Appendix (1) Sequence of PCR product of amplification of 16S rDNA by two sets of primer (B16SF and B16SR) of the *Bacillus* isolate B1.

ATGGGAGCTTGCTCCCTGATGTTAGCGGCGGACGGGTGAGTAACACGT GGGTAACCTGCCTGTAAGACTGGGATAACTCCGGGAAACCGGGGGCTAA TACCGGATGGTTGTTTGAACCGCATGGTTCAAACATAAAAGGTGGCTTC GGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGAGG TAACGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGATC GGCCACACTGGGACTGAGACACGGCCCAAACTCCTACGGGAGGCAGCA GTGGTCATAGCTGTTTCCTGAA

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GCATGGGAGCTTGCTCCCTGATGTTAGCGGCGGACGGGTGAGTAACAC

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GTGGGTAACCTGCCTGTAAGACTGGGATAACTCCGGGAAACCGGGGGCT AATACCGGATGCTTGTTTGAACCGCATGGTTCAAACATAAAAGGTGGC TTCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA GGTAACGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGA TCGGCCACACTGGGACTGAGACACGGCCCAAACTCCTACGGGAGGCAG CAGTGGTCATAGCTGTTTCCTGA Appendix (3) Sequence of PCR products of amplification of 16S rDNA by two sets of primer (B16SF and B16SR) of the *Bacillus* isolate B3.

GAATGGGAGCTTGCTCCCTGATGTTAGCGGCGGACGGGTGAGTAACAC GTGGGTAACCTGCCTGTAAGACTGGGATAACTCCGGGAAACCGGGGCT AATACCGGATGGTTGTTTGAACCGCATGGTTCAAACATAAAAGGTGGT TCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA GGTAACGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGA TCGGCCACACTGGGACTGAAACACGGCCCAAACTCCTACGGGAGGCAG CAGTGGTCATAGCTGTTTCCTGA

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Spendre (4) Sequence of PCR products of amplification of 16S rDNA by two sets

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of primer (B16SF and B16SR) of the *Bacillus* isolate B4.

GAATGGGAGCTTGCTCCCTGATGTTAGCGGCGGACGGGTGAGTAACAC GTGGGTAACCTGCCTGTAAGACTGGGATAACTCCGGGAAACCGGGGCT AATACCGGATGGTTGTTTGAACCGCATGGTTCAAACATAAAAGGTGGC TTCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA GGTAACGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGA TCGGCCACACTGGGACTGAAACACGGCCCAAACTCCTACGGGAGGCAG CAGTGGTCATAGCTGTTTCCGGAA Appendix (5) Sequence of PCR products of amplification of 16S rDNA by two sets of primer (B16SF and B16SR) of the *Bacillus* isolate B5.

GAATGGGAGCTTGCTCCCTGATGTTAGCGGCGGACGGGTGAGTAACAC GTGGGTAACCTGCCTGTAAGACTGGGATAACTCCGGGAAACCGGGGCT AATACCGGATGCTTGTTTGAACCGCATGGTTCAAACATAAAAGGTGGC TTCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA GGTAATGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGA TCGGCCACACTGGGACTGAGACACGGCCCAAACTCCTACGGGAGGCAG CAGTGGTCATAGCTGTTTCCGGAA

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GAATGGGAGCTTGCTCCCTGATGTTAGCGGCGGACGGGTGAGTAACAC

GTGGGTAACCTGCCTGTAAGACTGGGATAACTCCGGGAAACCGGGGCT AATACCGGATGCTTGTTTGAACCGCATGGTTCAAACATAAAAGGTGGC TTCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA GGTAATGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGA TCGGCCACACTGGGACTGAGACACGGCCCAAACTCCTACGGGAGGCAG CAGTGGTCATAGCTGTTTCCGGAA Appendix (7) Sequence of PCR products of amplification of 16S rDNA by two sets of primer (B16SF and B16SR) of the *Bacillus* isolate B8.

GAATGGGAGCTTGCTCCCTGATGTTAGCGGCGGACGGGTGAGTAACAC GTGGGTAACCTGCCTGTAAGACTGGGATAACTCCGGGAAACCGGGGGCT AATACCGGATGGTTGTTTGAACCGCATGGTTCAAACATAAAAGGTGGC TTCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA GGTAACGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGA TCGGCCACACTGGGACTGAGACACGGCCCAAACTCCTACGGGAGGCAG CAGTGGTCATAGCTGTTTCCGGA

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GTGGGTAACCTGCCTGTAAGACTGGGATAACTCCGGGAAAACCGGGGGCT AATACCGGATGCTTGTTTGAACCGCATGGTTCAAACATAAAAGGTGGC TTCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA GGTAATGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGGTGA TCGGCCACACTGGGACTGAGACACGGCCCAAACTCCTACGGGAGGCAG CAGTGGTCATAGCTGTTTCCGGAA Appendix (9) Sequence of PCR products of amplification of 16S rDNA by two sets of primer (B16SF and B16SR) of the *Bacillus* isolate B10.

GAATGGGAGCTTGCTCCCTGATGTTAGCGGCGGACGGGTGAGTAACAC GTGGGTAACCTGCCTGTAAGACTGGGATAACTCCGGGAAACCGGGGCT AATACCGGATGCTTGTTTGAACCGCATGGTTCAAACATAAAAGGTGGC TTCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA GGTAATGGCTCACCAAGGCAACGATGCGTAGCCGACCTGAGAGGGTGA TCGGCCACACTGGGACTGAGACACGGCCCAAACTCCTACGGGAGGCAG CAGTGGTCATAGCTGTTTCCGGAA

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TTGCAGACTGCGATCCGAACTGAGAACAGATTTGTGGGATTGGCTTAA CCTCGCGGTTTCGCTGCCCTTTGTTCTGTCCATTGTAGCACGTGTGTAGC CCAGGTCATAAGGGGCATGATGATGATTGACGTCATCCCCACCTTCCTCCG GTTTGTCACCGGCAGTCACCTTAGAGTGCCCAACTGAATGCTGGCAACT AAGATCAAGGGTTGCGCTCGTTGCGGGGACTTAACCCAACATCTCACGA CACGAGCTGACGACAACCATGCACCACCTGTCACTCTGCCCCCGAAGG GGACGTCCTATCTCTAGGATTGTCAGAGGATGTCAAGACCTGGTAAGG TTCTTCGCGTTGCTTCGAATTAAACCACATGCTCCACCGCTTGTGCGGGG CCCCCGTCAATTCCTTTGAGTTTCAGTCTTGCGACCGTACTCCCCAGGC GGAGTGCTTAATGCGTTAGCTGCAGCACTAAGGGGCGGAAACCCCCTA ACACTTAGCACTCATCGTTTACGGCGTGGACTACCAGGGTATCTAATCC TGTTCGCTCCCCACGCTTTCGCTCCTCAGCGTCAGTTACAGACCAGAGA GTCGCCTTCGCCACTGGTGTTCCTCCACATCTCTACGCATTTCACCGCTA CACGTGGAATTCCACTCTCCTCTTCTGCACTCAAGTTCCCCAGTTTCCAA TGACCCTCCCCGGTTGAGCCGGGGGGCTTTCACATCAGACTTAAGGAACC GCCTGCGAGCCCTTTACGCCCAATAATTCCGGACAACGCTTGCCACCTA CGTATTACCGCGGCTGCTGGCACGTAGTTAGCCGTGGCTTTCTGGTTAG GTACCGTCAAGGTACCGCCCAATAACCTTCACCTCACGCGCGCCCTTCCCTACC

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CGCCGCTAACATCAGGGAGCAAGCTCCCATCTGTCCGCTCGACTTGCAT GTATTAGGCA.

Appendix (11) Sequence of PCR products of amplification of 16s rDNA by two sets of primers (27F and 1492R) of the *Bacillus* isolate B9.

CGGCTGGCTCCTAAAAGGTTACCTCACCGACTTCGGGTGTTACAAACTC TCGTGGTGTGACGGGCGGTGTGTACAAGGCCCGGGAACGTATTCACCG CGGCATGCTGATCCGCGATTACTAGCGATTCCAGCTTCACGCAGTCGAG TTGCAGACTGCGATCCGAACTGAGAACAGATTTGTGGGGATTGGCTTAA CCTCGCGGTTTCGCTGCCCTTTGTTCTGTCCATTGTAGCACGTGTGTAGC CCAGGTCATAAGGGGCATGATGATTGACGTCATCCCCACCTTCCTCCG

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GTTTGTCACCGGCAGTCACCTTAGAGTGCCCAACTGAATGCTGGCAACT

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ACACTTAGCACTCATCGTTTACGGCGTGGACTACCAGGGTATCTAATCC

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TGTTCGCTCCCACGCTTTCGCTCCTCAGCGTCAGTTACAGACCAGAGA GTCGCCTTCGCCACTGGTGTTCCTCCACATCTCTACGCATTTCACCGCTA CACGTGGAATTCCACTCTCCTCTTCTGCACTCAAGTTCCCCAGTTTCCAA TGACCCTCCCCGGTTGAGCCGGGGGGCTTTCACATCAGACTTAAGAAACC GCCTGCGAGCCCTTTACGCCCAATAATTCCGGACAACGCTTGCCACCTA CGTATTACCGCGGCTGCTGGCACGTAGTTAGCCGTGGCTTTCTGGTTAG GTACCGTCAAGGTACCGCCCTATTCGAACGGTACTTGTTCTTCCCTAAC AACAGAGCTTTACGATCCGAAAACCTTCATCACTCACGCGGCGTTGCTC CGTCAGACTTTCGTCCATTGCGGAAGATTCCCTACTGCTGCCTCCCGTA GGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCCGATCACCCTCTCAGG TCGGCTACGCATCGTTGCCTTGGTGAGCCATTACCTCACCAACTAGCTA ATGCGCCGCGGGTCCATCTGTAAGTGGTAGCCGAAGCCACCTTTTATGT TTGAACCATGCGGTTCAAACAAGCATCCGGTATTAGCCCCGGTTTCCCG GAGTTATCCCAGTCTTACAGGCAGGTTACCCACGTGTTACTCACCCGTC CGCCGCTAACATCAGGGAGCAAGCTCCCATCTGTCCGCTCGACTTGCAT GTATTAGGC

# Appendix (12) Sequence of PCR products of amplification of 16s rDNA by two sets of primers (27F and 1492R) of the *Bacillus* isolate B1.

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ACGATGCGTAGCCGACCTGAGAGGGTGATCGGCCACACTGGGACTGAG ACACGGCCCAGACTCCTACGGGAGGCAGCAGCAGTAGGGAATCTTCCGCAA TGGACGAAAGTCTGACGGAGCAACGCCGCGTGAGTGATGAAGGTTTTC GGATCGTAAAGCTCTGTTGTTAGGGAAGAACAAGTACCGTTCAAATAG GGCGGTACCTTGACGGTACCTAACCAGAAAGCCACGGCTAACTACGTG CCAGCAGCCGCGGTAATACGTAGGTGGCAAGCGTTGTCCGGAATTATT GGGCGTAAAGGGCTCGCAGGCGGTTTCTTAAGTCTGATGTGAAAGCCC CCGGCTCAACCGGGGAGGGTCATTGGAAACTGGGGAACTTGAGTGCAG AAGAGGAGAGTGGAATTCCCCGTGTATCGGTGAAATGCGAAGAGATGT 

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CCAGGGTATCTAATCCTGTTCGCTCCCACGCTTTCGCTCCTCAGCGTC AGTTACAGACCAGAGAGTCGCCTTCGCCACTGGTGTTCCTCCACATCTC TACGCATTTCACCGCTACACGTGGAATTCCACTCGCCTCTTCTGCGTCA ATGTTCCCCAGTTTCCAATGACCCTCCCCGGTTGAGCTGGGGGGTTTTCA CTTCAACTTAAGAAACCGCCAGCGAGCCTTTACGCCCATATTTGG Appendix (13) Sequence of PCR products of amplification of 16s rDNA by two sets of primers (27F and 1492R) of the *Bacillus* isolate B2.

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GAGGAGAGTGGAATTCCACGTGTAGCGGAGAAATGCGTAGAGATGTG

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TAGCCGTGCTTTCAGTAAGAACGCAAGCACCGCCTATTCCAACGG

Appendix (14) Sequence of PCR products of amplification of 16s rDNA by two sets of primers (27F and 1492R) of the *Bacillus* isolate B3.

GACCCGCGGCGCATTAGCTAGTTGGTGAAGGTAACGGCTCACCAAGGC AACGATGCGTAGCCGACCTGAGAGGGGTGATCGGCCACACTGGGACTGA GACACGGCCCAGACTCCTACGGGAGGCAGCAGCAGTAGGGAATCTTCCGCA ATGGACGAAAGTCTGACGGAGCAACGCCGCGTGAGTGATGAAGGTTTT CGGATCGTAAAGCTCTGTTGTTAGGGAAGAACAAGTACCGTTCGAATA GGGCGGTACCTTGACGGTACCTAACCAGAAAGCCACGGCTAACTACGT GCCAGCAGCCGCGGTAATACGTAGGTGGCAAGCGTTGTCCGGAATTAT TGGGCGTAAAGGGCTCGCAGGCGGTTTCTTAAGTCTGATGTGAAAGCC CCCGGCTCACCCGGGGAGGGGTCTTTGGAAACTGAGGATTTTGTGCGTAT

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CTATGATCGAGGGGTGGGCTCGTTTCTTGTCGTAACCCAAAAAGAAAA

GACGCGCTTTGAATACAACCATGCAAGGCCTGACGTTATGGCTCCAAA GGGTAGGGGATATGTATAAGTTTTTCCGATTGCGTTTTGGCCTGTGCAG TCTAGACGCATTGCTGATTTTTTGAGCATTCACTACAGGTATTGGTCCG GCACCGATTCCTCATAACTGTTATTCCCCTTCTTTACTCCCTGTAAGATC CCTTCGCGCCACCCGAAC Appendix (15) Sequence of PCR products of amplification of 16s rDNA by two sets of primers (27F and 1492R) of the *Bacillus* isolate B4.

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TGGAGGAACCTAAGTGGATAAACGCGACCTCTCTGT////CGGCTGGCTCC TAAAAGGTTACCTCACCGACTTCGGGTGTTACAAACTCTCGTGGTGTGA CGGGCGGTGTGTACAAGGCCCGGGAACGTATTCACCGCGGCATGCTGA TCCGCGATTACTAGCGATTCCAGCTTCACGCAGGCGAGTTGCAGACTGC GATCCGAACTGAGAACAGATTTGTGGGATTGGCTTAACCTCGCGGTTTC GCTGCCCTTTGTTCTGTCCATTGTAACACGTGTGTAGCCCAGGTCATAA GGTGCATGATGATTTGACGTCATCCCCACCTTCCTCCGGTTTGTCACCG GCAGTCACCTTACAGTGCCCAATTGAATGCTGGCTACTAAAATCAGGGT

## 

Appendix (16) Sequence of PCR products of amplification of 16s rDNA by two sets of primers (27F and 1492R) of the *Bacillus* isolate B5.

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#### GCGGTACCTTGACGGTACCTAACCAGAAAGCCACGGCTAACTACGTG

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CCAGCAGCCGCGGTAATACGTAGGTGGCAAGCGTTGTCCGGAATTATT GGGCGTAAAGGGCTCGCAGGCGGTTTCTTAAGTCTGATGTGAAAGCCC CCGGCTCAACCGGGGAGGGTCATTGGAAACTGGGGGAACTTGAGTGCAG AAGAGGAGAGAGTGGAATTCCACGTGTAGCGGTGAAATGCGTAGAGATGT GGAGGAACACCAGTGGCGAAGGCGACTCTCTGGTCTGTAACTGACGCT GAGGAGCGAAAGCGTGGGGGAGCGAACAGGATTAGATACCCCTGTTAGT CCCCGCCGTAAACGATGAGAGCTAAGTGTAGGGGGGTTTCCGCCCCTTA GCGCTGCAGCTAACACAATAAGAACTCCCGCCTGGGGAGTACGGGTC/// TCGTGGTGTGACGGGCGGTGTGTACAAGGCCCGGGAACGTATTCACCG CGGCATGCTGATCCGCGATTACTAGCGATTCCAGCTTCACGCAGTCGAG TTGCAGACTGCGATCCGAACTGAGAACAGATTTGTGGGATTGGCTTAA CCTCGCGGTTTCGCTGCCCTTTGTTCTGTCCATTGTAGCACGTGTGTAGC CCAGGTCATAAGGGGCATGATGATGATTTGACGTCATCCCCACCTTCCTCCG GTTTGTCACCGGCAGTCACCTTAGAGTGCCCAACTGAATGCTGGCAACT AAGATCAAGGGTTGCGCTCGTTGCGGGGACTTAACCCAACATCTCACGA CACGAGCTGACGACAACCATGCACCACCTGTCACTCTGCCCCGAAGG GGACGTCCTATCTCTAGGATTGTCAGAGGATGTCAAGACCTGGTAAGG

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ATGACCCTCCCGGGGTGAGCCGGGGGGCTTTCACATCAGCTTAAAAACC GCCTGCGAGCCTTTACGCCAATAATTCCGGACAACGTTTGCCACTACGT ATTACAAGGTGCTGGCACTAG

Appendix (18) Sequence of PCR products of amplification of 16s rDNA by two sets of primers (27F and 1492R) of the *Bacillus* isolate B7.

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#### GAGGAGAGTGGAATTCCACGTGTAGCGGTGAAATGCGTAGAGATGTG

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GAGGAACACCAGTGGCGAAGGCGACTCTCTGGTCTGTAACTGACGCTG AGGAGCGAAAGCGTGGGGGGGGCGAACAGGATTAGATACCCTGGTAGTCC ACGCCGTAAACGATGAGTGCTAAGTGTTAGGGGGGTTTCCGCCCCTTAGT GCTGCAGCTAACGCATTAAGCACTCCGCCTGGGGAGTACGGTCGCAAG ACTGAAAACTCAAAGGAATTAGACGGGGGGGCCCGCACAAGCGGTGGA GCATGTGGTTTATTTCGAAGCAACGCGAAGAACTTTACCAGGTCTTGAC ATCCTCTGACAATTCTAGAAGATAAGACGTCCCCCTTCGGGGGCAGAGT GACAGGTGGAGCATGGTTGTCGTTCAGCTCGTCGTCGTAGATTGTTGGG TTTAAGTCCCCGCAACGAAGCACCACCTCCTGAATCCTTAGCTTGTCAA GCCATTCATGTTTGGCTACTTCTGATGGTGGACTTGGCCGGTTGACACA ACCCGGAAGGAAGGGAGGAGAGATGAACGTAGAAATCATCTGTCTCCTTA TTGATCCTGCGTTCTCTCAGCTCGTCTTTCACAAGTCGCAGCAAGTAAT CTAGATTG///GCGGCTGGCTCCTAAAAGGTTACCTCACCGACTTCGGGT GTTACAAACTCTCGTGGTGTGACGGGCGGTGTGTACAAGGCCCGGGAA CGTATTCACCGCGGCATGCTGATCCGCGATTACTAGCGATTCCAGCTTC ACGCAGTCGAGTTGCAGACTGCGATCCGAACTGAGAACAGATTTGTGG GATTGGCTTAACCTCGCGGTCTTCGCTGCCCTTTGTTCTGTCCATTGTAGC ACGTGTGTAGCCCAGGTCATAAGGGGCATGATGATTTGACGTCATCCCC

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GTATCTAATCCTGTTCGCTCCCACGCTTTCGCTCCTCAGCGTCAGGTAC AGACAGAGAGTCGCCTTCGCCACTGGTCGTCTTCACATCTTAACGCATT TCACCGCACCAGTGAAATGCACTCCTCTCTTCGGCATCAAGTCCCAAAT CCATGAACCTGCCCGGTGATCGAGCTTCACATCAAACTAAGAACGCCT GCAGACCTTTAGCCAAAATTCGGAAAAGATTGGCCCTACGATACCGCG CTGCTGCACGATA Appendix (19) Sequence of PCR products of amplification of 16s rDNA by two sets of primers (27F and 1492R) of the *Bacillus* isolate B8.

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GGAGGAACACCAGTGGCGAAGGCGACTCTCTGGTCTGTAACTGACGCT GAGGAGCGAAAGCGTGGGGGGGGGCGAACAGGATTAGATACCCTGGTACTC CACGCCGTAAACGATGAGTGCTAAGTGTTAGGGGGGTTTCCGCCCCTCA GTGCTGCAGCTAACGCATTAAGCACTCCGCCTGGGGAGTACGGGCGCA AGACTGAACTCATAAGAATTTGACGGAGGGCCCGCACAAGCGGTGGAG CATGTGGTTTTATTTAAAGCAACGAGCAAGAAACTTACTAAGGTCTGG ACATCCTCTGACATTCCTAGAGATATGTACGGCCCCTTCACGGGCAGAG TGAACAGGGTAGAGGCAAGCTGTCGTCATCTCTCTGCCATGAAGATGA TGACTTGGCATCTGGAAGATATACTCGTCAGTGAAGCACGGAGTATTG GATTAGGCTGCACTCACATGACGTTGAACCTGCTACACGGGTCGCATAC GTCATGACTCAGGTA///GCGGCTGGCTCCTAAAAGGTTACCTCACCGAC TTCGGGTGTTACAAACTCTCGTGGTGTGACGGGCGGTGTGTACAAGGCC CGGGAACGTATTCACCGCGGCATGCTGATCCGCGATTACTAGCGATTCC AGCTTCACGCAGTCGAGTTGCAGACTGCGATCCGAACTGAGAACAGAT TTGTGGGATTGGCTTAACCTCGCGGTTTCGCTGCCCTTTGTTCTGTCCAT TGTAGCACGTGTGTAGCCCAGGTCATAAGGGGGCATGATGATTTGACGT CATCCCCACCTTCCTCCGGTTTGTCACCGGCAGTCACCTTAGAGTGCCC A A CTICA A TICOTICICA A CTIA A CATICA A COOTTICOCOTTICOCOTTICOCOCA CTI

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CAGTTACAGACCAGAGAGTCACCTTCGCCACTGGTGTTCCTCCACATCT CTACGCATTTCACCGCTACACGTGGAATTCCACTCTCCTCTTCTGCACTC AAGTTCCCCAGTTTCAATGACCCTCCCCGGTTGAGCGGGGGGCTTTCACA TCAGACTTAAGAAACCGCCTGCGAGCCCTTTACGCCCATATTCCGGACA CGATGCCACTACATATTACGCGGCTGCTGGCACGTAGTAGCGTGGCTTC AGGTAGGTACCGTCAGGACCGCCTATTCGACGGTACTGTCTCGCTACAT CGAGCTTGCGATCGAAACTTCATCACTCACGCGCGTTGCTCCGTCGGAC TTTCGTCATTGCGTATATCCTACTGCTGCTCCCGAGGAGCTTGAGCGGG

TCAAGCCGATGTGGCCTATCCACTGCTCAGGTCGCCACTGTCGTGCCTA GGAGGCGTTACCCGACACTAGTATGTCCCGAGTCCTGGGAAGTGAGCC CATACTCACTTTTATAGTGAACTGT

Appendix (20) Sequence of PCR products of amplification of 16s rDNA by two sets of primers (27F and 1492R) of the *Bacillus* isolate B10.

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GGATCGTAAAGCTCTGTTGTTAGGGAAGAACAAGTACCGTTCGAATAG

GGCGGTACCTTGACGGTACCTAACCAGAAAGCCACGGCTAACTACGTG CCAGCAGCCGCGGTAATACGTAGGTGGCAAGCGTTGTCCGGAATTATT GGGCGTAAAGGGCTCGCAGGCGGTTTCTTAAGTCTGATGTGAAAGCCC CCGGCTCAACCGGGGGAGGGTCATTGGAAACTGGGGGAACTTGAGTGCAG AAGAGGAGAGTGGAATTCCACGTGTAGCGGTGAAATGCGTAGAGATGT GGAGGAACACCAGTGGCGAAGGCGACTCTCTGGTCTGTAACTGACGCT GAAGAGCGAAAGCGTGGGGGAGCGTAACAGGACTAGATACCCTGGTAG TCCACGCCCTAAACAGATGAGAGCAATGTGATAGGAGGTTTCCGCCCC CTTAGTGCCGCAGCTAATACATTAATAACTCTGGCCCCGGGATTACGGT GCAAGAATGACACTTAAAAGGAATTGTCCGTCTGCCCCCAACAGTCAG CGGATCTTGTGTTTTCTTTAAAACAATTCGACACAACTTACCATCTGTCT ACATTCATCTGATAATT////CGGCGGCTGGCTCCTAAAAGGTTACCTCAC CGACTTCGGGTGTTACAAACTCTCGTGGTGTGACGGGCGGTGTGTACAA GGCCCGGGAACGTATTCACCGCGGCATGCTGATCCGCGATTACTAGCG ATTCCAGCTTCACGCAGTCGAGTTGCAGACTGCGATCCGAACTGAGAA CAGATTTGTGGGATTGGCTTAACCTCGCGGTTTCGCTGCCCTTTGTTCTG TCCATTGTAGCACGTGTGTAGCCCAGGTCATAAGGGGGCATGATGATTTG ACGTCATCCCCACCTTCCTCCGGTTTGTCACCGGCAGTCACCTTAGAGT

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#### ACTACCAGGGTATCTAATCCTGTTCGCTCCCCACGCTTTCGCTCCTCAG

CGTCAGTTACAGACCAGAGAGTCGCCTTCGCCACTGGTGTTCCTCCACA TCTCTACGCATTTCACCGCTACACGTGGAATTCCACTCTCCTCTTCTGCA CTCAAGTTCCCCAGTTTCCAATGACCTCCCCGGTTGAGCCGGGGGGCTTT CACATCAGACTTAAGAAACCGCCTGCGAGCCCTTTACGCCCAATAATCC TGACAACGCTTGCCACCTACGAATACCGCGGGCTGCTGGTACGTAGTAG ACGTGGCTTTTCTGGTTAGGTACGTCTAGGTACCGCCCTATCGAACGGA ATTGTCTTCCTAACACCAGAGTTTTCGATCCAAAACCTCATCACTCCGC GGCGTTCTTCGTCCAGACTTCTTCATGGGGAAAATCCTACTGCTGCTCT

## CTCAGTGCTCGGGGACGGGGGGGATATTCCCGGGGGGCGTATACCTCTCAGG CGGCTCCCT

Appendix (21) Mass spectrum obtained from the MS1 analysis of peak retained at 8.4min for standard surfactin after HPLC analysis.





Appendix (23) Mass spectrum obtained from the MS1 analysis of peak retained at 9.9 min for standard surfactin after HPLC analysis.



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Appendix (25) Mass spectrum obtained from the MS1 analysis of peak retained at 13 min for standard surfactin after HPLC analysis.



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Appendix (27) Mass spectrum obtained from the MS1 analysis of peak retained at 16.9 min for standard surfactin after HPLC analysis.



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Appendix (29) Mass spectrum obtained from the MS1 analysis of peak retained at 22.4 min for standard surfactin after HPLC analysis.



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**Remove Watermark Now** 1072.9 679.6 1 701.6 815.8 758.5 914.5 700 800 1000 900 1100 m/z Appendix (31) Mass spectrum of *B*.B6 lipopeptide obtained from the MS2 analysis of the peak of  $[M+Na]^+$  at m/z 1017.



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Appendix (33) Mass spectrum of *B*.B6 lipopeptide obtained from the MS2 analysis of the peak of  $[M+Na]^+$  at m/z 1045.



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Appendix (35) Effect of *B*. B1 lipopeptide on K562 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (37) Effect of *B*. B1 lipopeptide on Hep-2 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (39) Effect of *B*. B3 lipopeptide on Hep-2 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (41) Effect of *B*. B5 lipopeptide on K562 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (43) Effect of *B*. B5 lipopeptide on Hep-2 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (45) Effect of *B*. B6 lipopeptide on L1210 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (47) Effect of *B*. B6 lipopeptide on Lo2 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (49) Effect of *B*. B7 lipopeptide on L1210 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (51) Effect of *B*. B8 lipopeptide on K562 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (53) Effect of *B*. B8 lipopeptide on Hep-2 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (55) Effect of *B*. B9 lipopeptide on L1210 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (57) Effect of *B*. B10 lipopeptide on Hep-2 cell proliferation for different incubation times (24,48,72)hrs.



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Appendix (59) Effect of *B*. B10 lipopeptide on L1210 cell proliferation for different incubation times (24,48,72)hrs.



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Biological Activity of Biosurfactants Produced by Locally Isolated *Bacillus* spp. College of Science/Al –Nahrain University By Sura Ali Ibrahim PhD \2012

Biological Activity of Biosurfactants Produced by Locally Isolated *Bacillus* spp. College of Science/Al –Nahrain University By Sura Ali Ibrahim PhD \2012

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# Conclusions

1- Ten locally *Bacillus* isolates were capable of producing biosurfactant with high efficiency that can lower (ST) of the cell free supernatant (40-43)%.

2- Molecular identification of these isolates showed that they belong to *Bacillus subtilis* group with similarity (97-100)%.

3- According to TLC and HPLC/MS results, biosurfactant produced by *Bacillus* isolates were belonged to the lipopeptide family, and that produced by *Bacillus* B6 was surfactin with partial sequence of Val-Asp -Leu-Leu-OH<sub>2</sub>.

4- Purified lipopeptides produced by Bacillus isolates (B1, B3, B5, B6, B7, B8,

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58.62% inhibition of the cell line growth at the same concentration, however,

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purified biosurfactans produced by (*B. subtilis* isolates B6, B8, B9) were the most cytotoxic on Hep 2 proliferation with inhibition rate (69.219,86.043,64.6)% at concentration 80mg/l, while purified biosurfactat produced by B.B6 had low effect on the proliferation of normal cell line L02.

<sup>7</sup>- Purified surfactin produced by B. subtilis B6 induces apoptosis in Hep-2 cell line through mitochondrial depolarization and caspase -3 activation.

7- Optimum conditions for surfactin production by *Bacillus* B6 isolate were achieved in a medium containing (1%) sucrose, (0,3%) potassium nitrate (1g/l), at pH 8, 30°C, 180rpm of shaking for 72 hrs. The biosurfactant yield was 1.4 g/l of culture medium under the optimum conditions.

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# Dedication

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Sura

#### Acknowledgement

First of all Praise to Allah , Lord of the Whole Creation . Mercy and Peace are to the Prophet Mohammed and his Family.

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Hatim Al- Rubaiee , Dr. Rami, Dr. Hussam, Hiam, Kawther, Raid.

I'm grateful to my friend Dr. Ghassan for his continuous advising through my work, and to all my colleagues, Dr. Khalel, Dr. Raghad, Dr. Asmaa, Dr. Rana for their friendship and encouragement.

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Sura

Republic of Iraq Ministry of Higher Education and scientific Research Al- Nahrain University, College of Science, Department of Biotechnology



# Biological Activity of Biosurfactants Produced by Locally Isolated *Racillus* spn

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#### **1.1 Introduction**

Biosurfactants are surface – active compounds produced by a wide variety of microorganisms such as bacteria, yeasts and fungi as membrane components or secondary metabolites (Gautam and Tyagi, 2006). They are a structurally diverse group of amphipathic molecules with both hydrophilic and hydrophobic moieties (Van Hamme *et al.*,2006).

The major classes of biosurfactants include glycolipids, lipopeptides and lipoproteins, phospholipids and fatty acids, polymeric surfactants and particulate surfactants (Cameotra and Makkar, 2004; Salihu *et al.*, 2009).

In recent years, the interest in biosurfactants has been remarkably increasing due to many advantages compared with chemical surfactants, including lower This is a watermark for the trial version, register to get the full one!

activity at extreme conditions (temperature, pH, salinity) (The

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personal care products, food processing, textile manufacturing, laundry supplies

metal treatment and processing, pulp and paper processing and paint industries (Singh *et al.*,2007; Rahman and Gakpe, 2008).

Biosurfactants have various interesting biological activities such as antiviral, antitumor, antibacterial and antifungal activities, also they can be used as immunoregulators, adhesive agents, ligands for binding immunoglibulins, adjuvants for antigen ..etc (Banat *et al.*, 2010 ; Fathabad, 2011).

Among the many classes of biosurfactants, lipopeptides produced by *Bacillus* spp. strains are the most powerful ones that possess many biological activities especially the antitumor activity that include the inhibition of the

carcinoma cells proliferation and induction of (differentiation, cytotoxicity, cell cycle arrest, and apoptosis) (Wang *et al.*, 2007; Cao *et al.*,2009a).

In Iraq, cancer is a growing problem that increases considerably every year, one of the treatment regiments is the chemotherapy which is hampered by the problem of drug-resistance, so a call for discovery of more effective agents to treat cancer is becoming increasingly urgent, for this purpose, new drugs have being synthesized and tested (WHO, 2006).

New trials for cancer treatment have been performed by many researchers in various countries including Iraq; these trials included using gene therapy, immunotherapy, biological therapy and bacterial byproducts (Mulherkar, 2001;Al-

Qadoori, 2004 ; Al-Saffar, 2010; Visagie and Joubert , 2010;). However, using

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Ance this study aimed to achieve the following targets:

• Isolation of *Bacillus spp*. from different localities contaminated with hydrocarbons.

- Selection and identification of efficient isolates of *Bacillus* spp. producing biosurfactant .
- Purification of biosurfactant to homogeneity level.
- Application of the biosurfactants as antitumor agents

#### **1.2 Literature review**

#### 1.2.1 Genus Bacillus

#### **1.2.1.1 General features**

It is a large Genus of Gram-positive rods that were first recognized and named by Ferdinand Cohn in 1872, they may occur singly or in chain, some are strict aerobes and others are facultative anaerobes. The vegetative bacilli are large  $(0.5 \times 1.2 \ \mu m$  to  $2.5 \times 10 \ \mu m)$  and straight, they can form highly resistant dormant endospores in response to nutrient deprivation and other environmental stresses.

The majority of vegetative cells are mesophiles with temperature optimum 30 °C or 45 °C, but the genus contains also a number of thermophilic species with This is a watermark for the trial version, register to get the full one!

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remperature of incubation, humidity,etc.), Some *Bacillus* species tend to swarm on solid media, especially if plates are not dried to remove the surface moisture before inoculation (Duguid ,1996 ; Rudner *et al.*,1998).

Endospores in pure culture are regular in size, shape and position within the sporangium, in most species they are oval (ellipsoidal) in shape, and central or subterminal in position. In a few species the spores are spherical in shape and in a few they are terminal in position, when they viewed under microscope unstained they appear edged in black and very bright and refractile (Holt *et al.*, 1994; Duguid, 1996; Piggot and Hilbert, 2004).

Endospores formation is affected by some factors including the temperature of growth, the pH ,aeration, presence of minerals , presence of certain carbon or nitrogen compounds and the concentration of the carbon or nitrogen source, in some circumstances a starvation for phosphorus source ,population density, cell cycle (Errington,1993; Piggot and Hilbert,2004; Goesselsberger *et al.*,2009).

Endospores are highly resistant to environmental stress such as high temperature, low temperature, irradiation, presence of strong acids, disinfectants, nisin, , high sodium chloride concentration etc. Mature spores have no detectable metabolism , a state that is described as cryptobiotic, they retain viability indefinitely such that under appropriate environmental conditions, they germinate

into vegetative cells (Nicholson, 2002 ; Faille et al., 2002; Cartman et al., 2008)

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hith *et al.*, (1952) which based on the shape of the endospore and its position in the mother cell or sporangium (Priest *et al.*, 1988; Priest, 2008).

This taxonomic was accepted until the introduction of modern taxonomic techniques such as numerical phonetics, DNA base composition determination and DNA reassosiation experiments, 16S rDNA sequence analysis, which found that *Bacillus* comprise more heterogenous species than scientists expected as shown in the GC content variation of known species of *Bacillus* which range from 32%-69% as listed in Bergeys Manual of Systemic Bacteriology (1986) by Claus and Berkeley. This indicates considerable genetic diversity among species and suggests that the genus should perhaps be split into several, more homogeneous taxa, so in the 1990s the genus *Bacillus* comprises in excess of 60 species as listed by

Konemann *et al.*(1997), but this taxonomy is not stable as sequence analysis has led and continues to lead, on the one hand, to the separation of groups of species from the core genus *Bacillus* to form new genera and, on the other hand, to the definition of novel genera to allocate new isolate.

Recently, according to Garrity *et al.* (2004), *Bacillus* is only one out of a whole series of genera of aerobic endospore-forming bacteria( AEFB) in which it can be classified as follows:

#### Phylum BXIII. Firmicutes phy. nov.

Class III. Bacilli

**Order I.** *Bacillales* 

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upon morphological, biochemical, physiological, and chemotaxonomical traits and on the results of various molecular genetic techniques.

#### 1.2.1.3 Natural products of *Bacillus* spp.

Members of the *Bacillus* genus are often considered as microbial factories for the production of a vast array of biologically active molecules with the potential for technical and scientific applications, on the other hand, several strains of *Bacillus* species also produce compounds toxic to mammalian cells (Lugan ,1988; Outtrup and Jörgensen 2002). For example *Bacillus* genus produce exocellular enzymes (e.g. amylases and proteases ) that form over one-half of the total commercial enzyme volume (Crueger and Crueger ,1982; Fogarty and Kelly ,1990), also some *Bacillus* spp produce different kinds of surface active compounds which known as Biosurfactants (e.g. surfactin, iturins, lichenysin A) that have a wide range of applications , also many strains of *Bacillus* produce several bacteriocin-like substances, e.g. some strains of *B. cereus* produce the bacteriocin cerein 7, a poreforming peptide of 3940 Da (Oscariz *et al.* 1999, Oscariz and Pisabarro 2000).

*Bacillus* spp. can also produce a vast array of antibiotics that active against important pathogens such as methicillin-resistant *Staphylococcus aureus* (Chatterjee *et al.* 2005).

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#### **1.2.2 Surfactants**

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Leir amphiphlilic composition of two functional moieties: polar with the main

hydrophilic group and non-polar with lipophilic trait that tend to partition preferentially at the interface between fluid phases with different degrees of polarity and hydrogen bonding such as oil / water or air / water interfaces (Desai and Banat, 1997; Rahman and gakpe,2008). The formation of such an ordered molecular film at the interface lowers the interfacial energy (interfacial tension) and surface tension which is responsible for the unique properties of surfactant molecules (Lin, 1996).

Surfactants are defined by their capability of reducing the surface tension (ST), critical micelle concentration (CCM) (which is the minimum surfactant

concentration required for reaching the lowest interfacial or surface tension values) , interfacial tension and hydrophilic-lipophilic balance (HLB) , (surfactants with HLB values less than 6 are more soluble in the oil phase ; while those with HLB values between 10 and 18 have the opposite characteristics) (Cooper and Zajic, 1980 ; Parkinson, 1985).

The hydrophobic part of the surfactant is a long – chain of fatty acids , hydroxy fatty acids , hydroxyl fatty acid or  $\alpha$ - alkyl – $\beta$ -hydroxy fatty acids which is usually a C8 to C22 alkyl chain or alkylaryl that may be linear or branched while the water soluble end (hydrophilic) can be a carbohydrate, amino acids, cyclic peptide, phosphate, carboxylic acid or alcohol (Oberbremer *et al.*, 1990 ; lin *et al.*, 1994a).

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**.2.1 Petroleum – based chemical surfactants** 

Surfactants under this category are synthesized from petroleum refining byproducts that are found in light cuts (gasoline and kerosene) coming from atmospheric distillation and catalytic cracking or by polymerization of short chain olefin, particularly in C3 and C4 (Salager, 2002).

They can be classified into anionic, cationic, amphoteric, and nonionic classes depending on the nature of their polar grouping, regarding the hydrophobic moiety of the molecule, it is a hydrocarbon chain in most common surfactants; however, in some more specialized surfactants, this hydrophobic part can be a nonhydrocarbon chain such as a polydimethylsiloxane or a perfluorocarbon (Van Ginkel, 1989).

**Anionic surfactants** are negatively charged which is usually due to a sulphonate ,sulphur , carboxylates , or phosphates group. They are the most commonly used surfactants that accounting for about 50 % of the world production. (Int .2)

A general formula may be ascribed to anionic surfactants as follows:

- \* Carboxylates:  $C_nH_{2n+1}COO^-X$
- \* Sulphates:  $C_nH_{2n+1}OSO_3^-X$ , (Figure 2-1)
- \* Sulphonates:  $C_nH_{2n+1}SO_3^-X$
- \* Phosphates:  $C_n H_{2n+1} OPO(OH)O^-X$

With n = 8-16 atoms and the counter ion X is usually Na<sup>+</sup>.

Several other anionic surfactants are commercially available such as alphosuccinates, isethionates and taurates and these are sometimes used for

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Figure (1-1) Alkyl ether sulphate as an example of anionic surfactant (Tadros, 2005)

**Nonionic surfactants** that lack ionic constituent, they come as a close second with about 45% of the overall industrial production. They do not ionize in aqueous solution, because their hydrophilic group is of a non dissociable type, such as alcohol, phenol, ether, ester, or amide (Sarney and Vulfson, 1995).

There are two major classes of anionic surfactants:

- \* Ethoxylated surfactants such as fatty acid ethoxylates with the general formula RCOO-(CH<sub>2</sub>CH<sub>2</sub>O)*n*H.
- \* Multihydroxy products such as glycol esters, glycerol (and polyglycerol) esters(Tadros, 2005).



#### Figure (1-2) Non ionic surfactants (Sarney and Vulfson, 1995)

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#### oups. (Tadros , 2005)

These surfactants are in general more expensive than anionics, because of the high pressure hydrogenation reaction to be carried out during their synthesis. (Salager, 2002)



Figure (1-3) Cationic surfactants (Salager, 2002)

• Amphoteric surfactants that have both positively and negatively charged moieties in the same molecule such as Alkyl betaines, alkyldimethylamines, imidazonilinum derivatives (Salager, 2002).



Figure (1-4) Amphoteric surfactants (Salager, 2002)

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Typical examples of surfactants derived from m

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constituent of animals and plants, Lecithin is usually produced from egg yolk and

consists of zwitterionic phosphatidylethanolamine and phosphatidylcholine, its mainly used as a pharmaceutical recipient for drug delivery and intravenous nutrition (Wang and Wang, 2008).

Gelatin is a high – molecular – eight polymer that obtained through the partial hydrolysis of collagen with dilute acid or base, it's a relatively poor protein surfactant, but its emulsifying properties can be improved by enzyme – catalyzed attachment of hydrophobic side chains (Dickinson, 1993). The main sources of commercial gelatin are bovine skin and bones and pigskin. It is mainly used as a stabilizer, thickener and texturizer in food and non – food applications (Karim and Bhat, 2008).



Figure 1-5 Chemical composition of lecithin (Wang and Wang, 2008)

#### 1.2.2.3 Plant – derived surfactants

Many surface-active compounds are derived from renewable plant resources.

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detary saponins are the legumes: soybeans, chickpeas, mung beans, peanuts, broad

beans, kidney beans and lentils; the saponin content in soybeans is 5–6% (Oakenfull, 1981).

Another example are the plants surface-active proteins. Soy protein is one of the most important plant-derived protein surfactants. Soybeans contain about 40% protein and 20% oil. Soy proteins are mainly globulins and can be classified into 2S, 7S, 11S, and 15S fractions. Soy proteins are available in three major forms that vary in protein content: soy flours, soy protein concentrates and soy protein isolates (Iwabuchi and Yamauchi, 1987).

#### **1.2.2.4 Biosurfactants**

Biosurfactants are surface active metabolites produced by microorganisms (Bacteria, yeasts and fungi) when grown on water miscible or oily substrates or both (Arino *et al.*, 1996; Kiran *et al.*, 2009; Rufino *et al.*, 2011). They either remain adherent to microbial cell surfaces or are secreted in the culture broth (Olivera *et al.*, 2009; Fathabad,2011). Their molecular masses generally range from 500 to 1500 Da.(Van Hamme *et al.*,2006).

Although, most biosurfactants are considered to be secondary metabolites, some may play essential roles for the survival of biosurfactant – producing microorganisms through facilitating nutrients uptake or microbial – host

interactions or by acting as biocide agents or promoting the swarming motility of

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differentiation (Ahimou *et al.*, 2000; Kearns and Losick, 2

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#### Imposition and microbial origin. For example Biermann et al. (1987) group

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biosurfactans based on their composition into glycolipids, lipopeptides , phospholipids, fatty acids, neutral lipids , polymeric and particulate compound. While Kosaric (1992) classify them depending on their structure namely; hydroxylated and cross- linked fatty acids, polysaccharide – lipid complexes, glycolipids, lipoproteins – lipopeptides, phospholipids and complete cell surfaces while Healy *et al.*, (1996) group biosurfactants into four main categories namely, glycolipids, phospholipids, lipoproteins / lipopeptides and polymeric. Lastly, Rosenberg and Ron (1999) suggested that biosurfactants can be divided into two categories: low – molecular – mass molecules, which efficiently lower surface and interfacial tension, and high – molecular – mass polymers, which are more effective as emulsion – stabilizing agents. The major classes of low – mass surfactants include glycolipids, lipopeptides and phospholipids, whereas high – mass surfactants include polymeric and particulate surfactants.

#### **1.2.2.4.2** Types of biosurfactants

There are many types of biosurfactants that produced by various microorganisms, the following are some of the various types of biosurfactants.

#### 1.2.2.4.2.1 Glycolipid biosurfactants

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The most known biosurfactants are glycolipids, they are carbohydrates in combination with fatty acids, the linkage is by means of either ether or an ester

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Chalolipids and sophorolipids (Desai and Banat, 1997; Morita et al., 2006)

#### • Rhamnolipids

They are generally a mixture of homologous species of glycolipids produced by the genus *Pseudomonas*, they composed of one or two molecules of rhamnose linked to one or two molecules of  $\beta$ -hydroxydecanoic acid usually, but other fatty acids may be found depending on the *Pseudomonas* species or growth conditions (Figure 1.6)(Desai and Banat, 1997), they have the ability to lower the interfacial tension against *n* – hexadecane to mN/m and the surface tension to 25 -30 mN/m (Guerra –Santos *et al.*, 1986). Also two unusual rhamnolipids, designated myxotyrosides A and B, have been isolated from a *Myxococcus sp*, they have a rhamnose unit linked to tyrosine and hence to a fatty acid (Ohlendorf *et al.*, 2009).



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#### group, such as Mycobacterium, Rhodococcus ,Corynebacterium, Arthrobacter,

*Nocardia* and *Gordonia*, different structures have been elucidated particularly in *Rhodococcus* genus(Franzetti *et al.*, 2010). They are composed of Trehalose (is a non-reducing disaccharide in which the two glucose units which linked together linked either to Mycolic acids in the *Mycobacterium* and most species of *Crynebacterium* and *Nocardia* (Silva *et al.*, 1979 ;Gautam and Tyagi, 2006) or to corynomycolic or nocardomycolic in the case of rest species of *Corynebacterium* and *Nocardia* (Goodfellow *et al.*, 1973; Shimakata and Minatogawa ,2000).





• Sophorolipids

They are group of biosurfactants produced by some yeast species, and in

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(sophorose) and a long chain hydroxyl fatty ocid linked on allyois the body
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SLs forms a macrocyclic lactone ring with the 4"- hydroxyl group of the sophorose by intramolecular esterification. (Figure 1.8)



Figure 1.8 Types of Sophorolipid biosurfactants (Hu and Ju ,2001)

#### **1.2.2.4.2.2 Fatty acids biosurfactants**

Biosurfactants under this category are produced from alkane as a result of microbial oxidations (Rehn and Reiff, 1981). These fatty acids are either straight chain acids, or complex fatty acids containing OH groups and alkyl branches such as Corynomucolic acids (Kretschner *et al.*, 1982) but the most active saturated fatty acids in lowering surface and interfacial tensions are in the range of C12-C14 because the hydrophilic or lipophilic balance of fatty acids is related to the length of the hydrocarbon chain (Rosenberg and Ron, 1999).

#### 1.2.2.4.2.3 Phospholipids

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#### Emulsan produced by Acinetobacter calcoaceticus was the first studied

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polymeric biosurfactant (Rosenberg *et al.*, 1979), its polyanionic amphipathic heteropolysaccharide bioemulsifier as mentioned in (Figure 1.9), It's a very effective emulsifying agent for hydrocarbons in water even at a concentration as low as 0.001-0.01% (Zosim *et al.*, 1982). Another example of this type of biosurfactants is liposan which was synthesized by using *Candida lipolytica*, its composed of 83% carbohydrate and 17% protein with the carbohydrate portion being a heteropolysaccharides consisting of glucose, galactose, galactosamine and galactoronic acid (Ciriglian and Carman, 1984).



Figure 1-9 Emulsan (Zosim *et al.*, 1982)

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uphylococcus aureus, Layer A in Aeromonas salmonicida (Desai and Banat,

1997).

#### 1.2.2.4.2.6 Lipopeptide biosurfactants

Biosurfactants of this type possess remarkable surface- active properties and produced by a wide variety of microorganisms such as *Agrobacterium tumefaciens* that produced lysine- containing lipids (Tahara *et al.*, 1976), *Pseudomonas rubescens* produced Ornithine- containing lipids (Yamane,1987), but the majority of these biosurfactants are produced from several species of the genus *Bacillus* that can be classified into three families:(Ongena and Jacques, 2007).

• Lipopeptides of the surfactin family

- Lipopeptides of the iturin family
- Fengycins and various lipopeptides

*Bacillus* lipopeptides consist of a peptide part containing 7-11 amino acids, either cyclic or linear or a combination of these while the lipid part is composed of  $\beta$ -hydroxy or  $\beta$ -amino fatty acids connected to the peptide backbone. The hydrocarbon length of the fatty acids and amino acid composition may vary, depending on the nutrition of the bacteria, and affect the properties of the lipopeptides. The cyclic structure of the peptide part protects the lipopeptide from enzymatic cleavage and maintains its general stability (Desai and Banat, 1997; Peypoux *et al.*, 1999; Muthusamy *et al.*, 2008).

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Lipopeptides of the surfactin family (surfactin, little vsu

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akimov et al., 1996). These peptides are powerful biosurfactants, produced by

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strains of several *Bacillus* species such as *Bacillus subtilis* (Arima *et al.*, 1968), *Bacillus coagulans* (Huszcza and Burczyk, 2006), *Bacillus pumilus* and *Bacillus licheniformis* (Naruse *et al.*, 1990; Thaniyavarn *et al.*, 2003).

Surfactin produced by *Bacillus subtilis* ATCC 21332 is one of the most powerful biosurfactants was discovered by Arima *et al.* (1968), it was named surfactin because of its exceptional surfactant activity, its structure was elucidated as that of a macrolide lipopeptide by Kakinuma *et al.*, (1969) ,Surfactin contains a heptapeptide (Glu1-Leu2-Leu3-Val4-Asp5-Leu6-Leu7), it lowers the surface tension from 72 to 27.9 mN/m at concentration as low as 0.005% (Arima *et al.*, 1968).

The natural analogue is lichenysin A has Gln in position 1 and Ileu at position 7, It is produced by *B. licheniformis* during anaerobic and aerobic growth, it was first isolated from oil wells and its structure was elucidated by Yakimov *et al.* (1995) and Yakimov *et al.* (1999), another analog is pumilacidin which has Leu at position 4, whereas other variations are located at position 7.

*Bacillus coagulans* has been found to produce several surfactins, four main components with molecular weights 1007, 1021 and 1035 Da were separated.

Their structures have been confirmed by spectrometric and spectroscopic studies and by acid hydrolysis. The compounds were found to represent two pairs of surfactin isoforms in which beta-hydroxy-iso-C14 or anteiso-C15 fatty acids are

linked to the [Leu7] or [Val7] heptapeptide molety by both an amide group and a This is a watermark for the trial version, register to get the full one!

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an contain a guelic hortenantida, aculated with 8 amino fatty

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acids with a chain length of C14 to C16 (Peypoux *et al.* 1978, Hourdou *et al.* 1989). Iturin A contains the heptapeptide Asn1-Tyr2-Asn3-Gln4-Pro5-Asn6-Ser7, whereas in the other members the amino acid residues in the heptapeptides vary slightly; e.g. mycosubtilin has Asn1-Tyr2-Asn3-Gln4-Pro5-Ser6-Asn7 (Yu *et al.* 2002).

#### • Fengycins and various lipopeptides

Fengycins (Figure 1.12) and the structurally similar plipastatins are distinguished from the other lipopeptides by the way in which the  $\beta$ -hydroxy fatty acid is linked with the polar dipeptide which is associated with the cyclic

octapeptide. The cyclic structure of fengycin  $cyclo(D-allo-Thr1-L-Glu2-D-Ala3/D-Val3-L-Pro4-LGln5-L-Tyr6-L-Ile7-D-Tyr8)D-Orn9-L-Glu10-<math>\beta$ -OH-FA ) is formed when *D*-Tyr8 is connected with amide bonds to *D-allo*-Thr1 and *D*-Orn9 and forms a lactone bond with *L*Ile7. The structure of fengycin A contains *D*-Ala3 instead of the *D*-Val3 of fengycin B (Volpon *et al.*,2000).



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Variants		Lenght and branching of the acyl chain
Esperin**	L-Glu-L-Leu-D-Leu-L-Val-L-Asp-D-Leu-L-Leu-COOH	
Lichenysin*** Pumilacidin	L-XL <sub>1</sub> -L-XL <sub>2</sub> -D-Leu-L-XL <sub>4</sub> -L-Asp-D-Leu-L-XL <sub>7</sub> L-Glu-L-Leu-D-Leu-L-Leu-L-Asp-D-Leu-L-XP-	$i$ - $C_{13}$ , $ai$ - $C_{13}$ , $n$ - $C_{14}$ , $i$ - $C_{15}$ , $ai$ - $C_{15}$
Surfactin	L-Glu-L-XS <sub>2</sub> -D-Leu-L-XS <sub>4</sub> -L-Asp-D-Leu-L-XS <sub>7</sub>	i-C <sub>14</sub> , $n$ -C <sub>14</sub> , $i$ -C <sub>15</sub> , $ai$ -C <sub>15</sub>
** the β-carboxy *** or halobacill	l of $Asp_5$ is engaged in the lactone	n, linear
$XL_1 = Gln \text{ or } Glu \text{ ; } XL_2 = Leu \text{ or } Ile \text{ ; } XL_4 \text{ and } XL_7 = Val \text{ or } Ile \text{ ; }$		ai anteiso

Figure (1.10) Surfactin family (Ongena and Jacques, 2007)



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Figure (1.11) Iturin Family (Ongena and Jacques, 2007)

Fengycins have stereoisomeric composition different from those of plipastatin. Fengycins contain *D*-Tyr8 instead of the *L*-Tyr8 of plipastatins and *L*-Tyr6 instead of the *D*-Tyr of plipastatins (Volpon *et al.* 2000). While Hathout *et al.* (2000) reported from *Bacillus thuringiensis kurstaki* HD-1 an antifungal compound structurally resembling plipastatin and fengycins.

Analogues of iturin and fengycin were reported with a double bond in the fatty acid part such as in *B. thuringiensis* strain CMB26 that produced an analogue of fengycin with a double bond in the fatty acid. It was fungicidal, bactericidal, and

insecticidal, and more effective against fungi than was iturin or surfactin (Kim *et al.* 2004). The fatty acids in iturins were predominantly C16 and C17 (Vater *et al.* 2002, Deleu *et al.* 2005). Recently *B. subtilis* strain GA1 was described as producing three lipopeptides: surfactin, fengycin, and iturins (Toure *et al.* 2004).

Another group of cyclic hexapeptide lipopeptides contains six amino acids acylated with 15-guanidino-n-hydroxypentadecanoic acid (bacillopeptins, fusaridins and LI-F. The fusaricidins (Kajimura and Kaneda 1997, Beatty and Jensen, 2002) and LI-F (Kurusu *et al.* 1987, Kuroda *et al.* 2000) from *Paenibacillus polymyxa* were fungicidal and antibacterial. LI-F compounds containing an azole group were toxic to ddY mice (Kuroda *et al.* 2001). The target

of the azole group containing fungicidals target is cytochrome P450 which is also

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when administered orally (Mclean *et al.* 2002) and impair micoche idri

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Several methods for screening and estimating of biosurfactants have been

developed. These methods were as follows:

• Drop-collapse method is one of the qualitative methods used to determine the presence of biosurfactant. Tugrul and Cansunar (2005) conducted experiments to confirm the reliability of the method using polystyrene microwell plate; oilcoated wells collapse was observed when the culture broth contained biosurfactant and there was no change in the shape of the droplets in the absence of biosurfactant.



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Figure (1.12) Fengycins family (Ongena and Jacques, 2007)

• Thin layer chromatography (TLC) is also used in preliminary characterization of the biosurfactant where the cell free extract containing biosurfactant is separated on a silica gel plate using chloroform: methanol: water (70:10:0.5, v/v/v); this is then followed by using color developing reagents. Lipopeptide biosurfactant showed red spots in the presence of ninhydrin reagent, while glycolipid biosurfactant is detected as yellow spots when anthrone is used as the color reagent (Yin *et al.*, 2009).
• Additionally, blood agar hemolysis tests is another method used; where the organisms with biosurfactant ability are streaked on blood agar plates and incubated at 40 °C. The plates are visually monitored for the presence of clearing zone around the colonies which is indicative of surfactant biosynthesis. The diameter of the clear zones depends on the concentration of the biosurfactant produced (Youssef *et al.*, 2004; Ghojavand *et al.*, 2008).

• Surface tension measurement by a du Nöuy ring-type tensiometer is one of the simplest techniques used. The surface tension measurement is carried out at room temperature after dipping the platinum ring in the solution for a while in order to attain equilibrium conditions. A higher biosurfactant concentration in the

test sample provides a lower surface tension until the critical micelle concentration

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despite the changes in concentration (Desai and Banate 1

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(UV, MS, evaporative light scattering detection, ELSD) for identification and quantification of biosurfactants (Heyd *et al.*, 2008).

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#### 1.2.2.4.4 Fermentation and production of surfactin

Most nutritional and production studies were done with batch cultures, usually in flasks with shaking , but occasionally in small-scale or large-scale fermentors (Cooper *et al.*1981; Mulligan and Gibbs 1990; Lin *et al.* 1994b).

A laboratory-scale cyclone column fermentor was used for continuous, phased growth with feedback control, based on the concentration of dissolved oxygen (Sheppard and Cooper, 1990). As the earlier studies carried out in nutrient broth gave a very low yield (0.1 g/ l)(Arima et al. 1968), in subsequent studies a minimal mineral salts medium, containing NH<sub>4</sub>NO<sub>3</sub> (0.05 M) as the inorganic nitrogen source and glucose (4%) as the carbon source, was defined by Cooper (Cooper's medium).Studies of the mineral requirement clearly established the need for and the stimulatory effect of iron and manganese (Cooper *et al.* 1981).

For continuous operation, a critical nitrogen/iron/manganese molar ratio of 920:7.7:1.0 was determined and was found to sustain surfactin production for at least 36 generations (Sheppard and Cooper 1991). Other work has been done in the semisynthetic medium of Landy *et al.* (1948), which contains L-glutamic acid (5 g/l) as the organic nitrogen source, glucose (2%) as the carbon source and a

trace of metal cations (Landy's medium) (Nakano *et al.* 1988; Sandrin *et al.* 1990)

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sources while the presence of glycerol greatly decrea

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#### ferences in the ability of different amino acids to support surfactin production

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has been found, yields being( $0.6\pm0.8$  g /l) (Sandrin *et al.* 1990). When they are used as the sole nitrogen source of a culture medium, some hydrophobic amino acids insert themselves directly into selected positions of the peptide sequence, thus amplifying the original structural microheterogeneity via the production of variants (Peypoux and Michel 1992).

Competition with cellular growth is probably one of the reasons for the rather disappointing product yields. In fact, in most of the studies the levels of increase are marginal and are probably due to differences in the process conditions, i.e.physicochemical and engineering parameters or nutritional factors; conversely, the replacement of Cooper's nitrogen source and the introduction of O2 limitation,

which redirects the energy efflux into product synthesis, have led to a productivity of(7 g /l), about 10 times as high as Cooper's basal yield (Kim *et al.* 1997).

*B. subtilis* was cultivated under aerobic conditions and at temperatures from  $30 \ ^{\circ}C$  to  $37 \ ^{\circ}C$ .

de Roubin *et al.* (1989) study the correlation between the primary metabolism of the cell and the secondary metabolism using UV mutagenesis, it was possible to isolate an overproducing strain of *B. subtilis* ATCC 21332, the mutant ATCC 51338. This mutant lowers the isocitrate dehydrogenase (a Krebs cycle enzyme) activity to 30 times less than that of the parent and produces 4 times more surfactin, i.e. (1.1 g/ l). The decrease of enzyme activity can also be achieved

by O2 limitation or by addition of citric acid to the production medium, and they

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increased. These observations suggest that the overall ra

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#### **1.2.2.4.5 Genes for surfactin synthesis**

The organization of the biosynthetic gene cluster of surfactin was published in the early 1990s by different researcher groups (Cosmina *et al.*, 1993; Fuma *et al.*, 1993; Nakano *et al.*, 1988).

It composed of a large operon of 25 kb, named srfA ,(Figure 2-13), which is also responsible for sporulation and competence development (Nakano *et al.* 1991; Hamoen *et al.* 1994). (Figure 1.13)

It contains four modular open reading frames, ORF1 (srfA-A), ORF2 (srfA-B), ORF3 (srfA-C) and ORF4 (srfA-D), the first three of them encoding the three

respective enzymes E1A, E1B and E2 needed to make the heptapeptide sequence (Vollenbroich *et al.* 1994).

The two nucleotide regions for the L-leucine epimerases are located at the 3<sup>°</sup> portion of the srfA-A and srfA-B genes. The two coded polypeptides show sequence similarities to other putative bacterial racemases (Marahiel *et al.* 1997).

At the end of the srfA-C gene, one region (TE, thioesterase) codes for an enzyme homologous to fatty acid thioesterases type I, The terminal gene srfD encodes a protein with high homology to external thioesterases of type II, It has been reported that SrfD is a repair enzyme, which regenerates with acetyl-CoA or with incorrect amino acids mischarged peptidyl carrier protein (PCP) domains

during non ribosomal peptide synthetases assembly (Schwarzer et al., 2002; Yel

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enzyme, termed sfp, belongs to the superfamily of 4`PPTases that function as primers of the non-ribosomal peptide and siderophore synthesis via a posttranslational phosphopantetheinylation of thiotemplates (Walsh *et al.* 1997).

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#### 1.2.2.4.6 Synthesis of Surfactin

Surfactin as one of the lipopeptide is synthesized by non ribosomal peptide synthetases (NRPSs), they are megaenzymes organized in iterative functional units called modules that catalyze the different reactions leading to peptide transformation (Stein, 2005).



Figure (1-13) Surfactin operon (Ongena and Jacques, 2007)

Each module is subdivided into several catalytic domains responsible for each biochemical reaction (Sieber *and* Marahiel , 2003; Walsh, 2004).

The NRPS is responsible for one reaction cycle of selective substrate recognition and activation as an adenylate (A-domain), tethering of a covalent This is a watermark for the trial version, register to get the full one!

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ribosomal peptide biosynthesis (Walsh, 2004).

(ii) The thiolation or peptidyl carrier domain (80 aa) is equipped with a 4<sup>-</sup>-phosphopantetheine (PPan) prosthetic group to which the adenylated amino acid substrate is transferred and thioesterified under release of AMP. Thus, the PPan cofactor acts as thiotemplate and as a swinging arm to transport intermediates between the various catalytic centers. The peptidyl carrier proteins are post-translationally converted from inactive apoforms to their active holoforms by dedicated PPan transferases (Lambalot *et al.*, 1996).

(iii) The formation of a new peptide bond is catalysed by condensation domains
 (450 aa) located between each pair of adenylation and peptidyl carrier domains (Walsh , 2004).

The linear assembly line-like arrangement of multiple of such core units (i– iii) ensure the co-ordinated elongation of the peptide product.

Mechanism of peptide biosynthesis has been outlined in the concept of the 'Multiple Carrier Model of Nonribosomal Peptide Biosynthesis at Modular Multienzymatic Templates' (Stein *et al.*, 1996).

So the surfactin biosynthesis starts with the acylation of the first amino acid glutamate, which is activated by the three-modular enzyme SrfAA (Menkhaus *et al.* 1993; Steller *et al.* 2004). These ten domains comprising synthetase exhibits

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module. However, NRP-synthetases producing lipopeptides us

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#### d the third module of the synthetase SrfAA as well. The third module converts

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L-Leu into the D-configurated isomer by its C-terminal epimerization domain (Figure 1.15). SrfAB incorporates the following three amino acids (Val-Asp-D-Leu) while SrfAC is responsible for the activation and incorporation of the last leucine residue and catalysis of product release by cyclization (Cosmina *et al.*, 1993; Tseng *et al.*, 2002).



Figure (1.14) Non ribosomal peptide synthetases model(Int.3)

#### **1.2.2.4.7 Biosurfactant purification**

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The methods, which are frequently adopted, for the recovery of biosurfactants, include solvent extraction, adsorption followed by solvent extraction, precipitation, crystallization, centrifugation and foam-fractionation (Kowall *et al.*, 1998; Davis *et al.*, 2001; Hsieh *et al.*, 2004; Chen and Juang, 2008).

The optimal approaches for biosurfactant recovery depend on the type and nature of the substrates and fermentation technique and on the type and physicochemical properties of the desired biosurfactants (Lin, 1996).



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Most of the microbial surfactants are amphiphilic molecules with a higher ratio of hydrophobic to hydrophilic character and hence, can be easily isolated by the traditional methods of solvent extraction, precipitation and crystallization (Desai and Banat, 1997; Mukherjee *et al.*, 2006).

Membrane separation processes are also well suited to those downstream processing steps that involve the recovery, concentration and purification of the biomolecules (Muthusamy *et al.*, 2008).

Ultrafiltration is a pressure-driven membrane separation technique for dissolved and suspended materials based on size and molecular scale; It is a very simple procedure and requires no phase change, no chemical addition and little energy. For molecules from 10 A  $^{\circ}$  to 500–1000 A  $^{\circ}$  diameter, ultrafiltration is useful both for product concentration by solvent removal and purification by removal of low molecular weight impurities under hydrostatic pressure (Chtioui *et al.*,2005).

The major filtration characteristics of a solute on passing through a membrane are usually judged by such parameters as permeate flux, rejection coefficient, concentration factor, solute yield or purification factor and hence, the recovery efficiency (Sen and Swaminathan ,2005).

The recovery process including concentration and purification of the product constitutes a major part in the economics of the whole process of biosurfactant

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foam-forming ability of biosurfactants pose serious

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flity to form micelles or miceller aggregates at concentrations higher than the

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critical micelle concentration (CMC) and hence can easily retained by high molecular weight cut-off ultrafiltration membranes (Chtioui *et al.*,2005).

The lipopeptide biosurfactant, surfactin, produced by various strains of *Bacillus subtilis* has been reported to be concentrated and purified from the cell-free fermentation broth using ultrafiltration method (Mulligan and Gibbs , 1990 ; Lin and Jiang , 1997).

A relatively higher molecular weight cut-off membrane was found to be quite effective in the recovery of surfactin, since it forms micelle aggregates of about 50–100 molecules at the CMC (Sen and Swaminathan ,2005).

#### **1.2.2.4.8** Applications of biosurfactants

Surfactants are one of the most frequently used chemicals in our daily lives due to their unique properties, they had applications in an extremely wide variety of industrial processes (Table 1-1), in spite of that there are many advantages of the biosurfactants as compared to their chemically synthesized counterpart as they characterized by: (Lin, 1996 ; Vollenbroich *et al.*, 1997; Kitamoto *et al.*, 2002; Singh and Cameotra, 2004; Ongena and Jacques, 2007; ; Rahman and Gakpe , 2008 ; Xu *et al.*, 2011).

### • Biodegradability

Biosurfactants are easily biodegradable and thus particularly suited for environmental applications such as bioremediation and dispersion of oil

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• Chemical diversity

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**Bio- compatibility and digestibility** 

Which allow their application in cosmetics, pharmaceuticals and as functional food additives (Gautam and Tyagi, 2006; Williams, 2011).

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### • Availability of raw materials

Biosurfactants can be produced from microorganisms grown on raw materials which are available in large quantities (Mukherjee *et al.*, 2006; Singh *et al.*, 2007).

#### • Acceptable production economics

Biosurfactants can be produced from industrial wastes and by – products and this is of particular interest for bulk production (Mukherjee *et al.*, 2006; Salihu *et al.*, 2009).

#### • Use in environmental control

Biosurfactants can be used in handling industrial emulsions, control of oil spills, biodegradation and detoxification of industrial effluents and in bioremediation of contaminated soil (Banat, 1995).

### 1.2.2.4.9 Biological activity of biosurfactants produced by Bacillus spp.

Many lipopeptides that produced by *Bacillus spp.* have attained increasing

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ntiviral and antitumor agents, immunomodulators or speci

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#### e lipopeptides have amphiliphilic structures. Thus, lipopeptides are capable of

penetrating into cells, with the lipophilic hydrocarbon chain interacting with the plasma membrane lipid moiety while the polar amino acids in the peptide part interact with the polar phosphatidyl moieties. Whether lipopeptides are able to damage the integrity of the plasma membrane or create ion selective pores depends on the nature of the lipopeptides and on the phospholipids of the membranes.(Cameotra and Makkar , 2004 ; Singh and Cameotra, 2004 ;Ongena and Jacques, 2007 ; Fathabad, 2011).

Industry	Application	Role of surfactants	
Petroleum Enhanced oil recovery Petroleum			
	De-emulsification	De-emulsification of oil emulsions; oil solubilization; viscosity reduction, wetting agent	
Environmentel	Bioremediation	Emulsification of hydrocarbons; lowering of interfacial tension; metal sequestration	
Environmental	Soil remediation and flushing	Emulsification through adherence to hydrocarbons; dispersion; foaming agent;	
		detergent; soil flushing	

Table (1-1) Industrial applications of chemical surfactants and biosurfactants (Muthusamy *et al.*, 2008).

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#### Functional ingredient

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	Biocontrol	Facilitation of biocontrol mechanisms of		
Agricultural microbes such as parasitism,				
		competition, induced systemic resistance and		
hypovirulence				
	Downstream	Biocatalysis in aqueous two-phase systems and		
Dionrogaging	processing	microemulsions; biotransformations; recovery		
ыоргосезsing		of intracellular products; enhanced production		
		of extracellular enzymes and fermentation		
		products		
Cosmotio	Health and beauty products	Emulsifiers, foaming agents, solubilizers,		
Cosmetic		wetting agents, cleansers, antimicrobial agent,		
		mediators of enzyme action		

#### **1.2.2.4.9.1** The biological activity of surfactin family of lipopeptides

Surfactin and its derivatives is one of the most effective biosurfactants that have many interesting biological activitites. They can inhibit fibrin clot formation, induces formation of ion channels in lipid bilayer membranes, inhibits cyclic adenosine monophosphate (cAMP), inhibits platelet and spleen cytosolic phospholipase A2 , exhibits antiviral and antitumor activities and show antimycoplasma properties (Singh and Cameotra, 2004).

It has been found that Surfactin in concentrations of 30-64  $\mu$ M was cytotoxic to several human and animal cell lines (Vollenbroich *et al.* 1997) and provoked hemolysis (Dufour *et al.* 2005).

Lichenysin A had a stronger surfactant effect and was hemolytic at 10-times

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Surfactin was reported to lyse protozoan membranes (Could ?

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lelanogaster (Assie et al. 2002).

Inactivation of enveloped viruses such as vesicular stomatitis virus (VSV), simian foamy virus (SFV), and suid herpesvirus 1 (SHV 1) by surfactin depended on its hydrophobicity: the C14 and C15 isoforms were more antiviral than C13 (Kracht *et al.* 1999).

The biological effects of surfactin and lichenysin A are probably due to their strong surfactant properties. However, surfactin also forms cation-selective K+ > Na+ channels in black-lipid membrane (BLM )(Sheppard *et al.* 1991). Surfactin-producing *B. subtilis* strains have high swarming motility and biofilm formation, whereas surfactin-nonproducing strains did not swarm or form biofilm (Connelly

*et al.* 2004). Surfactin promotes bacterial cell motion by lowering the surface tension (Kinsinger *et al.* 2003, Hofemeister *et al.* 2004, Mukherjee and Das; 2005).

The amphiphilic structures of the lipopeptides surfactin and lichenysin A explain their ability to form micelles and to penetrate the plasma membrane but in spite of all these properties surfactin has no remarkable fungitoxicity which may be due to the different in the sterol contents of the biological membranes to which surfactins can react with (Carrillo *et al.*, 2003).

### • The antitumor activity of the surfactin family

Over the past years, only a few studies have examined the actual effect of lipopeptide on tumour cells. Studies by Kameda's group (1974) proved that This is a watermark for the trial version, register to get the full one!

irst workers to provide evidence that lipopeptide extract and

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In addition, Wakamatsu *et al.* (2001) discovered that lipopeptide induces neuronal differentiation in rat adrenal phaeochromocytoma PC12 cells and

provided the groundwork for the use of microbial extracellular lipopeptide as a novel reagent for the treatment of cancer cells.

Another studies were conducted by kim *et al.*, (2007) to study the effect of surfactin on the proliferation of LoVo cells, a human colon carcinoma cell line, they found that Surfactin strongly blocked the proliferation of LoVo cells by inducing pro-apoptotic activity and arresting the cell cycle, according to several lines of evidence on DNA fragmentation, Annexin V staining, and altered levels of poly (ADP-ribose) polymerase, caspase-3, p21WAF1/Cip1, p53, CDK2 and cyclin E. The anti proliferative activity of surfactin was mediated by inhibiting

extracellular- related protein kinase and phosphoinositide 3-kinase/ Akt activation, as assessed by phosphorylation levels. Therefore, surfactin may have anti-cancer properties as a result of its ability to down regulate the cell cycle and suppress its survival, this study was followed by the work of Cao *et al.*, (2009a) who isolate surfactin from *Bacillus natto* TK-1 and test its antitumor activity on MCF-7 human breast-cancer proliferation, they found that the inhibition effect of surfactin was a dose- and time-dependent and the antitumour activity was associated with cell apoptosis determined by typical morphological changes.

In 2009 another group of researchers (Wang *et al.*, 2009) published their work twhich studied the molecular mechanisms involved in surfactin( isolated from *B*.

natto T-2) -induction of apoptosis in human leukemic K562 cells, they found that

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protein kinase ERK activation which evoked ERK phore

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#### **.2.4.9.2** The Biological activity of iturin family of lipopeptides

Biological effects of the iturin family peptides are due to their capability of forming ion-conducting pores (Maget-Dana and Peypoux, 1994). Iturin A and bacillomycin L provoked hemolysis and released potassium from erythrocytes (Latoud *et al.* 1986, Aranda *et al.* 2005). Iturin A induced morphological changes in human erythrocytes (Thimon *et al.* 1994). Iturin A, bacillomycins, and mycosubtilin formed channels in BLM (Maget-Dana *et al.* 1985a,b, Maget-Dana and Ptak 1990). Mycosubtilin altered the permeability of the plasma membrane, releasing nucleotides, proteins, and lipids from yeast cells (Besson and Michel 1989) and lysing erythrocytes (Besson *et al.* 1989). The lipopeptides of the iturin family are more active in membranes containing cholesterol, such as mammalian



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acavely from the cells of Saccharomyces cerevisiae mutants containing cholesterol

in the membrane instead of ergosterol (Latoud *et al.*1990). Nucleotides, proteins, polysaccharides, and lipids leaked from *S. cerevisiae* cells exposed to iturin A (Latoud *et al.* 1987) while the lethal concentration in *S. cerevisiae* was 10-60  $\mu$ g/ml (Besson *et al.* 1984). Iturin and bacillomycin L form ion pores by aggregation in the membranes and interacting with sterols (Quentin *et al.* 1982, Maget-Dana and Peypox 1994, Volpon *et al.* 1999). Mycosubtilin formed pores in dimyristoylphosphatidylcholine (DMPC) membranes by interacting with the phospholipids, forming a (1:2) complex with cholesterol, thus stabilizing the ion pore (Maget-Dana and Ptak 1990).

### **1.2.2.4.9.3** The Biological activity of fengicin family of lipopeptides

Fengycins and plipastatins inhibit phospholipase A2, an enzyme affecting inflammation, acute hypertensions, and blood platelet aggregation (Volpon *et al.* 2000). *Bacillus thuringiensis* strain CMB26 produced an analogue of fengycin with a double bond in the fatty acid. It was fungicidal, bactericidal, and insecticidal, and more effective against fungi than was iturin or surfactin (Kim *et al.* 2004). In low molar ratios from 0.1 to 0.5 of fengycin/dipalmitoylphosphatidylcholine (DPPC) membrane fengycin forms pores and at a ratio of > 0.66 it acts as a detergent that solubilize membrane (Deleu *et al.* 2005).

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### **List of Abbreviations**

Abbreviations	Meaning
A <sub>260</sub>	Absorbance at 260nm
A <sub>280</sub>	Absorbance at 280nm
AMP	Adenosine mono phosphate
A-domain	Adenylation - domain
AEFB	Aerobic endospore-forming bacteria
Ala	Alanine
α	Alpha
ATCC	American Type Culture Collection
aa	Amino acids
A °	Ancistrom
Asn	Asparagine

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Deoxyribonucleic acid DNA D.W Distilled water **Evaporative Light Scattering Detection ELSD** Extracellular-related protein kinase ERK Glu Glutamic acid Gln Glutamine Gram per liter g/1 GC ratio Guanine- Cytosine ratio High performance liquid chromatography **HPLC** Hydrophilic-lipophilic balance HLB Ileu isoleucine kb Kilo base

KDa	Kilo dalton
Leu	Leucine
μM	Micromolar
μl	Micro litter
μg	Microgram
μm	micrometer
mg/ml	Milligram per milliliter
mm	millimeter
mN/m	Millinuton per meter
М	Molar
nm	nanometer
NCBI	National centre for biotechnology
NRPS	Non ribosomal peptide synthetases

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Sixteen s ribosomal deoxy ribonuclease acid

SLs	
SHV 1	Suid herpesvirus 1
ST	Surface tension
TLC	Thin layer chromatography
TE	Thioesterase
Thr	Threonine
Tyr	Tyrosine
UV	Ultra violet
U/ml	Unit per milliter
Val	Valine
VSV	Vesicular stomatitis virus
v/v/v	Volume by volume

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Effect of incubation period on biosurfactant production

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### 2. Materials and methods

### **2.1 Materials**

### **2.1.1 Equipments and apparatus**

The following equipments and apparatus were used in this study:-

Equipment	Company /Origin
A garage get tank with power supply	Beijing Six One Instrument
Agarose ger tank with power suppry	Company /China
Autoclave	Shenan /China
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Fluorescence microscope	Olympus
Gel casting tray	Bio Rad/Germany
HPLC-ESI/MS (Agilent 1100 series) with Zorbax Eclips XDB-C18 Analytical (4.6x150) mm	Agilent Technologies /USA
Ice maker	Shenan
Incubator	Memmert /Germany
Incubator with CO <sub>2</sub> supply	Memmert
Laminar air flow	Jia Jie /china

Followed -----

Luminometer	Promega / USA
Lyophylizer	Sartorius /Germany
Magnetic stirrer	Fuhe 78-1
96 well microtiter plate, 6 well plate	Hangzhou Yanhui /China
Microfuge	Heraeus sepatech /Germany
Variable micropipettes( 10, 100, 1000)µl	Eppendorf /Germany
Microwave oven	Media /china
Milipore filter unit	Millipore Company
Minicentrifuge	Crystal MLX -206 /Germany
Flasks of tissue culture: plastic disposable of	Iwaki /Ianan

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	Shaker incubator	Memmert
	Tensiometer	Laryee/China
	Thermal cycler	Bio Rad /USA
	TLC tank	Sigma/ USA
	UV transilluminator	Beckman Instruments/USA
	Vortex mixer	Qilinbeirer
	Water bath	Atom/ Eryland

### 2.1.2 Chemicals and biological materials

The following chemicals and biological materials were used in this study:-

Materials	Company / Origin
Agarose	Sigma /USA
Ammonium chloride(NH <sub>4</sub> Cl)	Sigma
Ammonium nitrate (NH <sub>4</sub> NO <sub>3</sub> )	Sigma
Ammonium sulphate $((NH_4)_2SO_4)$	Sigma
Boric acid	Sigma
Calcium chloride dihydrate (CaCl <sub>2</sub> .2H <sub>2</sub> O)	

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#### Ethylene diamine tetra acetic acid (EDTA) Sigma Ferrous sulphate heptahydrate (FeSO<sub>4</sub>.7H<sub>2</sub>O) Sigma Glucose Sigma Glycerol Sigma Atago Company /China Machine oil Hydrochloric acid (HCl) Sigma Isopropanol Sigma Magnesium sulphate (MgSO<sub>4</sub>) Sigma

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Followed —

Materials	Company / Origin
Manganese sulphate (MnSO <sub>4</sub> )	Sigma
Manganese sulphate monohydrate (MnSO <sub>4</sub> .H <sub>2</sub> O)	Sigma
Manganese sulphate tetrahydrate (MnSO <sub>4</sub> .4H <sub>2</sub> O)	Sigma
Methanol	Sigma
Mineral oil	Al Dorah oil refineries
Lithium chloride hexa hydrate NiCl <sub>2</sub> .6H <sub>2</sub> O	Sigma
Olive oil	Wuhan for food
Penicillin	Sigma
Peptone	

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#### Streptomycin powder Sigma Sigma Surfactin standard Sodium molbdate dihydrate (Na<sub>2</sub>MoO<sub>4</sub>.2H<sub>2</sub>O) Sigma Disodium selenoxide (Na<sub>2</sub>SeO<sub>4</sub>) Sigma Sigma Sucrose Sigma Urea Sigma Yeast extract Zinc sulphate heptahydrate (ZnSO<sub>4</sub>.7H<sub>2</sub>O) Sigma Aluminium potassium sulfate(AlK(SO<sub>4</sub>)<sub>2</sub> Sigma
### 2.1.3 Reagents and solutions

Boric acid

NiCl<sub>2</sub>.6H<sub>2</sub>O

 $Na_2SeO_4$ 

### 2.1.3.1 Solution EA (Youssef et al., 2004)

This solution was prepared by dissolving 25 g of  $MgSO_4$  in 950 ml of D.W., then volume was completed to 1000 ml in a volumetric flask, and sterilized by autoclaving.

## 2.1.3.2 Solution EB (Youssef et al., 2004)

To prepare this solution 100g of  $(NH_4)_2SO_4$  was dissolved in 950 ml of D.W., then volume was completed to 1000 ml in a volumetric flask, and sterilized by autoclaving.

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0.01

0.005

0.003

All components were dissolved in 950 ml of D.W., then volume was completed to 1000 ml in a volumetric flask, pH was adjusted to 7 and sterilized by filtration.

# 2.1.3.4 Manganese and magnesium sulfate solution (Jacques et al., 1999)

This solution was prepared by dissolving 0.4 g of  $MnSO_4.H_2O$  and 5 g of  $MgSO_4$  in 950 ml of D.W., then volume was completed to 1000 ml in a volumetric flask.

# 2.1.3.5 Trace elements solution (Jacques et al., 1999)

It was prepared to be consisting of the following components:

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	EDTA		1	
	NiCl <sub>2</sub> .6H <sub>2</sub> O		0.004	
	KI		0.66	

All components were dissolved in 950 ml of D.W., then volume was completed to 1000 ml in a volumetric flask, pH was adjusted to 7 and sterilized by filtration.

# 2.1.3.6 Heat inactivated fetal bovine serum (Wang et al., 2007)

It was prepared by incubating fetal bovine serum in the water bath at 56  $^{\circ}\mathrm{C}$  for 30 min.

# 2.1.3.7 PCR mixture

PCR master mixture was supplied by Takara company. It was composed of the followings:

ComponentVolume(μl)Template DNA<0.5</td>(10×) Easy Tag Buffer (Mg++)5

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Then volume was completed to  $50 \,\mu$ l with D.W.

# 2.1.3.8 TLC solvent mixture (Symmank et al., 2002)

It was prepared by mixing chloroform , methanol and deionized D.W. at ratios of 65:25:4 (v/v/v).

# 2.1.3.9 Biosurfactant standard solution

It was prepared according to Symmank *et al.* (2002) by dissolving 50 mg of biosurfactant in 1 ml of methanol.

#### 2.1.3.10 Rhodamine B reagent (0.25%)

This reagent was prepared by dissolving 0.25 g of rhodamine B in 100 ml of absolute ethanol.

#### 2.1.3.11 Trypsin solution

It was prepared according to Seo –young *et al.*, (2007) by dissolving 0.25 g of trypsin and 0.02 g EDTA in 100 ml of phosphate buffer solution (pH 7.4), mixed gently and sterilized by filtration.

### 2.1.3.12 Phosphate buffer saline (PBS 1%)

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 $\mathbf{V}$  ater was mixed with JC-1 stain by vortexing followed by the addition of 2 ml JC-

1 staining buffer (5x).

### 2.1.3.14 TBE buffer solution (Lema et al., 1994)

To prepare stock solution of TBE buffer  $(5\times)$ , 54 g of Tris base, 27.5 g of boric acid were dissolved in 900 ml D.W., then 20 ml of 0.5 M EDTA was added, pH was adjusted to 8.3 and the volume was completed to 1000 ml with D.W.

### 2.1.3.15 Ethidium bromide (10mg/ml)

Stock solution of ethidium bromide was prepared according to Maniatis *et al.*, (1982) in a concentration of 10 mg/ml by dissolving 100 mg of ethidium bromide in 10 ml of D.W.

### 2.1.3.16. Hanks balanced salt solution (HBSS):

It was prepared to be consisting of the following components:

Component	Weight (g/l)
CaCl <sub>2</sub>	0.14

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NанСО <sub>3</sub>	
Na <sub>2</sub> PO <sub>4</sub>	0.09
D-glucose	1.00
Hepes	2.08

Each of the above constituents was dissolved separately.  $CaCl_2$  was added last, and made up in 1L. pH adjusted to 7,the solution was sterilized by filtration (Freshney,2000).

# 2.1.3.17 Trypan Blue Stain:

Trypan blue powder (1g) was dissolved in (100ml) Hank's solution (2.1.3.17). The solution was filtered by Watman filter paper stored at 4°C until use then diluted 1:10 in Hank's solution for using (Freshney, 2000).

# 2.1.4 Culture media (Fluka/ Germany)

The following media were used in this study

# 2.1.4.1 Ready made culture media

- Nutrient agar
- Nutrient broth

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ese media were prepared as recommended by the metalia, u

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This medium was consisted of the following components

Component	Weight (g)
KH <sub>2</sub> PO <sub>4</sub>	2.1
K <sub>2</sub> HPO <sub>4</sub>	13.9
Sucrose	10
NaCl	0.5
Yeast extract	50
NaNO <sub>3</sub>	1

All components were dissolved in 950 ml D.W., pH was adjusted to 6.9, then volume was completed to 1000 ml and sterilized by autoclaving. After cooling, 10 ml of solution EA (item 2.1.3.1), solution EB (item 2.1.3.2), and solution EC (item 2.1.3.3) were added to 1 liter of the above medium.

# 2.1.4.2. 2 Jacques medium (Jacques et al.,1999)

This medium was consisted of the following components

Component	Weight (g)	
KH <sub>2</sub> PO <sub>4</sub>	1.9	
Sucrose	20	

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completed to 1000 ml and sterilized by autoclaving.

# 2.1.4.2.3 Bushnell –Hass medium (Patel and Desai,1997)

This medium was consisted of the following components

Component	Weight (g)	
KH <sub>2</sub> PO <sub>4</sub>	1	
K <sub>2</sub> HPO <sub>4</sub>	1	
NH <sub>4</sub> NO <sub>3</sub>	1	
MgSO <sub>4</sub> .7H <sub>2</sub> O	0.2	
CaCl <sub>2</sub> .2H <sub>2</sub> O	0.02	
FeSO <sub>4</sub> .7H <sub>2</sub> O	0.05	
Yeast extract	0.5	

All components were dissolved in 950 ml D.W., before 1 ml of trace element solution (item 2.1.3.5) was added, pH was adjusted to 7, then volume was completed to 1000 ml and sterilized by autoclaving.

## 2.1.4.2.4 Blood agar medium

This medium was prepared according to manufacturing company (Fluka / Germany) by dissolving 31 g of blood agar base in 900ml D.W., pH was adjusted to 7.3, then volume was completed to 950 ml with D.W., and sterilized by autoclaving . After cooling, 50 ml of defibrinated sheep blood was added and mixed thoroughly then poured into petridishes.

#### \_\_\_\_\_2.1.4.2.5 RPMI 1640 medium

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This medium was enriched with 10% heat inactivated fetal bovine serum

(item 2.1.3.6), 100U/ml penicillin and 100µg Streptomycin.

#### **2.1.5 Biological kits**

#### 2.1.5.1 Genomic DNA extraction kit ver. 2

Takara Minibest bacterial genomic DNA extraction kit ver. 2 (Takara Biotechnology,Japan) was consist of the following solutions and buffers:

• SP buffer, RNAase AI, Lysozyme, EDTA buffer, Solution A, Solution B, Solution C, DB buffer, Rinse A, Rinse B, Elution Buffer

2.1.5.2 Cell titer 96 non- radioactive cell proliferation assay (MTT assay) (Promega, USA)

This kit was consist of two solutions:

- Dye solution (colorimetric 3-(4, 5- dimethylthiazol-2-yl)-2, 5diphenyl tetrazolium bromide)
- Solubilization solution / stop mix

#### 2.1.5.3 Caspase –G10 3/7 assay kit: (Promega, USA)

This kit was composed from the following components:

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Jc-1 (2000×), JC-1 staining buffer, positive control

### 2.2 Methods

### 2.2.1 Samples collection

After removing the soil surface layer, 45 soil samples heavily contaminated with engine oils were collected from different fuel stations in addition to Al- Dorah oil refinery in Baghdad Province from Feb. to Aug., 2008.

53

### 2.2.2 Isolation of Bacillus spp.

Isolation of *Bacillus* spp. was carried out according to Claus and Barkeley (1986) by adding 4 g from each soil sample to 20 ml of sterile D.W., mixed thoroughly then heated to 80 °C with gentle agitation for 10 min. and left to cool at room temperature, then 0.1 ml aliquot was taken from each sample and spreaded on a nutrient agar plates and incubated aerobically for 24 hrs. at 30 °C.

After incubation, colonies appeared with different shapes and sizes were selected for further identification.

#### 2.2.3 Identification of bacterial isolates

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Single colony of each bacterial isolate was fixed on a clean slide and stained

with Gram stain to study its Gram reaction and spore formation under the light compound microscope (Atlas *et al.*, 1995).

#### 2.2.3.3 Molecular identification

Molecular identification of bacterial isolates were achieved according to the analysis of DNA sequences and degree of similarity of 16S rDNA.

Bacterial isolates suspected to be *Bacillus* spp. were selected and propagated individually in nutrient broth for 18 hrs. at 30°C to obtain fresh cultures for extraction of genomic DNA.

### 2.2.3.3.1 Extraction of Genomic DNA

Genomic DNA was extracted from local isolates of *Bacillus* spp. using Takara Minibest Bacterial Genomic DNA Extraction Kit ver. 2 (2.1.5.1) as follows:

Aliquots of 4 ml from fresh culture of each bacterial isolates was centrifuged, at 10000 rpm (4 °C) for 2 min., then 150µl aliquots of sp buffer and RNAase AI solutions were added respectively to cells precipitate followed by the addition of 20 µl of lysozyme solution and incubated at 25°C for 10 min.

• After incubation, 30 µl of EDTA buffer solution was added, and left at 25 This is a watermark for the trial version, register to get the full one!

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• Supernatant was discarded; recentrifuged at 12000 rpm for another 1 min to

obtain pellet.

- Aliquot of 400 µl of solution DB was added, shacked, then contents were transferred to spin column in filter cup and centrifuged at 12000 rpm for 1 minute.
- The spin column was transferred to another filter cup, then 500 µl of Rinse A solution was added to the column and centrifuged at 4°C, 12000 rpm for 1minute.
- Aliquot of 700 µl of Rinse B solution was added and centrifuged at 12000 rpm for 1 minute.

 Spin column was placed in a new filter cup, then 60µl of elution buffer was added and incubated at 25 °C for 10 min., centrifuged at 12000 rpm for 1 minute, then filtrate which containing genomic DNA was collected.

#### 2.2.3.3.2 Quantization of DNA concentration (Maniatis et al., 1982)

Purity and concentration of DNA solution was measured by using Biophotometer device, by adding 8  $\mu$ l of DNA sample to 72  $\mu$ l of D.W. in a quartz cuvete and the absorbency at 260 nm and 280 nm was measured after calibration with D.W. at 260 nm and 280 nm respectively, Pure DNA has an A<sub>260</sub> /A<sub>280</sub> ratio of 1.7-1.9.The concentration of double strand DNA ( $\mu$ g/ml)was measured directly by the device

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2.2.3.3.3 Gel electrophoresis (Maniatis et al.,

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# y heating and solution is clear, after cooling to about 55°C, Ethidium bromide

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solution (2.1.3.15) was added at a concentration of 0.5  $\mu$ g/ml. Finally gel solution was poured into tray to a depth of about 5mm and allowed to a solidify for about 20 min at room temperature followed by removing the comb from the tray and placed in the electrophoresis chamber , submerged with TBE buffer (1×) , DNA samples were prepared by mixing 1µl of the (6×) gel loading dye with 5 µl of the DNA then loaded in the wells, electrophoresis was done at 72 v until dye markers have migrated in appropriate distance, then gel was removed and visualized under UV light in the transiluminator.

#### 2.2.3.3.4 Amplification of 16S rDNA

Two sets of primers were used as listed in table (2-1) to amplify 16S rDNA sequence for identification of bacterial isolates (Goto *et al.*, 2000; Thomas, 2004).

Table (2-1) Types of primers used to amplify the 16S rDNA sequence of bacterial isolates.

Primer	type	Sequence	Size	Tm
			(bp)	(°C)
27F	Forward primer	5'-AGAGTTTGATCCTGGCTCAG-3'	20	61
	Reverse primer	5'-ACGGTTACCTTGTTACGACTT -3'		59
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#### PCR reaction was performed by adding 3µl of the genomic DNA isolated

from locally isolated *Bacillus* spp. as DNA template to 27 µl of a PCR mixture (2.1.3.7). The initial denaturation step was done at 94°C for 1 min. followed by 30 cycles at 94 °C for 30 sec. at 47 °C for 30 sec. and at 72 °C for 1.5min, then a final extension reaction was achieved at 72 °C for 10 min. PCR products were sequenced by Huazhong agricultural company using ABI310 DNA sequencer and the ABI PRISM Big Dye Terminator Cycle Sequencing kit version 3.1 (Perkin-Elmer Applied Biosystems,CA,U.S.A.). DNA sequences were analyzed to determine the degree of similarity using the National Centre for

Biotechnology Information (NCBI; Bethesda, MD,USA) Blast system (http://www.ncbi.nlm.nih.gov/BLAST).

# 2.2.4 Sterilization methods (Atlas *et al.*, 1995)

# 2.2.4.1 Dry heat sterilization

Glassware were sterilized in an oven at 160°C for 3 hrs.

## 2.2.4.2 Moist heat sterilization

Media and solutions were sterilized by autoclaving at  $121^{\circ}C$  (15 pounds/ inch<sup>2</sup>)

for 30 min

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then wrapped with a parafilm, these plates can be store at 4 °C for a week as stock culture.

## 2.2.5.2 Medium term storage

Bacterial isolates were maintained by streaking on nutrient agar slants in screw- capped tubes containing 5 ml of the medium and incubated at 30 °C for 24 hrs., these slants can be stored for a few months at 4 °C.

#### 2.2.5.3 Long term storage

Aliquetes of 8.5 ml of the exponential phase of the bacterial growth was added to 1.5 ml of sterilized glycerol in a screw – capped tubes and stored at - 20 °C.

### 2.2.6 Screening of biosurfactant production by local isolates of *Bacillus* spp.

### 2.2.6.1 Preparation of inoculums

Single colony of each bacterial isolates was selected and used to inoculate 10 ml of nutrient broth medium, and incubated in shaker incubator (180 rpm) for 18 hrs. at 30 °C, then 10 ul of fresh culture was used to inoculate 10 ml of the

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medium and Jacques medium, separately, in Erlenmyer flasks (250 ml) in triplicate, and incubated aerobically (without shaking) for E medium and with shaking at 180 rpm for Jacques medium at 30 °C for three periods (24, 48 and 72 hrs.) .Each culture was then centrifuged at 4 °C, 10000 rpm. for 15 min. Production of biosurfactant was investigated in cell- free supernatant.

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#### 2.2.6.3 Qualitative screening

Qualitative screening for the ability of local isolates of *Bacillus* spp.of biosurfactants production were achieved by two methods:

#### 2.2.6.3.1 Detection of hemolytic activity (Banat, 1993)

Haemolytic activity was detected on blood agar medium by streaking each local isolate on blood agar plate and incubated for 48 hrs. at 30 °C, then plates were visually inspected for the formation of clear zones around the growing colonies.

#### 2.2.6.3.2 Oil spreading method (Morikawa et al., 2000)

This method was performed by adding 20  $\mu$ l of crude oil to the surface of water layer in large petridishes (25 cm diameter) (50ml /petridish). 10  $\mu$ l of cell – free supernatant were then added to the surface of oil layer. The diameter of clear zone on the oil surface was measured in centimeters which represent the

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*Bacillus* spp. was achieved by measuring surface tension of cell – free supernatant according to McInerney *et al.* (1990) using Du Nouy ring tensiometer and by estimation of biosurfactant dry weight .

#### 2.2.6.4.1 Surface tension measurement

Surface tension for each sample was measured by equilibrating 15 ml of cell- free supernatants for 15 min in a small weighing dish prior to measuring the surface tension in which the reading was recorded at the moment when the instrument's ring was detached from the surface of supernatant that immersed in it.

After calibration with D.W., glycerol and isopropanol, respectively, the percentage of lowering the surface tension was calculated according to the following equation.

Percentage (%) of lowering the surface tension = ST1-ST2 \*100

ST1

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ST1= Surface tension of the medium or water without any addition.

ST2= Surface tension of the medium after cell growth or addition of biosurfactant.

#### 2.2.6.4.2 Estimation of biosurfactant dry weight

Cell –free supernatants were subjected to acid precipitation by adding drops of 6N HCl to a final pH of 2 and allowing the precipitate to form at 4°C for This is a watermark for the trial version, register to get the full one!

Precipitate was collected by centrifugation with 1

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2.7 Optimization of biosurfactant production

Optimum conditions for biosurfactant production by the selected isolate of *Bacillus* spp. was determined according to Makkar and Cameotra (1998).

Optimum conditions included the optima of : carbon source, carbon source concentration , nitrogen source, nitrogen source concentration, dipotassium hydrogen phosphate concentration , pH , temperature, time of incubation and shaking speed respectively.

Each experiment was done in an Erlenmyer flask containing 50 ml of the production medium (Bushnell – Hass medium at pH 7) by inoculating with 0.5 ml of fresh culture of the selected isolate , then flask was incubated in shaker

incubator (180 rpm) at 30 °C for 72 hrs. After incubation, medium pH, surface tension, OD of bacterial growth and biosurfactant dry weight were measured.

### 2.2.7.1 Effect of carbon source

Five carbon sources (Sucrose, Glucose ,Olive oil, Corn Oil, Machine oil) were used to determine the optimum for biosurfactant production by the selected isolate of *Bacillus* spp. Each of these carbon sources was added to the production medium in a concentration of 2% (w/v) or (v/v).

#### 2.2.7.2 Effect of carbon source concentration

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Peptone and Tryptone) were used to determine the optimum for biosurfactant production by the selected isolate of *Bacillus* spp. These nitrogen sources were added to the production medium in a concentration of 0.1% (w/v) or (v/v).

#### 2.2.7.4 Effect of nitrogen source concentration

Different Concentrations (0.15, 0.3, 0.6, 1, 1.5 and 2%) of the optimum nitrogen source were added to the production medium to determine the optimum for biosurfactant production by the selected isolate of *Bacillus* spp.

#### 2.2.7.5 Effect of K<sub>2</sub>HPO<sub>4</sub> concentration

Different concentrations (0.03, 0.05, 0.1, 0.15, 0.2, 0.5, 1, 1.5, and 2 g/L) of  $K_2$ HPO<sub>4</sub> were examined to determine the optimum for biosurfactant production by the selected isolate of *Bacillus* spp.

### 2.2.7.6 Effect of medium pH

The production medium was adjusted to different pH values (5,6,7,8,9) to determine the optimum for biosurfactant production by the selected isolate of *Bacillus* spp.

#### 2.2.7.7 Effect of temperature

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#### .7.8 Effect of incubation period

To study the effect of incubation period on the ability of the selected isolate of *Bacillus* spp in biosurfactant production , production medium was incubated for 12, 24, 36, 48, 60, 72 and 96 hrs.

#### 2.2.8 Purification of biosurfactant

Biosurfactant, produced by the selected isolate of *Bacillus* spp., was purified according to Moran *et al.* (2002) and Das *et al.* (2008). This was achieved by centrifugation of the culture medium (10000 rpm) for 10 min at 4°C. Cell –free supernatant was subjected to acid precipitation by adding drops of 6N HCl to a final pH of 2 to allow precipitation at 4°C overnight. Acid precipitate was collected

by centrifugation (15000 rpm) at 4°C for 15 min., then pellet was resuspended 2 ml of D.W., pH was adjusted to 7 with agitation for complete dissolving, then biosurfactant precipitate was lyophilized and weighed . From the lyophilized biosurfactant precipitate ,200  $\mu$ g was dissolved in 4 ml of D.W. and filtered through a Centricon Centrifuge Filter (Mw Cut.off = 30 KDa) at 5000g.

Retentates were diluted in 50% methanol and filtered again. Filtrates were dried at 40  $^{\circ}$ C for 24 hrs. to obtain the biosurfactant.

#### 2.2.9 Characterization of the biosurfactant

2.2.9.1 Thin layer chromatography (Symmank et al., 2002)

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lid to allow the migration of solvent from the bottom to the top of the plates until

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reaches the end of the upper side (1.5 cm from the top), then plates were removed from the tank , air dried and sprayed with D.W. for detection of hydrophilic compounds or sprayed with rhodamine B prepared in (2.1.3.10) for detection of the presence of lipids under ultraviolet light. The relative mobility (Rm) of each spot was calculated according to the following equation.

Relative mobility (Rm) = Distance travelled by the compound Distance travelled by the solvent

#### 2.2.9.2 High Performance Liquid Chromatography

The molecular mass of pure biosurfactant produced by the selected isolate was determined by HPLC-ES-MS (electrospray-ionization MS, Agilent Technology, USA) (figure 2-1)equipped with an electrospray ion source. The electrospray needle and capillary voltage were operated at 4.5 and 10kV respectively.

Other conditions were in accordance with the manufacturer's recommendations. HPLC was equipped with a Zorbax Eclipse XDB-C<sub>18</sub> analytical column (4.6×150 mm; 5 $\mu$ m particle size). The mobile phase consisted of two parts .A and B. A was acctonitrile , and B was 0.1% formic acid, 5mmol/l ammonium acetate and D.W.. The parts were combined at an A/B ratio of 90:10(v/v). The This is a watermark for the trial version, register to get the full one!

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stage mass spectrometer (MS2). The spectra were recorded using a Macromass, Wythenshaw, Manchester, U.K.).The capillary voltage was 3.50 kV and the collision gas was argon.

#### 2.2.10 Studying the antitumor activity

#### 2.2.10.1 Cell lines

Four cell lines [ Human leukemia (K562), Human epidermal larynx carcinoma cell line (Hep-2), Mice leukemia (L1210)] were purchased from Cell Bank of Shanghai Institute for Cell Biology (Shanghai, China), while the normal liver cell line (Lo2) was purchased from cell bank of Chung Shan hospital (Shanghai, China).



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Figure (2-1) HPLC-ESI/MS (Agilent 1100 series) with Zorbax Eclips XDB-C18 Analytical (4.6x150) mm.

## • Human Leukemia (K562)

The human K562 cell line was established by Lozzio and Lozzio (1975) from a 53 year old patient with chronic myelogenous leukemia in blast crisis. The cells are non-adherent and rounded, in culture they exhibit much less clumping

than many other suspension lines (Lozzio and Lozzio ,1979). This cell line was used at a passage number 15.

### • Human epidermal Larynx carcinoma cell line (Hep-2)

The Hep-2 cell line was established in 1952 by Moore *et al.* from tumors that had been produced in irradiated-cortisonized weanling rats after injection with epidermoid carcinoma tissue from the larynx of a 56-year-old male, they form adherent cells. A hardy cell line, Hep-2 resists temperature, nutritional, and environmental changes without a loss of viability (Toolan, 1954).

It has high proliferation rate and a 23 hours cell cycle (Grem and Fisher, 1989).

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products for cytotoxic activity; preliminary testing for antitumor activity by the NCI in cancer chemotherapy screening studies. This cell line was used at a passage number 15.

### • Normal human liver cell line (Lo2)

The lo2 cell line was established from an adult Chines mail by Cell Biology of Shanghai institute of Cell bank of China (Liu *et al.*, 2007), this cell line was used at a passage number 7.

Cells were recovered by rapid thawing at  $37^{\circ}$ C in a water bath, centrifuged at 800 rpm for 10 min. at room temperature, then resuspended in 5ml culture medium [(2.1.4.2.5) for K562 and L1210, (2.1.4.2.6) for Hep-2 and Lo2] transferred to a T-25 tissue culture flask, which was incubated in a humidified atmosphere with 5% CO<sub>2</sub> and 95% air at 37°C for 24 hrs. After incubation, the medium was replaced with a fresh one.

Counting of viable cells was carried out using trypan – blue dye (0.4%).Dead cells take up the dye and appear blue under microscope while living cells exclude the dye and appear white.

#### 2.2.10.2 Cell Cultures (Wang *et al.*,2007)

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medium, washing the cell sheet twice with PBS (2.1.3.12), followed by detaching

the cell line from the flask surface by adding 2 ml trypsin solution (2.1.3.11) for each flask and incubated in a humidified atmosphere with 5%  $CO_2$  and 95% air at 37°C for few minutes until the cells were round up, then dispersed gently, trypsin was inactivated by adding 15 ml of the culture media since its supplemented with fetal bovine serum (2.1.4.2.6) and subdivided into two flasks which incubated under the same conditions.

#### 2.2.10.3 Cytotoxicity tests

#### **2.2.10.3.1** Preparing the cell lines (Cao *et al.*,2009b)

Cell lines were seeded in 96 well microtiter plate for the cell viability study (MTT) and incubated for (48-72) hrs. to a final concentration of 100000 cell/ ml ,then treated with different concentrations [2,4,8,16,32,64  $\mu$ g/ml] of biosurfactant in methanol produced and purified from the selected isolates and standard surfactin for 24,48 and 72 hrs. before testing with kit(2.1.5.2).

In order to study caspase activity, Hep-2 cell line was seeded in 96well microtiter plate and incubated for 72 hrs. to the final concentration of 20000 cell/ ml then treated with 8µg/ml of the biosurfactant preparation from the selected isolate for 6,12 and 18 hrs. before testing with kit(2,1,5,3), while the studying of **This is a watermark for the trial version, register to get the full one!** 

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## 2.2.10.3.2 Cell Titer 96 Non- Radioactive cell Proliferation Assay (MTT assay)

#### • Method:

A volume of 15  $\mu$ l of the dye solution was added to each well of the 96 well microtiter plate that contain 100  $\mu$ l of the treated cells or blank then incubated at 37°C for 4 hrs. in a humidified (5%) CO<sub>2</sub> atmosphere , after incubation 100  $\mu$ l of the solubilization solution / stop mix was added to each well and left for an hour, the contents of each well was mixed and finally the absorbance at 570 nm wavelength of the plate was recorded .

The cell inhibition rate (%) was calculated according to the following equation (Wang *et al.*, 2007):

Cell inhibition rate (%) = [(Average absorbance of control cells– Average absorbance of treated cells)/ Average absorbance of control cells] $\times 100$ 

Data were analyzed by 2-way analysis of variance with ANOVA- test followed by Duncan test. Data were presented as means of three replicates  $\pm$  SD. The level of significance P < 0.05 was used for analysis of variance test (ANOVA) (Al-Mohammed *et al.*, 1986).

#### 2.2.10.3.3 Caspase – G10 3/7 assay kit

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until the substrate was thoroughly dissolved to form the Caspase -G10 3/7

reagent.

#### • Reaction preparation

The following reactions were prepared to detect caspase -3 and -7 activities in cell Culture.

- \* Blank reaction :compose of the following Caspase G10 3/7 reagent, vehicle solvent of protein ,cell culture medium without cells.
- \* Negative control : Caspase –G10 3/7 reagent and vehicle –treated cells in medium.

Blank reaction was used to measure background luminescence associated with the cell culture system and Caspase -G10 3/7 reagent ,so the value for the blank reaction was subtracted from experimental values.

# • Method

- \* For each well of cells treated with biosurfactant (2.2.10) equal volume of the reagent was added and mixed well, then incubated for 3 hrs.
- \* luminescence from the plate was measured by luminometer

# 2.2.10.3.4 Mitochondrial membrane potential assay kit with JC-1

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pernatant sucked and washed twice with JC-1 staining buffer (1x). Finally 2 ml

of cell culture fluid was added then observed with fluorescence microscope (fluorescence wave length (529 nm and 590 nm).

# Recommendations

1- Genetic study on the biosurfactant producer locally isolated *Bacillus* isolates (B1,B2,B3,B4,B5,B6,B7,B8,B9,B10) to determine the genetic elements responsible for biosurfactant production.

2- Mutagenesis of *Bacillus* B6 to develop its ability in biosurfactant production using physical and chemical mutagens.

3- Studying the biological activity of the selected isolates as antimicrobial, antiviral, antifungal agents.

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Bo in Hep-2 cell line. Benefits for registered users: 1.No watermark on the output documents. 2.Can operate scanned PDF files via OCR. 3.No page quantity limitations for converted PDF files.

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#### 3. Results and discussion

# 3.1 Isolation and identification of biosurfactant – producing *Bacillus* species 3.1.1 Isolation and primary characterization of *Bacillus* spp.

In order to isolate biosurfactant producing *Bacillus*, 45 heavily oil contaminated soil samples were collected from fuel stations and Al- Dorah oil refineries in Baghdad Province, from which 109 bacterial isolates were isolated.

Oil contaminated soil were chosen for isolation of biosurfactant producing *Bacillus* since they create a selective niche to such microorganisms that have enhanced ability to utilize hydrocarbons as nutrition source (Rahman *et al.*, 2002; Rahman *et al.*, 2003; Priya and Usharani, 2009).

When these isolates were subjected to morphological and cultural This is a watermark for the trial version, register to get the full one!

88 isolates where suspected to be *Bacillus* isolates since the were

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wrinkled appearance, they vary greatly in translucence, opaqueness and in color so according to Collins and Lyne (1985), Claus and Berkeley (1986) and Fritz (2004) all these features were characteristics of *Bacillus* spp.

Such high percentage of *Bacillus* isolation (80.7 % of the total isolates) was expected due to the predominant of *Bacillus* spp in the soil (Earl *et al.*,2008) and to the selective isolation method was used that impose high temperature (80°C) on the sample which will select for *Bacillus* that can survive due to their spores formation (Claus and Berkeley ,1986).

#### **3.1.2 Screening of biosurfactant producing isolates**

To detect the biosurfactant production ability of the 88 *Bacillus* isolates, two detection methods were used , the surface tension measurements and haemolytic activity on sheep blood agar medium .

Results indicated in table (3.1) showed that 87 bacterial isolates were capable of lowering surface tension (ST) of the cell free supernatant after cultivation in E medium, while only 49 isolates were able to haemolyse blood when grew on blood agar plates. Although some negative isolates with the later detection method were capable of lowering (ST) with noticeable percentages (29-30) %, such observation was also detected by Youssef *et al.* (2004) who found

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Jpon quantitative screening for the highest biosurfactant producing isolates

in E medium, ten isolates were chosen for further studies named (B1, B2, B3, B4, B5, B6, B7, B8, B9, B10) as shown in table 3.2.

The ability of these ten isolates in lowering (ST) was also studied by another detecting method (oil spreading method), results in table (3-3) showed that diameter of the oil spreading was proportional with lowering of surface tension by tensiometer, such results were also recorded by Priya and Usharani, (2009).

Table (3-1) Detection of biosurfactant activity by measurement (ST) of E medium by Du Nouy ring tensiometer and blood haemolysis methods after cultivation in E medium for 24 hrs.

Number of isolates	Percentage of lowering the surface tension%*	Number of isolates	Blood haemolysis
18	28-30	14	+
		4	-
23	25-27	11	+
		12	-
40	14-24	19	+
		21	-
б	4.5-13	5	+

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Du Nouy ring tensiometer.

Symbol of isolates	Percentage of lowering surface tension%	
B1	29.10	
B2	29.54	
B3	29.72	
B4	28.60	
B5	29.55	
B6	29.56	
B7	29.55	
B8	28.40	
<b>B</b> 9	30.00	
B10	29.20	

Table (3-3) Detection of biosurfactant activity by measuring lowering of (ST) of Jacques medium by Du Nouy ring tensiometer and oil spreading method after cultivation in Jacques medium for 72 hrs (180 rpm).

Symbol of isolates	Percentage of lowering surface tension%	Oil spreading method (cm)
B1	43.3	6.3
B2	42.3	6.0
B3	41.5	5.8
B4	46.2	7.0
B5	44.6	6.5
B6	43.5	6.3

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#### 3.1.3.1 Isolation of Genomic DNA

In order to amplify16S rDNA for local isolates of *Bacillus* spp., genomic DNA of the selected isolates was extracted to provide a PCR template for the amplification.

Purity ratios of the extracted DNA samples were (1.8-1.9) which indicates a high purity since a pure DNA preparation has expected A260/A280 ratio of  $\approx$ 1.8. Which are based on the extinction coefficients of nucleic acids at 260 nm and 280 nm (Maniatis *et al.*, 1982).

Such results were also observed when the DNA samples were analyzed by gel electrophoresis, in which sharp DNA bands were detected indicating purified DNA samples as shown in figure (3-1) .

The concentrations of the DNA samples were range between (24-79)  $\mu g/\mu l$ .



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1. Bacillus B10, Lane (2): Bacillus B9 isolate , Lane (3): Bacillus B8, Lane (4): Bacillus B7 , Lane (5): Bacillus B6, Lane (6): Bacillus B5 , Lane (7):

Bacillus B4, Lane (8): Bacillus B3, Lane (9): Bacillus B2, Lane (10): Bacillus B1,

Lane (11): 15Kb DNA landmark ladder

## 3.1.3.2 Amplification of 16S rDNA

Two sets of primers were used to amplify the 16S rDNA.

Forward Primers:

```
    (B16SF) 5'-TGTAAAACGACGGCCAGTGCCTAATACATGCAAGTCGAGCG-3'
    (27F) 5'-AGAGTTTGATCCTGGCTCAG-3'
```

Reverse Primers:

(B16SR) 5'-CAGGAAACAGCTATGACCACTGCTGCCTCCCGTAGGAGT-3'

#### (1492R) 5'-ACGGTTACCTTGTTACGACTT -3'

The first set (B16SF and B16SR) were used to amplify the 5' end region (approx. 275 bp) which was the hypervariant region (HV region) in the16S rDNA gene and was highly specific for each type of bacterial strain of *Bacillus* spp.(Goto *et al.*, 2000).

In order to check the specifity of the amplification by the used primers, gel electrophoresis was carried out as shown in figure (3.2) which revealed single amplicon band with proximal size 300 bp while the other band of about 100bp size that appeared reflect primers polymerization.

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(Figure 3.2) Agarose gel electrophoresis for a PCR amplification of 16S rDNA by the set of primers (B16SF and B16SR) of the *Bacillus* isolates on (0.8%) agarose gel for 15 minutes, 72 voltage.

Lane (1) : *Bacillus* B10, Lane (2): *Bacillus* B9, Lane (3): *Bacillus* B8, Lane (4): *Bacillus* B7, Lane (5): *Bacillus* B6, Lane (6): *Bacillus* B5, Lane (7): *Bacillus* B4, Lane (8): *Bacillus* B3, Lane (9): *Bacillus* B2, Lane (10): *Bacillus* B1, Lane (11): 2Kb DNA landmark ladder. PCR products were purified and sequenced as illustrated in figure (3-3) and (appendices 1-9) which showed the complete nucleotide sequence of the HV region of the 16S rDNA gene.

GAATGGGAGCTTGCTCCCTGATGTTAGCGGCGGACGGGTGAGTAACAC GTGGGTAACCTGCCTGTAAGACTGGGATAACTCCGGGAAACCGGGGCT AATACCGGATGCTTGTTTGAACCGCATGGTTCAAACATAAAAGGTGGC TTCGGCTACCACTTACAGATGGACCCGCGGCGCATTAGCTAGTTGGTGA GGTAACGGCTCACCAAGGCAACGACGCCCAGACTCCTACGGGAGGGTG

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Isolates identities were established by comparing the sequences obtained with the NCBI data base, results showed that all the selected isolates were belong to the *Bacillus* species with (99- 100) % similarities with query coverage (92-93) % ,such results were also obtained by Goto *et al.*( 2000) who revealed that this (HV) region was highly conserved within the species.

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Also it was found that as shown in table (3-4) the partial 16S rRNA gene sequences of all isolates show sequence similarity to certain *Bacillus* strains mostly to *B. subtilis*.

One other set of primers was used (27F and1492R) which are universal primers for amplifying complete 16S rRNA gene of *Bacillus* to identify the

selected isolates to the strain level (Akhmaloka *et al.*, 2006 ;Thomas, 2004; 2006 ; Cerritos *et al.*,2008).

PCR products for the selected isolates were observed on agarose gel electrophoresis. The result showed an amplicon band for each one with the size of 1.5 kb respectively (Figure 3.4).



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(Figure 3.4) Agarose gel electrophoresis for a PCR amplification of 16S rDNA by the set of primers (27F and1492R) of the *Bacillus* isolates on (0.8%) agarose gel for 15 minutes, 72 voltage. Lane (1) : 2Kb DNA landmark ladder, Lane (2): *Bacillus* B10 isolate , Lane (3): *Bacillus* B9, Lane (4): *Bacillus* B8 , Lane (5): *Bacillus* B7, Lane (6): *Bacillus* B6 , Lane (7): *Bacillus* B5, Lane (8): *Bacillus* B4 , Lane (9): *Bacillus* B3 , Lane (10): *Bacillus* B2, Lane (11): *Bacillus* B1.

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Results from sequencing (Appendices10-11) showed the full length of the specified piece of 16S rDNA gene for two isolates (B6 and B9) while the other eight isolates were partially sequenced due to loop structure in the amplified amplicon so molecular analysis was done by using partial 16S rDNA sequences from the 5' (632bases) and 3' ends (725 bases) of first strand to these isolate, such results were also obtained by Thomas (2006).

Alignment of these sequences with NCBI data base revealed that strain B6 show sequence similarity 100% to *B. subtilis* (Genebank accession no. AB526464.1 and AB383135.1) with query coverage 100% and with the higher score, other strains show similarities range (96-100) % with (41) *B. subtilis*,

(7) Bacillus mojavensis,(2) Bacillus amyloliquefaciens, (2) Bacillus axarquiensis,

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l of these species are members of the *Bacillus* sub-

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and Bacillus sonorensis. (Fritz ,2004).

Other sequences of the rest studied isolates also show the greatest sequence similarity to a large number of *B. subtilis* as shown in table (3-4) in comparing with other *Bacillus* spp.

Lee *et al.*, (2007) depended partial sequencing of the 16S rRNA gene and partial *gyrA* gene to identify a strain of *B. amyloliquefaciens*, they found that the partial 16S rRNA gene of *B. amyloliquefaciens* LP03 exhibited over 98% homology with most *Bacillus* species, but the partial *gyrA* gene showed greater homology with *B. amyloliquefaciens* (95%) than other *Bacillus* species (less than 83%). While Joshi *et al.* (2008) got 98% sequence similarity to *B. subtilis* as per

NCBI, BLAST (Genbank accession no. DQ 922951) when align the partial 16S rRNA gene of the strain *B. subtilis* 20B which was already identified by biochemical tests as *B. subtilis*.

#### 3.2 Selection of the best biosurfactant production media

In order to select the best production media for biosurfactant production by the selected isolates, two media were used; E medium and Jacques Medium (Youseef *et al.*,2004; Jacques *et al.*,1999).

Results shown in table (3-5) illustrated that Jacques medium was the best one for biosurfactant production compared to E medium since a high percentage of

owering surface tension by produced biosurfactant was achieved.

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#### two production media.

Symbol of Isolates	Percentage of lowering (ST) in Jacques medium (%)	Percentage of lowering surface tension in E medium (%)
B1	43.3	29.10
B2	42.3	29.54
B3	41.5	29.72
B4	46.2	28.60
B5	44.6	29.55
B6	43.5	29.56
B7	43.7	29.55
B8	45.0	28.40
B9	42.5	30.00
B10	42.1	29.20
#### **3.3 Optimum period for biosurfactant production**

The selected B. subtilis isolates (B1, B2, B3, B4, B5, B6, B7, B8, B9, B10) were propagated in Jacques medium for different periods of time in order to determine the production time for biosurfactant production from each isolate.

Results from figure (3-5) revealed that maximum yields (g/l) of the surface active agents produced by B. subtilis isolates B1, B2, B6, B7, B9 and B10 were achieved after 48 hours of incubation, followed by drop in the yield which may result from the consumption of biosurfactant as a carbon source after depletion of nutrients in the broth (Shaligram and Singhal, 2010). It was also found that surfactin acts as end product inhibitor which lead to lower yield (Drouin and

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Figure (3-5) Dry weight of biosurfactant produced by *Bacillus subtilis* isolates propagated in Jacques medium for different incubation time.

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#### **3.4 Purification of biosurfactant produced by the selected isolates**

Combination of techniques were used to purify biosurfactants produced by the ten selected *B. subtilis* isolates , beginning with acid precipitation of biosurfactant in cell – free supernatant to crystallize biosurfactant since such compounds tends to become insoluble at low pH due to charge neutralization and protonation of carboxylic acid side chains of aspartic or glutamic acids in the peptide portions of these molecules (Maneerat and Phetrong, 2007).

The obtained brown precipitates were solubilized in alkaline D.W to retain their surface active lowering activity followed by lyophylization and weighing to calculate the primary yield of biosurfactant (g/L of broth) as shown in table (3-6).

These results were inagreement with Cooper et a.l (1981) who found that

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surface tension from 27 mN/m to 62 but when neutralized and the

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Table (3-6) Weight of partially purified biosurfactant from the selected

*Bacillus* isolates after cultivation in Jacques medium for the optimum period of incubation.

Isolates Symbol	Incubation periods (hrs.)	Partially purified biosurfactant (g/l)
B1	48	2.050
B2	48	3.050
B3	72	1.625
B4	72	3.600
B5	72	1.425
B6	48	1.725
B7	48	1.250
B8	72	1.875
B9	48	1.025
B10	48	1.100

For further purification, a two-step membrane ultrafiltration process was performed for each sample; the first step to remove low molecular weight contaminant, while the high molecular contaminant were removed during the second step of ultrafiltration ((Lin *et al*, 1998a, Isa *et al.*,2007).

Results in table (3-7) showed the yield of purified biosurfactant, Chen *et al.* (2007b) got 97% recovery of the initial concentration of surfactin by this method.

Table (3-7) Weight of purified biosurfactant from the selected *B. subtilis* isolates by two steps ultrafiltration method.

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	B7			
	<b>B</b> 9		0.125	
	B10		0.150	
	B8 B9 B10		0.100 0.125 0.150	

#### **3.5** Characterization of purified biosurfactant

#### **3.5.1** By thin layer chromatography (TLC)

Characterization of biosurfactants produced by the ten selected *B. subtilis* isolates was done by TLC analysis which was frequently employed for the characterization of biosurfactants (Javaheri *et al.*, 1985; McInerney *et al.*, 1990).

Each sample of purified biosurfactant was dissolved in absolute methanol and analyzed on silica gel plates by using chloroform-methanol-water (65:25:4v/v/v) as the mobile phase.

Thin-layer chromatography of the purified biosurfactant and standard surfactin revealed white spots when the plate was sprayed with water as indication of hydrophilic compound (Figures (3-6) and (3-7)). Same spots were appeared with the rhodamine B when examined under UV light, indicating the presence of lipids in the compounds. As a conclusion, purified biosurfactants were belonged to the lipopeptide family.

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Figure (3-6) TLC analysis of biosurfactants produced by *B. subtilis* isolates (B1,B2),st: standard surfactin

Results of table (3- 8) showed that the relative mobility (Rm) of standard surfactin was (0.6) with single spot. Regarding the isolates, purified biosurfactants from isolates (B1, B2) were separated into three and four spots, resprctively, and biosurfactants from isolates (B3.B7,B8,B5) were separated into two spots, while

(B4,B6,B9,B10) were separated into a single spot. However, almost all isolates were capable to produce one spot that align with the spot of standard surfactin , indicating that all isolates were capable in producing surfactin. This fact was also supported by the haemolytic activity of all the selected isolates, a characteristic of the surfactin (Moran *et al.*, 2002).



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#### B5,B7)

Separation of other spots indicated the presence of different species or isoforms of biosurfactants which were detected in many strains of *Bacillus* such as *B. subtilis* complex BC1212 that produced four surfactin isoforms (A,B,C and D) differed slightly in their physiological properties (Kim *et al.*, 2006). Sandrin *et al.* (1990) isolated *B. subtilis* S 499 strain that produced surfactin and antifungal lipopeptides belonged to the iturin family.

Table (3-8) Relative mobility of the purified surface active agents produced by the *B. subtilis* isolates (B1,B2,B3,B4,B5,B6,B7,B8,B9,B10) after TLC Separation .

sample	Number of spots	Relative mobility (Rm )
Standard surfactin	1	0.6
B1	3	0.4,0.5, 0.6
B2	4	0.1,0.4, 0.5, 0.6
B3	2	0.1,0.6
B4	1	0.6
B5	2	0.1, 0.6
B6	1	0.6
B7	2	0.1,0.6

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followed by ms/ms analysis to study the amino acids sequence in the purified biosurfactants.

Under same purification conditions, the purified biosurfactant produced by the isolate *B. subtilis* B6 were resolved into main five fractions on HPLC with the same retention time as standard surfactin (figures (3-8) and (3-9)).

Separation of standard surfactin into several fractions was due to the fact that natural surfactin produced by *B. subtilis* was a mixture of isoforms with slightly different properties as a result of substitutions in amino acids and the aliphatic chain (Grangemard *et al.*, 1997).



Figure (3-8) HPLC spectrogram of standard surfactin from Sigma-Aldrich

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Figure (3–9) HPLC spectrogram of the purified lipopeptide produced by *B*. *subtilis* B6.

In order to compare the molecular mass of standard surfactin and the purified lipopeptide, Ion extracted chromatogram (IEC) of fractions retained in the HPLC were studied by HPLC- mass .

Results of table (3-9) and chromatograms in appendices (21-30) showed that both standard surfactin and purified lipopeptide had the same molecular weight.

Retantion time (minutes) of peaks of standard surfactin	Retantion time (minutes) of peaks of purified lipopeptide	<i>m/z</i> [M+H] <sup>+</sup>	<i>m/z</i> [M+ Na] <sup>+</sup>	Molecular weight (Dalton)
8.4	8.4	995	1017	993.92
9.9	9.8	1009	1031	1007.9
13	12.8	1023	1045	1021.9
16.9	16.6	1037	1059	1035.9

Table (3-9) Results of IEC of the standard surfactin and purified lipopeptide .

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nolecules represent the ionized fragments so the analysis of those peaks would

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give important information about the connecting relationship of the peptide since the difference between any two peaks was the mass of the lost fragments so the differences can be used to determine the connection of some amino acid residue in a peptide chain (Wang *et al.*,2007 ; Cao *et al.*,2009b).

So each peak in table (3-6) was further fragmented by MS2 which generating an identifying mass spectrum for each component in the mixture.

The mass spectrum of  $[M+Na]^+$  at m/z 1059 (figure 3-10) showed the connection of the following amino acids Leu -Leu -Asp –Val ( as a result of peak

 $1059.3 \rightarrow 945.7 \rightarrow 833.6 \rightarrow 718.5 \rightarrow 619$ ) while the peaks (707.5  $\rightarrow 594.5 \rightarrow 481.4 \rightarrow 365.4$ ) suppose the connection of Leu- Leu –Asp.

In addition, according to the mechanism of (double hydrogen transfer), the C-terminus of the peptide was formed at peak 800.6 which represent the loss of Leu-Leu-OH<sub>2</sub>, which suggests that a leucine residue serves at the C-terminus of the peptide chain (Yang *et al.*,2006).



revealed the same amino acids connection for the studied lipopeptide as shown in appendices (31-34).

#### 3.6 Optimization of biosurfactant production by isolate B. subtilis B6

#### 3.6.1 Effect of carbon source

Carbon source represents an important factor in biosurfactant production; therefore, several kinds of carbon sources were tested to optimize the production of biosurfactant.



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sources Sucose and sucrose) were the best substrates for synthesis of

biosurfactant in reducing the surface tension (ST). Although glucose was better than sucrose in increasing the surfactin yield, the later was better in enhancing the percentage of (ST) reduction.

Addition of water insoluble carbon source (Olive oil, Corn oil, machine oil) didn't support the production of biosurfactant.

This characteristic for the carbon sources utilization was in accordance with Sandrin *et al.*, (1990) and Kim's *et al*, (1997) results that found *B. subtilis* prefer the water-soluble carbon sources especially glucose, fructose and sucrose as carbon sources to produce biosurfactants while, addition of the hydrocarbons to culture

medium completely inhibited surfactin production by *B. subtilis* (Davis *et al.*,1999).

Makkar and Cameotra (2002) found that *Bacillus subtilis* MTCC 2423 was unable to utilize sodium acetate as a carbon source for biosurfactant production, while Liu *et al.*, (2009) found that in medium containing olive oil, *B. subtilis* can grow and produce biosurfactant but was unable to lower the (ST). In contrast, glucose, sucrose and sodium supported the production of biosurfactant

When the biomass and biosurfactant production by *B. subtilis B6* grown on different carbon sources were compared, great variations in the O.D. of cells and concentration of lipopeptides were recorded. Maximum O.D. values were

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ly O.D. values (0.66,0.61 and 2.47), respectivel

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specially glucose.

As a consequence of these results sucrose was subsequently used.

#### 3.6.2 Effect of carbon source concentration

Different concentrations (0.5, 1, 2, 3, 4, 5%) of sucrose after was chosen as the best carbon source for surfactin production ,were used to propagate *B. subtilis* B6 to determine the optimum concentration.



Figure (3-12) Effect of sucrose (as carbon source) concentration on biosurfactant production from *B. subtilis* B6, grown at 30 °C in a shaker incubator (180 rpm) for This is a watermark for the trial version, register to get the full one!

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increased over 2%. This may be due to the toxic effect of sucrose on the growth of

producing isolates *Bacillus* B6, since the O.D of the bacterial growth dropped from 1.6 at 1% sucrose concentration to 1.3 at 2%.

Regarding lowering ST, highest percentage was obtained at sucrose concentration of (1%), while higher or lower concentrations caused decreased in the percentage.

These results were in accordance with Hartoto and Mangunwidjaja (2002) who studied the importance of sucrose and its concentration in surfactin production by *Bacillus sp*.BMN14 in a shaker flask. They found that an optimum surfactin production with lower ST and higher bacterial mass was reached at sucrose

concentration of 2%, but dropped when the concentrations increased or decreased from that.

#### 3.6.3 Effect of nitrogen source

Biosurfactant production was depending on the nitrogen source used in the fermentation. As seen from figure (3-13), inorganic nitrogen sources were more efficient in enhancing the isolate for production of biosurfactant than organic sources. Also, it was observed that among the inorganic nitrogen sources, potassium nitrate was the best as it caused the production of highest yield of biosurfactant (0.6 g/l) along with highest reduction in ST (54%), while ammonium nitrate and sodium nitrate were less efficient in producing biosurfactant.

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(4.051,4.018,2.045,1.69,1.67).

As a consequence of these results potassium nitrate was chosen for next experiments.

These results were in agreement with Makkar and Cameotra (2002) ) who noted potassium nitrate salts as a preferable nitrogen source for the biosurfactant production by *B. subtilis* MTCC 2423. However, Haddad *et al.* (2009) found itammonium sulphate ,while, Abushady *et al.*(2005) mentioned ammonium nitrate, and Ghribi and Ellouze-Chaabouni (In press ) mentioned urea as the best nitrogen source by different isolates of *B. subtilis*, furthermore, another group (Wu *et al.*,2007) use a mixture of two nitrogen sources ( ammonium sulphate :ammonium nitrate) in a ratio (1:2) to produce biosurfactant by *Bacillus* spp.



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#### **3.6.4 Effect of nitrogen concentration**

The effect of different concentrations of potassium nitrate on biosurfactant production which was chosen as the best nitrogen source was studied.

Results illustrated in figure (3-14) showed that the optimum concentration was 0.3% for both biosurfactant yield and lowering ST, which drop dramatically above this concentration while, lower concentration had lower effect on biosurfactant production .



Figure (3-14) Effect of potassium nitrate (as a nitrogen source) concentration on biosurfactant production from *B. subtilis* B6 grown with 1% sucrose at 30 °C in aThis is a watermark for the trial version, register to get the full one!

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concentration in all studied Bacillus strains.

#### 3.6.5 Effect of phosphate concentration

Effect of phosphate concentration on biosurfactant production was studied.

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Results in figure (3-15) showed that optimum concentration was 1g/l for biosurfactant production which decreased when phosphate concentration change above or below the optimum concentration.



Figure (3-15) Effect of KH<sub>2</sub>PO<sub>4</sub> concentration on biosurfactant production from

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incubator (180 rpm) for 72 hrs.

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#### 5g/100ml) high yield was obtained ,One explanation for the reduced surfactin

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biosynthesis at high concentration is that phosphate repression might play a role in the regulation of surfactin production, which is consistent with the negative effect of easy utilizable phosphate source on the biosynthesis of antibiotics and other secondary metabolites at the level of transcription (Liras *et al.*,1990).

While in some *B. subtilis* strains, high phosphate concentration was demanded to increase the yield of biosurfactant as found by Kim *et al.* (1997), *B. subtilis* C9 required as much as 13.5%  $K_2HPO_4$  for production of high yield biosurfactant.

#### 3.6.6 Effect of pH

The pH of medium plays an important role in biosurfactant production, so to different pH values the production media was adjusted.

Results from figure (3-16) revealed that the optimum pH for biosurfactant production was 8 and the yield drop when pH decreased or increased from that value.



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Figure (3-16) Effect of pH on biosurfactant production from *B. subtilis* B6, grown with 1% sucrose, 0.3% KNO<sub>3</sub> and 1g/L K<sub>2</sub>HPO<sub>4</sub> at 30 °C in a shaker incubator (180 rpm) for 72 hrs.

These results were in accordance with that of a study conducted on *B*. *licheniformis* byAli *et al.* (2010) and Huszcza and Burczyk (2003) when they studied the biosurfactant production by *B. coagulanse*.

Sepahy *et al.* (2005) found that very low yield of surfactin was recorded when the production medium of *B. subtilis* was adjusted at pH values lower than 6.5, and the maximum yields were achieved at a pH range of (6.5- 8.5).

Makkar and Cameotra (2002) found that *B. subtilis* strain MTCC 2423 was able to produce biosurfactant in a pH range of (4.5- 10.5), although the maximal yield of the biosurfactant was obtained at pH 7.0 (, 2002).

#### **3.6.7 Effect of temperature**

Different temperatures of incubation were investigated to study their effect on biosurfactant production.

The optimal incubation temperature was  $30^{\circ}$ C for biosurfactant production as illustrated in figure (3-17) followed by 37 °C and dropped significantly at 40°C and 28°C while the OD. of the bacterial growth was higher at 30°C followed by

 $37^{\circ}$ C then  $28^{\circ}$ C, with less value at  $40^{\circ}$ C as follows (4.1, 3.5, 3.1, 2.0)

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o studied the effect of temperature on biosurfactant

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above 40 °C (Makkar and Cameotra, 2002; Sepahy *et al.*, 2005).

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#### 3.6.8 Effect of incubation period

The surfactin production medium was inoculated with *B.subtilis* B6 and incubated for different incubation periods from 12- 96 hrs, figure (3-18) show that surfactin concentration increased with the increased in incubation period up to 72 hrs but surfactin concentration was decreased with the other incubation periods.

Also it was observed that reduction in ST was not highly effected by the incubation period.



Figure (3-17) Effect of temperature on the growth and biosurfactant production This is a watermark for the trial version, register to get the full one!

in a shaker incubator (180 rpm) for 7. Benefits for registered users:

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Fig (3-18) Effect of incubation period on biosurfactant production from *B. subtilis* B6, grown with 1% sucrose, 0.3% KNO<sub>3</sub> and 1g/L K<sub>2</sub>HPO<sub>4</sub>, pH 8, at 30  $^{\circ}$ C in a shaker incubator (180 rpm) for 72 hrs.

In this regard Abushadi *et al.* (2005) found that there was no biosurfactant production before 24 hrs,. but it increased until reached its maximum at 72 hrs. Above that, no further increases in the concentration was observed.

Ali *et al.* (2010) repoted that best incubation time for a strain of *B. licheniformis* was 72 hrs of incubation and biosurfactant production was decreased in other incubation periods used.

#### 3.7 Biological activity of purified biosurfactants

#### 3.7.1 The cytotoxicity assay by MTT kit

In order to evaluate the potential antitumor activity of the purified lipopeptide from *B. subtilis* isolates (B1,B2,B3,B4,B5,B6,B7,B8,B9,B10) in vitro.

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centrations of these lipopeptide samples for 24, 4

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regultant offect for each cell line was explained as the following:

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#### A. Cytotoxic effect on K562 cell line.

The results in tables (3-11),(3-12) and(3-13) revealed that lipopeptide from *B. subtilis* isolates (B2,B3,B4,) showed low effect on the proliferation of human leukemia (K562) cell line during all periods of exposures, while lipopeptide produced by *B. subtilis* isolates (B1,B5,B6,B7,B8,B9,B10) had cytotoxic effect on K562 proliferation as were shown in tables (3-10), (3-14), (3-15),(3-16), (3-17),(3-18) and (3,19) with high significant inhibition rate at highest concentrations which were being decreased gradually as concentrations decreased too..

These results were in accordance with Cao *et al.* (2009a) results which showed that lipopeptide produced by *Bacillus* natto TK-1induce significant reduction in K562 cell line viability in a dose dependent manner at 48 hrs. of exposure.

It was observed from tables (3-15), (3-16) and (3-18) that there were no significant differences at (P<0.05) between inhibition rate when lipopeptide concentrations 240mg/l and 200 mg/l were used from *B. subtilis* isolates (B6,B7,B9).

Table (3-10) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* 

#### B1 on K562 cell line.

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160  $97 \pm 0.65$ 19.625 ±4.589  $2.323 \pm 2.65$ D.a B.b C,a 80 5.58 ±0.535  $13.167 \pm 3.495$  $7.753 \pm 0.645$ E,a C,a D,a 40  $0.15 \pm 0.05683$  $0.192 \pm 0.118$  $0.119 \pm 0.019$ E.a C,a D.b  $0.172 \pm 0.09678$ 20  $0.11903 \pm 0.09119$  $0.3424 \pm 0.03704$ 

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Table (3-11) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B2 on K562 cell line.

Exposure time	Inhibition rate% (mean ±SD)				
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.		
240	A,a	A,b	A,c		
	0.24740±0.09787	0.0975±0.1147	0.11613±0.07716		
200	A,a	A,b	A,c		
	0.31703±0.05919	0.1468±0.1274	0.08280±0.05631		

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The effect of exposure time was also studied, it was observed that there was

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Generally it was observed that inhibition rate was increased gradually with

prolonged incubation period till 72 hrs. ,however , with prolonged incubation, lipopeptide from *B. subtilis* isolates (B1,B5,B6,B8,B10) became less inhibitor for malignant cell proliferation ,this may be due to the development of resistance cells since it was found that some malignant cells can either had repair system that became activated to overcome the damage done by the antitumor agent (Ledzewicz , 2006) or may be the antitumor agent was susceptible to ABC transport proteins which remove molecules out of the cell, thus an over –expression by gene amplification for example in these molecules is an important mechanism for resistance to various drugs and antitumor agents (Mao and Unadkat, 2005).

Malignant cancer cell populations were highly heterogeneous - the number of genetic errors present within one cancer cell lies in the thousands and fast duplications combined with genetic instabilities provided just one of several mechanisms which allow for quickly developing acquired resistance to anti-cancer drugs (Loeb,2001).

When all the results of cytotoxic effect exhibited by lipopeptide from *B. subtilis* isolates (B1,B5,B6,B7,B8,B9) on K562 cell line were compared together, *B.* B7, was found to be the most effective with IC50 at 80mg/l for 72 hrs. of exposure.

While lipopeptide from isolates (B1, B6, B9) and (B5, B8) induce IC50 at

#### 160 mg/l and 200mg/l respectively

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(wang *et al.*,2009).

Surfactin has also been reported to have an antitumor activity against Ehrlich's ascite carcinoma cells (Kameda *et al.*, 1974), another study on the effect of surfactin on the proliferation of a human colon carcinoma cell line showed that surfactin strongly blocked the cell proliferation which was due to apoptosis induction and cell cycle arrest via the suppression of cell survival regulating signals such as ERK and PI3K/Akt (Kim *et al.*,2007).

Table (3-12) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B3 on K562 cell line.

Exposure time	Inhibition rate% (mean ±SD)			
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.	
240	A,a	A,a	A,a	
	0.7117±0.5187	0.1762±0.2045	0.1223±0.0558	
200	A,a	A, a	A,a	
	0.7±0.5456	0.1658±0.1254	0.1148±0.1655	

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#### Lable (3-13) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B4

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Exposure time	Inhibition rate% (mean ±SD)			
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.	
240	A,a	A,b	A,c	
	0.24410±0.05604	0.10483±0.08625	0.6378±0.1081	
200	A,a	A,b	A,c	
	0.28580±0.04655	0.19417±0.06576	0.5567±0.1369	

Different upper case letters: significant difference (P<0.05) between means of rows Different lower case letters: significant difference (P<0.05) between means of columns Table (3-14) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B5 on K562 cell line.

Exposure time	Inhibition rate% (mean ±SD)			
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.	
	A,a	A,a	A,a	
240	$83.12 \pm 5.85$	91.307 ±3.36	85.27 ±2.29	
	B,a	B,b	B,b	
200	$25.993 \pm 6.655$	75.42 ±4.958	$66.26 \pm 8.793$	
	C,a	C,b	C,a	
160	$0.183 \pm 0.08904$	17.547 ±3.697	$1.450 \pm 0.268$	
	C,a	C,b	C,a	
80	0.2752 ±0.13912	$10.513 \pm 3.177$	$0.1725 \pm 0.03128$	
	C,a	D,b	C,a	

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Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,b	A,b
240	73.58±3.758	92.29±1.066	84.113±0.357
	B,a	A,b	A,b
200	64.1±3.09	89.597±1.974	84.823±1.372
	C,a	B,b	B,b
160	38.3±0.6	77.177±6.121	68.36±6.05
	D,a	C,b	C,ab
80	8.547±2.021	20.493±2.532	13.714±3.293
	E,a	D,b	D,b
40	$0.0908 \pm 0.03904$	7.097±0.616	3.377±3.129
	E,a	E,a	D,a
20	$0.06347 \pm 0.05484$	0.1315±0.00537	0.09243±0.0479

#### nhibition rate% (mean ±SD)

Different upper case letters: significant difference (P<0.05) between means of rows Different lower case letters: significant difference (P<0.05) between means of columns Table (3-16) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B7 on K562 cell line.

Exposure time	Inhibition rate% (mean ±SD)			
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.	
	A,a	A,a	A,a	
240	85.99 ±0.54	90.84 ±4.96	90.94 ±0.253	
	B,a	A,a	A,a	
200	$77.15 \pm 5.996$	87.31 ±5.44	$86.87 \pm 3.626$	
	C,a	B,a	B,b	
160	$26.443 \pm 3.301$	30.273 ±3.091	$70.907 \pm 2.057$	
	D,a	C,a	C,b	
80	7.813 ±4.033	$6.53 \pm 5.65$	57.17 ±1.254	
	DE,a	C,a	D,a	

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Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,b	A,a
240	79.103±2.96	89.86±1.65	82.11±0.139
	B,a	B,a	B,a
200	$40.42 \pm 8.08$	61.61±2.99	50.49±0.6
	C,a	C,b	C,a
160	0.023±0.008	11.797±3.939	$1.581 \pm 0.648$
	C,a	D,b	C,a
80	0.908±0.119	4.467±1.569	$0.2889 \pm 0.1182$
	C,a	D,a	C,a
40	$0.130 \pm 0.008$	0.19413±0.02441	0.152±0.1685
	C,a	D,a	C,a
20	$0.0897 \pm 0.04753$	$0.05697 \pm 0.05932$	$0.0187 \pm 0.0251$

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Table (3-18) Cytotoxic effect of the purified lipopeptide produced by B. sub	tilis B9
on K562 cell line.	

Exposure time	Inhibition rate% (mean ±SD)		
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,a	A,b
240	71.58±1.45	70.723±1.397	87.52±4.04
	B,a	A,a	B,b
200	60.945±6.739	65.873±2.079	80.91±2.935
	C,a	B,a	C,b
160	48.307±4.159	48.035±3.939	63.58±2.01
	D,a	C,b	D,c
80	4.133±0.531	$10.860 \pm 3.762$	23.07±0.762
	D,a	D,b	E,b
40	2.526±0.508	7.460+1.81	9.467±2.207

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time	minution rate 78 (mean ±SD)		
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,b	A,c
240	$17.54 \pm 2.89$	$55.865 \pm 1.6242$	$28.75 \pm 0.22$
	B,a	B,b	B,b
200	$0.23003 \pm 0.04322$	21.23±3.23	$14.57 \pm 0.28$

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#### B. Cytotoxic effect on L1210 cell line.

The effect of the purified lipopeptide on malignant animal cell line proliferation was studied through treating mice leukemia (L1210) cell line with different concentrations of lipopeptides produced by *B. subtilis* isolates (B1, B2, B3, B4, B5, B6, B7, B8, B9, B10) for 24, 48, 72 hrs.

Results illustrated in tables (3-21) and (3,23) revealed that lipopeptides from isolates *B*. (B2, B4) were ineffective in inhibition the growth rate of the tested cell line for all the exposure times while lipopeptides from *B. subtilis* isolates (B1,B3, B5,B6, B7, B8, B9, B10) inhibited the growth rate in a dose dependent manner as it increased towards the higher concentrations with significant differences at (P<0.05) as illustrated in tables (3-20), (3-22), (3-24), (3-25), (3-26), (3-27), (3-29).

Highest inhibition effect by *B. subtilis* isolates (B1,B3,B5,B6,B7,B8,B9)

was achieved 48 hrs. of exposure as were shown in appendices (36, 42, 45, 49,

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Concentrations	24 urs.	40 urs.	72 nrs.
mg/l			
	A,a	A,a	A,b
240	$87.68 \pm 1.64$	93.74 ±0.12	54.14 ±8.32
	B,a	A,b	AB,c
200	69.297 ±8.831	$88.72 \pm 1.98$	45.57 ±4.1079
	C,a	B,b	B,a
160	$44.24{\pm}1.834$	52.913 ±7.437	$37.92 \pm 1.133$
	D,a	C,a	C,a
80	$1.153 \pm 0.508$	2.123 ±0.799	1.857 ±0.4397
	D,a	C,a	C,a
40	$0.009 \pm 0.00052$	0.0143 ±0.02181	0.0201±0.00469
	D,a	C,a	C,a
20	$0.00703 \pm 0.00431$	$0.01463 \pm 0.01488$	$0.03873 \pm 0.03791$

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Table (3-21) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B2 on L1210 cell line.

Exposure time	Inhibition rate% (mean ±SD)		
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
240	A,a	A,b	A,c
	0.07353±0.02832	0.1681± 0.1104	0.6039±0.2718
200	A,a	B,b	A,a
	0.15283±0.03682	0.7993±0.1078	0.1171±0.0560

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#### Innutuon rate /0 (mean ±51) time Concentrations 24 hrs. 48 hrs. 72 hrs. mg/l A.a A.a A.b 60.113 ±8.887 240 $72.338 \pm 5.8$ $21.68 \pm 13.025$ B.a B.b A.a 200 1.480±0.156 6.765±1.888 $1.5 \pm 0.427$

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# troduced by *R* subtilis P2

Table (3-23) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B4 on L1210 cell line.

Exposure time	Inhibition rate% (mean ±SD)		
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,b	A,c
240	11.09 ±4.401	32.18 ±7.99	$0.3866 \pm 0.4223$
	A,a	B,a	A,b
200	$6.724 \pm 5.608$	6.61 ±2.669	$0.1612 \pm 0.1182$

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Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,a	A,b
240	84.09±6.886	86.63±2.498	69.35 ±13.18
	B,a	B,b	B,c
200	59.383 ±4.211	75.46 ±3.317	45.57 ±8.96
	C,a	C,a	B,a
160	36.748±2.539	37.55 ±6.11	34.087±3.301
	D,a	D,b	C,a
80	1.003 ±0.718	19.68 ±2.041	$0.887 \pm 0.105$
	D,a	E,b	C,a
40	0.174±0.17	1.753±0.767	0.329±0.103
	D,a	E,b	C,a
20	0.02±0.02	$1.032 \pm 1.429$	$0.014 \pm 0.008$

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Table (3-25) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B6 on L1210 cell line.

Exposure time	Inhibition rate% (mean ±SD)		
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,b	A,b
240	$71.685 \pm 5.805$	$88.51 \pm 3.5$	$76.86 \pm 4.8624$
	B,a	B,b	B,a
200	51.84 ±7.44	$82.62 \pm 0.44$	$61.97 \pm 7.894$
	C,a	C,b	B,c
160	37.717 ±5.958	$72.273 \pm 1.945$	53.813±2.564
	D,a	D,b	C,c
80	3.86±0.255	24.717 ±1.168	$11.955 \pm 2.898$
	D,a	D,b	D,a

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Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,b	A,a
240	$80.38 \pm 2.84$	$91.22 \pm 1.32$	79.71 ±2.1905
	B,a	B,b	B,c
200	$68.163 \pm 1.709$	80.375 ±2.114	46.33±6.675
	C,a	C,b	B,c
160	43.333 ±4.748	58.277 ±2.33	31.097 ±8.815
	D,a	D,a	C,b
80	2.415±0.53	2.875±0.813	$0.495 \pm 0.49$
	D,a	D,a	C,a
40	$0.143 \pm 0.078$	0.272±0.224	0.741±0.192
	D,a	D,a	C,a
20	$0.0254 \pm 0.0021$	0.0199 ±0.01345	$0.06575 \pm 0.00247$

Different upper case letters: significant difference (P<0.05) between means of rows Different lower case letters: significant difference (P<0.05) between means of columns Table (3-27) Cytotoxic effect of the purified lipopeptide produced by B. subtilis B8 on L1210 cell line.

Exposure time	Inhibition rate% (mean ±SD)		
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,b	A,b
240	39.35 ±1.81	84 ±5.16	71.33 ±8.567
	B,a	B,b	B,c
200	32.97 ±0.575	73.8 ±2.1	48.36 ±9.7
	B,a	B,b	C,b
160	14.34±4.257	24.826±1.127	20.173±1.728
	D,a	D,b	D,a
80	4.285±1.676	13.863±2.734	3.223±1.011
	D,a	E,b	D,a

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Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,b	A,a
240	86.457±1.662	92.9±0.42	$87.36 \pm 1.261$
	A,a	A,b	A,a
200	$85.24 \pm 1.869$	92.04±1.129	84.323 ±1.312
	B,a	A,b	A,b
160	$55.08 \pm 2.855$	$86.76 \pm 0.95$	$87.12 \pm 0.562$
	C,a	B,b	B,b
80	1.477±0.376	$58.62 \pm 0.666$	50.15 ±1.463
	C,a	C,b	C,a
40	1.063±0.501	5.593 ±0.5391	2.518±1.293
	C,a,	C,b	C,c
20	$0.01813 \pm 0.003$	1.651±0.257	$0.495 \pm 0.075$

Different upper case letters: significant difference (P<0.05) between means of rows Different lower case letters: significant difference (P<0.05) between means of columns Table (4-29) Cytotoxic effect of the purified lipopeptide produced by *Bacillus* B10 on L1210 cell line.

Exposure time	Inhibition rate% (mean ±SD)		
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,b	A,b
240	$33.75 \pm 1.89$	79.7 ±4.97	74.59 ±9.23
	B,a	B,a	B,a
200	$8.88 \pm 3.78$	$17.563 \pm 5.58$	$6.34 \pm 1.03$

Different upper case letters: significant difference (P<0.05) between means of rows Different lower case letters: significant difference (P<0.05) between means of columns

Comparison among the inhibition effect of all tested lipopeptides on L1210

cell line showed that lipopeptide from isolate *B*. B9 was the most effective one followed by *B. subtilis* isolates (B1, B6, B7).

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Results from tables (3-31) and (3-33) showed that lipopeptide from the two

*B.* isolates B2 and B4 had very low effect on the cell line proliferation, this may due to the fact that several surfactin isoforms usually coexist in the cell as a mixture of several peptidice variants (Kowall *et al.*, 1998; Bonmatin *et al.*,2003) with a different aliphatic chain length (Hue *et al.*, 2001) that differ in their biological activity (Seydlová and Svobodová, 2008).

The results from tables (3-30) - (3-39) showed that cell survival in Hep-2 treated cultures were progressively decreased with increasing the concentration and it was time dependent as increased with prolonged incubation this may explained

by the release of intracellurar components from dead cells that induce more cells to die as a results of toxicity increasing .

Cao *et al.*, (2009) found that surfactin from *B. natto* TK-1 was dose and time dependent in inhibiting the proliferation of human breast cancer cell line (MCF-7).

However, statistically , there were no significant differences (P<0.05) between the inhibition rate when lipopeptides from *B. subtilis* isolates (B1,B3, B6,B7,B8,B9) were used at concentrations 240 mg/l and 200 mg/l (dose independent) or between the two exposure times 48 and 72 hrs. of same concentrations as showed in tables (3-30), (3-32), (3-35), (3-36), (3-37) and (3-38).

This may due to the heterogeneity within the tumor cell line which lead to

the rising of drug resistant cells that will multiply and remain as a surviving cells

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The previous results showed that inhibition rate depends

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lular receptors in each type of cell lines; making the cells interacts at same

concentration in different manners. Moreover the metabolic pathways in response to each treatment differed from one line to another(Kim *et al.*, 2007). This fact was mentioned in different studies which investigated at different bacterial and plants extracts in treating several types of cell lines (Li *et al.*, 2003; Al-dulami , 2006; Darwish, 2007).

Table (3-30) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B1 on Hep-2 cell line.

Exposure time	Inhibition rate% (mean ±SD)			
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.	
	A,a	A,b	A,b	
240	70.917±0.815	86.573±2.026	88.19±0.908	
	A,a	A,b	A,b	
200	69.863±1.917	86.313±2.293	87.587±1.035	
	A,a	A,b	A,b	
160	66.8±1.46	81.83±4.713	86.437±0.182	
	B,a	AB,b	A,c	
80	$51.375 \pm 1.082$	70.64±4.952	$85.287 \pm 0.657$	
	C,a	C,a	B,b	

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#### lep-2 cell line.

Exposure time	Inhibition rate% (mean ±SD)		
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
240	A,a	A,ab	A,a
	8.433±0.41	13.783±3.726	8.043±2.202
200	A,a	A,a	A,a
	5.34±4.831	10.707±3.327	7.96±0.178

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Table (3-32) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B3 on Hep-2 cell line.

Exposure time	Inhibition rate% (mean ±SD)		
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.
	A,a	A,ab	A,b
240	74.685±4.054	87.179±1.555	90.433±0.519
	A,a	A,ab	A,b
200	$70.42 \pm 8.622$	87.133±2.513	89.967±0.047

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Hep-2 cell line.

Exposure time	Inhibition rate% (mean ±SD)			
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.	
	A,a	A,a	A,a	
240	5.613±3.65	4.165±0.346	$3.47 \pm 0.8485$	
	A,a	B,b	B,a	
200	$3.155 \pm 3.444$	39.335±0.445	$0.67 \pm 0.00$	

Different upper case letters: significant difference (P<0.05) between means of rows Different lower case letters: significant difference (P<0.05) between means of columns
Table (3-34) Cytotoxic effect of the purified lipopeptide produced by *B. subtilis* B5 on Hep-2 cell line.

Exposure time	Inhibition rate% (mean ±SD)				
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.		
	A,a	A,b	A,b		
240	52.04±4.8	82.533±0.242	87.097±0.447		
	B,a	A,b	A,c		
200	28.81±3.118	73.51±0.339	85.947±0.6		
	C,a	B,b	B,c		
160	9.93±1.247	19.855±0.474	75.085±4.632		
	C,a	C,a	B,b		
80	8.233±4.536	7.645±3.472	9.513±2.331		
	C,a	C,ab	C,ac		

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Concentrations 24 hrs. 48 hrs. 72 hrs. mg/l A.a A.b A.b 69.183±1.079 85.963±0.162 88.307±0.359 240 A.a A.b A.b 200 66.187±1.022 85.77±0.29 87.47±0.501 B.ab A,a A,b 160  $65.377 \pm 2.228$  $71.537 \pm 4.642$  $85.003 \pm 0.309$ B,a C,b B.b 80 30.477±6.751 57.757±7.199 69.219±1.419 C,b C,a D.b  $3.3{\pm}1.748$  $15.46 \pm 4.627$ 40  $14.955 \pm 3.062$ C.b C.a D.ab 20 6.36±2.818 10.45±1.852 14.337±3.888

Different upper case letters: significant difference (P<0.05) between means of rows Different lower case letters: significant difference (P<0.05) between means of columns

Table (3-36) Cytotoxic	effect of the purified	l lipopeptide produ	ced by B. s	ubtilis B7
on Hep-2 cell line.				

Exposure time	Inhibition rate% (mean ±SD)				
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.		
	A,a	A,b	A,c		
240	79.13±2.161	84.473±2.483	$90.6 \pm 0.67$		
	A,a	A,ab	A,b		
200	75.257±4.012	80.595±1.336	86.27±3.6		
	B,a	A,b	B,a		
160	52.12±4.78	74.177±10.842	61.5±4.37		
	C,a	B,a	C,b		
80	4.387±2.011	5.36±1.895	22.797±1.798		
	C,a	B,b	D,b		
40	4.927±1.163	0.813±0.147	0.447±0.202		

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mg/l				
	A,a	A,b	A,b	
240	74.593±1.979	86.737±3.055	88.91±4.14	
	A,a	A,b	A,b	
200	73.483±5.137	85.78±2.889	88.757±4.181	
	A,a	A,b	A,b	
160	72.677±5.596	84.967±3.71	$88.357 {\pm} 3.886$	
	B,a	B,b	A,c	
80	11.710±4.2	28.22±4.992	86.043±1.529	
	B,a	C,a	B,a	
40	9.820±2.036	10.047±3.605	16.87±5.94	
	C,a	D,a	C,b	
20	$0.106 \pm 0.074$	0.125±0.046	3.723±0.297	

Different upper case letters: significant difference (P<0.05) between means of rows Different lower case letters: significant difference (P<0.05) between means of columns

Table (3-38) Cytotoxic	effect of the purified	l lipopeptide pro	oduced by <i>B</i> .	subtilis B9
on Hep-2 cell line.				

Exposure time	Inhibition rate% (mean ±SD)				
lConcentrations mg/l	24 hrs.	48 hrs.	72 hrs.		
	A,a	A, ab	A,b		
240	75.497±3.581	$85.78 \pm 2.889$	90.6±0.28		
	A,a	A,b	A,c		
200	71.575±0.827	$84.407 \pm 0.802$	89.7±0.608		
	AB,a	A,ab	A,ab		
160	63.843±9.122	74.03±0.4	84.67±1.6		
	C,a	B,b	B,c		
80	6.16±0.663	20.21±0.99	64.6±7.07		
	C,a	BC,a	C,a		
40	5.407±2.813	8.857±8.252	6.77±5.8		

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#### 10 on Hep-2 cell line.

Exposure time	Inhibition rate% (mean ±SD)				
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.		
240	A,a	A,b	A,b		
	64.083±1.948	84.42±2.421	85.947±0.313		
200	B,a	B,ab	A,b		
	4.34±0.792	30.96±8.319	59.483±4.75		

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Exposure time	Inhibition rate% (mean ±SD)				
Concentrations mg/l	24 hrs.	48 hrs.	72 hrs.		
	A,a	A,ab	A,b		
240	$83.873 \pm 1.872$	88.048 ±0.512	90.542 ±1		
	A,a	A,b	A,b		
200	83.555 ±2.201	88.76 ±0.104	$88.014 \pm 0.836$		
	A,a	A,b	A,b		
160	$79.872 \pm 4.158$	$87.98 \pm 1.689$	$87.3406 \pm 0.663$		
	A,a	A,a	B,a		
80	72.031±13.032	79.603 ±7.411	$78.07 \pm 0.52$		
	B,a	B,a	C,b		
40	$39.142 \pm 8.482$	39.573±4.236	56.67 ±1.049		
	BC,a	C,a	D,a		
20					

Table (3-40) Cytotoxic effect of standard surfactin on Hep-2 cell line.

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#### fied lipopetide (surfactin) produced by the isolate B. B6 was selected to

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study its cytotoxic effect on normal human cell line since it inhibited the proliferation of K562, L1210, Hep-2 cell lines with high efficiency as were shown in appendices (44, 45,46).

Results from table (3-41), revealed very low cytotoxic affectivity of this surfactin on the normal cell line even at the highest concentrations. This considered as indication of the relative safety of *B*. B6 lipopeptide towards normal cells.

Such selective toxicity towards malignant cell lines was due to the differences in the malignant cellular physiology such as the present of some metabolic factors that found in the cancer cell lines but not found in normal cells, like the angiogenic promoters and inhibitors (Folkman,2000 ,Moteki *et al.*,2002). In addition DNA of tumor cell found in relaxant shape ,and the DNA molecule was found in a unstable figure because the far away between the H-bond which connect the both strand of DNA and this make easy for compound to interfere or associated to both strands of DNA, while DNA of normal cell has a strong H-bond connect the both strands to each other and make it more stable, so the compounds cannot interfere or associated with DNA strand (Belijanski, 2000).

The Hep-2 cell line was chosen for further studies since there is no known studies about the effect of *Bacillus* lipopeptide on this cell line.therefore, purified lipopeptide from *B. subtilis* B6 isolate (surfactin ) was chosen at concentration 40

mg/ L and 80 mg/L that cause IC50 to study the mechanism of proliferation

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Concentrations	24 nrs.	<b>40 IIIS.</b>	72 hrs.	
mg/l				
	A,a	A,b	A,c	
240	6.9767±2.0011	30.707±5.927	0.6991 ±0.228	
	B,a	B,b	B,c	
200	1.2133±0.9775	0.0793±0.013		
	B,a	C,a	B,b	
160	$0.3583 \pm 0.2852$	$0.550 \pm 0.043$	$0.0787 \pm 0.008$	
	B,a	C,a	B,a	
80	$0.0538 \pm 0.0562$	$0.051 \pm 0.002$	$0.0396 \pm 0.035$	
	B,a	C,a	A,a	
40	$0.2152 \pm 0.1515$	15 0.4842±0.044 0.1113 ±0.09		
	B,a	C,a	A,b	
20	$0.0124 \pm 0.0014$	$0.026 \pm 0.003$	0.1972±0.1322	

Different upper case letters: significant difference (P<0.05) between means of rows Different lower case letters: significant difference (P<0.05) between means of columns

#### 4.7.2 Mitochondrial membrane potential assay JC-1

In order to study mechanism of action of the purified surfactin from the isolate *B*. B6 in the inhibition of Hep-2 cell proliferation, mitochondrial membrane potential assay were conducted.

Results from figures (3-19) and (3-20) showed gradual loss of J-aggregates which represent the accumulation of the membrane permeate JC-1 dye within healthy mitochondria when cell line treated with it because the dye was bearing a delocalized positive charge that tend to enter into the negatively charged intact mitochondrial membrane and accumulate to form J-aggregates which become fluorescent red while In apoptotic cells, the mitochondrial membrane potential

collapses, and the JC-1 cannot accumulate within the mitochondria

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fluorescent monomeric form, then finally diffused to form av a nice

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Atochondrial membrane at concentration as low as 5000ng/ml.

#### 3.7.3 Caspase – G10 3/7 assay

Caspase -G10 3/7 assay were used to study the involvement of apoptosis in cellular inhibition when treated with *B*. B6 surfactin.

Results shown in table (3-42) revealed increasing caspase 3 activity with prolonged incubation with surfactin which indicate the onset of cellular apoptosis.



Figure (3-19) Hep-2 cell line treated with 40 mg/l of *B. subtilis* B6 surfactin for 8 hrs and stained with JC-1 under A: red light fluorescence microscope showed \* intact cells with J-aggregates, B: under green light fluorescence microscope showed • apoptotic cells that loss J-aggregates to the cytoplasm then diffuse to

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Figure (3-20) Hep-2 cell line treated with 40 mg/l of *B. subtilis* B6 surfactin for 18 hrs and stained with JC-1 under C: red light fluorescence microscope showed  $\star$  decreased number of intact cells with J-aggregates, D: green light fluorescence microscope showed  $\blacklozenge$  increased number of apoptotic cells that loss J-aggregates to the cytoplasm then diffuse to form green monomer (X40).

Table (3-42) Caspase activity in Hep-2 cell line treated with 40mg/L of purified surfactin produced by *B. subtilis* B6 for different incubation time.

sample	Caspase activity (unit/well)
control	9690.00
Treated cell for 6 hours	24848.0
Treated cell for 12 hours	48367.0
Treated cell for 18 hours	56407.3

Caspase 3 and caspase 7 form part of effector caspases which when activated cleave a large number of substrates in the cell that lead to disassembly of This is a watermark for the trial version, register to get the full one!

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arresting the cell cycle according to several lines of evidence on DNA

fragmentation, Annexin V staining, and altered levels of poly (ADP-ribose) polymerase, caspase-3 activation, while Cao *et al.*,(2010) found that surfactin produced by *Bacillus natto* TK-1 induce apoptosis in Human breast cancer (MCF-7) 1 by caspase cascade activation through induction of ROS/JNK-mediated mitochondrial/caspase pathway.

## Summary

Eighty eight local bacterial isolates of *Bacillus* spp. were obtained from 45 oil contaminated soil samples from different fuel stations in addition to Al- Dorah oil refinery in Baghdad Province /Iraq.

All these isolates were screened for their ability for biosurfactant production by surface tension measurement of cell free supernatant after cultivation in E medium, which referred that 81 of the 88 isolates were biosurfactant producer, while screening depended on the blood haemolysis activity on sheep blood agar indicated that only 49 isolates were biosurfactant producers.

Molecular identification based on the 16S rDNA by two sets of primers (27F and 1492R). (B16SF and B16SR) of the best ten biosurfactant producer isolates This is a watermark for the trial version, register to get the full one!

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nest Lency (40 -43.1)% in Jacques medium than in E medium (28-30)%.

Biosurfactants produced by the selected isolates were purified by acid

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precipitation followed by ultrafiltration with centricons.

Purified biosurfactants were found to be member of the lipopeptide family when characterized by the TLC that sprayed with water or with Rhodamine B, and all isolates were capable of producing surfactin with or without other types of biosurfactants depending on the isolate as showed by comparison with standard surfactin. HPLC/MS analysis of the purified biosurfactant from *B. subtilus* B6 revealed that it is a surfactin with partial sequence determination as follows: Val-Asp -Leu-Leu-OH<sub>2</sub>.

Cytotoxicity assay of the purified biosurfactant from the selected isolates on (K562, Hep-2, L1210) were tested, results showed that lipopeptides cytotoxicity were dependent on the type of tumor cell line, type of isolate, biosurfactant concentration and exposure time

Purified biosurfactant produced by *Bacillus* B7 was found to be the most cytotoxic one to K562 cell line that caused 57.17% inhibition of the cell line growth at concentration (80) mg/l, while purified biosurfactant produced by

isolate Bacillus B9 was the most cytotoxic one to L1210 cell line that caused

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purified biosurfactans produced by (B. subtilis isolates Be

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was found that it caused depolarization of mitochondria membrane which

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finally led to cell death. In addition, the caspase activity assay revealed that surfactin induced apoptosis in Hep-2 cells that associated with caspase 3.

Optimum conditions for biosurfactants production from *B. subtilis* B6 were determined . Results indicated that maximum biosurfactant production from this isolate was achieved by using Sucrose (1%) v/v , Potassium nitrate (0.3%) and Potassium dihydrogen phosphate (1 g/l), at pH 8, 30 °C, 180 rpm of shaking for 72 hrs. The biosurfactant yield was 1.4 g/l of culture medium under the optimum conditions.

Table (3-4) : Results of NCBI alignment of PCR amplification of 16S rDNA by two sets of primers (B16SF and B16SR) and (27F and 1492R) of the isolates (B1,B2,B3 ,B4,B5,B6,B7,B8,B9,B10).

Primer	<i>B</i> .	Bacillus	Bacillus	Bacillus	Bacillus	Bacillus	Bacillus	Bacillus	Bacillus
	spp	subtilis	tequilensis	licheniformis	amyloliquefaciens	methylotrophicus	mojavensis	axarquiensis	malacitensis
B1*	25	68	4	0	2		1	0	0
B1									0
(27F)								<b>C</b> II <b>C</b> I	TM
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(1492R)									
B2*		Benef	fits for registe	ered users:					0
B2		36	0	9			8	4	2
(27F)		LINO V	watermark or	i the output doc	suments.				
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(1492R)		3.No	page quantity	/ limitations for o	converted PDF files.				
B3*									0
B3									0
(27F)									
B3	16	٤٢	17	0	9	6	3	2	0
(1492R)									
B4*	29	59	4	0	2	0	1	0	0
B4	18	٧٥	3	1	0	0	0	0	0
(27F)									
B4	15	٤٢	17	0	8	6	3	4	0
(1492R)									
B5*	20	23	16	0	6	0	5	4	0
B5 (27F)	48	٤٧	1	2	0	0	1	1	0

B5	11	०٦	16	0	6	6	2	2	0
(1402P)			10	v	Ū	Ū	-	-	Ŭ
(1492K)									
B6 *	46	41	0	0	2	0	7	2	0
B6 <sup>◆</sup>	47	26	2	2	0	0	13	6	0
B7*	20	23	16		6	0	5	4	0
B7	43	48	1	1	0	0	1	3	0
(27F)									
B7		55	. 16	0	7	6	2	2	TN
(1402D)		This	s is a wa	termark for	the trial vers	ion, register t	o get the	e full one!	
(1492K)						· U			
B8*									0
B8		Bene	fits for registe	ered users:					0
(27F)		1 No.	watermark or	the output doc	uments				
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B9*		3.110	page quantity	minitations for	convented PDF mes.				0
B9 <sup>●</sup>									0
B10*									0
B10	49	46	1	2	0	0	1	1	0
(27F)		_			-	-			-
P10	20	50	15	0	2	0	2	2	0
DIU	20	39	15	U	Z	U	Z	Z	U
(1492R)									

\*: primer set (B16SF and B16SR), \*: primer set (27F and 1492R)

الملخص

عزلت ٨٨ عزلة محلية عائدة للجنس Bacillus من ٤٥ عينة تربة ملوثة بالمشتقات النفطية من مختلف محطات الوقود إضافة إلى مصفى الدورة.

تبين من إخضاع جميع هذه العزلات للتحري عن قابليتها في إنتاج المستحلب الحيوي بواسطة قياس الشد السطحي لوسط النمو E- medium الخالي من الخلايا بعد نموها فيه ان ٨١ عزلة كان منتجة بينما اظهر التحري عن الإنتاجية بواسطة الكشف عن تحلل الدم قابلية ٤٩ عزلة فقط على إنتاج المستحلب الحيوي . وعند تشخيصها بالاعتماد على دراسة Bidsr and 16S rDNA باستخدام زوجين من البرايمرات Bidsr and 27F and وعند تشخيصها بالاعتماد على دراسة العزلات المخالية العزلات المتحدام زوجين من البرايمرات المستحلبات

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Bacillus subtilis، وكونها تطابق Bacillus subtilis بنسبة تتراوح (۹۷–

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السطحي عند تتميتها في وسط Jacques medim و بنسبة (43-40)% بالمقارنة مع وس

E. medium بنسبة (۳۰-۳۰)%

نقيت المستحلبات الحيوية من العزلات البكتيرية المختارة بواسطة الترسيب بالحامض والترشيح بالاغشية ثم وصفت باستخدام كروماتوغرافيا الطبقة الرقيقة والتي اظهرت عائديتها لعائلة الببتيدات الدهنية، اذ كانت موجبة لفحص الماء وكاشف الرودامين B ، وعند مقارنتها مع السرفاكتين القياسي تبين انها منتجة للسرفاكتين والذي أنتج بصورة مفردة أو كمزيج مع غيره من الببتيدات الدهنية اعتمادا على نوع العزلة المنتجة. عند التنقية والتوصيف باستخدام جهاز كروماتوغرافيا السائل عالي الكفاءة لتحديد الكتلة للمستحلب الحيوي المنتج من قبل العزلة Bacillus B6 تاكد بأنه سرفاكتين كما تم تحديد تتابع الأحماض الأمينية المكونة له بصورة جزئية وكان بالتسلسل Val-Asp-Leu-Leu-OH2 .

أظهرت نتائج فحص السمية للمستحلبات الحيوية المنقاة المنتجة من قبل العزلات المختارة على خطوط الخلايا السرطانية (K562,Hep-2, L1210)، أن تأثير المستحلبات الحيوية اعتمد على نوع حط الحلايا السرطانية ونوع العزلة المنتجة وعلى تركيز المستحلب الحيوي ومدة التعريض.

لوحظ إن المستحلب الحيوي المنقى المنتج من العزلة Bacillus BV بأنه الأكثر تثبيطا لنمو خط الخلايا السرطانية K562 اذ تسبب بنسبة تثبيط بلغت 57.17% بتركيز ٨٠ ملغم /لتر) بينما كان المستحل المنتج من العزلة Bacillus B الأكثر تثبيطا لنمو خط الخلايا السرطانية L1210 بنفس

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#### الخلايا الذاتى

أظهرت دراسة الظروف المثلى لإنتاج المستحلب الحيوي من قبل العزلة Bacillus B6 إن أعلى إنتاجية للمستحلب الحيوي بلغت (1.4 غم/لتر) في وسط يحتوي على السكروز (١%) كمصدر كاربوني وعلى نترات البوتاسيوم (0.3%) كمصدر نايتروجيني وفوسفات الصوديوم ثنائية الهيدروجين (١غم/لتر) كمصدر فوسفاتي و pH مساوي لـ ٨ بدرجة حرارة حضن ٣٠م وبمعدل اهتزاز ١٨٠ دورة/دقيقة لمدة حضن ٢٢ ساعة.



جمهورية العراق وزارة التعليم العالي والبحث العلمي جامعة النهرين كلية العلوم

## الفعالية البيولوجية للمستحلبات الحيوية المنتجة من بكتيريا .*Bacillus* spp المعزولة محلياً

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باشراف

د.ناهي يوسف ياسين (أستاذ)

د. حميد مجيد جاسم (أستاذ)

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تشرين الثاني ٢٠١١

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## Chapter One

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# **Chapter Four**

## Conclusions

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