

LEG

1
8
4

SITE

1
1
4
4

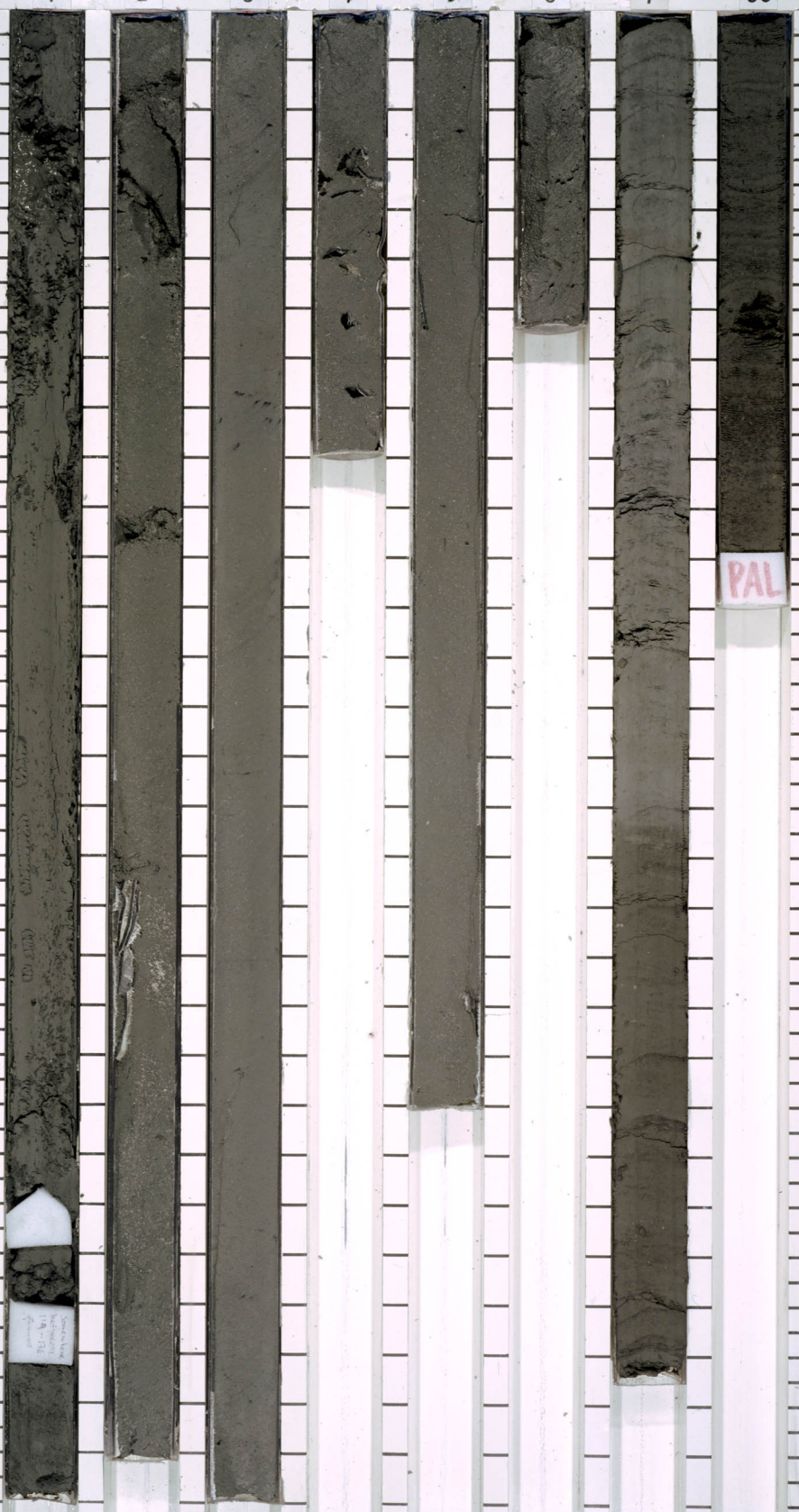
HOLE

B

CORE

2
1
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