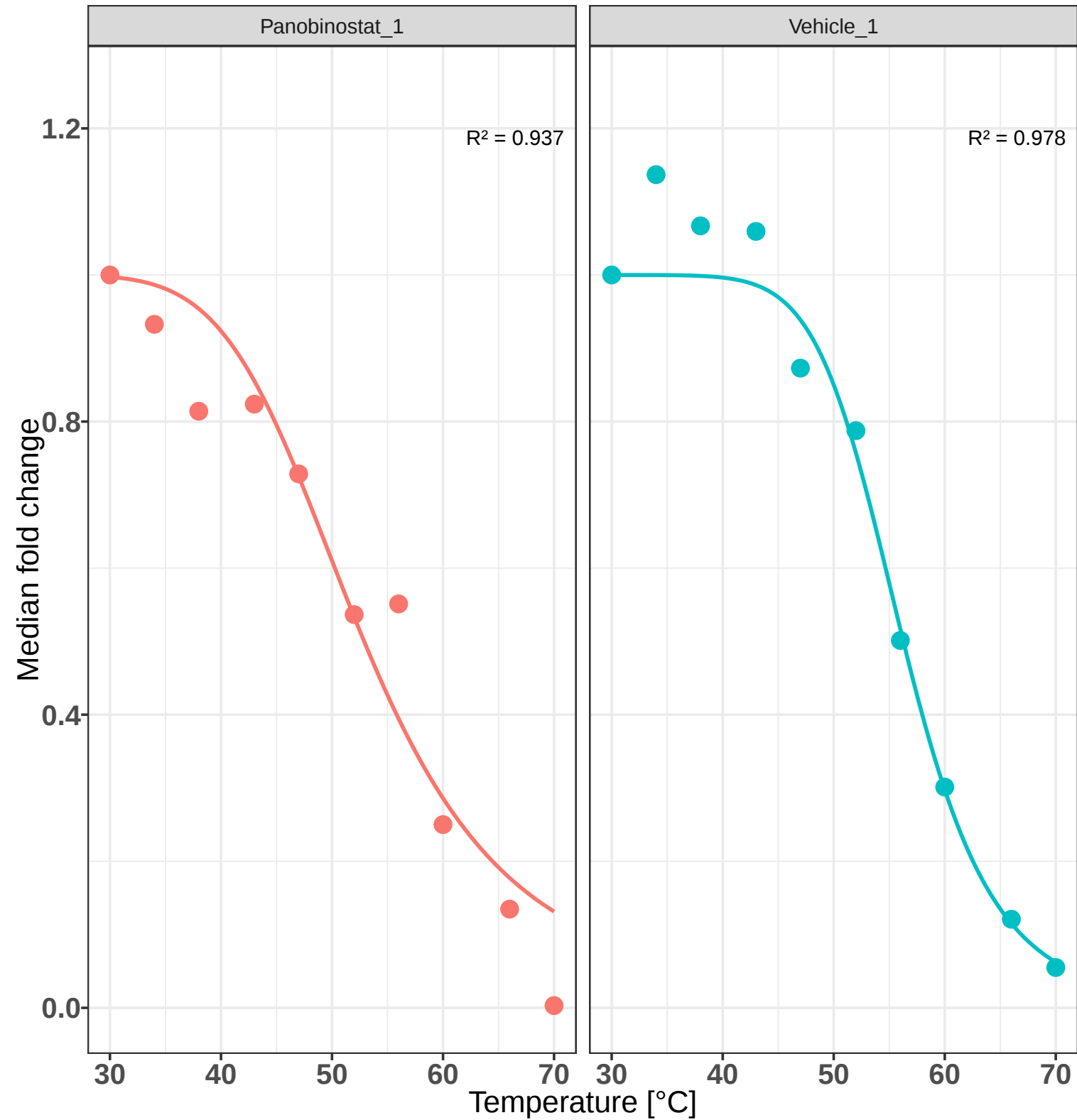
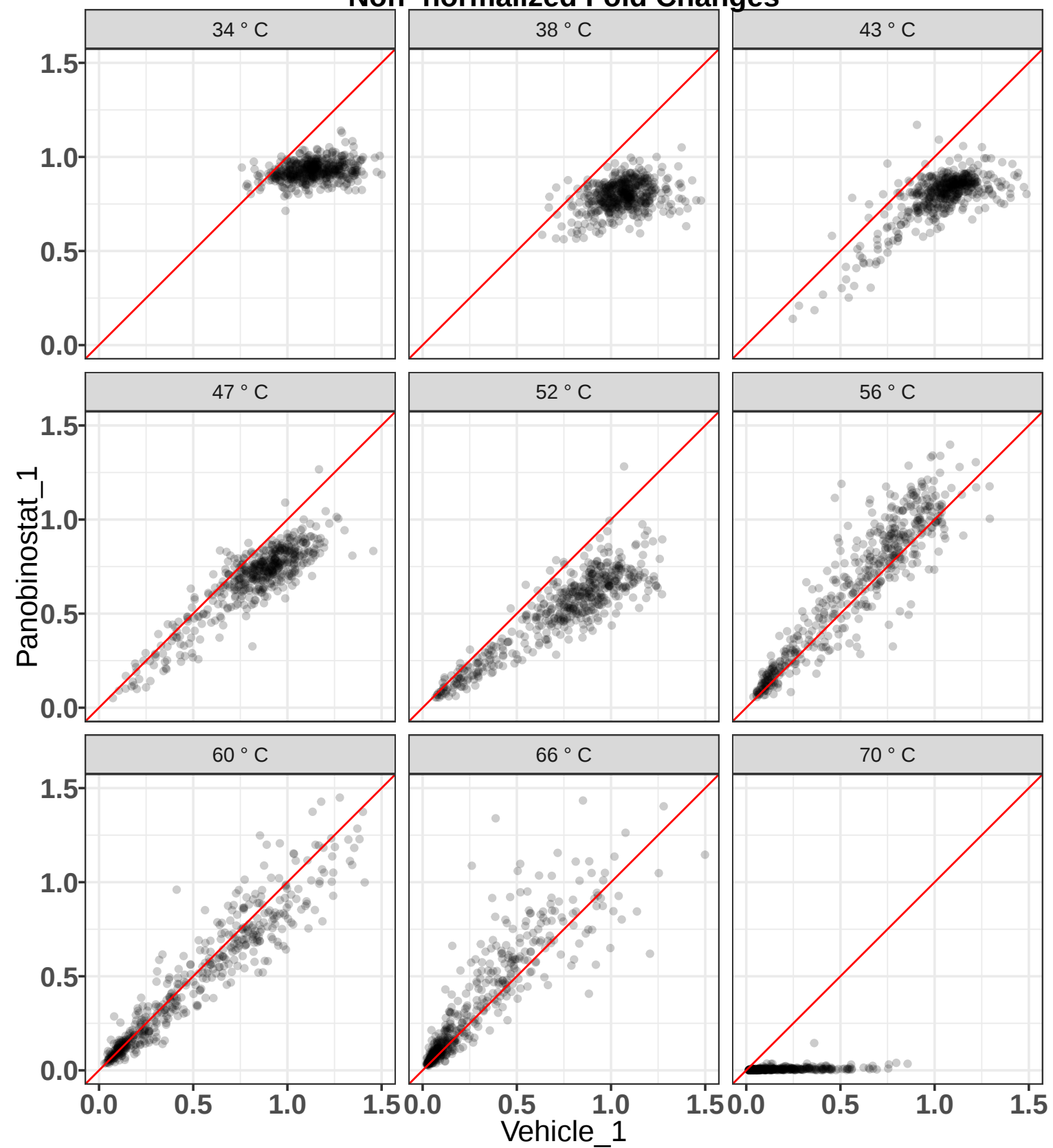


Normalization curves

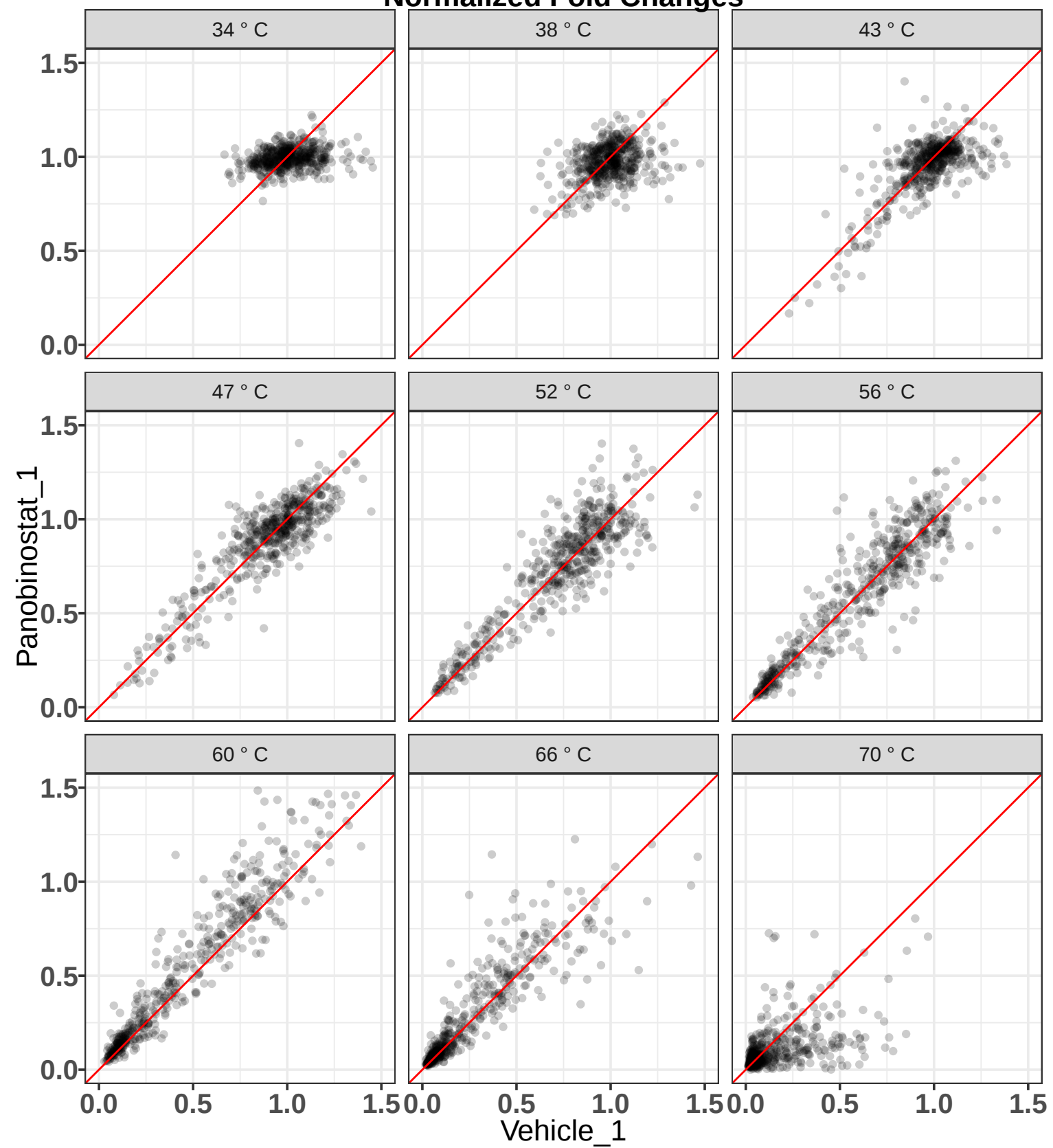
based on 62 proteins

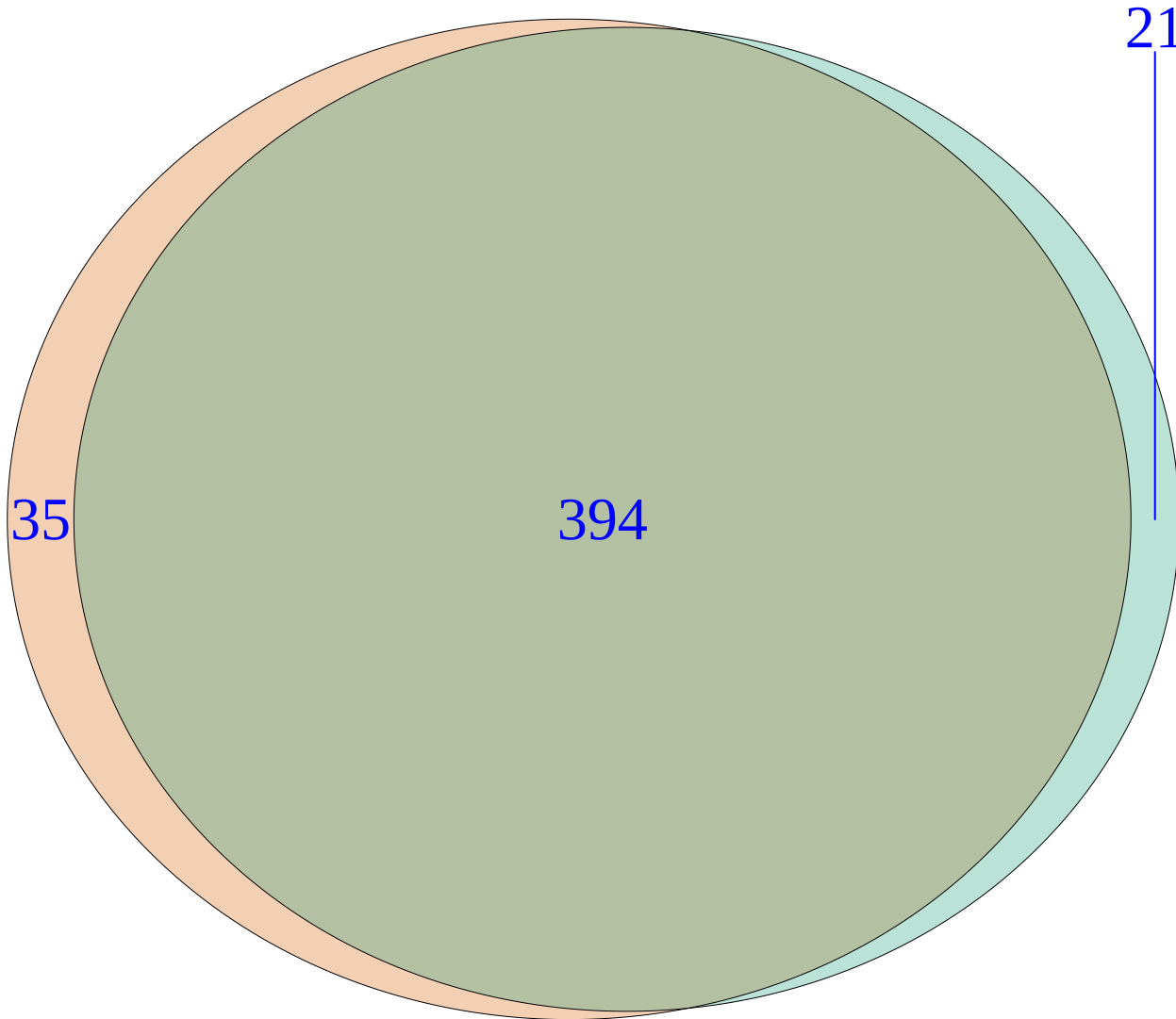


Non-normalized Fold Changes



Normalized Fold Changes





Vehicle_1
484 proteins identified overall

Panobinostat_1

Vehicle_1

425

59

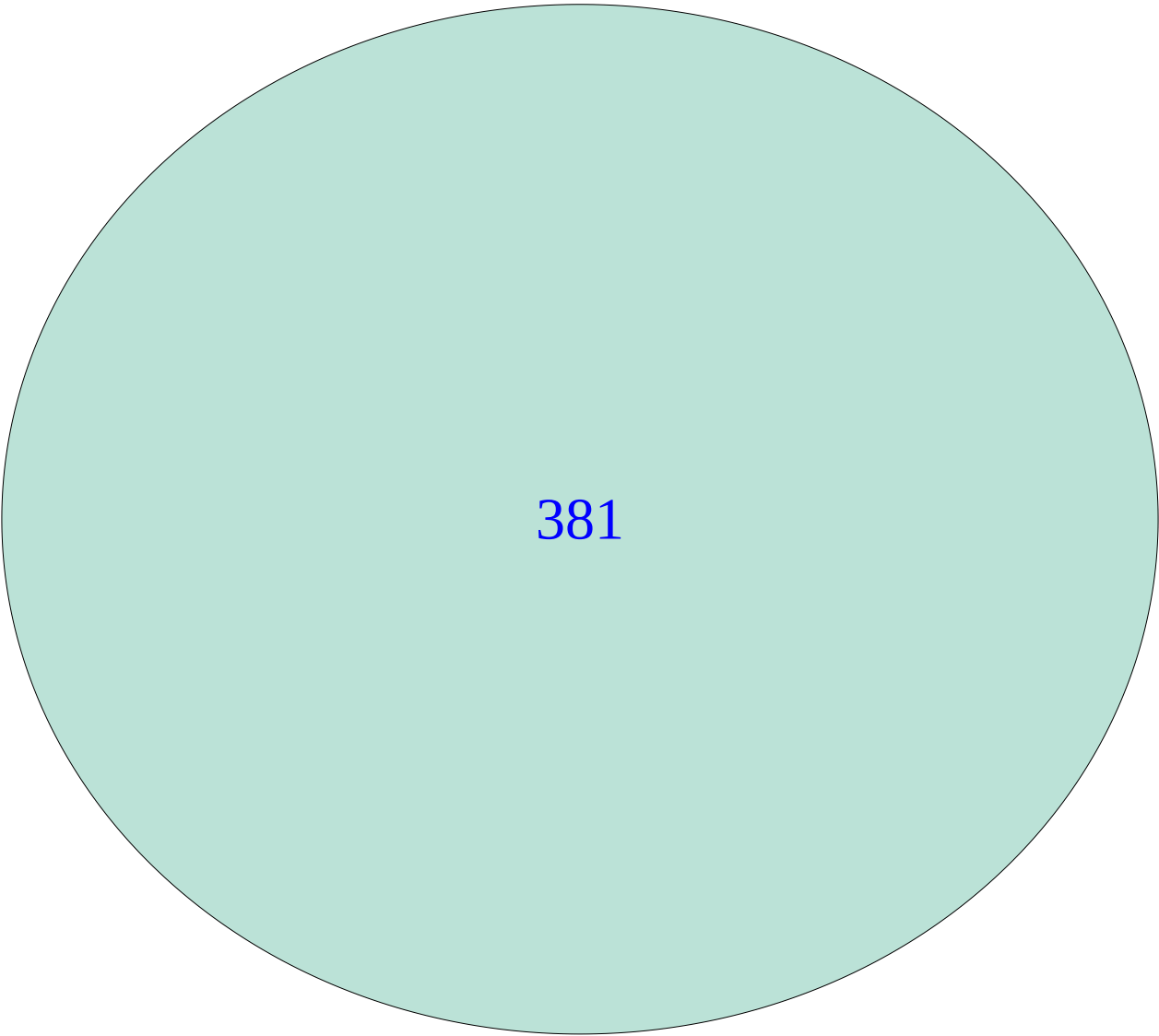
484

R2 >= 0.8

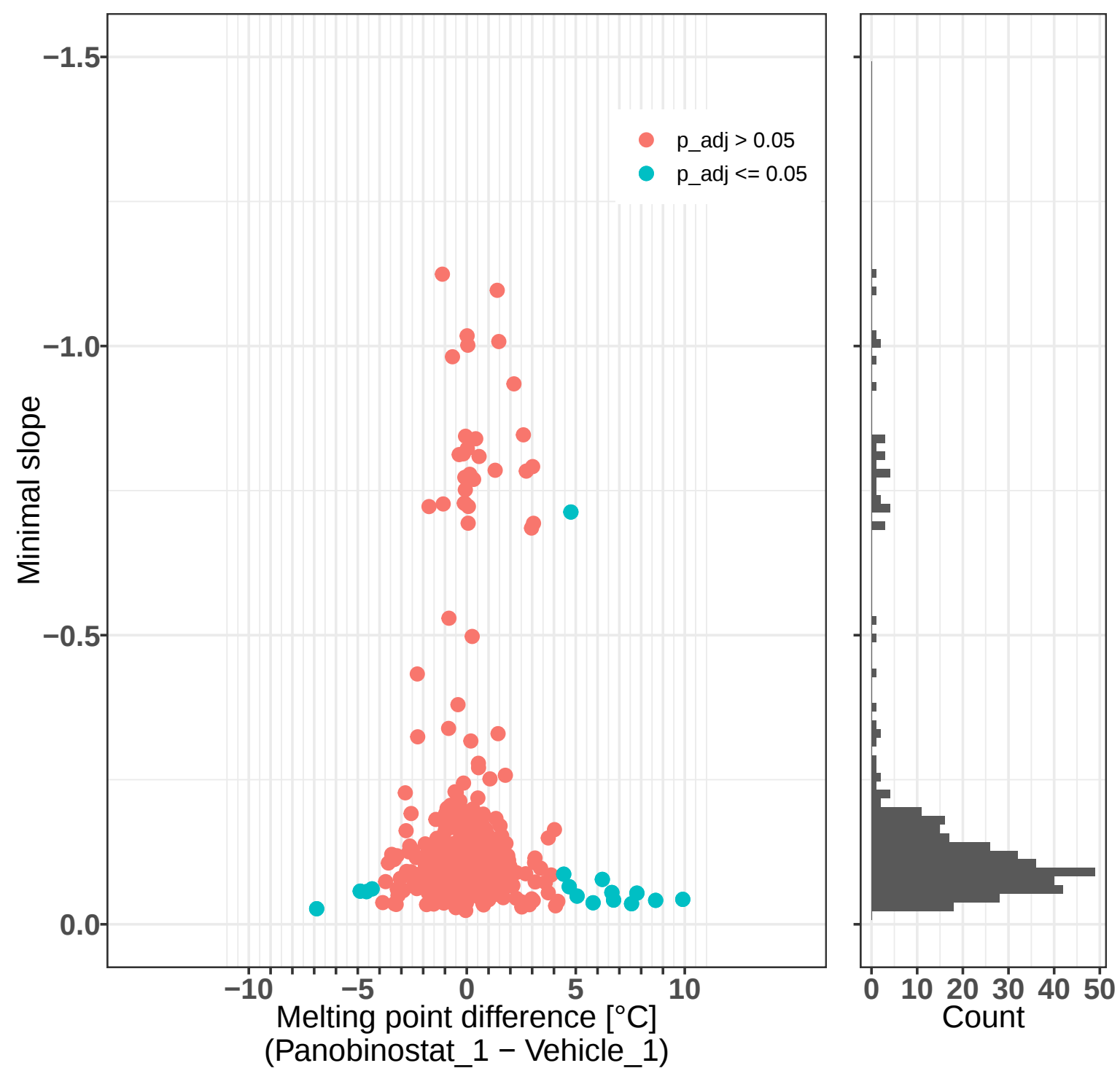
R2 < 0.8

Total

Panobinostat_1_vs_Vehicle_1



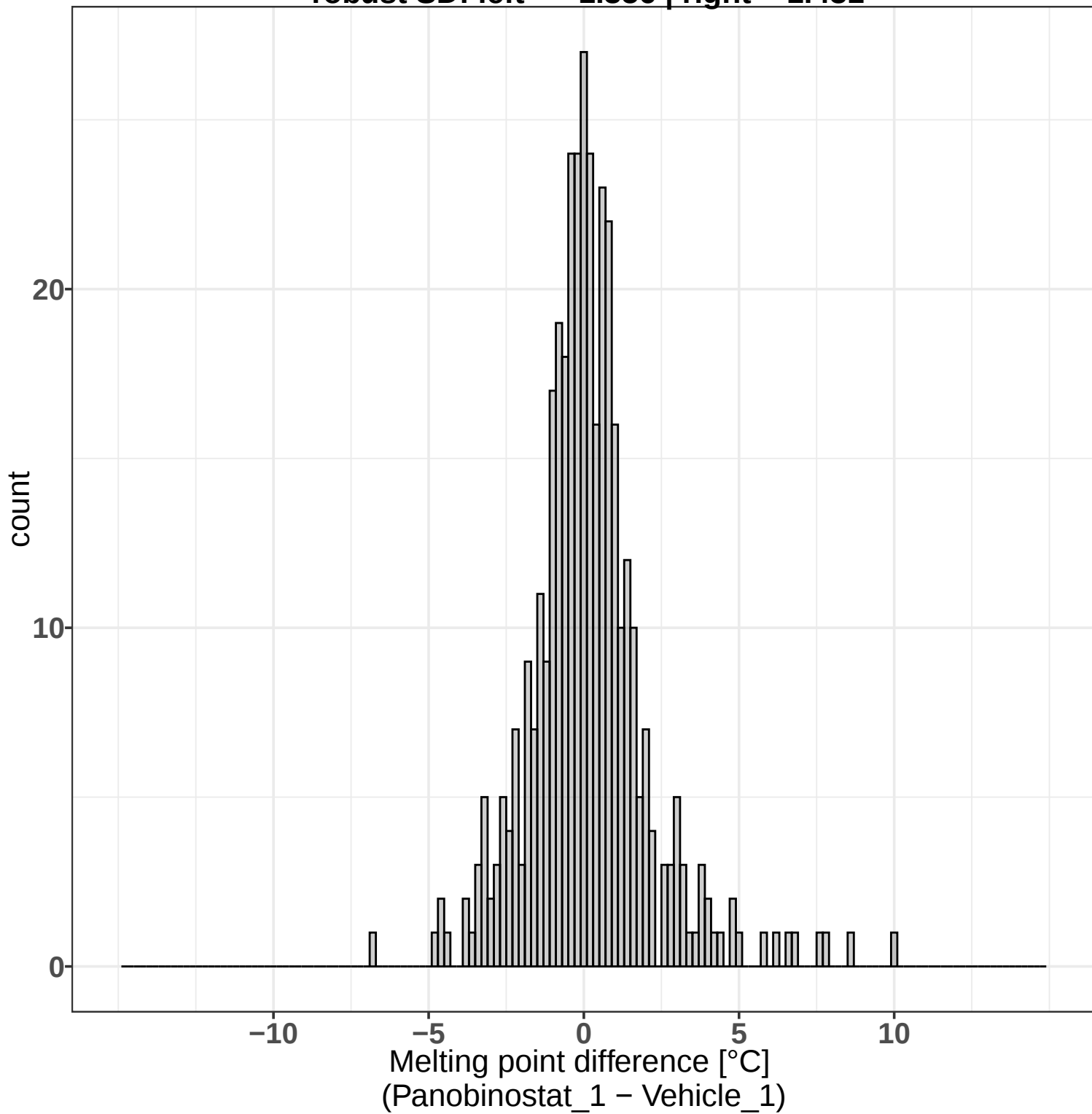
nobinostat_1_vs_Vehicle_1	103	484
passed_filter	filtered_out	Total



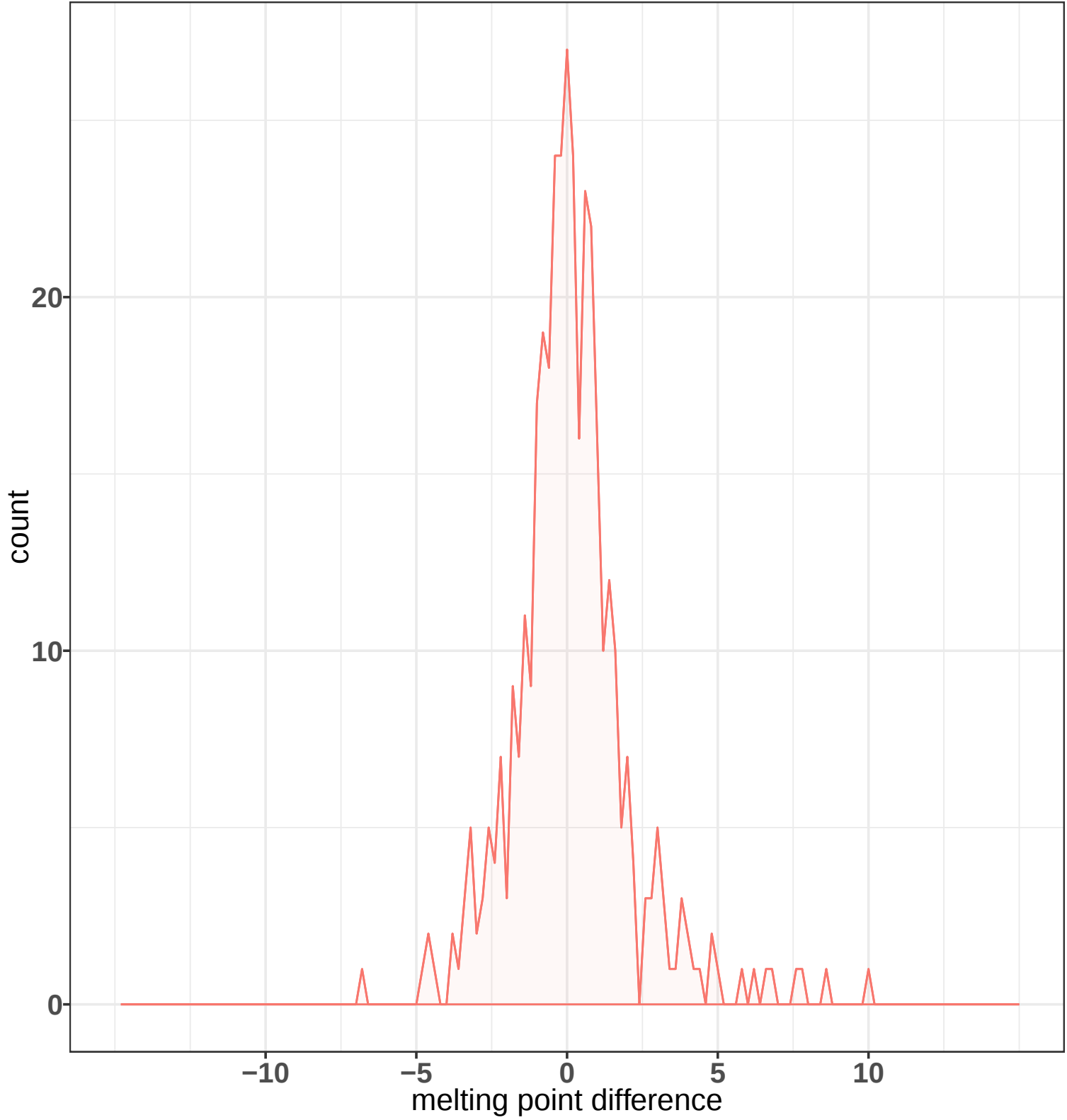
n = 379

Panobinostat_1 vs. Vehicle_1

**median = 0.018 ,
robust SD: left = -1.356 | right = 1.431**



n = 394 , R_sq >= 0.8



comparison Panobinostat_1 - Vehicle_1