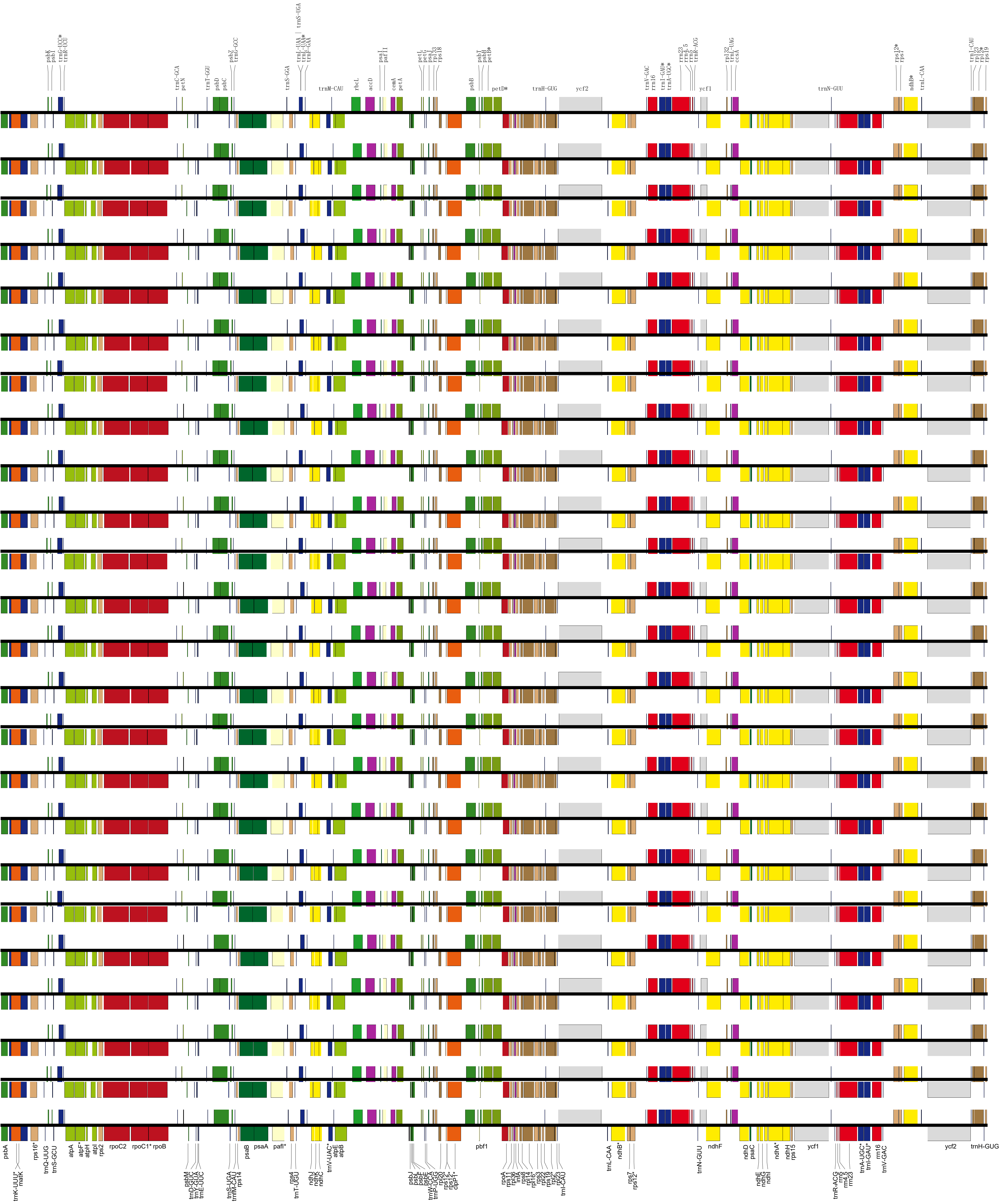




<i>Pholidota chinensis</i>	159,122 bp
<i>Otochilus fuscus</i>	160,053 bp
<i>Otochilus porrectus</i>	160,078 bp
<i>Pholidotaimbricata</i>	159,238 bp
<i>Neogyna gardneriana</i>	160,071 bp
<i>Coelogyne punctulata</i>	159,539 bp
<i>Coelogyne corymbosa</i>	159,732 bp
<i>Coelogyne cristata</i>	159,833 bp
<i>Coelogyne flaccida</i>	159,996 bp
<i>Coelogyne viscosa</i>	160,081 bp
<i>Coelogyne barbata</i>	160,093 bp
<i>Panisea tricallosa</i>	159,333 bp
<i>Ischnogyne mandarinorum</i>	159,487 bp
<i>Pholidota protracta</i>	159,781 bp
<i>Bulleyia yunnanensis</i>	159,581 bp
<i>Dendrochilum apoense</i>	160,175 bp
<i>Dendrochilum cootesii</i>	159,908 bp
<i>Chelonistele sulphurea</i>	160,280 bp
<i>Pholidota ventricosa</i>	159,418 bp
<i>Coelogyne rochussenii</i>	159,592 bp
<i>Pleione maculata</i>	158,394 bp
<i>Bletilla striata</i>	159,491 bp
<i>Thunia alba</i>	159,637 bp
<i>Thuniopsis cleistogama</i>	159,557 bp

- photosystem I

photosystem II

cytochrome b/f complex

ATP synthase

NADH dehydrogenase

RubisCO large subunit
- photosystem assembly/stability factors

RNA

polymerase

ribosomal proteins (SSU)

ribosomal proteins (LSU)

transfer RNAs

ribosomal RNAs
- clpP, matK

other genes

hypothetical chloroplast reading

frames (ycf) ORFs

origin of replication

polycistronic transcripts