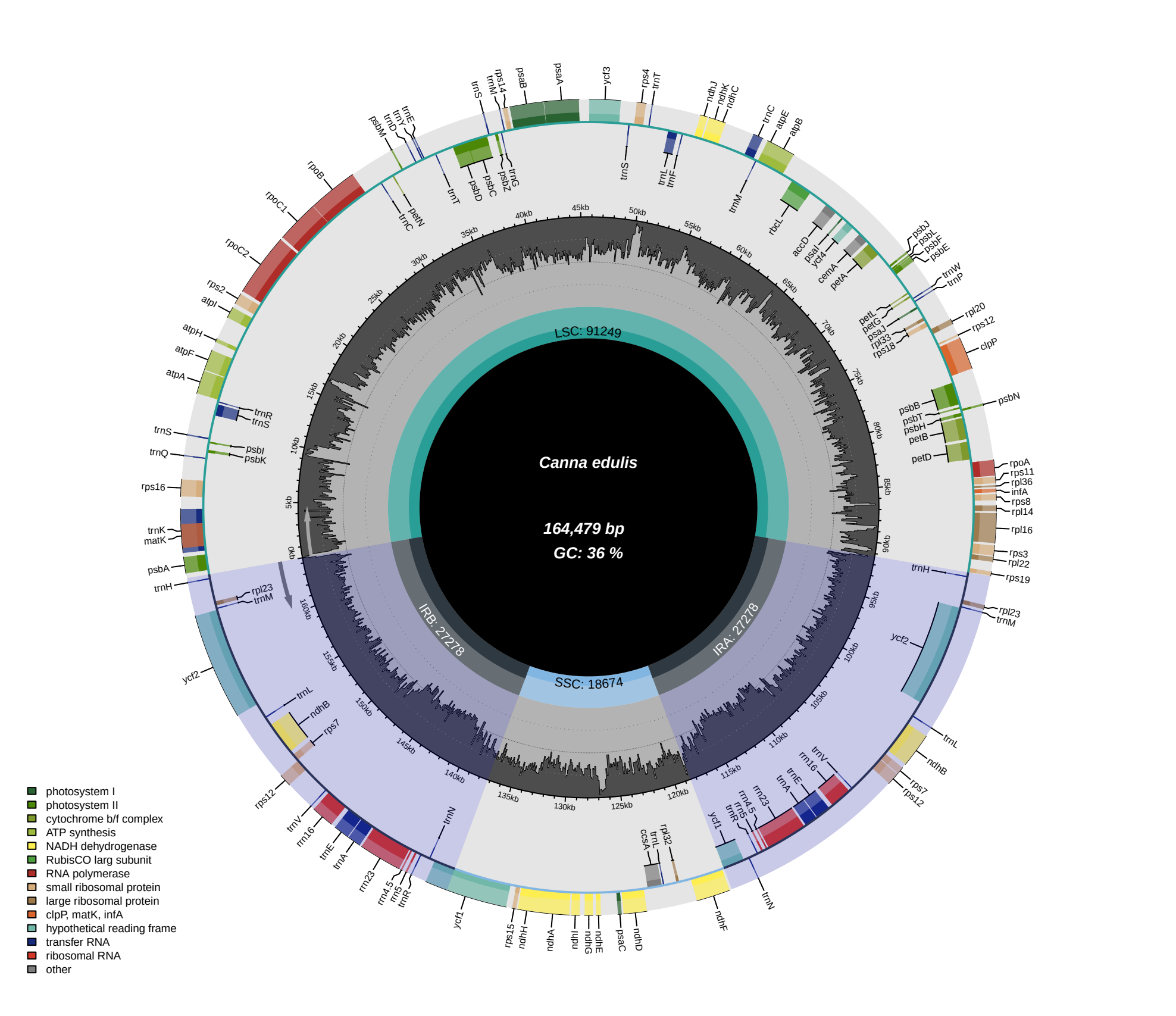
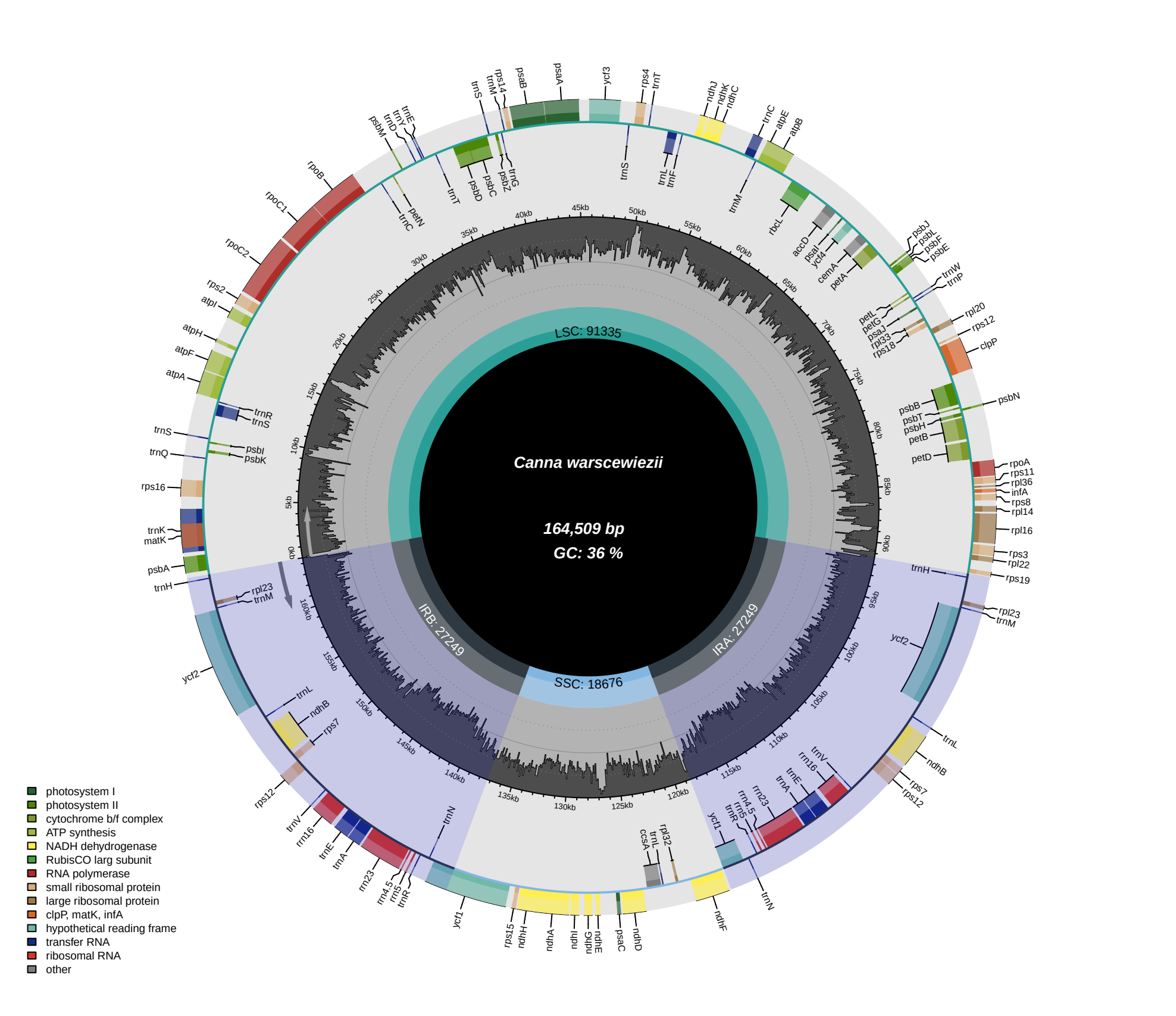
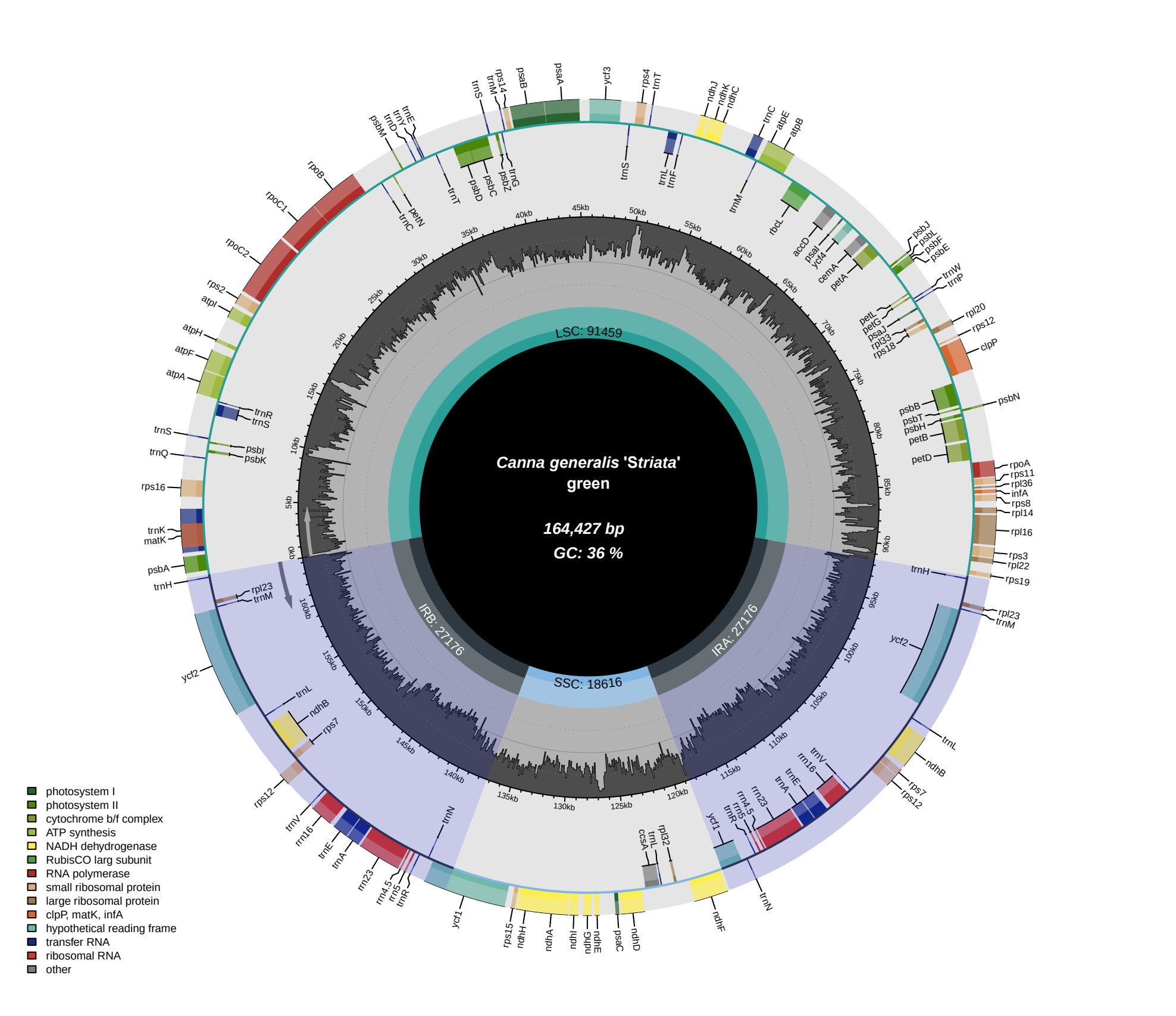
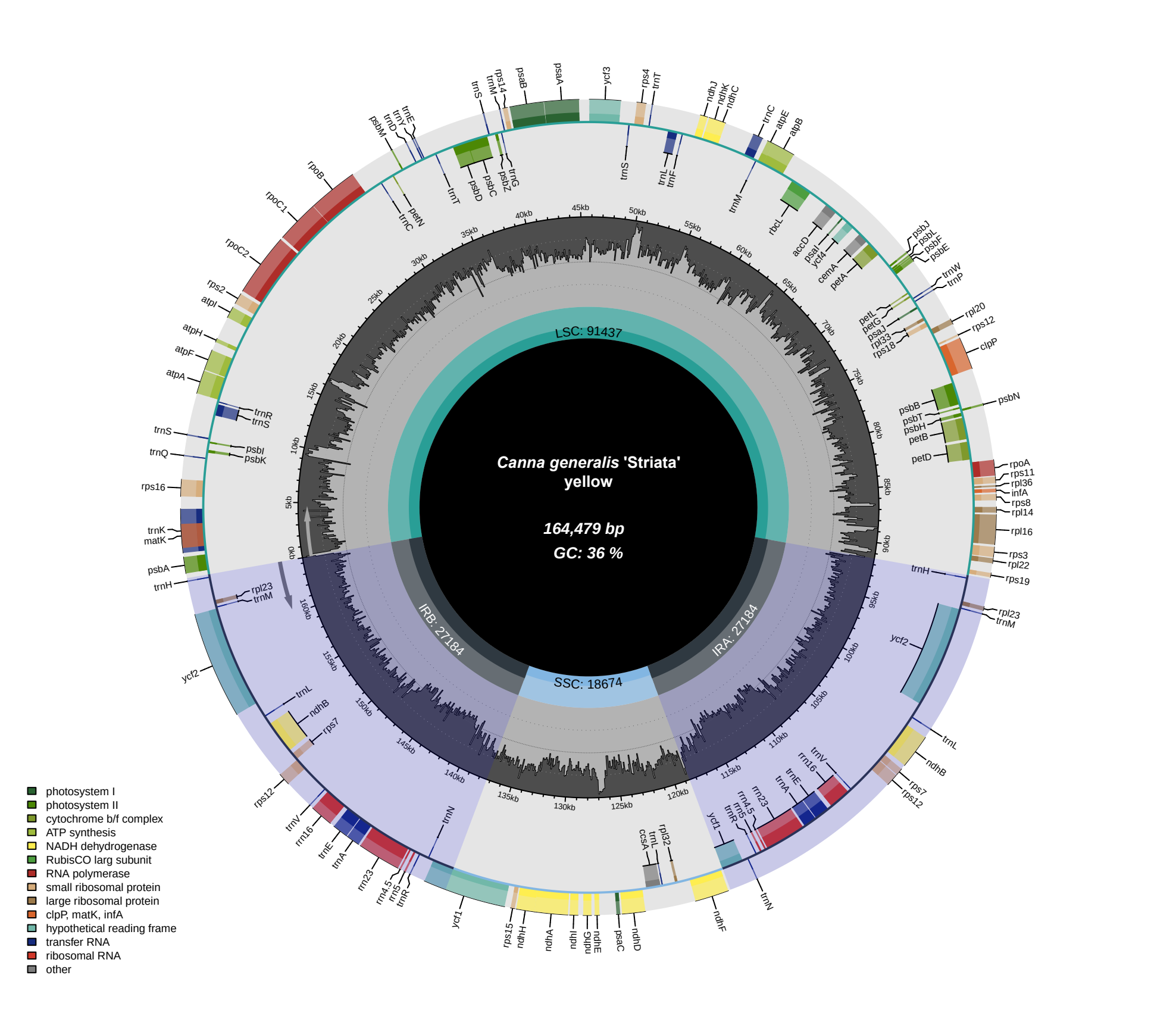

Canna edulis
164,479 bp
GC: 36 %
LSC: 91249
IRB: 27278
IRA: 27278
SSC: 18674
photosystem I
photosystem II
cytochrome b/f complex
ATP synthesis
NADH dehydrogenase
RubisCO larg subunit
RNA polymerase
small ribosomal protein
large ribosomal protein
clpP, matK, infA
hypothetical reading frame
transfer RNA
ribosomal RNA
other


Canna warscewiezii
164,509 bp
GC: 36 %
LSC: 91335
IRB: 27249
IRA: 27249
SSC: 18676
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


Canna generalis 'Striata'
green
164,427 bp
GC: 36 %
LSC: 91459
IRB: 27176
IRA: 27176
SSC: 18616
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


Canna generalis 'Striata'
yellow
164,479 bp
GC: 36 %
LSC: 91437
IRB: 27184
IRA: 27184
SSC: 18674
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