

How does it correlate?

In this tab you can explore the correlations between individual traits. The values representing individual accessions are based on the average calculated per accession per condition. The names of the accessions are plotted next to the points representing the individual line. The Pearson correlation coefficient (r) and p-value are plotted above the graphs

We hope that those correlation graphs will be helpful to select accessions showing extreme phenotypes, that can be used in, for example for bulked segregant analysis or in depth individual study of the accession.

Trait 1

average Lateral Root Length 75 mM NaCl

Trait 2

Total Root Size 75 mM NaCl

Change the phenotypes to correlate

The Pearson Correlation coefficient (r) is
0.6929243

The p-value is
6.032573e-51

Correlation strength and significance
are reported here

