

549.4-554.2 mbsf

1

2

3

4

LEG

3  
0  
5

SITE

1  
3  
0  
9



HOLE

D

CORE

1  
1  
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R

5  
10  
15  
20  
25  
30  
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115  
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130  
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150



CONT.

A  
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P  
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Y

CONT.

M  
T

M  
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