Table A5.2: The genes we analysed in our study, including the assignment of our subset of protein-coding genes to three clusters we identified in our topology-clustering analysis. Abbreviations: IGS = intergenic spacer.

Gene	Туре	Cluster ID
atpA	Coding	1
<i>atp</i> B	Coding	2
<i>atp</i> E	Coding	3
<i>atp</i> F	Coding	1
<i>atp</i> H	Coding	3 2
atpl	Coding	2
ccsA	Coding	_
cemA	Coding	_ 2
matK	Coding	1
ndhA	Coding	2 2
ndhB	Coding	2
ndhC	Coding	_ 2
ndhD	Coding	2
ndhE	Coding	_
ndhF	Coding	1
ndhG	Coding	3
<i>ndh</i> H	Coding	3
ndhl	Coding	3
ndhJ	Coding	_
ndhK	Coding	- - 2 3
<i>pet</i> B	Coding	2
<i>pet</i> D	Coding	3
petG	Coding	_
petL	Coding	_
<i>pet</i> N	Coding	3 2
psaA	Coding	2
<i>psa</i> B	Coding	1
<i>psa</i> C	Coding	3
psal	Coding	_
<i>psa</i> J	Coding	3
psbA	Coding	2
<i>psb</i> B	Coding	2 2 2
<i>psb</i> C	Coding	2
<i>psb</i> D	Coding	2
<i>psb</i> E	Coding	_
<i>psb</i> F	Coding	_
<i>psb</i> G	Coding	_
<i>psb</i> H	Coding	_
psbl	Coding	_
<i>psb</i> J	Coding	3
<i>psb</i> K	Coding	_
<i>psb</i> L	Coding	3
<i>psb</i> M	Coding	3

<i>psb</i> N	Coding	_
psbT	Coding	3
psbZ	Coding	3
<i>rbc</i> L	Coding	3
rp/14	Coding	_
rpl16	Coding	2
rpl2	Coding	3
rpl20	Coding	2 3 2 2
rp/22	Coding	2
rpl23	Coding	_
rp/32	Coding	_
rp/33	Coding	
rpl36	Coding	_ _
•	Coding	2
rpoA		1
rpoB	Coding	
rpoC1	Coding	2
rpoC2	Coding	1
<i>rps</i> 11	Coding	1
rps12	Coding	3
rps14	Coding	_
<i>rps</i> 15	Coding	3
<i>rps</i> 16	Coding	_
<i>rps</i> 18	Coding	3
<i>rps</i> 19	Coding	3
rps2	Coding	_
rps3	Coding	2
rps4	Coding	1
rps7	Coding	3
rps8	Coding	3
ycf2	Coding	1
ycf3	Coding	3
ycf4	Coding	_
atpF–atpA	IGS	_
<i>ccs</i> A– <i>trn</i> L_UAG	IGS	_
cemA-ycf4	IGS	_
matK-trnK UUU	IGS	_
ndhB–rps7	IGS	_
ndhD-psaC	IGS	_
ndhE-ndhG	IGS	_
ndhG-ndhl	IGS	_
ndhH–rps15	IGS	_
ndhl-ndhA	IGS	
petA-cemA	IGS	_
•	IGS	_
petB-petD		_
petD-rpoA	IGS	_
petG-petL	IGS	_
petN-trnC_GCA	IGS	_
psaB-rps14	IGS	_

psbN-psbH psbT-psbN	IGS IGS	
psbT-psbN	IGS	_
rbcL-atpB	IGS	_
rpl14-rpl16	IGS	_
rpl22-rps19	IGS	_
rpoA-rps11	IGS	_
rpoC1-rpoC2	IGS	_
rpoC2-rps2	IGS	_
rps11-rpl36	IGS	_
rps12–rpl20	IGS	_
rps12-rps7	IGS	_
rps12-trnV_GAC	IGS	_
rps14-trnfM_CAU	IGS	_
rps2-atpl	IGS	_
rrn16-trnV_GAC	IGS	_
rrn4.5-rrn23	IGS	_
rrn4.5–rrn5	IGS	_
rrn5-trnR_ACG	IGS	_
trnA_UGC-rrn23	IGS	_
trnA_UGC-trnl_GAU	IGS	_
trnD_GUC-psbM	IGS	_
trnE_UUC-trnY_GUA	IGS	_
trnl_CAU-rpl23	IGS	_
trnl_CAU-ycf2	IGS	_
trnl_GAU-rrn16	IGS	_
trnL_CAA-ndhB	IGS	_
trnQ_UUG-psbK	IGS	_
trnR_ACG-trnN_GUU	IGS	_
trnV_UAC-trnV_UAC	IGS	_
trnW_CCA-petG	IGS	_
trnY_GUA-trnD_GUC	IGS	_
ycf15-trnL_CAA	IGS	_
ycf2–ycf15	IGS	_
ycf4–psal	IGS	_
ndhB	Intron	_
rpl2	Intron	_
rpoC1	Intron	_
ycf3	Intron	_