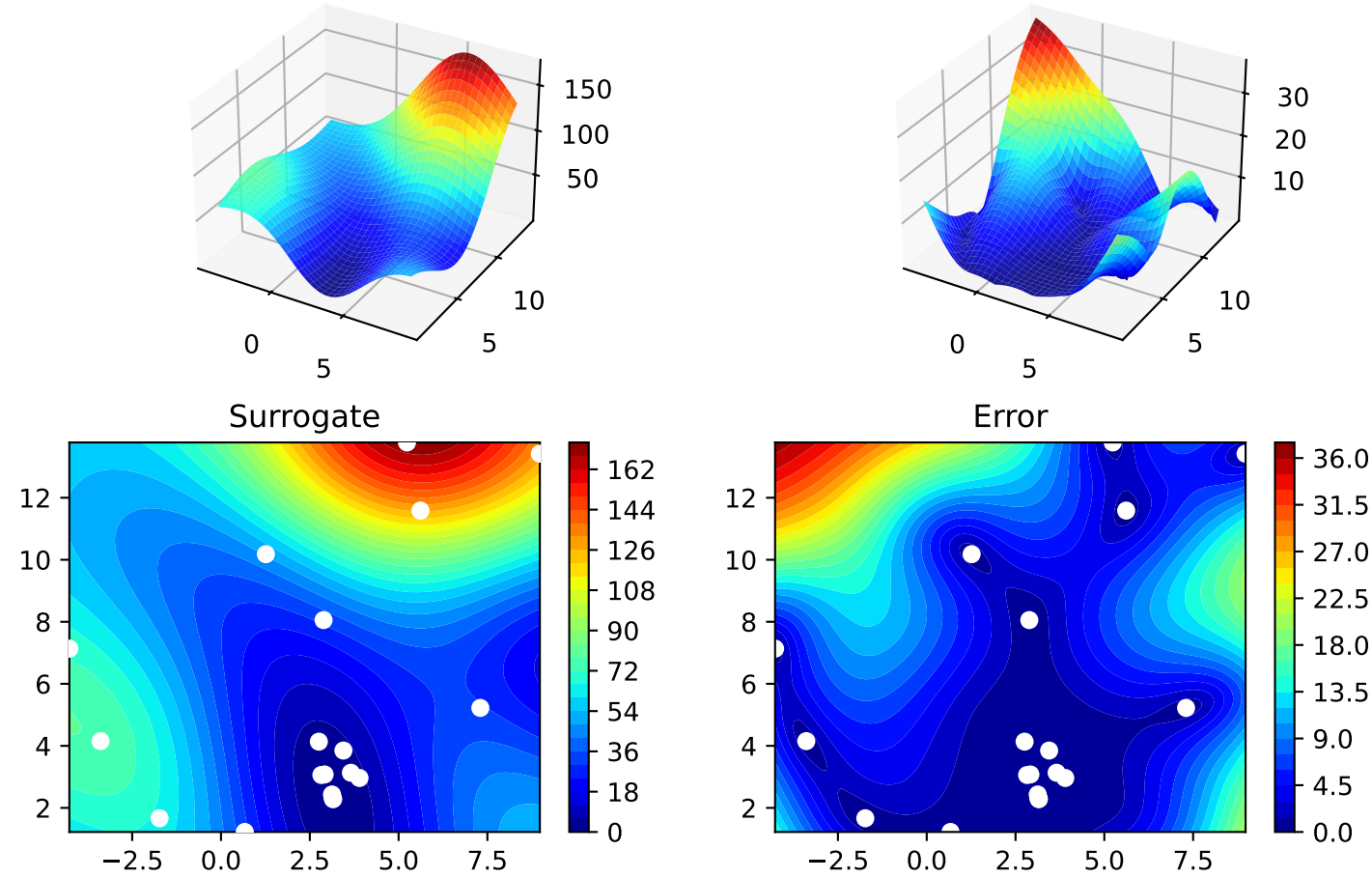
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5
10
5
30
20
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10
5
Surrogate
Error
12
10
8
6
4
2
−2.5
0.0
2.5
5.0
7.5
162
144
126
108
90
72
54
36
18
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36.0
31.5
27.0
22.5
18.0
13.5
9.0
4.5
0.0