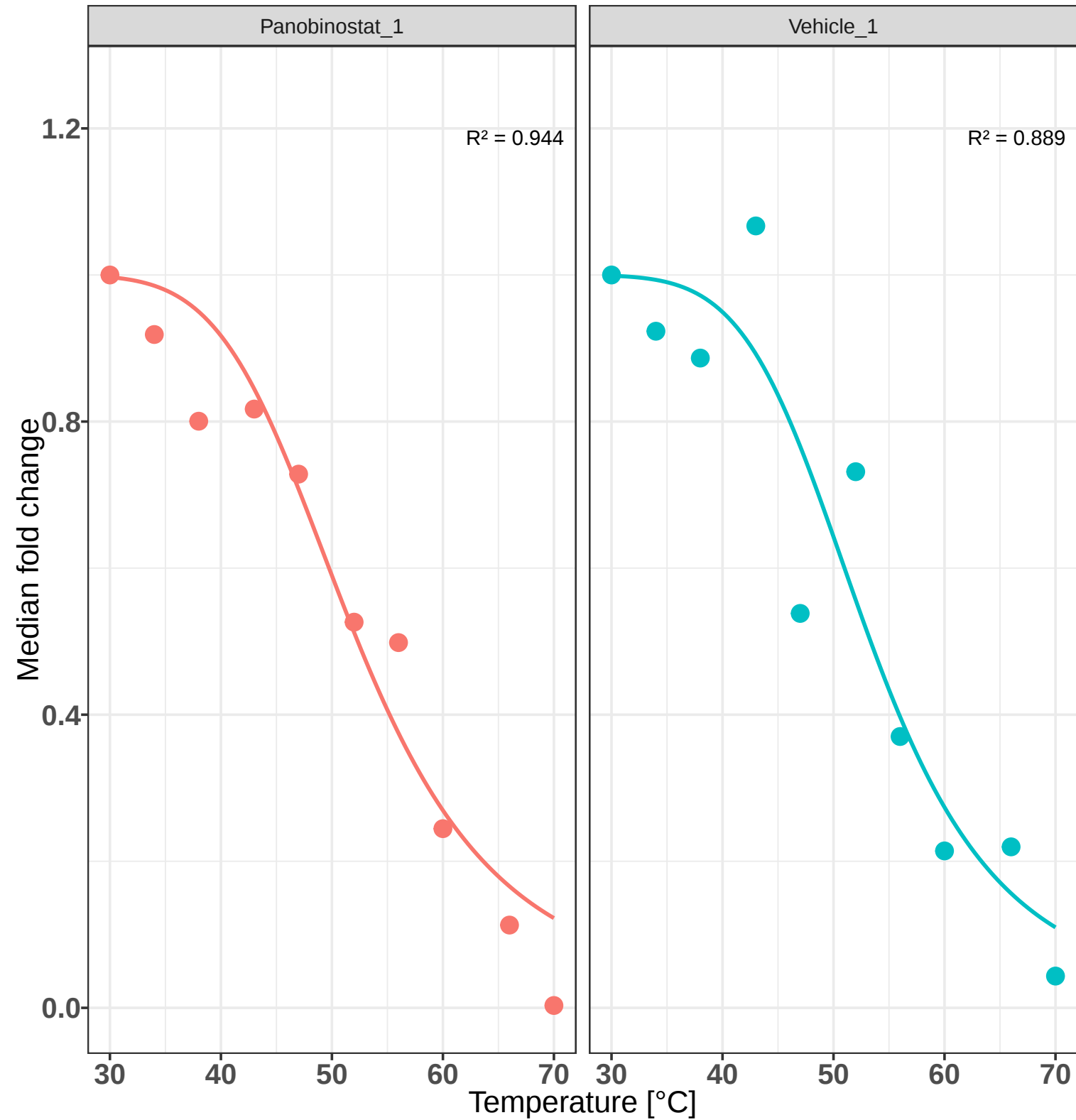
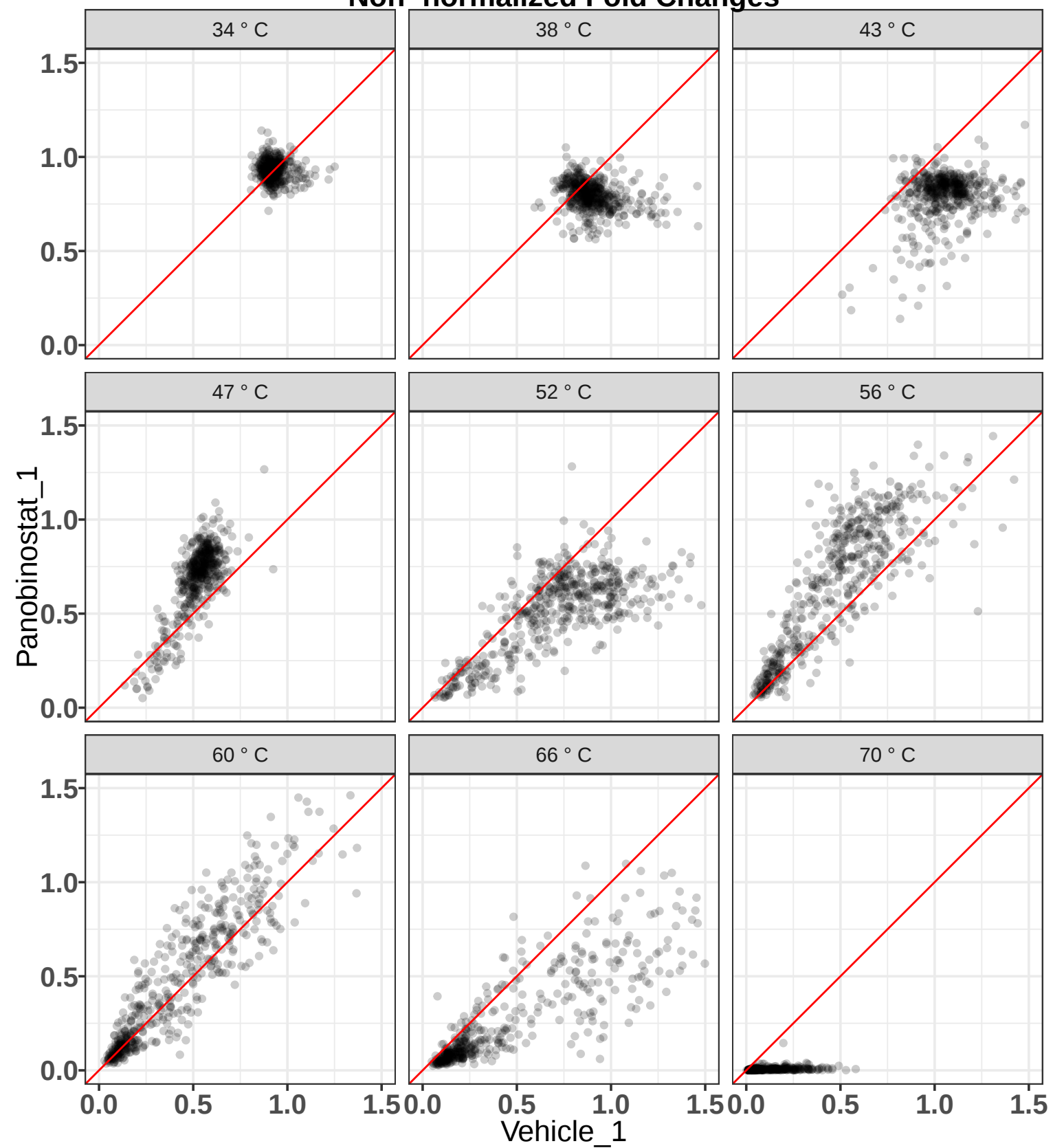


Normalization curves

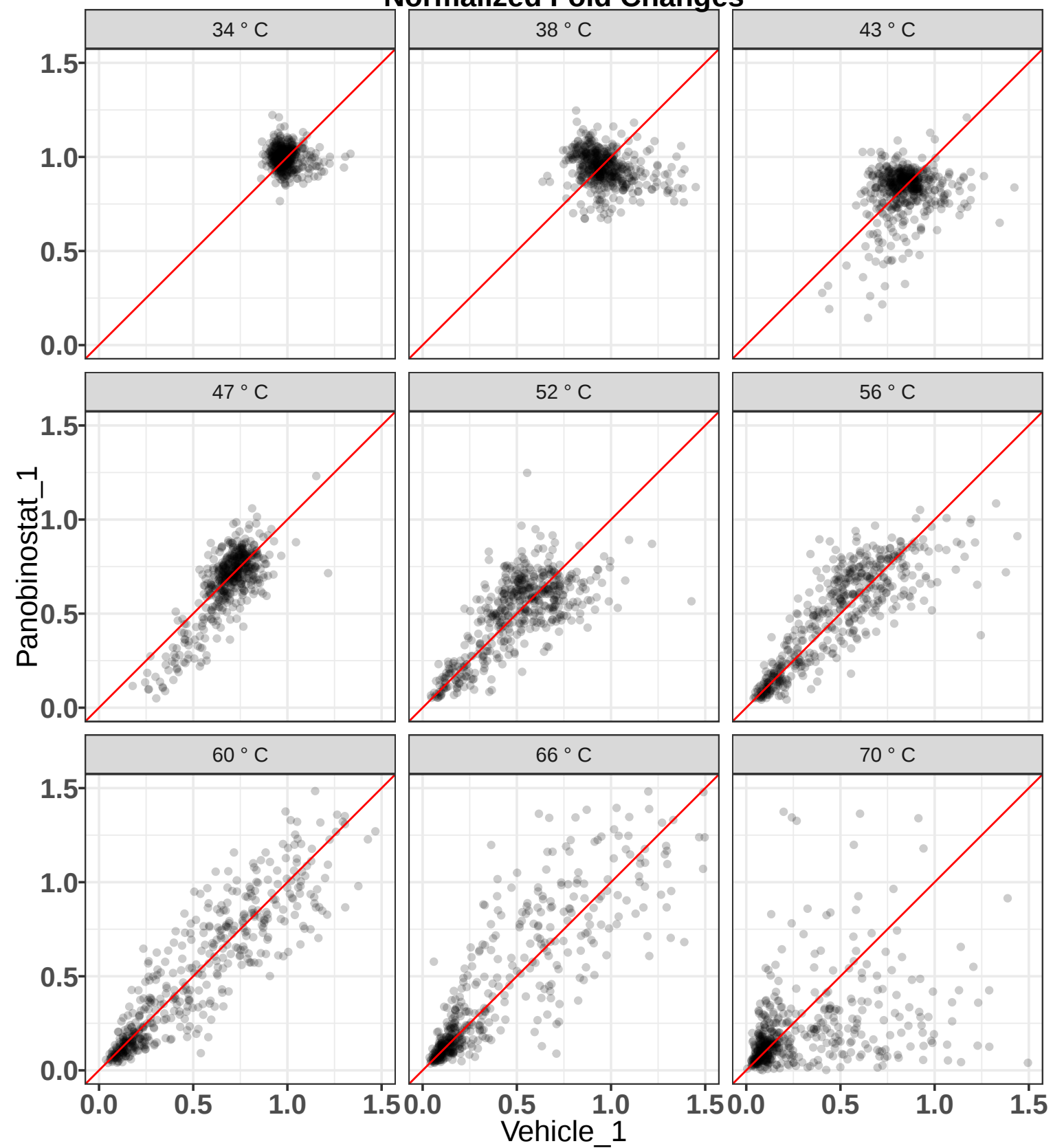
based on 45 proteins



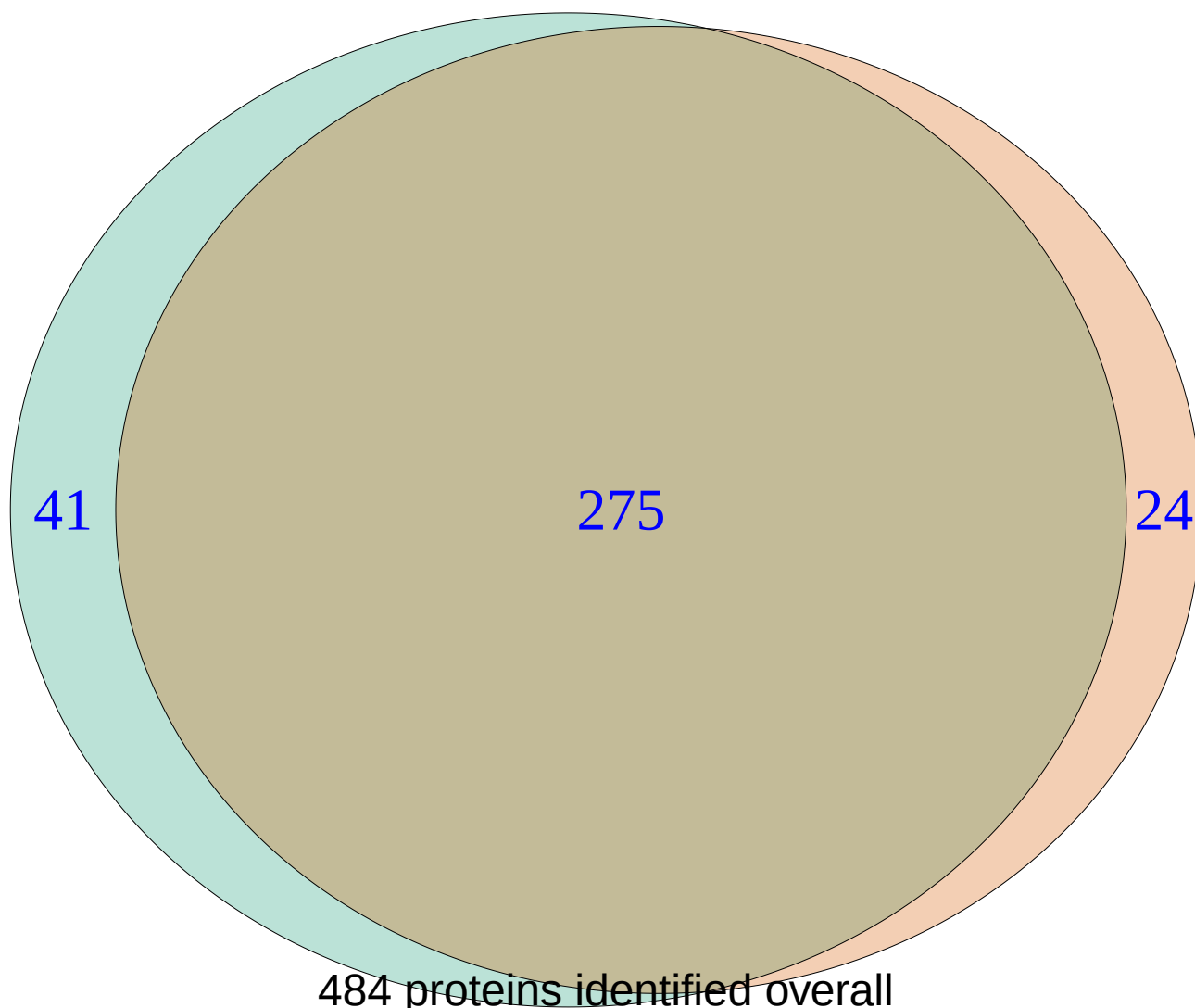
Non-normalized Fold Changes



Normalized Fold Changes



Panobinostat_1
Vehicle_1



Panobinostat_1

Vehicle_1

Intersection

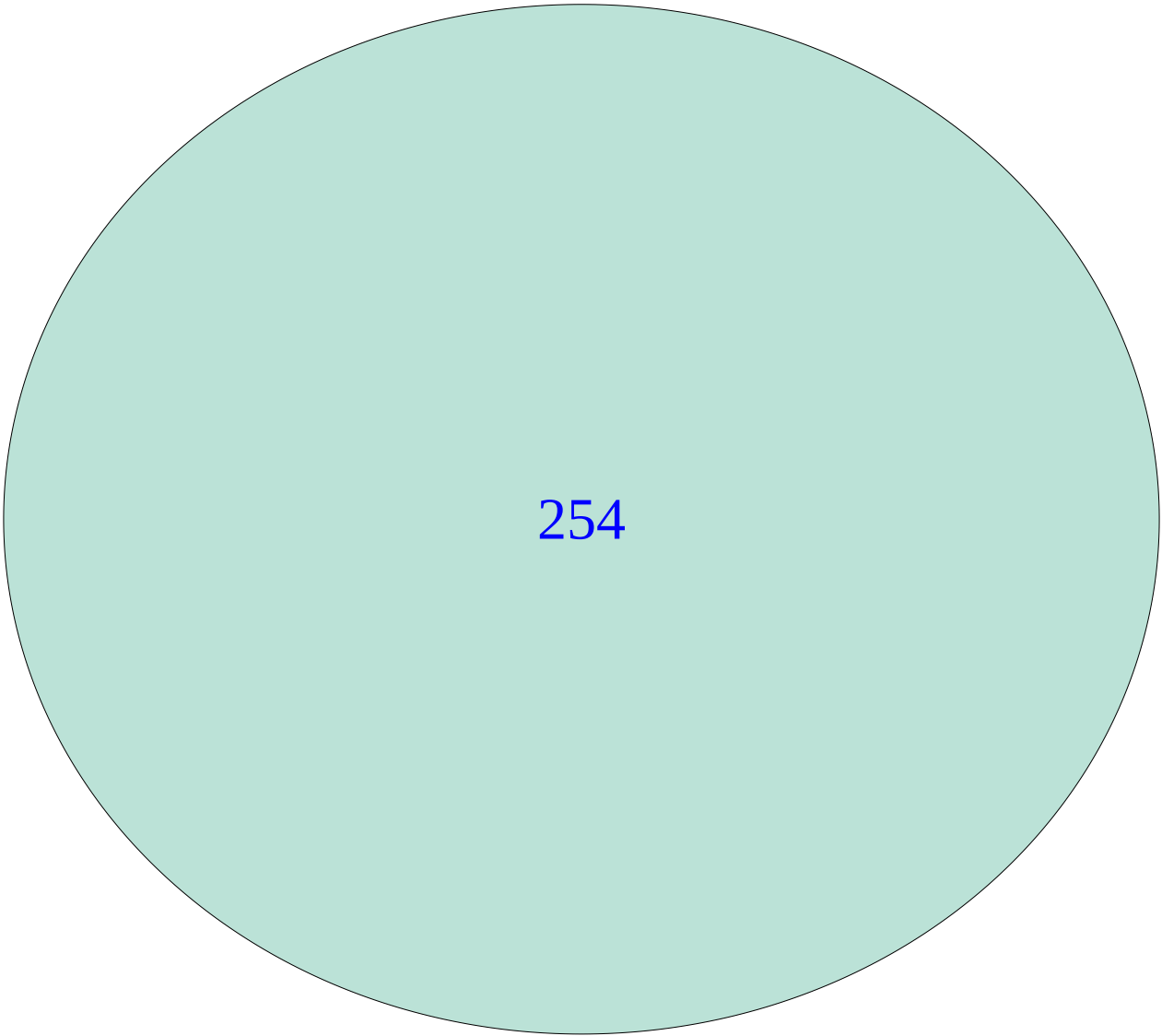
Total

R2 >= 0.8

R2 < 0.8

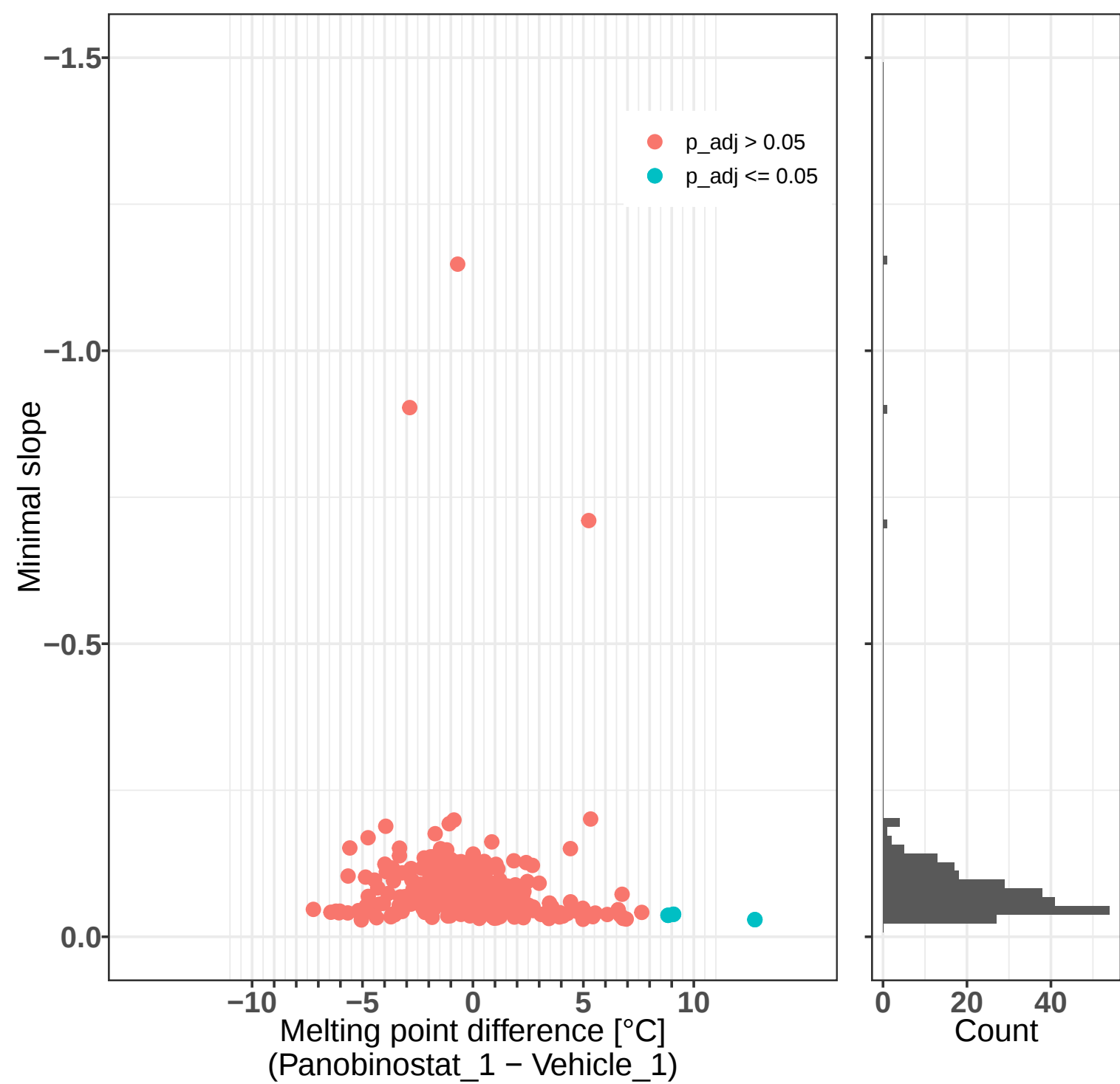
Total

Panobinostat_1_vs_Vehicle_1



254

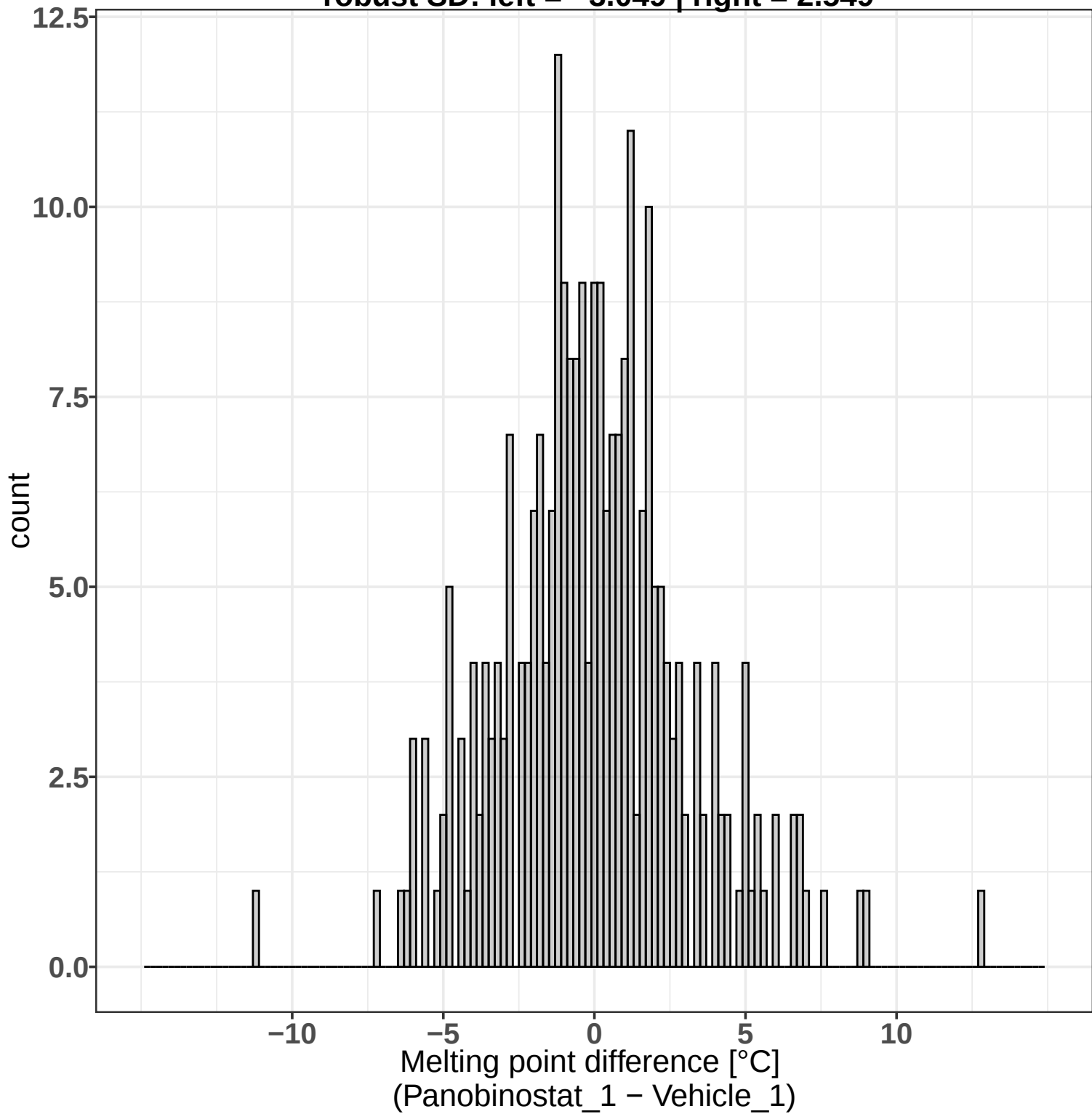
nobinostat_1_vs_Vehicle_1	254	230	484
passed_filter	filtered_out	Total	



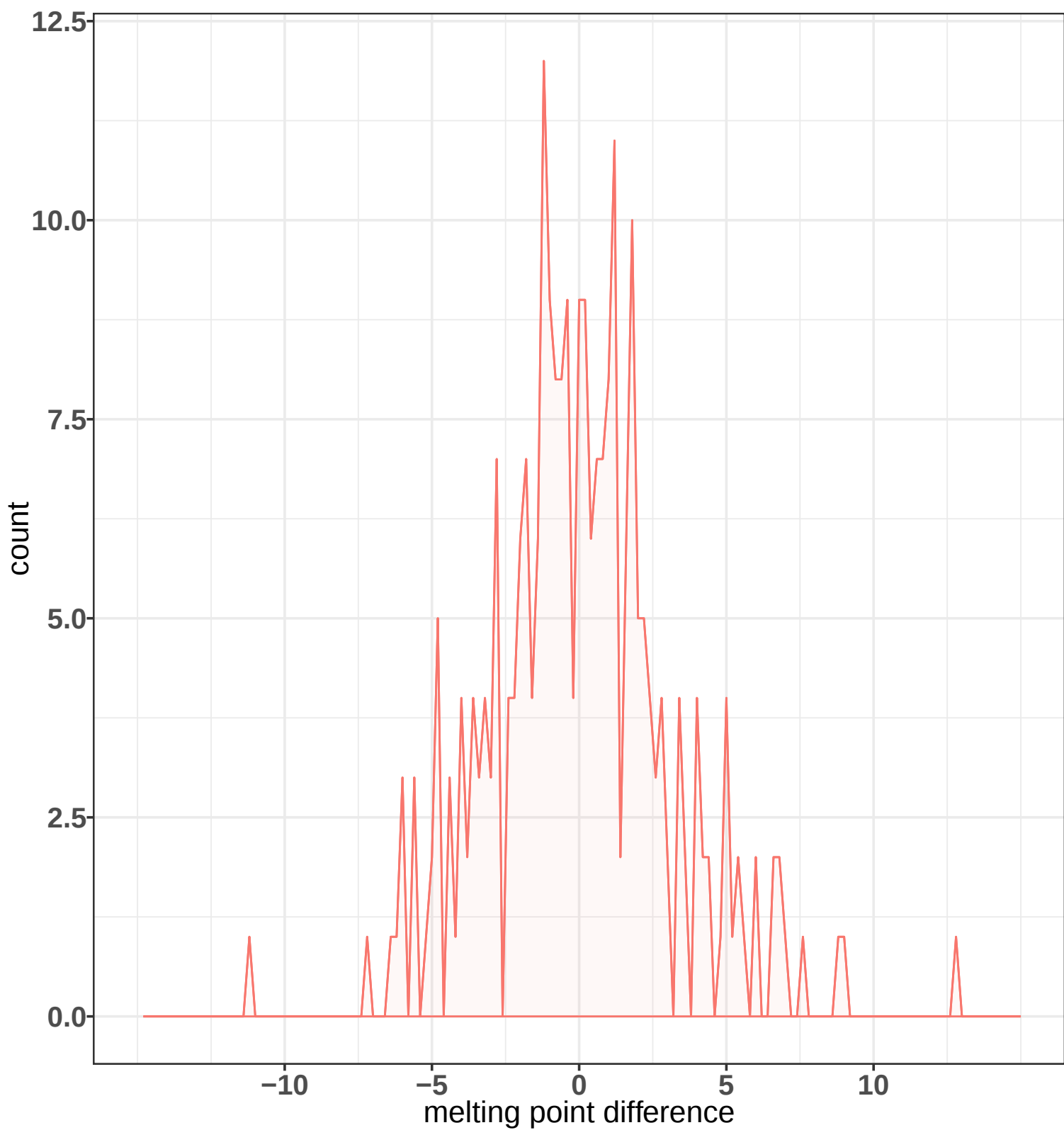
n = 252

Panobinostat_1 vs. Vehicle_1

median = -0.094 ,
robust SD: left = -3.049 | right = 2.549



n = 275 , R_sq >= 0.8



comparison ▬ Panobinostat_1 - Vehicle_1