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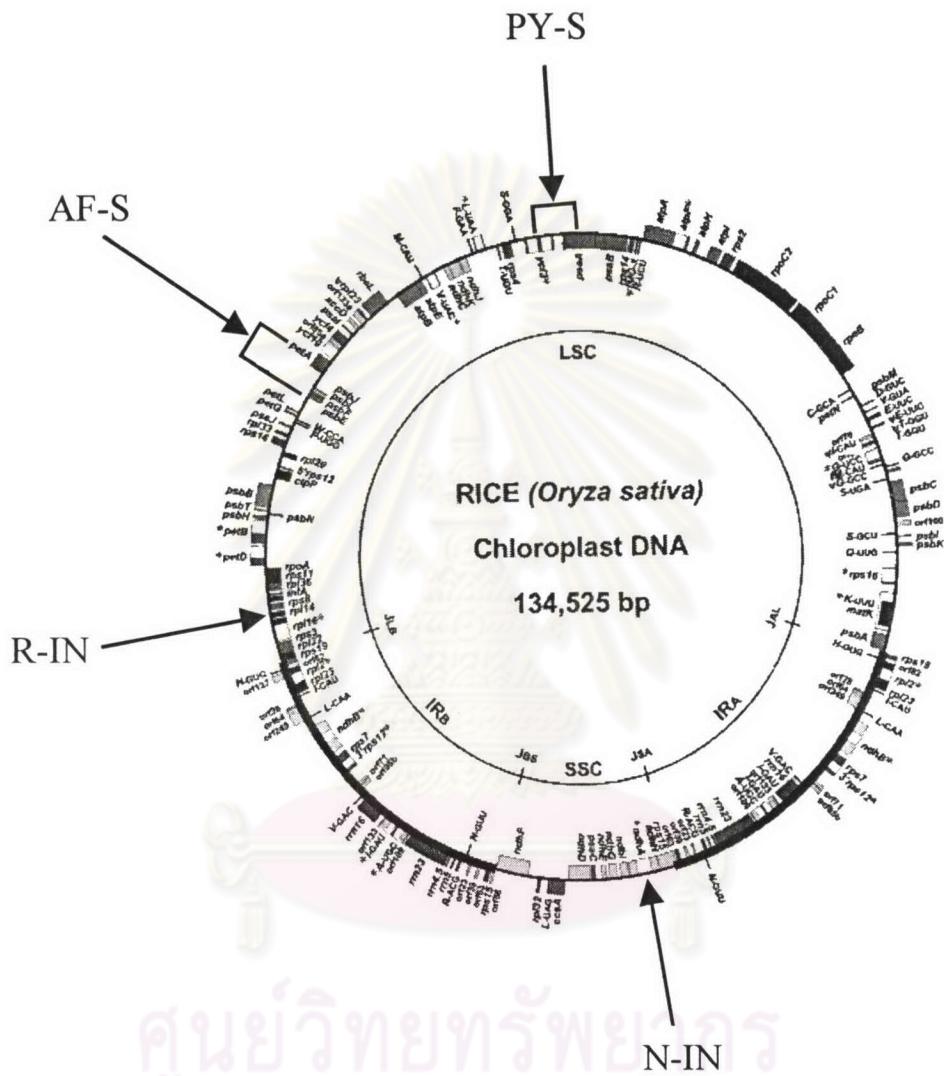


APPENDICES

ศูนย์วิทยทรัพยากร
จุฬาลงกรณ์มหาวิทยาลัย

APPENDIX A

RICE COMPLETE CHLOROPLAST GENOME



Locations of four selected fragments for chloroplast sequence analyses in *Musa* (Chapter 3) on rice complete chloroplast genome (Tsudzuki et al. in press). The fragments indicated by arrows are (from top to bottom, counter clock-wise) PY-S = *psaA*-*ycf3* intergenic spacer; AF-S = *petA*-*psbJ*-*psbL*-*psbF* intergenic spacers; R-IN = *rpl16* intron; and N-IN = *ndhA* intron.

APPENDIX B

ALIGNMENT OF *Musa* COMBINED CHLOROPLAST SEQUENCES¹

¹ Accession codes appeared according to Table 3.1.
 Dot symbol (.) show a site with the same character variant to that of the first sequence.
 See Table 3.4 for detail of each fragment.

TT_JCK	GCTATGCTTA	GTGTTGACT	CGTTAATTTC	TCTTTAGAGT	TTAGAGTTGG	GATT-----	-----	-----	-CA AAAA.....	[120]
TT_FAA	[120]
TT_FTL	[120]
TT_TXT	[120]
BB_NAN	[120]
BB_SKT	[120]
BB_PKW	[120]
BB_P28	[120]
BB_CMR	[120]
BB_LCK	[120]
BBA_PSB	[120]
BBBA_PLH	[120]
BBA_NWN	[120]
BBA_MNG	[120]
BBA_HIN	[120]
ABB_HIN	[120]
ABB_PGM	[120]
AB_AUK	[120]
AA_SAM	[120]
AA_MAL	[120]
AAA_HOM	[120]
AA_BKS	C.....	[120]

TT_JCK	GCTGTGTTCC AATACTCAGG AGCTTGATCG AACCAAGCCT CCGCAATTTC AGAATCACCT	TGTAGAAATGG CCTGTTCTCC TCGGGCTGCT	CAatCTATAA GTTATGAAAT ACCATTAAC	A.....	[2160]
TT_FAA
TT_FTL
TT_TXT
BB_NAN
BB_SKT
BB_PKW
BB_P28
BB_CMR
BBB_LCK
BBA_PSB
BBBA_PLH
BBA_NWN
BBA_MNG
BBA_HIN
ABB_HIN
ABB_PGM
AB_AUK
AA_SAM
AA_MAL
AAA_HOM
AA_BKS
TT_JCK	CTATGTGT TATCAAATTC TCTACGTGTG ATTCTGGTGA CATGAACCTT ATACTCTT CCTAGAAAGTA AAGGAAAGAA GTTGAATGTA	ATGTATTGAA TAGATATTT	CTTTTTATT	CTTTTTATT	[2280]
TT_FAA
TT_FTL
TT_TXT
BB_NAN
BB_SKT
BB_PKW
BB_P28
BB_CMR
BBB_LCK
BBA_PSB
BBBA_PLH
BBA_NWN
BBA_MNG
BBA_HIN
ABB_HIN
ABB_PGM
AB_AUK
AA_SAM
AA_MAL
AAA_HOM
AA_BKS

TT_JCK	ACGATTCCGA	TCTAGAGTAT	GCTCCTATTC	ACTGGTTAAA	GAATGACTA	TCCAGAACCGA	ATTAATCCCT	TATTCTTTTT	TTTT---AT	TTTTTTAAA	GTACCCCTTT	CTGAGATAGA	[2880]
TT_FAU	[2880]
TT_FTL	[2880]
TT_TXT	[2880]
BB_NAN	TTT...	[2880]
BB_SKT	-C...	[2880]
BB_PKW	-C...	[2880]
BB_P28	-C...	[2880]
BB_CMR	-C...	[2880]
BBB_LCK	-C...	[2880]
BBA_PSB	-C...	[2880]
BBBA_PLH	-C...	[2880]
BBA_NWN	-C...	[2880]
BBA_MNG	-C...	[2880]
BBA_HIN	-C...	[2880]
ABB_HIN	-TT...	[2880]
ABB_FGM	-TT...	[2880]
AB_AUK	-TT...	[2880]
AA_SAM	-C...	[2880]
AA_MAL	-C...	[2880]
AAA_HOM	-C...	[2880]
AA_BKS	-C...	[2880]
TT_JCK	AGAAZAGGAA	CAAAAGAAAT	GGAAATGTAAT	AATCGAATG-	-----	-----	-----	-----	-----	-----	-----	-----	[3000]
TT_FAU	[3000]
TT_FTL	[3000]
TT_TXT	[3000]
BB_NAN	T.....	[3000]
BB_SKT	G.....	[3000]
BB_PKW	G.....	[3000]
BB_P28	G.....	[3000]
BB_CMR	G.....	[3000]
BBB_LCK	G.....	[3000]
BBA_PSB	G.....	[3000]
BBBA_PLH	G.....	[3000]
BBA_NWN	G.....	[3000]
BBA_MNG	G.....	[3000]
BBA_HIN	G.....	[3000]
ABB_HIN	G.....	[3000]
ABB_FGM	G.....	[3000]
AB_AUK	G.....	[3000]
AA_SAM	G.....	[3000]
AA_MAL	G.....	[3000]
AAA_HOM	G.....	[3000]
AA_BKS	G.....	[3000]

[3120] TTT_JCK	TAATGGACTA	-----AATT CTTAAATTCTT TAGGATTA TTGATAAC GAGTATTAA TTGAAGGCT CGAACAAAGA AAAATAAATT TTGAT-----CA TAAGGTGATG	[3120]
[3120] TTT_FAA	[3120]
[3120] TTT_FTL	[3120]
[3120] TTT_TXT	[3120]
[3120] BB_NAN	C..	ATTCCCT..	[3120]
[3120] BB_SKT	C..	ATTCCCT..	[3120]
[3120] BB_PKW	C..	ATTCCCT..	[3120]
[3120] BB_P28	C..	ATTCCCT..	[3120]
[3120] BB_CMR	C..	ATTCCCT..	[3120]
[3120] BB_LCK	C..	ATTCCCT..	[3120]
[3120] BBA_PSB	C..	ATTCCCT..	[3120]
[3120] BBA_PLH	C..	ATTCCCT..	[3120]
[3120] BBA_NWN	C..	ATTCCCT..	[3120]
[3120] BBA_MNG	C..	ATTCCCT..	[3120]
[3120] BBA_HIN	C..	ATTCCCT..	[3120]
[3120] ABB_HIN	.	ATTCCCT..	[3120]
[3120] ABB_PGM	.	ATTCCCT..	[3120]
[3120] AB_AUK	.	ATTCCCT..	[3120]
[3120] AA_SAM	.	ATTCCCT..	[3120]
[3120] AA_MAL	.	ATTCCCT..	[3120]
[3120] AAA_HOM	.	ATTCCCT..	[3120]
[3120] AA_BKS	.	ATTCCCT..	[3120]
[3240] T-T_JCK	GATCAATTCC	GAAGCACTTT TCTTATTCT AGCAAAACAGA ATTCCATTTG TTTAATTTG GACTCTCTGA TATAACTTAA TTCTAGTAC CTCCAAAGAA AGGGGGTAT AATATCGAAC	[3240]
[3240] T-T_FAA	[3240]
[3240] T-T_FTL	[3240]
[3240] T-T_TXT	[3240]
[3240] BB_NAN	T..	[3240]
[3240] BB_SKT	T..	[3240]
[3240] BB_PKW	T..	[3240]
[3240] BB_P28	T..	[3240]
[3240] BB_CMR	T..	[3240]
[3240] BB_LCK	T..	[3240]
[3240] BBA_PSB	T..	[3240]
[3240] BBA_PLH	T..	[3240]
[3240] BBA_NWN	T..	[3240]
[3240] BBA_MNG	T..	[3240]
[3240] BBA_HIN	T..	[3240]
[3240] ABB_HIN	T..	[3240]
[3240] ABB_PGM	T..	[3240]
[3240] AB_AUK	T..	[3240]
[3240] AA_SAM	T..	[3240]
[3240] AA_MAL	T..	[3240]
[3240] AAA_HOM	T..	[3240]
[3240] AA_BKS	T..	[3240]

CGATCCTTAC ATTATTGTG GGCCATAGAG GAGCCGTATG AAGCTGAGGT CTCATGTACG GTTTGGAA AGCGATGGAA ACAGTAAGT TATAATTATC TAACAGTTCA [3360]
 TT JCK ...
 TT FAA ...
 TT FTL ...
 TT TXT ...
 BB NAN G...
 BB SKT G...
 BB PKW G...
 BB P28 G...
 BB CMR G...
 BBB LCK G...
 BBB PSB G...
 BBB PLH G...
 BBB NWN G...
 BBB MNG G...
 BBB HIN G...
 BBB HIN TAG...
 BBB PGM TAG...
 BBB AUK TAG...
 AA SAM TAG...
 AA MAL TAG...
 AAA HOM TAG...
 AAA BKS TAG...
 AGTAGCTT JCK ...
 TT FAA ...
 TT FTL ...
 TT TXT ...
 BBB NAN ...
 BBB SKT ...
 BBB PKW ...
 BBB P28 ...
 BBB CMR ...
 BBB LCK ...
 BBB PSB ...
 BBB PLH ...
 BBB NWN ...
 BBB MNG ...
 BBB HIN ...
 BBB HIN ...
 BBB PGM ...
 BBB AUK ...
 AA SAM ...
 AA MAL ...
 AAA HOM ...
 AAA BKS ...

TT_JCK	GTCAGAAATA TAGTGAATC GTCTTTCCCA TTCTTTCCC CGACCCCTGCT ACGAAGAAAAG ACGTTCACTT CTTAAATAT CCCATATACA TAGGGGAA CAGAGGAAGG GGTCAAATT [3840]
TT_FAA
TT_FTL
TT_TXT
BB_NAN
BB_SKT
BB_PKW
BB_P28
BB_CMR
BB_LCK
BBA_PSB
BBBA_PLH
BBA_NWN
BBA_MNG
BBA_HIN
ABB_HIN
ABB_PGM
AB_AUK
AA_SAM
AA_MAL
AAA_HOM
AA_BKS
TT_JCK	ATCCGTGATGG TAGAAAAGT AACAAATACAG TCTATATGC TACATCAGCA GGTATAGTA GCAGAATAGT ACGTAAAGAA AAAGGGGAT ATGAAATAAC CATAGTGAT GTATGGATG [3960]
TT_FAA
TT_FTL
TT_TXT
BB_NAN
BB_SKT
BB_PKW
BB_P28
BB_CMR
BBB_LCK
BBA_PSB
BBBA_PLH
BBA_NWN
BBA_MNG
BBA_HIN
ABB_HIN
ABB_PGM
AB_AUK
AA_SAM
AA_MAL
AAA_HOM
AA_BKS

[4080] GACATCAAGT GGTTGATATT ATACCTCCAG GACCAGAACT TCTTGTTCAG GAGGGTAAT CCATCAAGCT TGATCAACCA TTAACAAGCA ATCCCATAATGT AGGAGCTTT GGTCAGGGAG [4080]
TTT_FAA ... [4080]
TTT_FTL ... [4080]
TTT_TXT ... [4080]
BB_NAN ... [4080]
BB_SKT ... [4080]
BB_PKW ... [4080]
BB_P28 ... [4080]
BB_CMR ... [4080]
BBB_LCK ... [4080]
BBB_PSB ... [4080]
BBBA_PLH ... [4080]
BBBA_NWN ... [4080]
BBA_MNG ... [4080]
BBA_HIN ... [4080]
ABB_HIN ... [4080]
ABB_PGM ... [4080]
AB_AUK ... [4080]
AA_SAM ... [4080]
AA_MAL ... [4080]
AAA_HOM ... [4080]
AA_BKS ... [4080]
[4200] ATGAGAAAT AGTGCTTCAA GACCCATTAC GGGTCCAAGG TCTTTTGTTC TTCTTGGCAT CTGTTATTT GGCACAAAGT TTTTTGGTCA TTAAAAAGAA ACAGTTGAA AAGGTTCAAT [4200]
TT_JCK ... [4200]
TT_FAA ... [4200]
TT_FTL ... [4200]
TT_TXT ... [4200]
BB_NAN ... [4200]
BB_SKT ... [4200]
BB_PKW ... [4200]
BB_P28 ... [4200]
BB_CMR ... [4200]
BBB_LCK ... [4200]
BBB_PSB ... [4200]
BBBA_PLH ... [4200]
BBBA_NWN ... [4200]
BBA_MNG ... [4200]
BBA_HIN ... [4200]
ABB_HIN ... [4200]
ABB_PGM ... [4200]
AB_AUK ... [4200]
AA_SAM ... [4200]
AA_MAL ... [4200]
AAA_HOM ... [4200]
AA_BKS ... [4200]

TT_JCK	CGAATAAT ATTGATTAA TGTGACTGG CCAATTGTA TGTATGAA TAGATAAAA CCCCTCTA--	--TAAGA GTAAGAAAAG AACTCAACGG GACCTTA--	--TCGCTC
TT_FAA	[5280]
TT_FTL	[5280]
TT_TXT	[5280]
BB_NAN	[5280]
BB_SKT	[5280]
BB_PKW	[5280]
BB_P28	[5280]
BB_CMR	[5280]
BBB_LCK	[5280]
BBA_PSB	[5280]
BBB_PLH	[5280]
BBA_NWN	[5280]
BBA_MNG	[5280]
BBA_HIN	[5280]
ABB_HIN	[5280]
ABB_PGM	[5280]
AB_AUK	[5280]
AA_SAM	[5280]
AA_MAL	[5280]
AAA_HOM	[5280]
AA_BKS	[5280]
TT_JCK	TAATTAGACA AAGGAGAGTA AGGTCCGGTT GAGTTCTTT CATCTCTTT ATCCGATTACT AGAGAGATGA ACCAACCCA GAATATGAAC CGTAAAGAA	CGTAAAGAA	[5400]
TT_FAA	[5400]
TT_FTL	[5400]
TT_TXT	[5400]
BB_NAN	[5400]
BB_SKT	[5400]
BB_PKW	[5400]
BB_P28	[5400]
BB_CMR	[5400]
BBB_LCK	[5400]
BBA_PSB	[5400]
BBB_PLH	[5400]
BBA_NWN	[5400]
BBA_MNG	[5400]
BBA_HIN	[5400]
ABB_HIN	[5400]
ABB_PGM	[5400]
AB_AUK	[5400]
AA_SAM	[5400]
AA_MAL	[5400]
AAA_HOM	[5400]
AA_BKS	[5400]

รายงานผลการตรวจพยานร่องรอยทางชีววิทยาลักษณะ

รายการ	ชนิดของพยาน	รหัสพยาน	ผลการตรวจ
TTT_JCK	TTT_FAA	[57760]	AGTACAAAAA TGAGTAATAA ACCCCAGTAT AGACTGGTAC GATTCAATT AACATTTC TAATTCAAAT TAGGTTTAC TTGATTTGT CATAGCTCTA TCATTGGGT TCATTGGGT
BBB_NAN	BBB_SKT	[57760]
BBB_PKW	BBB_P28	[57760]
BBB_CMN	BBB_LCK	[57760]
BBA_NWN	BBA_MNG	[57760]
BBA_HIN	AAB_HIN	[57760]
AAB_PGM	AB_AUK	[57760]
AA_SAM	AA_MAL	[57760]
AA_HOM	AAA_BKS	[57760]
BB_CMN	BB_LCK	[5774]	CTGCATTGCT GATA [5774]
BBA_PSB	BBA_PLH	[5774]
BBA_NWN	BB_NWN	[5774]
BBA_MNG	BB_MNG	[5774]
BBA_HIN	BB_HIN	[5774]
AAB_HIN	AAA_HOM	[5774]
ABB_PGM	AA_BKS	[5774]
AB_AUK	AA_SAM	[5774]
AA_MAL	AA_HOM	[5774]
AAA_BKS	BB_NWN	[5774]

APPENDIX C

ALIGNED GBSS SEQUENCE DATA MATRIX OF *Musa*¹

¹ Accession codes appeared according to Table 4.1.

Dot symbol (.) shows a site with the same character variant to that of the first sequence; Question mark (?) indicates missing data.

BBA_NWN_20	???????????	???????????	?????????GTG	CAGGGTGGAG	TATTGCCGGC	TAGCTAGCTA	[60]
BBA_NWN_29	CACTACAAGG	ATGCATGGGA	CACAAGT	[60]
AAB_NGA_03	CACTACAAGG	ATGCATGGGA	TACAAGC	.G.	T.T.	[60]
AAB_NGA_10	CACTACAAGG	ATGCGTGGGA	TACAAGC	T.T.	[60]
BBA_NWN_64	CACTACAAGG	ATGCCTGGGA	TACAAGC	T.T.	[60]
AAB_NGA_82	CACTACAAGG	ATGCCTGGGA	CACAAGC	T.T.	[60]
ABB_HMO_95	???????????	???????????	???????????	???????????	???????????	???????????	[60]
BBA_NWS_13	???????????	???????????	???????????	???????????	???????????	???????????	[60]
AAB_KPL_31	???????????	???????????	???????????	???????????	???????????	???????????	[60]
BBA_NWM_96	CACTACAAGG	ATGCATGGGA	TACAAGC	T.T.	[60]
BBA_NWN_23	CACTACAAGG	ATGCATGGGA	TACAAGC	[60]
AAB_NGA_17	CACTACAAGG	ATGCCTGGGA	TACAAGC	[60]
AAB_NGA_14	CACTACAAGG	ATGCATGGGA	CACAAGC	----	[60]
AAB_NGA_02	CACTACAAGG	ATGCATGGGA	CACAAGC	T.T.	[60]
ABB_HMI_89	???????????	???????????	???????????	???????????	???????????	-----	[60]
BB_CMR_05	???????????	???????????	???????????	???????????	???????????	???????????	[60]
BBA_NWN_18	CACTACAAGG	ATGCCTGGGA	TACAAGC	[60]
BBA_HIN_99	???????????	???????????	???????????	-----	[60]
AAB_TBK_09	???????????	???????????	???????????	???????????	???????????	???????????	[60]
BBA_NWN_22	CACTACAAGG	ATGCATGGGA	CACAAGC	[60]
BBA_NWN_46	CACTACAAGG	ATGCGTGGGA	TACAAGC	[60]
BBA_NWN_66	CACTACAAGG	ATGCATGGGA	TACAAGC	[60]
BB_PKW_03	CACTACAAGG	ATGCGTGGGA	CACAAGC	[60]
BB_NAN_06	???????????	???????????	???????????	???????????	???????????	???????????	[60]
BB_NAN_11	???????????	???????????	???????????	???????????	???????????	???????????	[60]
BBA_NWN_20	GCTAG----	-AGATTGCAA	CACAAAGCAA	CCTGC-TGCA	TTTTGAAGTT	TGTGGTTATG	[120]
BBA_NWN_29	-----	[120]
AAB_NGA_03TCT	-----	[120]
AAB_NGA_10TCT	-----	[120]
BBA_NWN_64TCT	-----	[120]
AAB_NGA_82TCT	---	[120]
ABB_HMO_95	-----	-----	[120]
BBA_NWS_13	-----	-N..N...	[120]
AAB_KPL_31	???????????	???????????	?????????.CT	..C.N..N.	..GN..N.N.	..AN..	[120]
BBA_NWM_96CTATC	T.....	[120]
BBA_NWN_23	-----	[120]
AAB_NGA_17	-----	-A.	[120]
AAB_NGA_14TCT	-----	[120]
AAB_NGA_02	-----	-A.	[120]
ABB_HMI_89	???????????	???????????	?????????..A.-N.A.	[120]
BB_CMR_05	???????????	???????????	??.CC..N..N.G.G..	[120]
BBA_NWN_18	-----	[120]
BBA_HIN_99	???????????	???????????	???????????	-----	[120]
AAB_TBK_09	???????????	???????????	??.C..C.N..A..N..G..	[120]
BBA_NWN_22	-----	[120]
BBA_NWN_46	-----	[120]
BBA_NWN_66	-----	[120]
BB_PKW_03	-----	[120]
BB_NAN_06	???????????	???????????	???????????	-----	[120]
BB_NAN_11	???????????	???????????	???????????	-----	[120]

BBA_NWN_20	TGCAGTTGAA	AGTTGGGAAC	AGAGTCGAAA	CAGTTCGCTT	CTTCCACTGC	TACAAAAGGG	[180]
BBA_NWN_29	[180]
AAB_NGA_03	[180]
AAB_NGA_10	[180]
BBA_NWN_64	[180]
AAB_NGA_82	[180]
ABB_HMO_95	[180]
BBA_NWS_13	[180]
AAB_KPL_31	G..N..NN.	NN.N.....	N...G.A.C.	...N..N.N	T..T.....	.N.C.NN..	[180]
BBA_NWM_96	[180]
BBA_NWN_23	M.....	[180]
AAB_NGA_17	[180]
AAB_NGA_14	[180]
AAB_NGA_02	[180]
ABB_HMI_89	T .. T.....	[180]
BB_CMR_05	N.....	N.....	TTT.....	A .. GN.....	[180]
BBA_NWN_18	[180]
BBA_HIN_99	?.....?	?.....?	?.....?	?.....?	?.....?	?.....?	[180]
AAB_TBK_09NA.....	..NN.....	.C..N.....	..NA.....	T..N.....	N..NN.....	[180]
BBA_NWN_22	G.....	[180]
BBA_NWN_46	[180]
BBA_NWN_66	[180]
BB_PKW_03	[180]
BB_NAN_06	?.....?	?.....?	?.....?	?.....?	?.....?	?.....?	[180]
BB_NAN_11	?.....?	?.....?	?.....?	?.....?	?.....?	?.....?	[180]
BBA_NWN_20	GAGTCGACAG	GGTCTTCGTT	GATCACCTA	TGTTTCTTGC	CAAGGTTCGA	ACTATTTTAT	[240]
BBA_NWN_29	[240]
AAB_NGA_03	[240]
AAB_NGA_10	[240]
BBA_NWN_64	C.....	[240]
AAB_NGA_82	C.....	[240]
ABB_HMO_95	[240]
BBA_NWS_13	Y.....	[240]
AAB_KPL_31NNNNNGNN.NGGN.....NNC..NNNNN....GNCNN.....	[240]
BBA_NWM_96	[240]
BBA_NWN_23	C.....	[240]
AAB_NGA_17	[240]
AAB_NGA_14	[240]
AAB_NGA_02	[240]
ABB_HMI_89	...NN.N.....NNN.....NN.G.....C.....	[240]
BB_CMR_05N.....NNN.....NN.G.....C.....	[240]
BBA_NWN_18	[240]
BBA_HIN_99	.TCA.C.NG.	TT.T.CGT..C.....N..TC.....T.....	[240]
AAB_TBK_09TN.....	..G..N..NG.....N..N.N.....NN..N.....	[240]
BBA_NWN_22	[240]
BBA_NWN_46	[240]
BBA_NWN_66	[240]
BB_PKW_03	[240]
BB_NAN_06	[240]
BB_NAN_11	[240]
BBA_NWN_20	ATCTGATGCC	GACTTCAGAT	AACATTTCGA	GTGGCATAAA	-GCTGACCAC	GGATCTGTGC	[300]
BBA_NWN_29	T.....	[300]
AAB_NGA_03	-T.T.....	[300]
AAB_NGA_10	-..T.....	[300]
BBA_NWN_64	-..T.....	[300]
AAB_NGA_82	-..T.....	[300]
ABB_HMO_95	-..T.....	[300]
BBA_NWS_13	-..T.....	[300]
AAB_KPL_31	C.N....N.	NN..NNNN.	N..N.NNAN.	C..NN..N	-GN.....	T.NNNNNNN	[300]
BBA_NWM_96	-..T.....	[300]
BBA_NWN_23	-..T.....	[300]
AAB_NGA_17	-	[300]
AAB_NGA_14	-	A.....	[300]
AAB_NGA_02	-	[300]
ABB_HMI_89	-	[300]
BB_CMR_05	N.....	-	[300]
BBA_NWN_18	-	[300]
BBA_HIN_99	N.....	N..-G..	AG.....	A..N.....	A.....	[300]
AAB_TBK_09N.....	..N..N..K.....	-..N..N.....	[300]
BBA_NWN_22	-	[300]
BBA_NWN_46	-	A.....	[300]
BBA_NWN_66	-	[300]
BB_PKW_03	-	[300]
BB_NAN_06	-	[300]
BB_NAN_11	-	[300]

BBA_NWN_20	GT-----GTA	AGGTGTGGGG	AAAGACCGGA	GGAAAGATAT	ATGGTCCTGC	CACAGGAACA	[360]
BBA_NWN_29	[360]
AAB_NGA_03	..GTAAG..	T ..T.....	[360]
AAB_NGA_10	..GTAAG..	T ..T.....	[360]
BBA_NWN_64	..GTAAG..	[360]
AAB_NGA_82	..GTAAG..	T ..T.....	[360]
ABB_HMO_95	A..GTAAG..	T ..T.....	[360]
BBA_NWS_13	..GTAAG..	T ..T.....	[360]
AAB_KPL_31	.NNTTAG..NG..NN	A.N..NC..	N..G..NN	TNN..NNN.C	[360]
BBA_NWM_96	..GTAAG..	T ..T.....	[360]
BBA_NWN_23	N.....	N.N..N..	[360]
AAB_NGA_17	[360]
AAB_NGA_14	[360]
AAB_NGA_02	[360]
ABB_HMI_89	[360]
BB_CMР_05	N..	[360]
BBA_NWN_18	[360]
BBA_HIN_99	N..C..NC..	[360]
AAB_TBK_09	R.....NN..	[360]
BBA_NWN_22	[360]
BBA_NWN_46	[360]
BBA_NWN_66	[360]
BB_PKW_03	[360]
BB_NAN_06	[360]
BB_NAN_11	[360]

BBA_NWN_20	GATTACGAAG	ACAACCAGCA	AAGATTCAAGC	CTTTCTGTC	AGGTCAGTGA	TTATTAATTA	[420]
BBA_NWN_29	[420]
AAB_NGA_03	G..T..	[420]
AAB_NGA_10	G..T..	[420]
BBA_NWN_64	[420]
AAB_NGA_82	G..T..	[420]
ABB_HMO_95C.....	[420]
BBA_NWS_13	G..T..	[420]
AAB_KPL_31	NNN.N..NC..	..T...NN.	C.NNN..	N..TT..T..	NG.....N.N	[420]
BBA_NWM_96	T...C	[420]
BBA_NWN_23	[420]
AAB_NGA_17	C..	[420]
AAB_NGA_14	[420]
AAB_NGA_02	[420]
ABB_HMI_89	[420]
BB_CMР_05	[420]
BBA_NWN_18	[420]
BBA_HIN_99T.NN..N..	C..	[420]
AAB_TBK_09N..	N..	[420]
BBA_NWN_22	[420]
BBA_NWN_46	[420]
BBA_NWN_66	[420]
BB_PKW_03	[420]
BB_NAN_06	...C..	[420]
BB_NAN_11	[420]

BBA_NWN_20	CTAAACAAA	TTCAGCATCG	ATTTCTAAGT	TACACTGCGA	AGCAATTA--	--GTG	[480]
BBA_NWN_29	--	--	[480]
AAB_NGA_03	T..	T..	[480]
AAB_NGA_10	T..	[480]
BBA_NWN_64	G--	[480]
AAB_NGA_82	T..	[480]
ABB_HMO_95	[480]
BBA_NWS_13	T.C..	[480]
AAB_KPL_31	..NN..NNC..	NN.C..NNN	N..N..N..A..N..N..N..	CN.TG..	[480]
BBA_NWM_96	[480]
BBA_NWN_23	...R.Y..	CC	GAACCTTT..	[480]
AAB_NGA_17	...G..	CT	GAACCTTT..	[480]
AAB_NGA_14	...G..	CT	GAACCTTT..	[480]
AAB_NGA_02	...G..	CT	GAACCTTT..	[480]
ABB_HMI_89	...G..	CT	GAACCTTT..	[480]
BB_CMР_05	...G..	CC	GAACCTTT..	[480]
BBA_NWN_18	.C.G..	G..	CC	GAACCTTT..	[480]
BBA_HIN_99	N.G..	T..	G..T..N.N..	N..CC	GAACCTTT.A	[480]
AAB_TBK_09	...R.T..	Y..	CC	GAACCTTN..	[480]
BBA_NWN_22	...G.T..	CC	GAACCTTT..	[480]
BBA_NWN_46	...G.T..	CC	GAACCTTT..	[480]
BBA_NWN_66	...G.T..	CC	GAACCTTT..	[480]
BB_PKW_03	...G.T..	CC	GAACCTTT..	[480]
BB_NAN_06	...G..	CT	GAACCTTT..	[480]
BB_NAN_11	...G..	CT	GAACCTTT..	[480]

BBA_NWN_20	AAACCTACAT	CTACTACTCT	TATGGATCTC	TCTGCCTTG	ACAAAATCAT	GGATGACCTT	[540]
BBA_NWN_29	[540]
AAB_NGA_03	.	.	.	--.	.	.	[540]
AAB_NGA_10	.	.	.	--.	T.	.	[540]
BBA_NWN_64	.	.	.	--.	.	.	[540]
AAB_NGA_82	.	.	.	--.	.	.	[540]
ABB_HMO_95	.	.	.	--.	.	.	[540]
BBA_NWS_13	.	R.	.	T.G.N.T	T.	.	[540]
AAB_KPL_31	.	N.	.	T.T	-N.	-	[540]
BBA_NWM_96	.	.	.	--.	.	.	[540]
BBA_NWN_23	.	-	-	-	-	-	C [540]
AAB_NGA_17	.	G-	-	-	-	-	C [540]
AAB_NGA_14	.	G-	-	-	-	-	C [540]
AAB_NGA_02	.	G-	-	-	-	-	C [540]
ABB_HMI_89	.	G-	-	-	-	-	C [540]
BB_CMR_05	.	Y-	-	-	-	N..C	[540]
BBA_NWN_18	.	-	-	-	-	-	[540]
BBA_HIN_99	.	-	-	-	-	A..NN	[540]
AAB_TBK_09	.	-	-	-	-	-	C [540]
BBA_NWN_22	.	-	-	-	-	-	C [540]
BBA_NWN_46	.	-	-	-	-	-	C [540]
BBA_NWN_66	.	-	-	-	-	-	C [540]
BB_PKW_03	.	-	-	-	-	-	C [540]
BB_NAN_06	.	G-	-	-	-	-	C [540]
BB_NAN_11	.	G-	-	-	-	-	C [540]
BBA_NWN_20	TCCAGGCAGC	TTTAGAACCG	CCAAGGTCC	TACATCTAA	CAACAGTGAA	TACTATTCCG	[600]
BBA_NWN_29	.	.	.	T.	.	.	[600]
AAB_NGA_03	.	.	T.	T.	.	G.	[600]
AAB_NGA_10	.	.	T.	T.	.	.	[600]
BBA_NWN_64	.	.	.	T.	.	.	[600]
AAB_NGA_82	.	.	T.	T.	.	.	[600]
ABB_HMO_95	.	.	T.	T.	.	.	[600]
BBA_NWS_13	.	.	T.	T.	.	.	[600]
AAB_KPL_31	.	.	T.	T.	.	.	[600]
BBA_NWM_96	.	.	T.	T.	.	.	[600]
BBA_NWN_23	G.	.	.	YT.	.	.	[600]
AAB_NGA_17	G.	.	.	T.	.	.	[600]
AAB_NGA_14	G.	.	.	T.	.	.	[600]
AAB_NGA_02	G.	.	.	T.	.	.	[600]
ABB_HMI_89	G.	.	.	T.	.	.	[600]
BB_CMR_05	G.	.	.	T.	.	.	[600]
BBA_NWN_18	G.	.	.	T.	.	.	[600]
BBA_HIN_99	NG..N.	N.	.	T.	T.	NN	[600]
AAB_TBK_09	G.	.	.	TG	.	.	[600]
BBA_NWN_22	G.	.	.	T.	.	.	[600]
BBA_NWN_46	G.	.	.	T.	.	.	[600]
BBA_NWN_66	G.	.	.	T.	.	.	[600]
BB_PKW_03	G.	.	.	T.	.	.	[600]
BB_NAN_06	G.	.	.	T.	.	.	[600]
BB_NAN_11	G.	.	.	T.	.	.	[600]
BBA_NWN_20	GGCCATATGG	TACGATTCTT	ATTTGTTGC	TTCTCCCGAT	GGCTTATGTC	GTAAACATTG	[660]
BBA_NWN_29	[660]
AAB_NGA_03	G.	[660]
AAB_NGA_10	[660]
BBA_NWN_64	[660]
AAB_NGA_82	[660]
ABB_HMO_95	T.	.	[660]
BBA_NWS_13	[660]
AAB_KPL_31	C.C.	C..	[660]
BBA_NWM_96	C.C.	C..	[660]
BBA_NWN_23	G..	[660]
AAB_NGA_17	G..	[660]
AAB_NGA_14	G..	[660]
AAB_NGA_02	G..	[660]
ABB_HMI_89	G..	[660]
BB_CMR_05	N.	GN..	[660]
BBA_NWN_18	G..	[660]
BBA_HIN_99	.	N..N.	N..	N..	N..	N..	[660]
AAB_TBK_09	.	N..	.	N..	N..	N..	[660]
BBA_NWN_22	G..	[660]
BBA_NWN_46	G..	[660]
BBA_NWN_66	G..	[660]
BB_PKW_03	G..	[660]
BB_NAN_06	G..	[660]
BB_NAN_11	G..	[660]

BBA_NWN_20 TGAAAA----- -CTCATATGA TGTGTTCCAT GCAGGTACGA CAAGCC????? [1080]
 BBA_NWN_29G.TTCT CTTGATGAAA A..... .G....CGTG [1080]
 AAB_NGA_03G.TTCT CTTGATGAAA A..... .G....CGTG [1080]
 AAB_NGA_10G.TTCT CTTGATGAAA A..... .G....CGTG [1080]
 BBA_NWN_64G.TTCT CTTGATGAAA A..... .GGTG [1080]
 AAB_NGA_82 -..... .GGTG [1080]
 ABB_HMO_95G.TTCT CTTGATGAAA AN..N... N..NGG? ?????????? ?????????? [1080]
 BBA_NWS_13G.TTCT CTTGATGAAA AN..N... N..NG? ?????????? ?????????? [1080]
 AAB_KPL_31G.TTCT CTTGATGAAA A..N... A.GG? ?????????? ?????????? [1080]
 BBA_NWM_96G.TTCT CTTGATGAAA A..... .G....CGTG [1080]
 BBA_NWN_23G.TTCT CTTGATGAAA A..... .G....CGTG [1080]
 AAB_NGA_17G.TTCT CTTGATGAAA A..... G... .G....CGTG [1080]
 AAB_NGA_14G.TTCT CTTGATGAAA A..... G... .G....CGTG [1080]
 AAB_NGA_02 -..... .GGTG [1080]
 ABB_HMI_89A--- -C...N ..NGG? ?????????? ?????????? [1080]
 BB_CMR_05 -C.N.N ..N.A.G?? ?????????? ?????????? [1080]
 BBA_NWN_18 -..... .GGTG [1080]
 BBA_HIN_99 ...??????? ?????????? ?????????? ?????????? ?????????? ?????????? [1080]
 AAB_TBK_09 ...??????? ?????????? ?????????? ?????????? ?????????? ?????????? [1080]
 BBA_NWN_22 -..... .GGTG [1080]
 BBA_NWN_46 -..... .GGTG [1080]
 BBA_NWN_66 -..... .GGTG [1080]
 BB_PKW_03 -..... .GGTG [1080]
 BB_NAN_06 ?????????? ?????????? ?????????? ?????????? ?????????? ?????????? [1080]
 BB_NAN_11 ?????????? ?????????? ?????????? ?????????? ?????????? ?????????? [1080]
 BBA_NWN_20 ?????????? ?????????? ?????????? ?? [1112]
 BBA_NWN_29 AAAGGAAGGA AAATTAACGT GATGAAGGCT AA [1112]
 AAB_NGA_03 AAAGGAAGGA AGATAAATTG GATGAAGGCT GG [1112]
 AAB_NGA_10 AAAGGAAGGA AAATCAATTG GATGAAGGCT GG [1112]
 BBA_NWN_64 AAAGGAAGGA AAATCAACGT GATGAAGGCT GG [1112]
 AAB_NGA_82 AAAGGAAGGA AAATTAACGT GATGAAGGCT GG [1112]
 ABB_HMO_95 ?????????? ?????????? ?????????? ?? [1112]
 BBA_NWS_13 ?????????? ?????????? ?????????? ?? [1112]
 AAB_KPL_31 ?????????? ?????????? ?????????? ?? [1112]
 BBA_NWM_96 AAAGGAAGGA AGATAAATTG GATGAAGGCT GG [1112]
 BBA_NWN_23 AAAGGAAGGA AGATCAACGT GATGAAGGCT GG [1112]
 AAB_NGA_17 AAAGGAAGGA AAATAAATTG GATGAAGGCT GG [1112]
 AAB_NGA_14 AAAGGAAGGA AAATAAATTG GATGAAGGCT GG [1112]
 AAB_NGA_02 AAAGGAAGGA AGATAAATTG GATGAAGGCT GG [1112]
 ABB_HMI_89 ?????????? ?????????? ?????????? ?? [1112]
 BB_CMR_05 ?????????? ?????????? ?????????? ?? [1112]
 BBA_NWN_18 AAAGGAAGGA AGATAAATTG GATGAAGGCT GG [1112]
 BBA_HIN_99 ?????????? ?????????? ?????????? ?? [1112]
 AAB_TBK_09 ?????????? ?????????? ?????????? ?? [1112]
 BBA_NWN_22 AAAGGAAGGA AGATCAACGT GATGAAGGCT GG [1112]
 BBA_NWN_46 AAAGGAAGGA AAATAAACGT GATGAAGGCT GG [1112]
 BBA_NWN_66 AAAGGAAGGA AGATAAACGT GATGAAGGCT GG [1112]
 BB_PKW_03 AAAGGAAGGA AAATCAATTG GATGAAGGCT GG [1112]
 BB_NAN_06 ?????????? ?????????? ?????????? ?? [1112]
 BB_NAN_11 ?????????? ?????????? ?????????? ?? [1112]

ศูนย์วิทยาพรพยากร
จุฬาลงกรณ์มหาวิทยาลัย

BIOGRAPHY

Sasivimon Swangpol (maiden name, Chomchalow) was born in Bangkok, Thailand, on 13 August 1966. She earned her Bachelor Degree in Science with Honors in Botany from the Department of Botany, Faculty of Science, Chulalongkorn University, Bangkok, in 1988. In 1991, she received her Master of Science Degree in Horticulture from the Horticulture Department, Institute of Florida Agricultural Science, University of Florida, Gainesville, Florida, USA. After graduation, she worked as a general manager at the Siam Flower Lab Company Limited for 3 years. Since March 1995, she has joined the Department of Plant Science, Faculty of Science, Mahidol University. In 1999, she pursued her Ph.D. study in Biological Science Ph.D. Program, Faculty of Science, Chulalongkorn University. She was appointed for a research assistantship by the Biodiversity Research and Training Program (BRT), the Thailand Research Fund (TRF) in 2002.

Sasivimon Swangpol is married to Mr. Jackrawut Swangpol and has two daughters, Malaivee, 8, and Mapa, 6. They reside in Bangkok.

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