2021.12.30 DS\_FINAL condon usage

1. introduce all features

[1]Kindom: Dependent variable

[2]DNA type: dummy variables

[3]Species ID: It’s just a ID so we remove it.

[4]Ncodons: remove

[5]Speices name: It’s just name so we remove it.

[6] – [69]: All are numerical features and named by short form gene code. We consider it as independent variable.