

Figure 6A. Descriptive statistical analysis of annotated NCI60 gene predictor localization

The label on each part of the pie indicates the number of genes that were annotated with a particular cellular localization. 476 genes have “unknown” cellular localization.

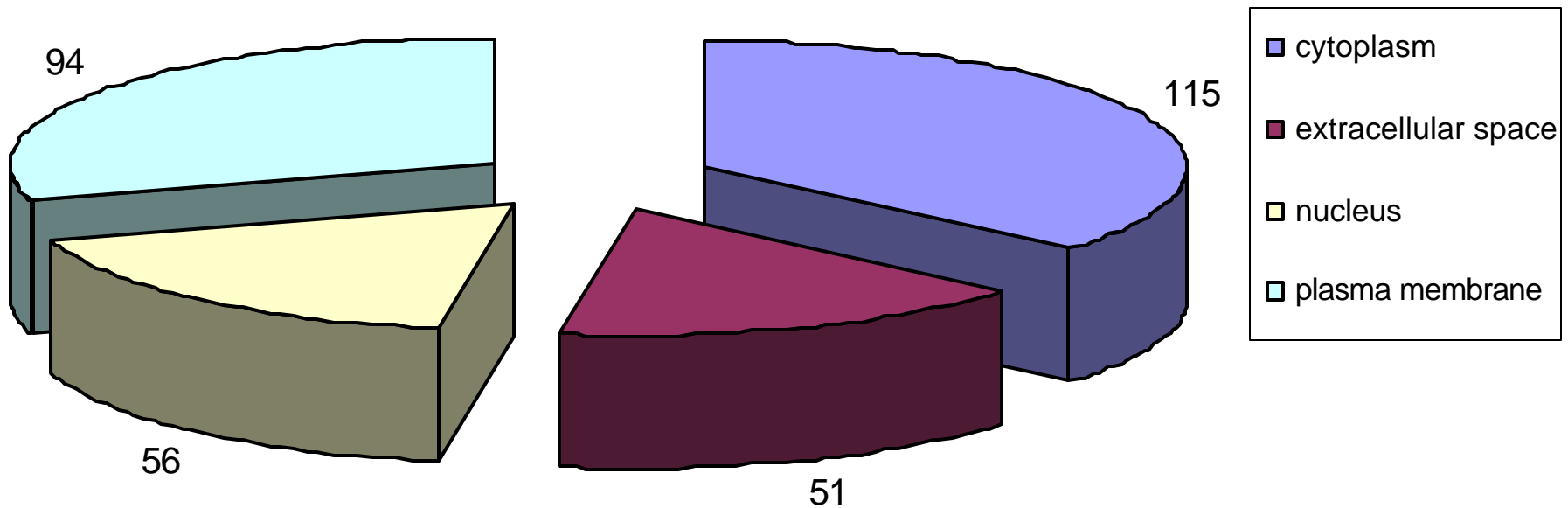


Figure 6B. Descriptive statistical analysis of annotated NCI60 gene predictor function

The label on each part of the pie indicates the number of genes that were annotated with a particular function. 605 genes have “unknown” function.

