The diagram illustrates the distribution of taxa and character counts across different regions. Here are the details:

- **18S rDNA**: 567 taxa, 4592 characters
- **atpB**: 170 taxa, 2379 characters
- **rbcL**: 119 taxa, 3428 characters
- **matK**: 119 taxa, 3428 characters
- **26S rDNA**: 170 taxa, 3428 characters

The total number of taxa across all regions is 856, and the total number of characters is 11,340.