

1177.5-1182.3 mbsf

1

2

3

LEG

3
0
5

SITE

1
3
0
9

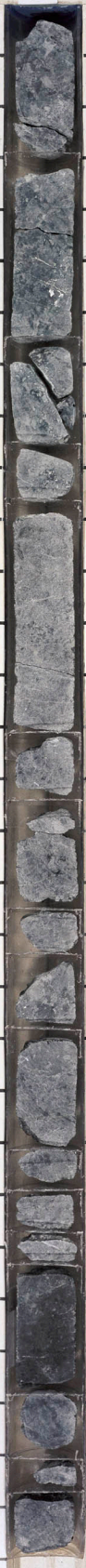
HOLE

D

CORE

2
4
5
R

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



E
M
P
T