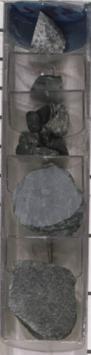


LEG

1
7
8

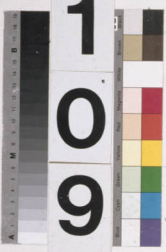
5
10
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SITE

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9
7

35
40
45
50
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60
65
70
75



HOLE

A

80
85
90
95
100
105

CORE

3
8
R

110
115
120
125
130
135
140
145