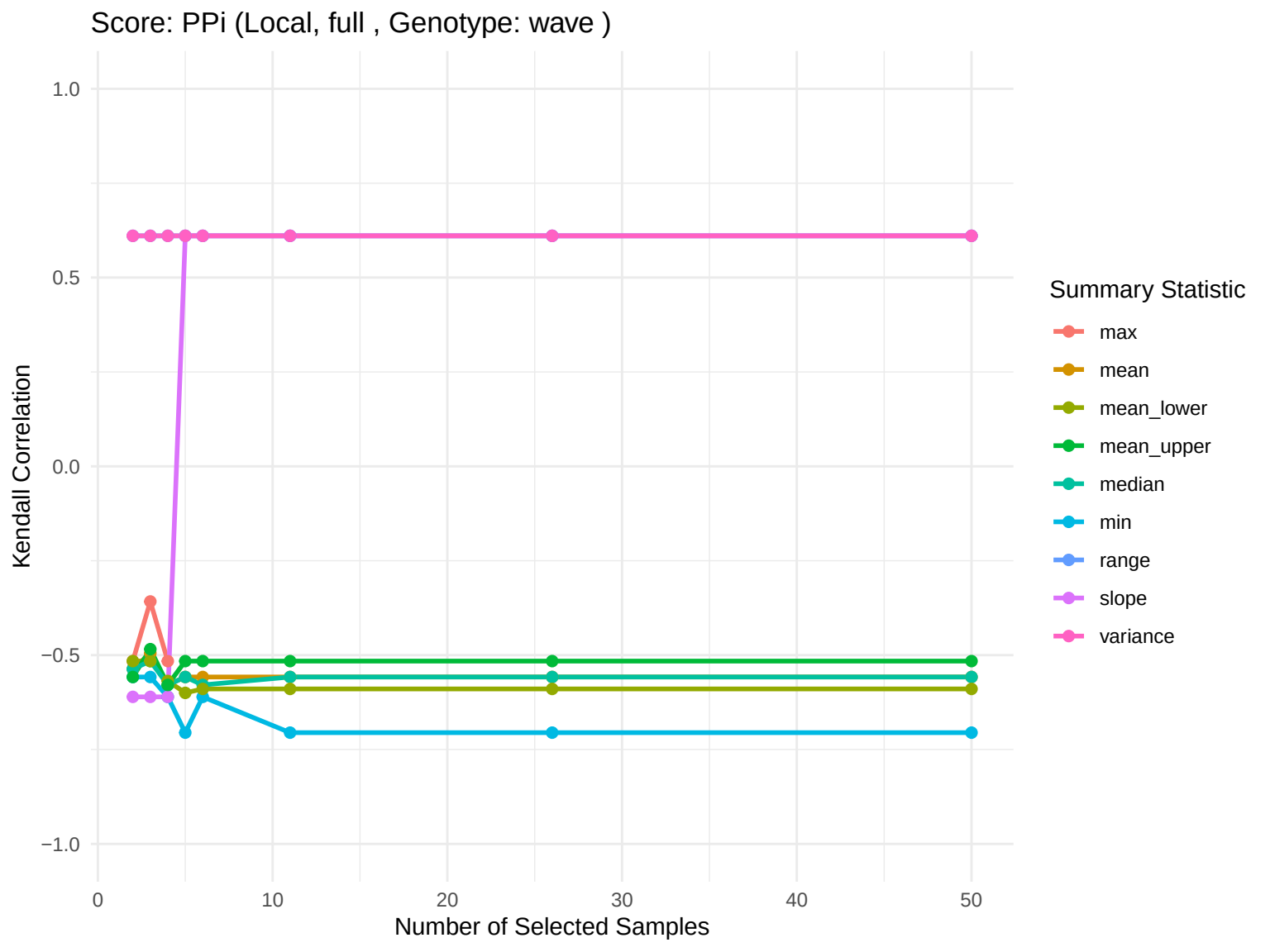
Score: PPi (Local, full , Genotype: wave)
Kendall Correlation
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0.0
−0.5
−1.0
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Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
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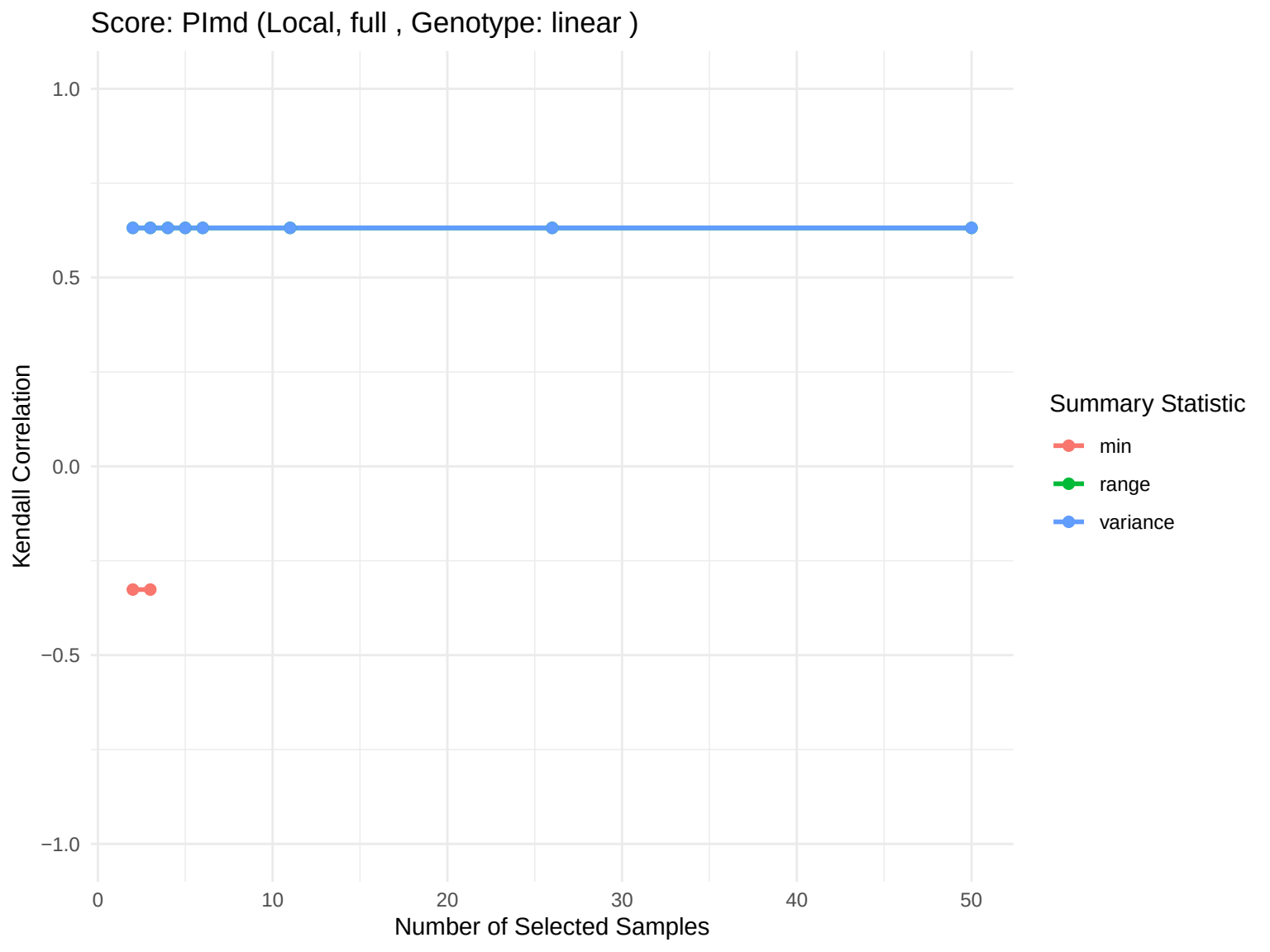
slope
variance


Score: PImd (Local, full , Genotype: linear)
Kendall Correlation
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−1.0
0
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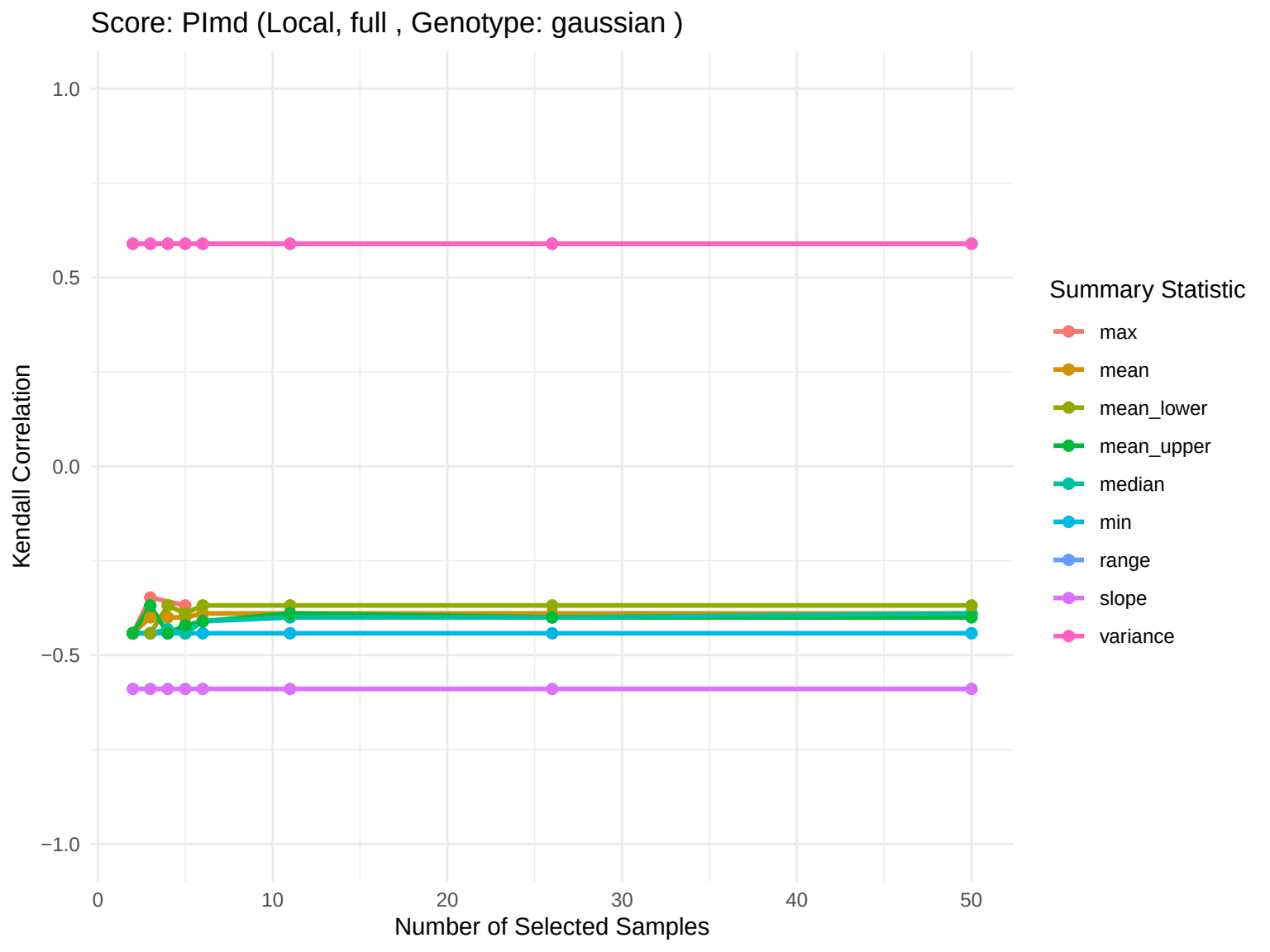
Number of Selected Samples
Summary Statistic
min
range
variance


Score: PImd (Local, full , Genotype: gaussian)
Kendall Correlation
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0.0
−0.5
−1.0
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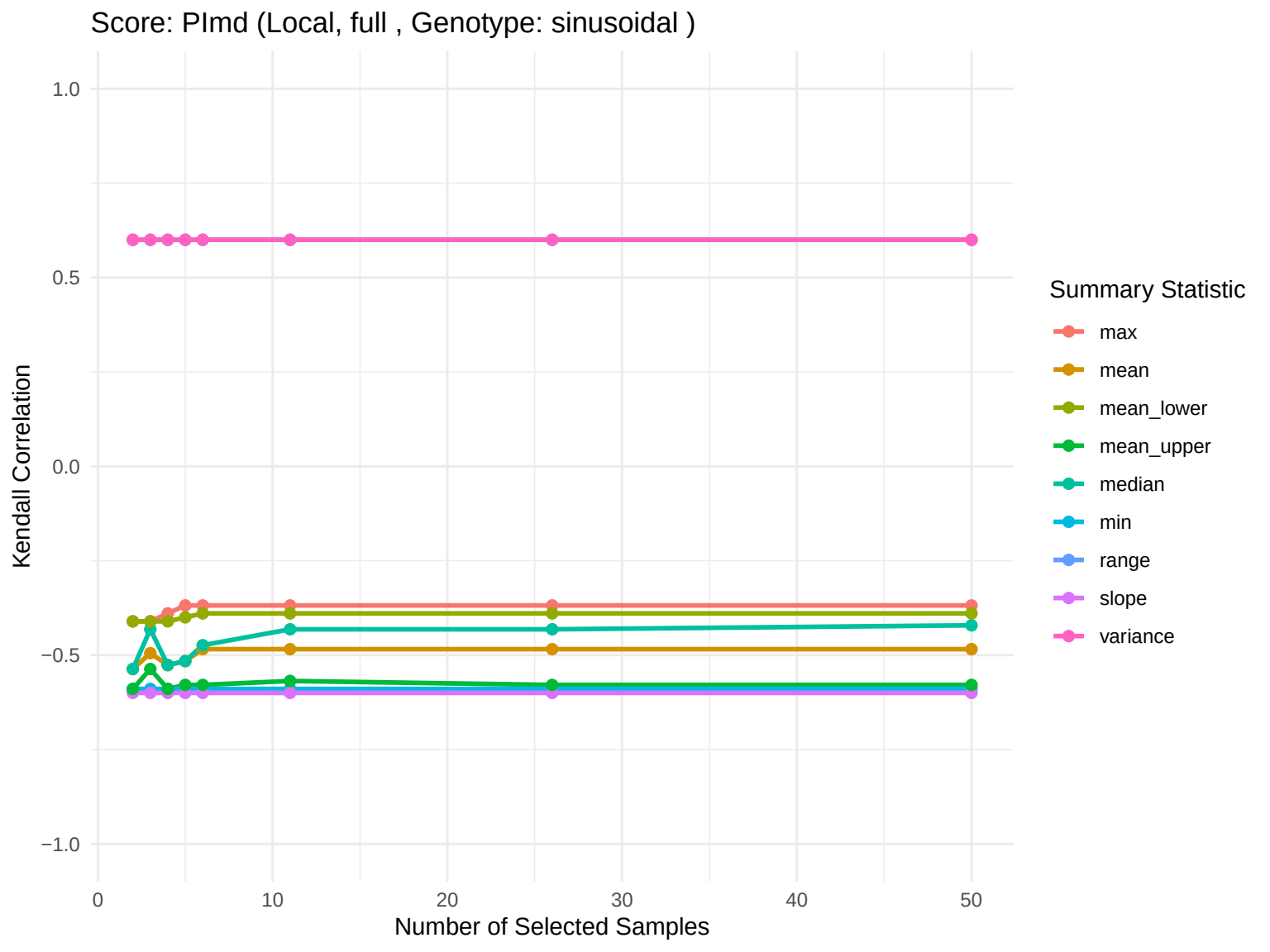
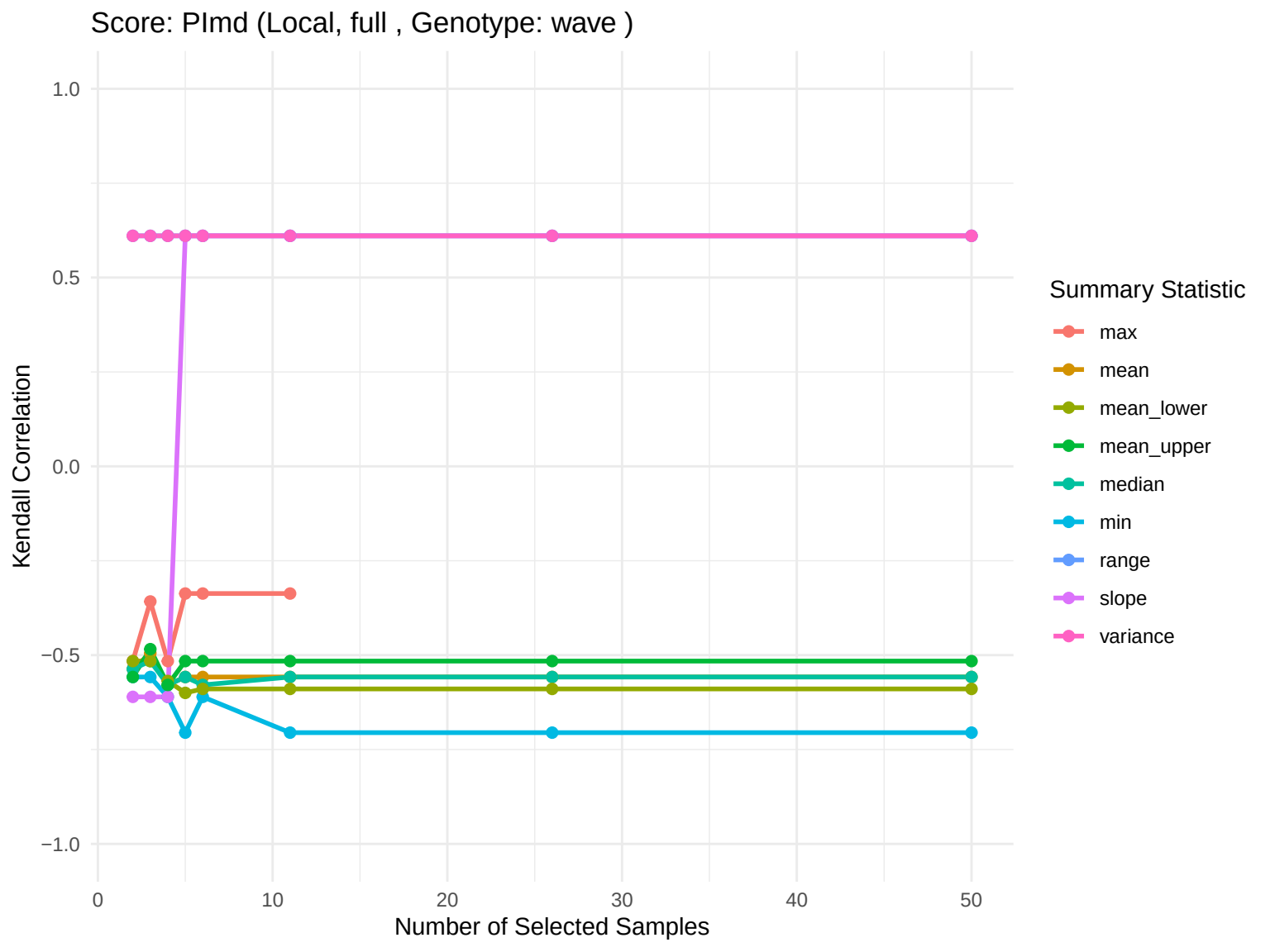
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PImd (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PImd (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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20
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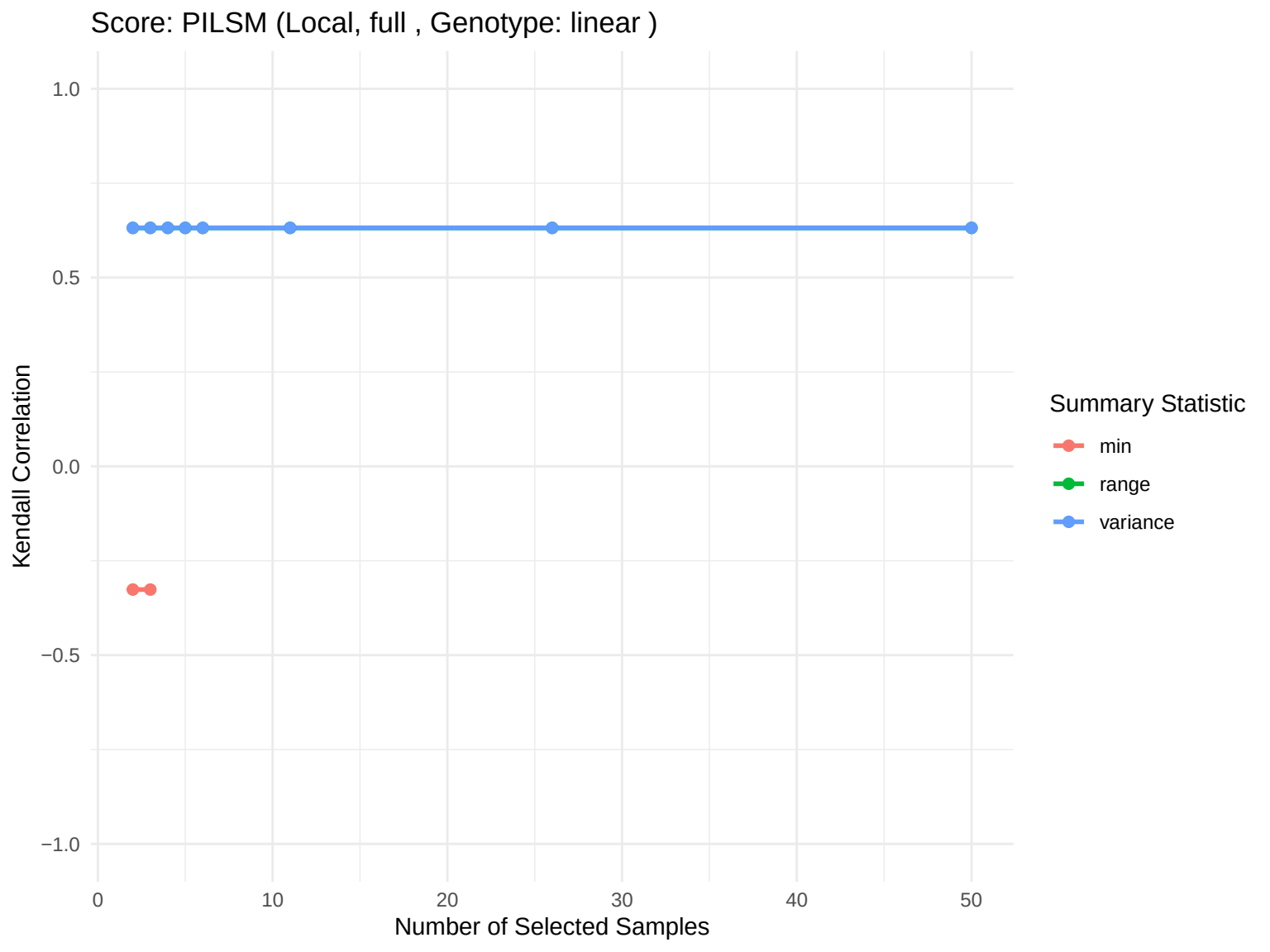
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PILSM (Local, full , Genotype: linear)
Kendall Correlation
1.0
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−1.0
0
10
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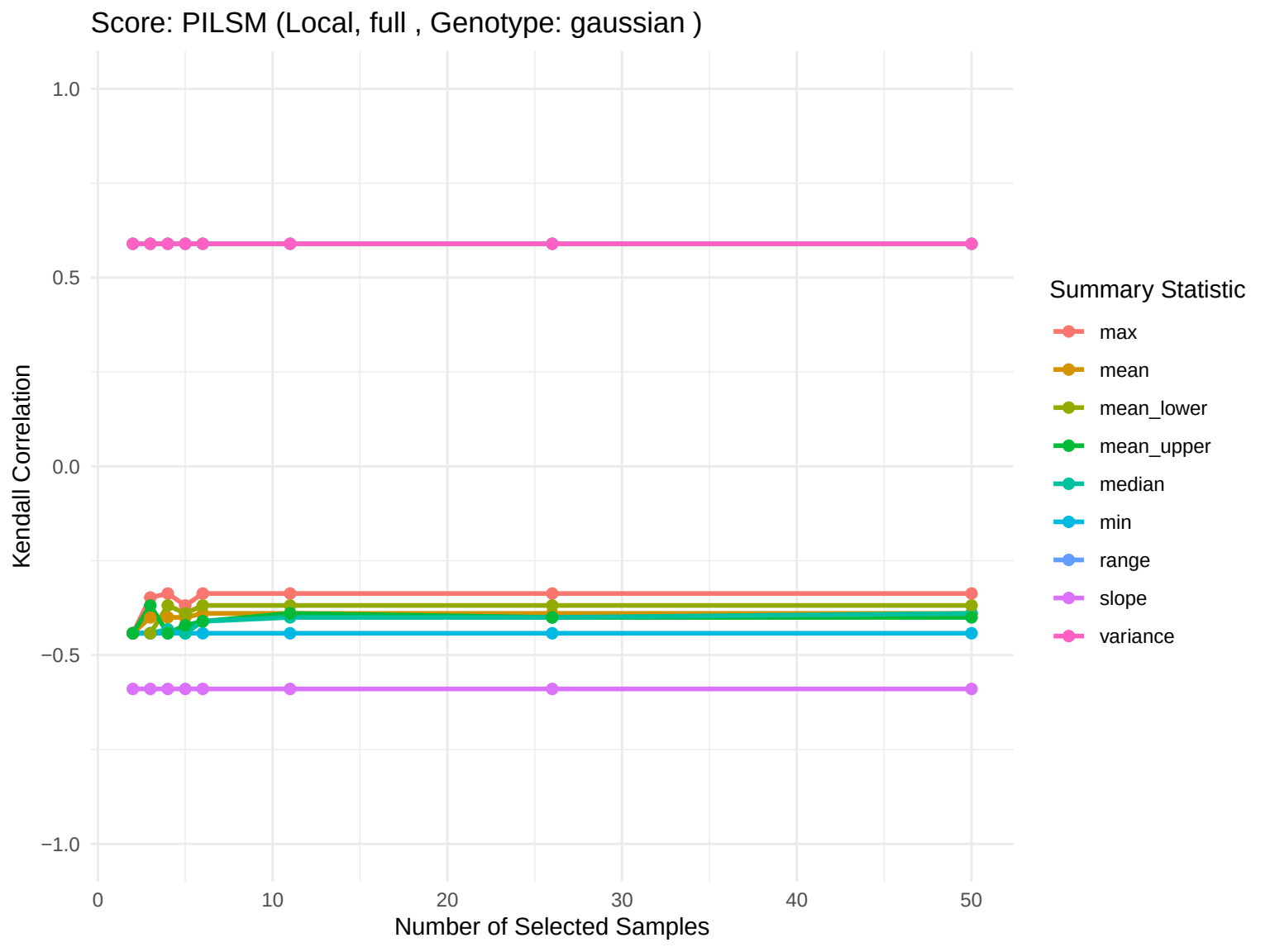
40
50
Number of Selected Samples
Summary Statistic
min
range
variance


Score: PILSM (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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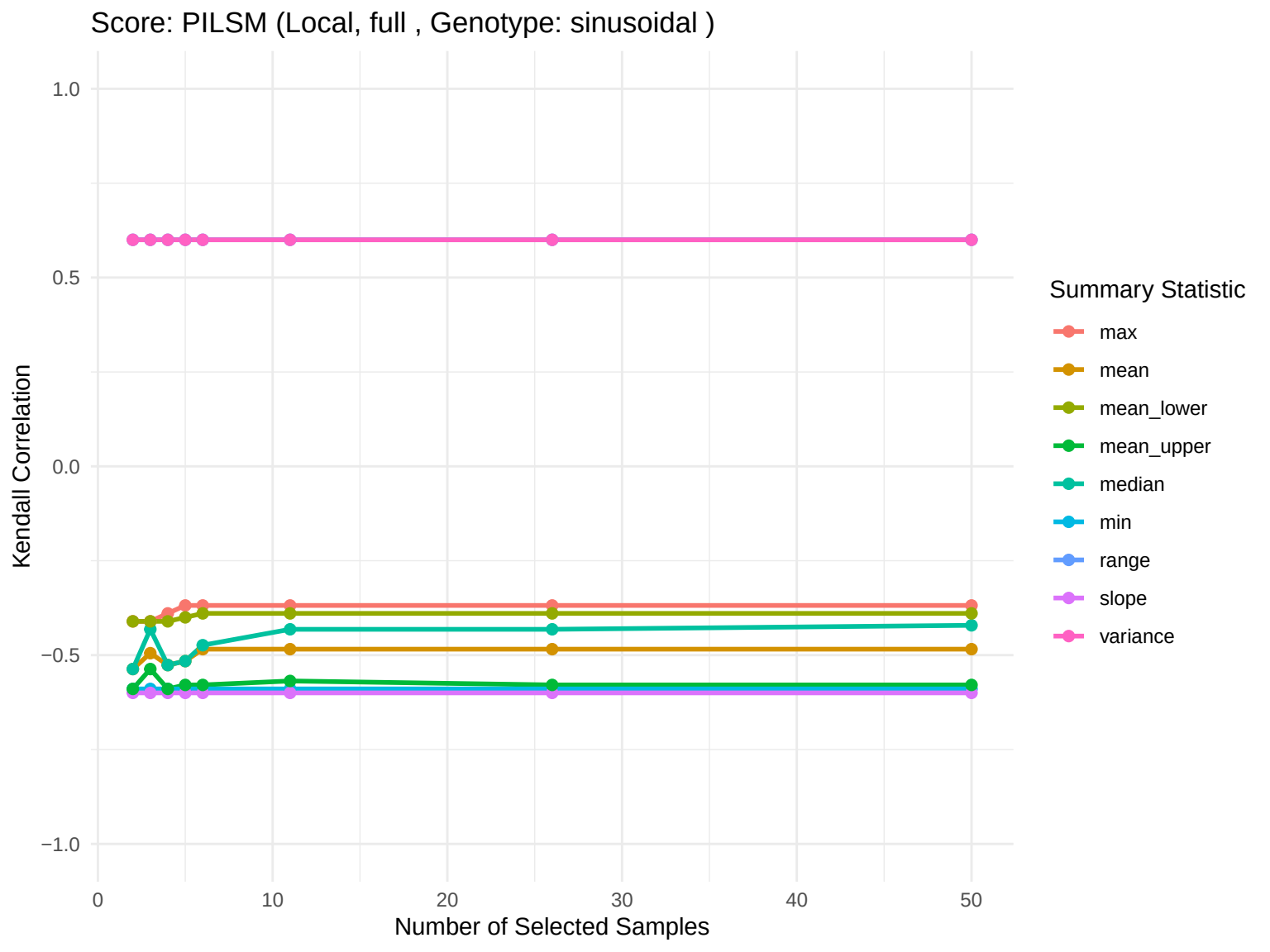
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PILSM (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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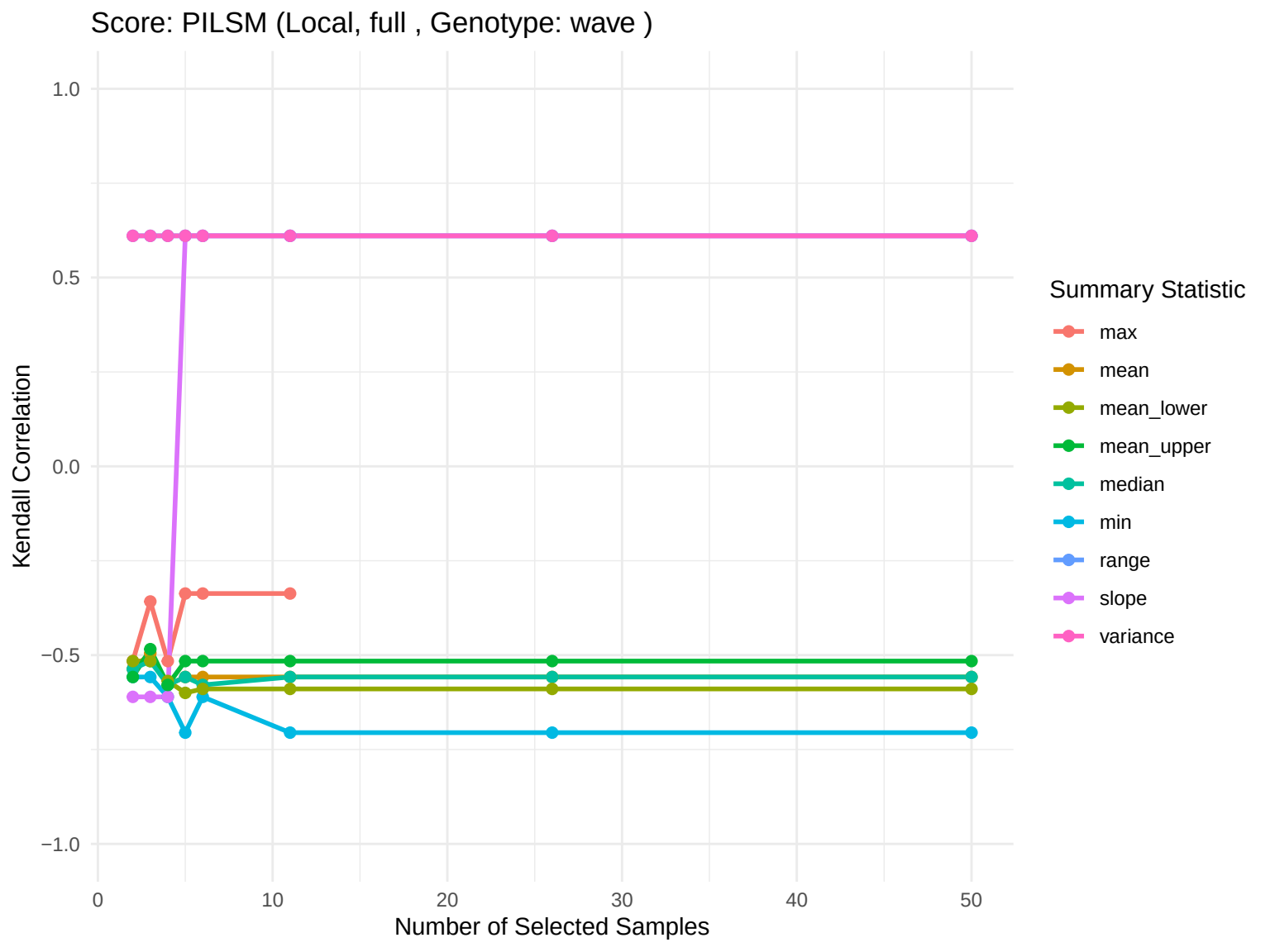
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PILSM (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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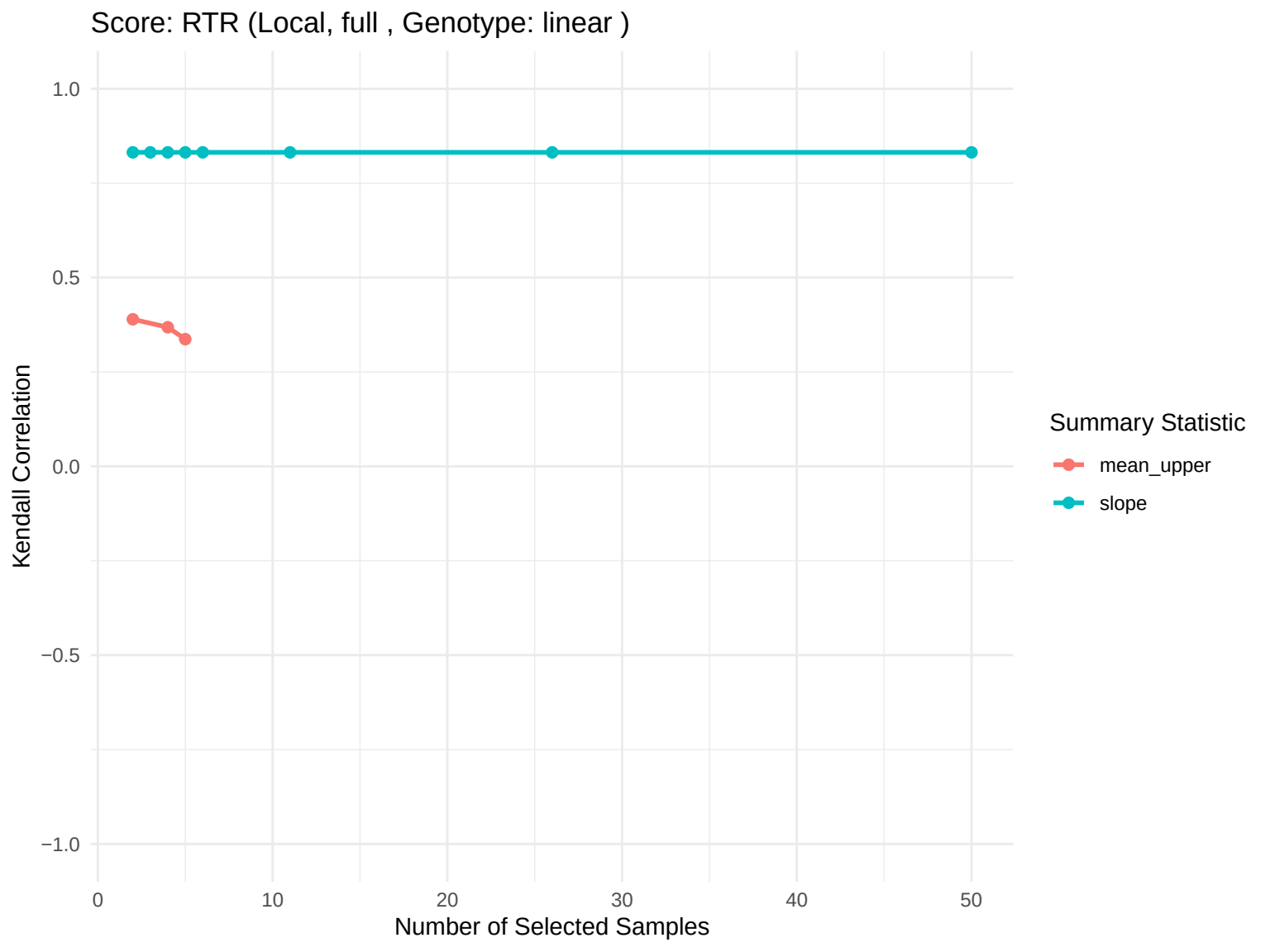
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RTR (Local, full , Genotype: linear)
Kendall Correlation
1.0
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0.0
−0.5
−1.0
0
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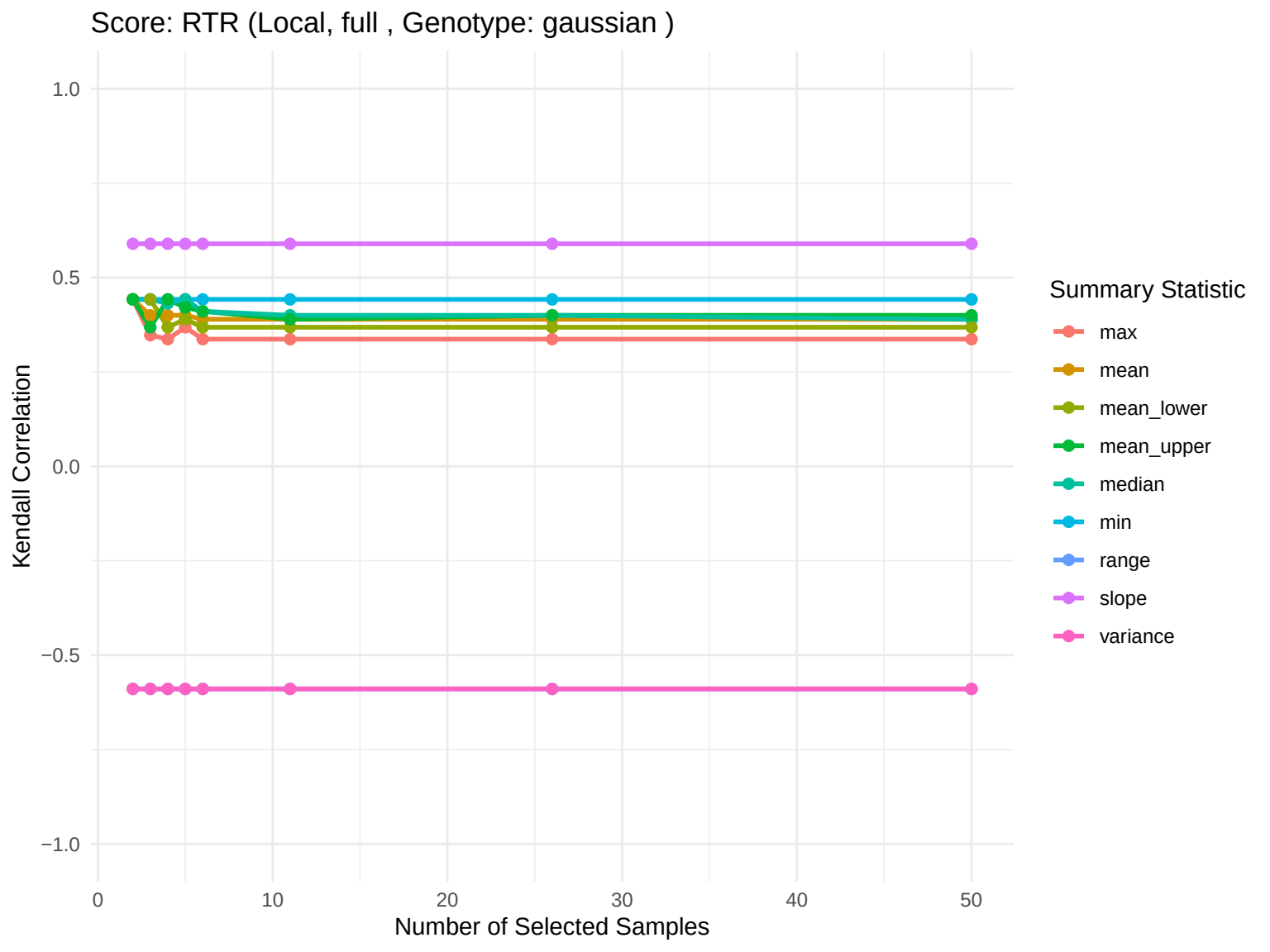
40
50
Number of Selected Samples
Summary Statistic
mean_upper
slope


Score: RTR (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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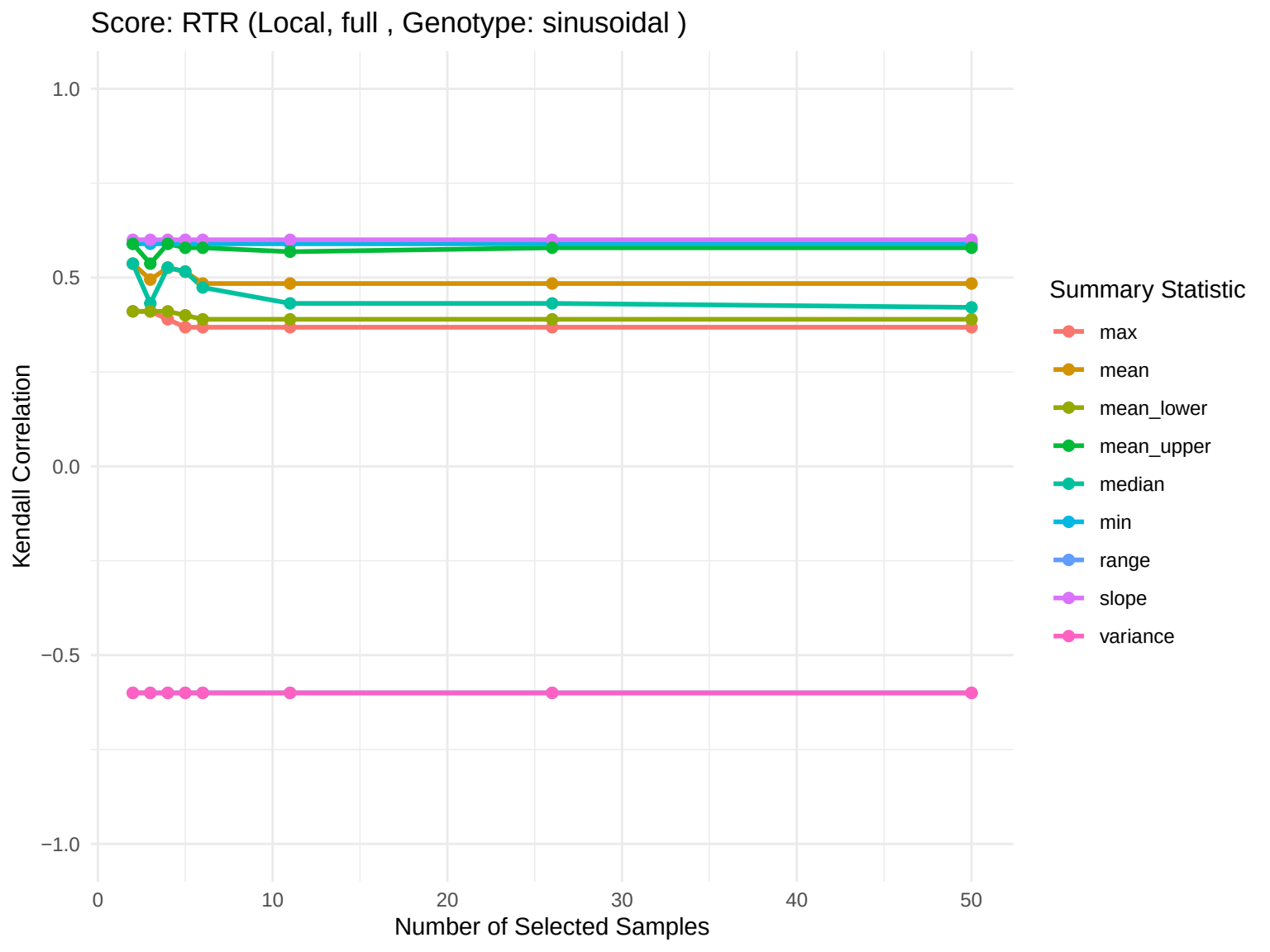
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RTR (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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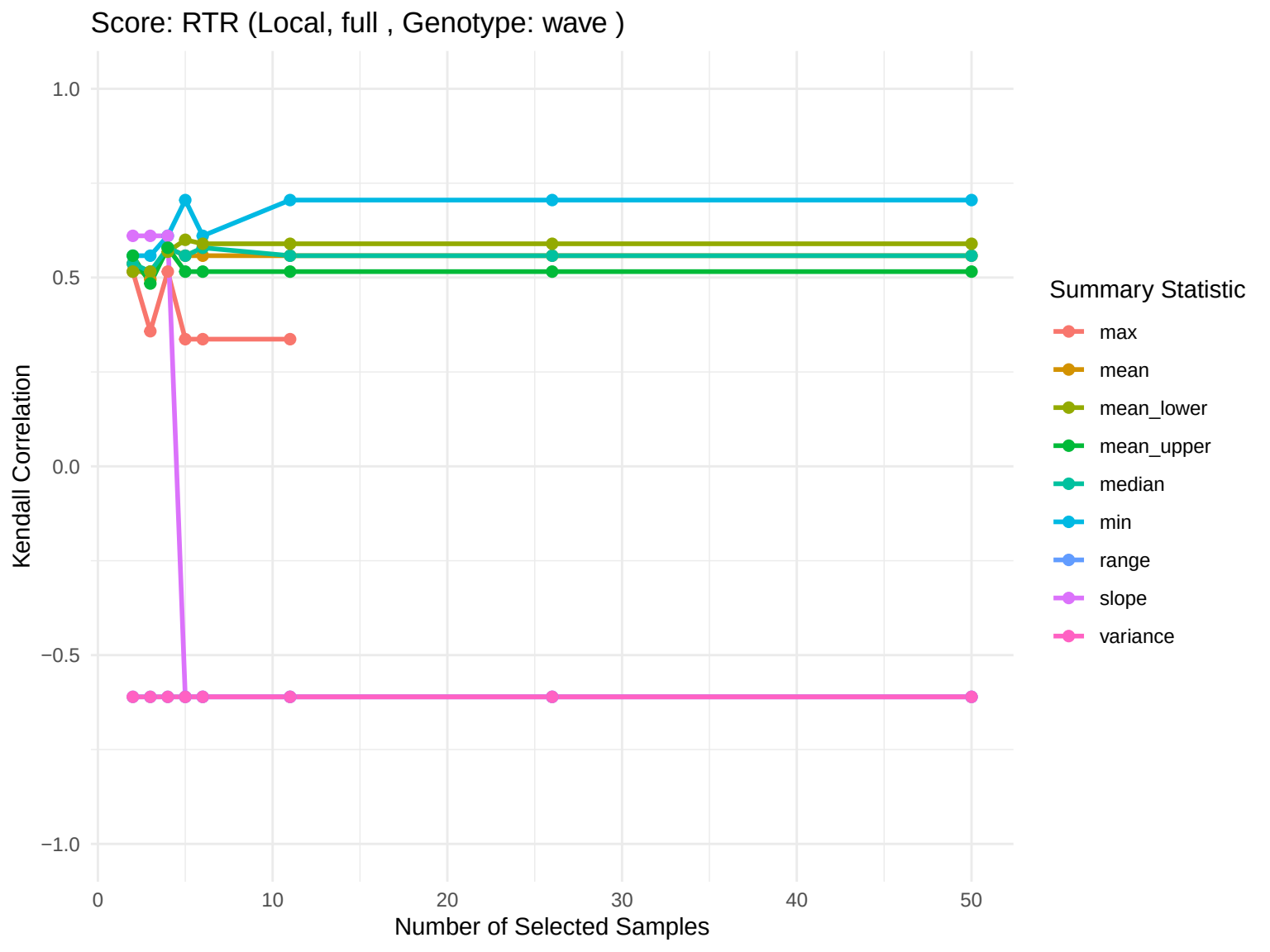
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RTR (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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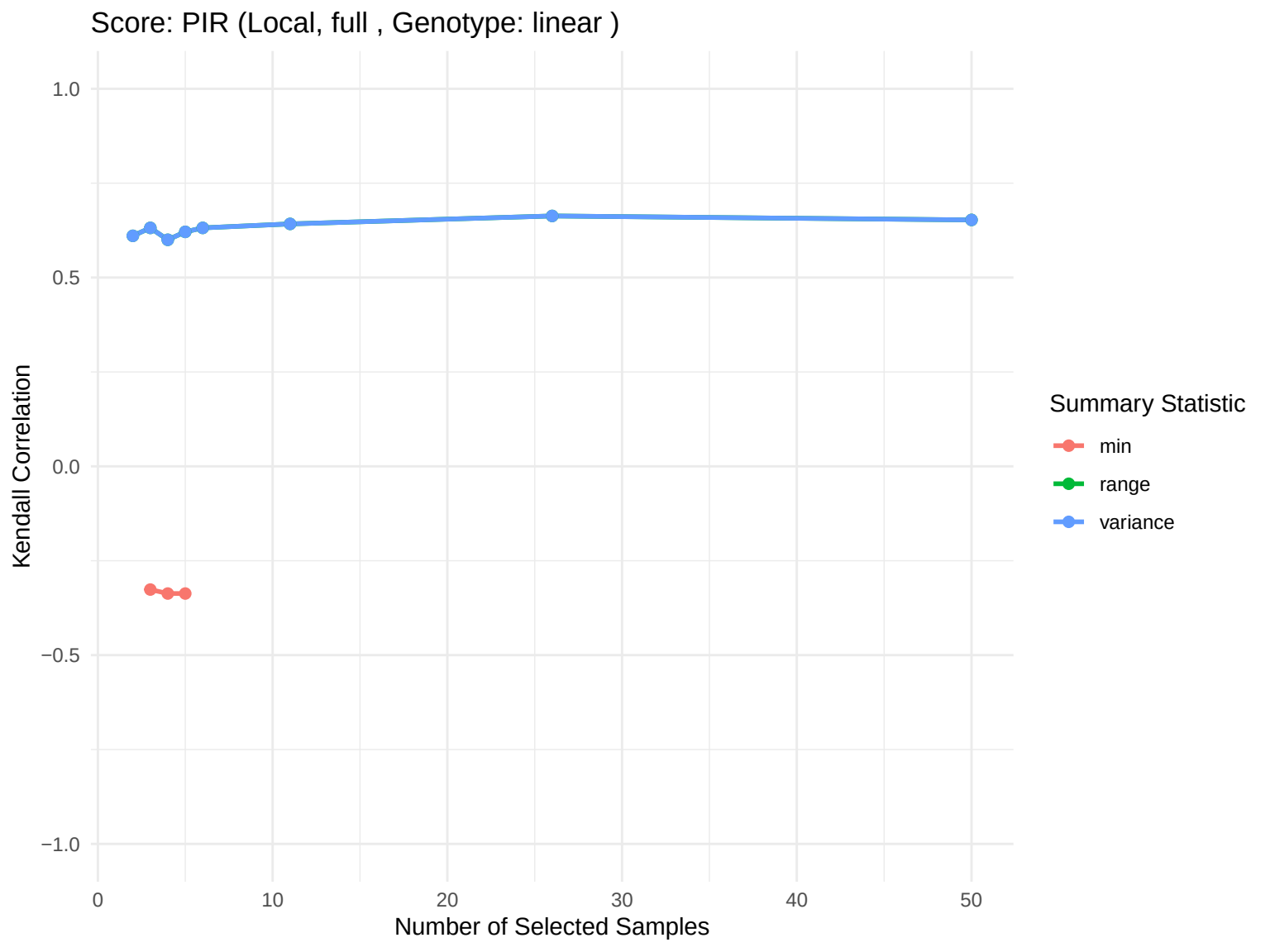
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PIR (Local, full , Genotype: linear)
Kendall Correlation
1.0
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0.0
−0.5
−1.0
0
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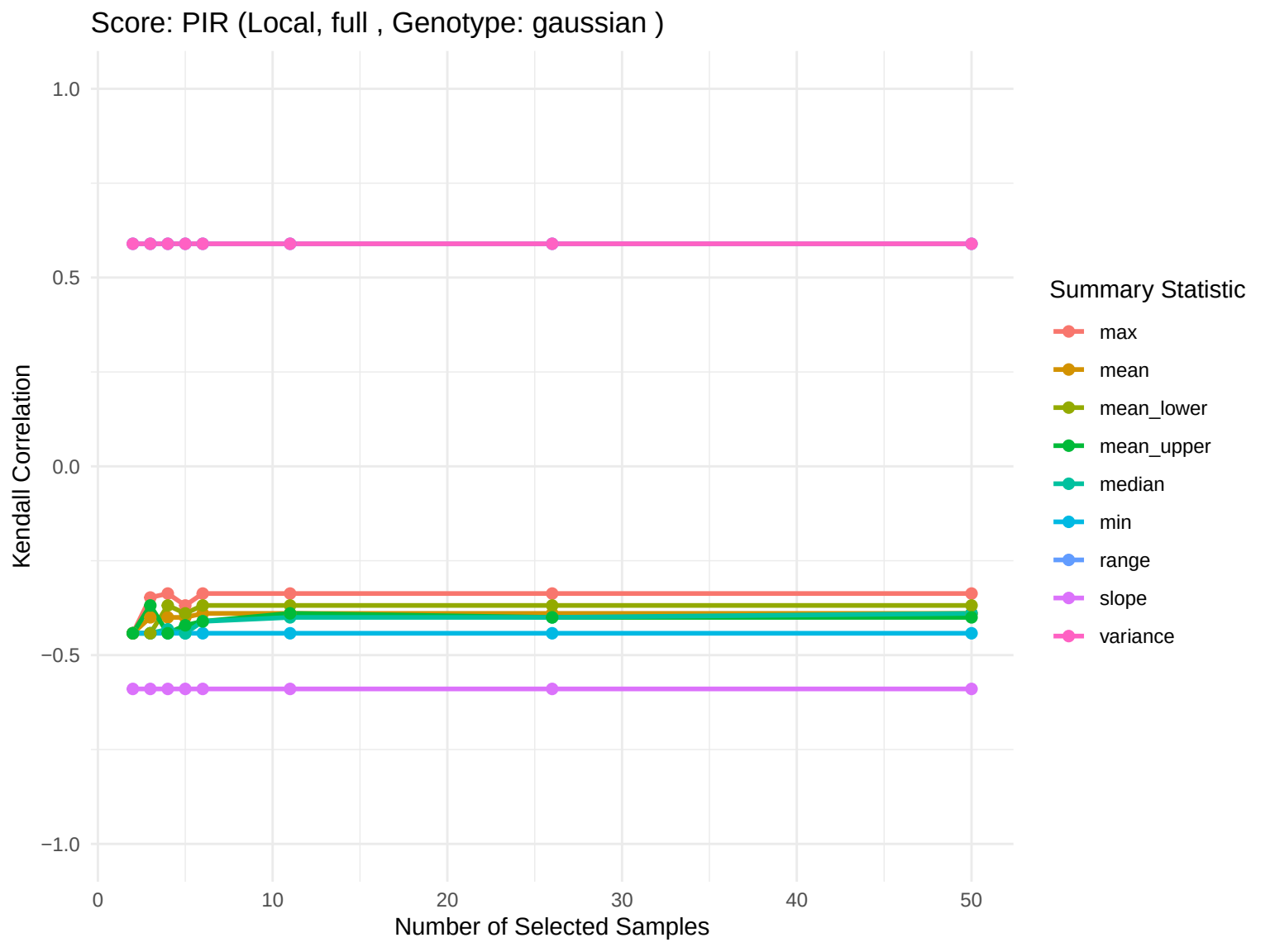
40
50
Number of Selected Samples
Summary Statistic
min
range
variance


Score: PIR (Local, full , Genotype: gaussian)
Kendall Correlation
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0.5
0.0
−0.5
−1.0
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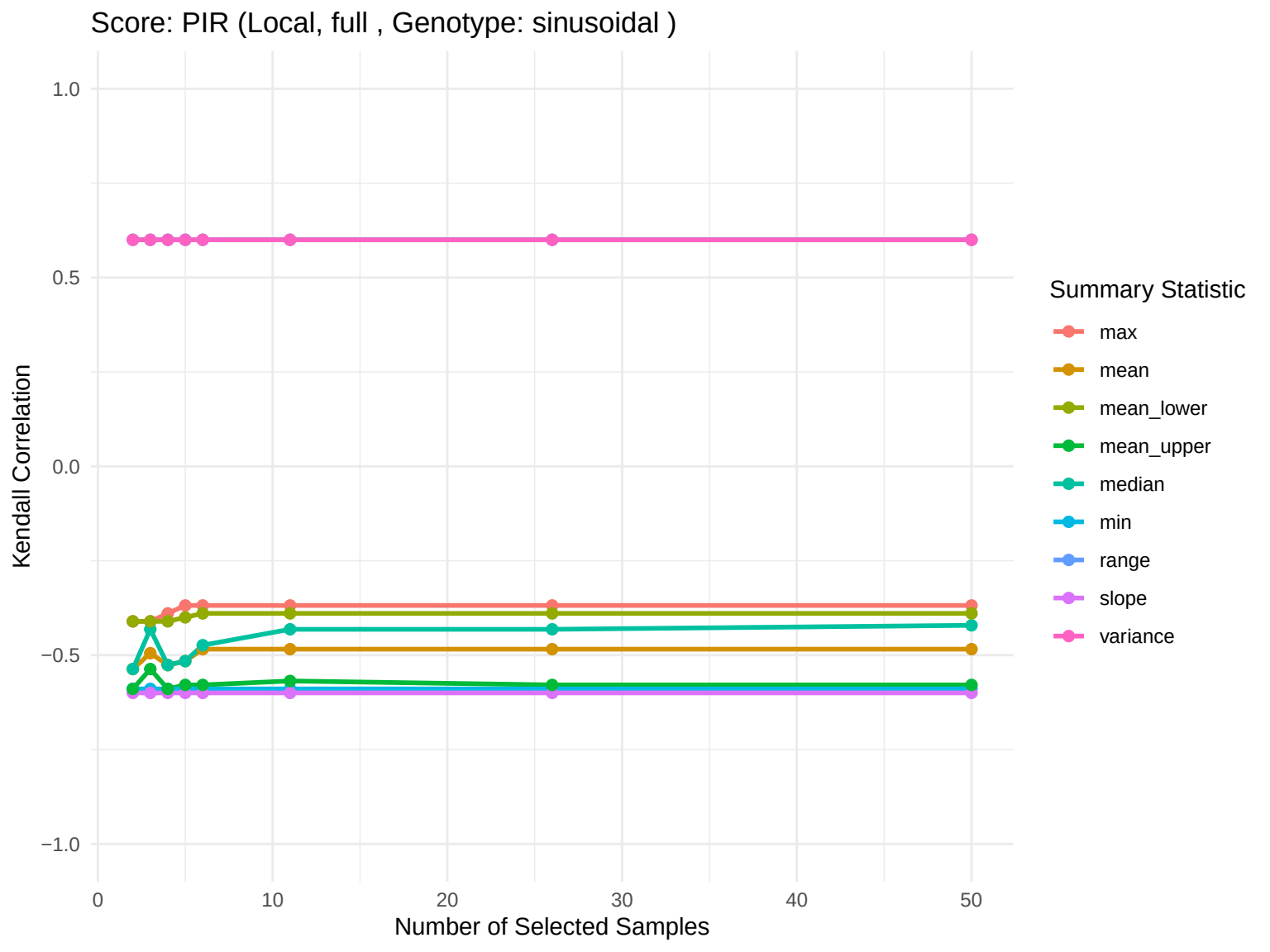
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PIR (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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20
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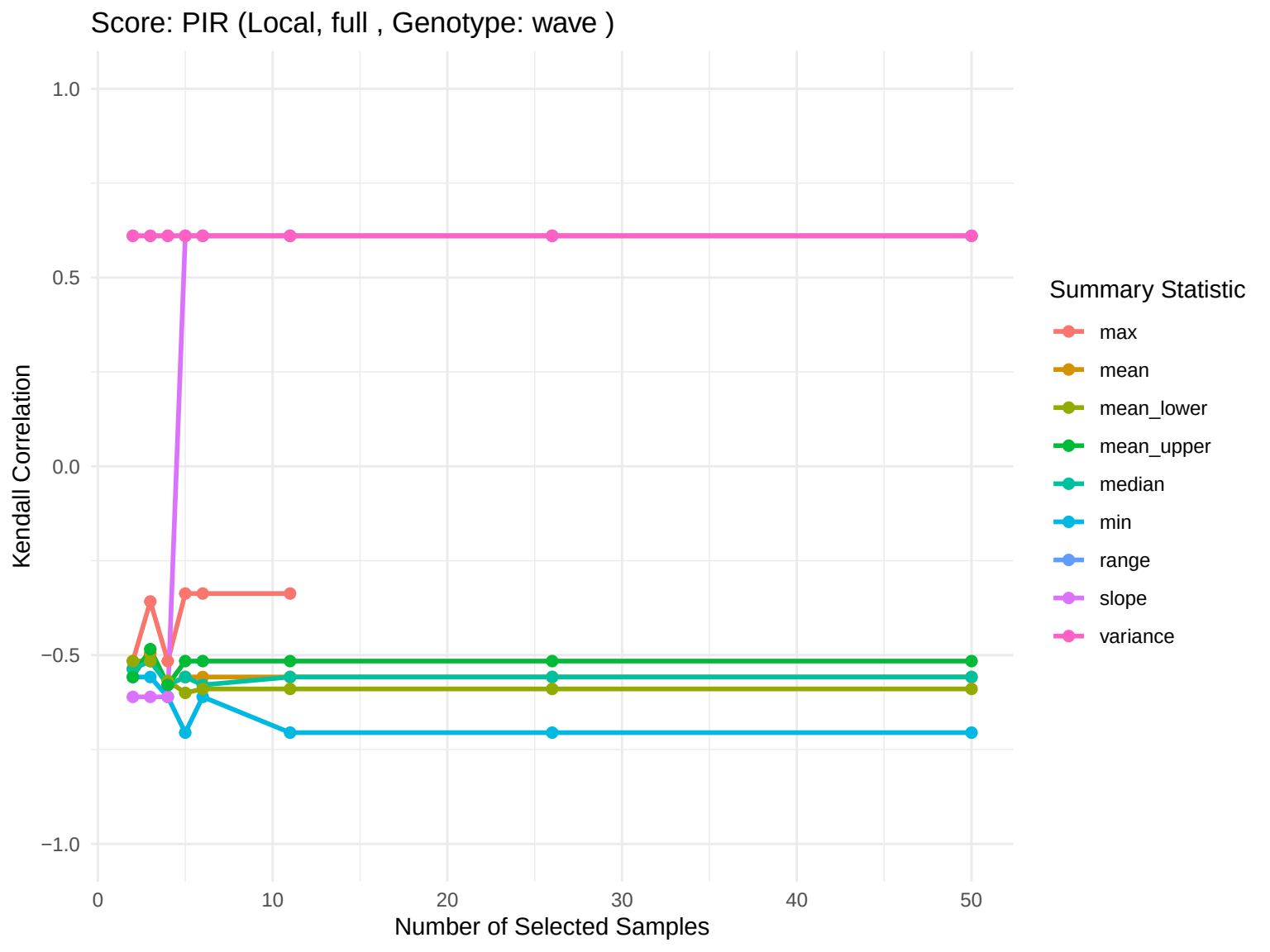
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PIR (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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20
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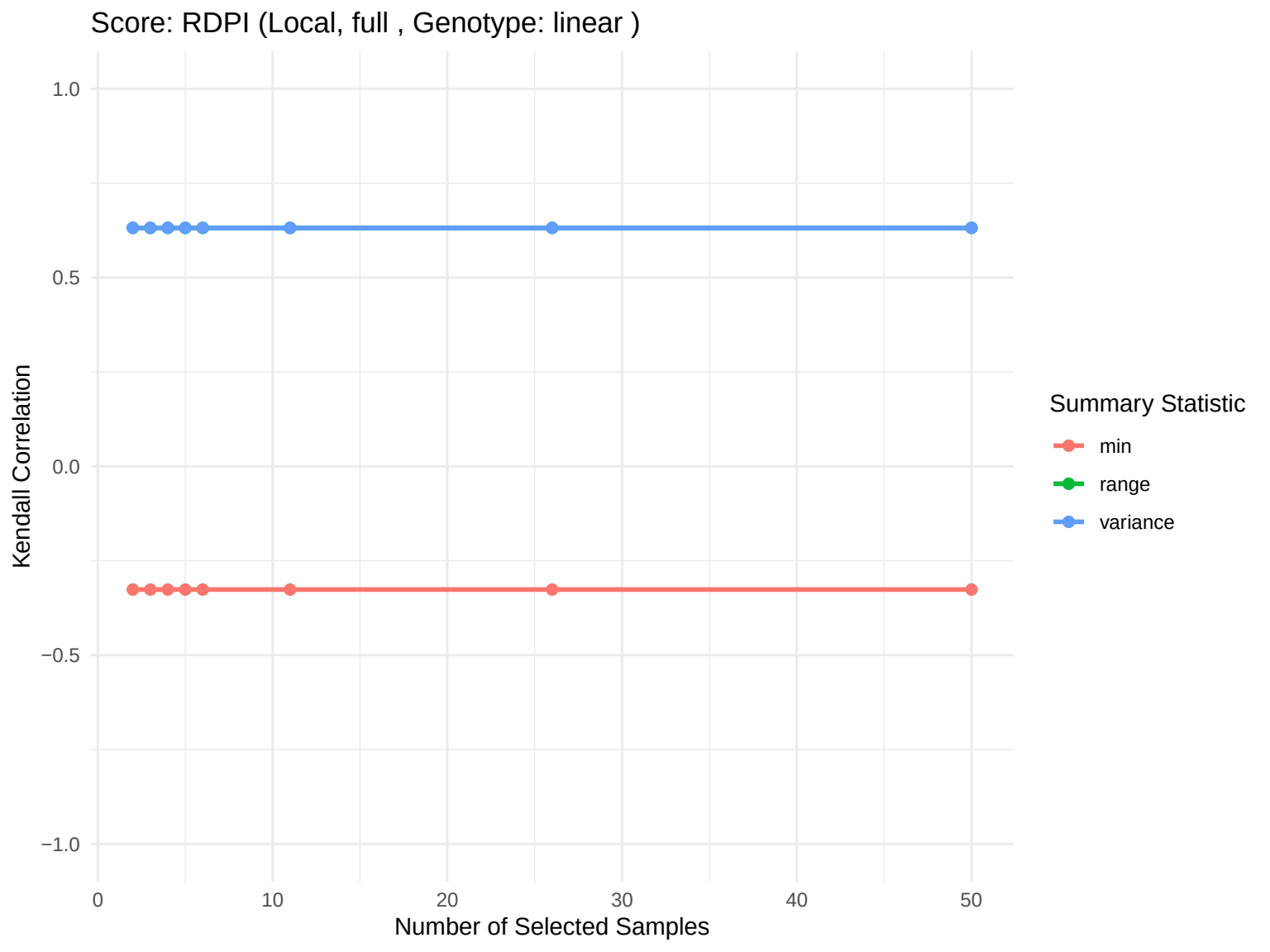
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RDPI (Local, full , Genotype: linear)
Kendall Correlation
1.0
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−1.0
0
10
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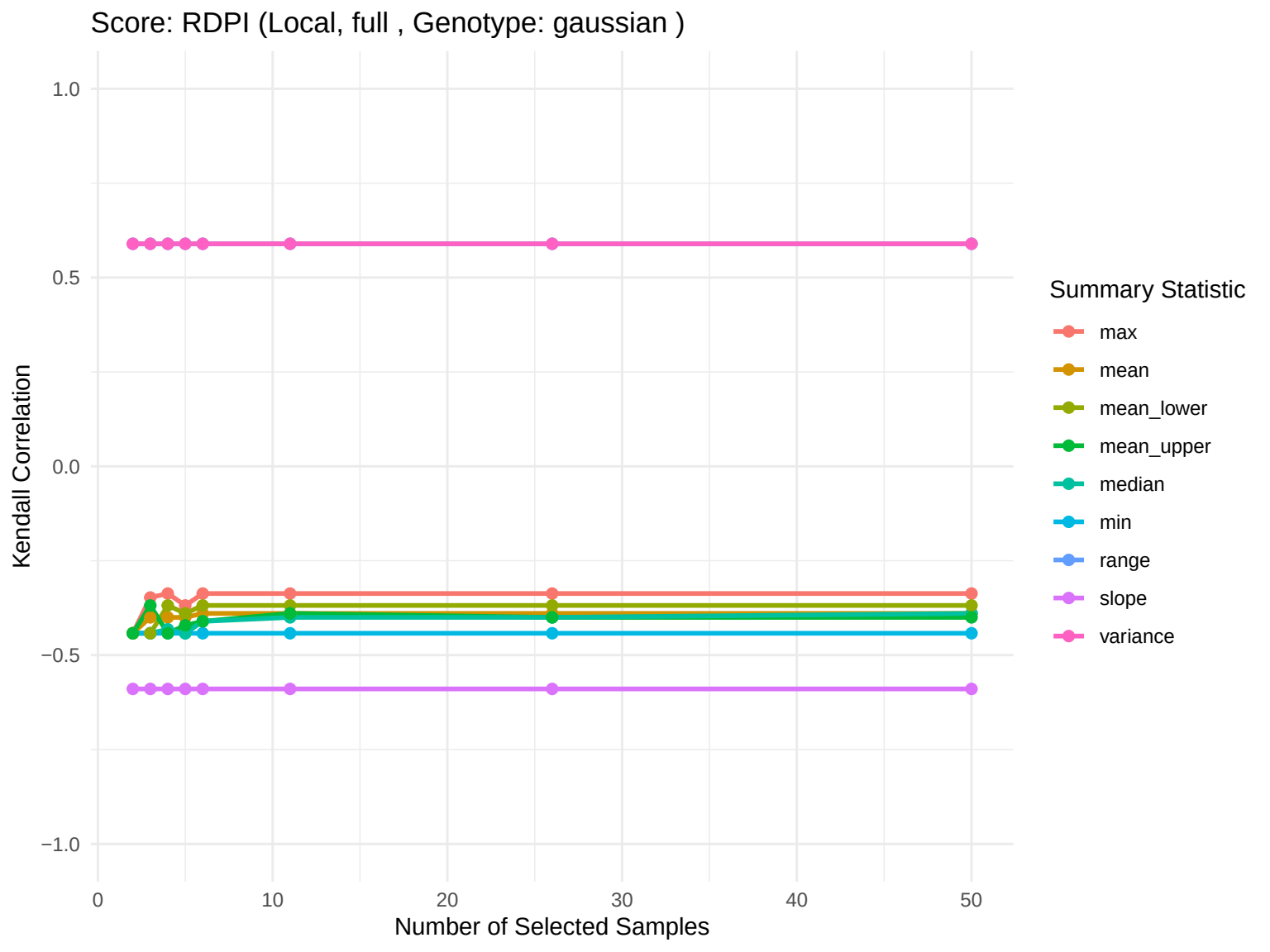
50
Number of Selected Samples
Summary Statistic
min
range
variance


Score: RDPI (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
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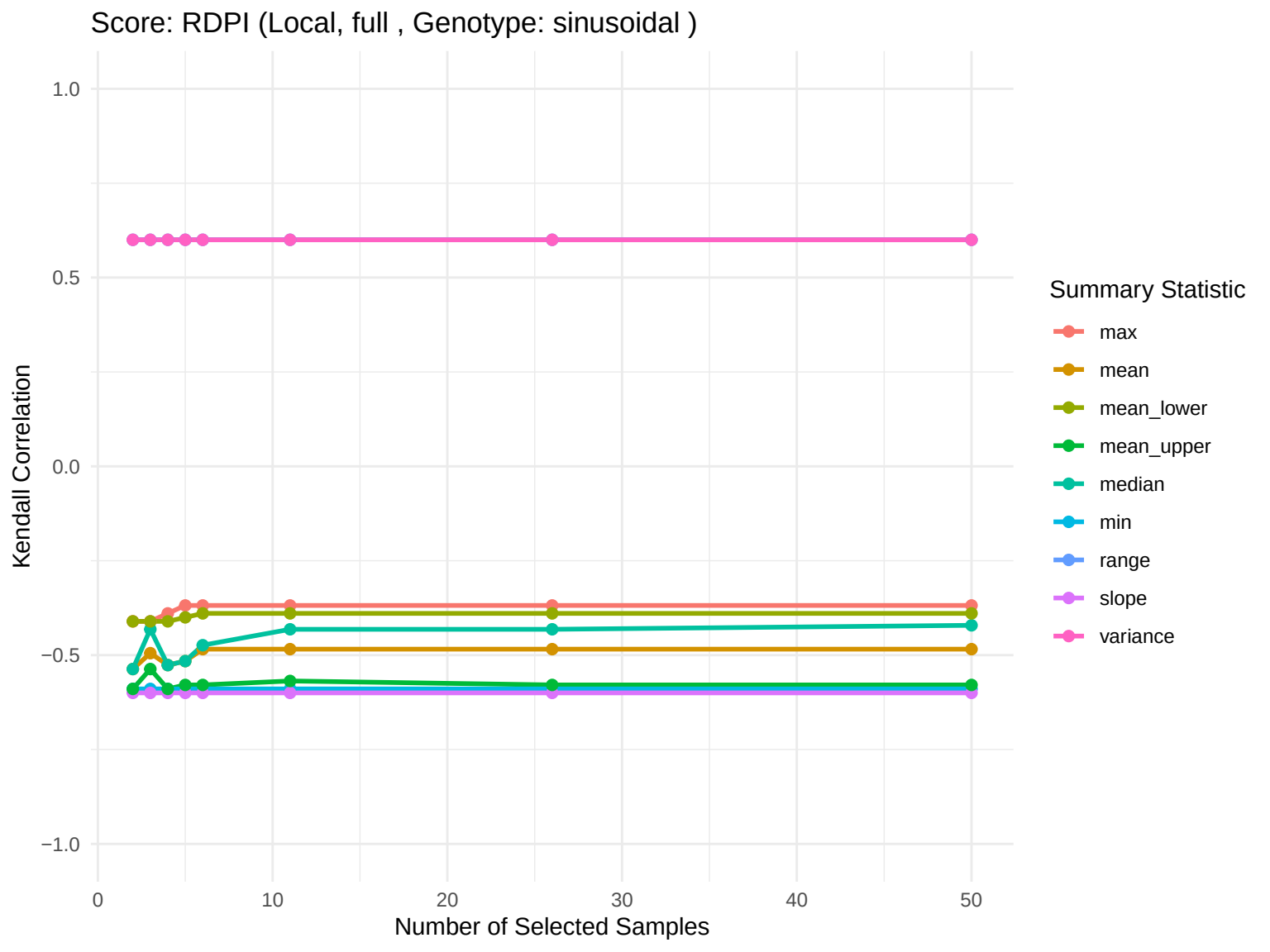
slope
variance


Score: RDPI (Local, full , Genotype: sinusoidal)
Kendall Correlation
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0.0
−0.5
−1.0
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Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
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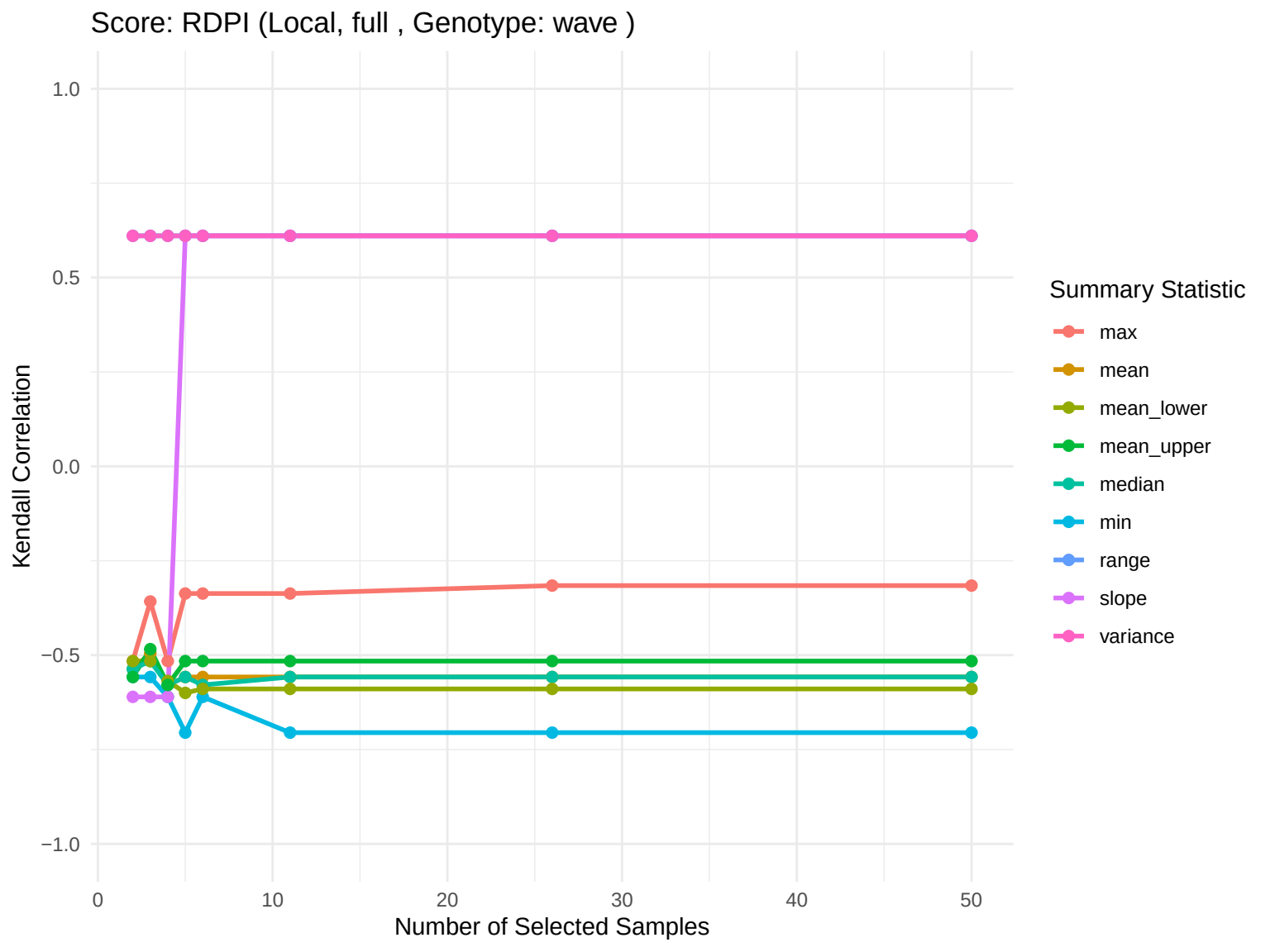
range
slope
variance


Score: RDPI (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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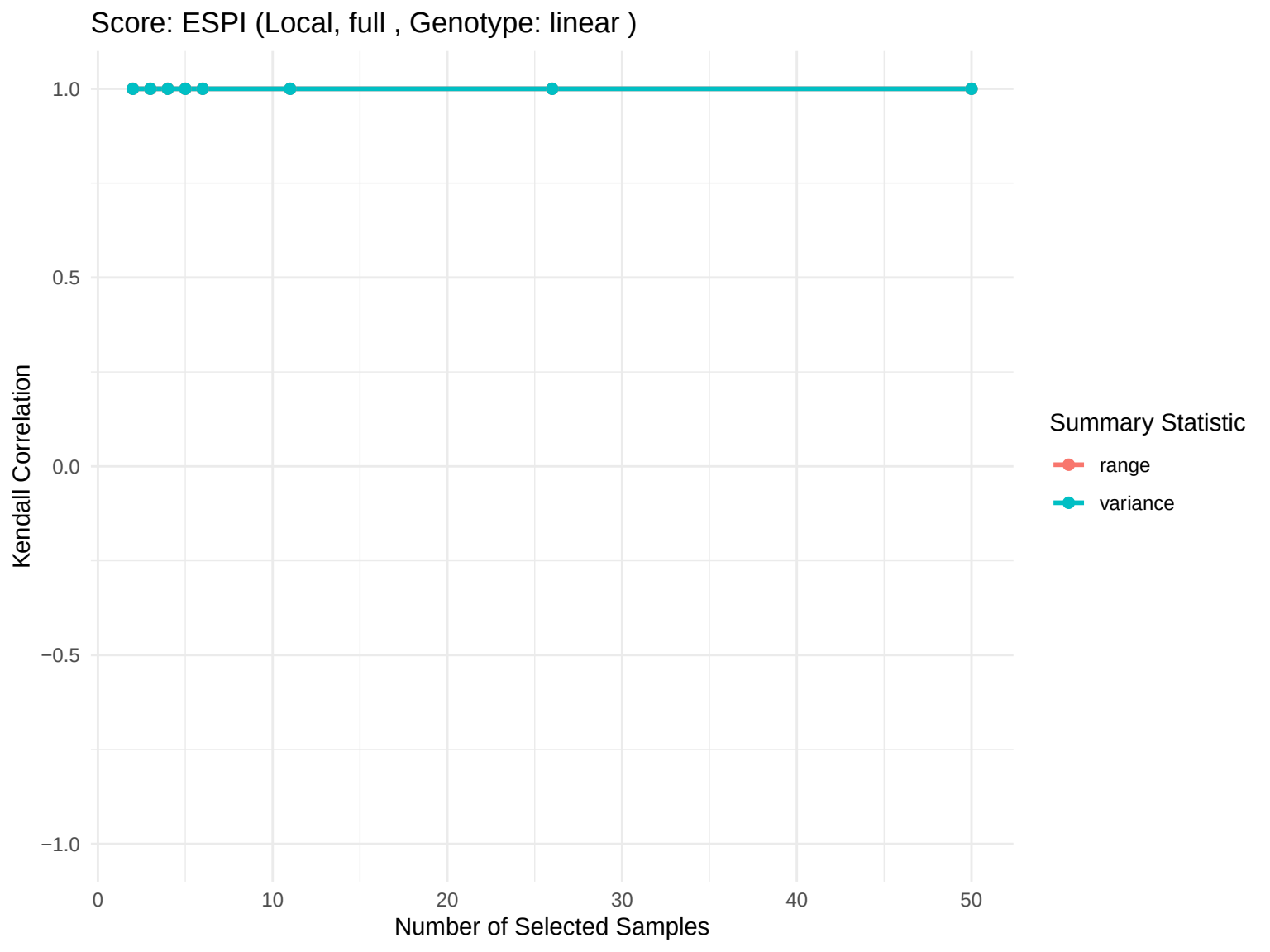
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: ESPI (Local, full , Genotype: linear)
Kendall Correlation
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−0.5
−1.0
0
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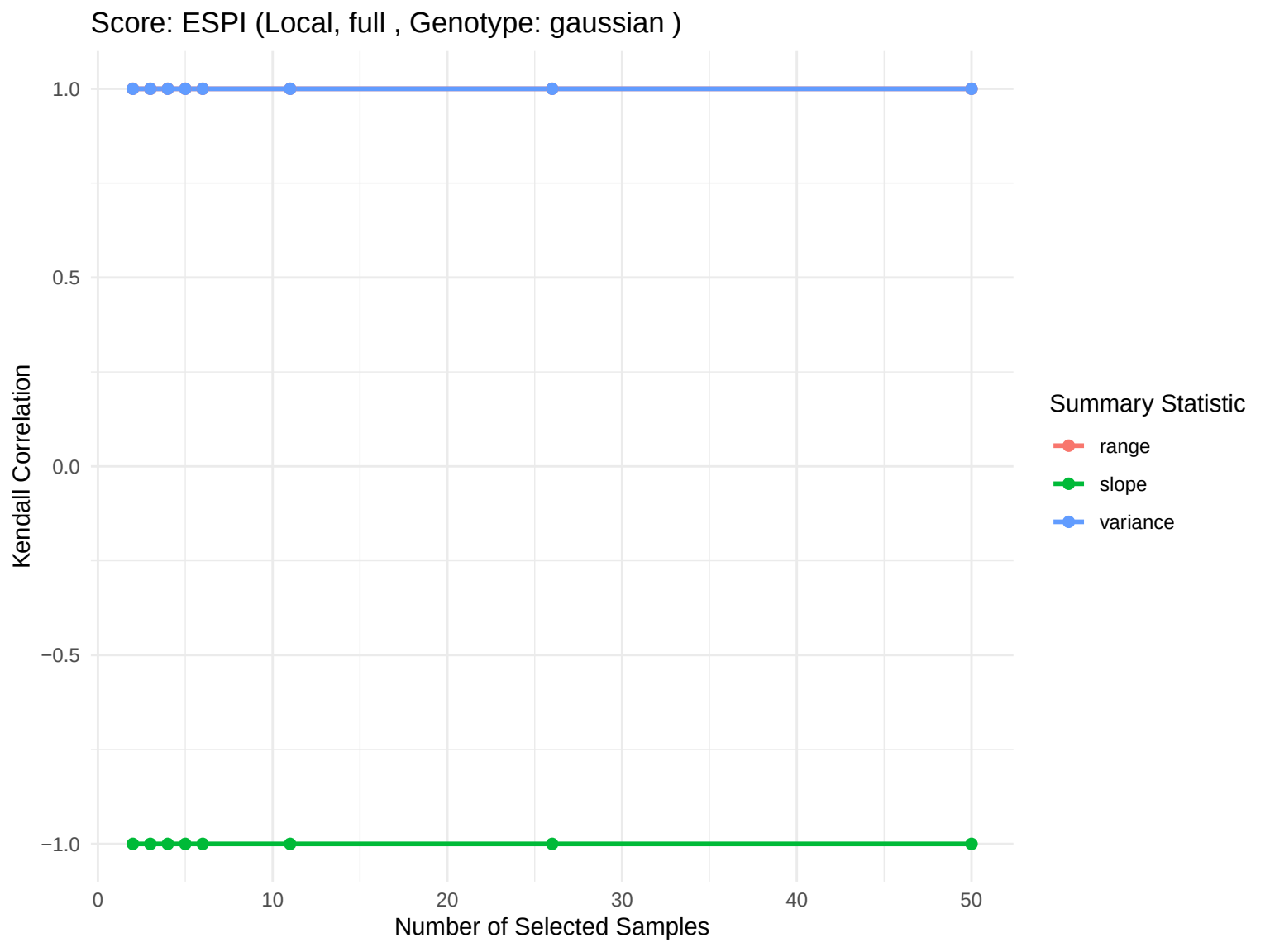
40
50
Number of Selected Samples
Summary Statistic
range
variance


Score: ESPI (Local, full , Genotype: gaussian)
Kendall Correlation
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−1.0
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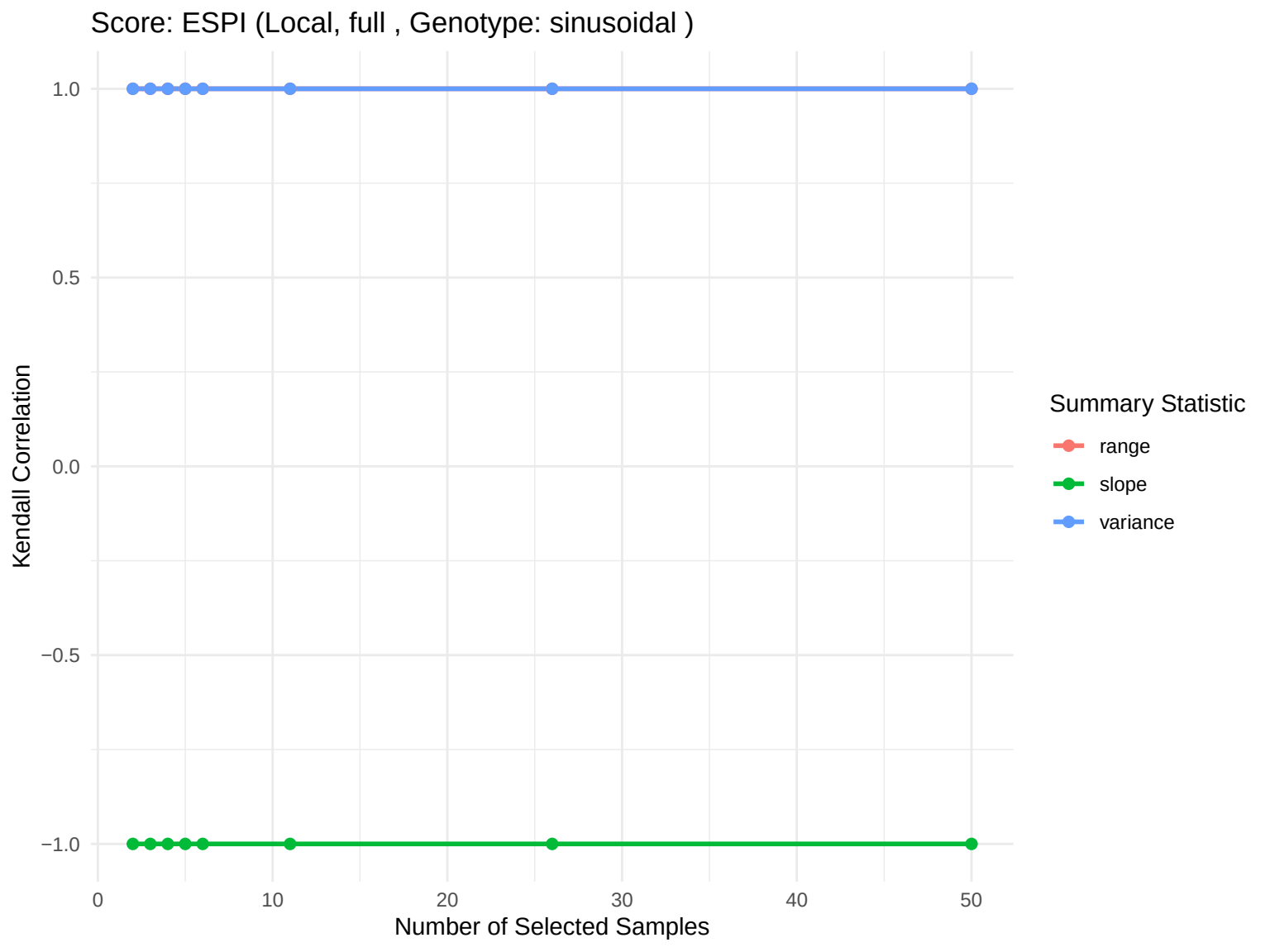
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: ESPI (Local, full , Genotype: sinusoidal)
Kendall Correlation
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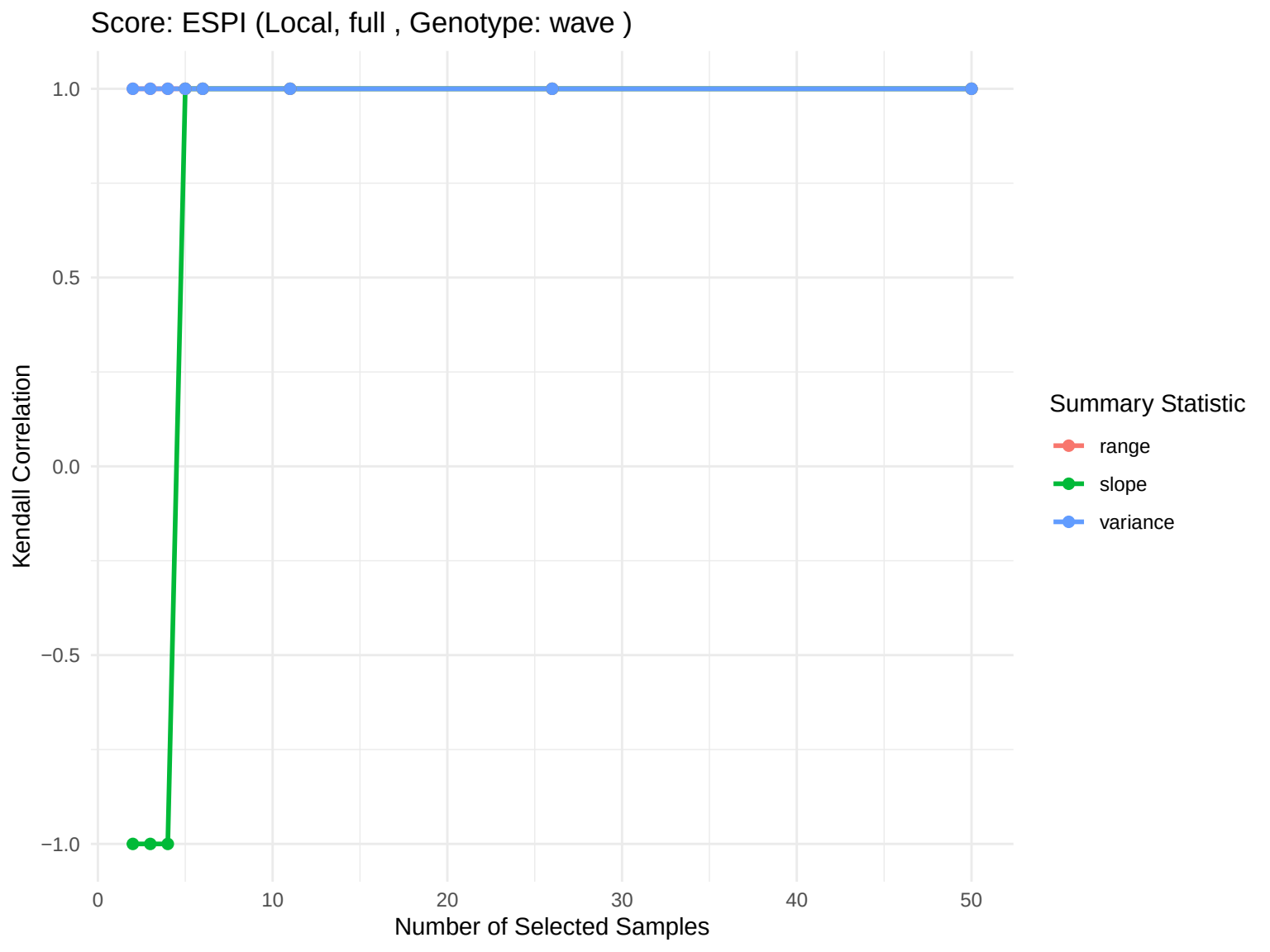
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: ESPI (Local, full , Genotype: wave)
Kendall Correlation
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−1.0
0
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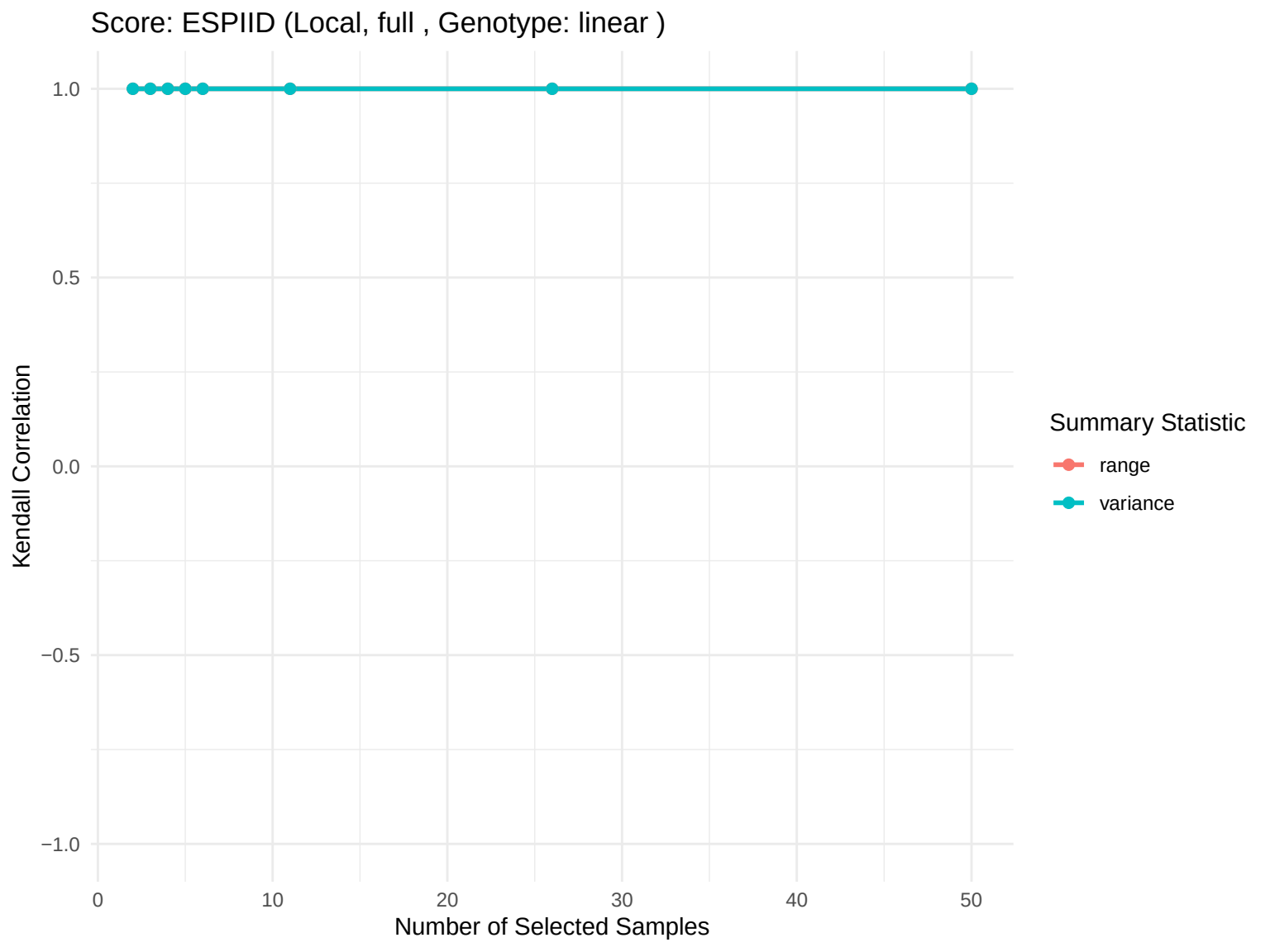
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: ESPIID (Local, full , Genotype: linear)
Kendall Correlation
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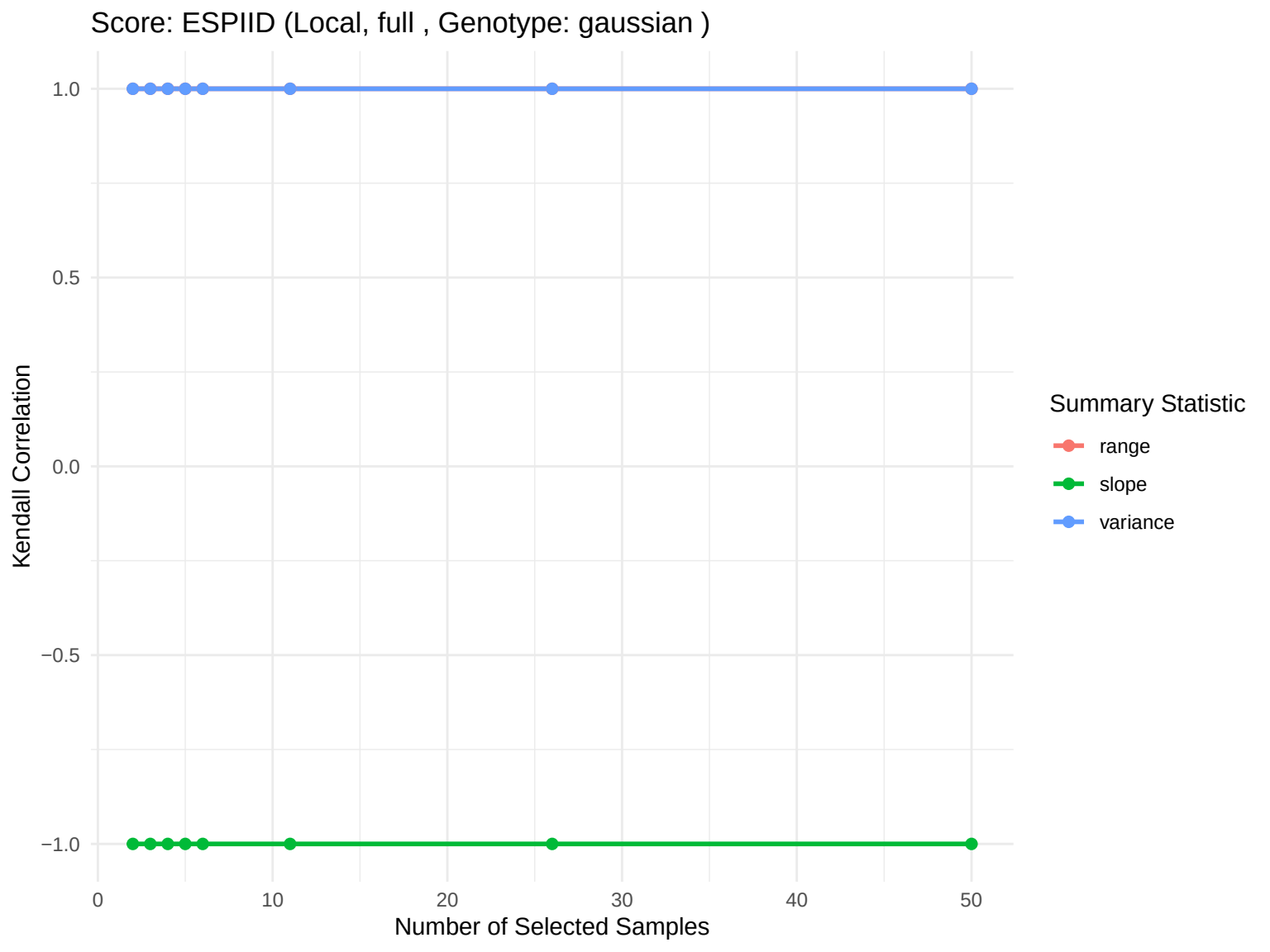
Number of Selected Samples
Summary Statistic
range
variance


Score: ESPIID (Local, full , Genotype: gaussian)
1.0
0.5
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−1.0
Kendall Correlation
0
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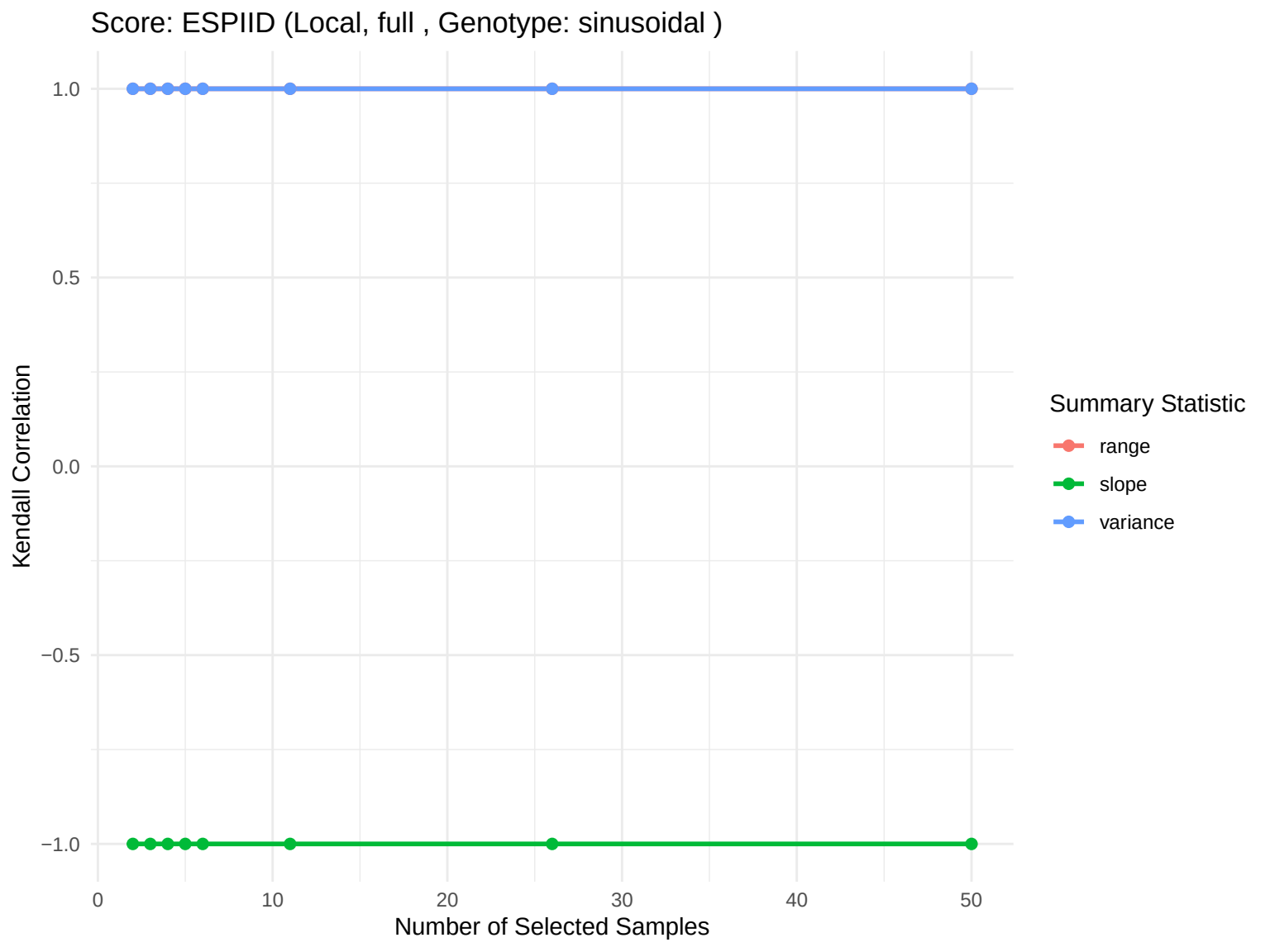
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: ESPIID (Local, full , Genotype: sinusoidal)
Kendall Correlation
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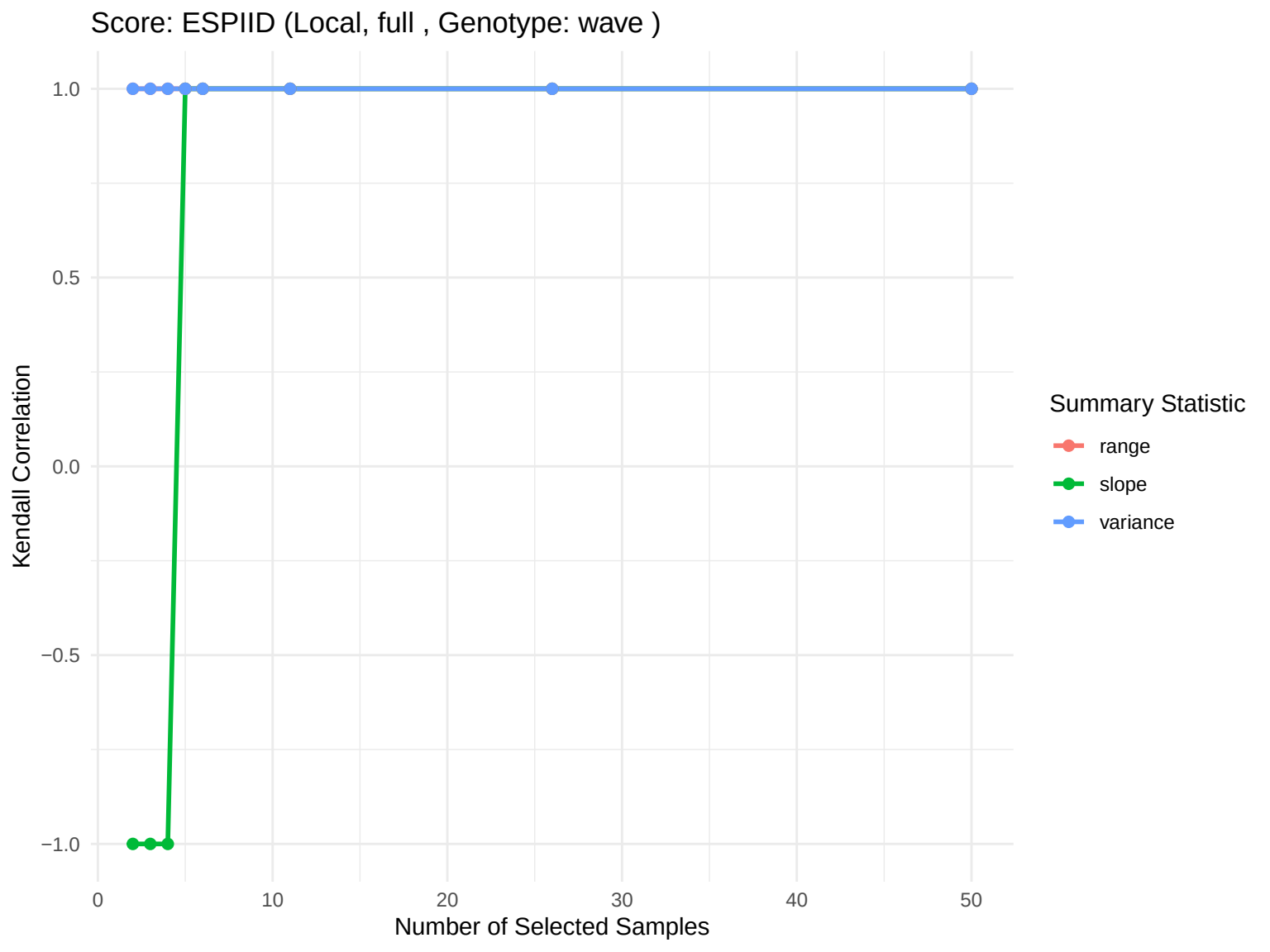
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: ESPIID (Local, full , Genotype: wave)
Kendall Correlation
1.0
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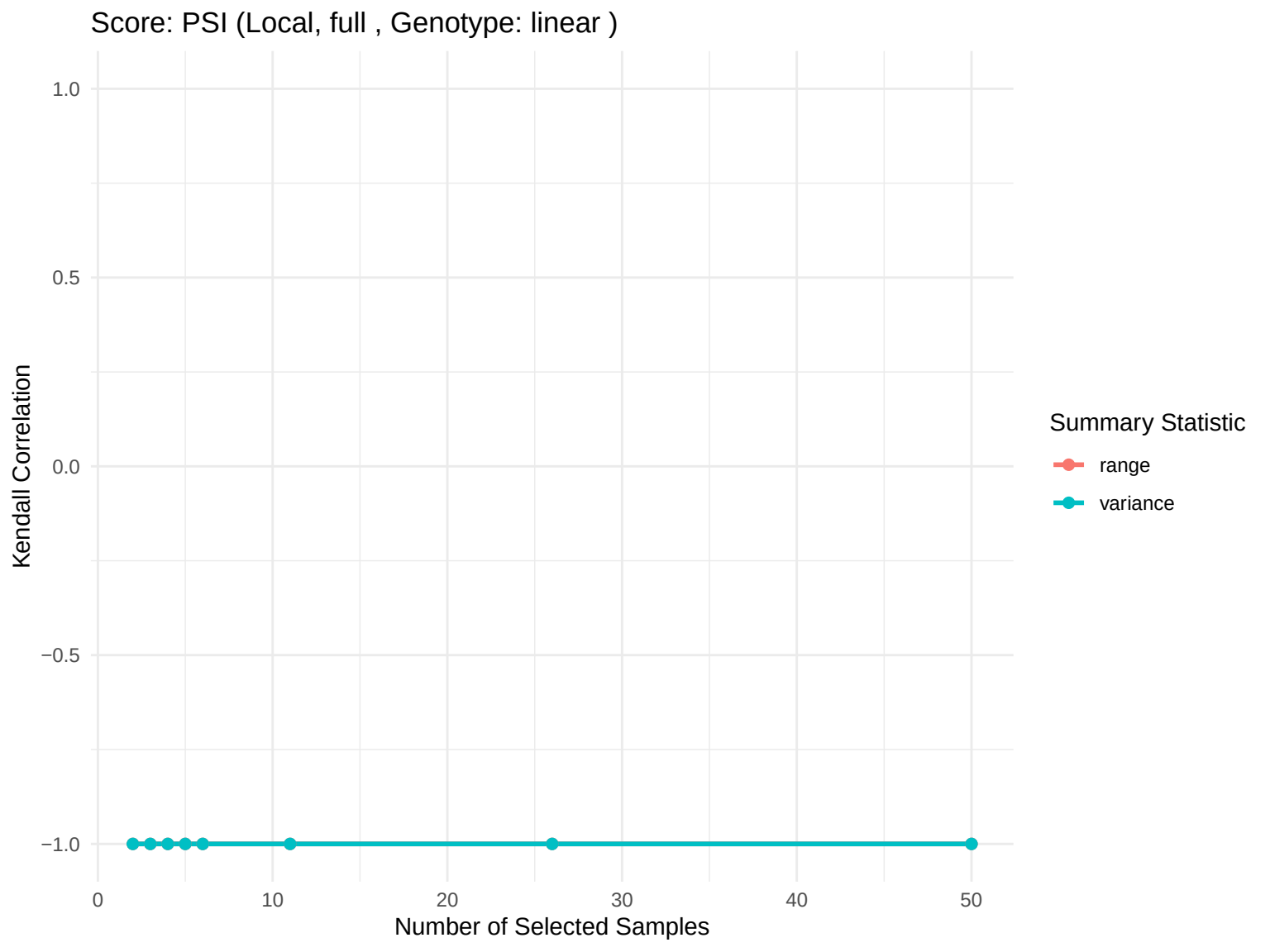
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: PSI (Local, full , Genotype: linear)
Kendall Correlation
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−0.5
−1.0
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40
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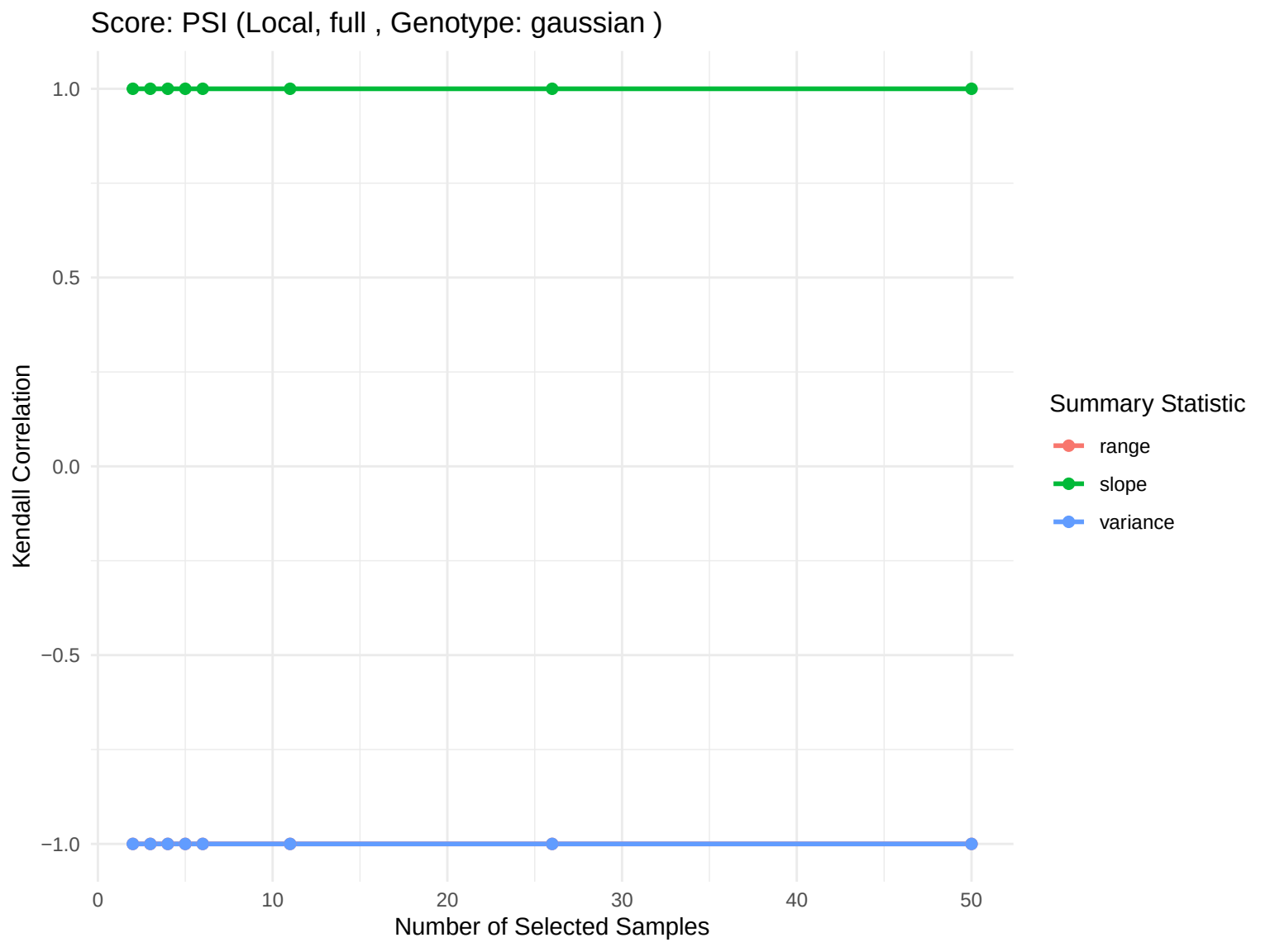
Number of Selected Samples
Summary Statistic
range
variance


Score: PSI (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
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−1.0
0
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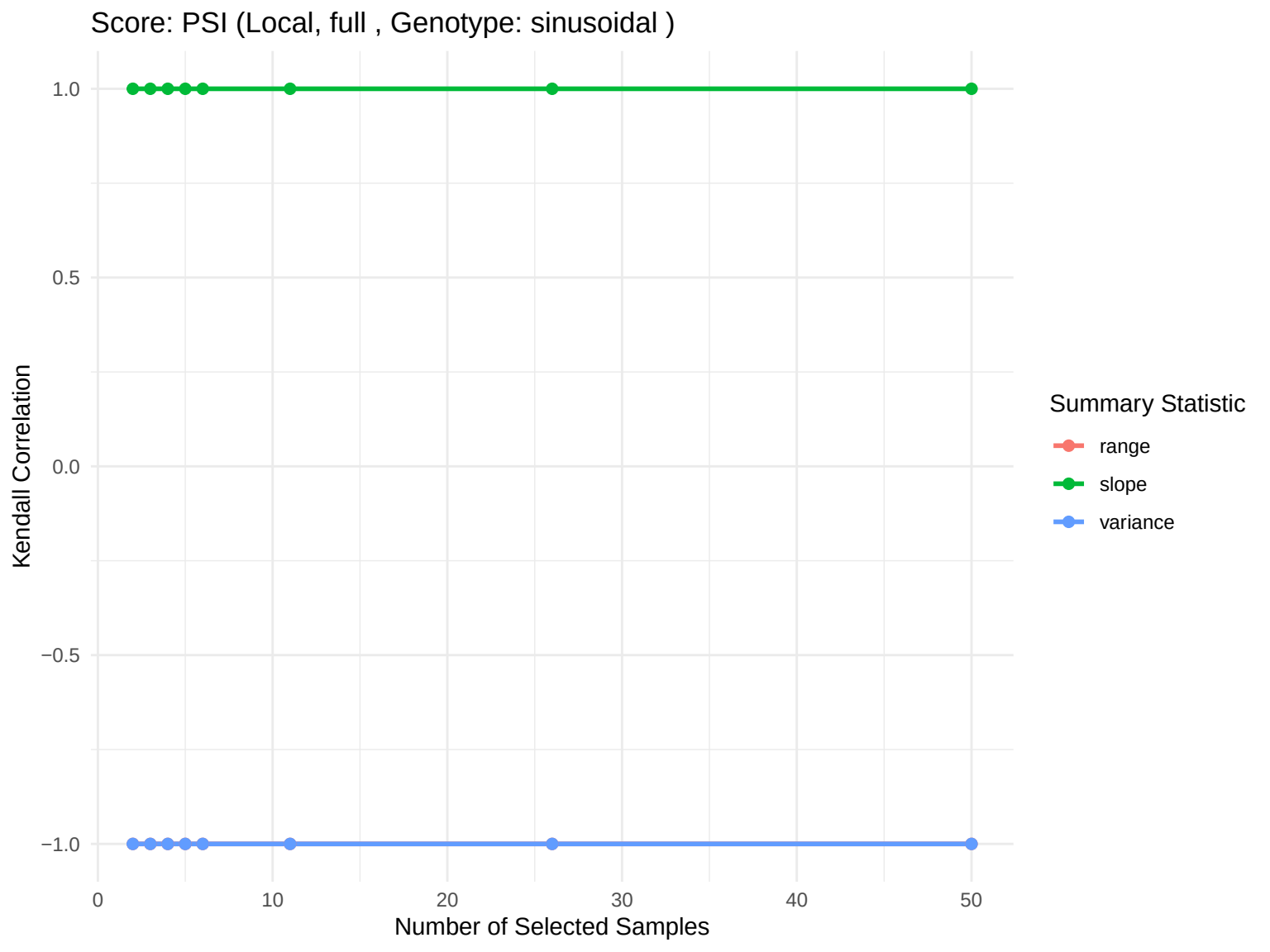
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: PSI (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
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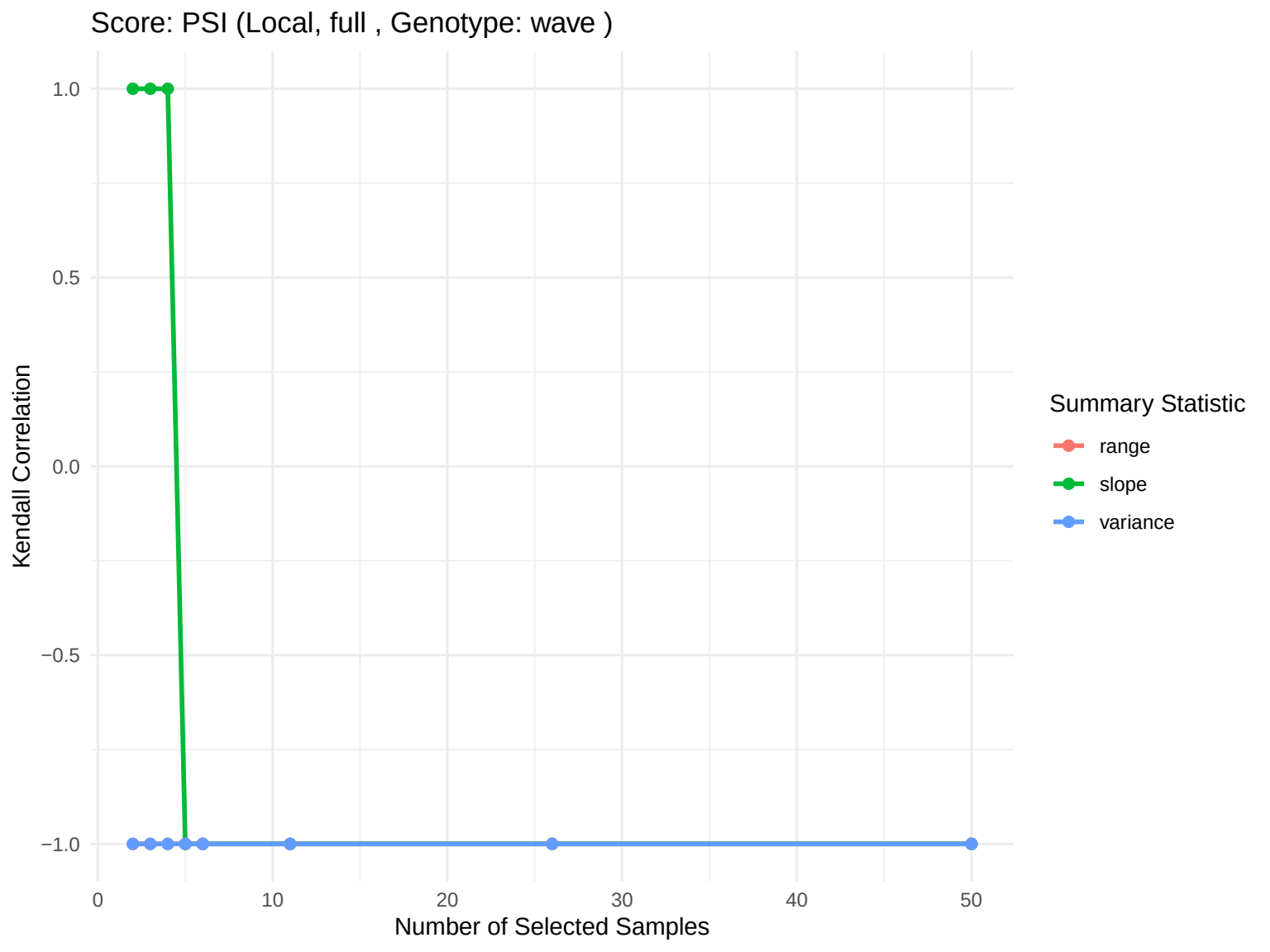
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: PSI (Local, full , Genotype: wave)
Kendall Correlation
1.0
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0.0
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−1.0
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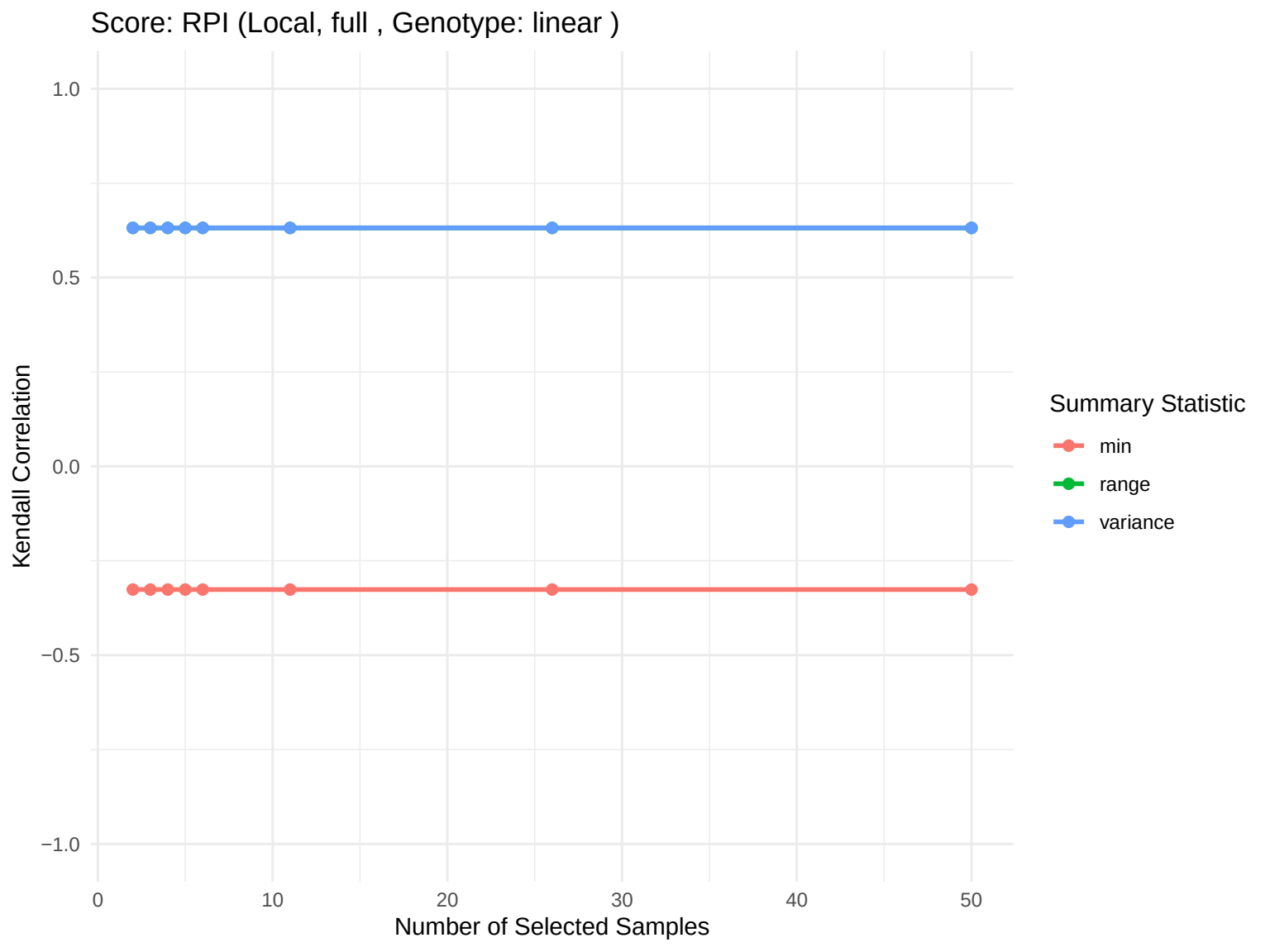
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: RPI (Local, full , Genotype: linear)
Kendall Correlation
1.0
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0.0
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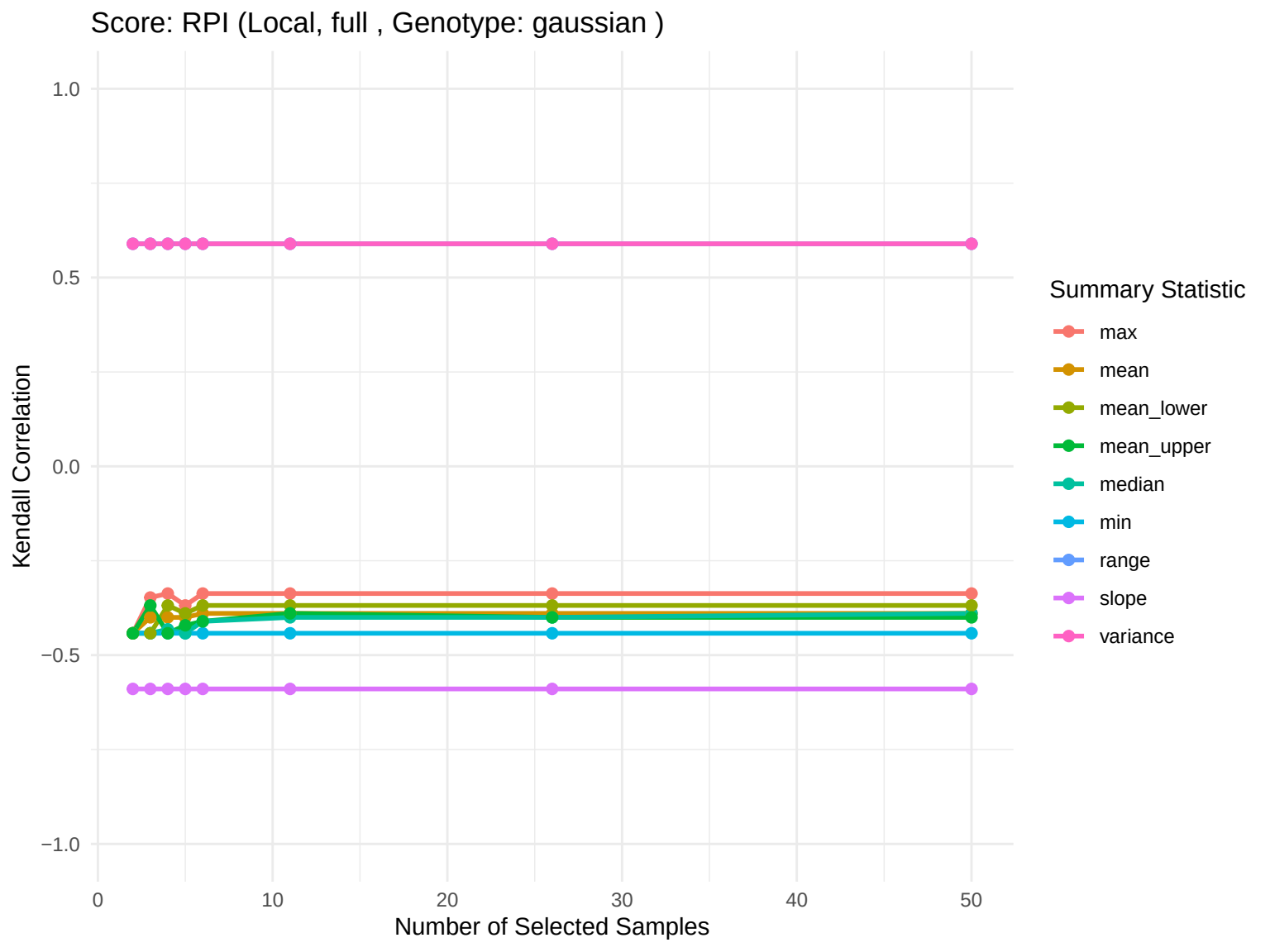
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
min
range
variance


Score: RPI (Local, full , Genotype: gaussian)
Kendall Correlation
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0.5
0.0
−0.5
−1.0
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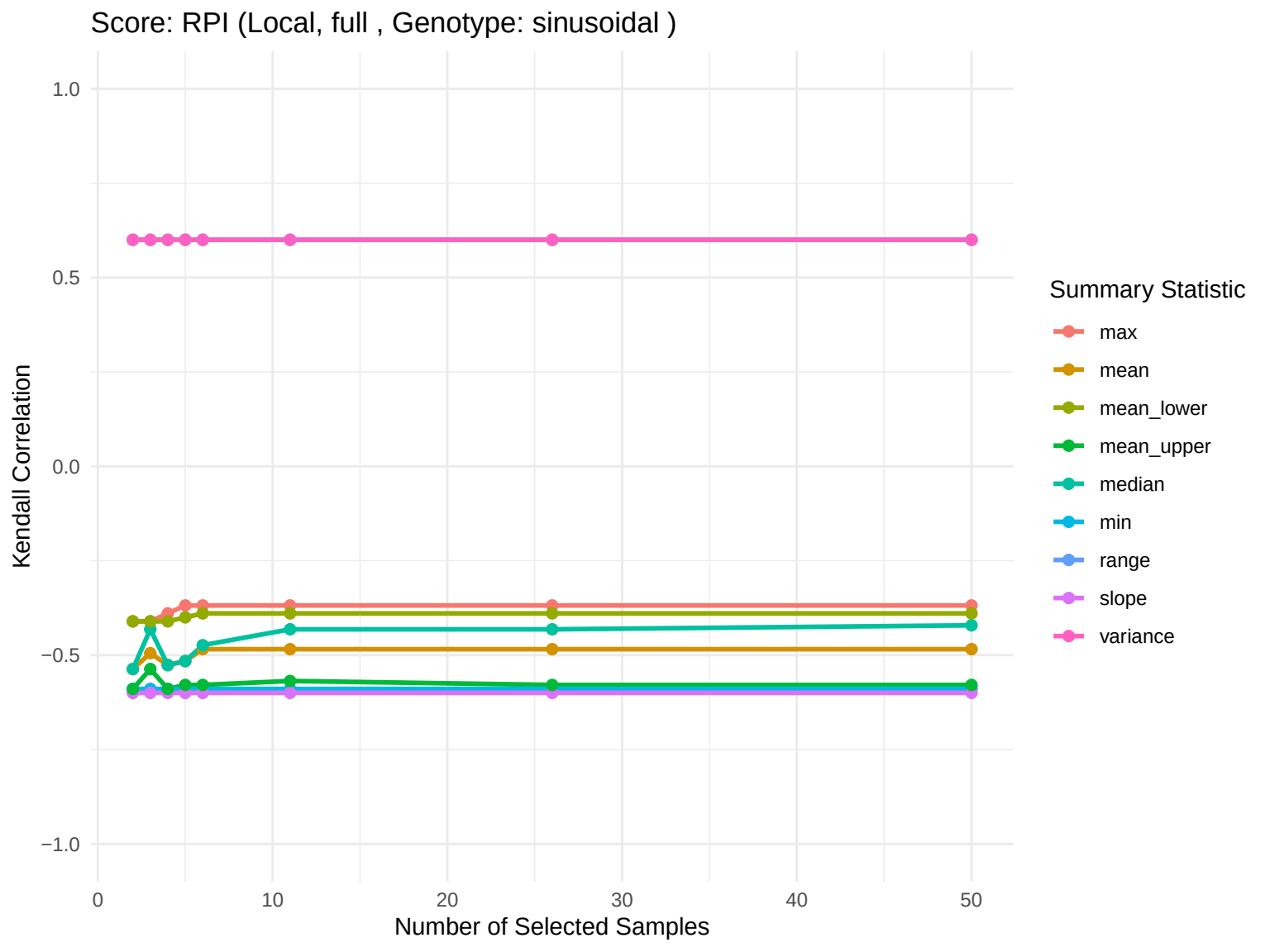
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RPI (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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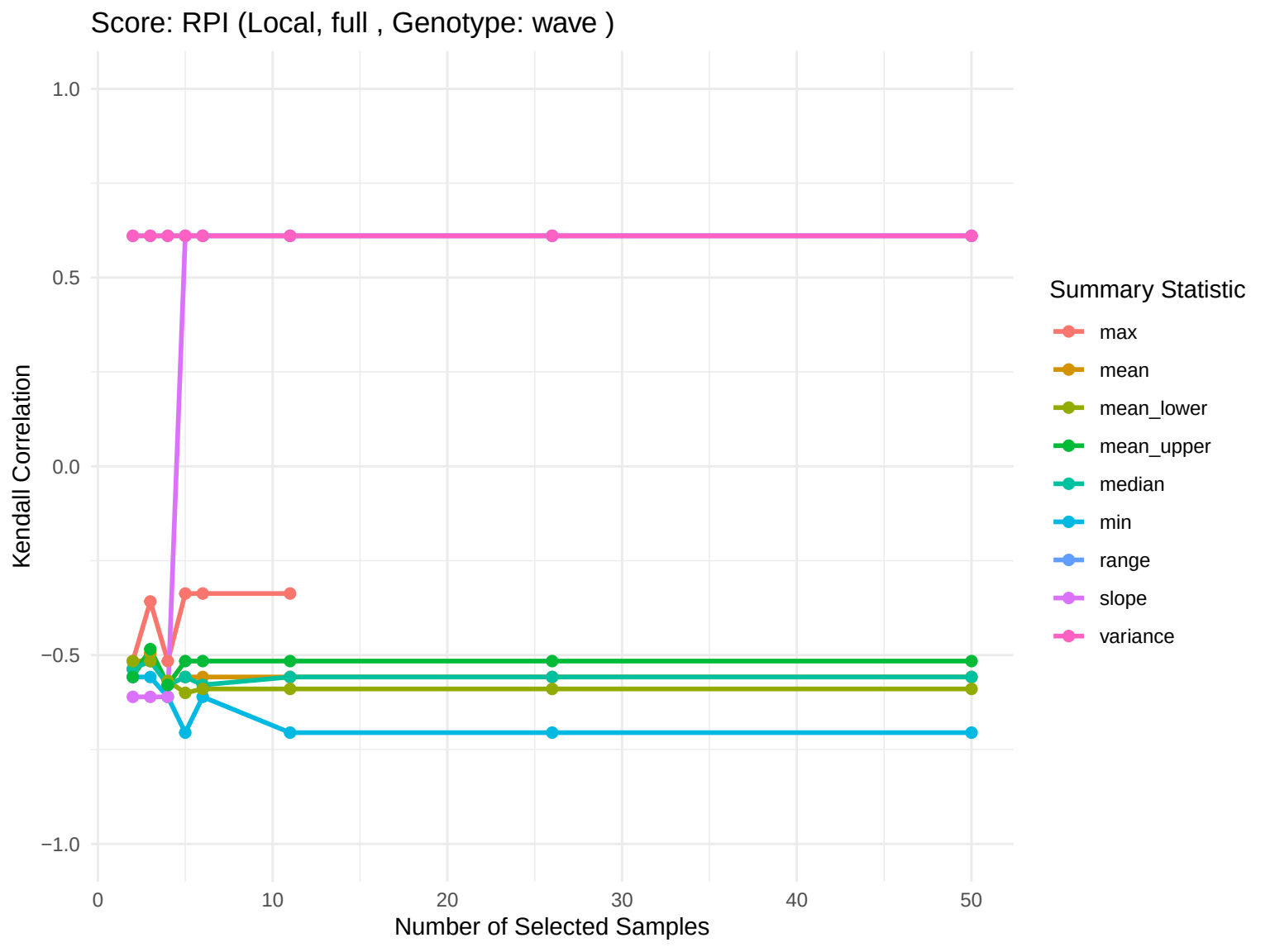
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RPI (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
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40
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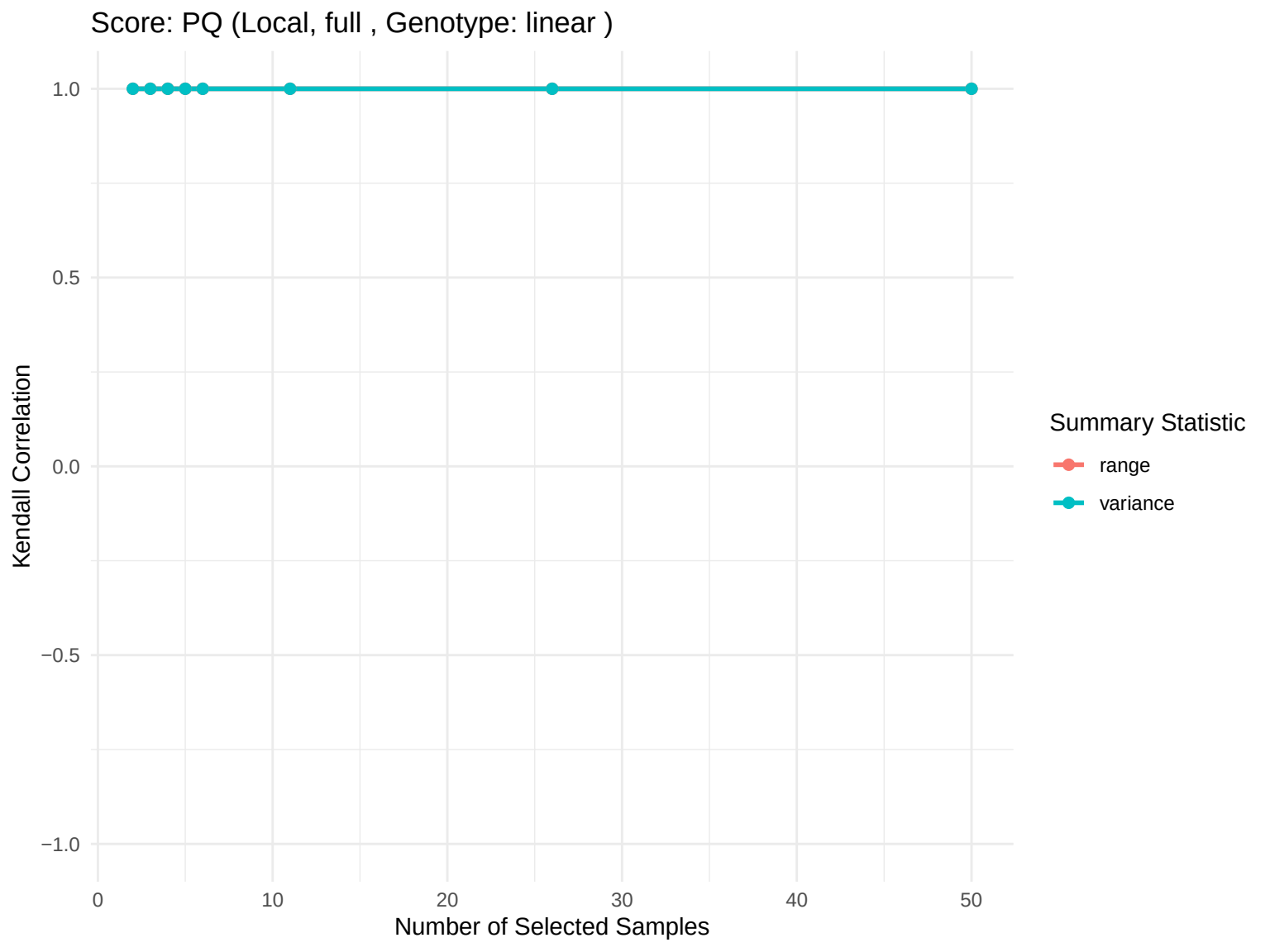
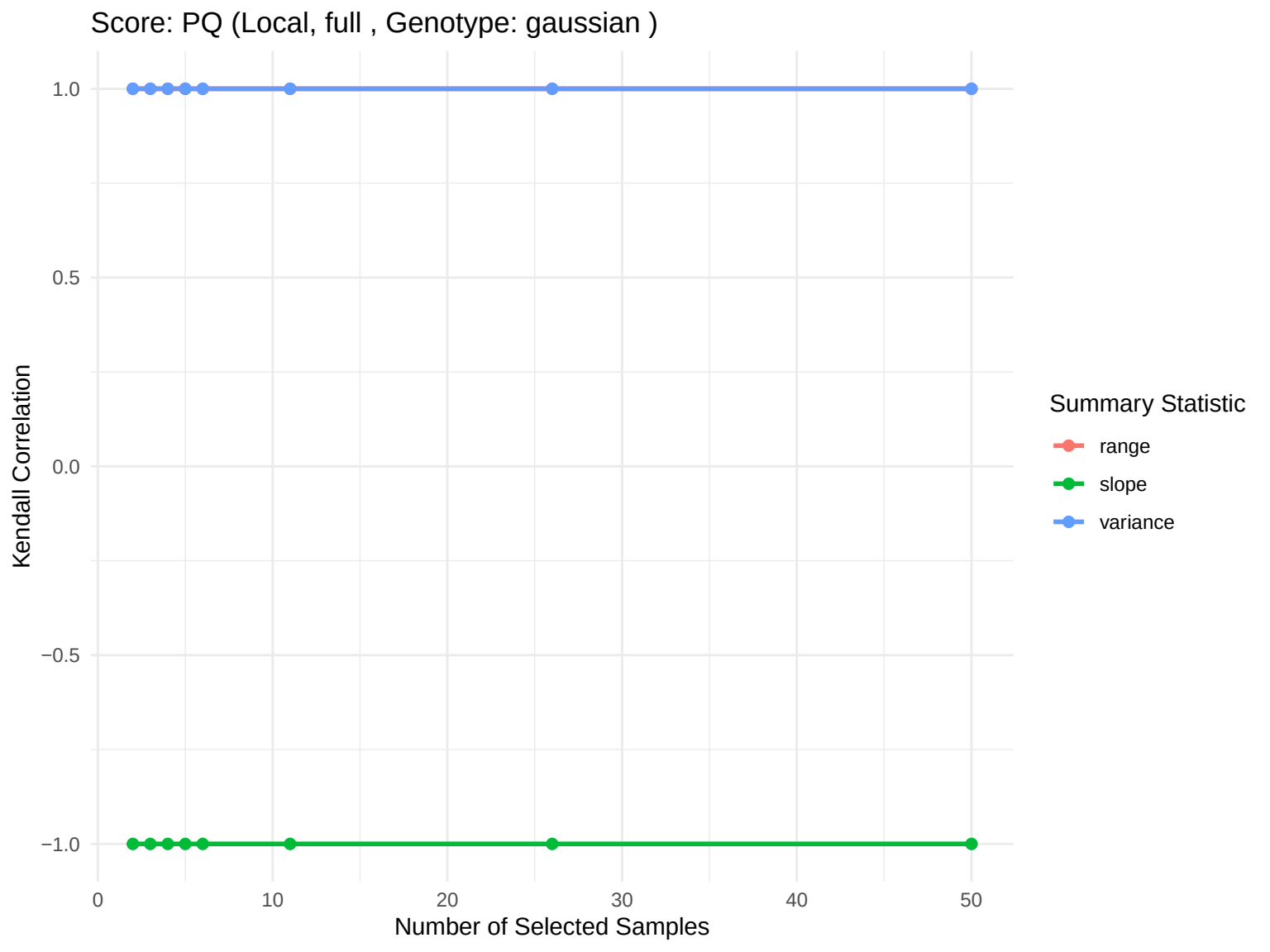
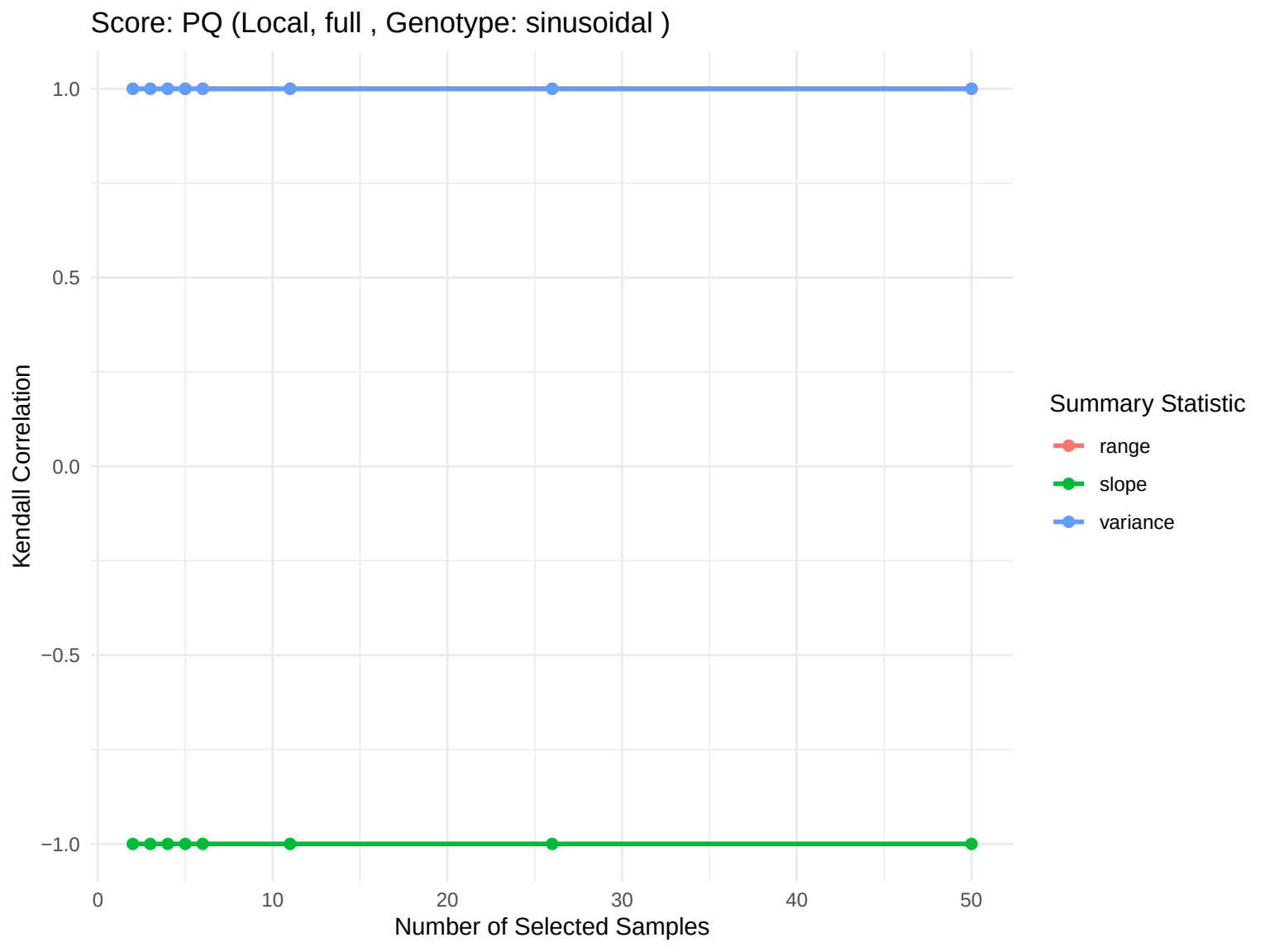
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PQ (Local, full , Genotype: linear)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
variance


Score: PQ (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: PQ (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
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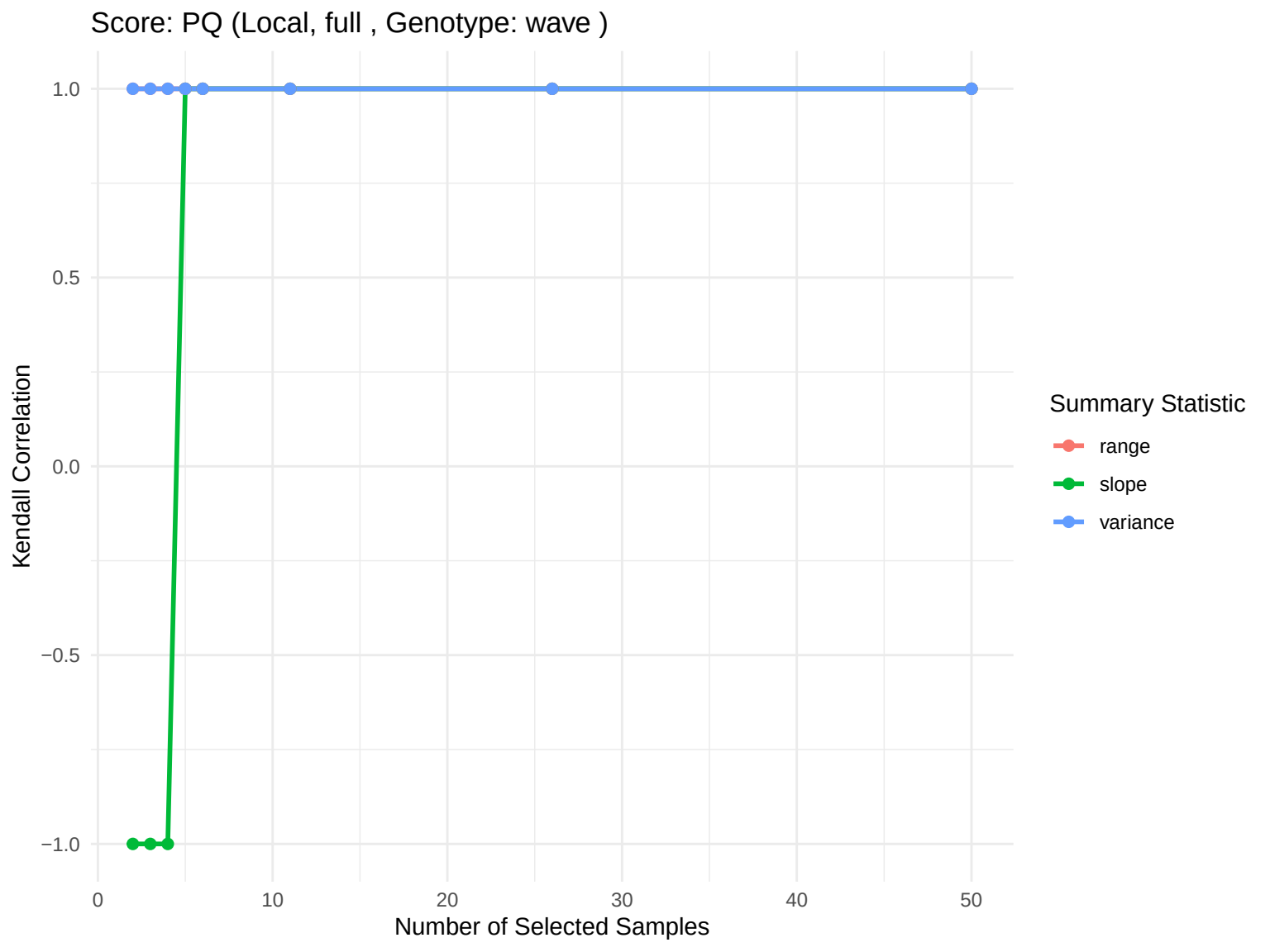
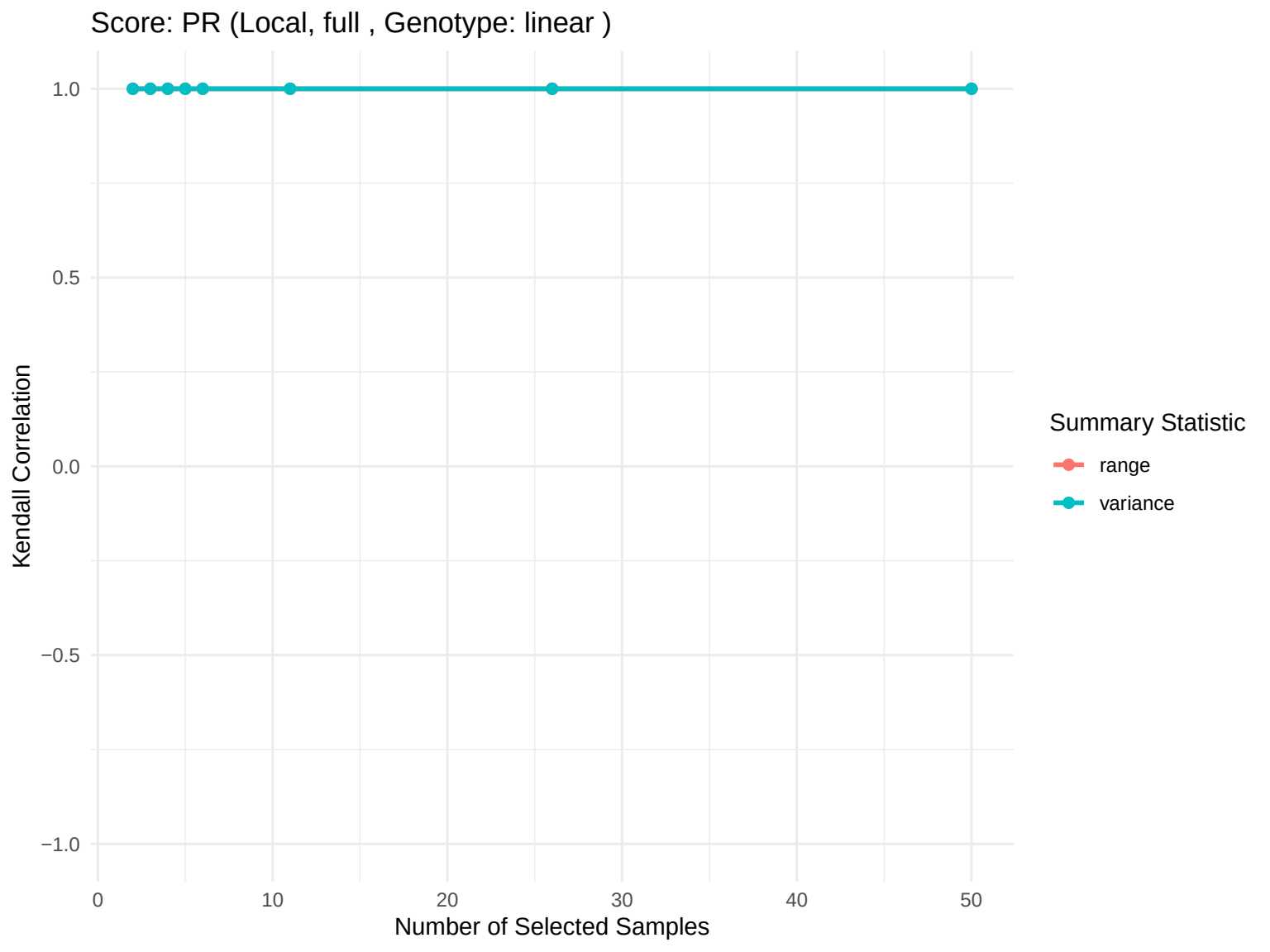
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: PQ (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: PR (Local, full , Genotype: linear)
Kendall Correlation
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0.5
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−0.5
−1.0
0
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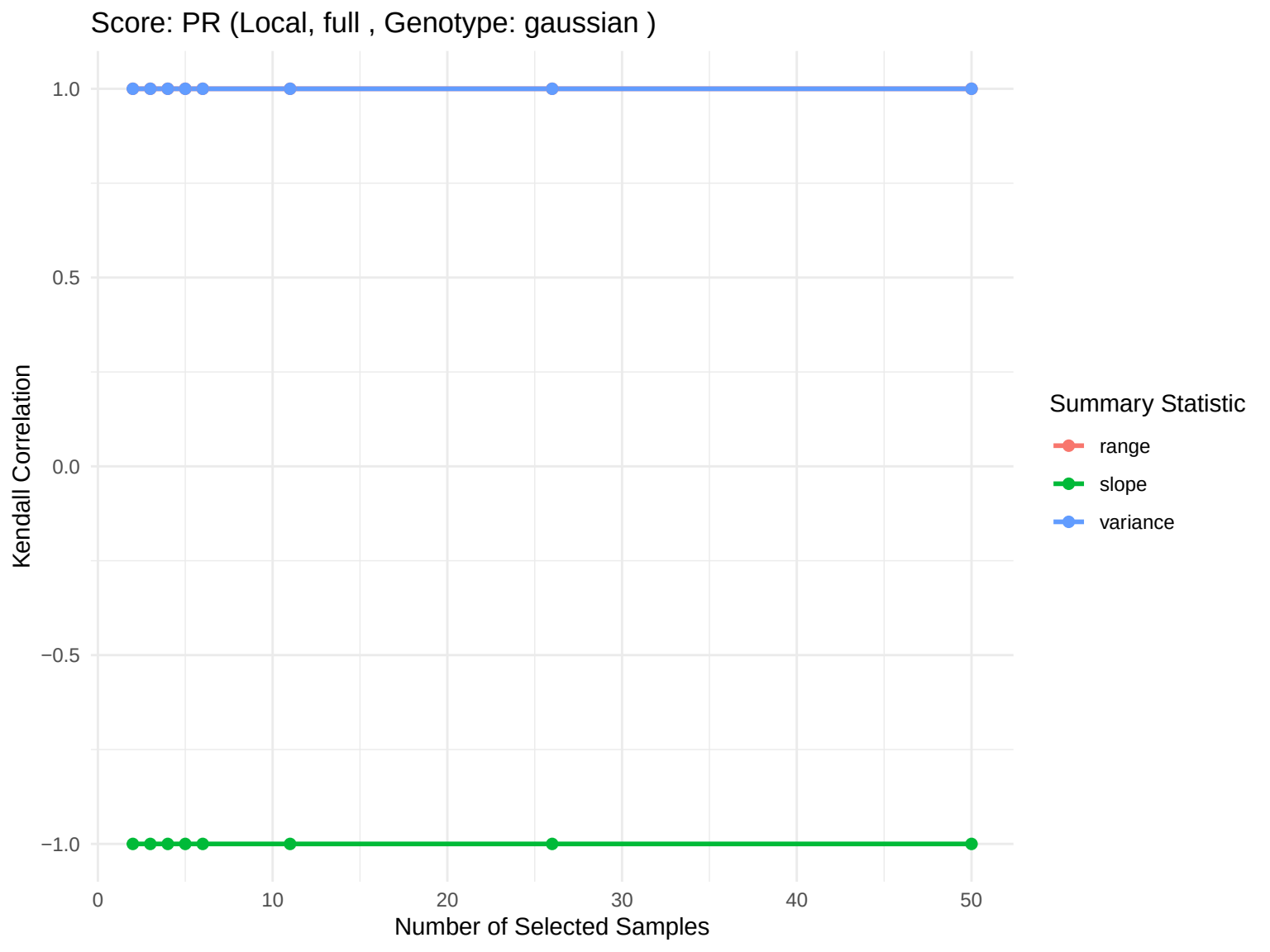
40
50
Number of Selected Samples
Summary Statistic
range
variance


Score: PR (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
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−1.0
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10
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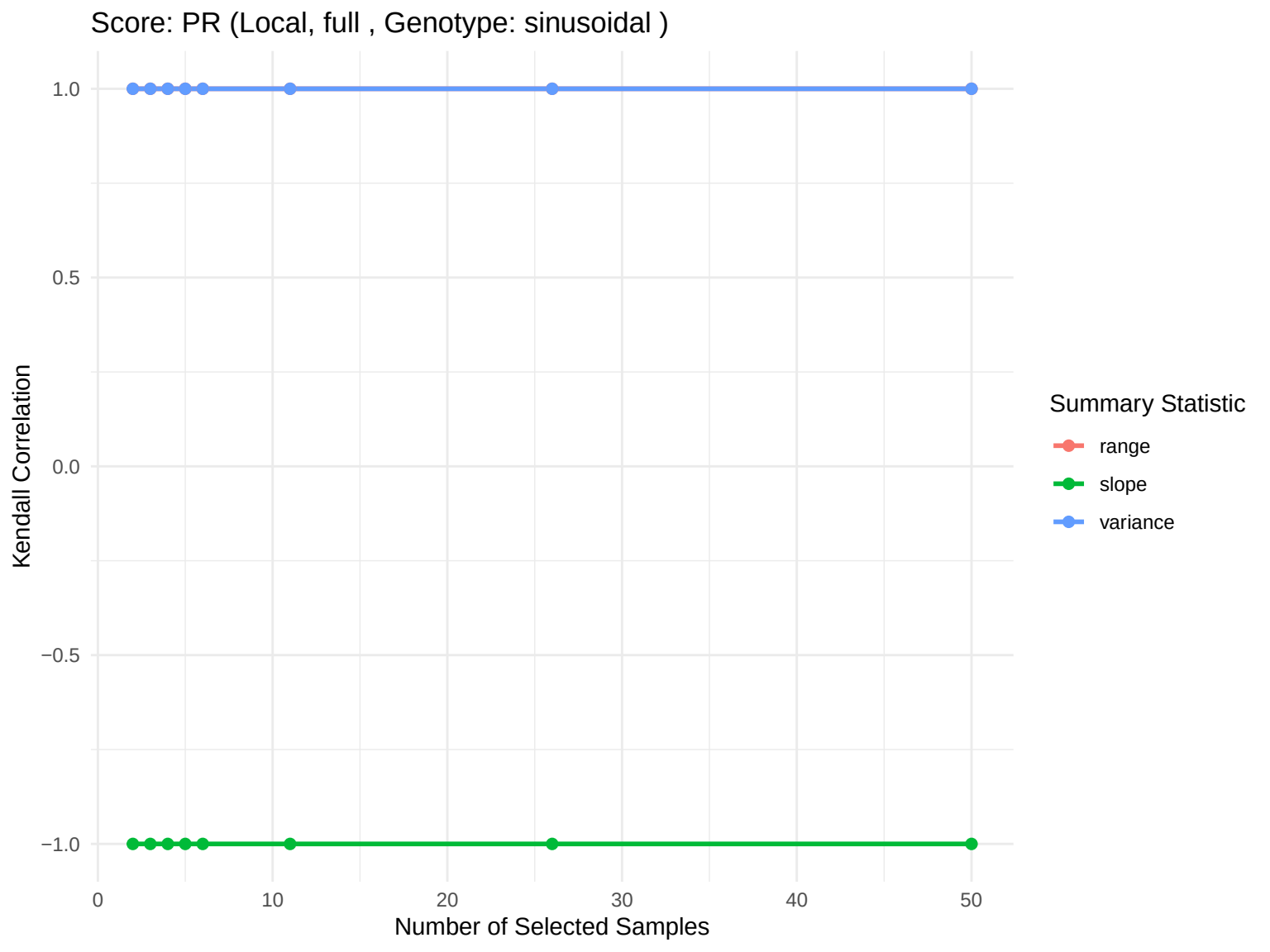
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: PR (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
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−1.0
0
10
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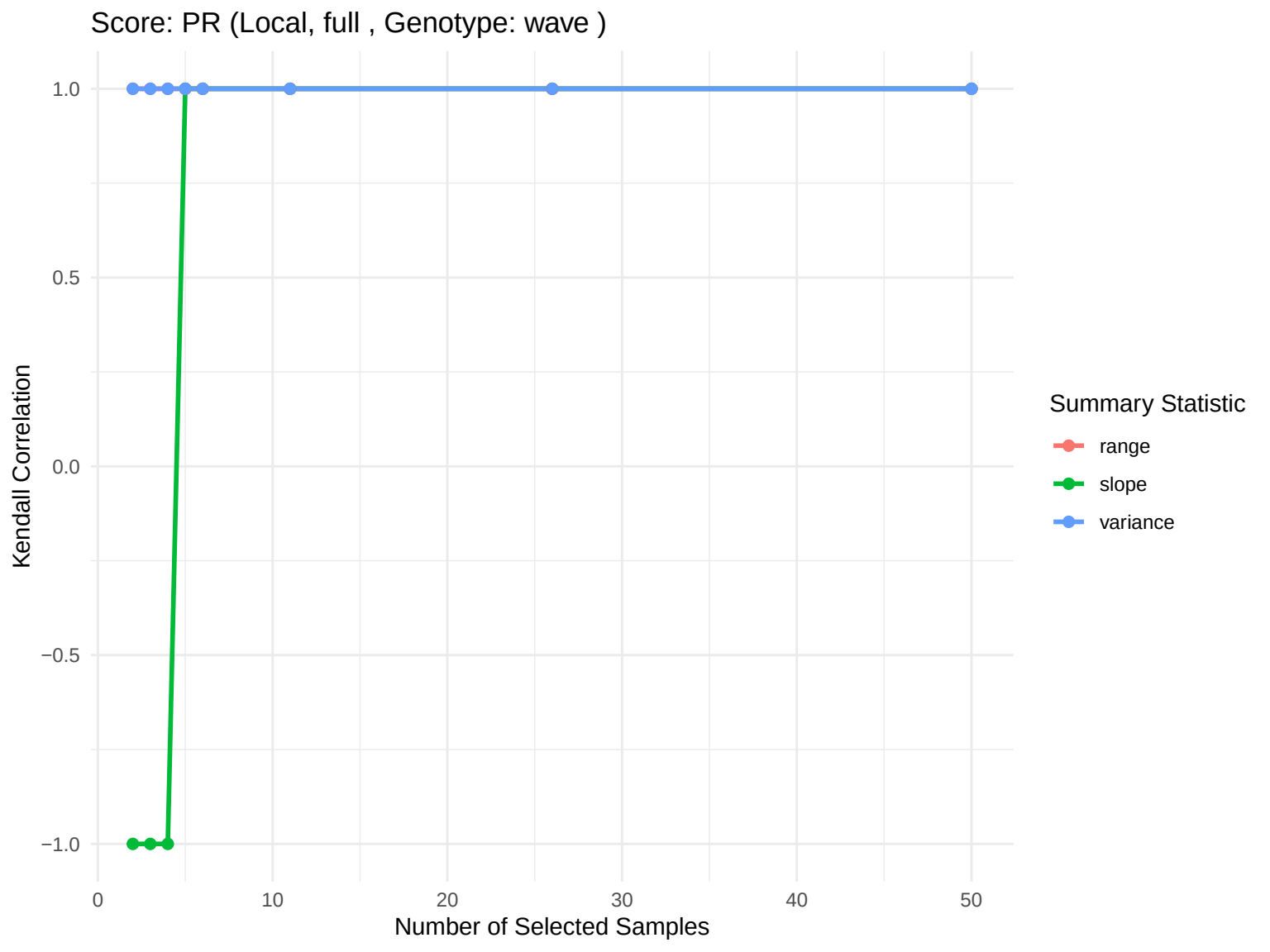
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: PR (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
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−1.0
0
10
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30
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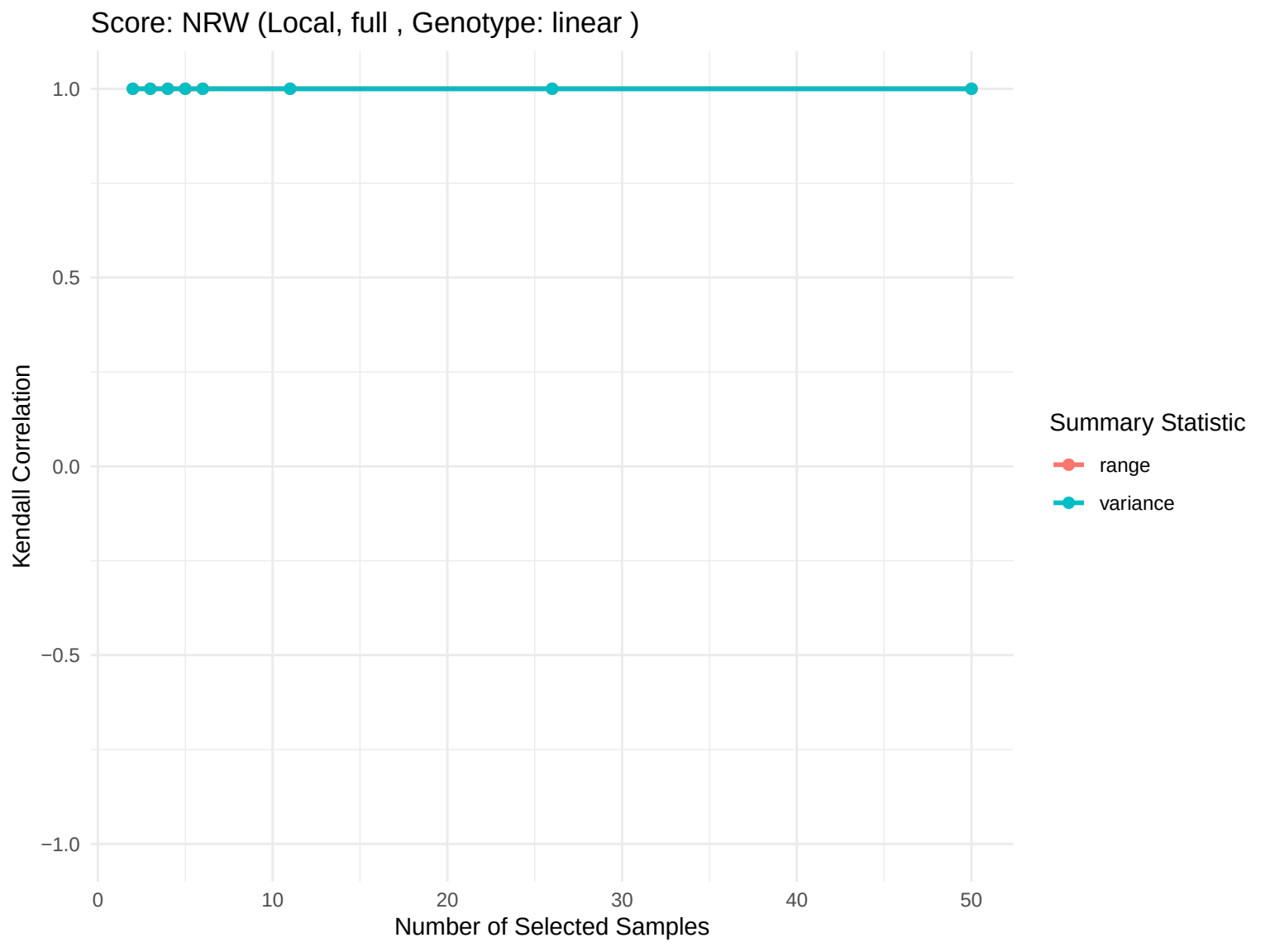
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: NRW (Local, full , Genotype: linear)
Kendall Correlation
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−0.5
−1.0
0
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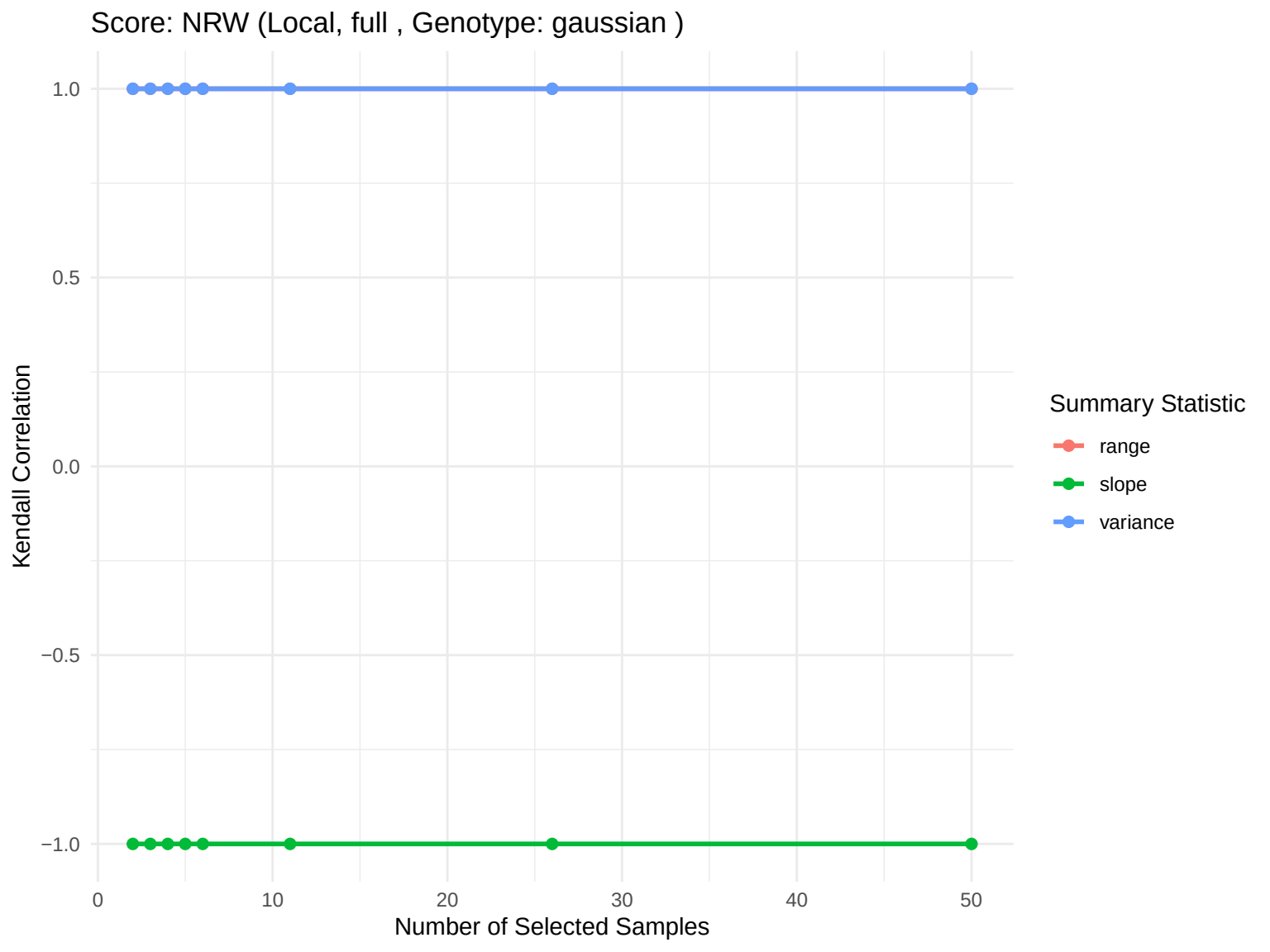
40
50
Number of Selected Samples
Summary Statistic
range
variance


Score: NRW (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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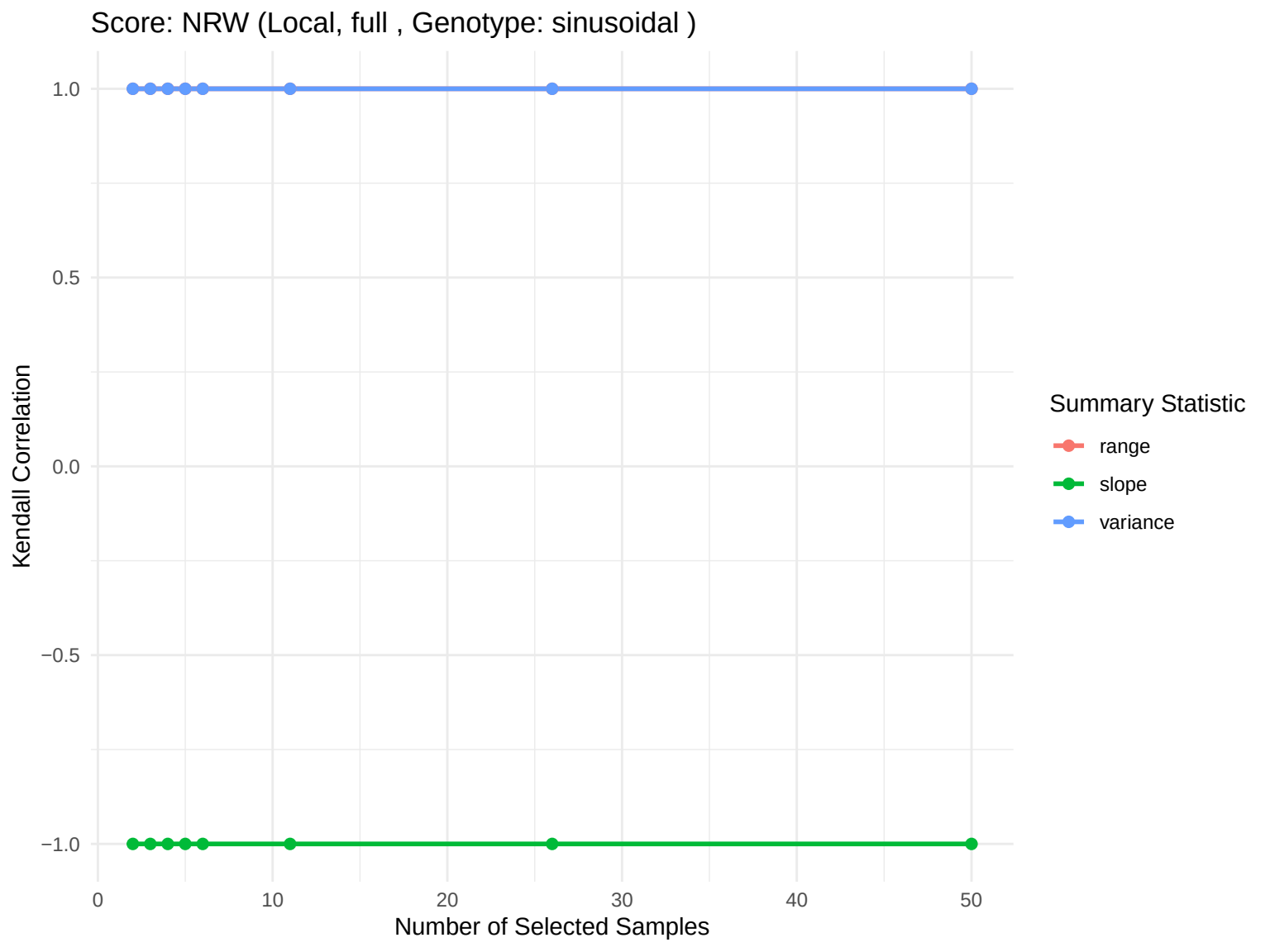
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: NRW (Local, full , Genotype: sinusoidal)
Kendall Correlation
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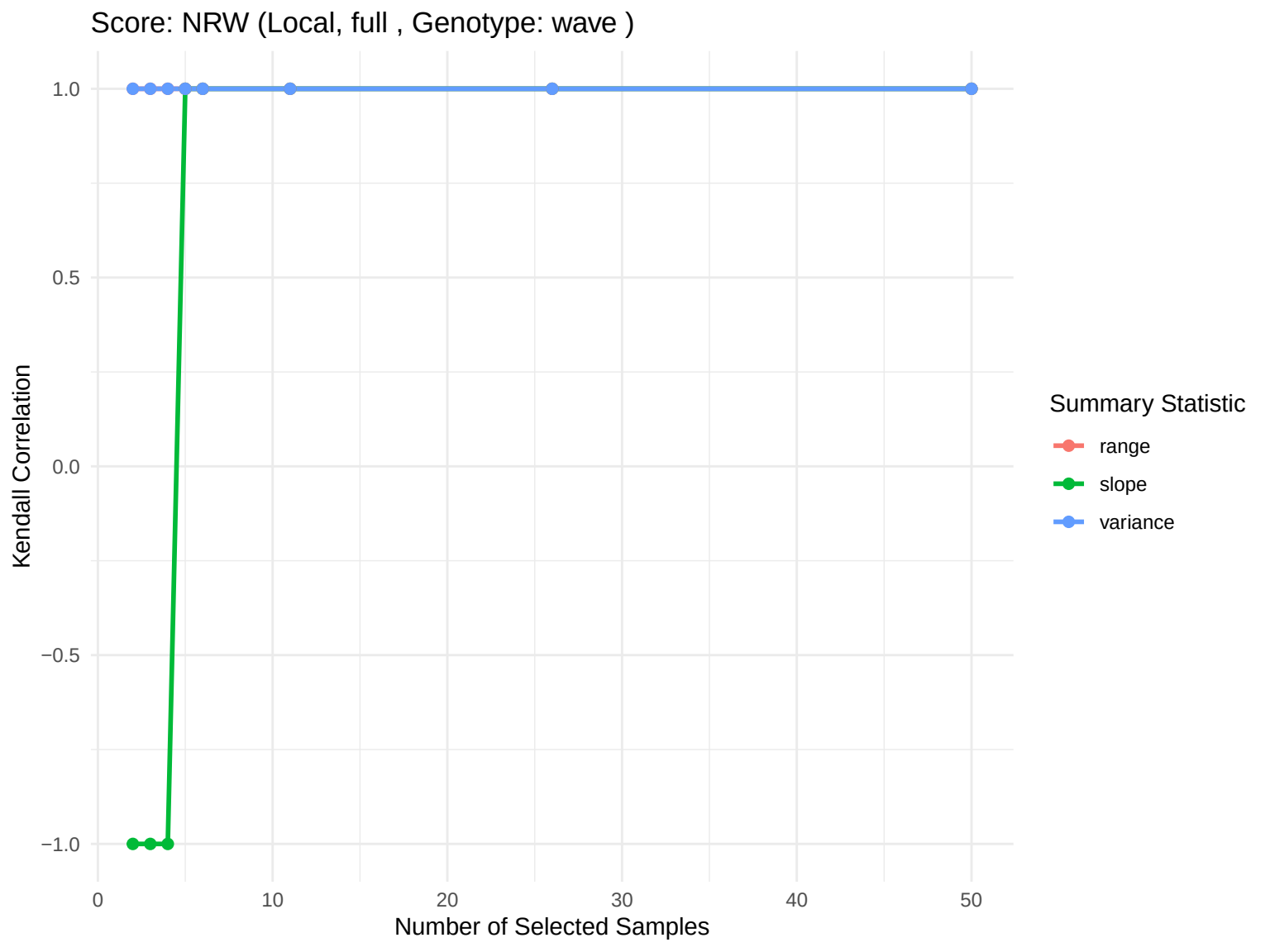
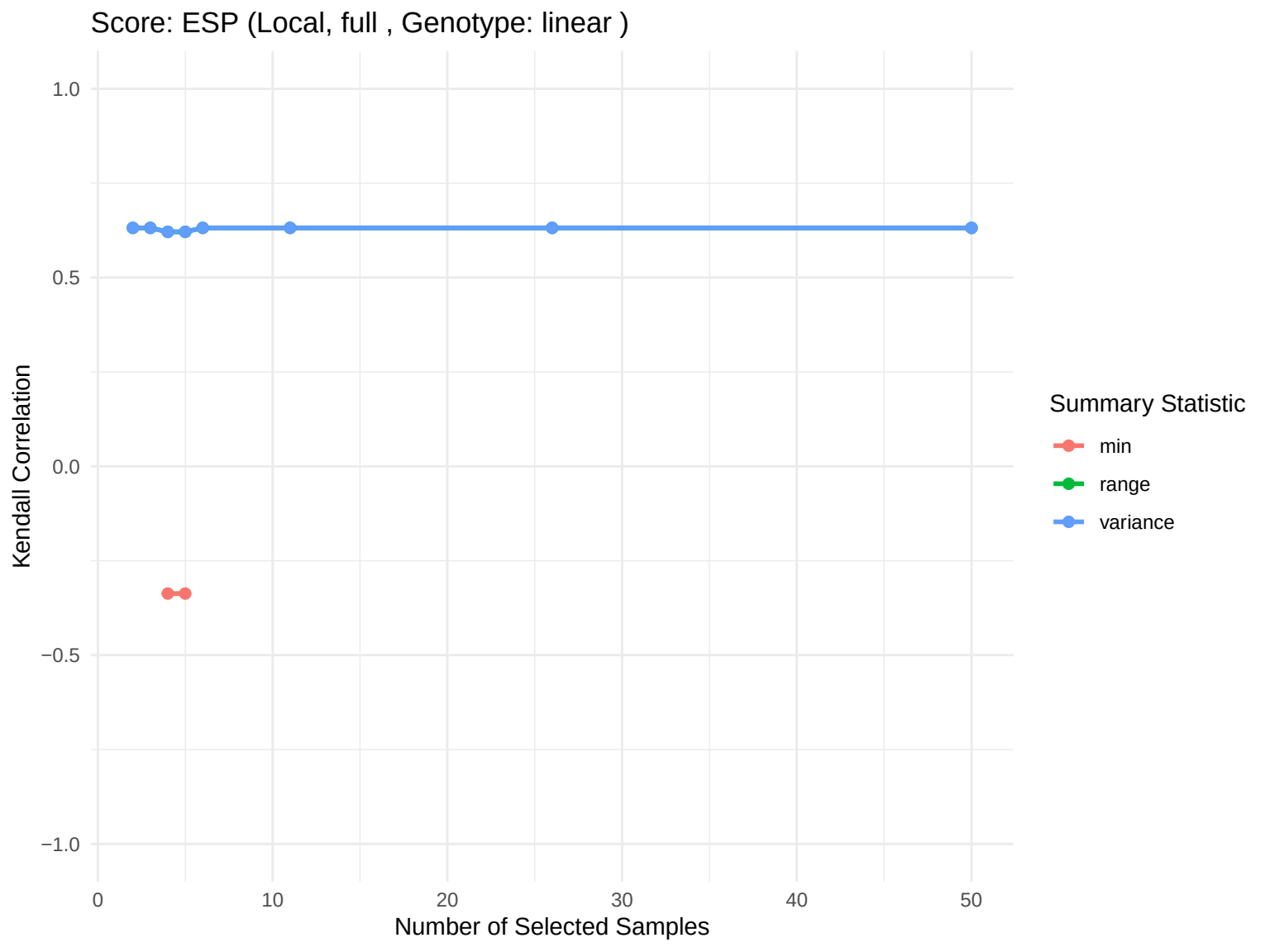
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: NRW (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: ESP (Local, full , Genotype: linear)
Kendall Correlation
1.0
0.5
0.0
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−1.0
0
10
20
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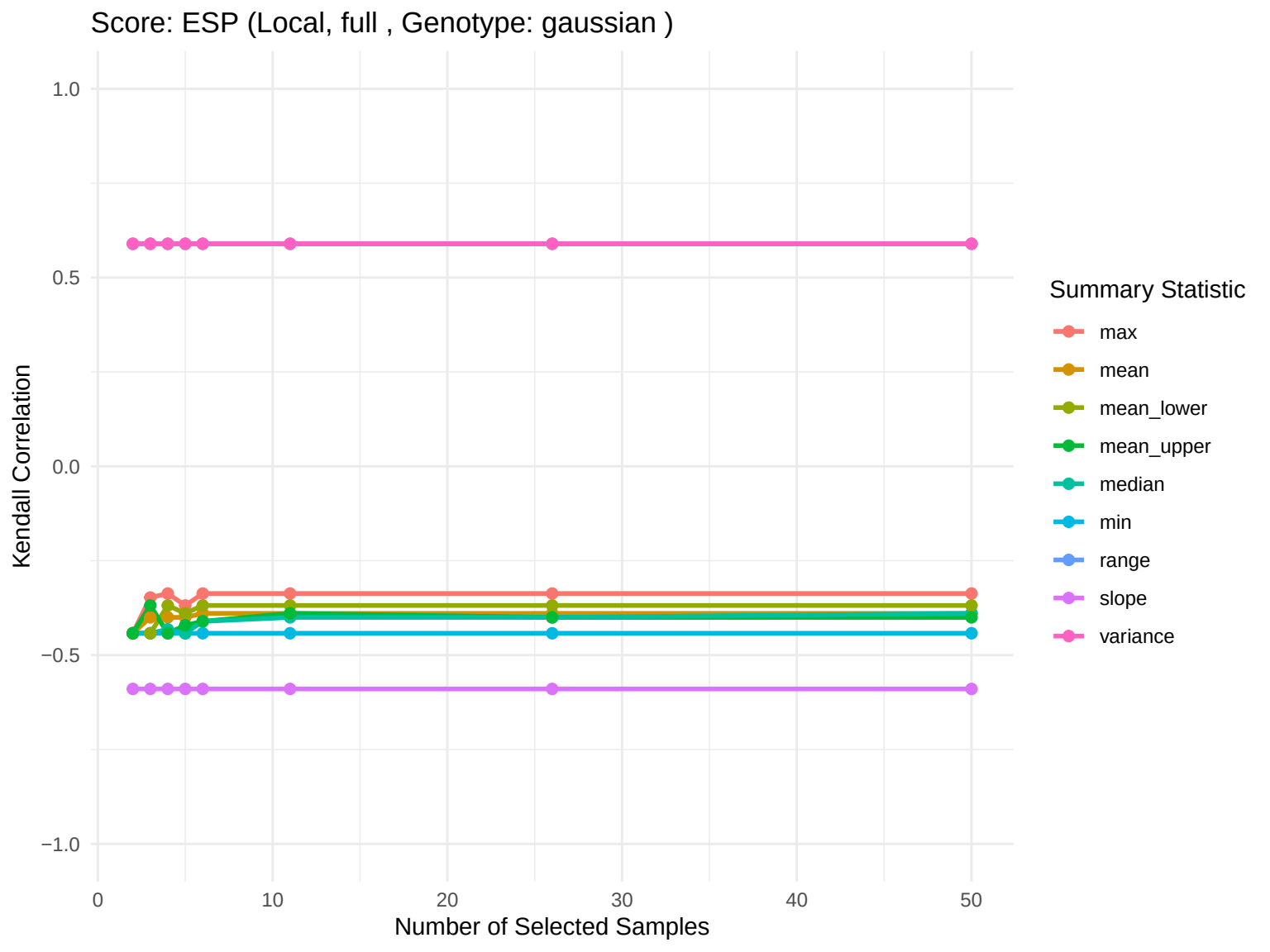
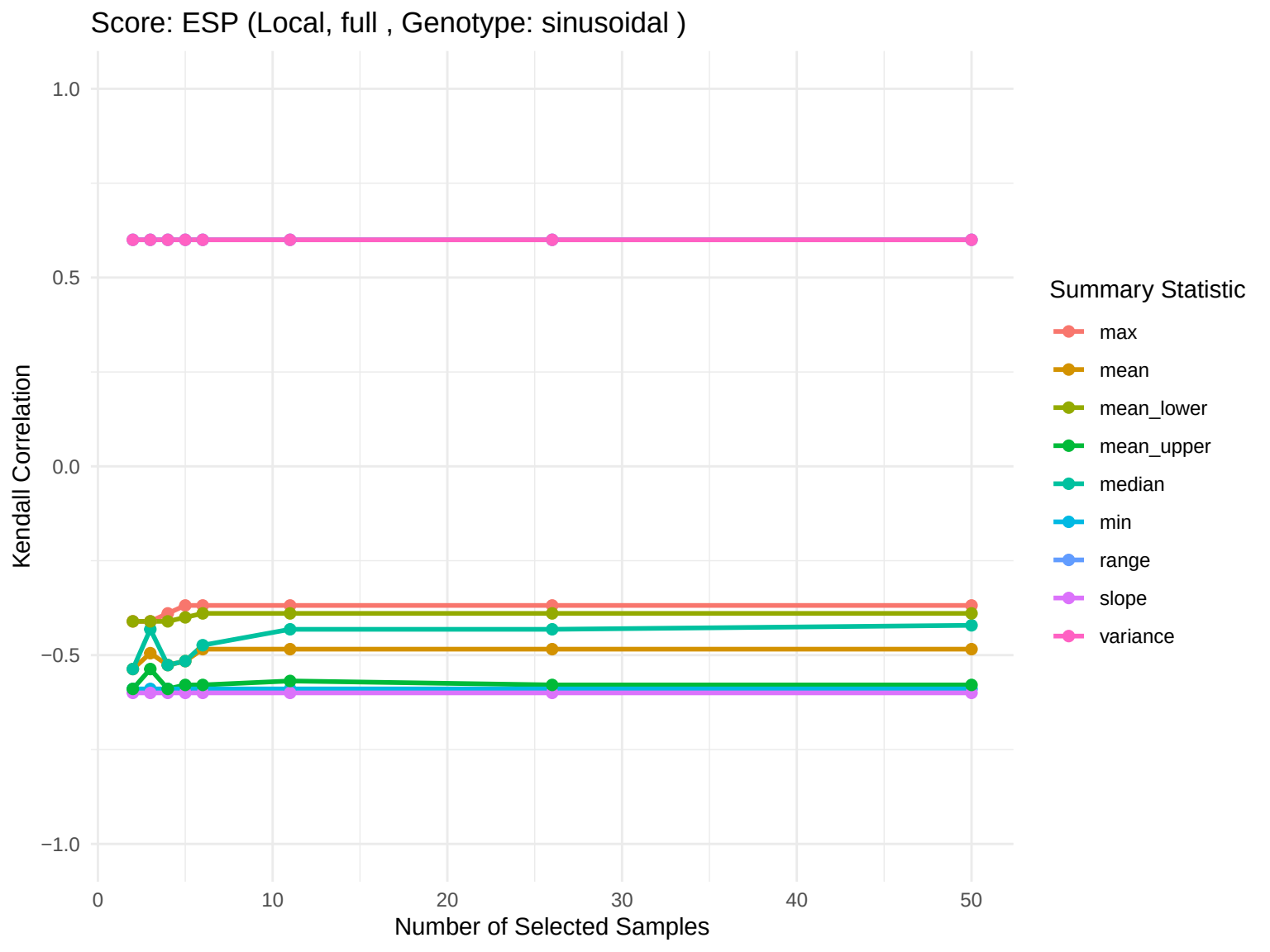
40
50
Number of Selected Samples
Summary Statistic
min
range
variance


Score: ESP (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: ESP (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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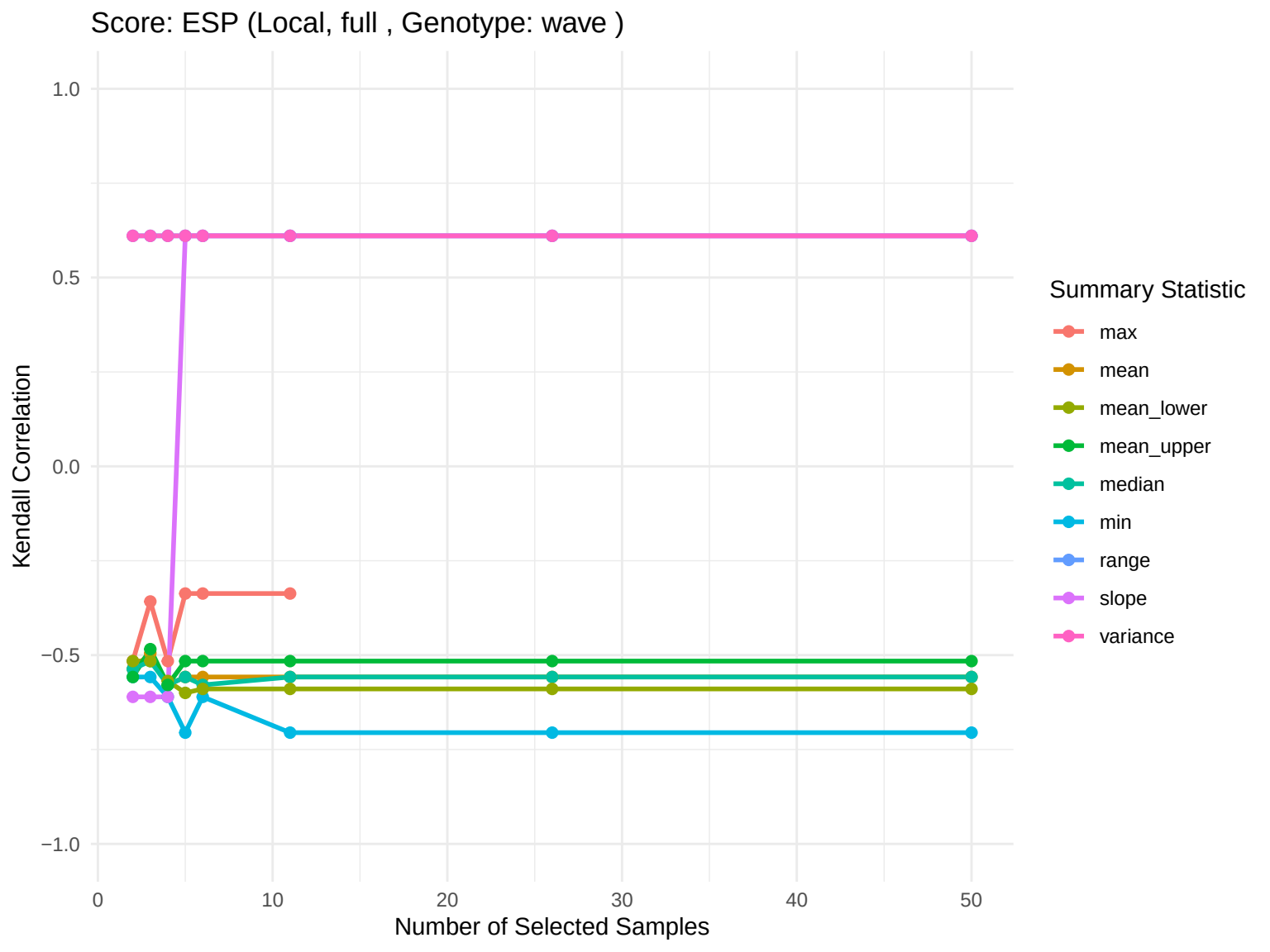
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: ESP (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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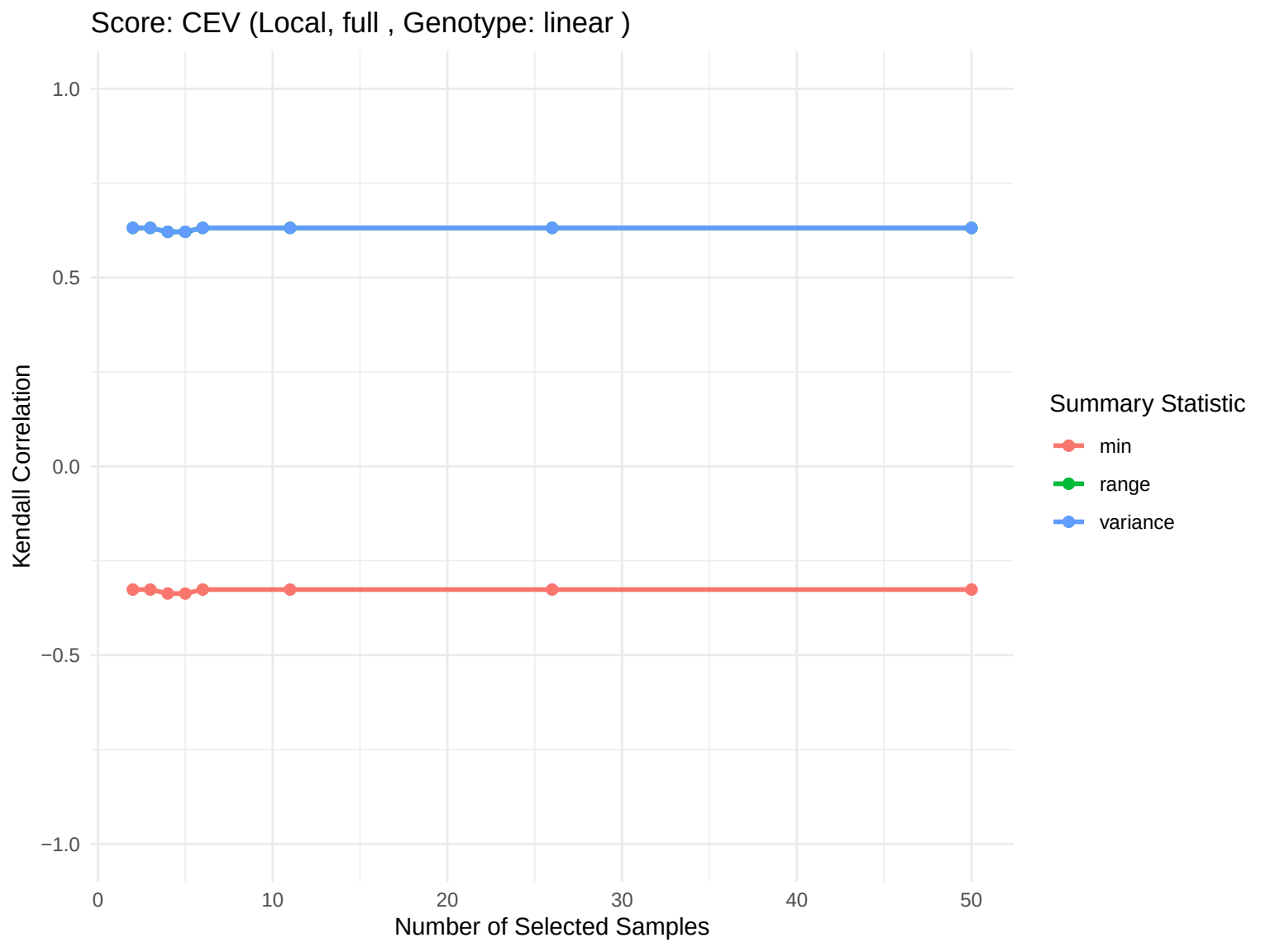
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: CEV (Local, full , Genotype: linear)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
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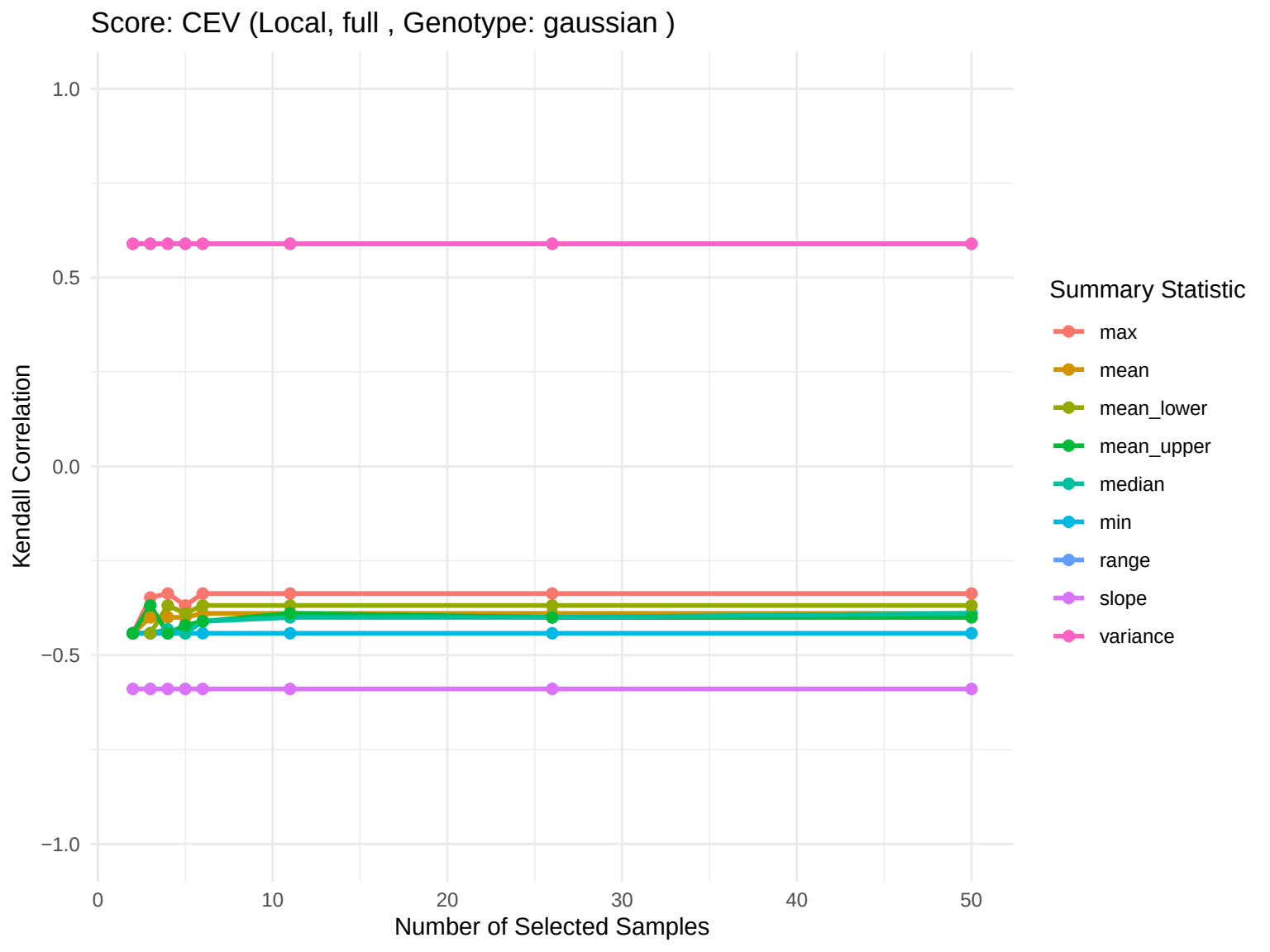
50
Number of Selected Samples
Summary Statistic
min
range
variance


Score: CEV (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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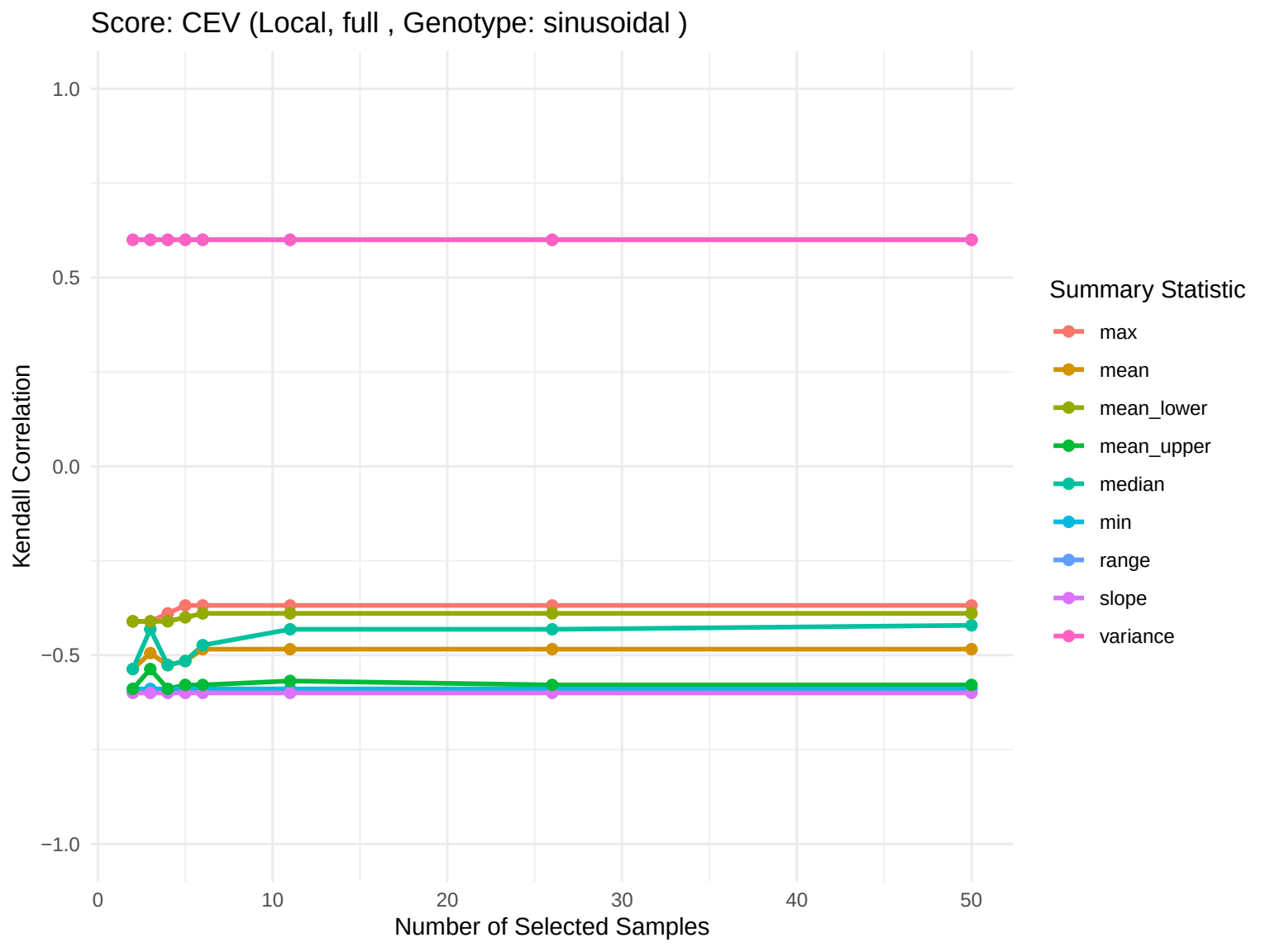
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: CEV (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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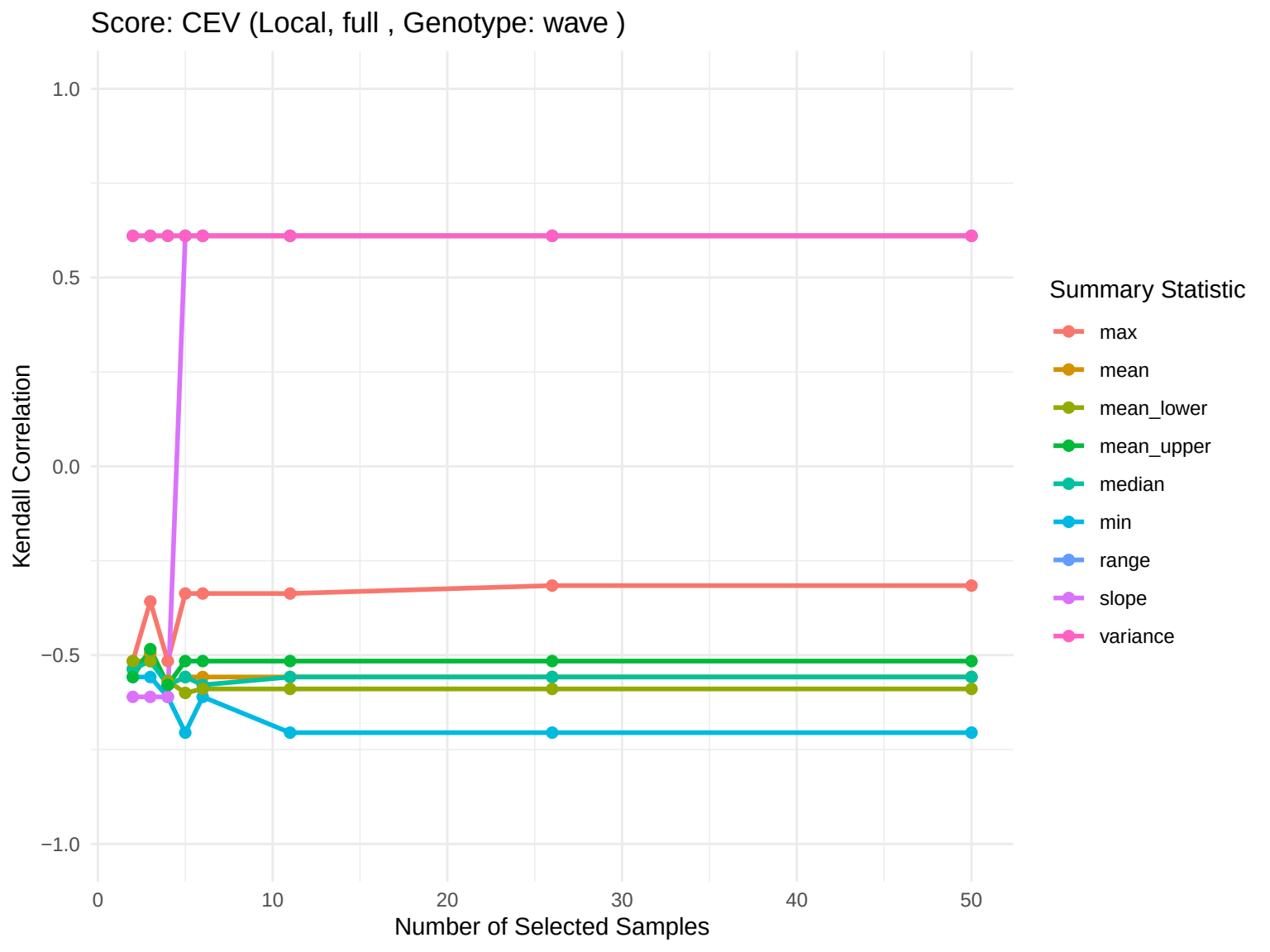
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: CEV (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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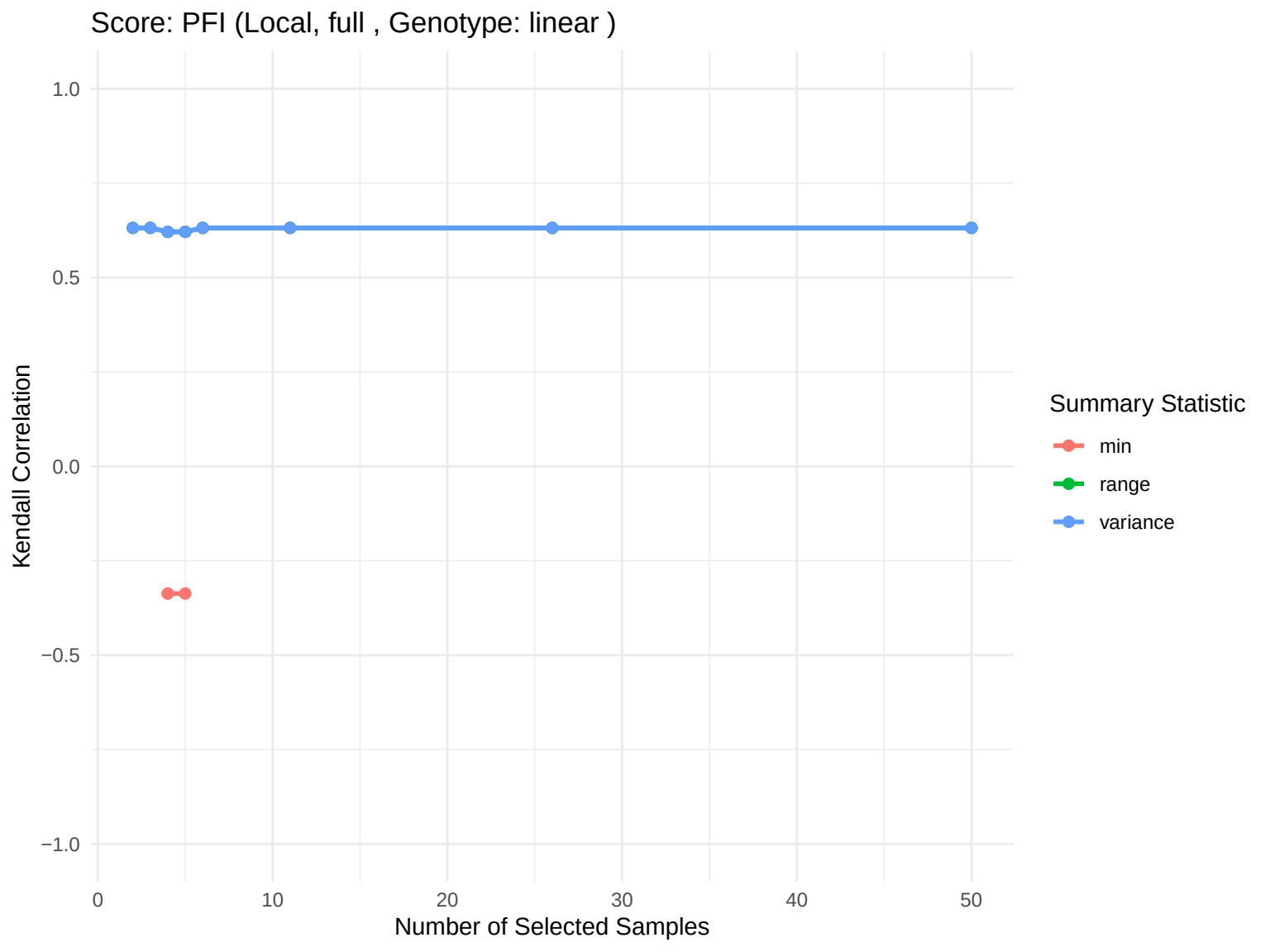
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PFI (Local, full , Genotype: linear)
Kendall Correlation
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0.0
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−1.0
0
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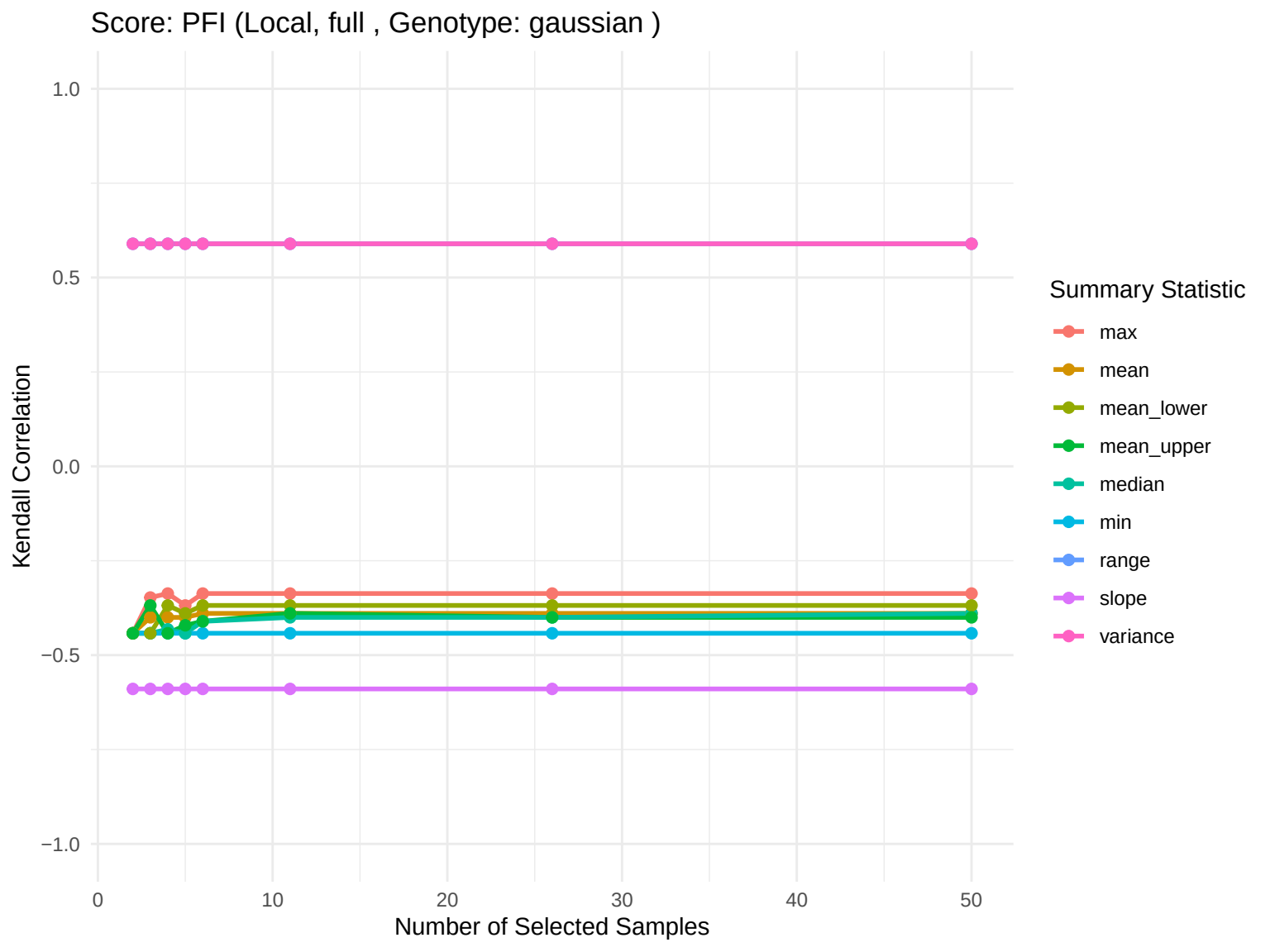
30
40
50
Number of Selected Samples
Summary Statistic
min
range
variance


Score: PFI (Local, full , Genotype: gaussian)
Kendall Correlation
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0.0
−0.5
−1.0
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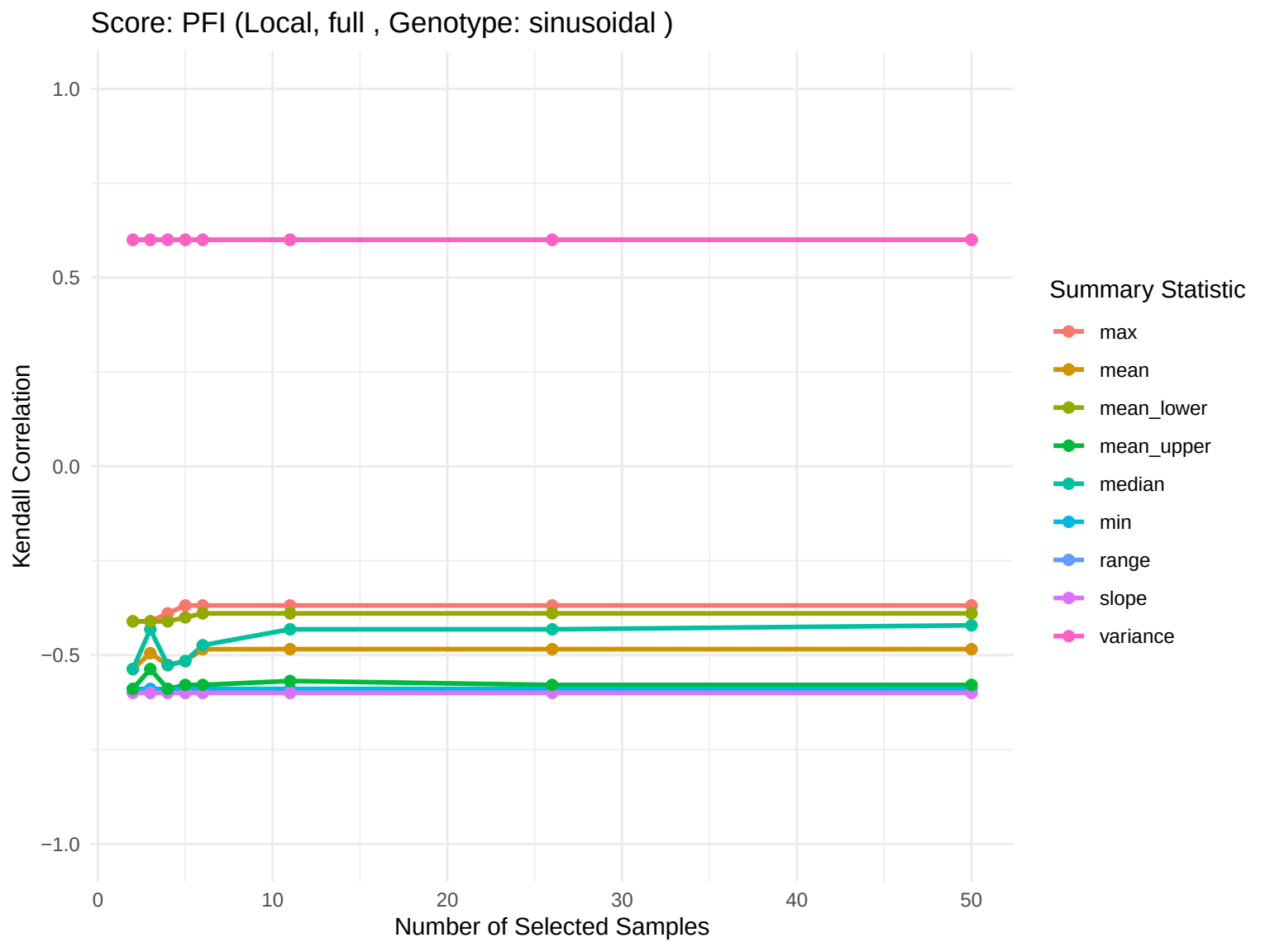
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PFI (Local, full , Genotype: sinusoidal)
Kendall Correlation
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0.0
−0.5
−1.0
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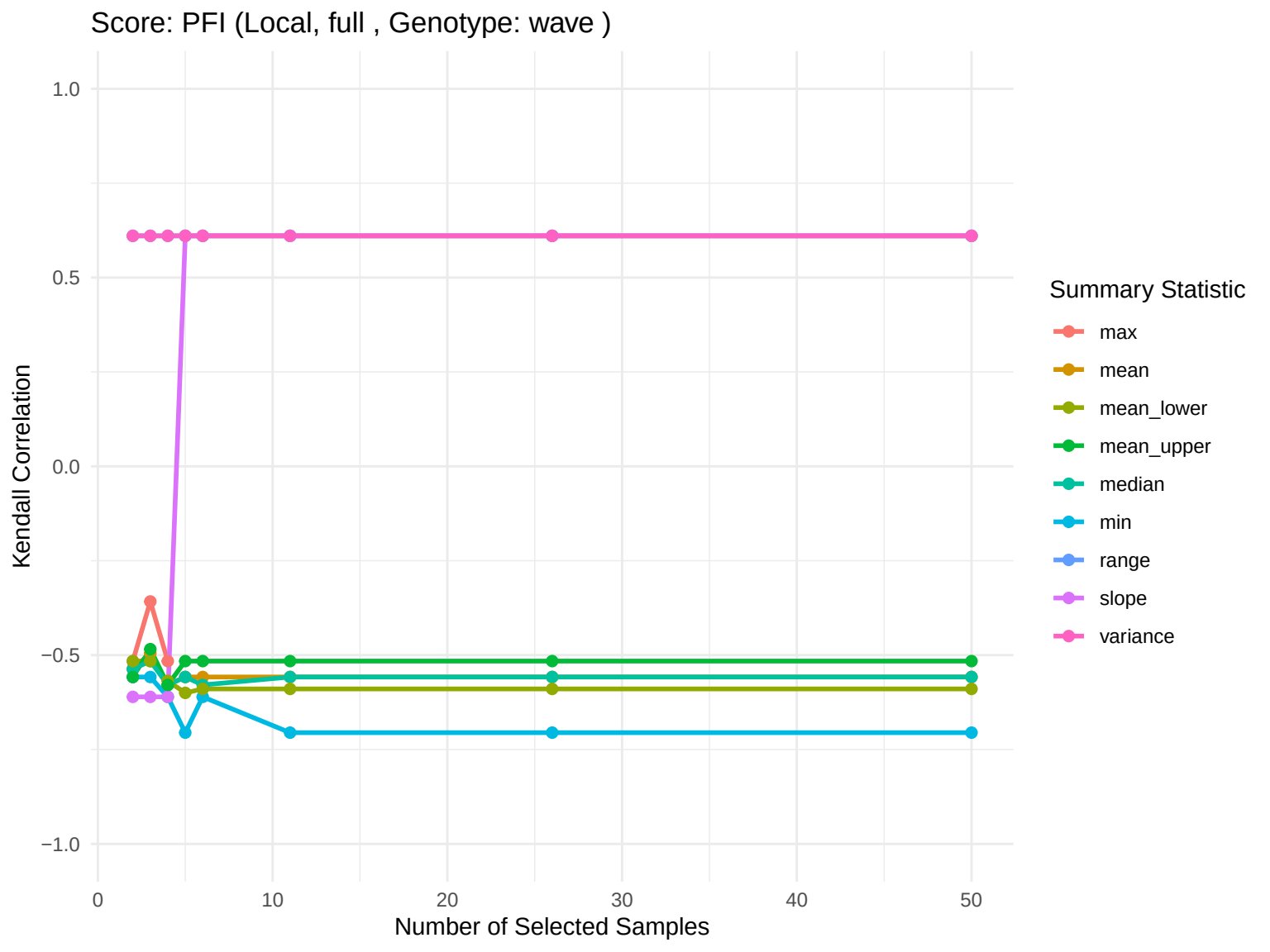
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: PFI (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
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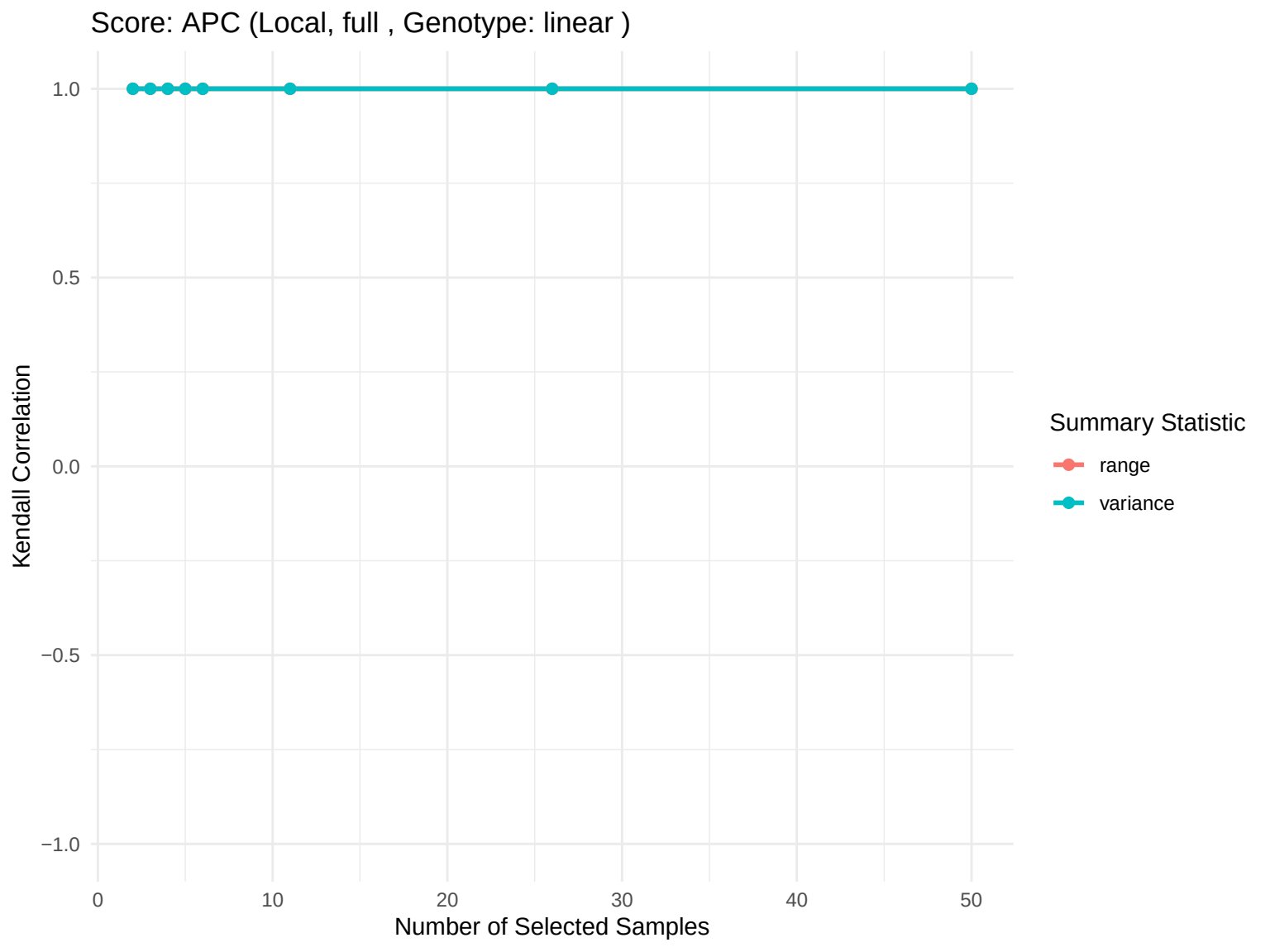
range
slope
variance


Score: APC (Local, full , Genotype: linear)
Kendall Correlation
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−0.5
−1.0
0
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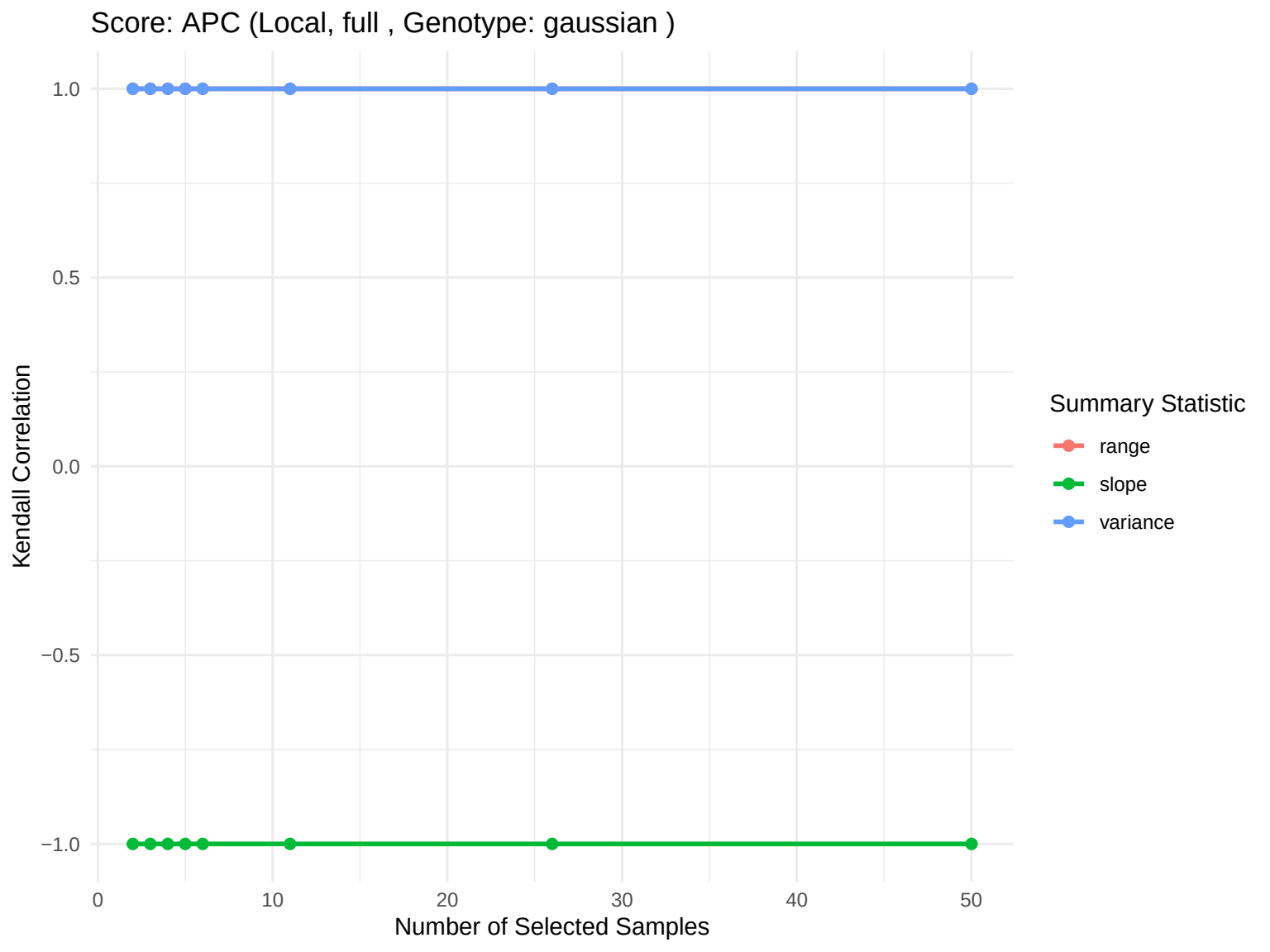
40
50
Number of Selected Samples
Summary Statistic
range
variance


Score: APC (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
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0.0
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−1.0
0
10
20
30
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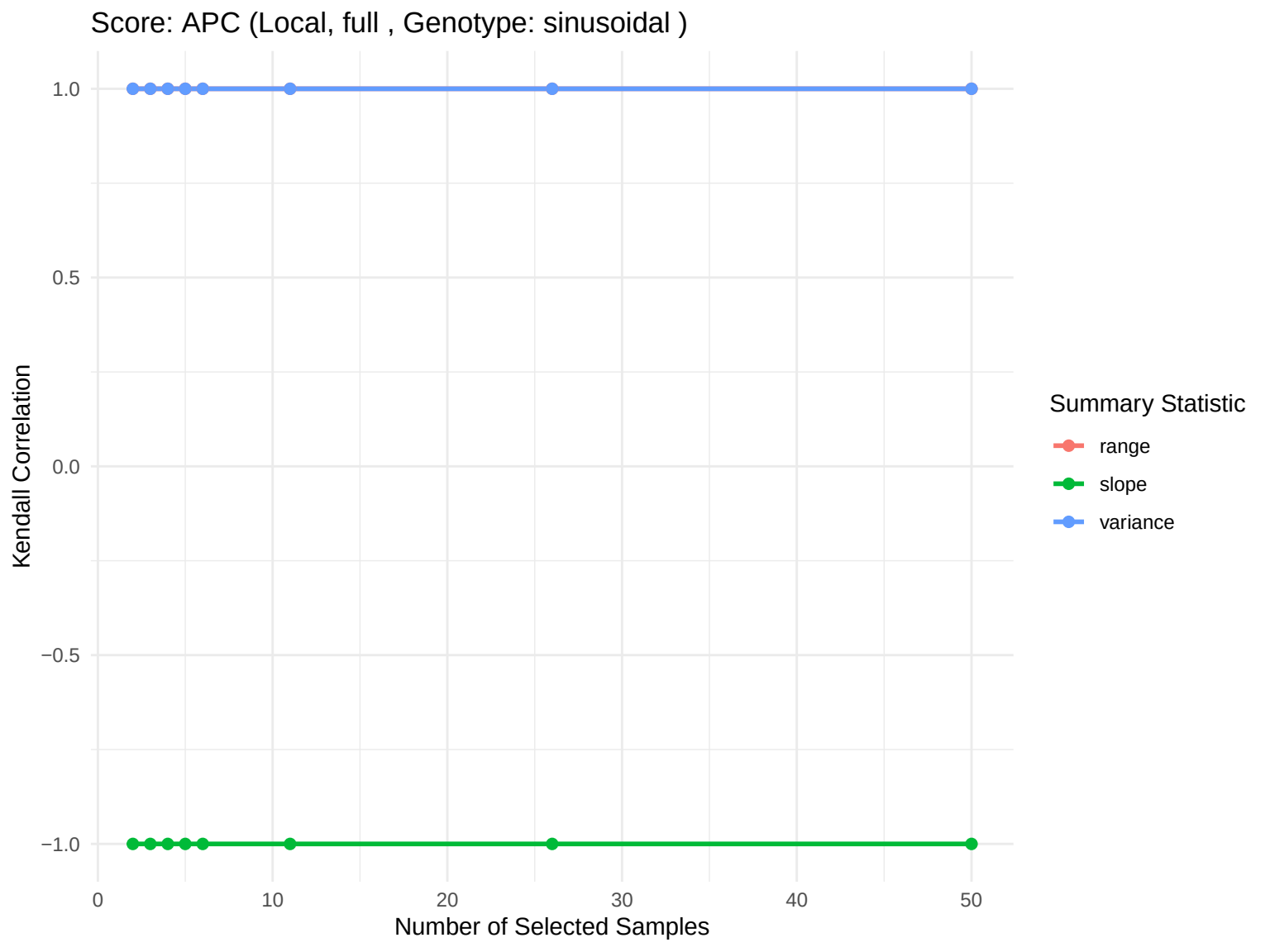
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: APC (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
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−1.0
0
10
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30
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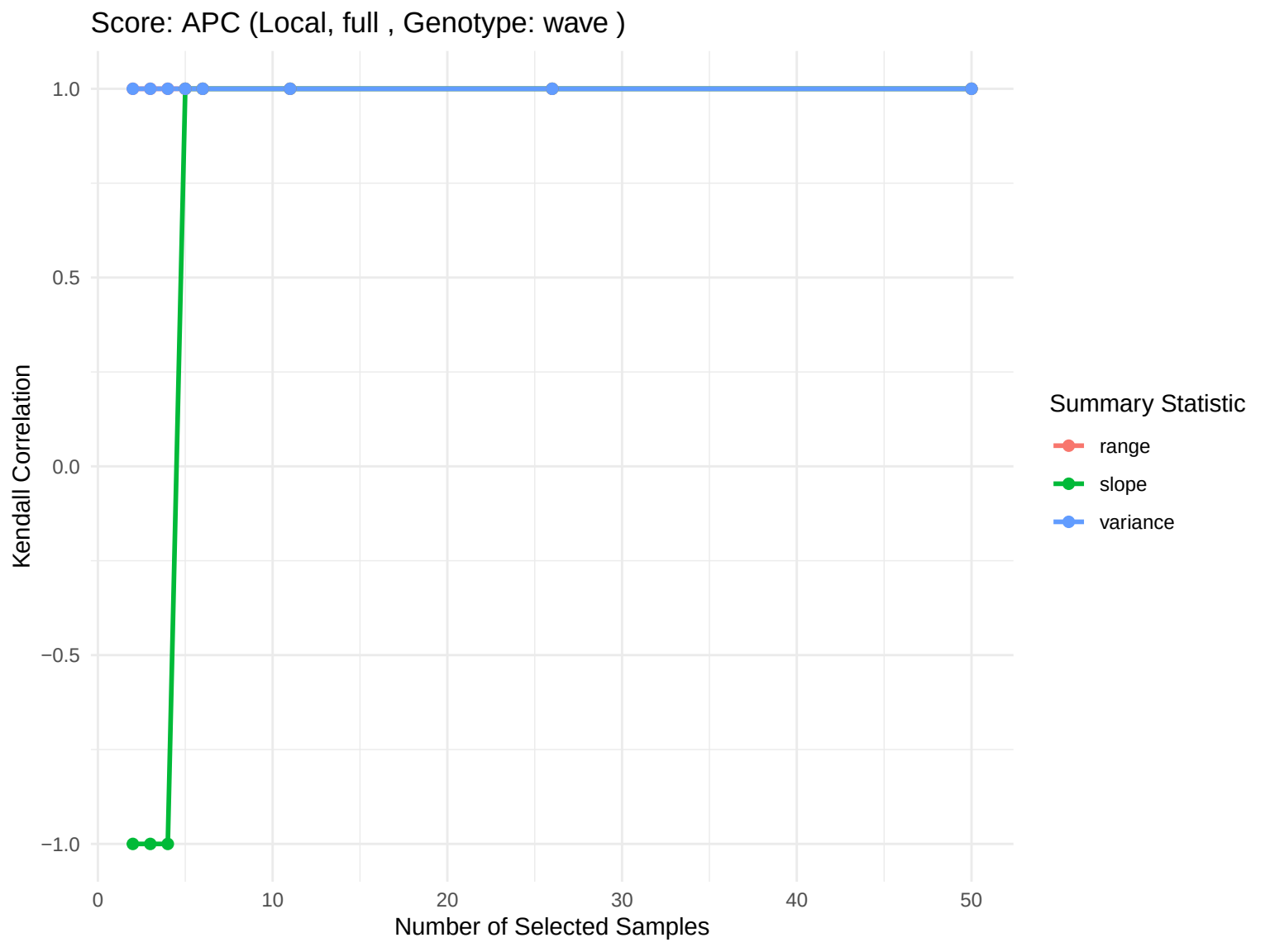
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: APC (Local, full , Genotype: wave)
Kendall Correlation
1.0
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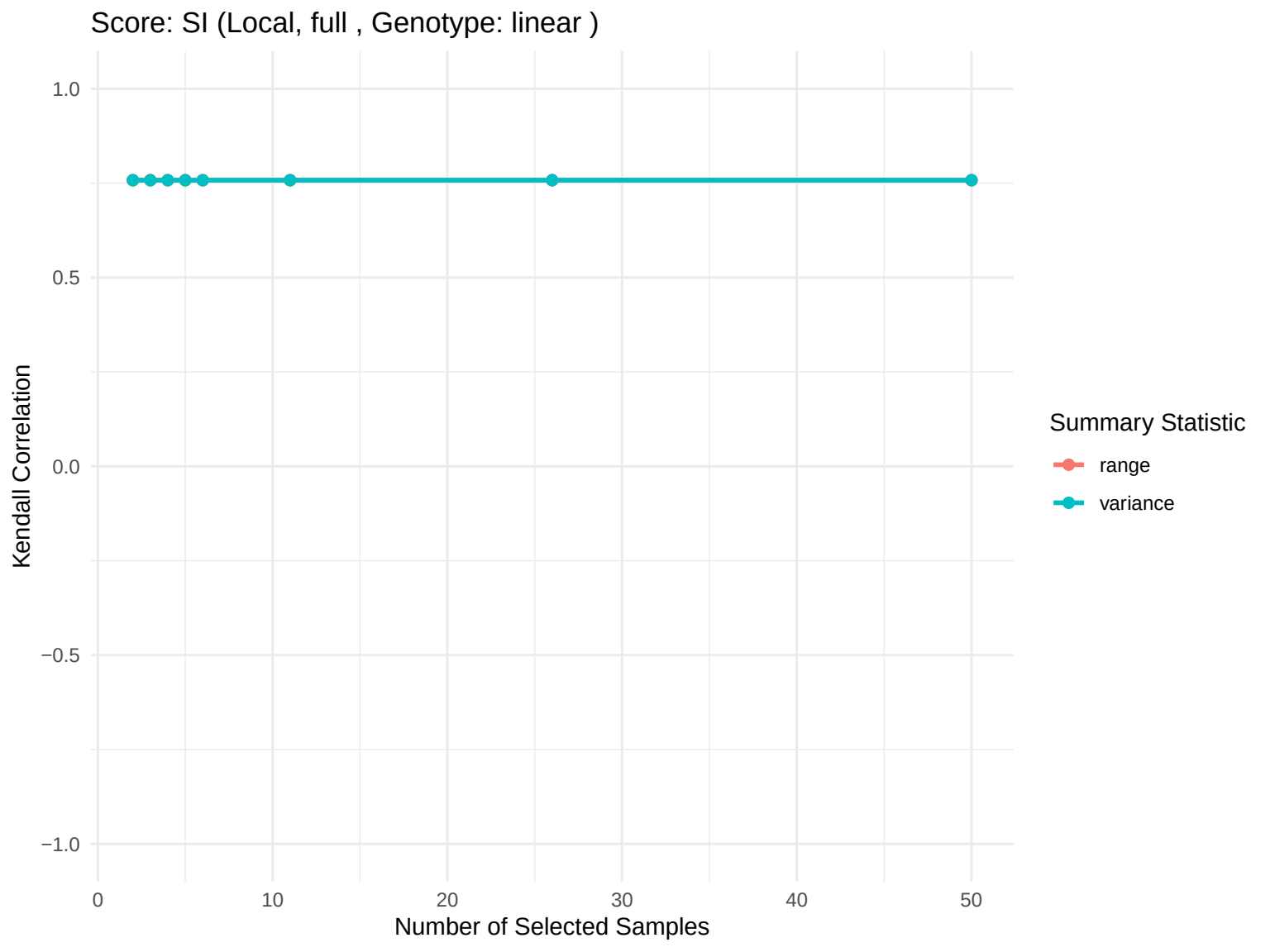
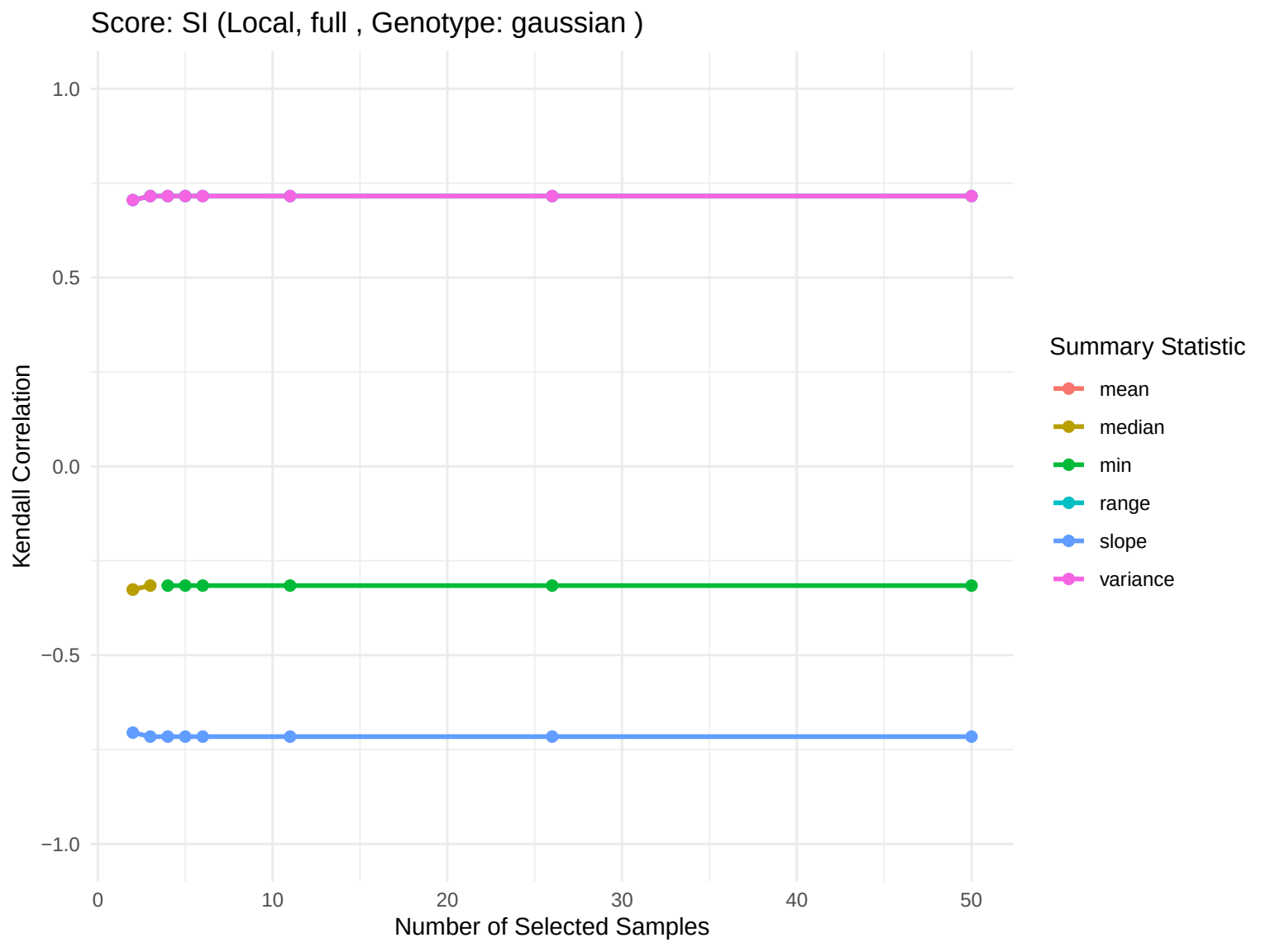
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: SI (Local, full , Genotype: linear)
1.0
0.5
0.0
−0.5
−1.0
Kendall Correlation
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
variance


Score: SI (Local, full , Genotype: gaussian)
Kendall Correlation
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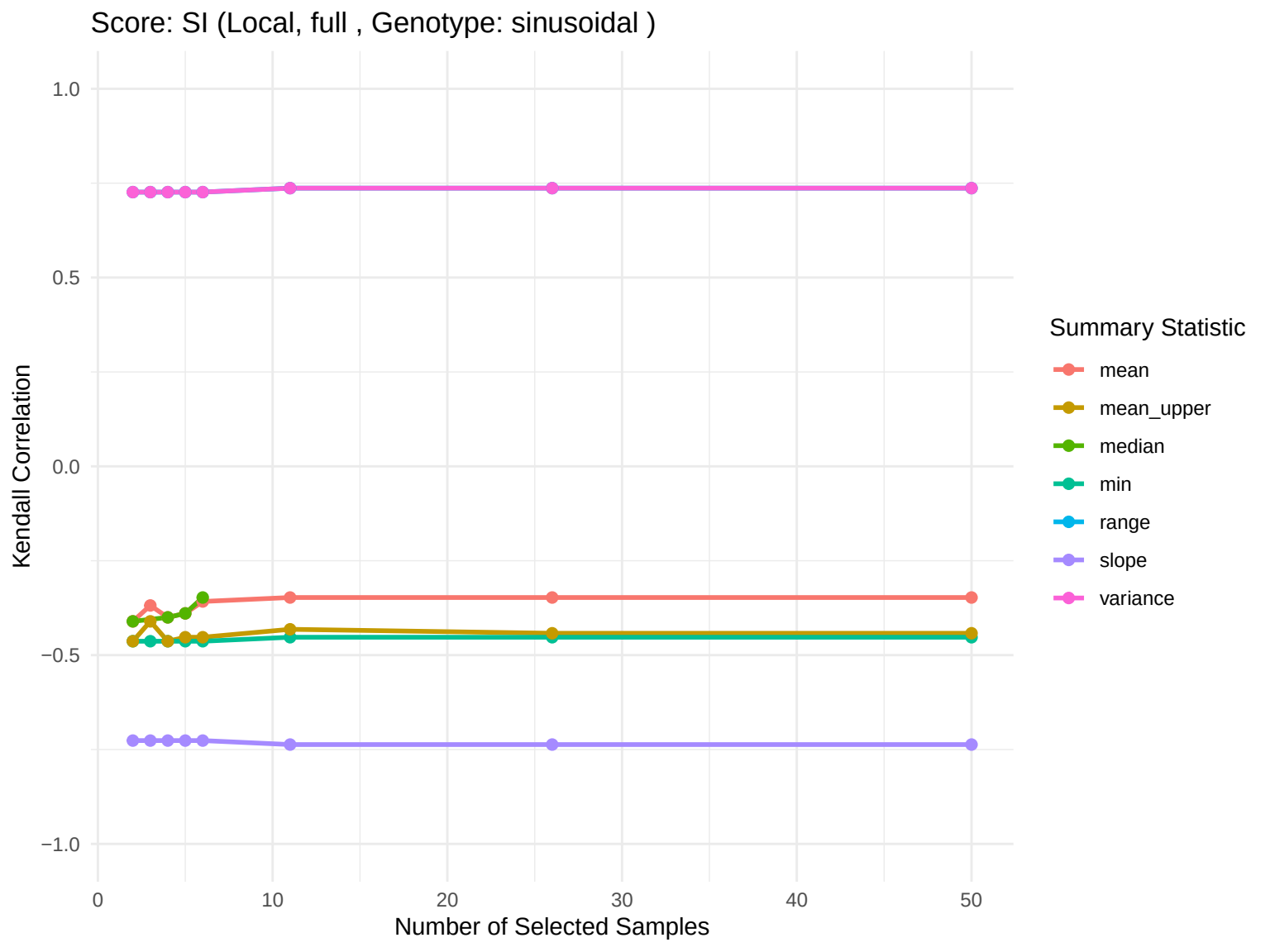
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
mean
median
min
range
slope
variance


Score: SI (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
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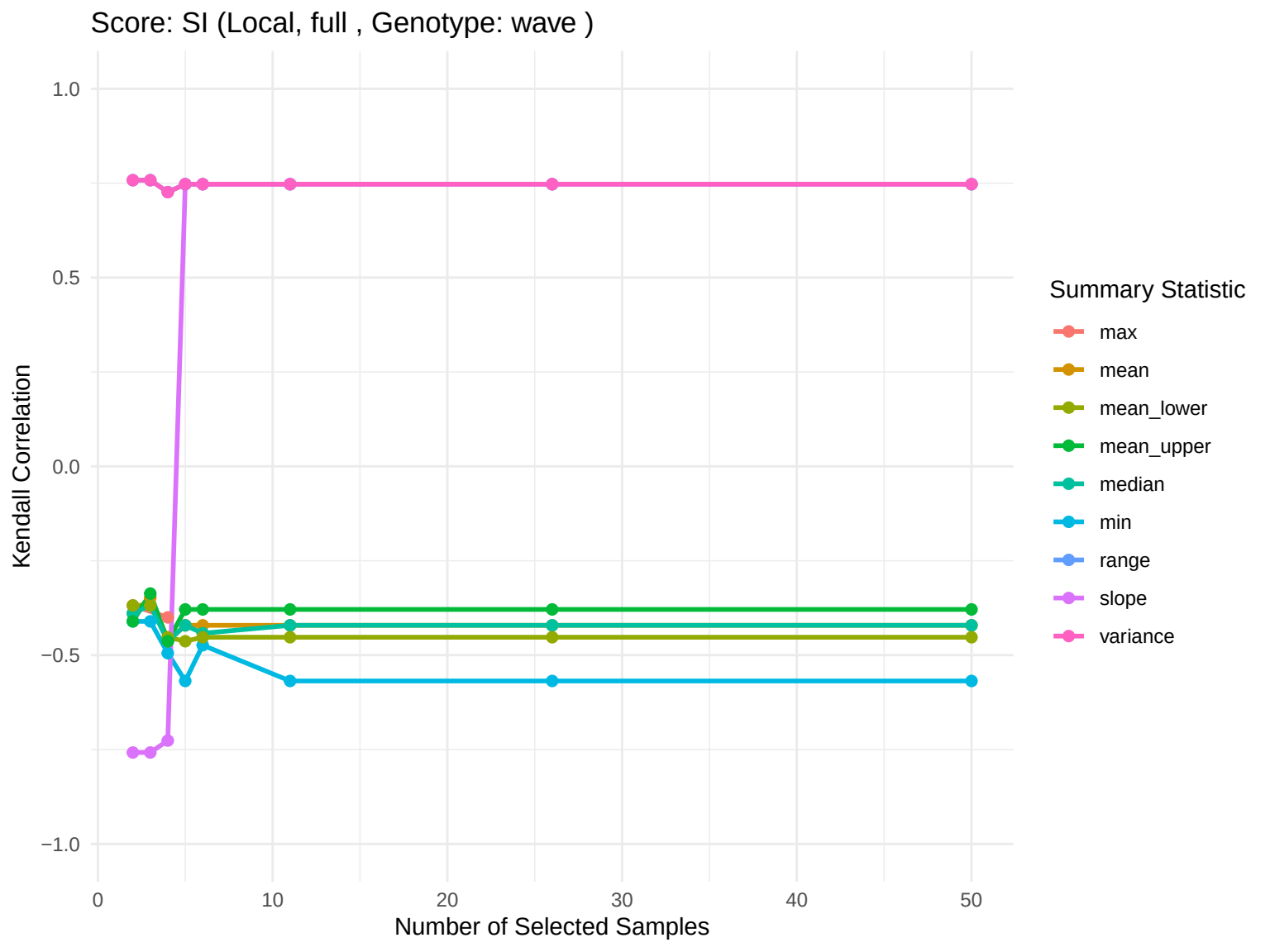
20
30
40
50
Number of Selected Samples
Summary Statistic
mean
mean_upper
median
min
range
slope
variance


Score: SI (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
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20
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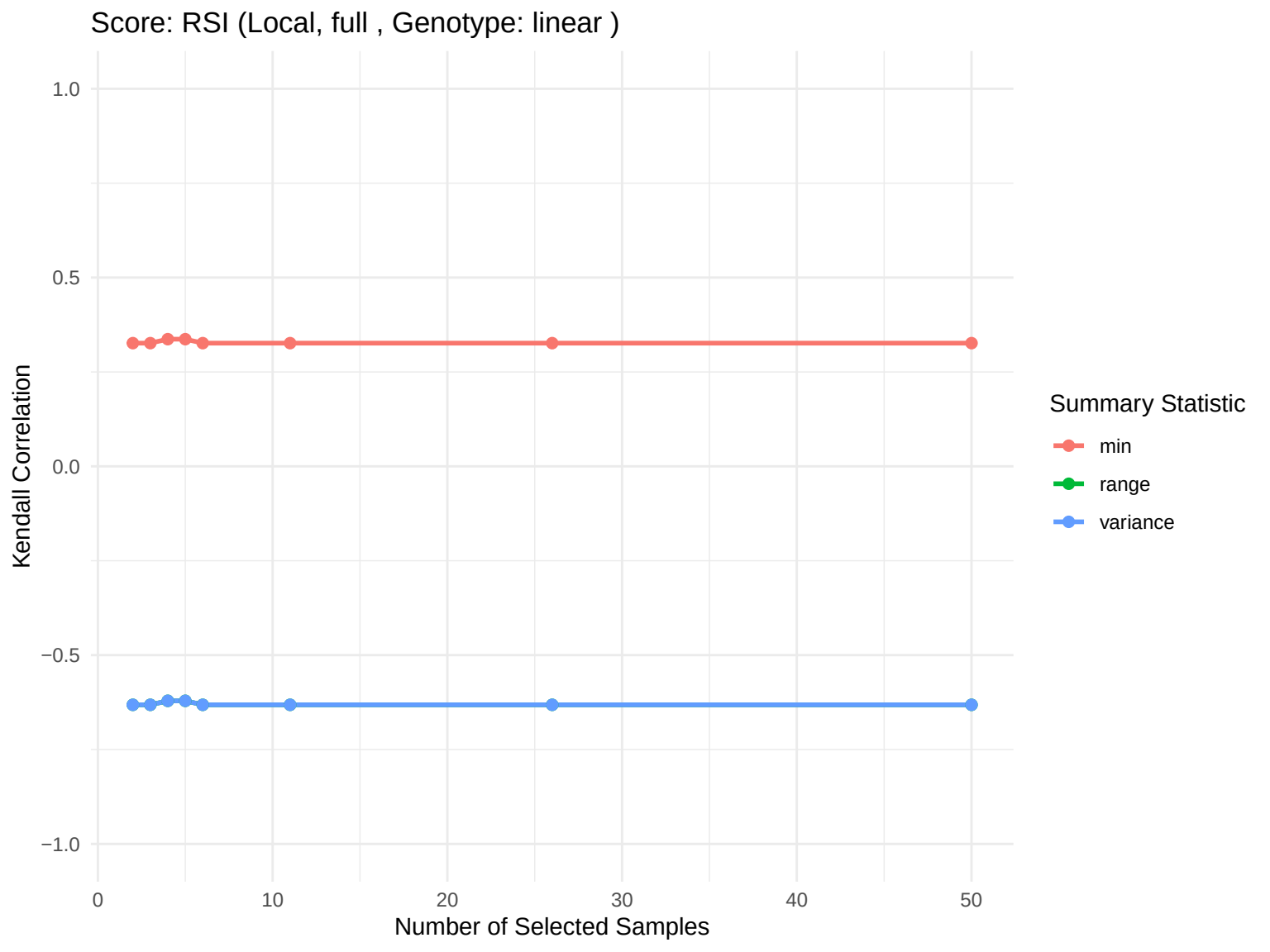
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RSI (Local, full , Genotype: linear)
Kendall Correlation
1.0
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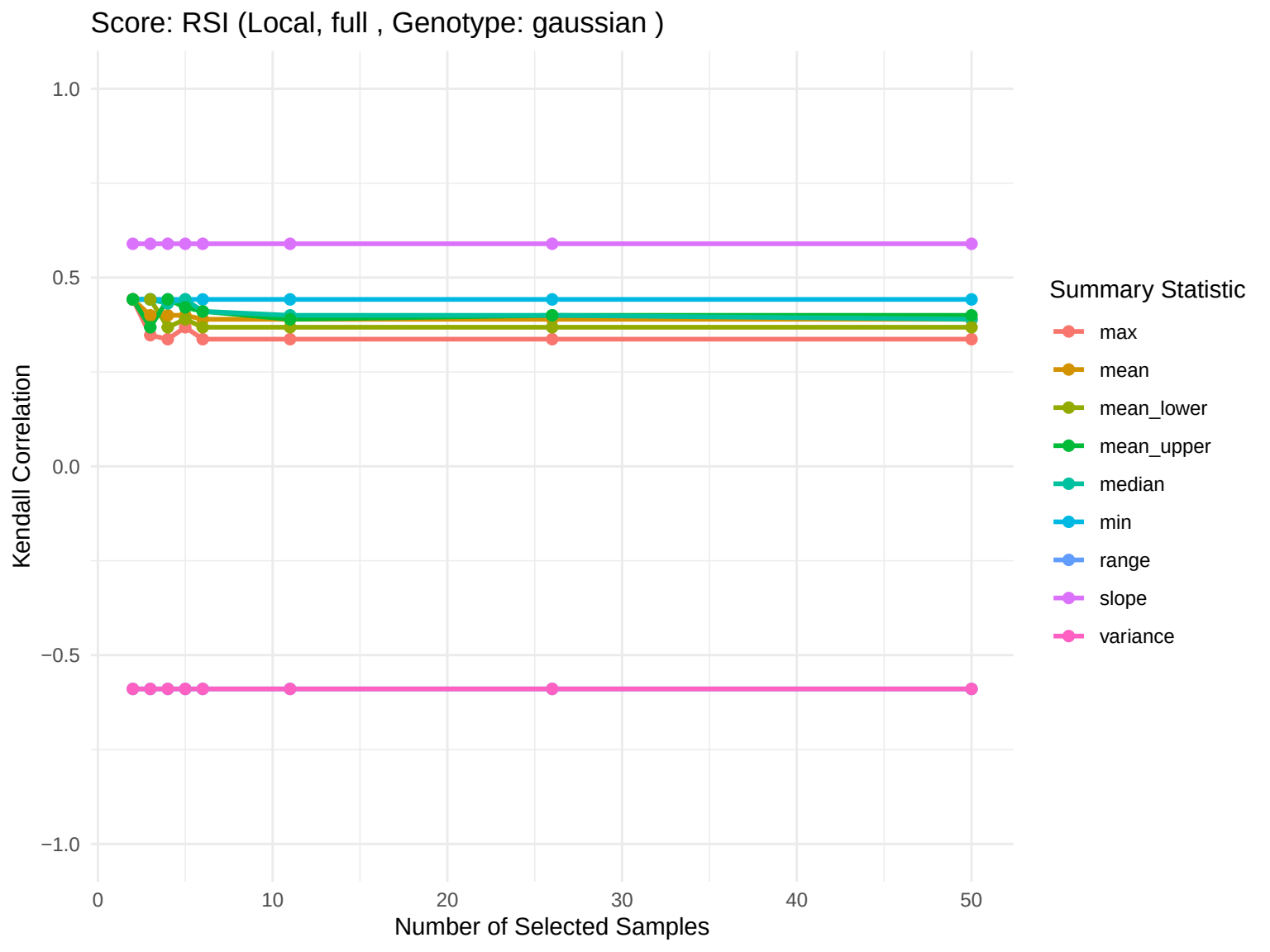
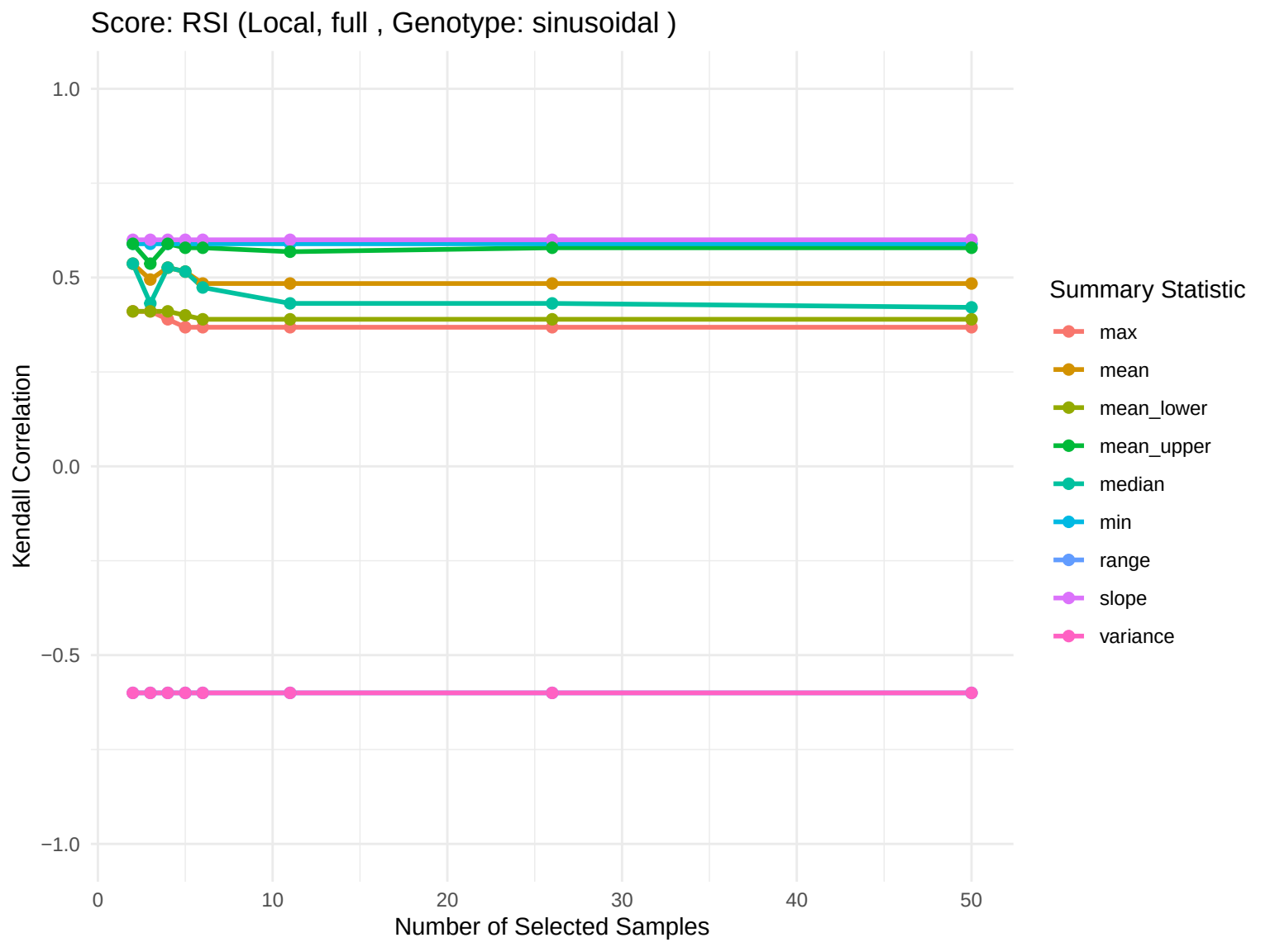
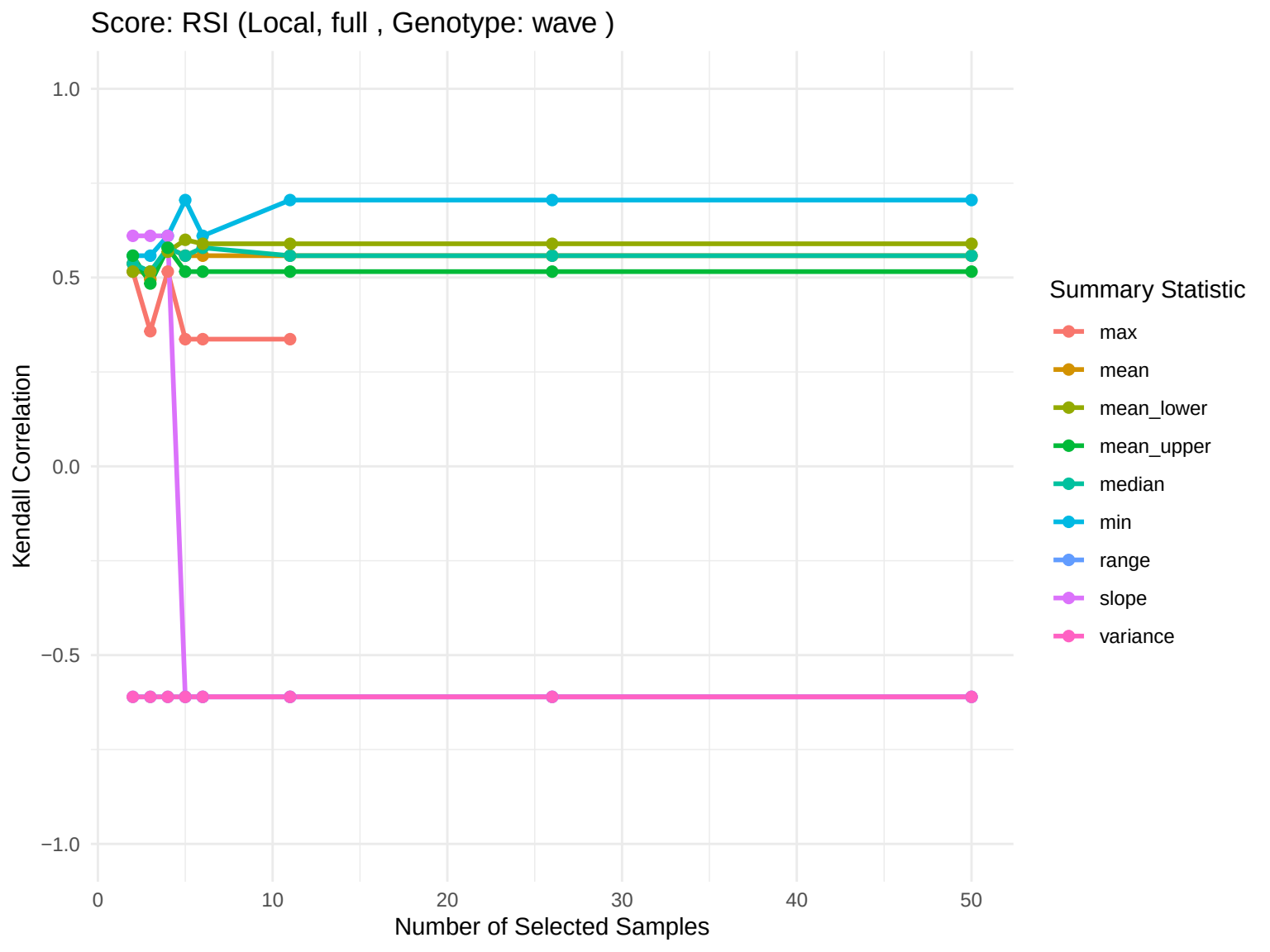
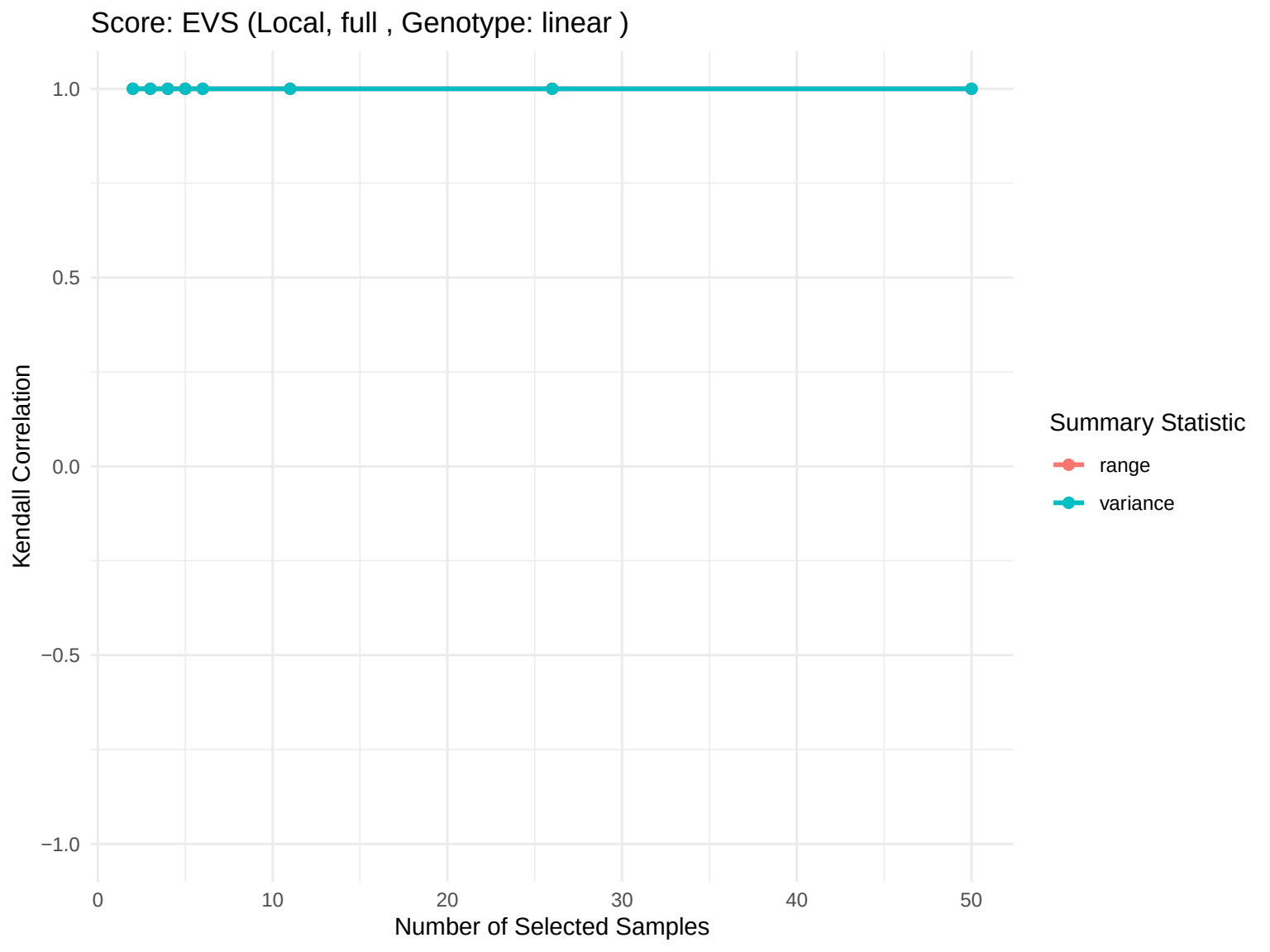
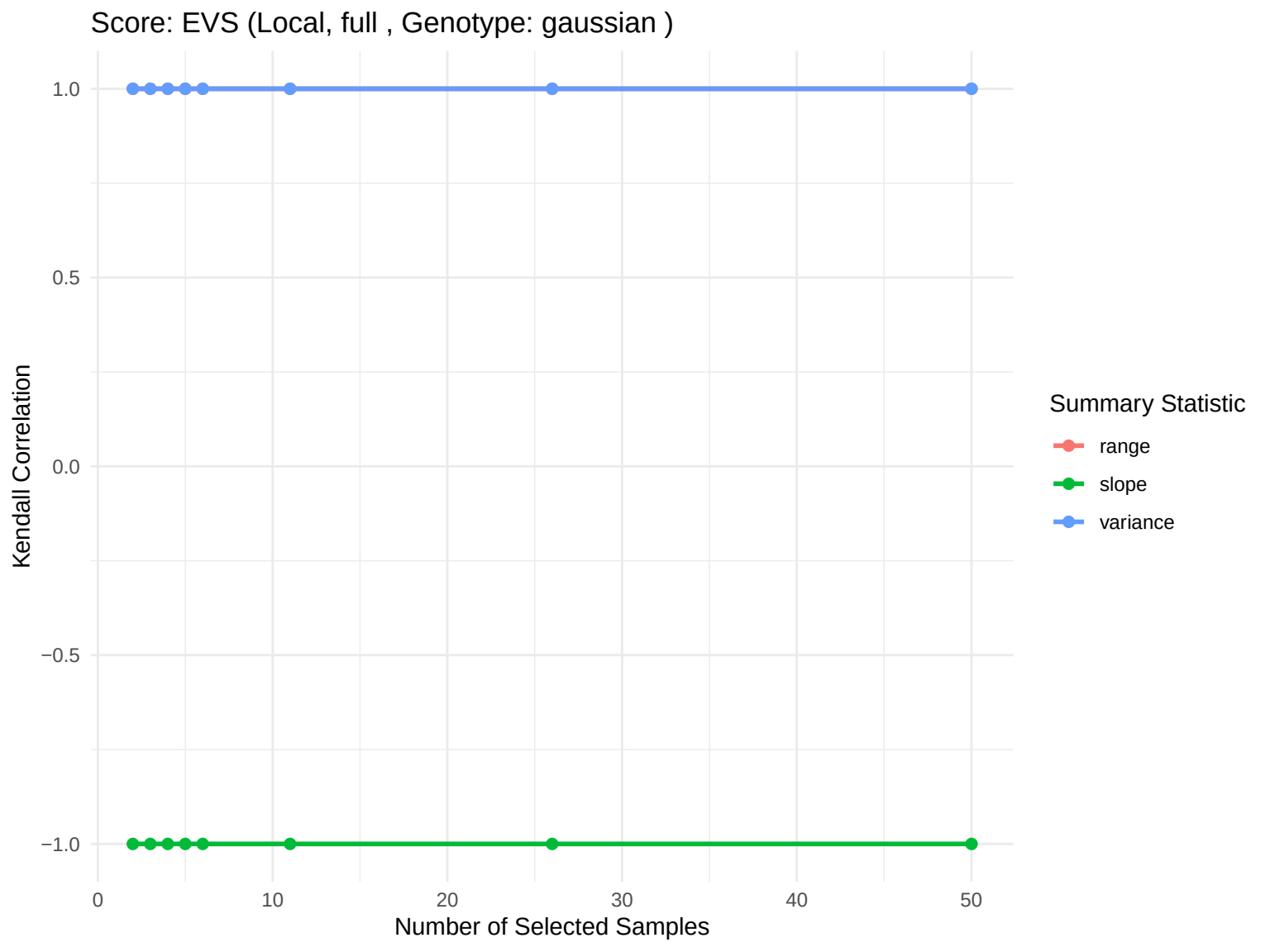
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
min
range
variance

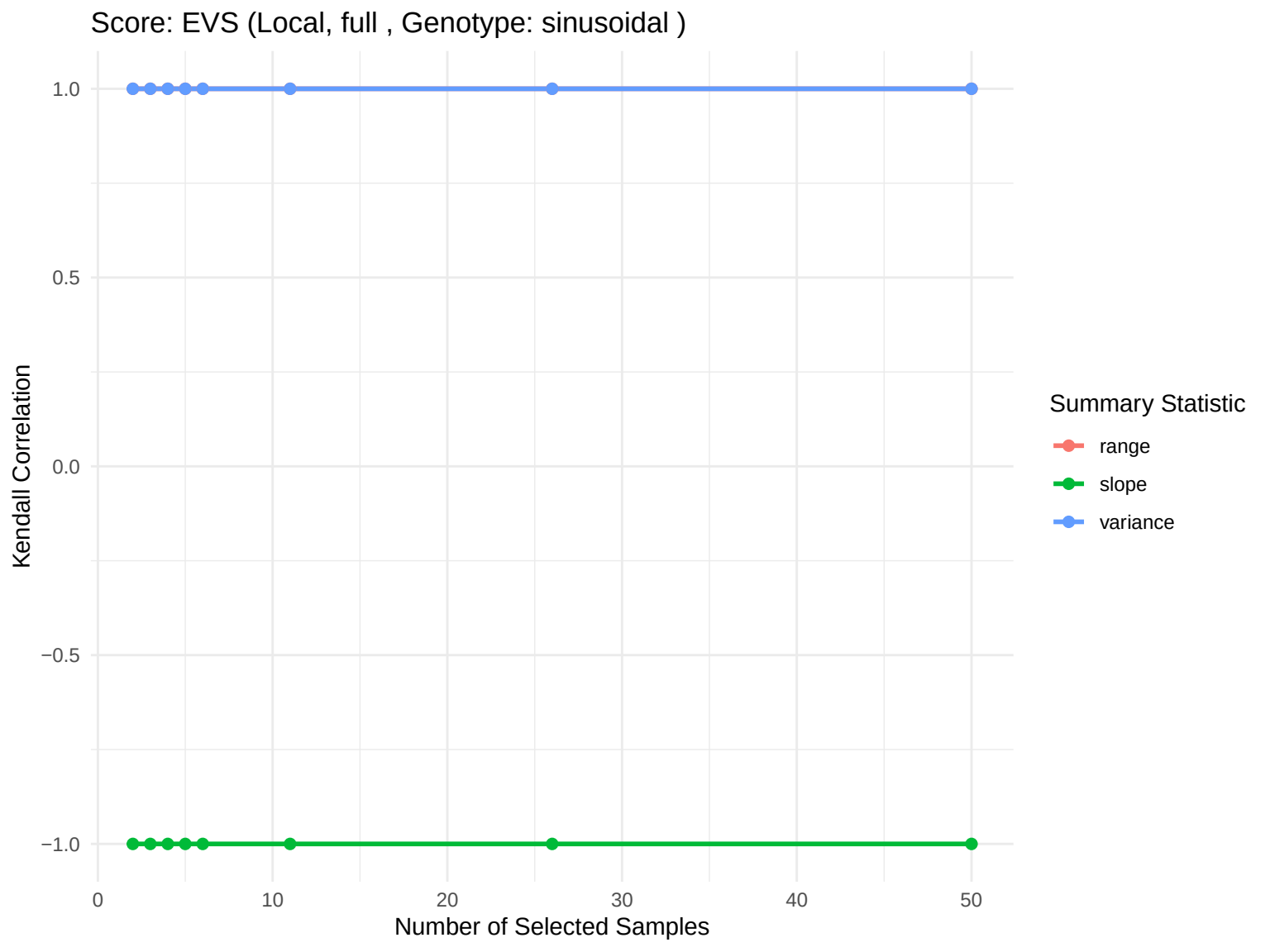

Score: RSI (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance

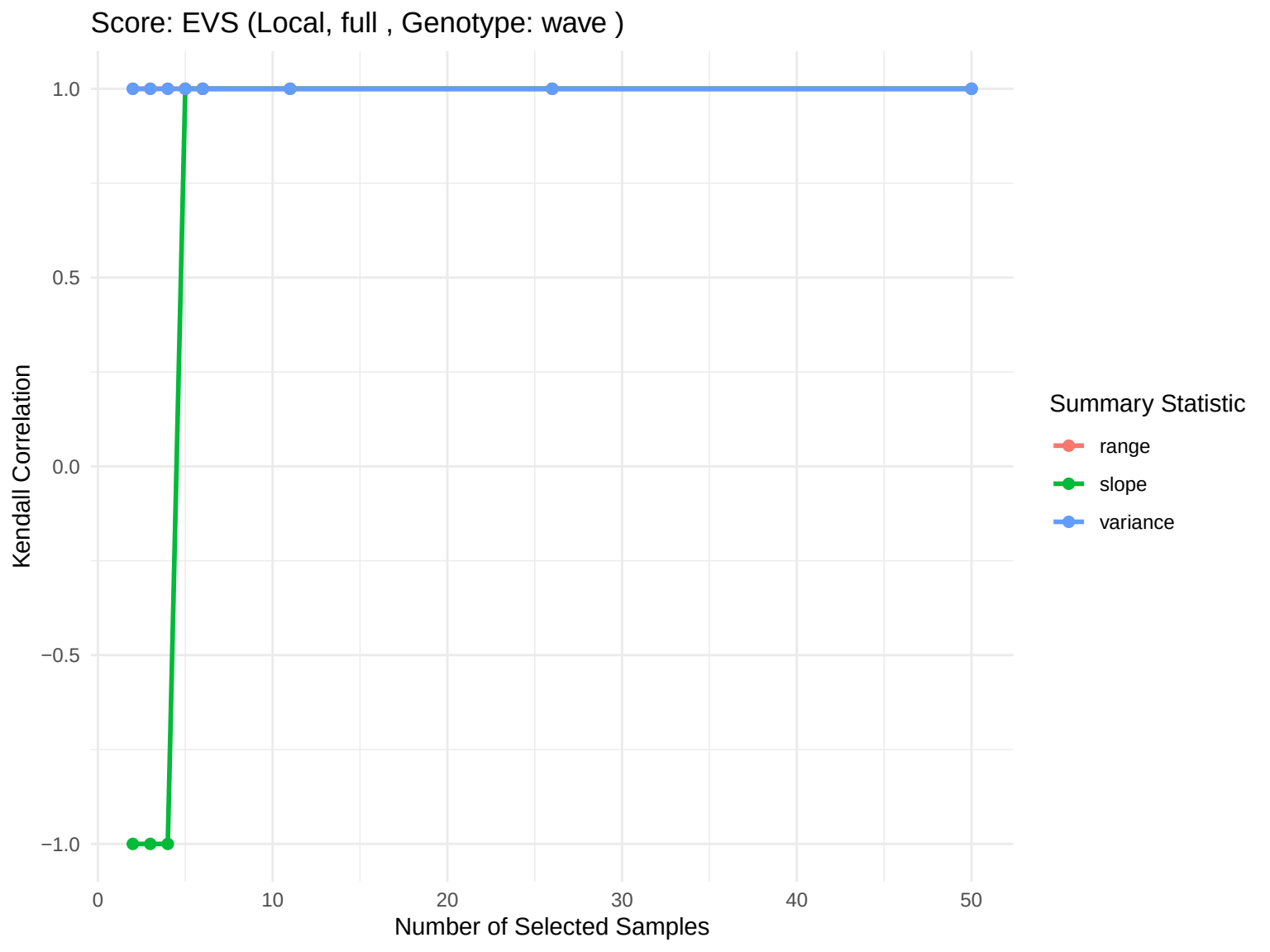

Score: RSI (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: RSI (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
max
mean
mean_lower
mean_upper
median
min
range
slope
variance


Score: EVS (Local, full , Genotype: linear)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
variance


Score: EVS (Local, full , Genotype: gaussian)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: EVS (Local, full , Genotype: sinusoidal)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance


Score: EVS (Local, full , Genotype: wave)
Kendall Correlation
1.0
0.5
0.0
−0.5
−1.0
0
10
20
30
40
50
Number of Selected Samples
Summary Statistic
range
slope
variance