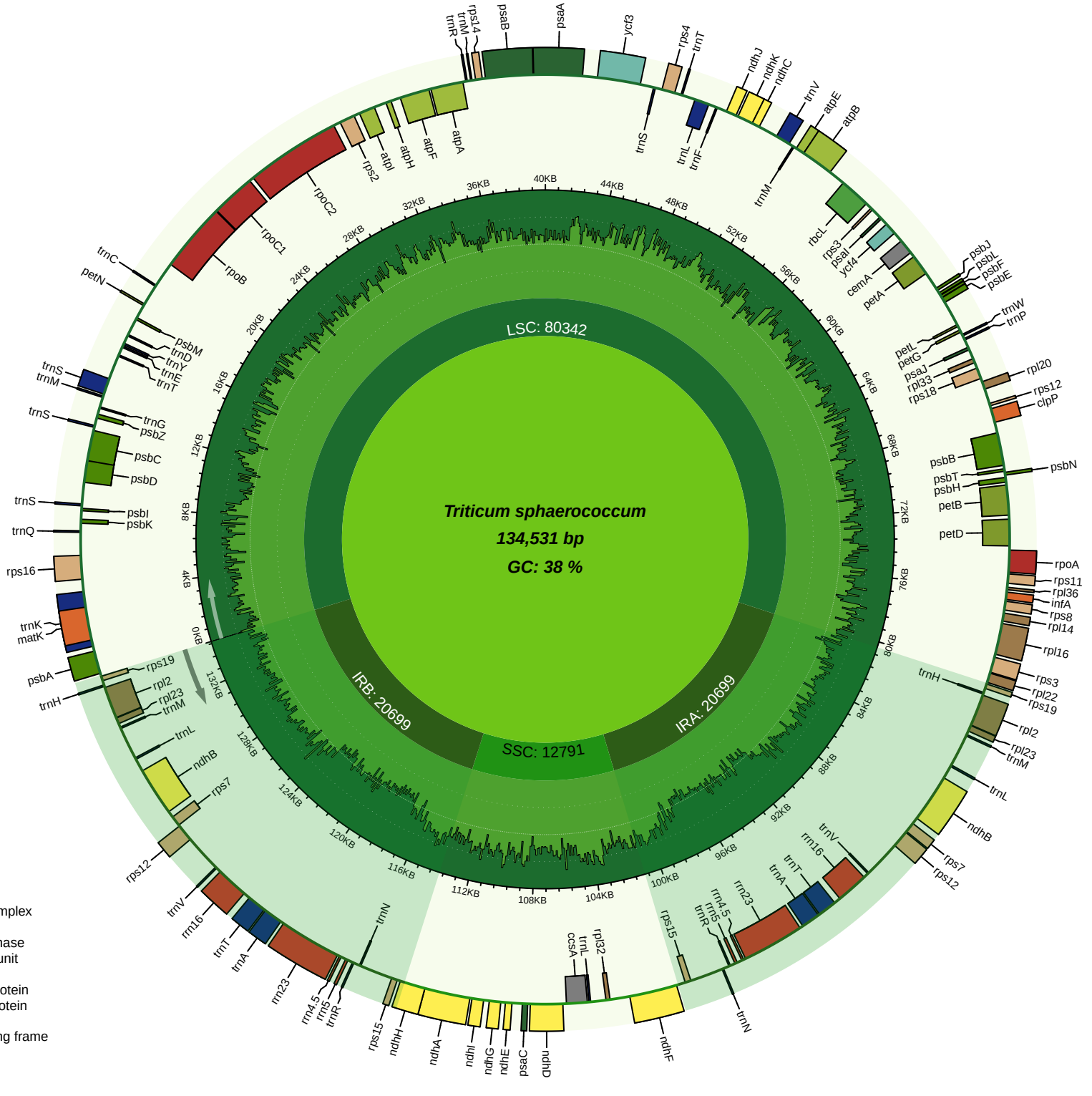
Triticum sphaerococcum
134,531 bp
GC: 38 %
LSC: 80342
IRB: 20699
IRA: 20699
SSC: 12791
0KB
4KB
8KB
12KB
16KB
20KB
24KB
28KB
32KB
36KB
40KB
44KB
48KB
52KB
56KB
60KB
64KB
68KB
72KB
76KB
80KB
84KB
88KB
92KB
96KB
100KB
104KB
108KB
112KB
116KB
120KB
124KB
128KB
132KB
psaB
psaA
ycf3
rps4
trnT
ndhJ
ndhK
ndhC
trnV
atpE
atpB
trnS
trnL
trnF
trnM
rbcL
rps3
psaI
ycf4
cemA
petA
psbJ
psbL
psbF
psbE
trnW
trnP
petL
petG
psaJ
rpl33
rps18
rpl20
rps12
clpP
psbB
psbT
psbH
petB
petD
psbN
rpoA
rps11
rpl36
infA
rps8
rpl14
rpl16
trnH
rps3
rpl22
rps19
rpl2
rpl23
trnM
trnL
ndhB
rps7
rps12
trnV
rrn16
trnT
trnA
rrn23
rrn4.5
rrn5
trnR
trnN
rps15
ndhF
rpl32
trnL
ccsA
ndhD
psaC
ndhE
ndhG
ndhI
ndhA
ndhH
rps15
trnN
rrn4.5
rrn5
trnR
rrn23
trnA
trnT
rrn16
trnV
rps12
rps7
ndhB
trnL
rpl23
trnM
rpl2
rps19
rps22
rpl16
rpl14
rps8
infA
rpl36
rps11
rpoA
petD
petB
psbH
psbT
psbB
psbN
clpP
rps12
rpl20
rpl33
rps18
psaJ
petG
petL
trnP
trnW
psbE
psbF
psbL
psbJ
petA
cemA
ycf4
psaI
rbcL
rpoC2
rpoC1
rpoB
rps2
atpI
atpH
atpF
atpA
trnR
trnM
rps14
psaB
psaA
trnC
petN
psbM
trnD
trnY
trnE
trnT
trnS
trnM
trnS
trnG
psbZ
psbC
psbD
trnS
psbI
psbK
trnQ
rps16
trnK
matK
psbA
trnH
rps19
rpl2
rpl23
trnM
trnL
ndhB
rps7
rps12
trnV
rrn16
trnT
trnA
rrn23
rrn4.5
rrn5
trnR
trnN
rps15
ndhH
ndhA
ndhI
ndhG
ndhE
psaC
ndhD
ccsA
trnL
rpl32
rps15
ndhF
trnN
rrn4.5
rrn5
trnR
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trnA
trnT
rrn16
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ndhB
trnL
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trnM
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rps8
infA
rpl36
rps11
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petD
petB
psbH
psbT
psbB
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clpP
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rpl20
rpl33
rps18
psaJ
petG
petL
trnP
trnW
psbE
psbF
psbL
psbJ
petA
cemA
ycf4
psaI
rbcL
rpoC2
rpoC1
rpoB
rps2
atpI
atpH
atpF
atpA
trnR
trnM
rps14
psaB
psaA
photosystem I
photosystem II
cytochrome b/f complex
ATP synthesis
NADH dehydrogenase
RubisCO larg subunit
RNA polymerase
small ribosomal protein
large ribosomal protein
clpP, matK, infA
hypothetical reading frame
transfer RNA
ribosomal RNA
other