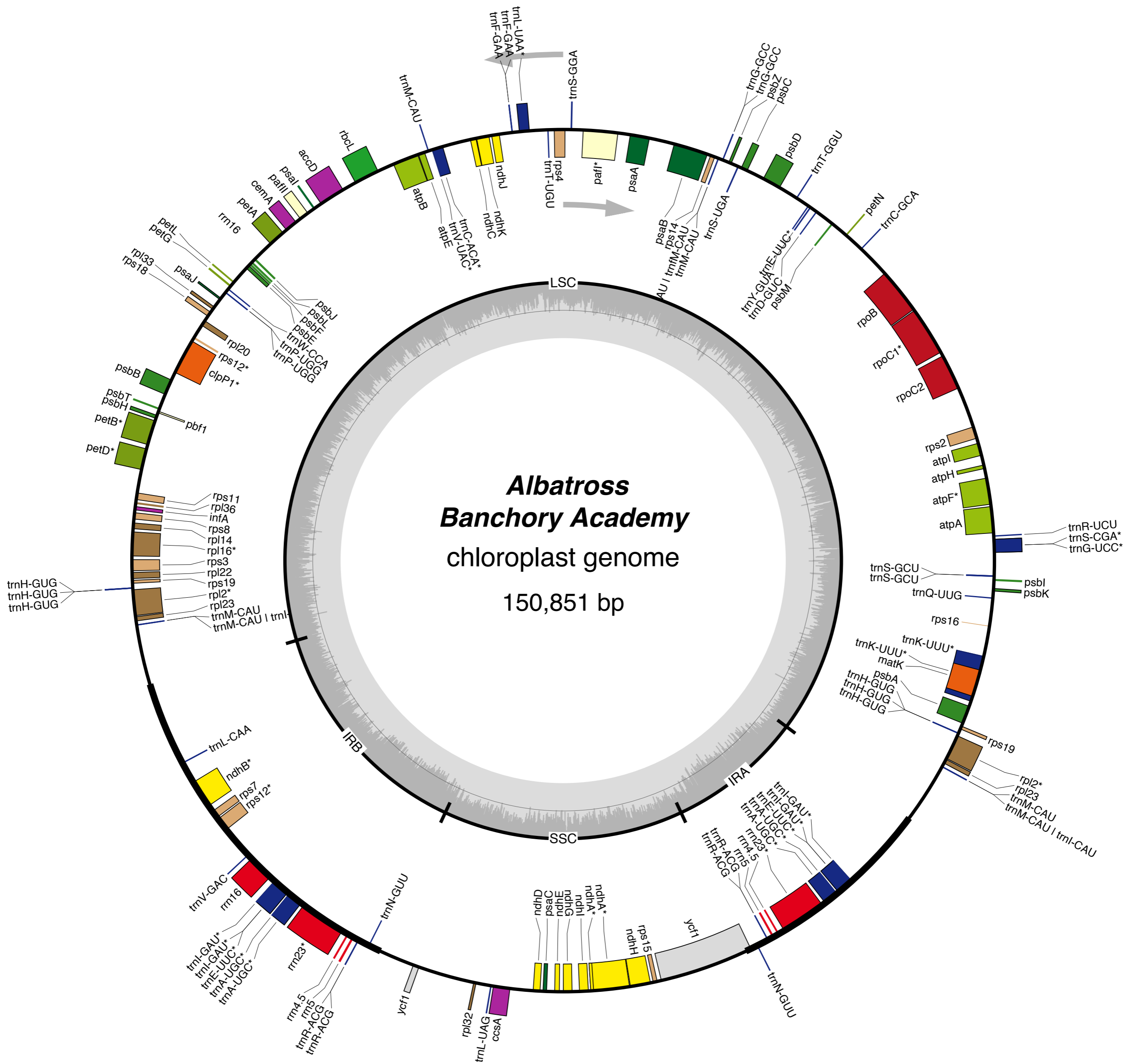


**Albatross  
Banchory Academy  
chloroplast genome**  
150,851 bp



- |   |  |  |   |   |
|---|--|--|---|---|
| <span style="color: green;">■</span> photosystem I          | <span style="color: lightgreen;">■</span> ATP synthase     | <span style="color: yellow;">■</span> photosystem assembly/stability factors | <span style="color: brown;">■</span> ribosomal proteins (LSU) | <span style="color: purple;">■</span> other genes                                 |
| <span style="color: darkgreen;">■</span> photosystem II     | <span style="color: yellow;">■</span> NADH dehydrogenase   | <span style="color: red;">■</span> RNA polymerase                            | <span style="color: red;">■</span> ribosomal RNAs             | <span style="color: grey;">■</span> hypothetical chloroplast reading frames (ycf) |
| <span style="color: olive;">■</span> cytochrome b/f complex | <span style="color: green;">■</span> RubisCO large subunit | <span style="color: tan;">■</span> ribosomal proteins (SSU)                  | <span style="color: orange;">■</span> clpP, matK              |   |