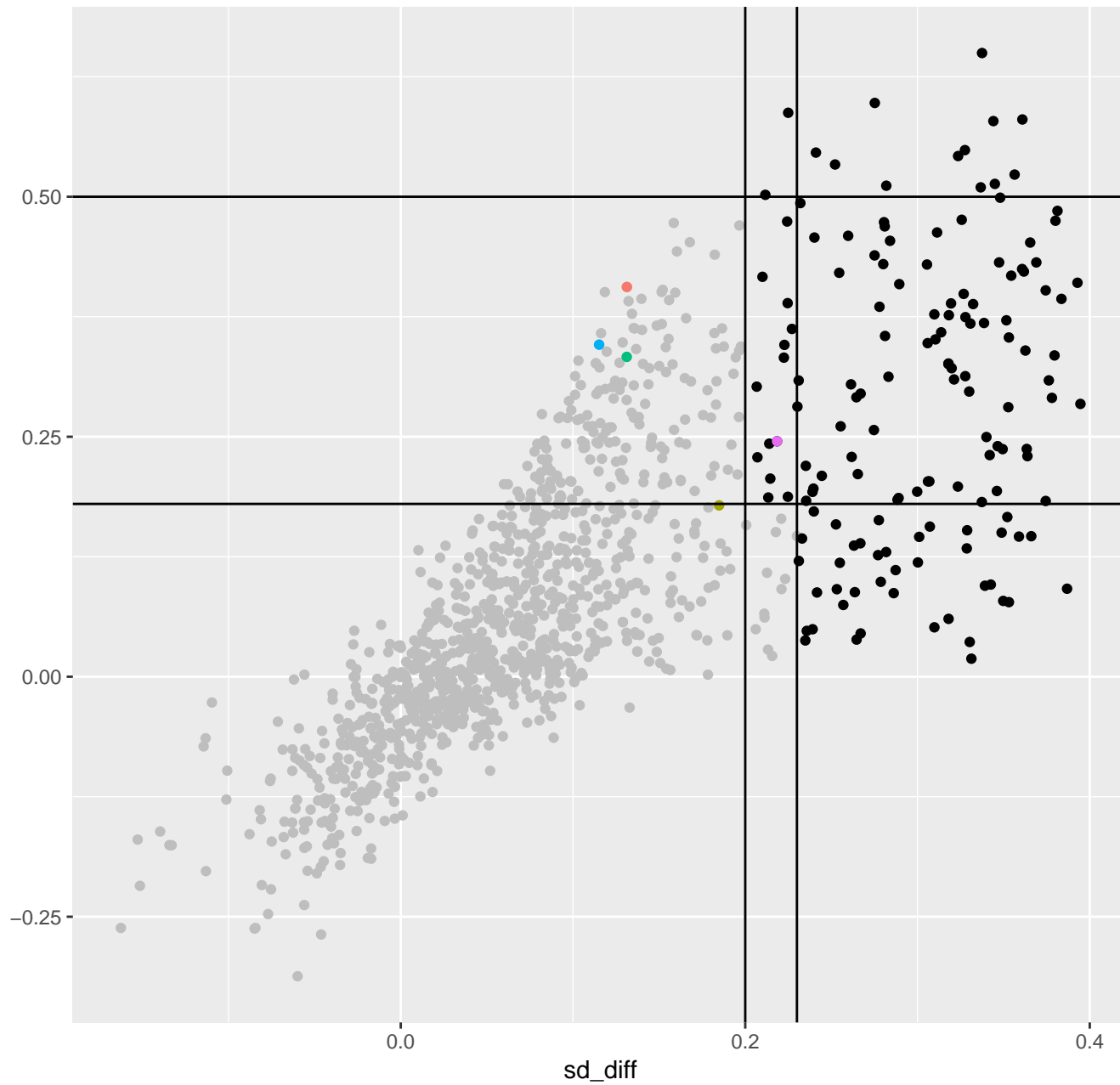


w_corr_random_high

diff_of_upper_whiskers



cluster

- enzyme regulator activity_GO_30_1_all
- lipid binding
- organelle fission_GO_30_2_mut
- protein phosphorylation
- structural constituent of ribosome