

LEG

1
7
5

SITE

1
0
8
1

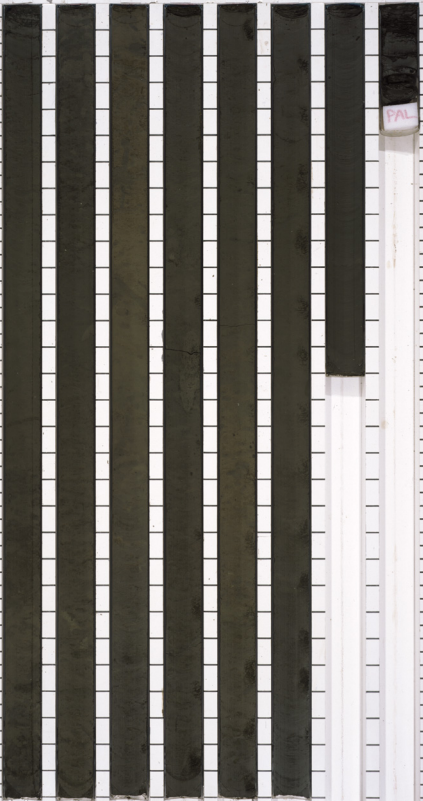
HOLE

B

CORE

1
3
H

5
10
15
20
25
30
35
40
45
50
55
60
65
70
75
80
85
90
95
100
105
110
115
120
125
130
135
140
145
150



PAL