

MMETSP0253-20130528|12949_1
DVNP2_sequenced

```
1      10      20      30      40      50      60
CCGTCAAAGCTGATGCCGAAGGGGGTTGTCAAAGGGAAGTGCGCAGCCAAGGTCGCCCCGA
.....
```

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DVNP2_sequenced

```
70      80      90      100     110     120
GGGCGTGGTGCGAAGGCCGCTGGTCCTCAATGGTCGCAGGGAGAGAACGGCGGGAGGTCTG
.....
```

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DVNP2_sequenced

```
130     140     150     160     170     180
AAGGCCGACATGTTGATGAGGAACCGCAGTGGAAAAGTGGTCTCGAAGCGGGCCAGTGCA
.....GATGAGGAACCGCAGTGGAAAAGTGGTCTCGAAGCGGGCCAGTGCA
```

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DVNP2_sequenced

```
190     200     210     220     230     240
GCTGGGCACGCAATTTTCGCCAATAATTGAGCCATGGCTGCAGGCTGTGATGACTGCCCGC
GCTGGGAACGCAATTTTCGCCACAATTGAGCCATGGTGCAGGCTGTGATGACTGCCCGC
```

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DVNP2_sequenced

```
250     260     270     280     290     300
GAGTGTCTCCGGGTGACGGGTTTCGTTGCAATCAACGGGCGCACCTTGCGAGGGCAAGGCA
GAGTGTCTCCGGGTGACGGGTTTCGTTGCAATCAACGGGCGCACCTTGCGAGGGCAAGGCA
```

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DVNP2_sequenced

```
310     320     330     340     350     360
CTGTACGTCAAGTCGAGGGCCATCTACGGATCGGGCTCAAGGGGACTGAGGCTGATCCA
CTGTACGTCAAGTCGAGGGCCATCTACGGATCGGGCTCAAGGGGACTGAGGCTGATCCA
```

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```
370     380     390     400     410
GTGTCATCTTCAGCGGGCGGCCAGCGCGGGTGCTGACACCATGTCGAGCTTGGCGAAGCG
GTGTCATCTTCAGCGGGCGGCCAGCGCGGGTGCTGACACCTTGTGTCGAGCTTGGCGAAGCG
```

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DVNP2_sequenced

```
420     430     440     450     460     470
GGGCTGTTCTGATGGAGGCCTCATCTTGAATTGGCGCGTGGTTGTTTCGAGGGTGGTCGAG
GGGCTGTTCTGATGGAGGCCTCATCTTGAATTGGCGCG.....
```

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DVNP2_sequenced

```
480     490
CGGCCACATTTGCCNN
.....
```