

1196.7-1201.5 mbsf

1

2

3

LEG

3  
0  
5

SITE

1  
3  
0  
9



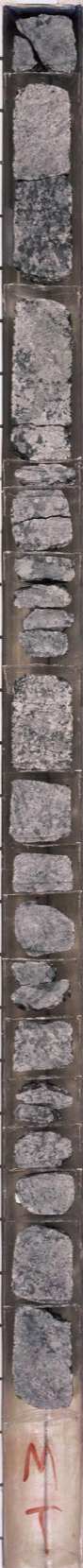
HOLE

D

CORE

2  
4  
9  
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  
Y

M  
T

M  
T