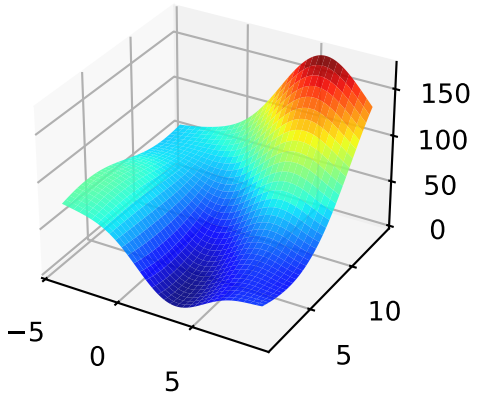
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100
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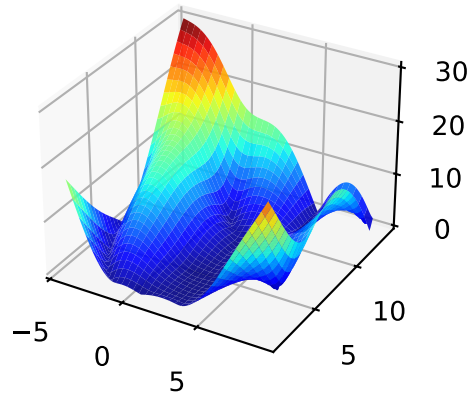
−5
0
5
10
5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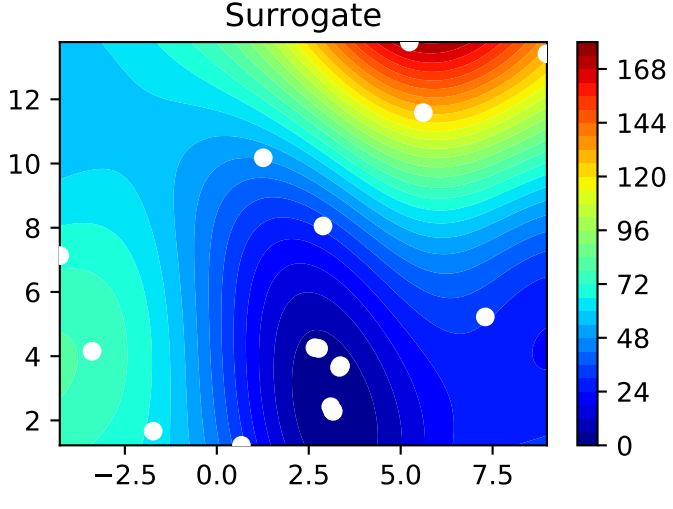
20
10
0
−5
0
5
10
5


Surrogate
12
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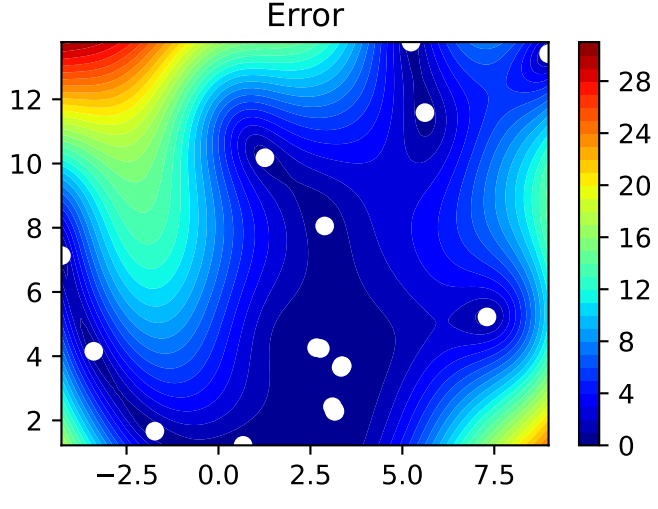
8
6
4
2
−2.5
0.0
2.5
5.0
7.5
168
144
120
96
72
48
24
0


Error
12
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8
6
4
2
−2.5
0.0
2.5
5.0
7.5
28
24
20
16
12
8
4
0