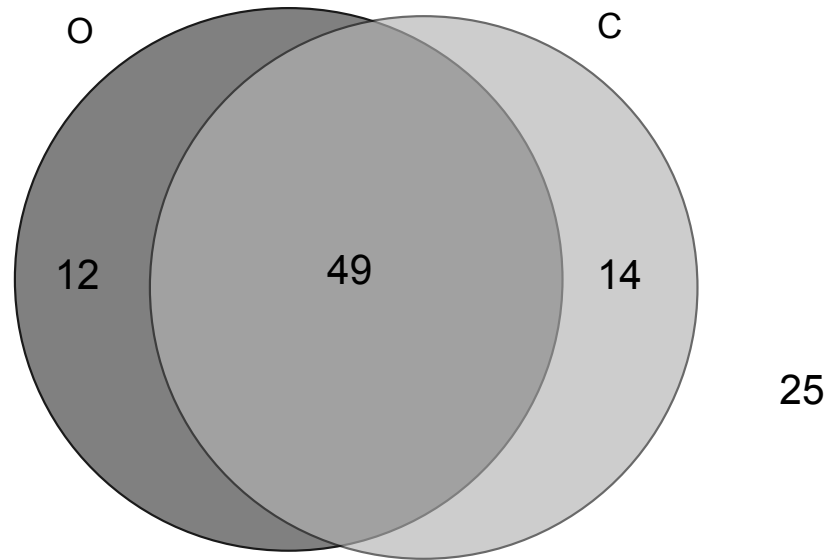


STE up-regulated genes-1850 genes

TPs from predictions using O and C motif set (%)



STE specific up-regulated genes-362 genes

TPs from predictions using O and C motif set (%)

