

Demascota G-32
Sample Overview

SAMPLES

| Sample Type | Interval (m) | # of Samples |
|-------------------------|---------------------|---------------------|
| Washed Cuttings | 313.9 – 4,672.5 | 1,166 |
| Unwashed Cuttings | 313.9 – 4,672.5 | 1,157 |
| Sidewall Core | 303.2 – 3,395.4 | 143 |
| Canned Cuttings (dried) | 1,103.3 – 4,633 | 247 |

Slides

| | Interval (m) | # of Samples | Sample Source |
|---------------------|---------------------|---------------------|----------------------|
| Micropaleo slides | 304.8 – 4,660.4 | 147 | cuttings |
| Palynology slides | 304.8 – 4,660.4 | 212 | cuttings |
| Palynology slides | 1,066.8 – 3,578.4 | 77 | sidewall core |
| Nannofossil slides | 2,301.2 – 4,048.0 | 40 | cuttings |
| Nannofossil slides | 303.3 – 3,373.0 | 143 | sidewall core |
| Nannofossil slides | 3,425.6 – 4,048.0 | 4 | core |
| Thin Section slides | 3,795.7 – 3,874.3 | 2 | core |
| Thin Section slides | 3,697.2 – 4,544.6 | 9 | cuttings |

Core:

| | Interval (m) | Recovery (m) |
|---------|---------------------|---------------------|
| Core #1 | 3,422.3 – 3,434.5 | 7.3 |
| Core #2 | 3,607.6 – 3,617.7 | 7.3 |
| Core #3 | 3,872.2 – 3,881.3 | 5.0 |
| Core #4 | 4,045.6 – 4,054.8 | 3.6 |
| Core #5 | 4,389.1 – 4,398.3 | 7.2 |