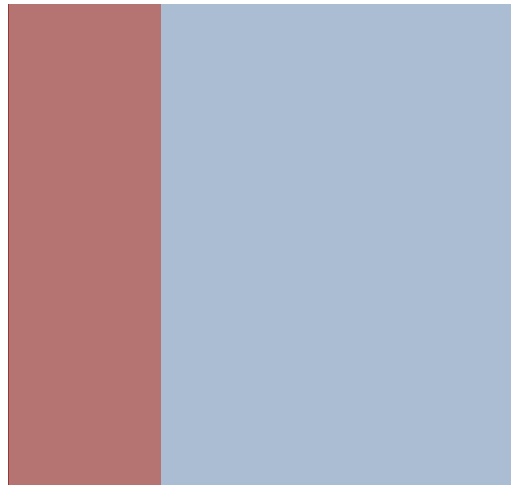
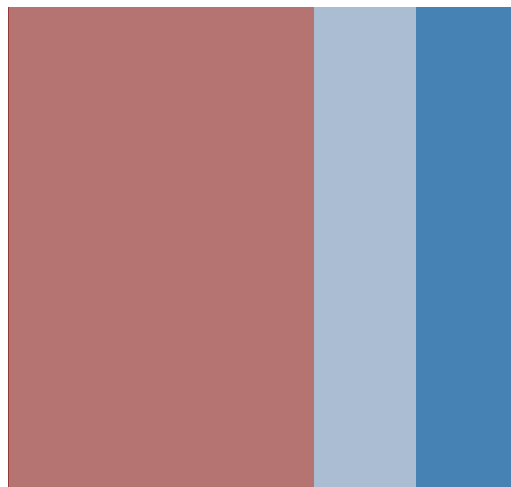


chaperone binding
(23 genes)

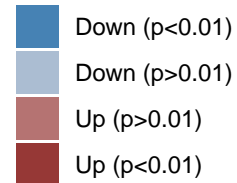
JFYL007_U_3_0.1vsJFYL007_AS_3_0.1



OKYL029_U_3_0.1vsOKYL029_AS_3_0.1



Direction



0% 25% 50% 75% 100%
% of Genes

