Supplemenatary (S2)

Egyptian Journal of Botany
http://ejbo.journals.ekb.eg/

Supplementary (S2). Sequence alignment of the ITS region

Data 1 Sequence alignment for the ITS region of 10 selected species of Malvaceae *s.l*

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4       ----------------------------------------------GGTGAACCTGCGGAAG    16
9       ------------------------------------------------------------    0
10      ------------------------------------------------------------    0
14      ------------------------------------------------------------    0
19      ---------------------------------------CGTAGGTGACCTGCGGAAG    19
1       CTTGGTCATTTAGAGGAAGTAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAG    60
16      ------------------------------------------------------------    0
22      ------------------------------------TCCGTAGGTGAACCTGCGGAAG    22
13      --------------GGAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAG    46
17      ------------------------------------TCCGTAGGTGAACCTGCGGAAG    22

4       GATCATTTTCGAAACCTGCCTAGCAGAACGACCCGCGAACGAGTTATCAACACACC----    72
9       --------------ACTGTCCCGCAGACCGACCCGTGAACTAGTTGTGTACACAAACATC    46
10      --------------CTCGGACCCTAAAACGACCCGTGAACGCGTTAA--GTACAAACATC    44
14      --------------CCTGCATAGCAGAACGACCCGTGAATGTGTTATCATACAAAACAAC    46
19      GATCATTGTCGAAACCTGCCTAGCAGAACGACCCGCGAACGYGTTATCGAACAACCGATC    79
1       GATCATTGTCGAAACCTGCCCAGCAGAGCGACCCGCGAACTAGTAACACAACACTCGGGG    120
16      --------------CCTGCCTAGCAGAACGACCCGCGAACGCGTTGCAAACAACACCGGA    46
22      GATCATTTCCGAAACCTGCCCAGCAGAACGACCCGCGAACGCGTTGTAAACAACACCGGA    82
13      GATCATTGTCGAAGCCTGCCCAGCAGAACGACCCGCGAACGTGTTATCGAAAAACAACGG    106
17      GATCATTGTCGAAACCTGCCTAGCAGAACGACCCGTGAACGTGTTATCWAACATCAAAGG    82
                        *      * * ******* ***   **

4       CGAAGGGGGCCAGGATGCCACCACGCTCCGAACCCCCTTCGAGTCGCGGCGAGCGTGACG    132
9       GCGGGGGGGCGAGGGTGCGTACGTGCCCCGAGCCCCCTCGGTGCCCTGGCGTTCGCGACC    106
10      GCGGGGGGGTGAGGGAGCGTCCGCGCCCCGATCTCCCTCGCGGCCCTTGCGCGCGGGACC    104
14      GAGAGGGCGCGGGTTTAGACCGTCCC--AACTTTC--TCGATGCCTTGGTGCGTTTGGTC    102
19      GAGGGGGTGCGGATGCATCCTYGCCCCGAGCCCCC--TCGACGCCTCGGCGCGCCGAGCC    137
1       GGCGAGGGCGCACAAGTGTG----CTCTACTCCCC--CCCATGGCCCGGG--TGCTCGTG    172
16      GGTGGTGCGGGTGC--ATCCTCGCCTCTCGCCACC-CC-CGTGTCTCGGAGCGGCCAGTC    102
22      GGTGGTGGGGGTGC--ATCCTTGCCTCTCGCTACC-CTCCACGTCTCGGAGCGATCGGTC    139
13      GTGGGGCGAGGCGGCATCCC-CGCCCATCGTCC----------------------------    138
17      GAGGGGGTGCGGTCGCATCATTGCGCCCCGTCCCC--TTCCTGA----------------    124

4       GTTGCTTCG--CTTTGCCCTCGAGGCAGGTGGGGTGCCCGCTCTCCTCGCCCGCGGCACA    190
9       TCGA-CCCG--TCAGGCCTTCGTGGCGGGCGGGATGCGTGGTCG-CGTCGCTTGCGGCAG    162
10      AGCA-CCCG--CCACGCCCCCCGGGCGTGCGGGACGGTTGGTTCCCGTCGCTAGCGGCCG    161
14      TTGCCACATCCCCCGTTT-----TGGAGGGTGACAAGCATGTTCCATCCACACAAAGGCA    157
19      TCTCCGCATCCGTCCTCG-----GGC-GGGTGT-CCCGGGTCTCGTCGTGCTCCGGGGCA    190
1       ACCCCGTGTCGTTCCGTGGCCTCTGGGCATGGG---ATGCGCGAGGCCGTCGCCTGTGGC    229
16      TCGTCGTCCCTTTGCCCGTCGGGTGGGGTGAGATGCCGGGATCAACCTCTTCGAGGCAA-    161
22      TTGCAGTCCCTCCGCCCATCGGGCGGGGTGAGATGCCAGGTTCAACCTCTCCGAGGCAAA    199
13      ------------------------------------CCCTACCCTCGCCTCTCGGGGCGA    162
17      ---------------------------CCCGGTGACATGGGAACCCGTGTCCCCGTGGCA    157
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Data 1 (continous)

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4       CAATAACGAACCCCCGGCGCGAATCGCGCCAAGGAACCG--TAAAGAAAAAGAGCACGTT    248
9       AACCAACGAACCCCCGGCGCGAGTCGCGCCAAGGA--AACGGAATGACGAGGAGCACGTC    220
10      AAACAACG-AACCCCGGCGCGAGTCGCGCCAAGGAATACGAAAATGATGAGGAGCACGCC    220
14      AAACCAACAACCCCCGGCGTGAATTGCGCCAAGGAATTAAAACTAAAAGAGGG--CACGC    215
19      AAACGAACAACCCCCGGCGCGAATCGCGTCAAGGAATAAAAAATGAAAAGAGT--GCGTG    248
1       TTCATAACGAACCCCGGCGCGAATCGCGCCAAGGAACTCGAACTAAAGGACCACGACGG-    288
16      AGCGAACAAACCCCCGGCGCGAATCGCGCCAAGGAATCGAAACGAAAGAAGGGGCACGTC    221
22      AACGTACAAACCCCCGGCGCGAATCGCGCCAAGGAATCGAAACGAAAAAAAGGGCACGTC    259
13      AACGAACAAACCCCCGGCGCGAATCGCGCCAAGGAATCTGAATTGAAAGGA--GCACGTC    220
17      AAACAAACAACCCCCGGCGTGAATTACGCCAAGGAATCTGAATGAAAAGGT--GGTCGTC    215
             *   * ******** ** *  ** ******          *

4       GCTCGTCGCTGCCCCGTTCGCGGTGAGTA--A---GCGCGGCCTCGTGTCTCCTTTCCTA    303
9       CGCTGCCCCCGCCCCGCCCGCGGTGCGCGTGCGGGCAGCGGCGTGTCTCCTCCGTCG--T    278
10      GGCTGTCGCCGCCCCGTCCGCGGGGTCCGTGCGCGCAGCGTCGTGTCTCCTCCGTCGCGA    280
14      TACTGTCGCAGACCCGTTTGCGGTGTTTGTGCGG---CAGTGTCGTCTGCTACTTTTGTC    272
19      ------------------TTTCGTTGTCGTATGG---CAGCGAGGGCGTTACTCTCGTCG    287
1       -----AAGCCGCCCCGGTCTCGGTGTG------T-TGGCAACCGGCCGTGCCTAGTTTAT    336
16      TTCTGTCGCCGCACCGTTCGCGGTGTCGATGCTT-CAGTGATGTTGTTCTCTTGTCGC--    278
22      TCATGTCGCCGCACCGTTCGCGGTGTCGGCGCTT-CGGTGATGTTGTTCTTCCGTCGCGA    318
13      CCCCGTCGCCACCCCGTCCGCGGTGCGTGTGCCG-CGGGGA---CGCTGCGACTTCGTCG    276
17      TGTTGTCGCACCCCCGTTCGCGGTGCGTGTGCGG-CGGAGA---CGTGCCCACTTTGTCG    271
                             *

4       AGAAAATCTAAAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATGAAGAACGT    363
9       GAAAATTAAGAAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATGAAGAACGT    338
10      AAAAAATAACAAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATGAAGAACGT    340
14      GTGAAAAACAAAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATGAAGAACGT    332
19      TGAAAATTAAAAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATGAAGAACGT    347
1       TGAGAATCTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATGAAGAACGT    396
16      AAAATATACAGAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATGAAGAACGT    338
22      AACACATACAGAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATGAAGAACGT    378
13      TGAATACACAAAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATGAAGAACGT    336
17      TGAATACACAAAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATGAAGAACGT    331
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4       AGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACG    423
9       AGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACG    398
10      AGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACG    400
14      AGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGCGAACCATCGAGTCTTTGAACG    392
19      AGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACG    407
1       AGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACG    456
16      AGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACG    398
22      AGCGAAATGCGATACTTGGTGTGAATTGCAGGATCCCGTGAACCATCGAGTCTTTGAACG    438
13      AGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACG    396
17      AGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTTTTTGAACG    391
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Data 1 (continous)

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4       CAAGTTGCGCCCCAAGCCATTAGGCCGAGGGCACGCCTGCCTGGGTGTCACGCATCGTCG    483
9       CAAGTTGCGCCCCGAGCCATTAGGCTGAGGGCACGCCTGCCTGGGTGTCACGCATCGTCG    458
10      CAAGTTGCGCCCCAAGCCATTAGGCCGAGGGCACGTCTGCCTGGGTGTCACGCATCGTCG    460
14      CAAGTTGCGCCCCAAGCCATTAGGCCGAGGGCACGTCTGCCTGGGTGTCACGCATCGTCG    452
19      CAAGTTGCGCCCCAAGCCTTTAGGCCGAGGGCACGTCTGCCTGGGTGTCACGCATCGTCG    467
1       CAAGTTGCGCCCCAAGCCATTAGGCCGAGGGCACGTCTGCCTGGGTGTCACGCATCGTCG    516
16      CAAGTTGCGCCCCAAGCCATTAGGCCGAGGGCACGTCTGCCTGGGTGTCACGCATCGTCG    458
22      CAAGTTGCGCCCCAAGCCATTAGGCCGAGGGCACGTCTGCCTGGGTGTCACGCATCGTCG    498
13      CAAGTTGCGCCCCAAGCCGTTAGGCCGAGGGCACGTCTGCCTGGGTGTCACGCATCGTCG    456
17      CAAGTTGCGCCCCAAGCCATTAGGCCGAGGGCACGTCTGCCTGGGTGTCACGCATCGTTG    451
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4       CTCCCCCCAACGACCCTCACGGGTGAAC----------GGAAGGGACGGAAAATGGCCTC    533
9       CCCCCTCCAATCCCTTAGCCCCTCGGGGCG-GGGACGAGGTGGGGGCGGAAAATGGC-CT    516
10      CCCCTCCCCATCCCTGGCCCCTCGGGGCTCCGGGTGAGGCGGGGGGCGGAAAATGGC-CT    519
14      CCCCCATCAACACT---------------------G-AGTGTGGGCGGAAATTGG-CTT    488
19      CCCCCGTCAAACCCCGAGCCCTC-GGGCCGGGATCGA-CGCGCGGGCGGAAATTGG-CCT    524
1       CCCCCCCCAACTCCCGAGCCTCCAAGG-CTCCGGGTT-GCATCGGGCGGATAATGG-CCT    573
16      CCCCCATCCAACCATGAGCCCTC-GAGCCTCGGTTGG-ACCGCGGGCGGAAATTGG-CCT    515
22      CCCCCATCCAACCCCGAGCCCTC-GGGCCTCGGTTGA-ACTGTGGGCGGAAATTGG-CCT    555
13      CCCCCATCCAACCCTTTCCCCGGAGGGAACGGGTTGC-GTTGCGGGCGGACAATGG-CCT    514
17      CCCCCATCCAACCCTTACCCACA-GGGCATCGGTTGA-GGTGTGGGCGGAGAATGG-CCT    508
        * **   * *                                 ** **** * ***

4       CCGTGAGCCTCCTCGCTCGCGGTTGGCCCAAAT-GCGAGTCCTCGGCGACGGCAGCGCCG    592
9       CCCGTGCGCTCCCCGCTCGCGGTTGGCCTAAAA-TCGGGTCCCGGGCGGCGACAGCGTCG    575
10      CCCGTGAGCTCCGCGCTCGCGGTTGGCCTAAAA-TCGGGTCCTCGGCGACGATAGCGCCG    578
14      CCCGTGCGCTCATCGCTCGCGGTTGGCCTAAAA-ATGAGTCTTCGGCGAT-GAAGTGCCG    546
19      CCCGTGCGCTCACCGCTCGCGGTTGGTCTAAAT-TCGAGTCCTCGGCGAT-GAAGCGCCG    582
1       CCCGTGCGCTCTTGGCTCGCGGTTGGCCTAAATAGCCAGTCCTCGGCGATCATGGTGCCG    633
16      CCCGTGCGCTCACAGCCAGCGGTTGGCCTAAAT-TCGAGTCCTCGACGACATCATCGTCG    574
22      CCCGCGAGCTCACCGCCGGCGGTTGGCCTAAAT-ACGAGTCCTCGACGACGGTATCGTCG    614
13      CCCGTCCGCGCATCGCCCGCGGTTGGCCCAAAA-TCGAGTCATCGGCGACCACGGTGCCG    573
17      CCCGTGCGCTCACCGCTCGCGGTTGGCCTAAAA-TCGAGTCCTCGGCGACCACGGTGCCG    567
        **      * *   **  ******** * ***      ***   * **        * **

4       CGACGATCGGTGGTAACGCCTCCTAGGCTTCCCTCGTCCCTCGTCGCAAGCGCCCTTCGT    652
9       CGACGATCGGTGGTGCTGCCTCGGGCGCGCCTCGTTCGC--GGTCGCGCGCG---CTTTC    630
10      CGACAATCGGTGGTCATGCCTCGAGCGCTCC----------GGTCGCGTGCG---TTTTC    625
14      CGACAATCGGTGGGAATGCTTTCAGCTGCCTCGTTCGTAGTCG-----TGTGCGCTCGTC    601
19      CGACGATCGGTGGGAACGCCTTTGGCTGCCTCGTTCGGAGTCG-----CGCGCGCTCGTC    637
1       CGGCCTTCGGTGGTAACGCTATTATCGGCATGCCCCGGTCGCG-----TTCGCCCCTCGT    688
16      CGACGATCGGTGGTAATGCTGCAAGCAACCTCGTTCGGAGTCG-----TGCGCGTCCGT-    628
22      CGACGATCGGTGGTAATGCTATCGGCTGCCTCGTTCGGAGTCG-----TGCACGTTCGT-    668
13      CGACGATCGGTGGTAACGCTTCGGGCTGCCTCGTTCGTAGTCG-----CGCGCTACCGT-    627
17      CGACAATCGGTGGTGTTGCTTCGAGCTGCCTCGTTTTTTGTCG-----TGTGCTGCCGT-    621
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Data 1 (continous)

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4       CCGTCCGGACCCCCTTCGAGACCCTGAACGCATCGTTTCGACGATGCTCGCATCGCGACC    712
9       GCTCGGCCGGACCCATCGAGACCCTACTCGCGTCGCACGAGCGATGCTCGCATCGCGACC    690
10      GTTCGTTCGGACCCTTCGAGACCCTACTCGCGTCGCACGAGCGATGCTCGCATCGCGACC    685
14      GATT---AGGACATCATGACCCTT--TTTGCATCAAACTGTTGATGCTCGCATCGCGACC    656
19      GATC---GGGACGCTTTCGACCCTTTAAGGCATCGCGACGTCGATGCTCGCATCGCGACC    694
1       CCTGTTGGACCAATGAATAAGACCCTATGGCGTCGATCTGTCGATGCCCGCAATGCGACC    748
16      ----CGATCGAGACCCTTGAACCCTTTCGGCATCGCAAGGACGGTGCTCGCATCGCGACC    684
22      ----CGATCGAGGCCCCTGACCCCTTTTGGCATCGCAAGGACGGTGCTCGCATCGCGACC    724
13      ----CGACCCCGGCTCTCCGACCCTTCCTGCACCGCGAACACGGTGCTCGCGTCGCGACC    683
17      ----CGATCCGTGCTCTCTGACCCTTTCGGCACCGCAAGCACGGTGCTCGCATCGCGACC    677
                                  **  *        * *** ***   ******

4       CCAGGTCAGGCGGGATTACCCGCTGAATTTAAG-------------------    745
9       CCAGGTCAGGCGGGACCACCC-------------------------------    711
10      CCAGGTCAGGCGGGACTACCC-------------------------------    706
14      CCAGGTCAGGCGGGATTACCC-------------------------------    677
19      CCAGGTCAGGCGGGATCACCCG------------------------------    716
1       CCAGGTCAGGCGGGATTACCCGCTGAGTTTAAGCATATCAATAAGCGGAG--    798
16      CCAGGTCAGGCGGGATTACCC-------------------------------    705
22      CCAGGTCAGGCGGGATTACCCGCTGAGTTTA---------------------    755
13      CCAGGTCAGGCGGGACTACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA    735
17      CCAGGTCAGGCGGGATTACCCGCTGAGTTTAAGCATATCAATAAGCGGAGAA    729
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