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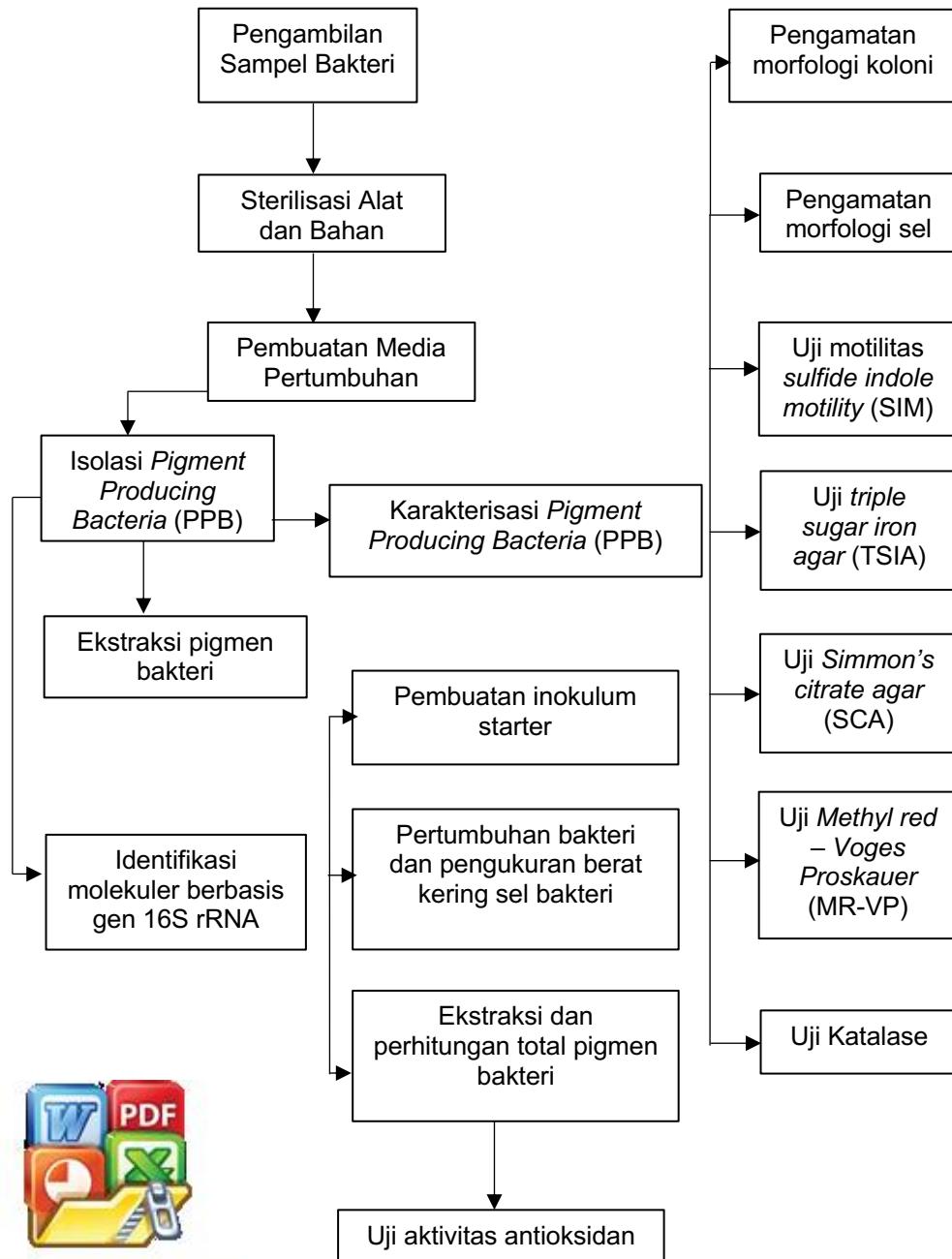
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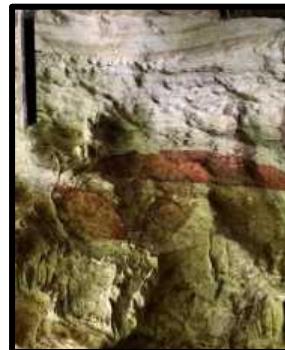
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LAMPIRAN

Lampiran 1. Alur Penelitian



Lampiran 2. Pengambilan Sampel**Lampiran 3.** Pengukuran Nilai OD untuk perhitungan total pigmen

Lampiran 4. Surat keterangan hasil uji aktivitas antioksidan DPPH

LABORATORIUM KIMIA PAKAN
JURUSAN NUTRISI DAN MAKANAN TERNAK
FAKULTAS PETERNAKAN
UNIVERSITAS HASANUDDIN

HASIL ANALISIS BAHAN

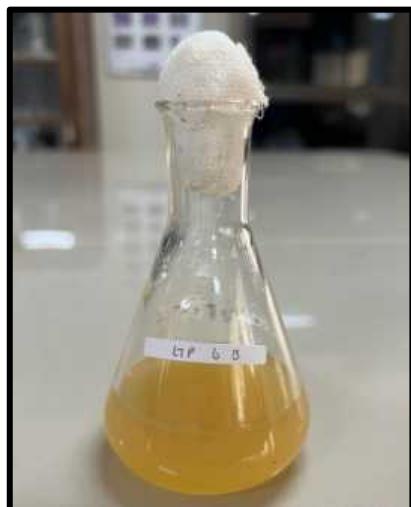
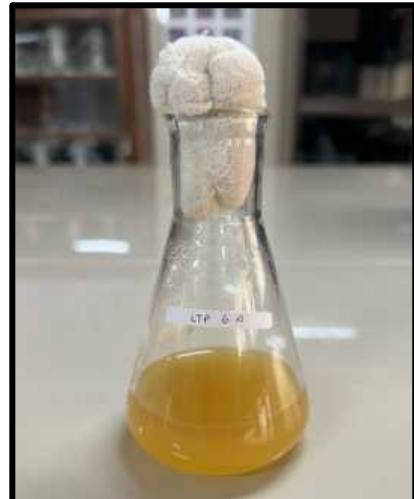
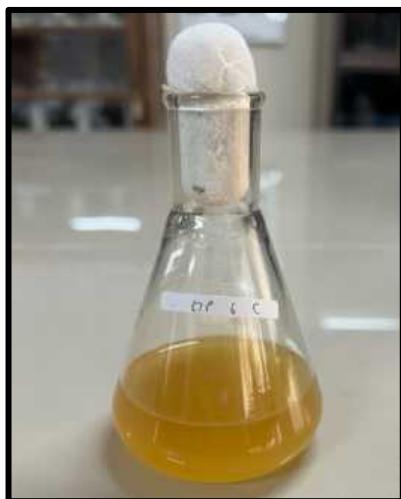
No	Kode Sampel	Antioksidan (%D _H DPPH)
1	LTP 6A	15,66
2	LTP 6B	15,80
3	LTP 6C	14,35

Malang, 3 Mei 2024

Muhammad Syahrul
Nip. 19790603 2001 12 1 001



Lampiran 5. Pertumbuhan Bakteri Inokulum pada medium cair setelah inkubasi



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Lampiran 6. Urutan Basa Nukleotida hasil BLAST NCBI *Exiguobacterium sp.*

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Exiguobacterium sp. strain Z1-19 16S ribosomal RNA gene, partial sequence
Sequence ID: [PPS94399.1](#) Length: 1435 Number of Matches: 1

Range 1: 11 to 1329 [GenBank](#) [Graphics](#) [Next Match](#) [Previous Match](#)

Score 2183 bits(1182)	Expect 0.0	Identities 1276/1319(97%)	Gaps 30/1319(2%)	Strand Plus/Minus
Query 11	TGCTGA-CTGCAGATTACTAGCGATTCCGACTTCATGCAGCGAGGTTGCAGCCCTGCAATCC			
Sbjct 1329	TGCTGACCTTGATTAATCAGCGATTCCGACTTCATGCAGCGAGGTTGCAGCCCTGCAATCC			
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Sbjct 1269	GAATCTGGAAACGGCTTTATGGGATTTGGCTCACCTCGCGGTCTCGCTGCCCTTTGTACCG			
Query 130	TCCATTGTAGCACCGTGTGAGGCCAACTCATAAGGGGCATGATGATTGACGTATCCCC			
Sbjct 1209	TCCATTGTAGCACCGTGTGAGGCCAACTCATAAGGGGCATGATGATTGACGTATCCCC			
Query 190	ACCTTCCCTCGGTTTGTGACCGGAGTCTCCCTAGAGTGCCAAGTGAATGCTGGCAACT			
Sbjct 1149	ACCTTCCCTCGGTTTGTGACCGGAGTCTCCCTAGAGTGCCAAGTGAATGCTGGCAACT			
Query 250	AAGGATAGGGTTGGCGCTGGTGCGGGACTTAACCCAAACATCTCACGACACGGAGCTGACG			
Sbjct 1089	AAGGATAGGGTTGGCGCTGGTGCGGGACTTAACCCAAACATCTCACGACACGGAGCTGACG			
Query 310	ACAACCATGCACCAACTGTACCCATTGTCCCCGAAGGGAAAACCTTGAATCTCAAGCGGT			
Sbjct 1029	ACAACCATGCACCAACTGTACCCATTGTCCCCGAAGGGAAAACCTTGAATCTCAAGCGGT			
Query 370	CAATGGGATGTCAGAAGTTGGTAAGGTTCTTGGCGTTGCTTGAATTAAACCATGTC			
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Query 430	CACCGCTTGTGGGGTCCCCGTCAATTCTCTTGGATTTCAGGCCCTGGGGCGTACTCCCC			
Sbjct 909	CACCGCTTGTGGGGTCCCCGTCAATTCTCTTGGATTTCAGGCCCTGGGGCGTACTCCCC			
Query 490	AGGGCGGAGTCTTAATGGTGTAGCTTACAGACTGAGGGGGGAAACCCCCAACACCTAG			
Sbjct 849	AGGGCGGAGTCTTAATGGTGTAGCTTACAGACTGAGGGGGGAAACCCCCAACACCTAG			
Query 550	CACTCATCGTTACGGGGTGGAACCTACACGGGTATCTAACTCTGTGTTGCTCCCCAGCTTT			
Sbjct 789	CACTCATCGTTACGGGGTGGAACCTACACGGGTATCTAACTCTGTGTTGCTCCCCAGCTT			
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Sbjct 669	CTCTACGEATTTCACCGCTACACGTGGAAATTCCACTCTTCCTCTGACTCAAGCTTC			
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Query 850	GCTGCTGGCACGTAGTTAGCCGTGGCTTCTGGTAAGGTACCGTCAGGTAAGGCAATT			
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Sbjct 429	CTCTTCGTACGTGTTCTTCCCTTACAACAGAGTTTACGATCCGAAAAACCTTCATCACTC			
Query 970	ACGGCGGGTTGCTCCATCAGACTTCTGGCCATTGGAAAGATTCCCTACTGCTGGCTCCC			
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Query 1030	GTAGGAGTCGGGGCGGTGTCAGTCCCCAGTGTGGCCGATCACCTCTAGGTGGCTAT			
Sbjct 309	GTAGGAGTCGGGGCGGTGTCAGTCCCCAGTGTGGCCGATCACCTCTAGGTGGCTAT			



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Ornithinibacillus scapharcae.

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Ornithinibacillus scapharcae strain N-2-1-2 16S ribosomal RNA gene, partial sequence
 Sequence ID: [KU601305.1](#) Length: 1422 Number of Matches: 1

Range 1: 9 to 1327 [GenBank](#) [Graphics](#) [▼ Next Match](#) [▲ Previous Match](#)

Score 2178 bits(1179)	Expect 0.0	Identities 1275/1319(97%)	Gaps 30/1319(2%)	Strand Plus/Minus
Query 11	TGCTGA-CTGCAGATTACTAGCGATTCCGACTTCATGCAGCGAGTTGCAGCCTGCAATCC			
Sbjct 1327	TGCTGACCTGGATTACTAGCGATTCCGACTTCATGCAGCGAGTTGCAGCCTGCAATCC			
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Sbjct 727	CGGCCCTCAGGGTCAAGTACAGACAAAGAGTCGCTTGGCAACTGGTTCTCCACAT			
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Sbjct 667	CTCTACGGCATTTACCGCTACACGTGGAAATTCACCTTCTCTCTGTACTCAAGCTTC			
Query 730	CAGTTTCAATGGCCCTCCCCGGTTGAGCCGGGGCTTTCACATCAGACTAAAGGCCG			
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Lampiran 7. Hasil Perhitungan Jarak Genetik metode UPGMA MEGA XI

Spesies	LTP6-A	LTP6-B	LTP6-C
Ornithinibacillus scapharcae strain N-2-1-2	0,00143061614051	0,00071479640479	0,00143061614051
Exiguobacterium sp.2A11S4	0,0014306161405	0,0021474621414	0,0014306161405
Exiguobacterium sp.NIU-K2	0,0014299763755	0,0007146686713	0,0014299763755
Exiguobacterium sp.NIU-K4	0,0014299763755	0,0007146686713	0,0014299763755
Exiguobacterium sp.BDH26	0,0007147964047	0,00143061614	0,0021474621414
Exiguobacterium sp.IARI-R-140	0,0007147964047	0,0000000000	0,0007147964047



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