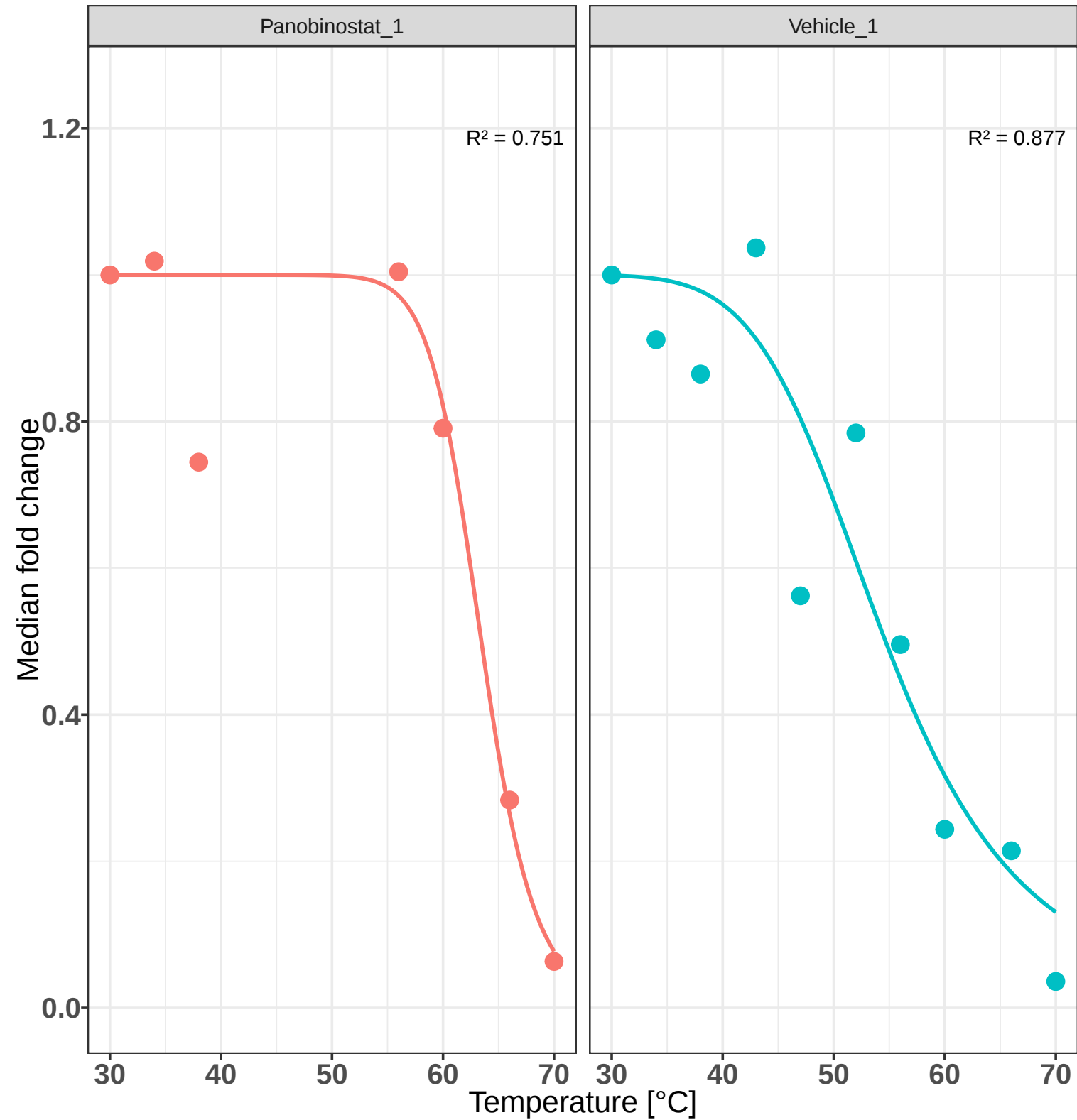
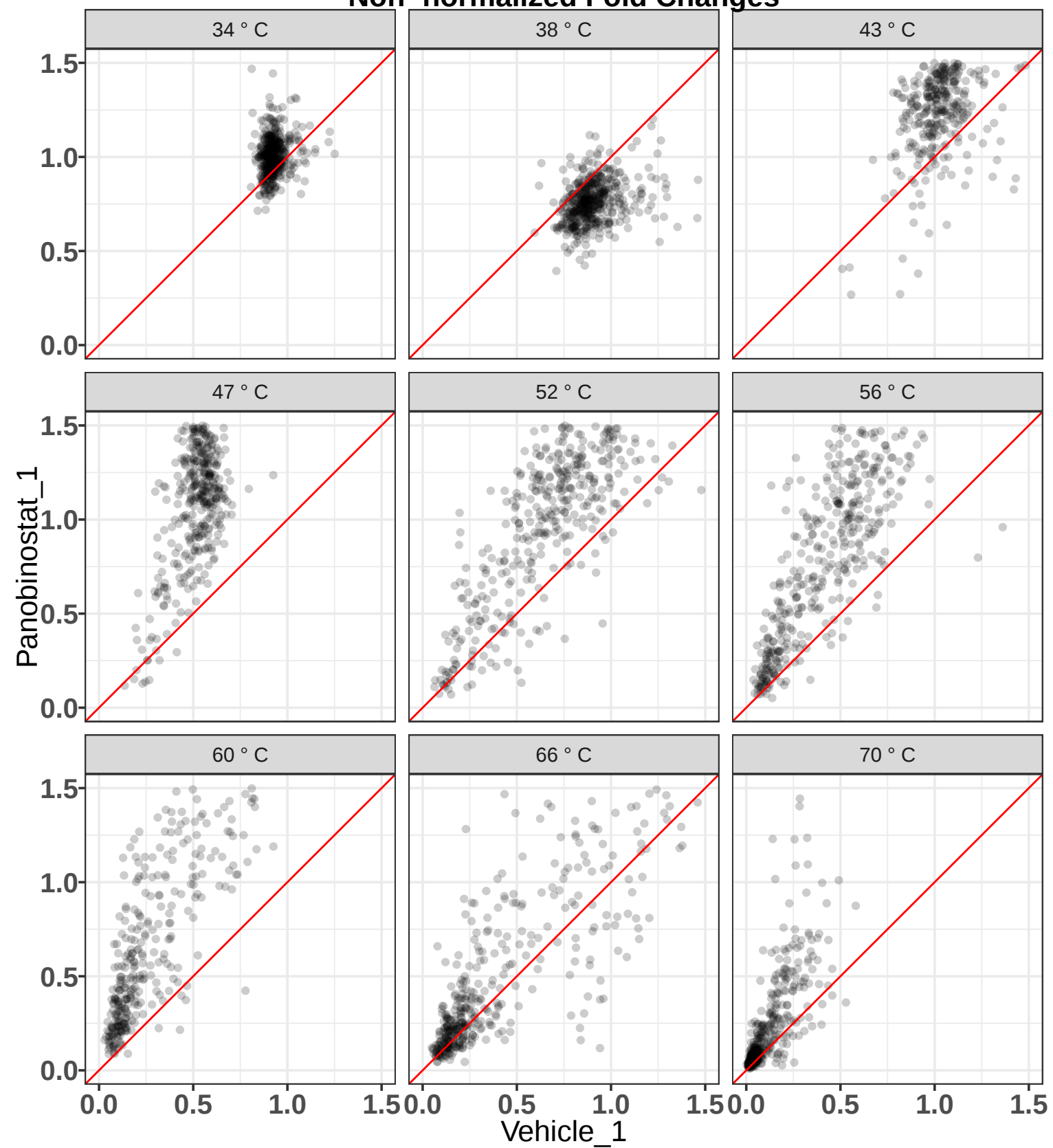


Normalization curves

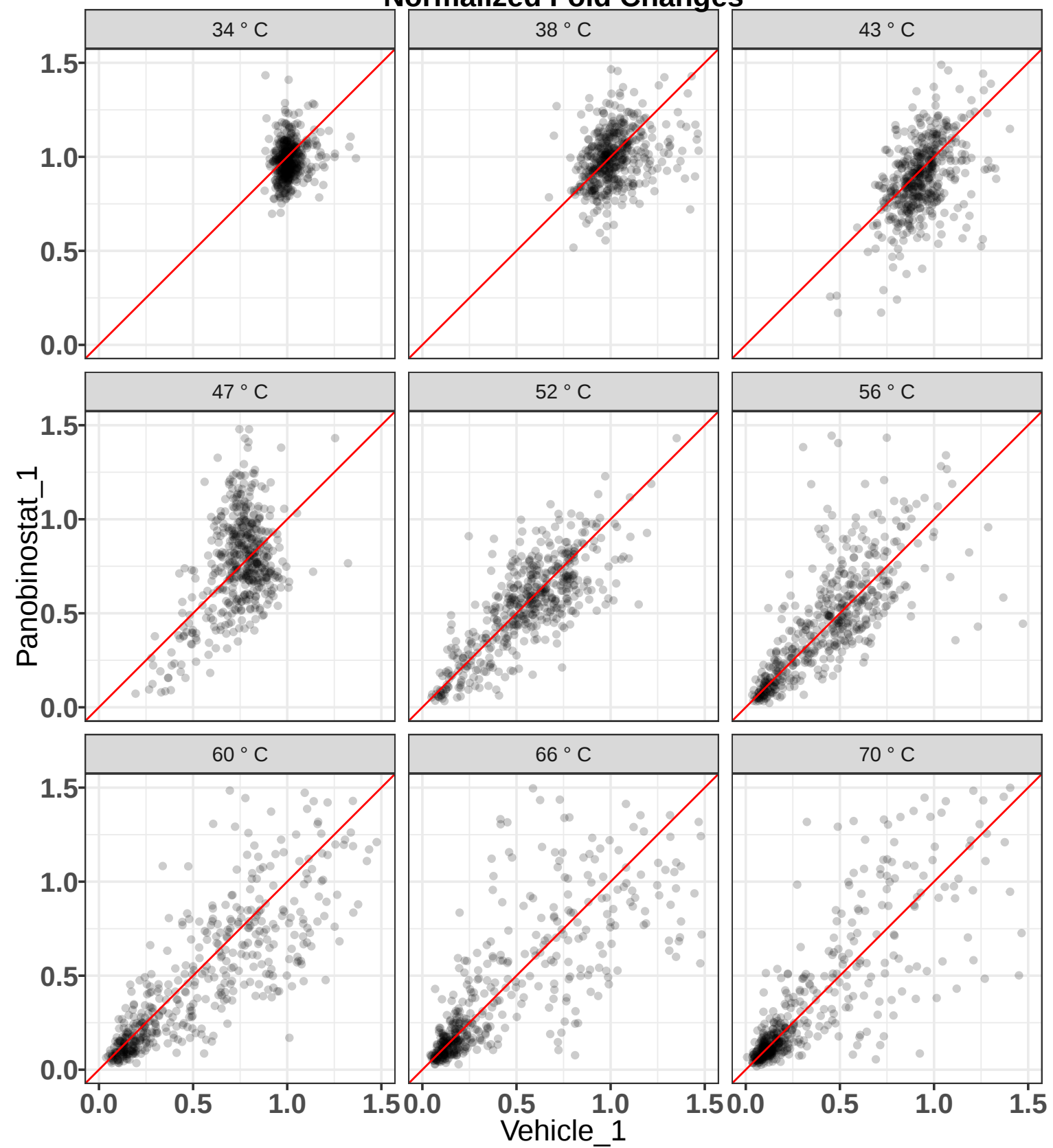
based on 37 proteins

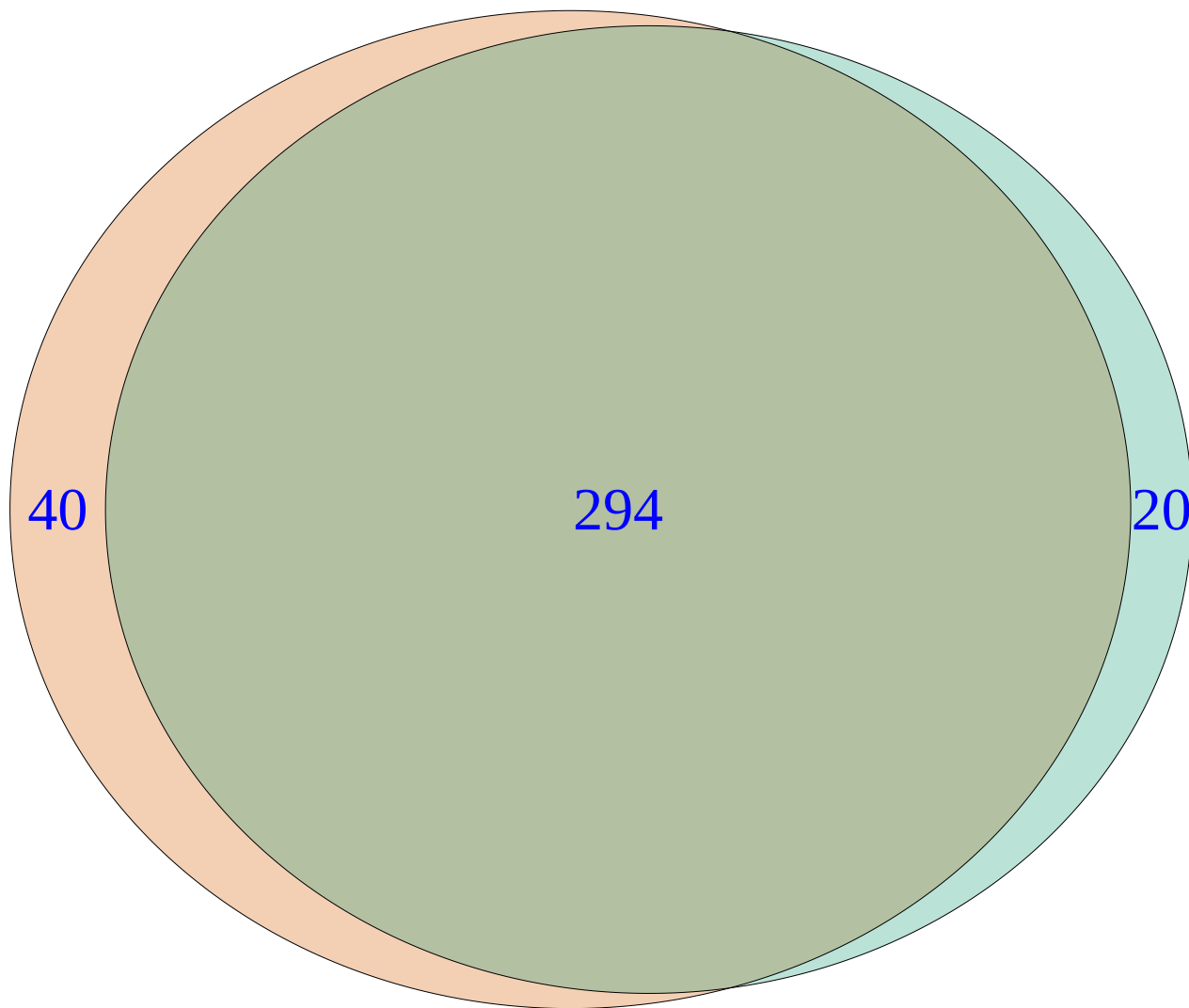


Non-normalized Fold Changes



Normalized Fold Changes





Vehicle_1
484 proteins identified overall
Panobinostat_1

Vehicle_1

334

150

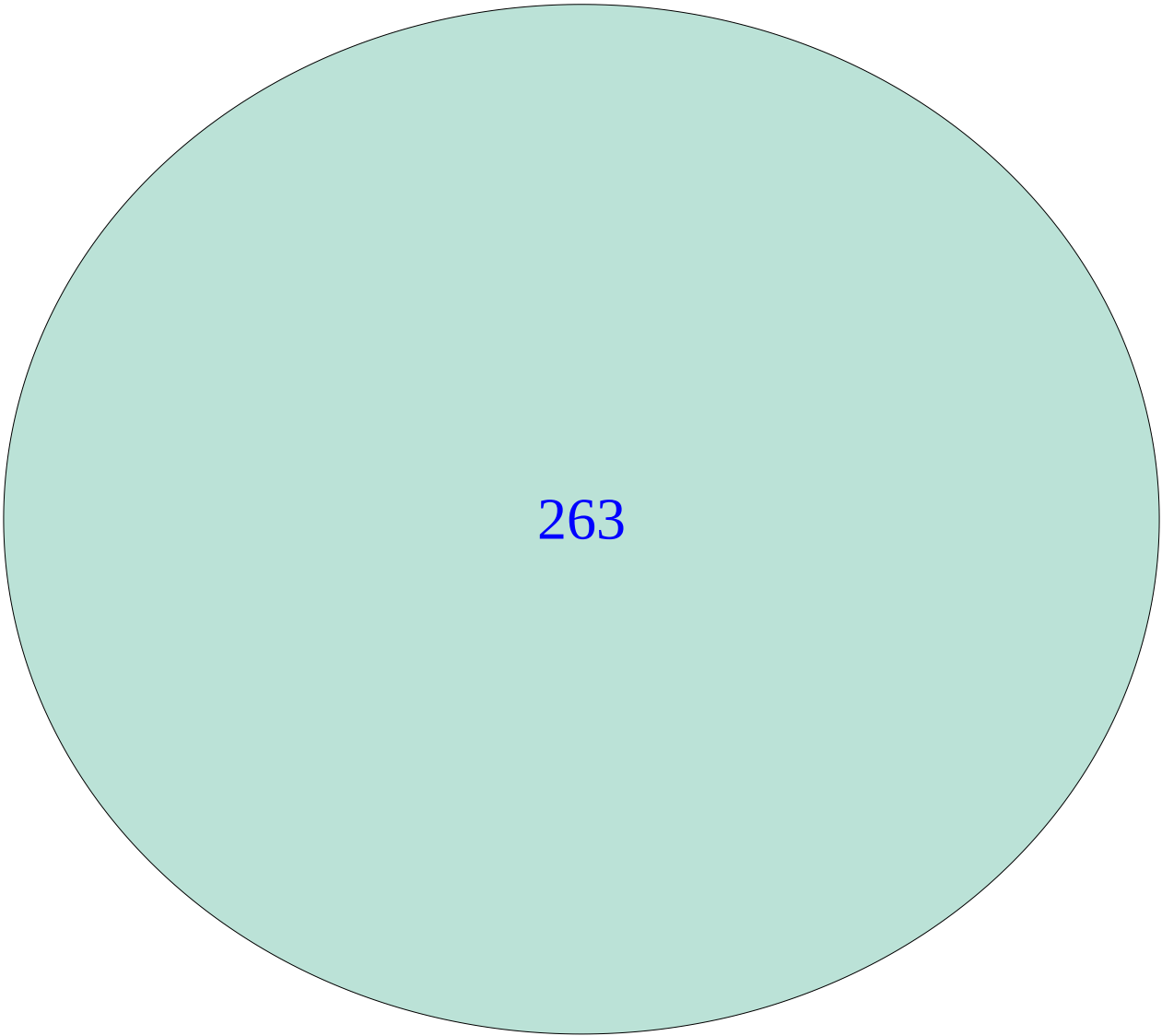
484

R2 >= 0.8

R2 < 0.8

Total

Panobinostat_1_vs_Vehicle_1



nobinostat_1_vs_Vehicle_1

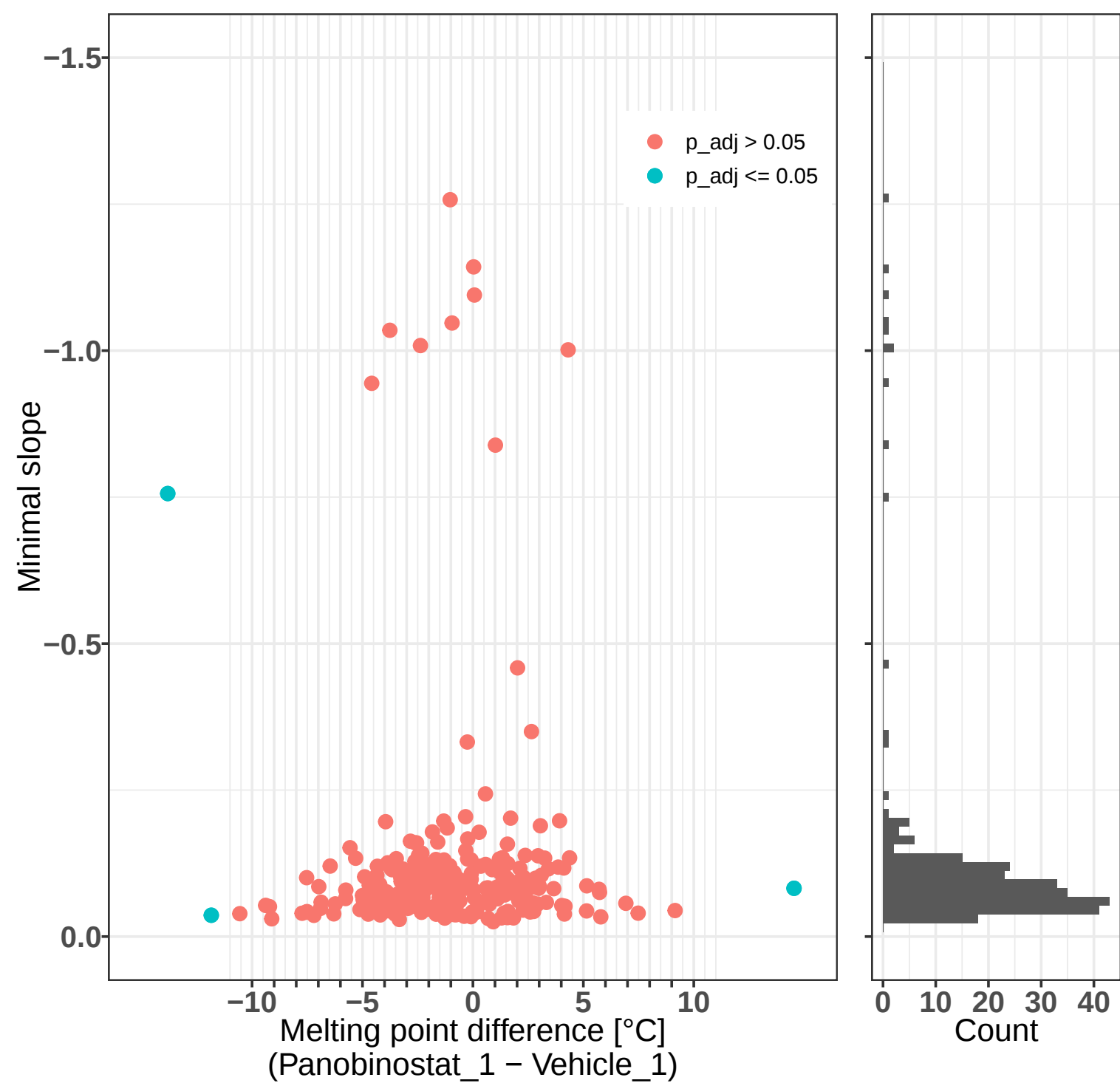
221

484

passed_filter

filtered_out

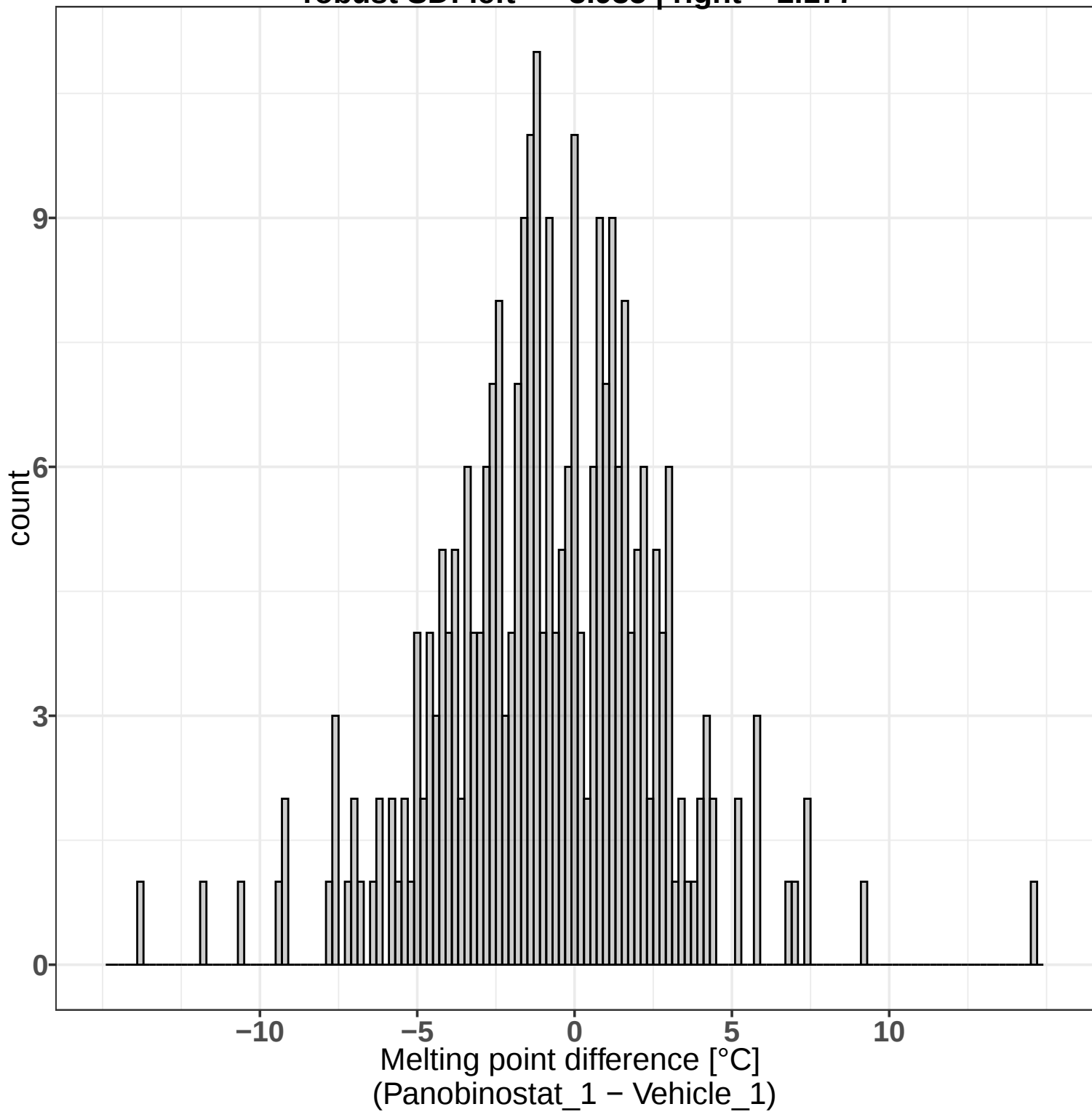
Total



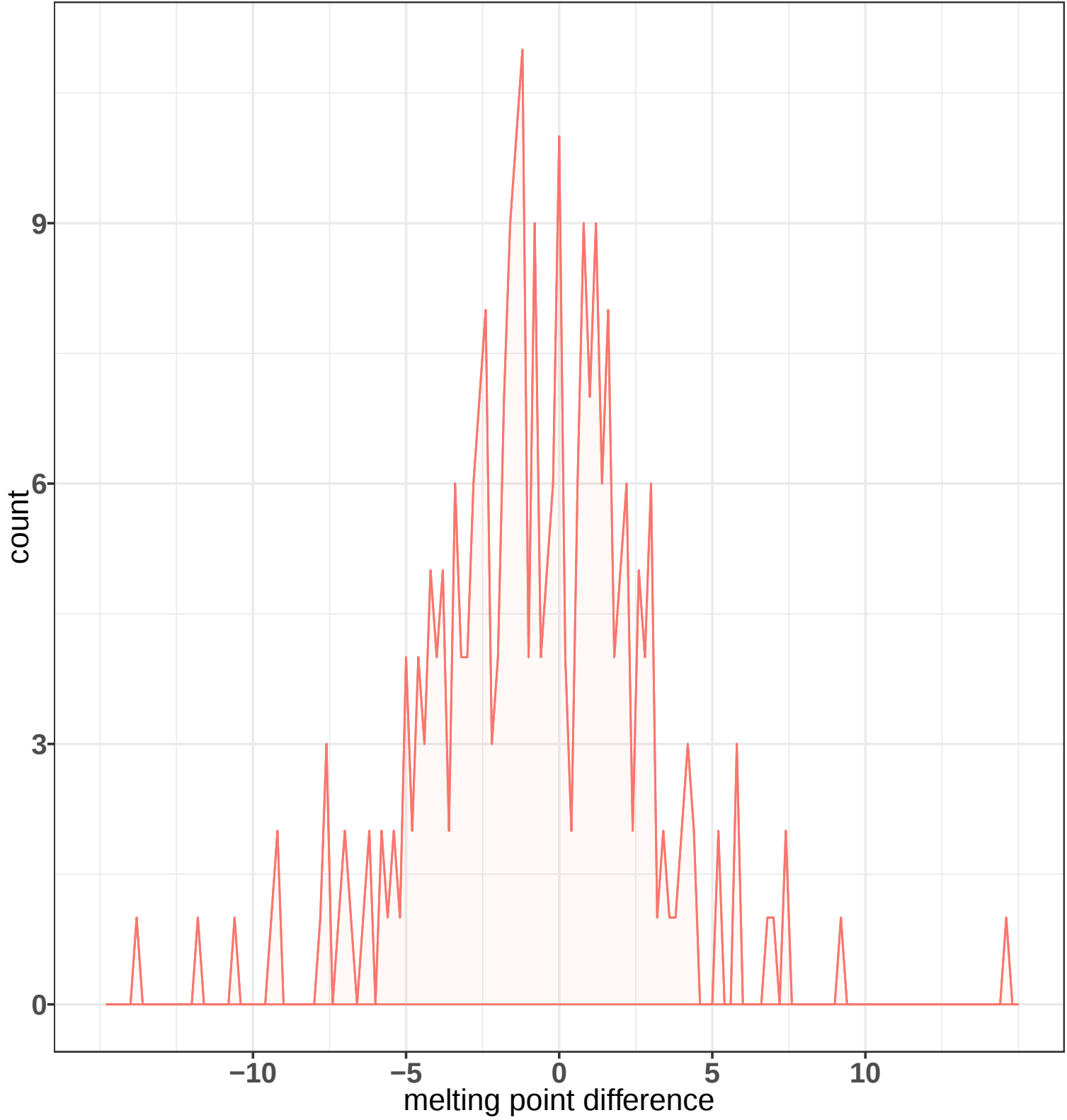
n = 263

Panobinostat_1 vs. Vehicle_1

**median = -0.858 ,
robust SD: left = -3.935 | right = 2.177**



n = 294 , R_sq >= 0.8



comparison Panobinostat_1 - Vehicle_1