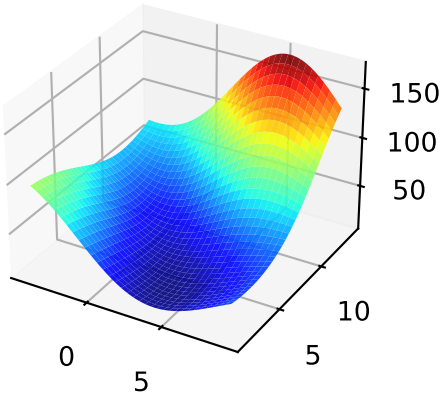
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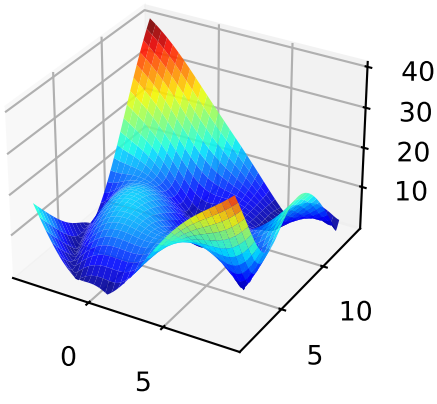
0
5
10
5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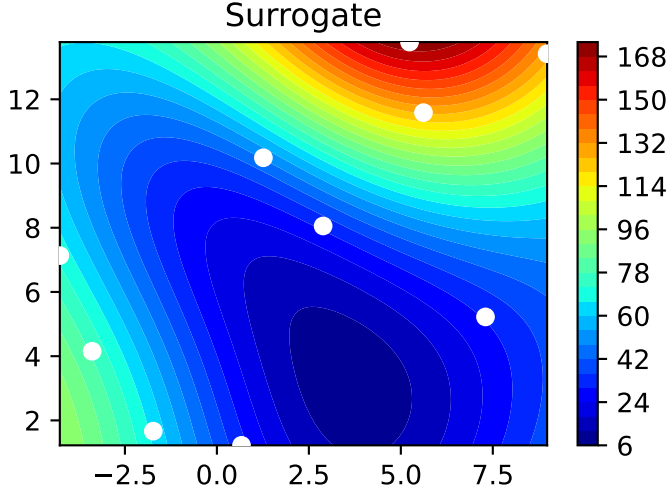
20
10
0
5
10
5


Surrogate
12
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8
6
4
2
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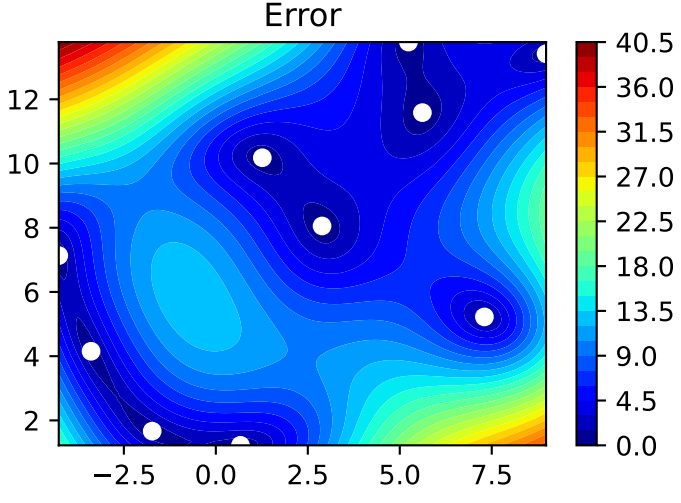
0.0
2.5
5.0
7.5
168
150
132
114
96
78
60
42
24
6


Error
12
10
8
6
4
2
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2.5
5.0
7.5
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36.0
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22.5
18.0
13.5
9.0
4.5
0.0