

Baccaurea ramiflora clone dauct5 satellite LANGDAU1 sequence

GenBank: MN519731.1

[FASTA Graphics](#)

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LOCUS MN519731 442 bp DNA linear PLN 12-NOV-2019

DEFINITION Baccaurea ramiflora clone dauct5 satellite LANGDAU1 sequence.

ACCESSION MN519731

VERSION MN519731.1

KEYWORDS .

SOURCE Baccaurea ramiflora

ORGANISM [Baccaurea ramiflora](#)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Tracheophyta;
Spermatophyta; Magnoliopsida; eudicotyledons; Gunneridae;
Pentapetalae; rosids; fabids; Malpighiales; Phyllanthaceae;
Antidesmatoideae; Scapeae; Baccaurea.

REFERENCE 1 (bases 1 to 442)

AUTHORS Nguyen, L.T.

TITLE Direct Submission

JOURNAL Submitted (27-SEP-2019) Genomic Mapping, Hight Agricultural
Technology Research Institute for Mekong Delta, G9-11, Campus 586,
Cai Rang, Can Tho 084, Vietnam

COMMENT ##Assembly-Data-START##
Sequencing Technology :: Sanger dideoxy sequencing
##Assembly-Data-END##

FEATURES Location/Qualifiers

source 1..442
/organism="Baccaurea ramiflora"
/mol_type="genomic DNA"
/db_xref="taxon:[566045](#)"
/clone="dauct5"

[repeat_region](#) 1..442
/rpt_type=tandem
/satellite="satellite:LANGDAU1"

ORIGIN

1 gttatitttcc caaagcggta gaagtcagag gagtgaatcg tagaagtatt ggtgtgaagg
61 gagtcttgac gcattctcat ttttacagc ttcccttcat aggcaaggct tttatatatt

121 aatcgcatTTt ttgcctgcgc aaacaatacc ttcgaagagt tttggTTTTt ggcaaagggc
181 ggacccctct catTTtggat gtaggaaggg acacttcact agTTtcgccc gtgctctTTt
241 gTTtatttgg tttccctgt cgtctcatTTt ggttgggttg tgtccgcgc ccgtcttact
301 tggtgagtgg ttgttgcat agttgttatt attatatcaa gtgtttgagc ctatatagtt
361 agtatccata gtcaaaacat tgcattgtac caaagaagaa acagcggggg aggaagggac
421 cgagaccgag gaaggtggtg ga

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