



TC_HDP2_PTs7S35F1+R1_599bp_Sangon Biotech
599 bp

[illegible][illegible]

[illegible][illegible]

Original Sequence: TC_HDP2_PTs7S35F1+R1_599bp_Sangon Biotech.dna

- 1: TC_HDP2_PTs7S35F1+R1_599bp_Sangon Biotech ➡
599 bases
1 .. 599
- 2: FM212965.1 Taenia saginata intergenic spacer ➡
1865 bases
1 .. 599 (1 mismatch)
- 3: FM212966.1 Taenia saginata intergenic spacer ➡
1865 bases
1 .. 599 (1 mismatch)
- 4: FM212967.1 Taenia saginata intergenic spacer ➡
1865 bases
1 .. 599 (2 mismatches)
- 5: FM212968.1 Taenia saginata intergenic spacer ➡
1865 bases
1 .. 599 (1 mismatch)
- 6: FM212969.1 Taenia saginata intergenic spacer ➡
1865 bases
1 .. 599 (2 mismatches)
- 7: FM212970.1 Taenia saginata intergenic spacer ➡
1865 bases
1 .. 599 (1 mismatch)
- 8: A1133740.1 Taenia saginata cross-reactive HDP2 fragment ➡
3954 bases
1055 .. 1653 (1 mismatch)
- 9: AJ582972.1 Taenia saginata asiatica GSS ➡
599 bases
1 .. 599 (39 mismatches)
- 10: FM212953.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (27 mismatches)
- 11: FM212954.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (3 mismatches)
- 12: FM212955.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (4 mismatches)
- 13: FM212956.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (4 mismatches)
- 14: FM212957.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (8 mismatches)
- 15: FM212958.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (3 mismatches)
- 16: FM212959.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (25 mismatches)
- 17: FM212960.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (4 mismatches)
- 18: FM212961.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (2 mismatches)
- 19: FM212962.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (9 mismatches)
- 20: FM212963.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (2 mismatches)
- 21: FM212964.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (3 mismatches)
- 22: FM212971.1 Taenia asiatica intergenic spacer ➡
1868 bases
1 .. 599 (4 mismatches)
- 23: EF043036.1 Taenia solium cross-reactive HDP2 fragment genomic sequence ➡
171 bases
1 .. 171 (25 mismatches)

DNA Type: Synthetic DNA
Laboratory Host Organism:

Description: TC_HDP2 (599 碱基对)

Created: 2022年10月24日
Last Modified: 2022年10月24日

Accession Number:
Code Number:

Sequence Author:

Comments:

References: