

621.4-626.2 mbsf

1

2

3

4

LEG

3
0
5

SITE

1
3
0
9



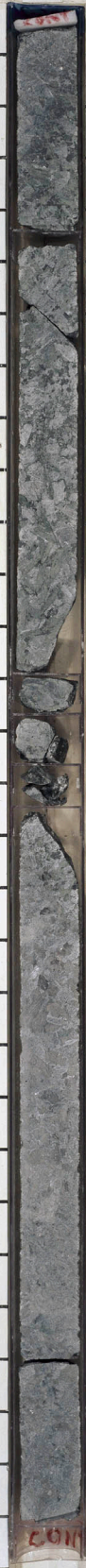
HOLE

D

CORE

1
2
6
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



CONT

14
T