

311.1-316.1 mbsf

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2

3

4

LEG

1

9

7

SITE

1

2

0

6

HOLE

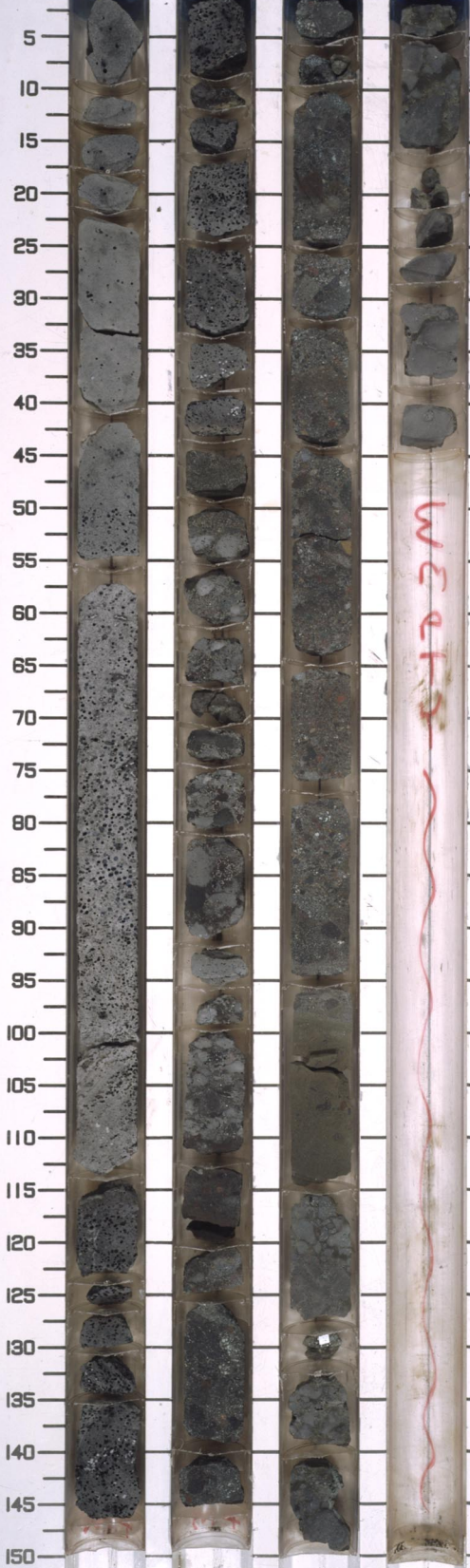
A

CORE

4

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R



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