

Figure 1 is a scatter plot showing the relationship between the number of amino acids (x-axis) and the number of residues (y-axis) for various proteins. The x-axis ranges from 0 to 200, and the y-axis ranges from 0 to 200. The plot includes data points for IVEGOR (green circles) and LUPARG (pink circles). Lines connect the data points for each protein, showing a linear relationship. The data points are labeled with their respective amino acid counts: IVEGOR (1742, 1050, 2519, 1060, 2364) and LUPARG (1071, 1069, 1060, 2364).

LUPARG
1050*A

SENCRA
 1071*A
 IVEGIR 1069*A

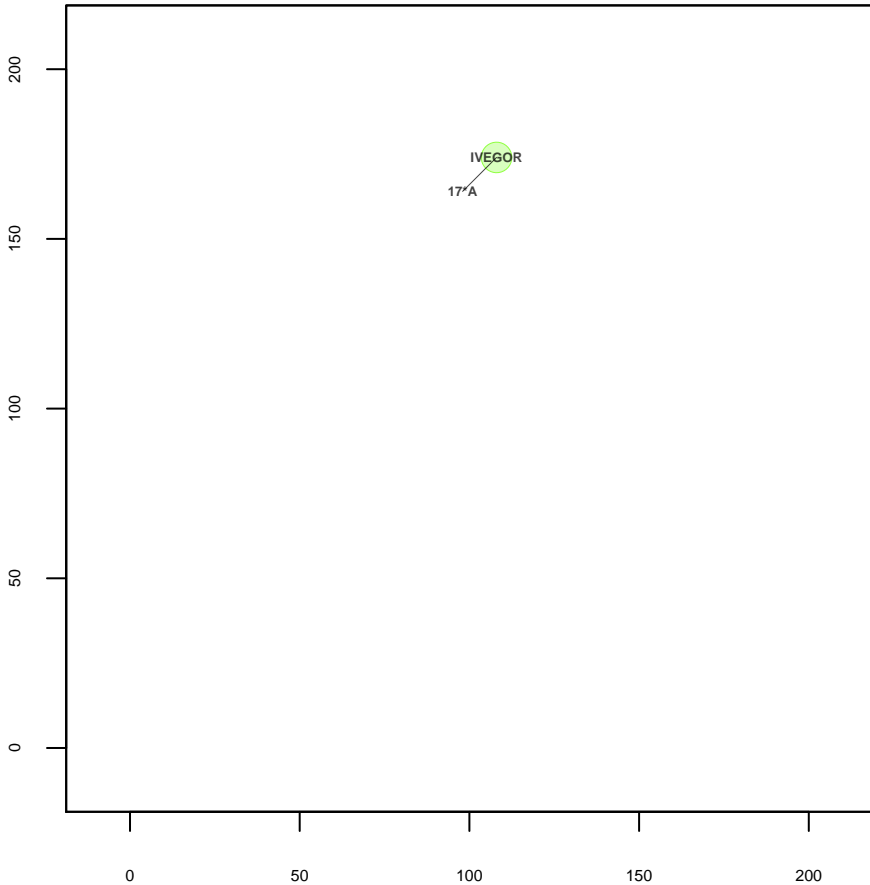
1060* A

2364*A

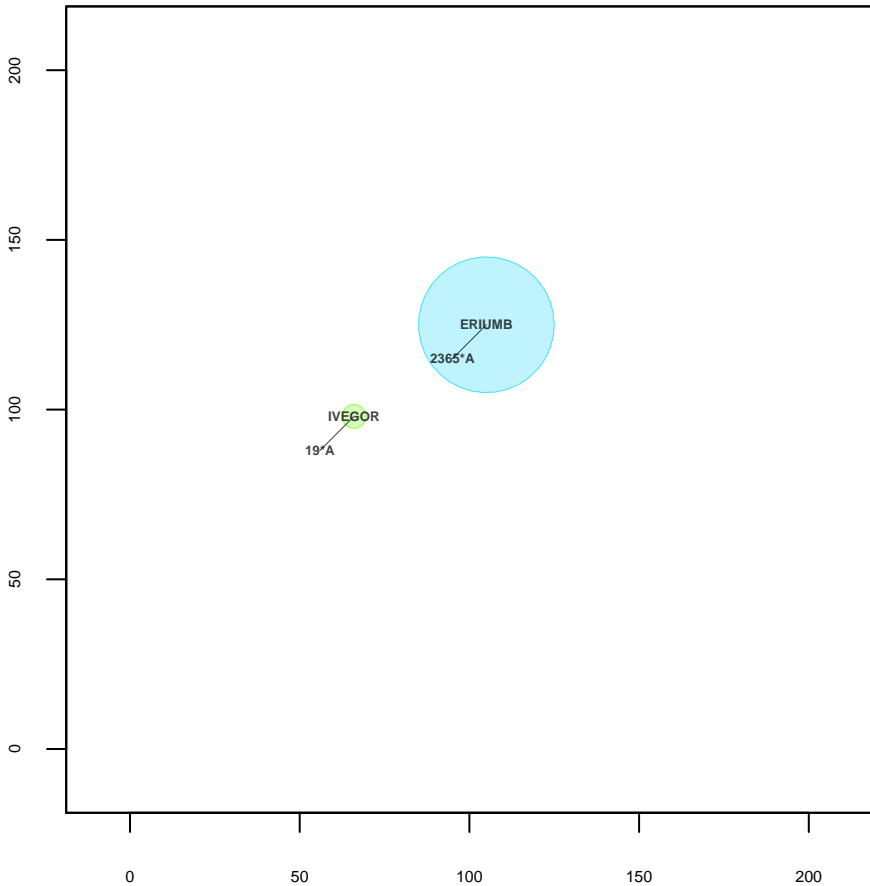
LUPARG

IVEGGR^{*}A

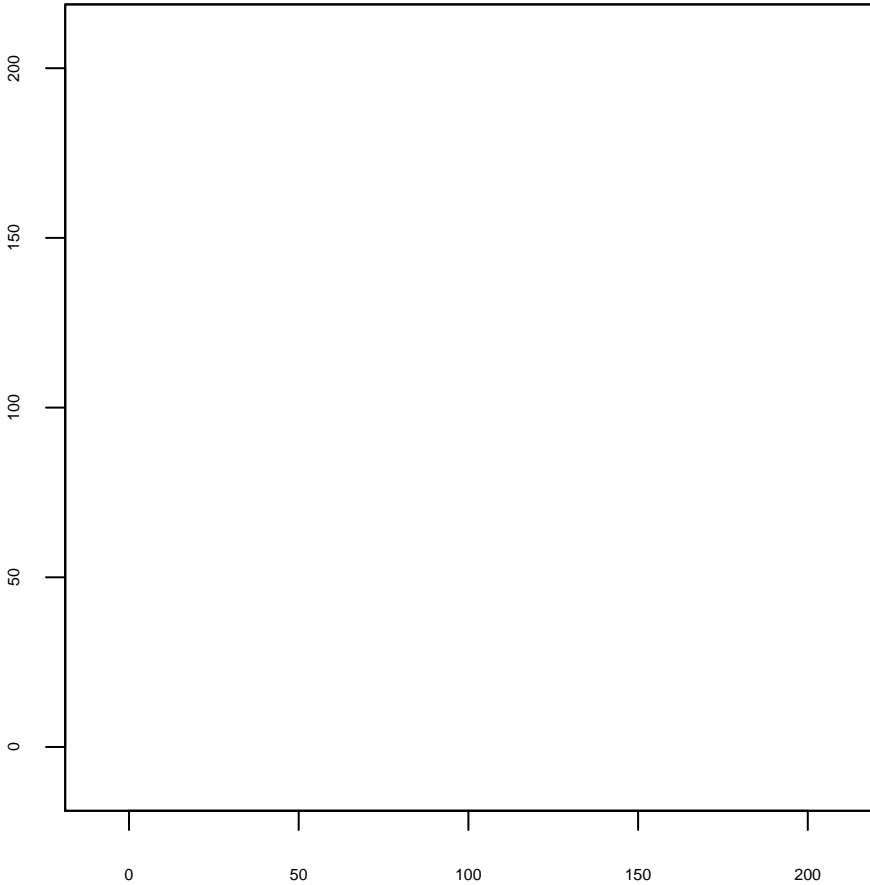
Plot 2



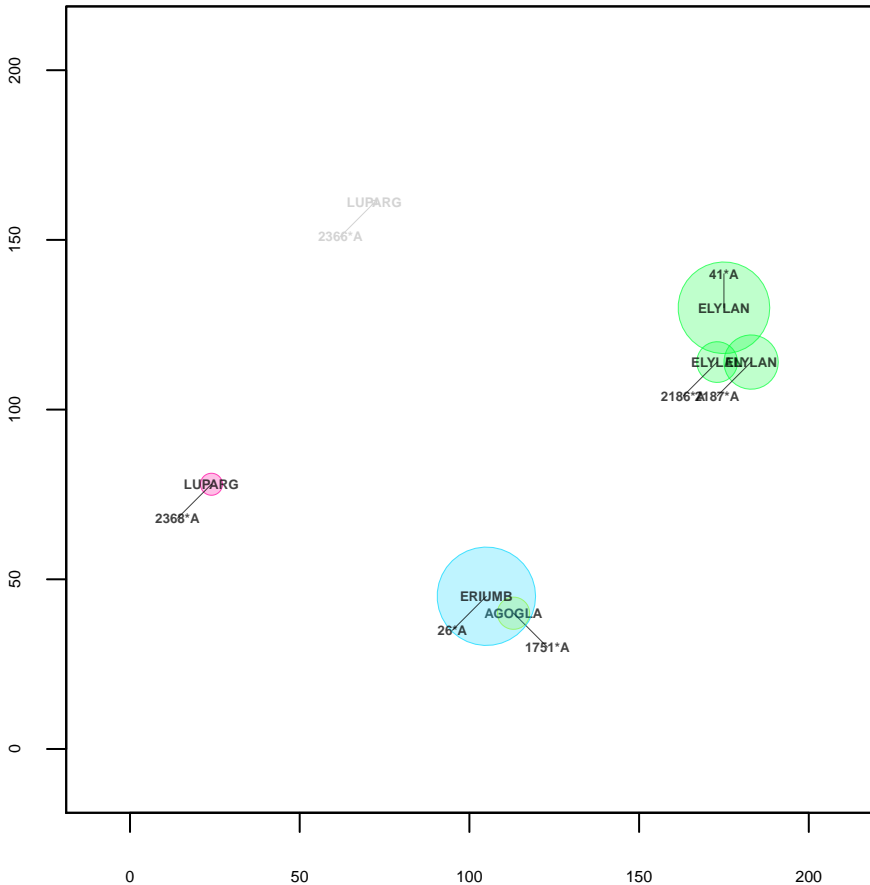
Plot 3



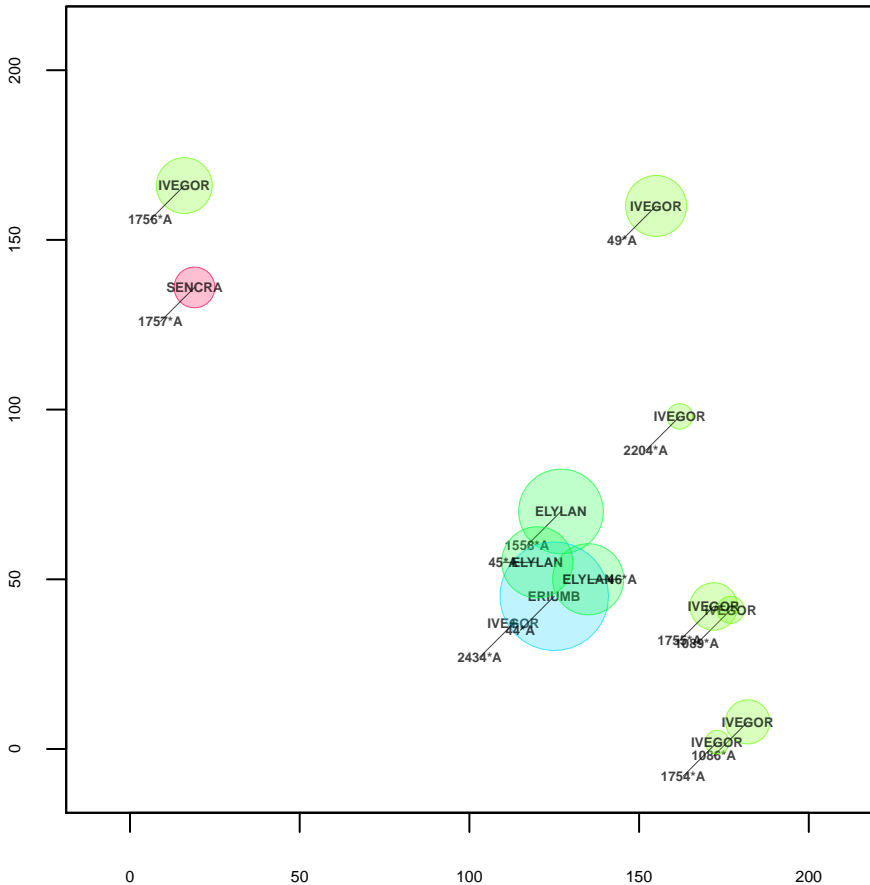
Plot 4



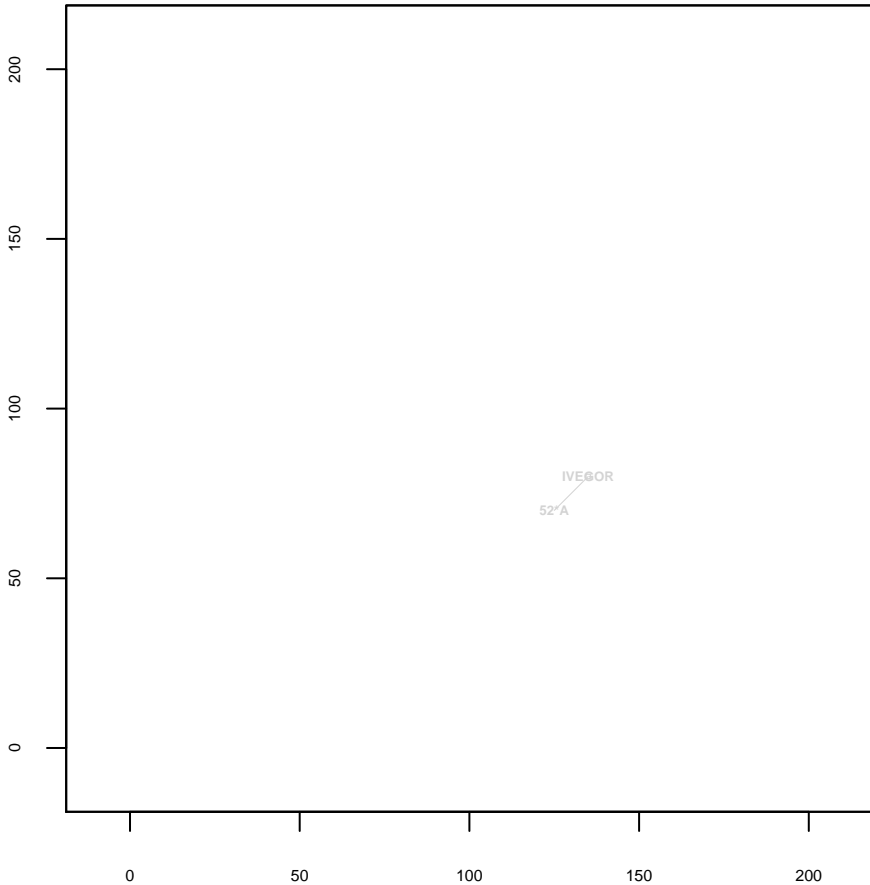
Plot 5



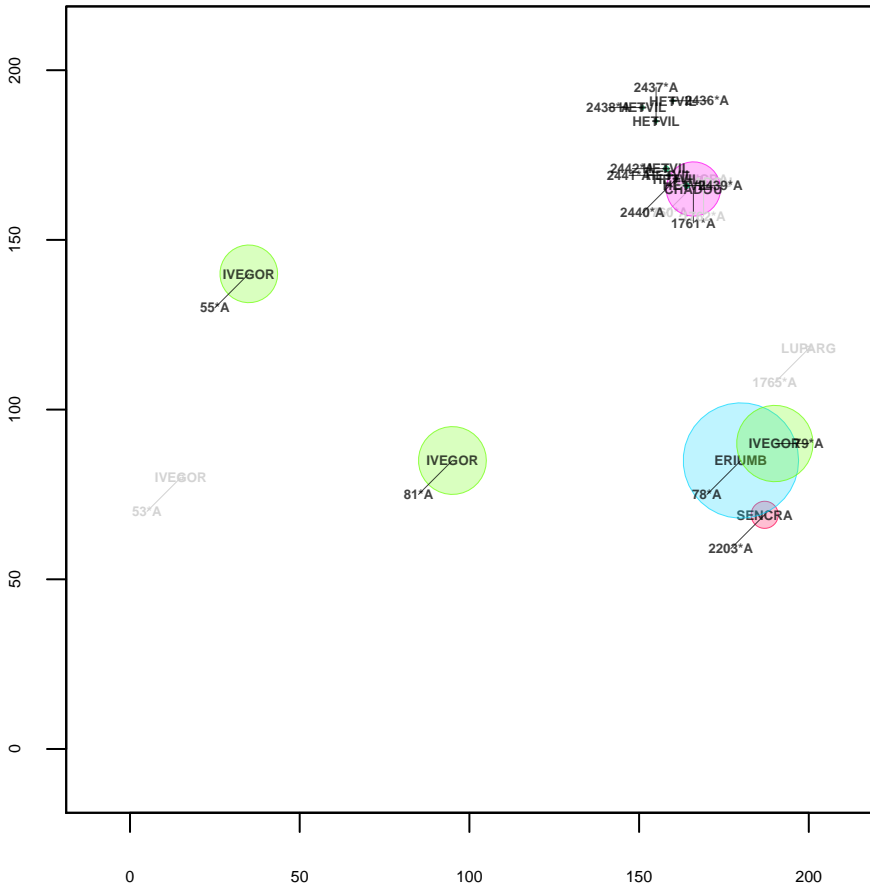
Plot 6



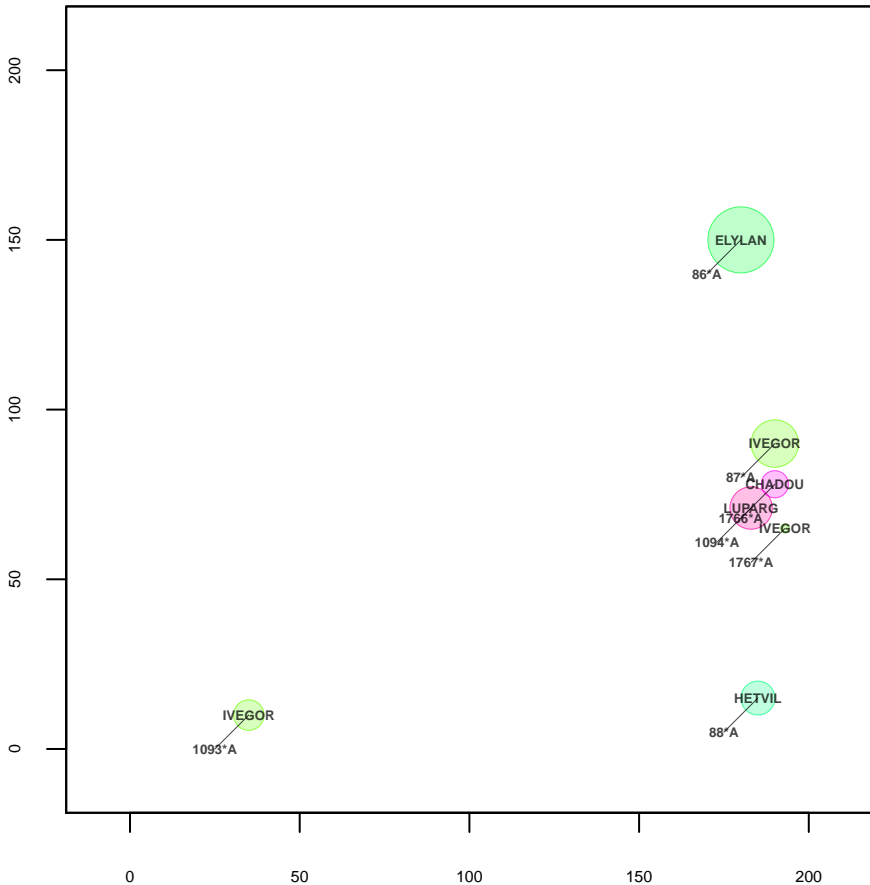
Plot 7



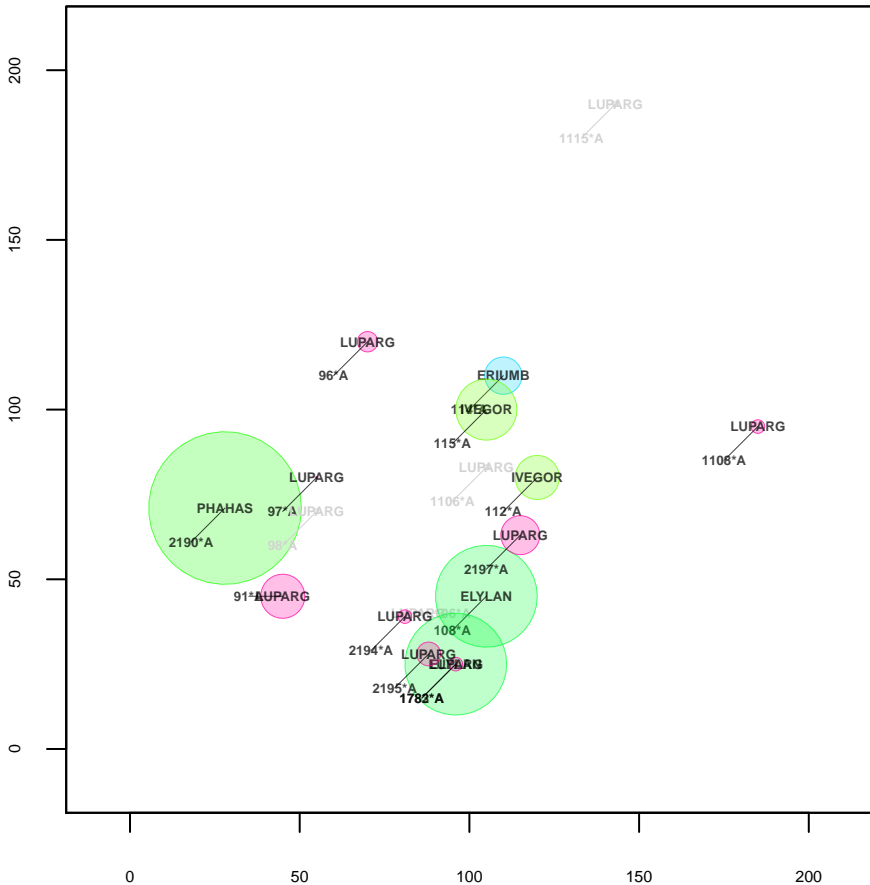
Plot 8



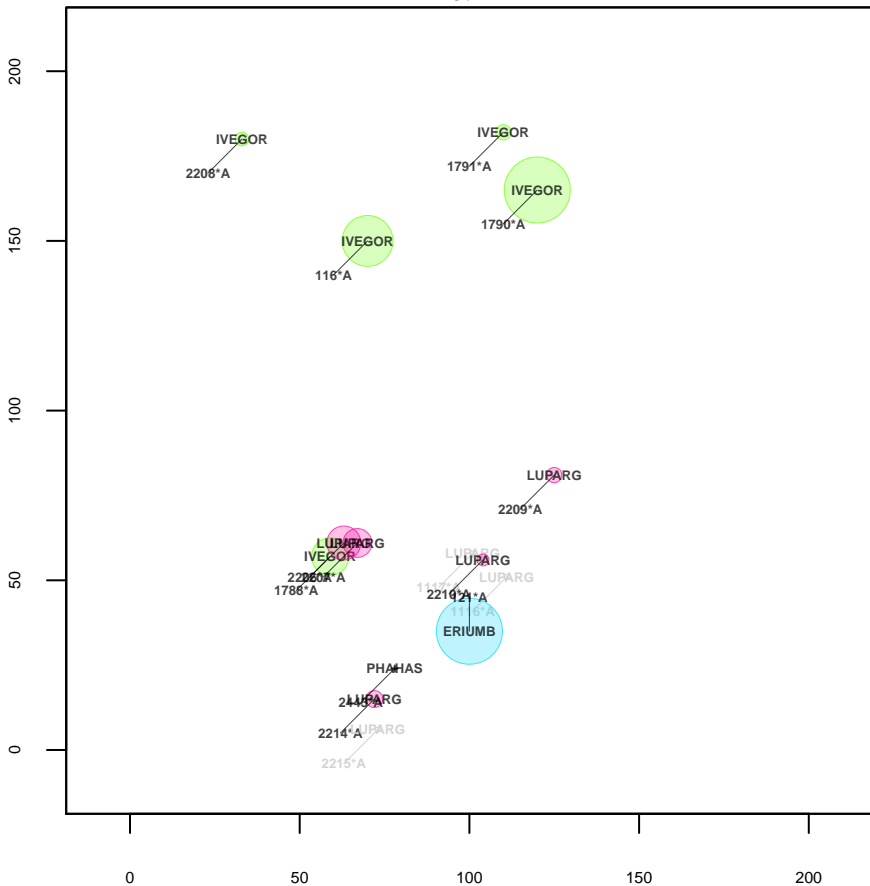
Plot 9



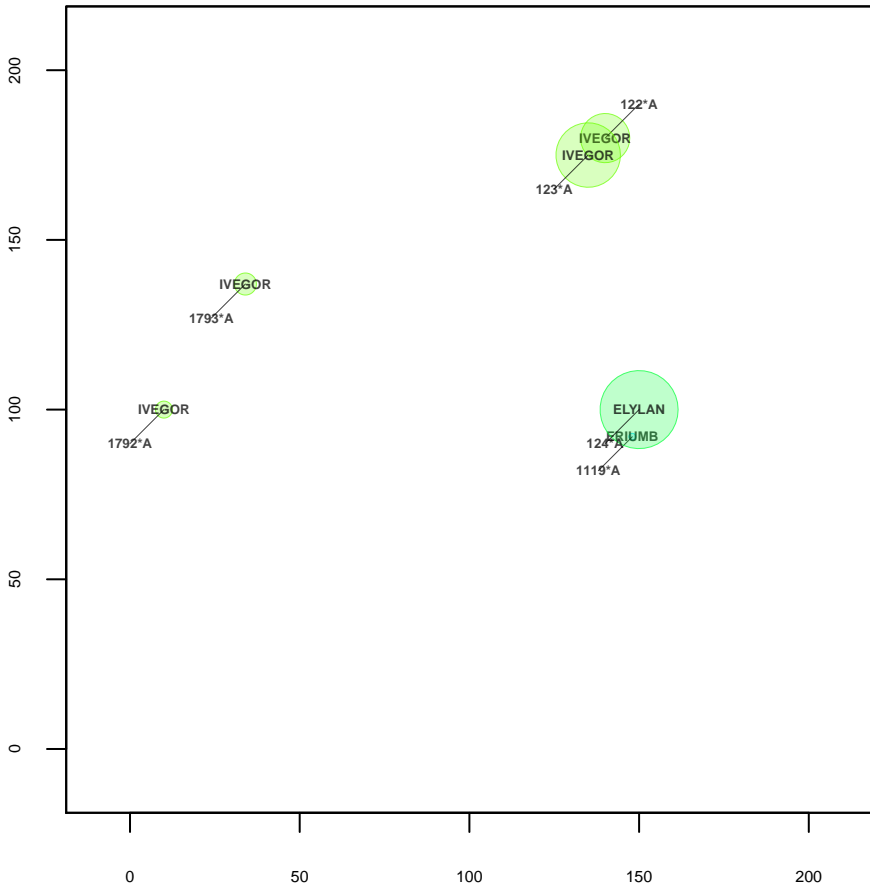
Plot 10



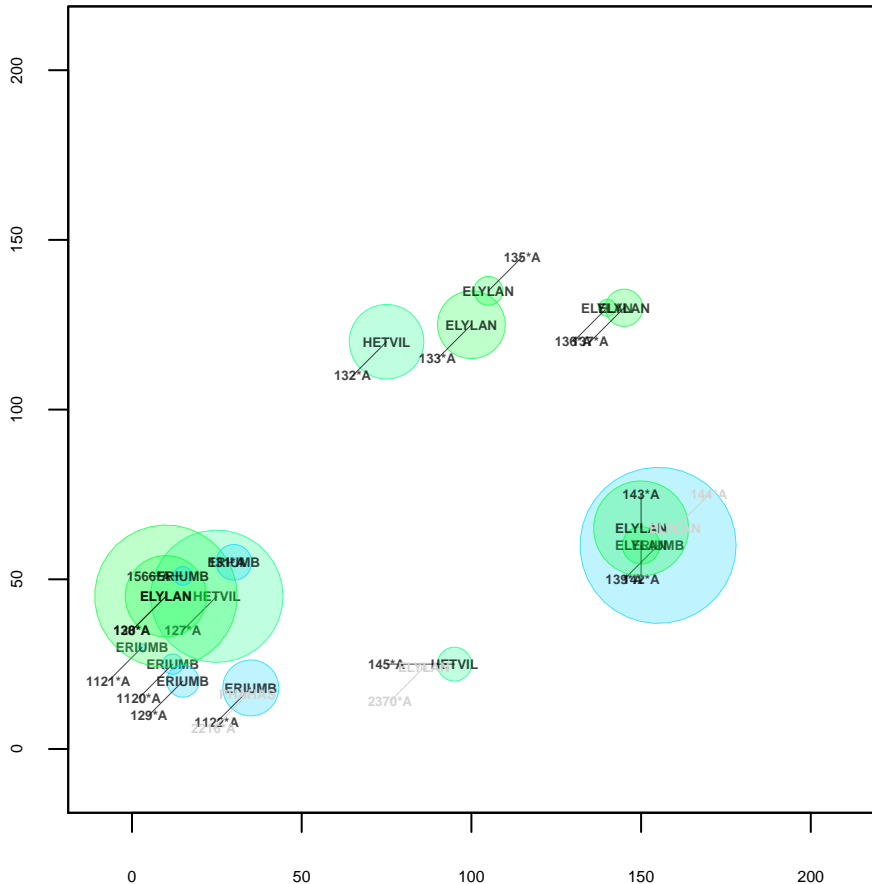
Plot 11



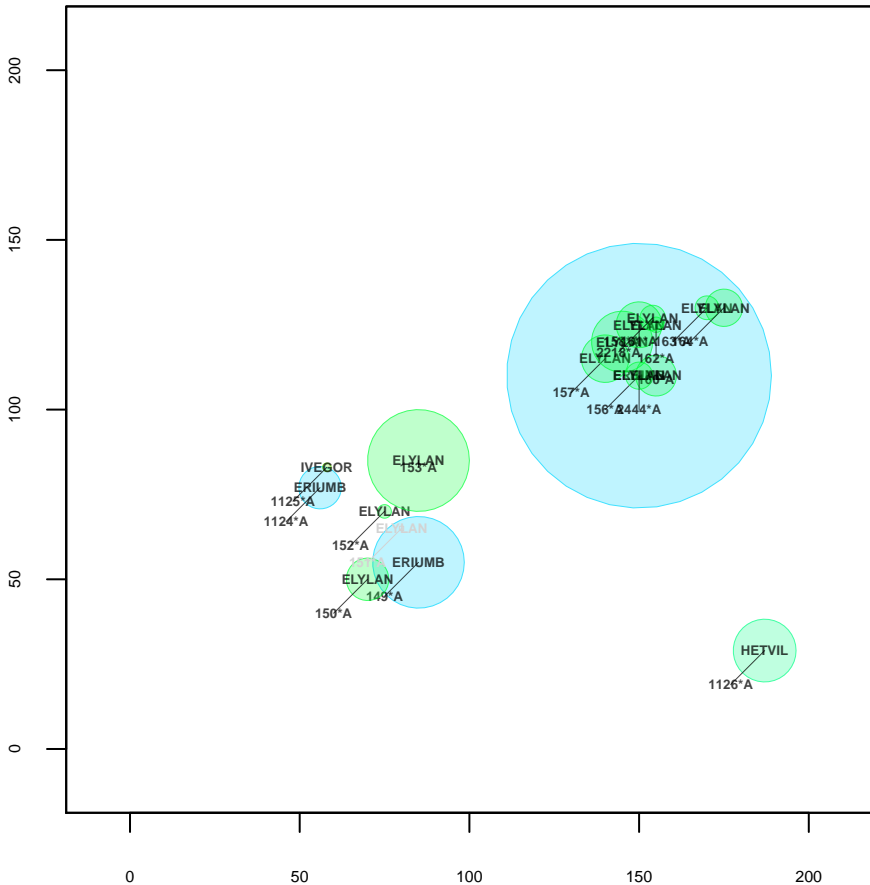
Plot 12



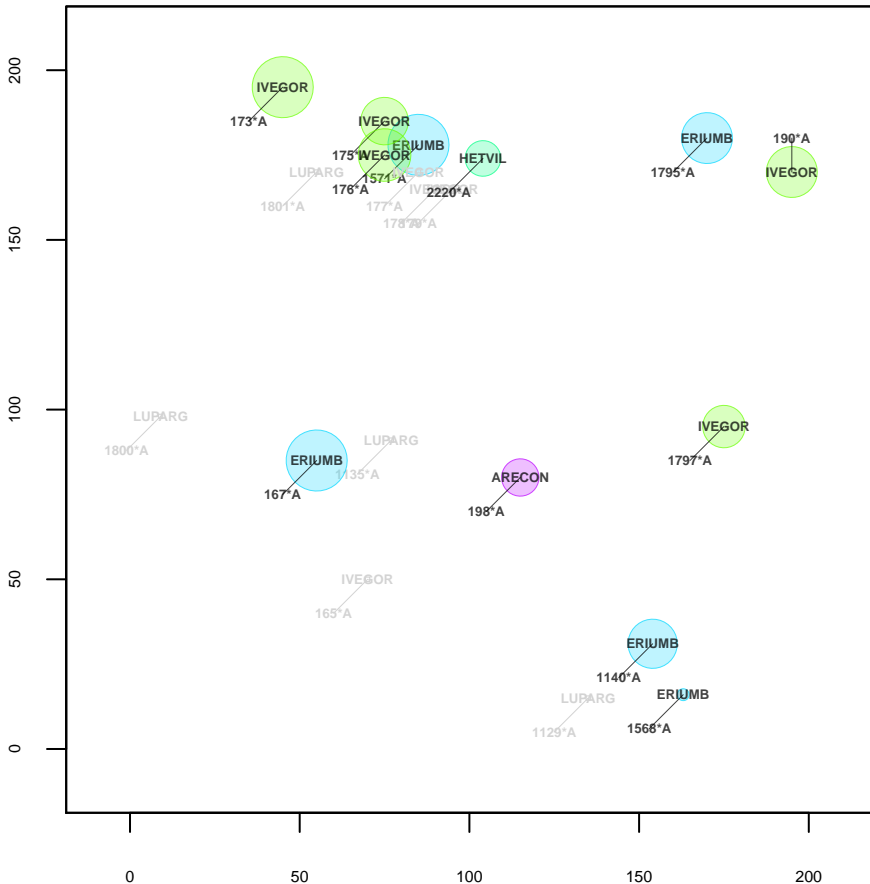
Plot 13



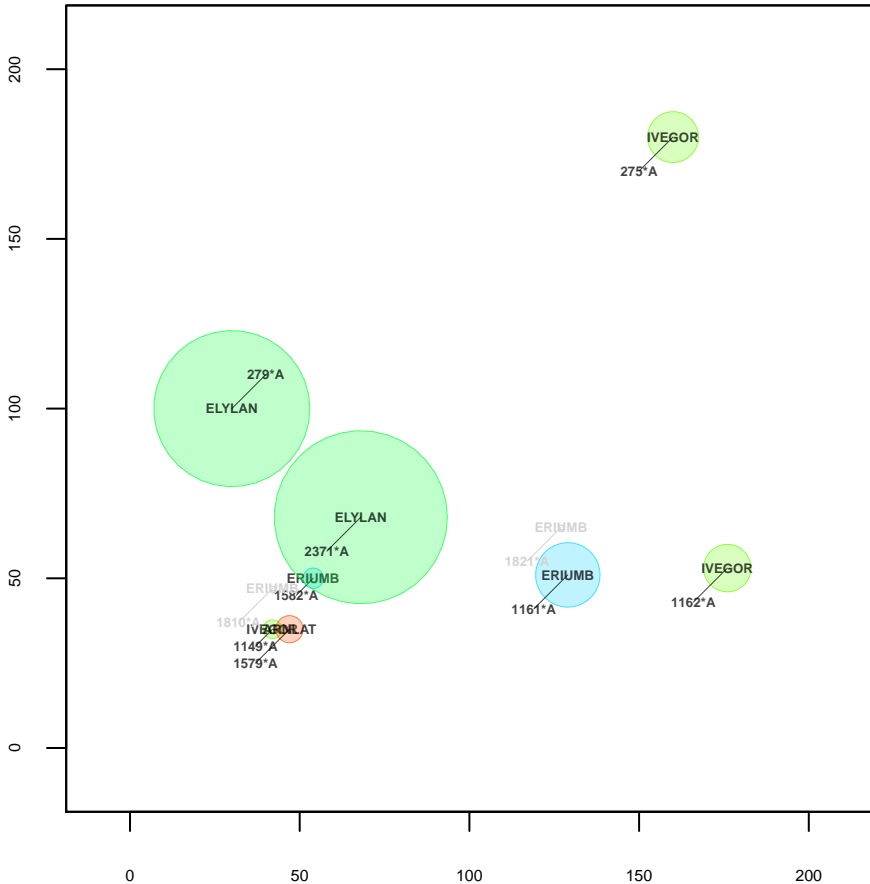
Plot 14



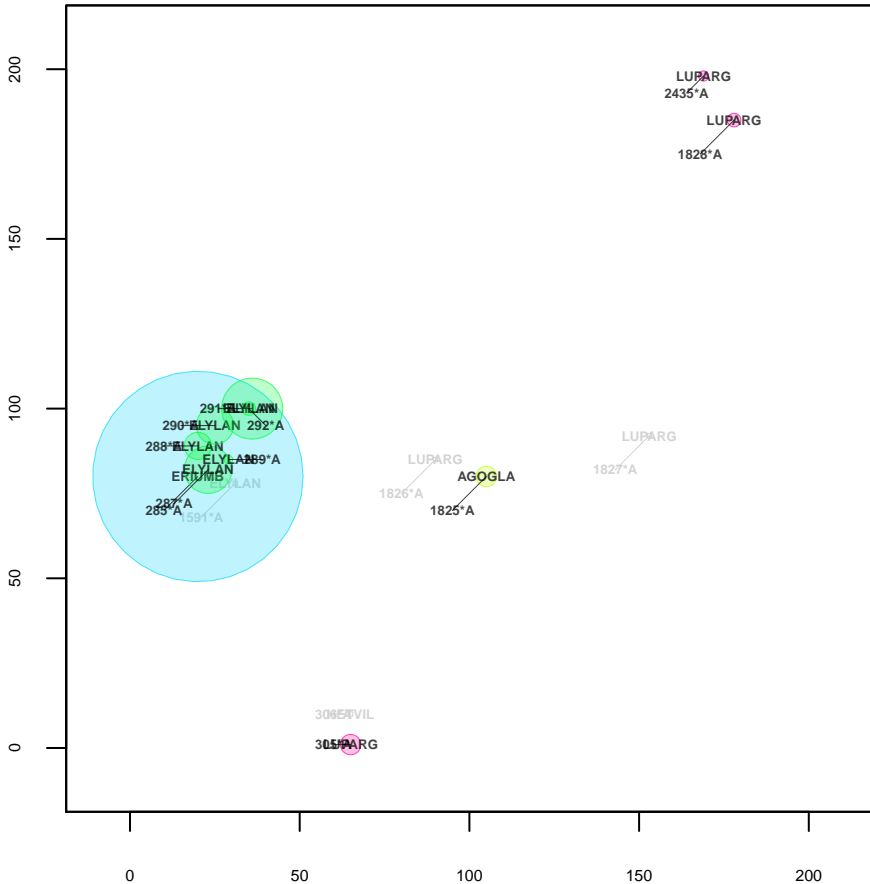
Plot 15



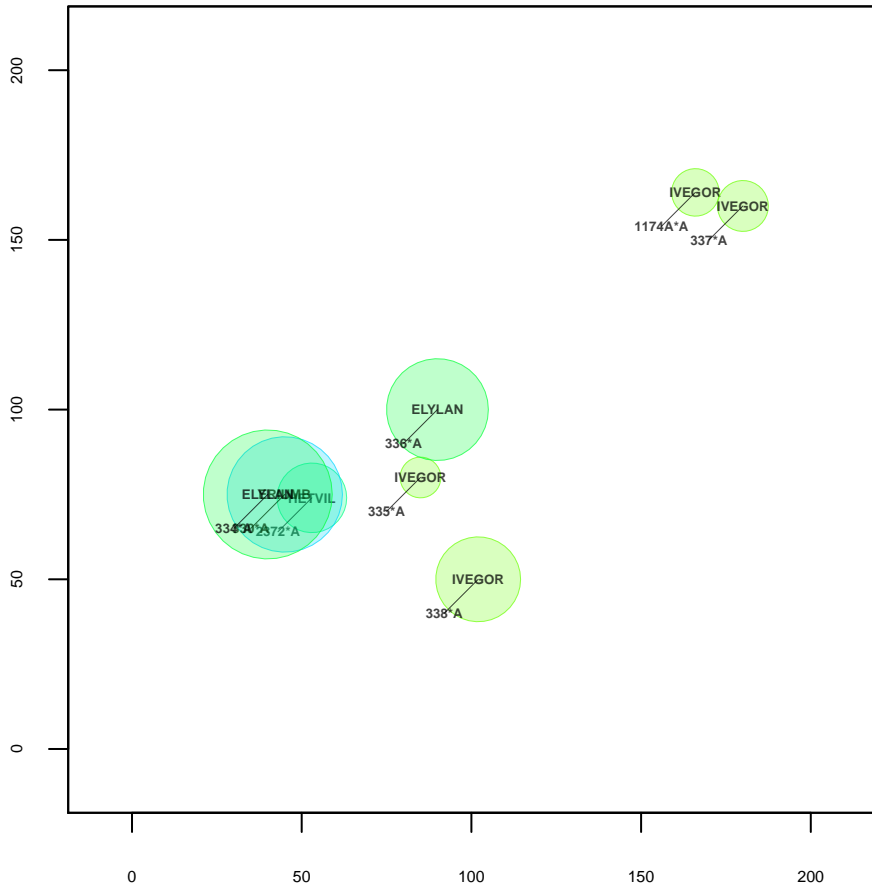
Plot 16



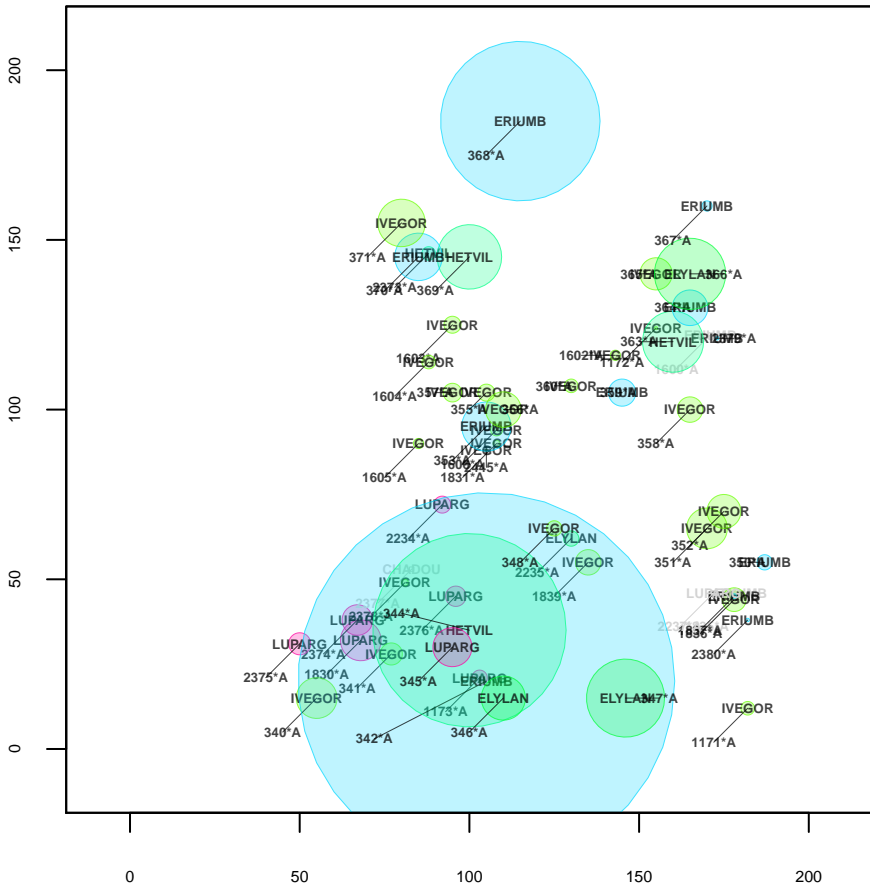
Plot 17



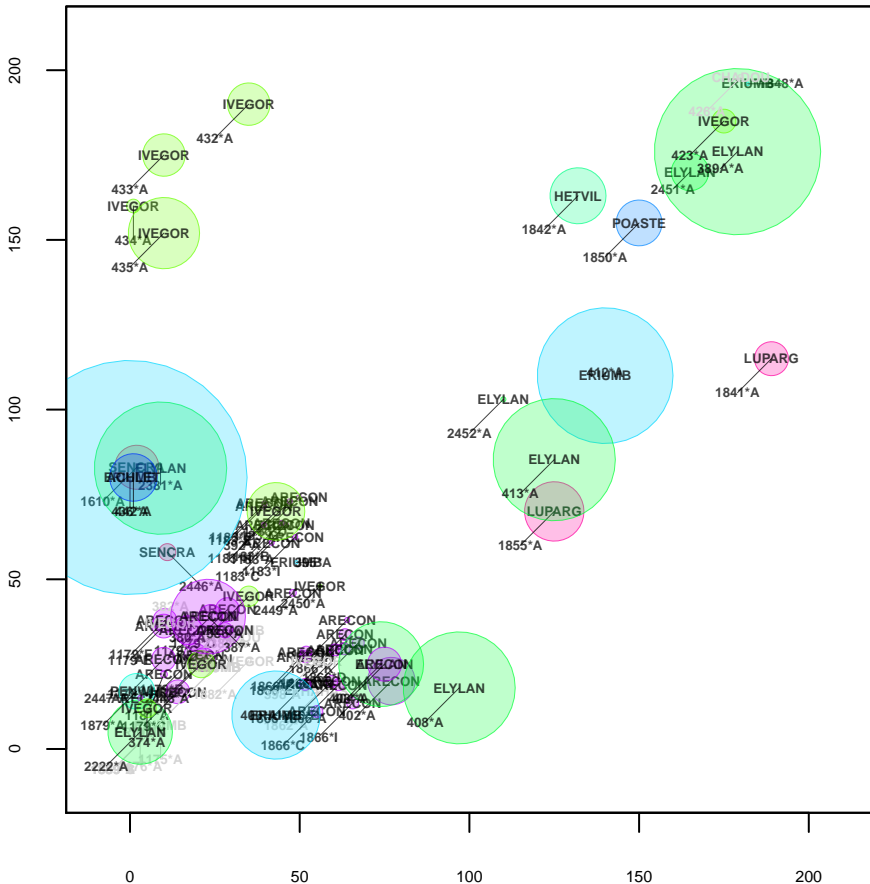
Plot 18



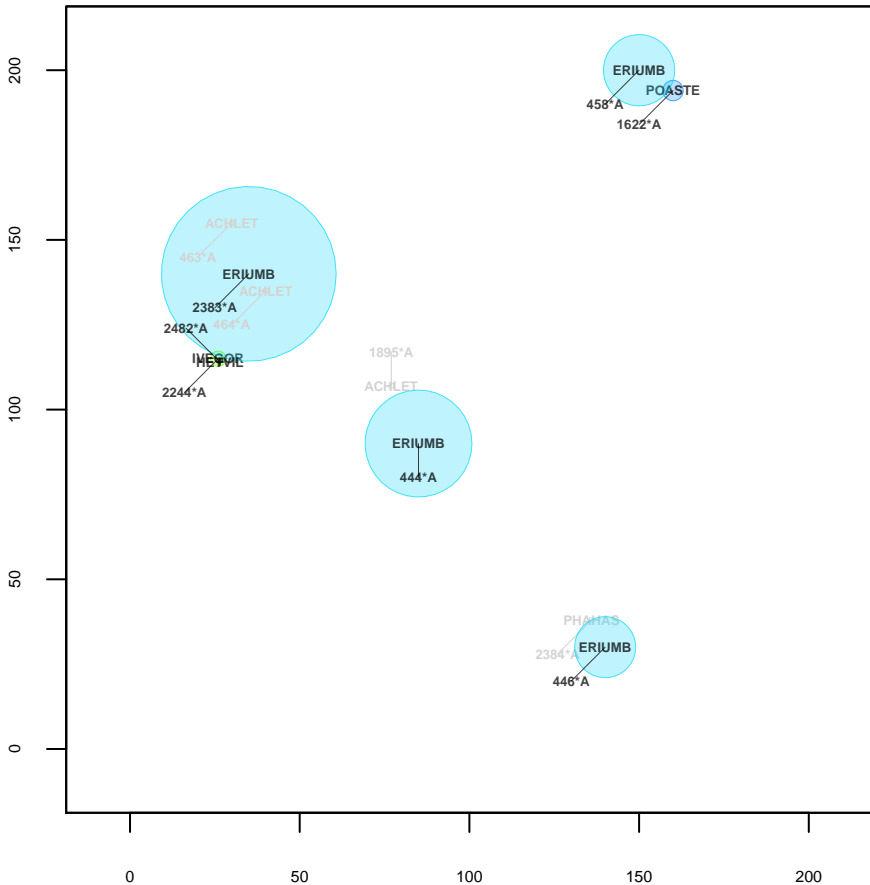
Plot 19



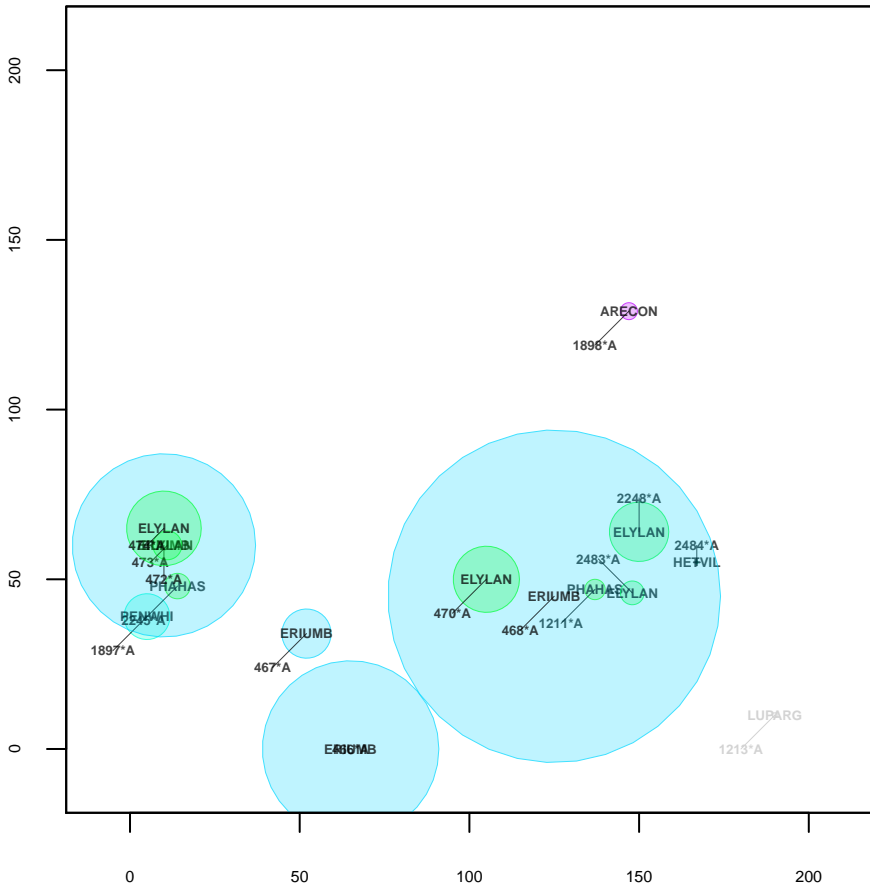
Plot 20



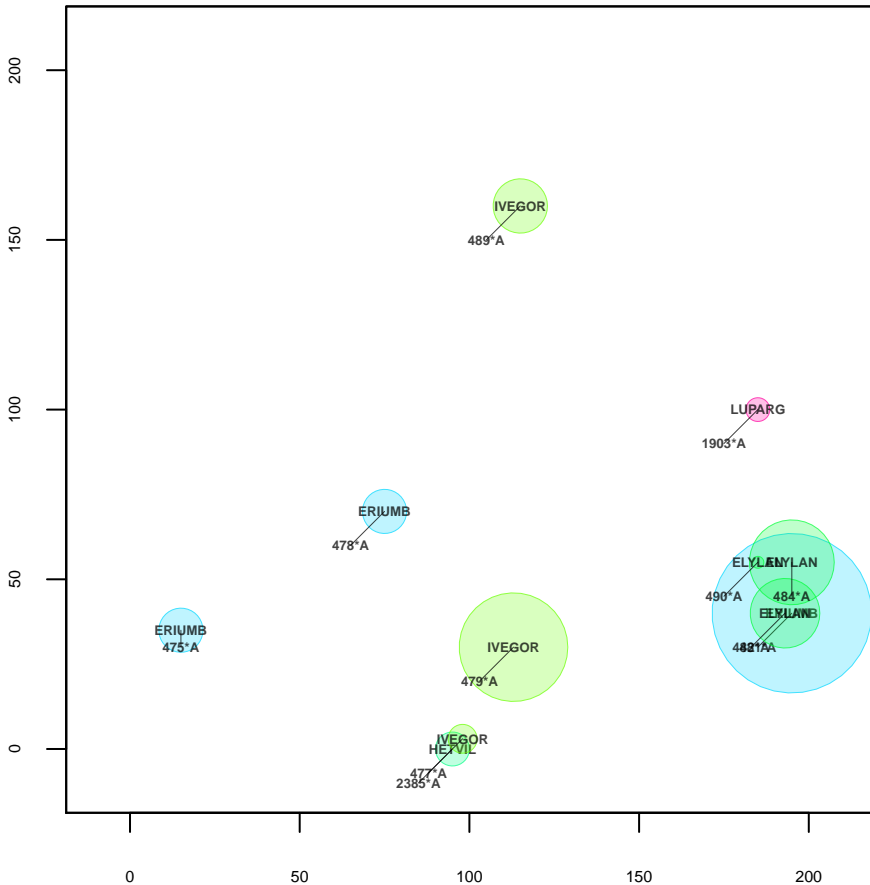
Plot 21



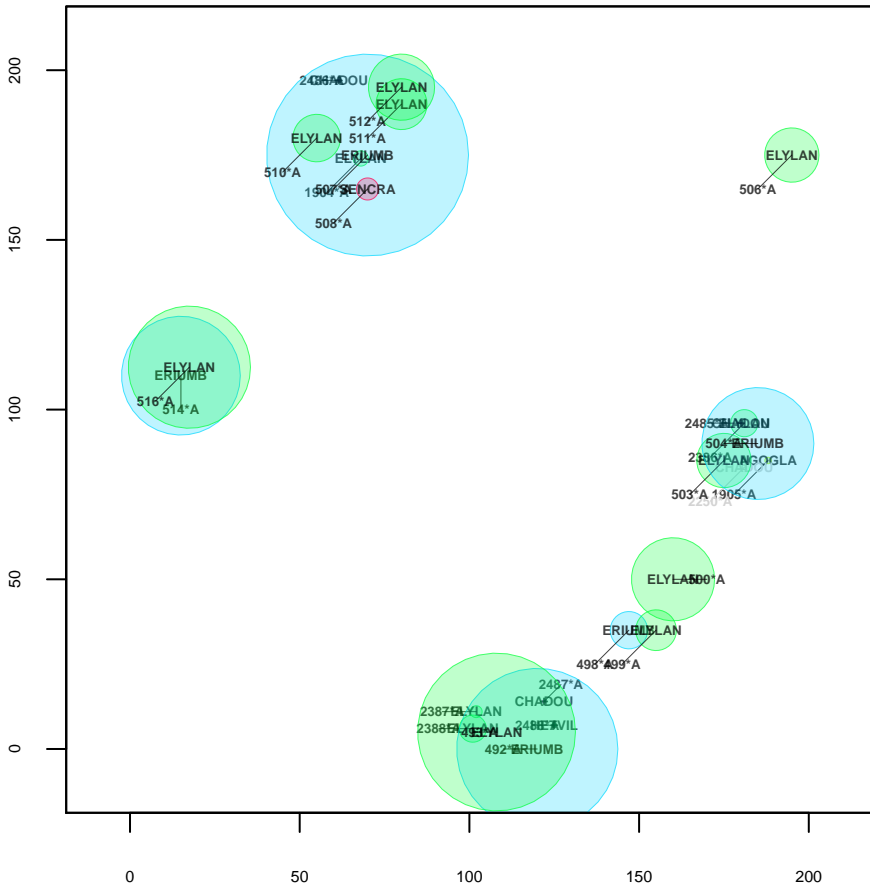
Plot 22



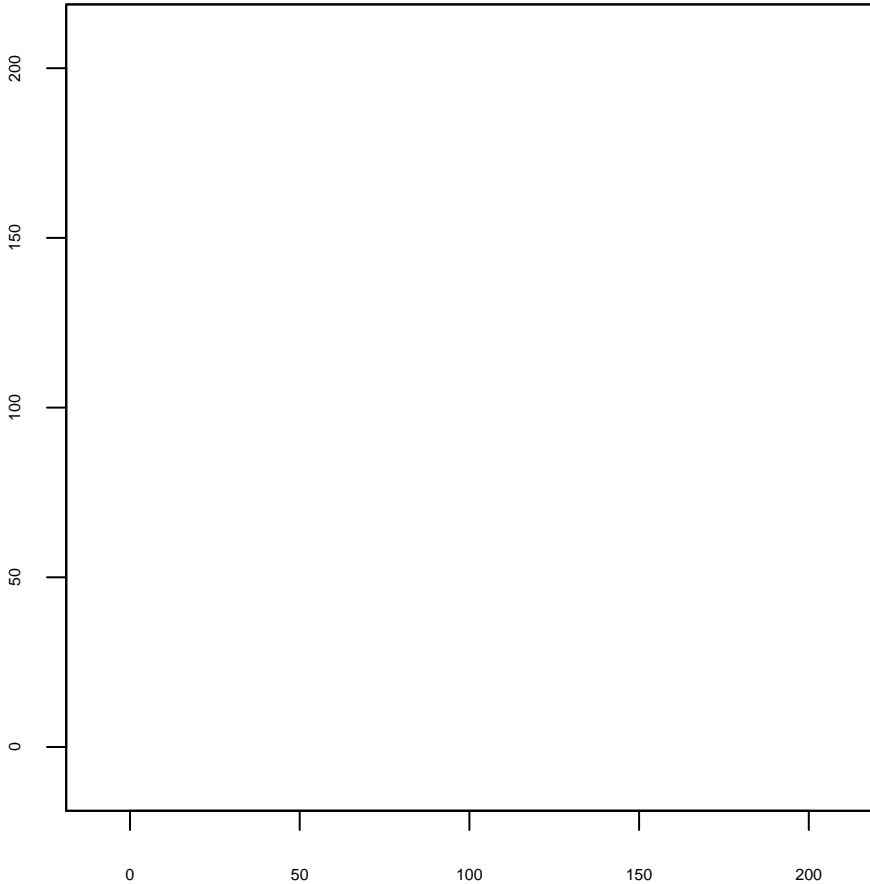
Plot 23



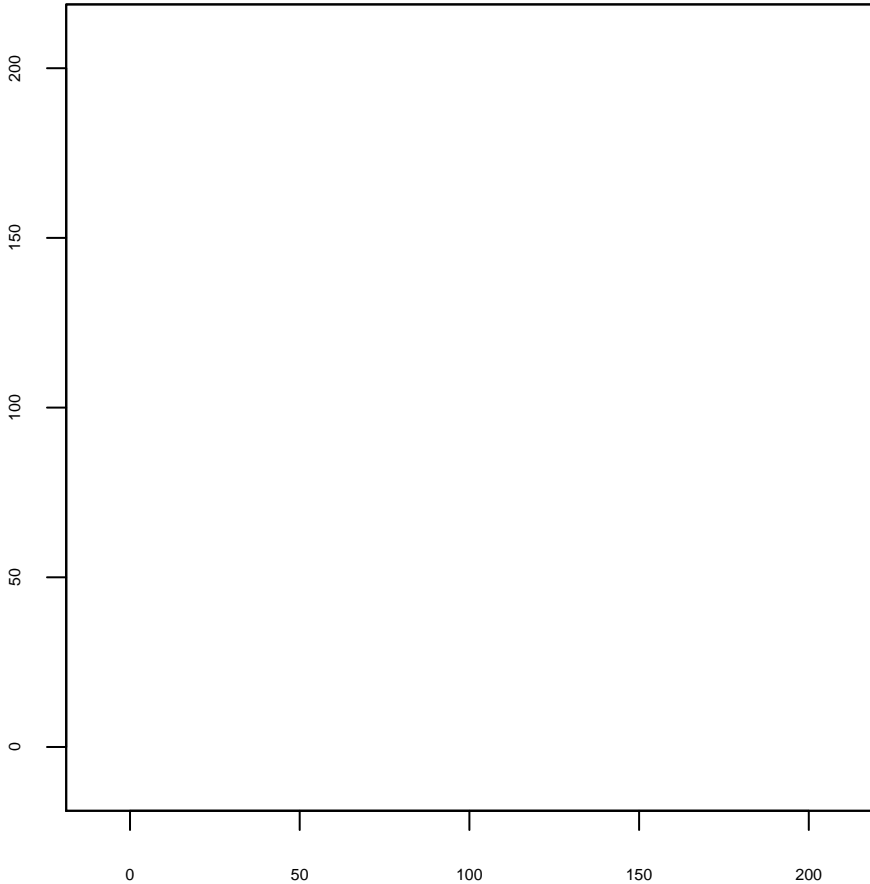
Plot 24



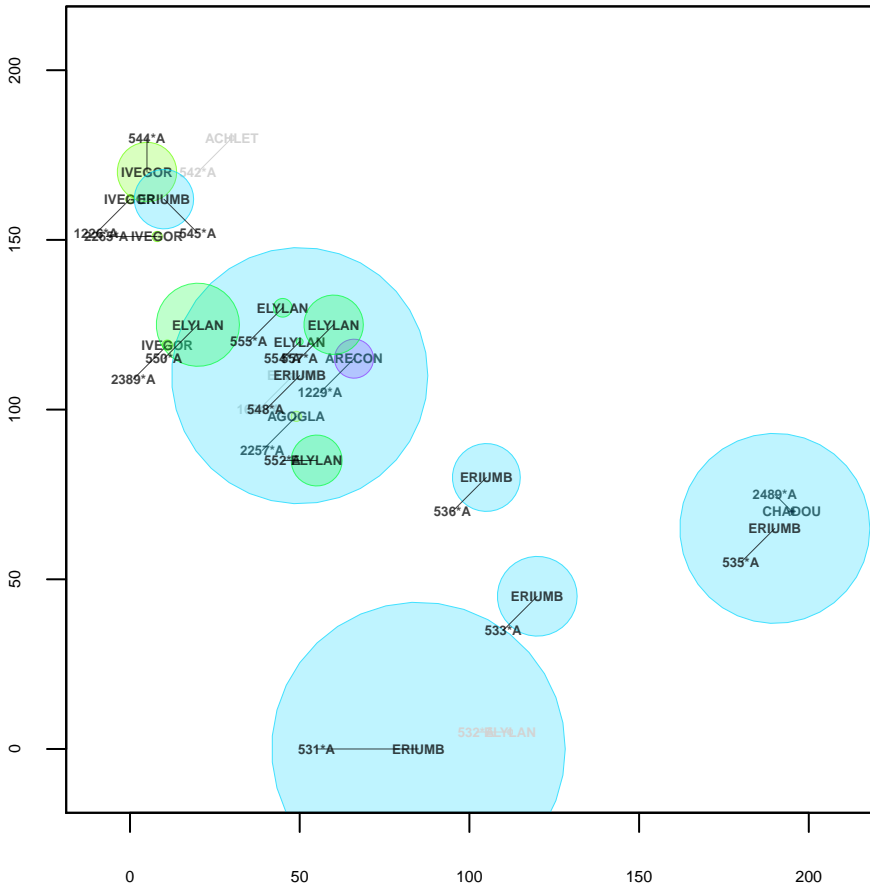
Plot 25



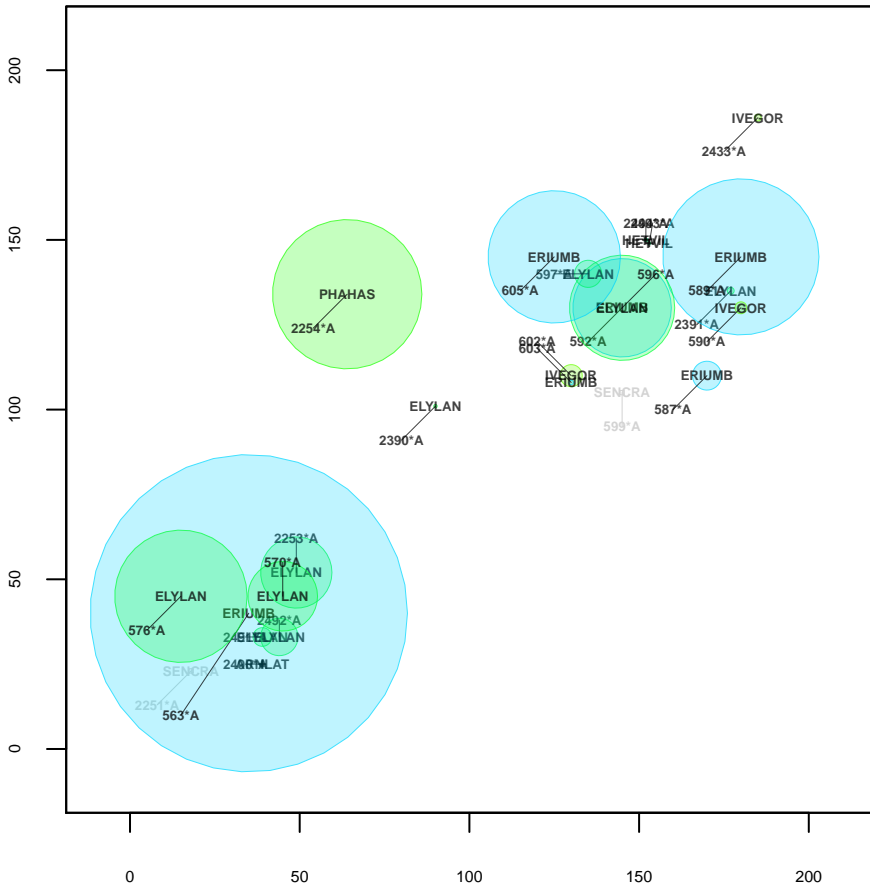
Plot 26



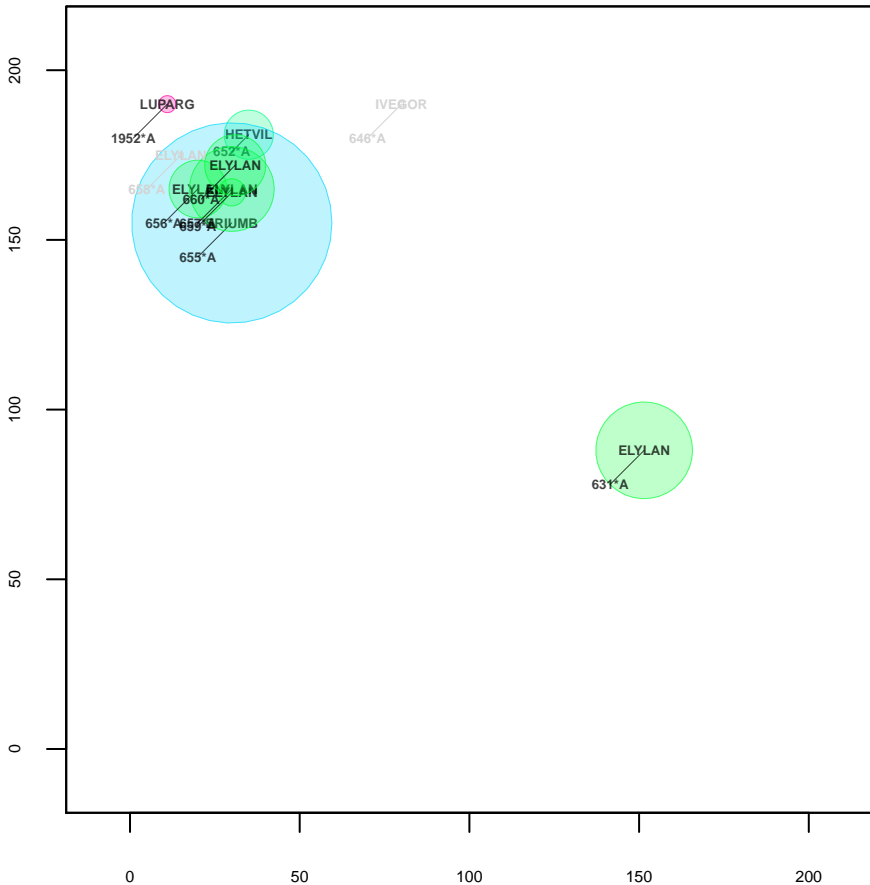
Plot 27



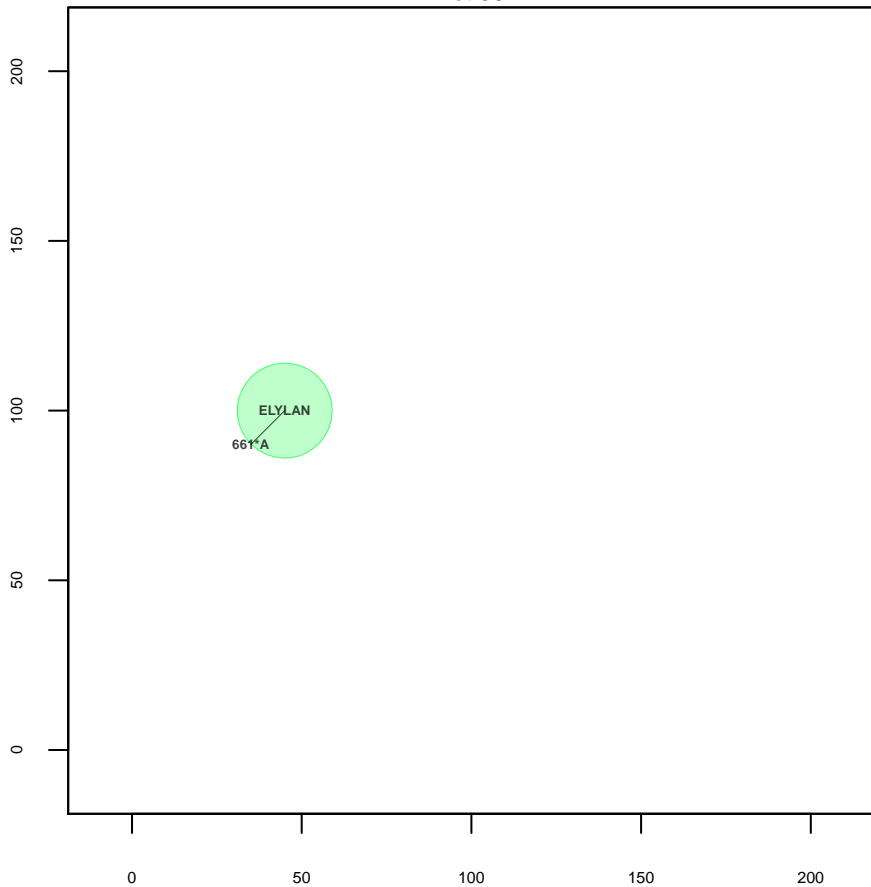
Plot 28



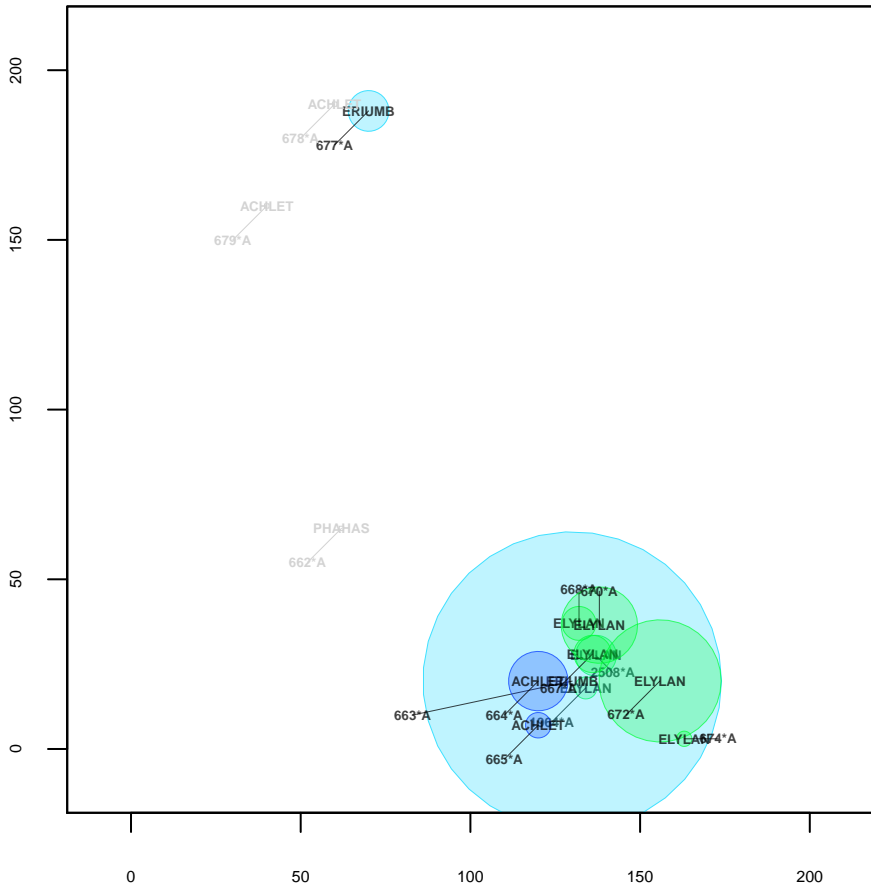
Plot 29



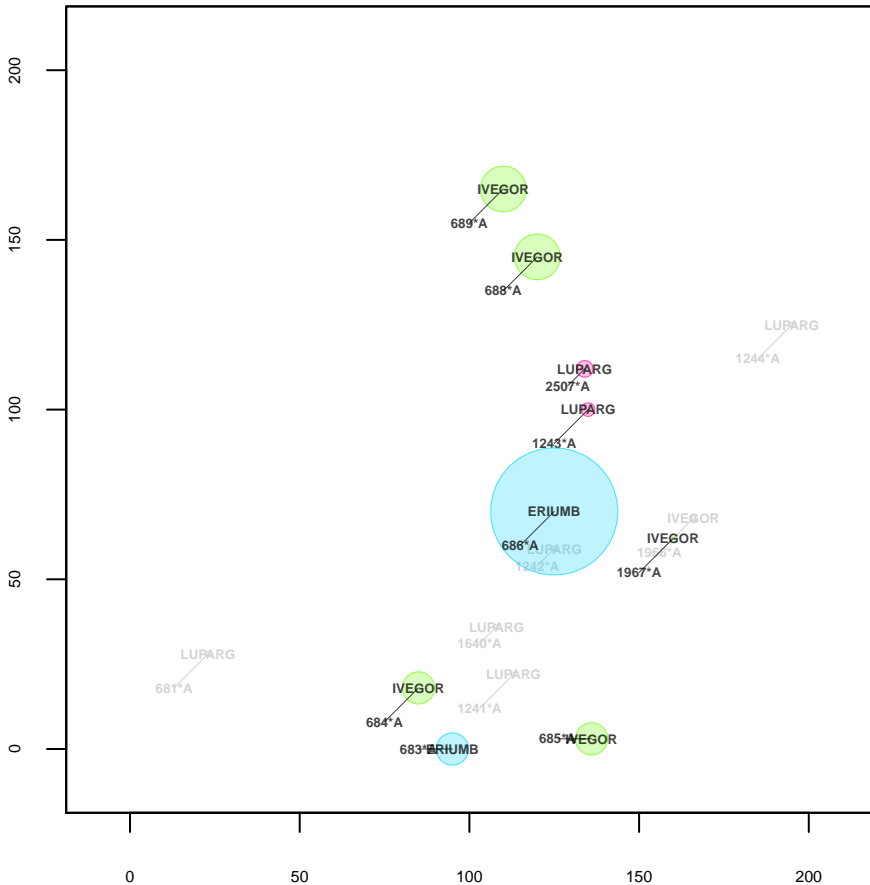
Plot 30



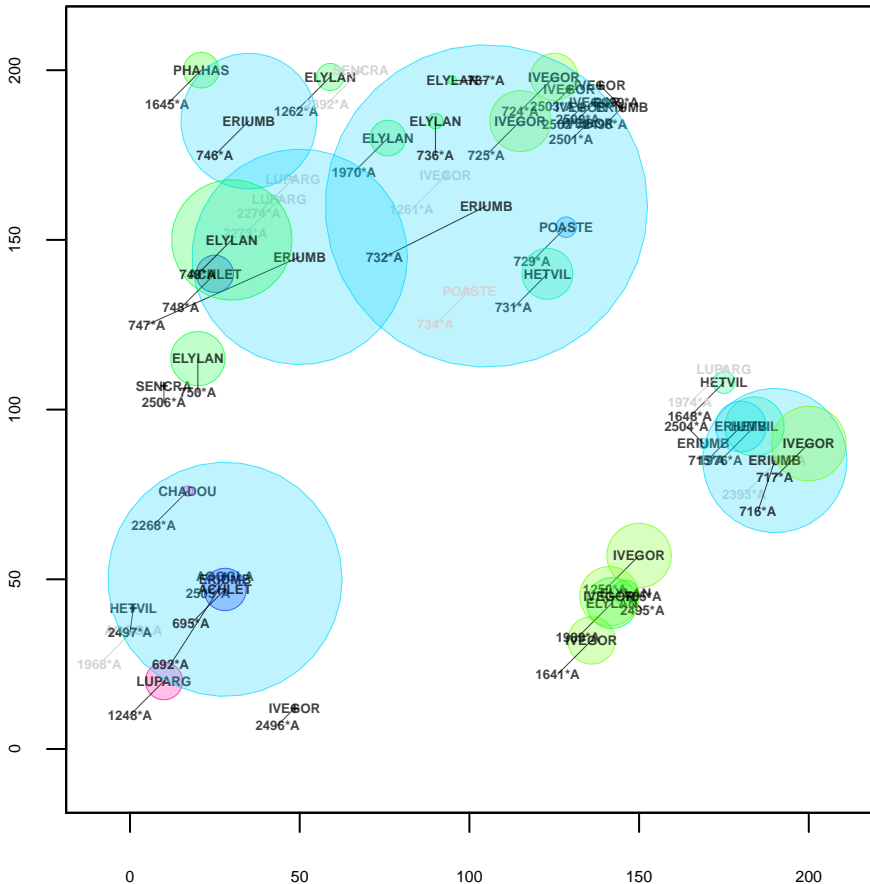
Plot 31



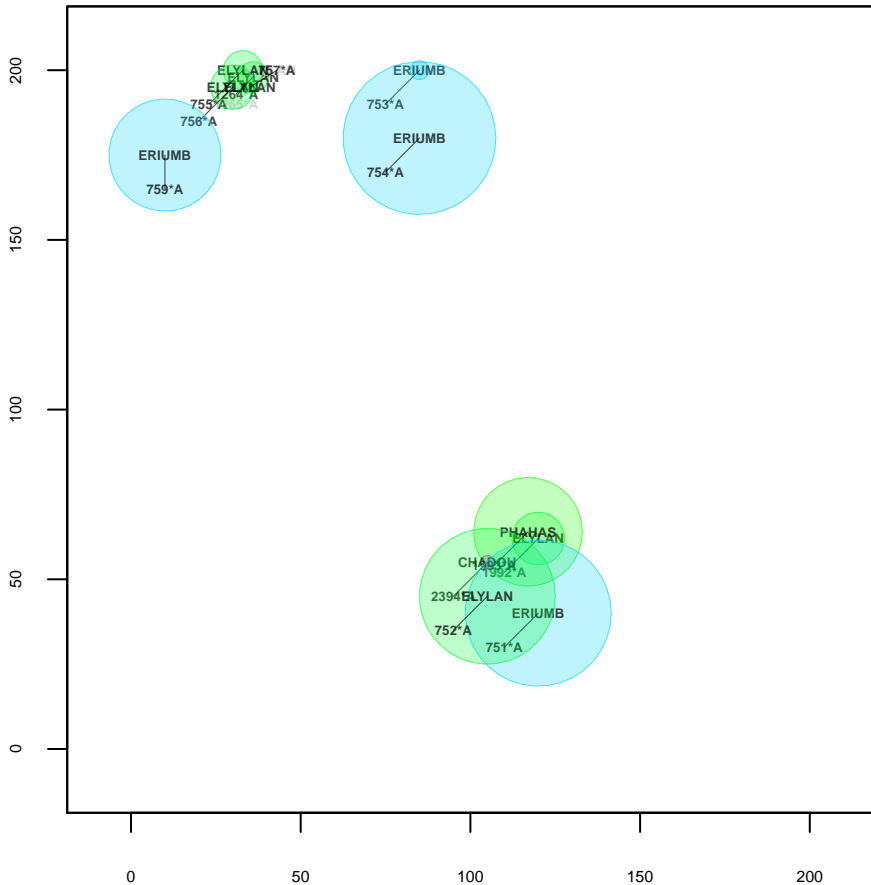
Plot 32



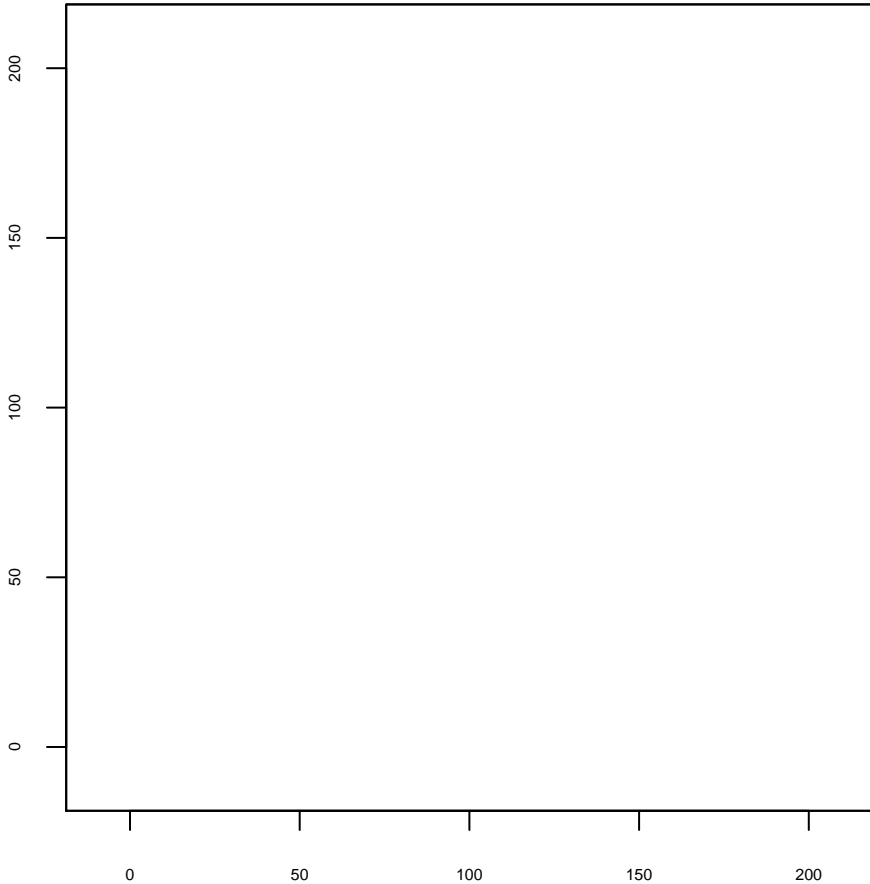
Plot 33



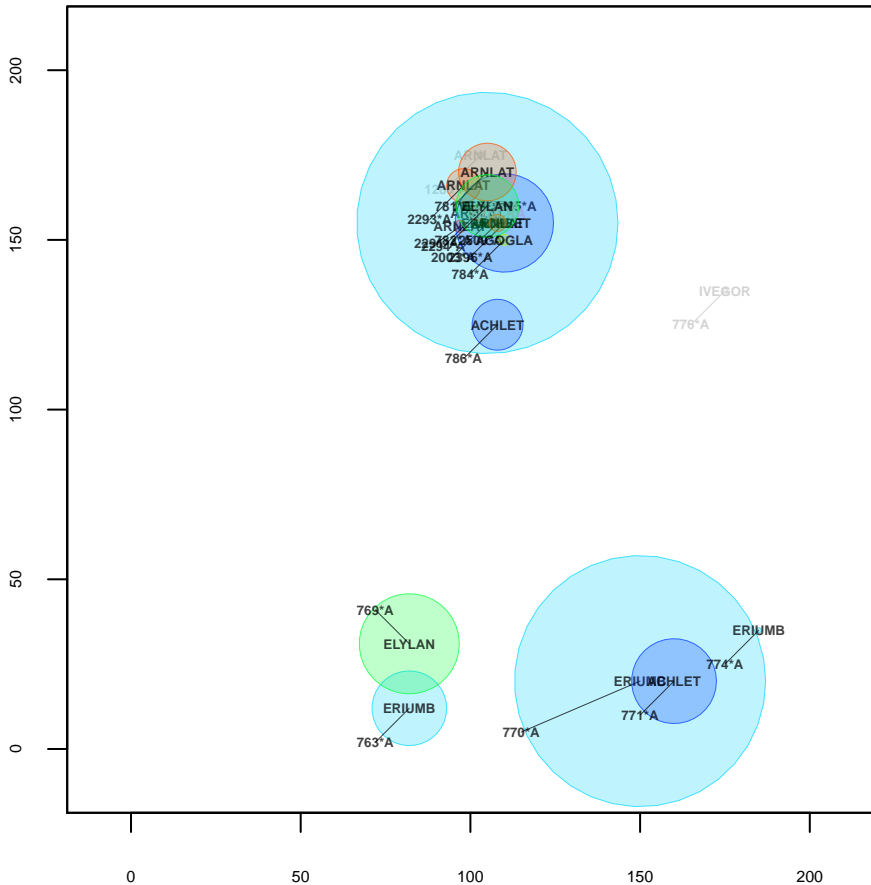
Plot 34



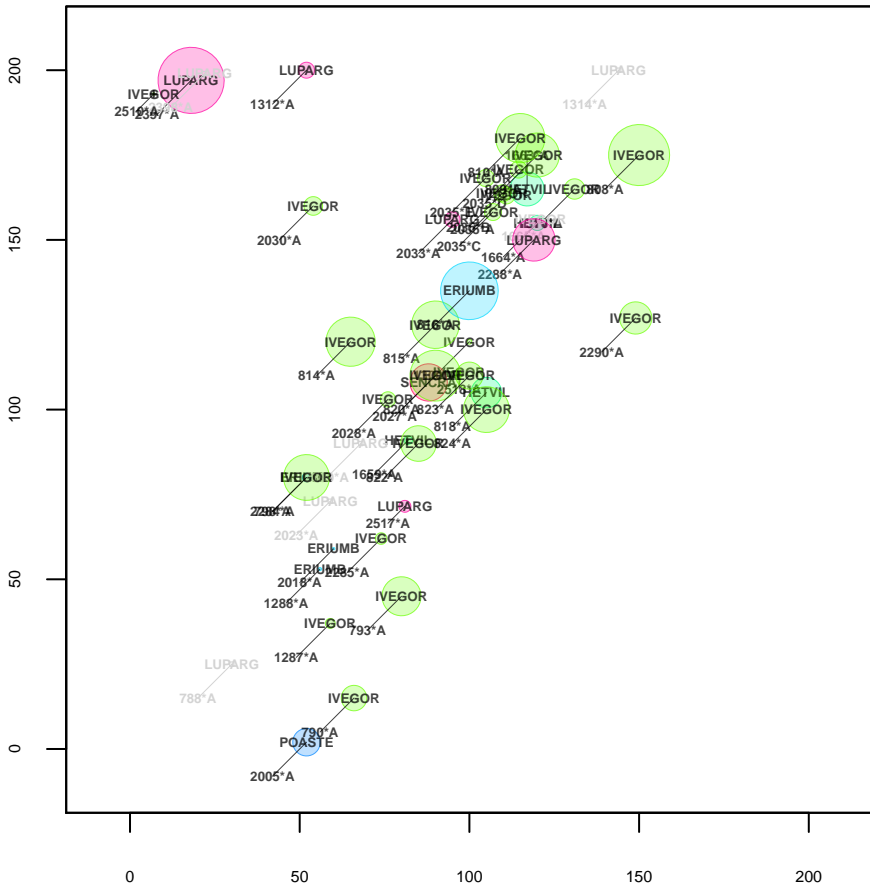
Plot 35



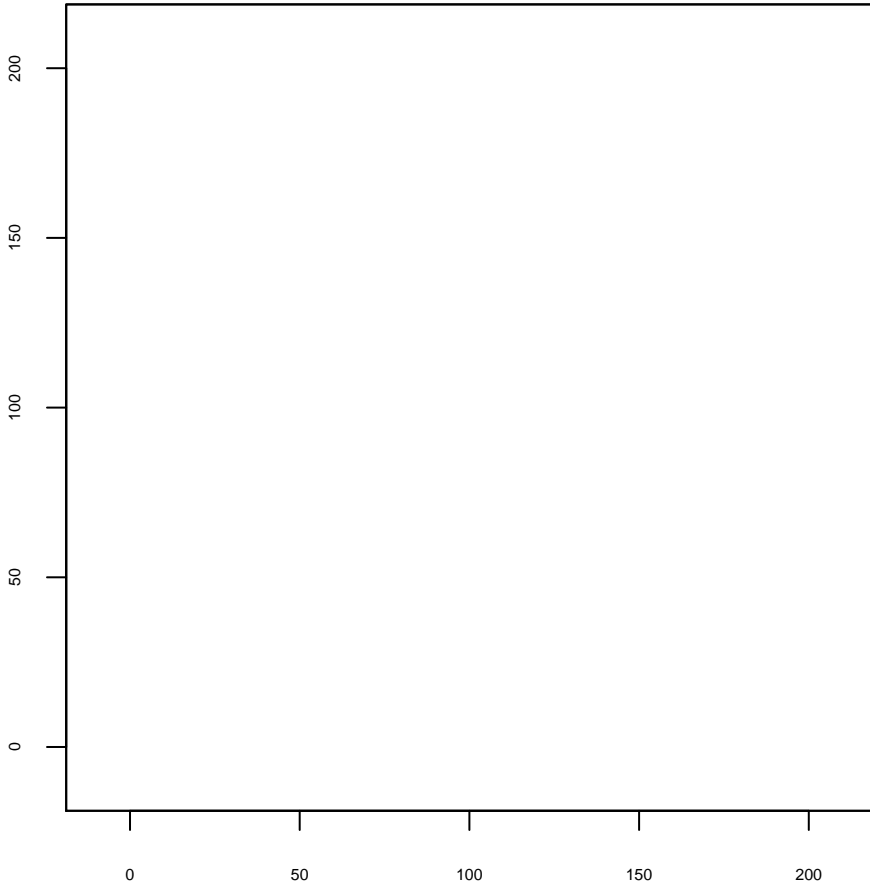
Plot 36



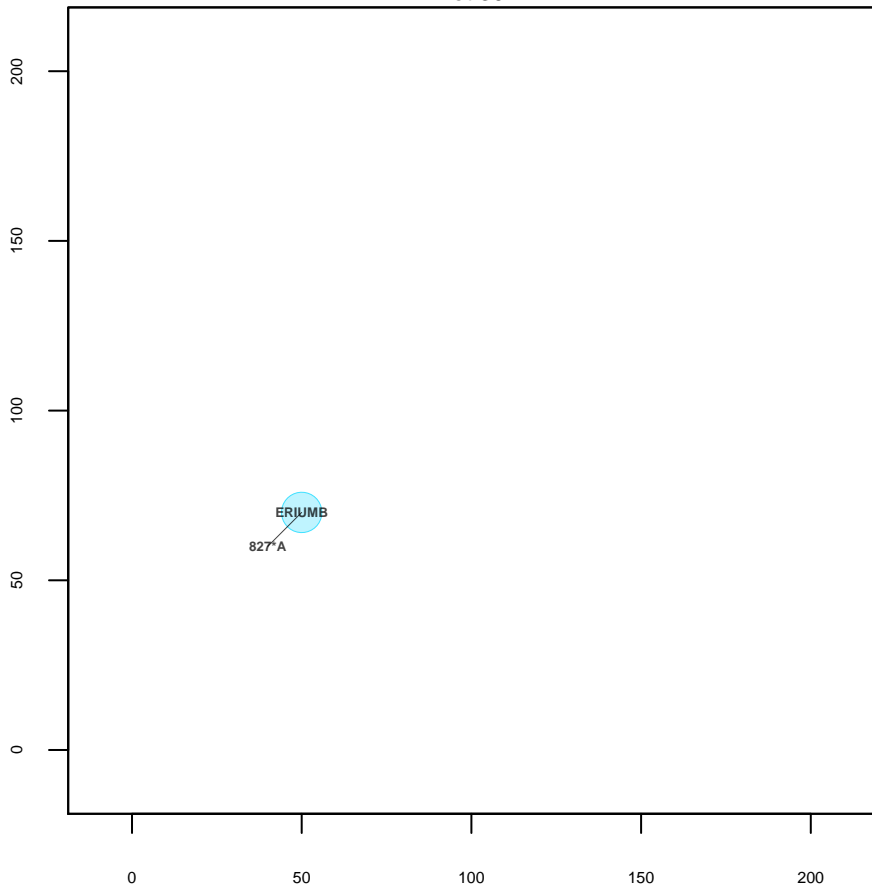
Plot 37



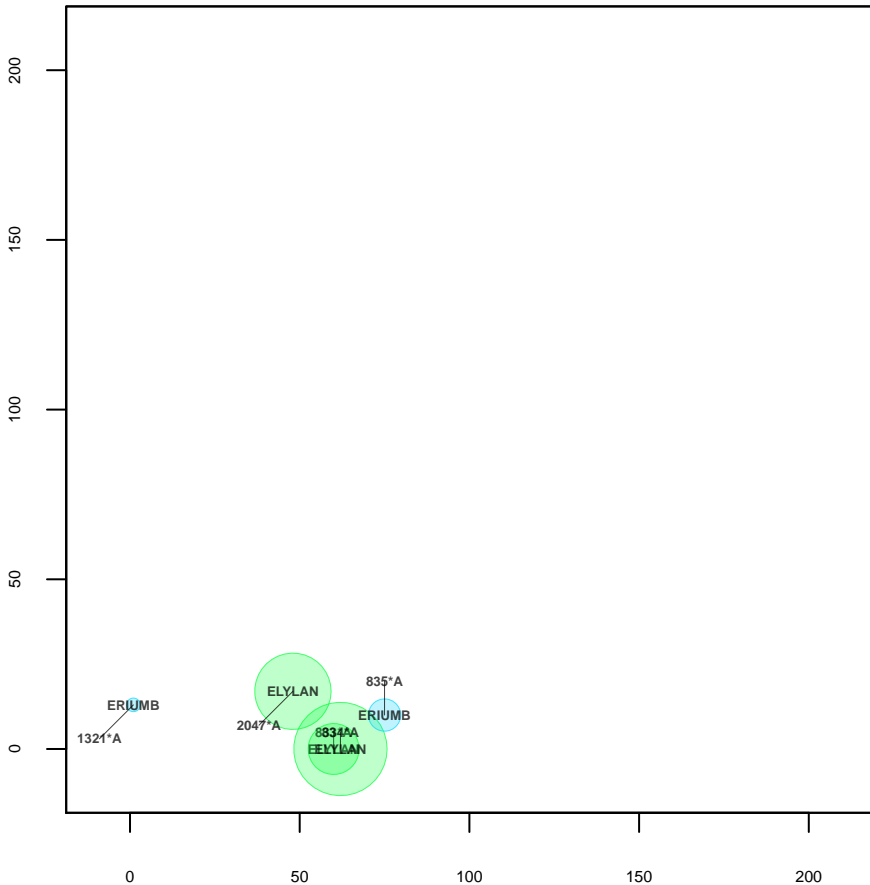
Plot 38



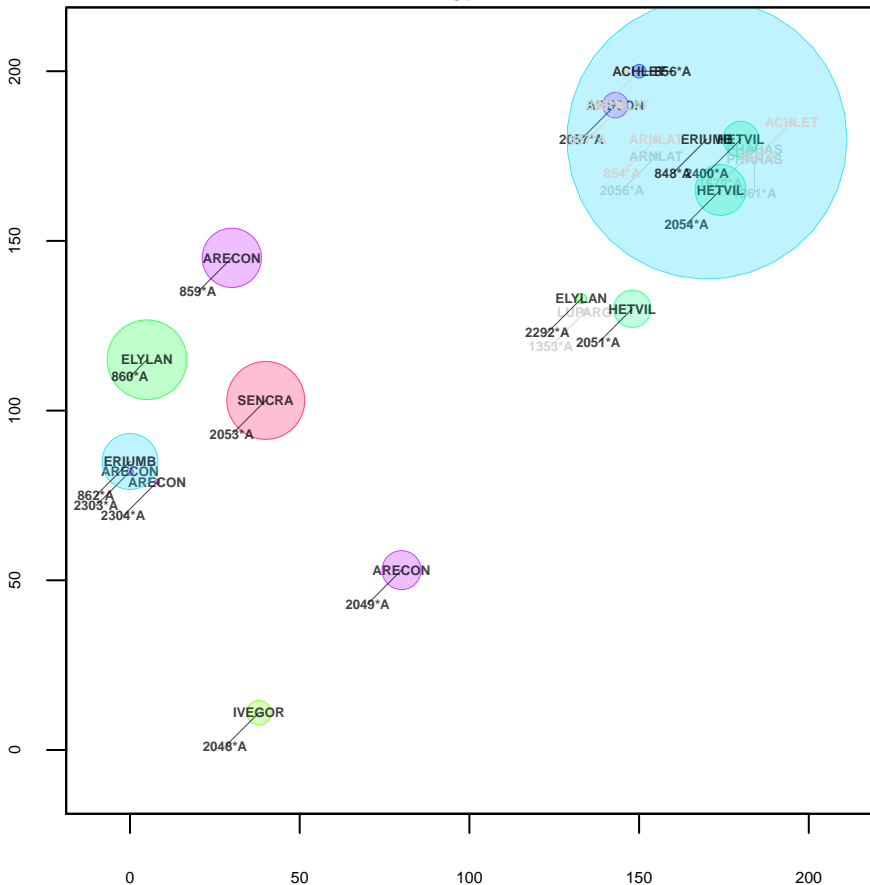
Plot 39



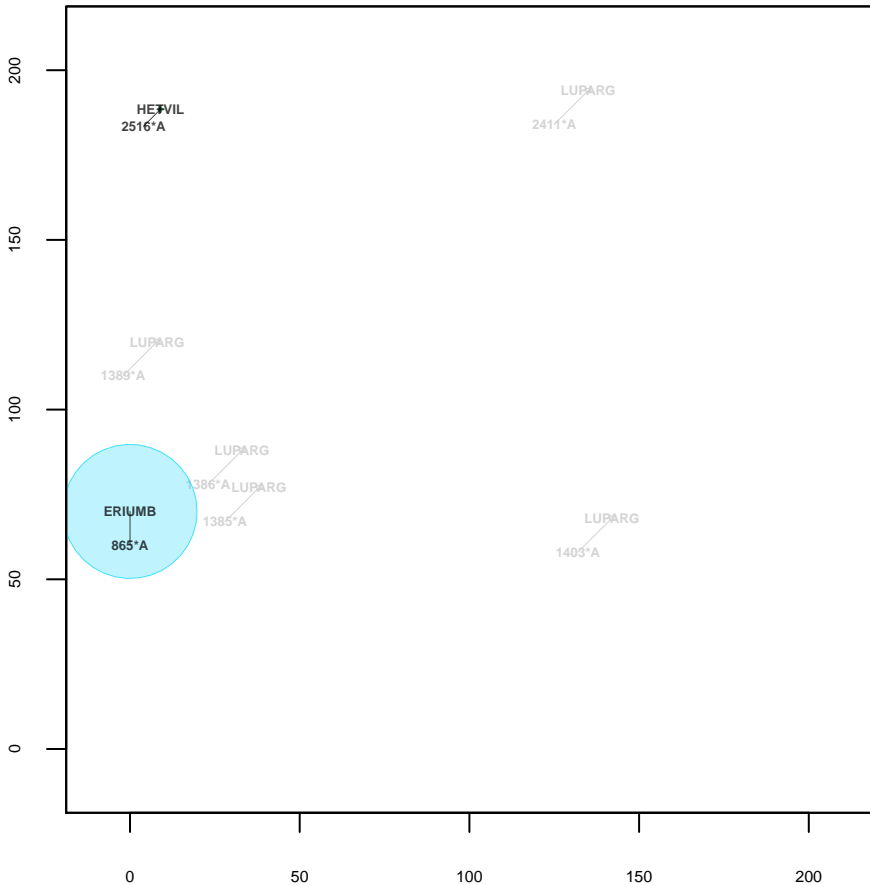
Plot 40



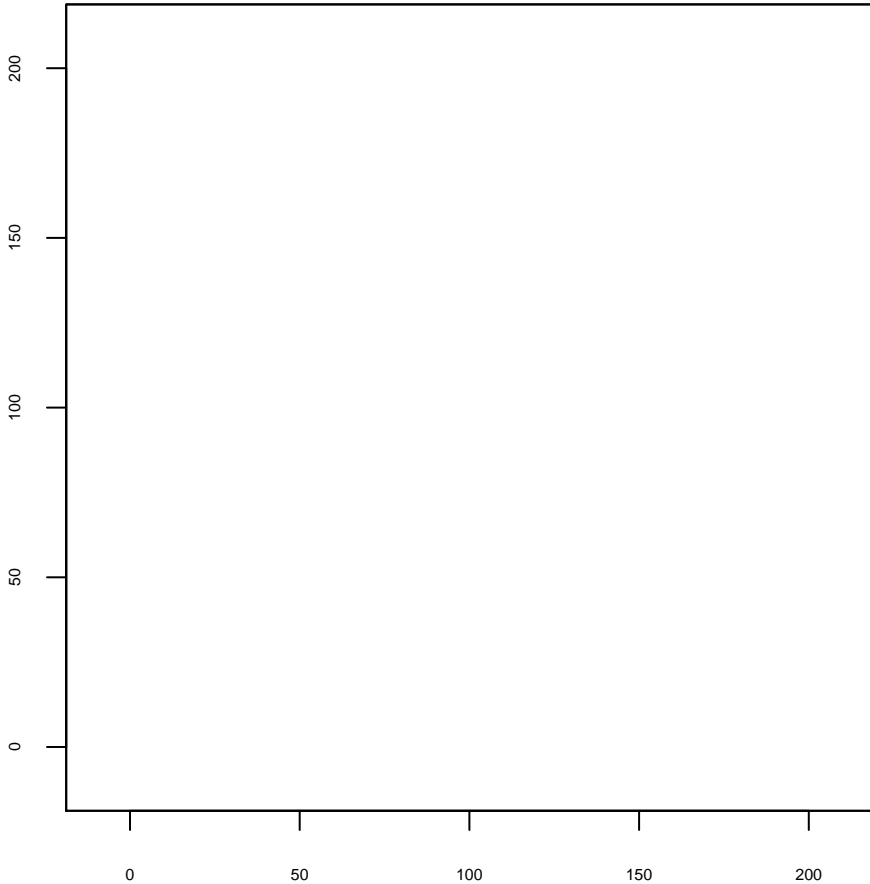
Plot 41



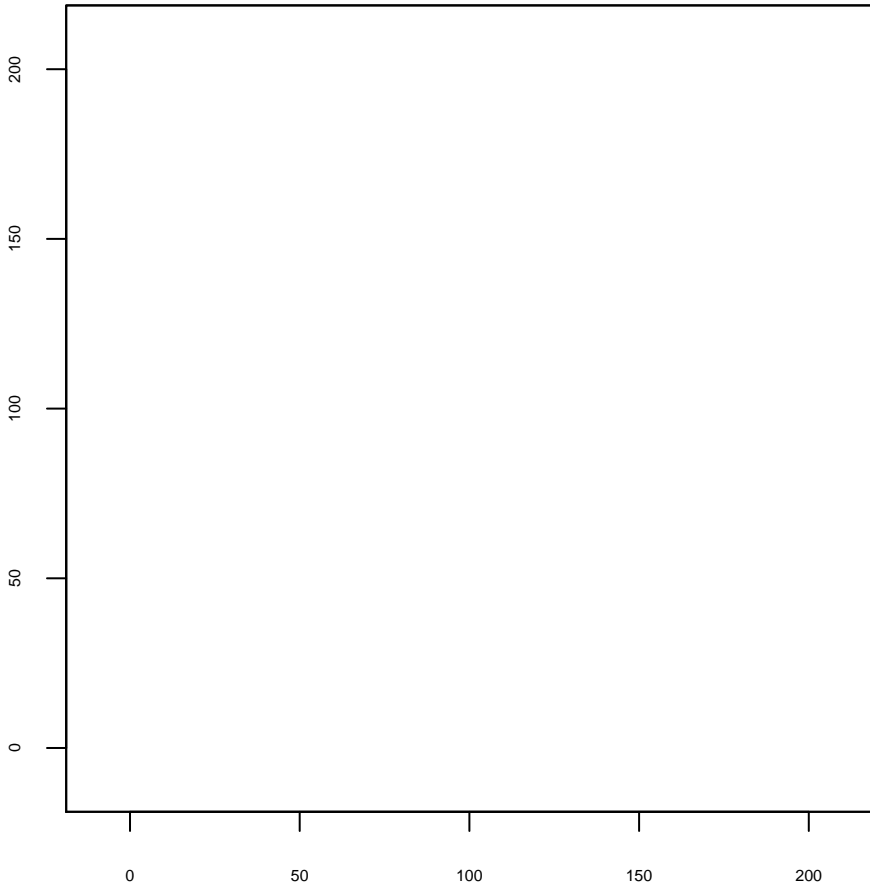
Plot 42



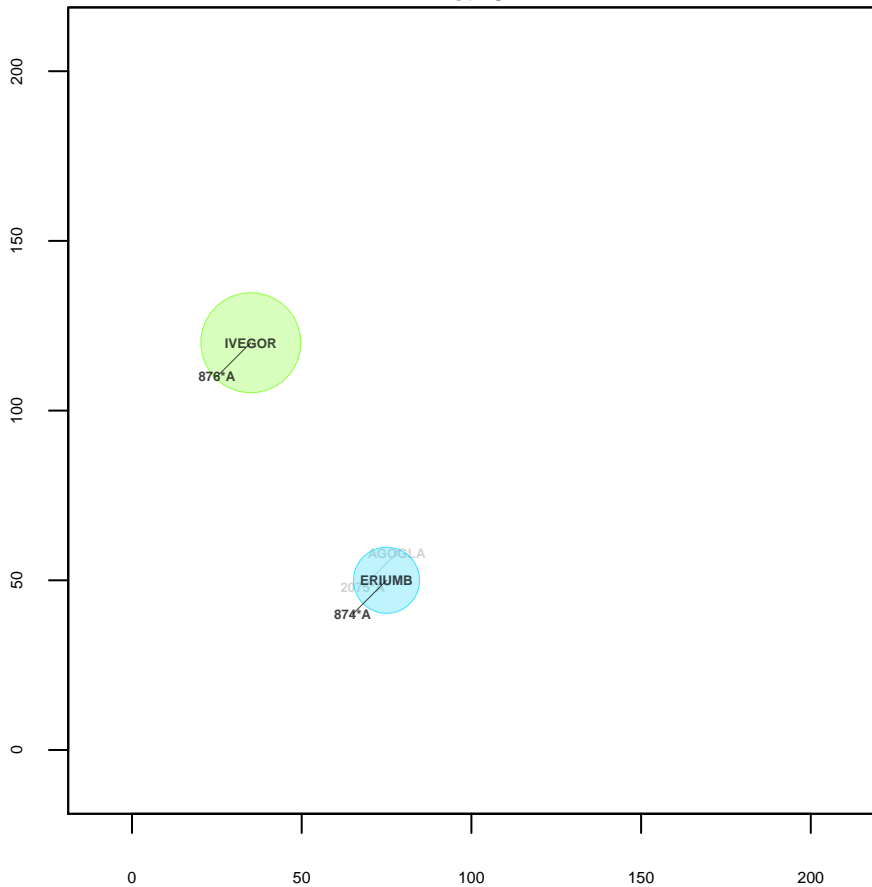
Plot 43



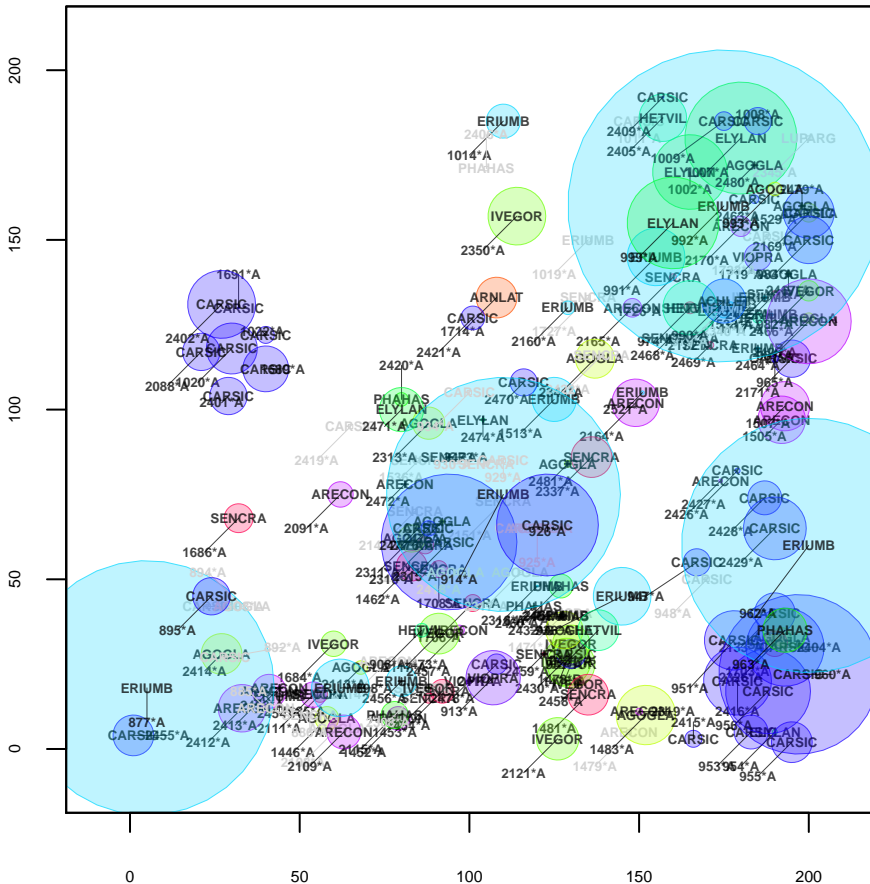
Plot 44



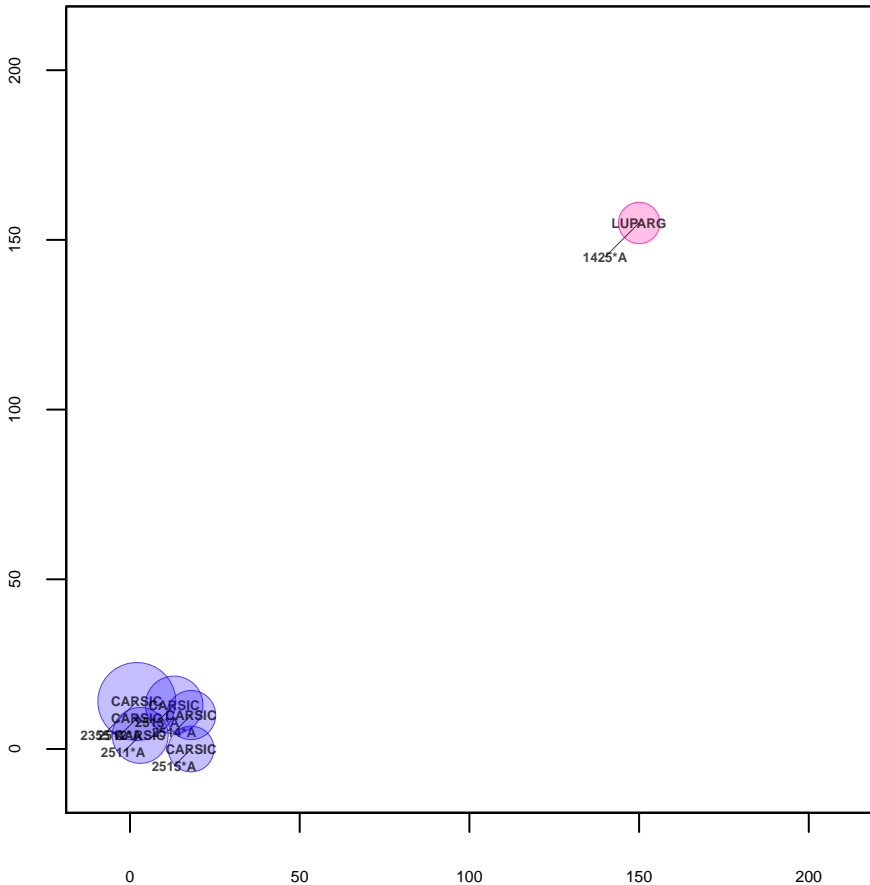
Plot 45



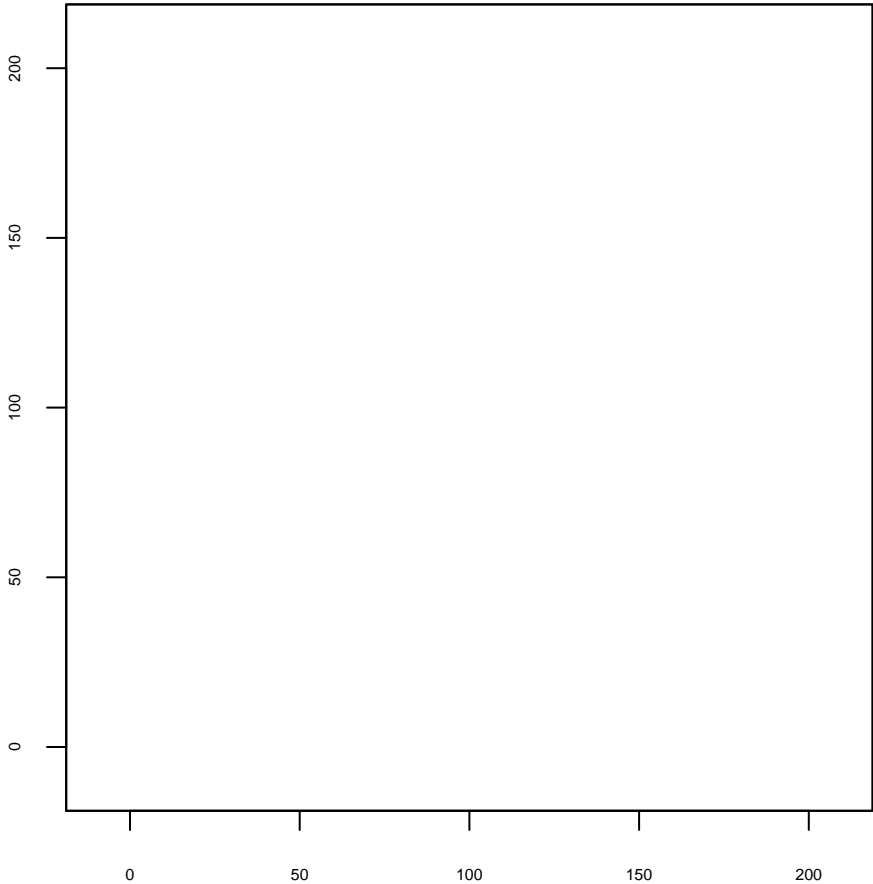
Plot 46



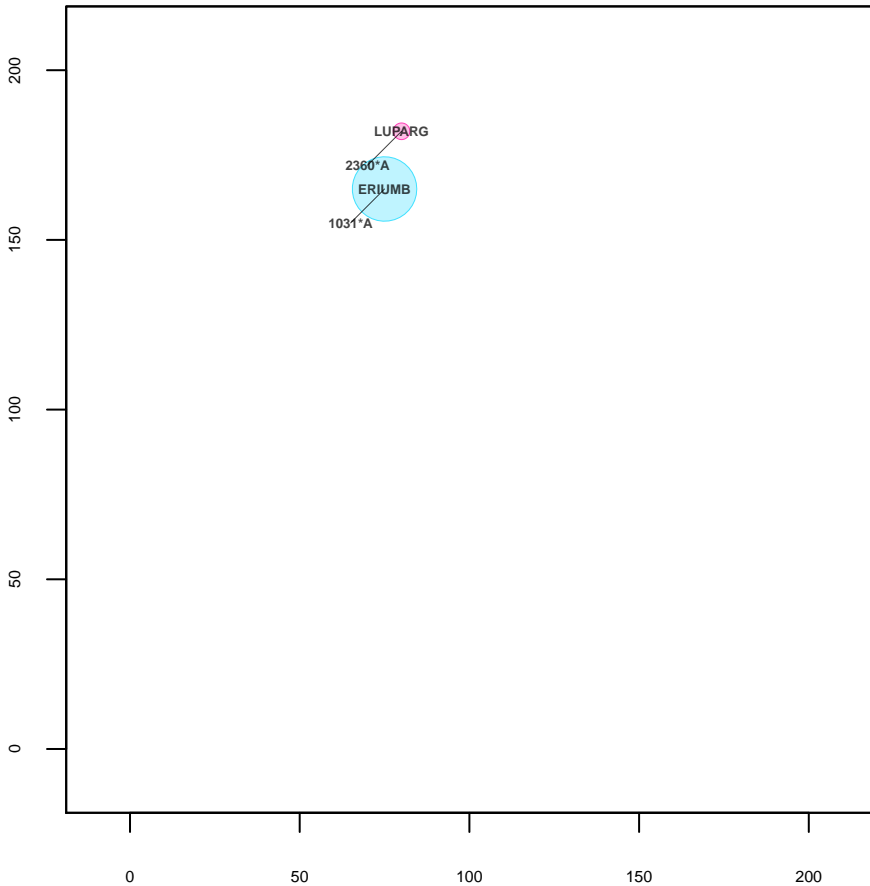
Plot 47



Plot 48



Plot 49



Plot 50

