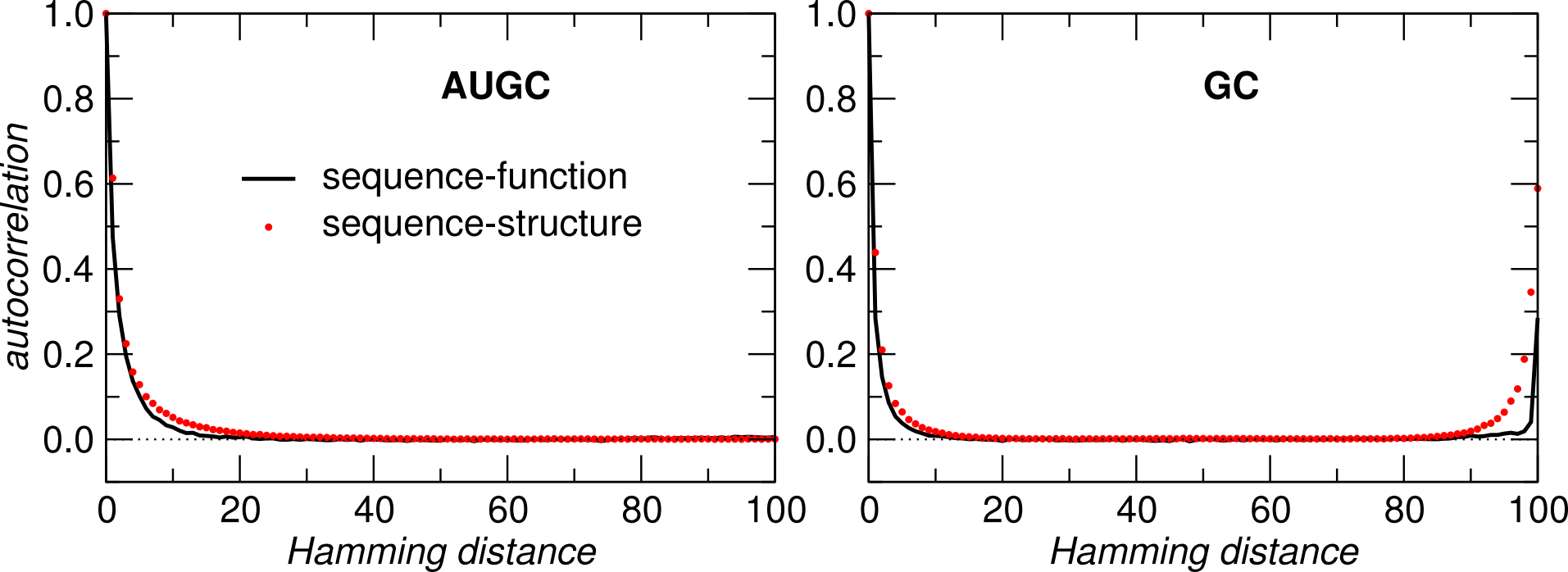
AUGC
GC
sequence-function
sequence-structure
autocorrelation
Hamming distance
Hamming distance
1.0
0.8
0.6
0.4
0.2
0.0
0
20
40
60
80
100