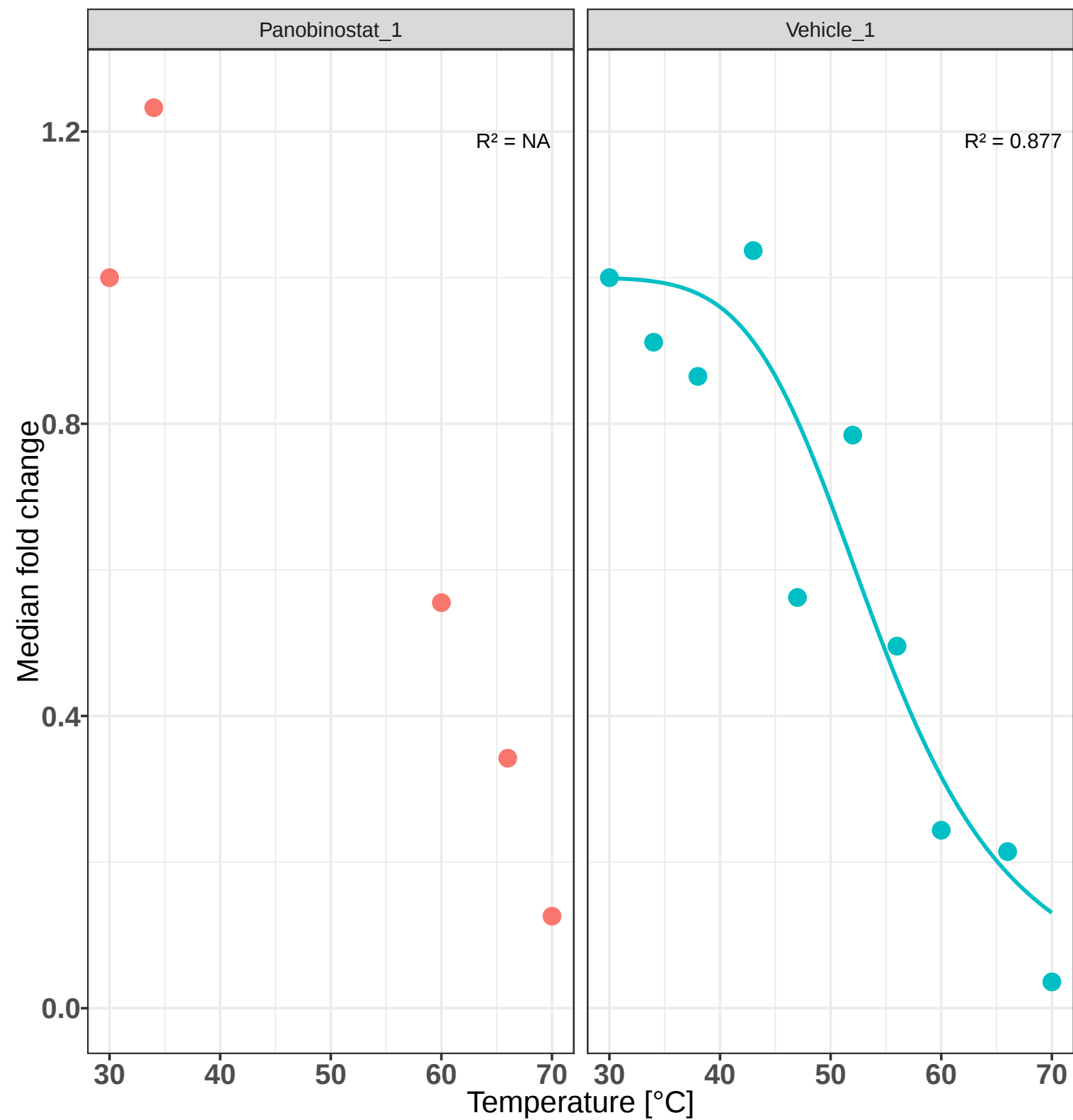
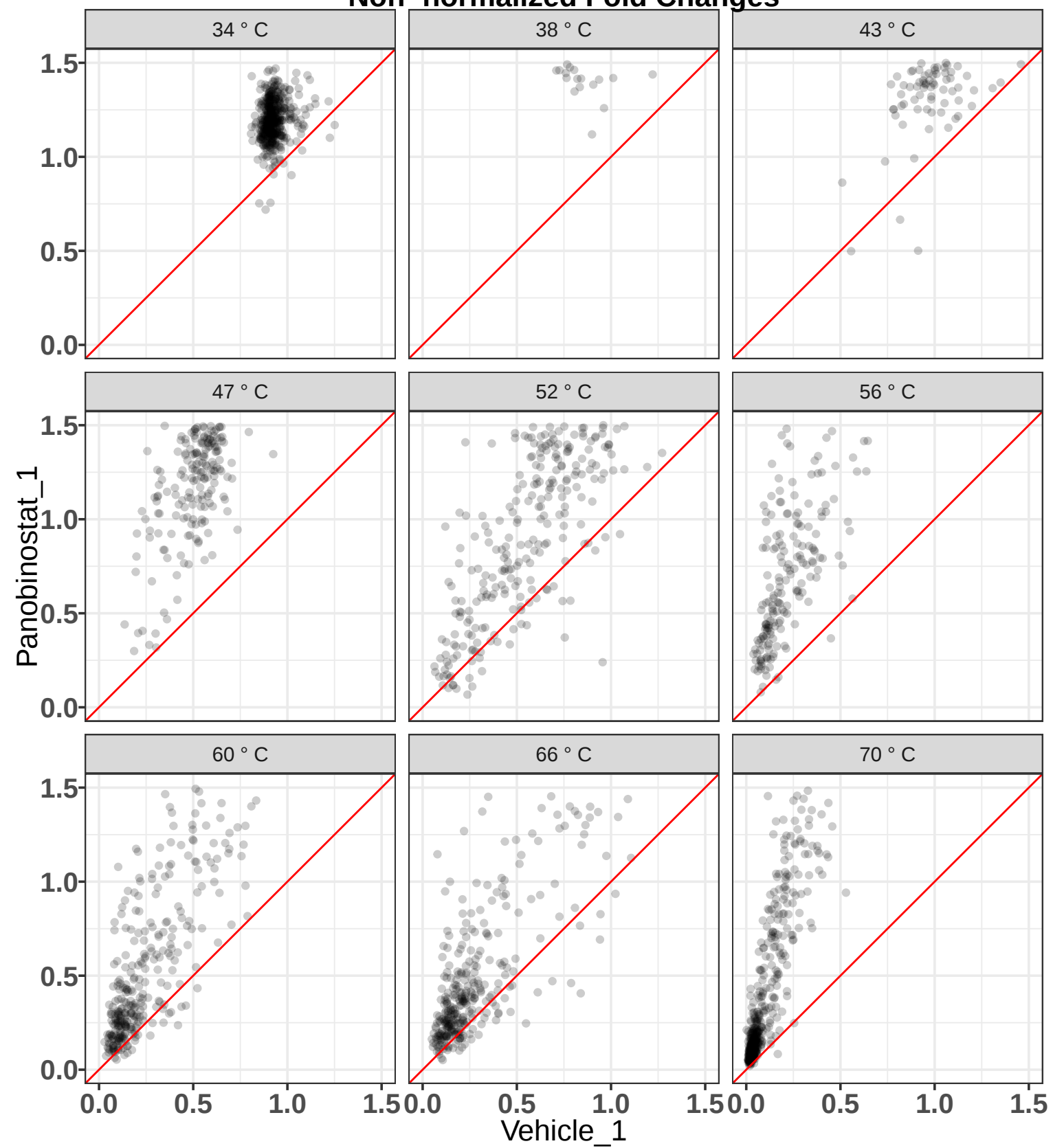


Normalization curves

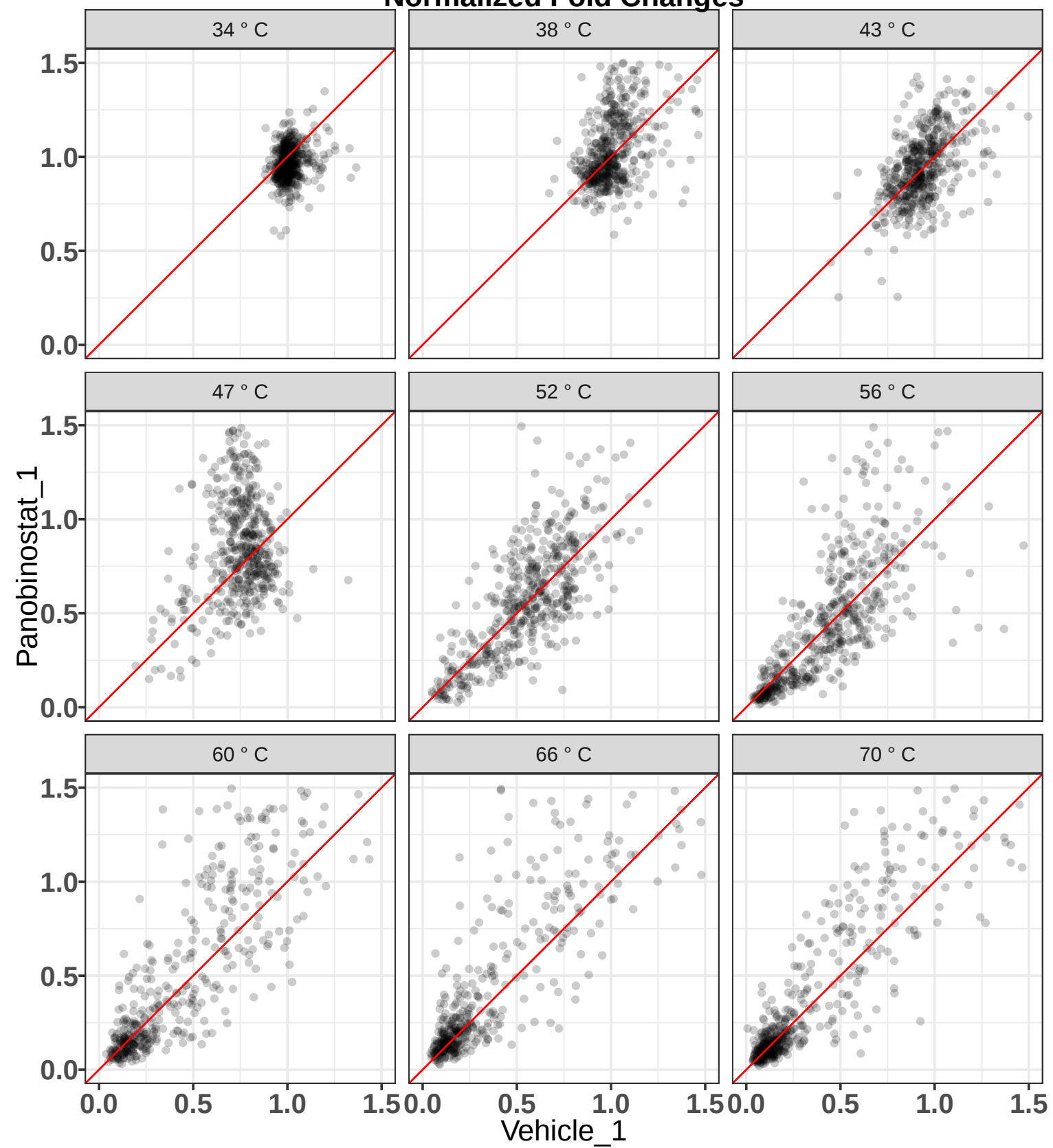
based on 37 proteins



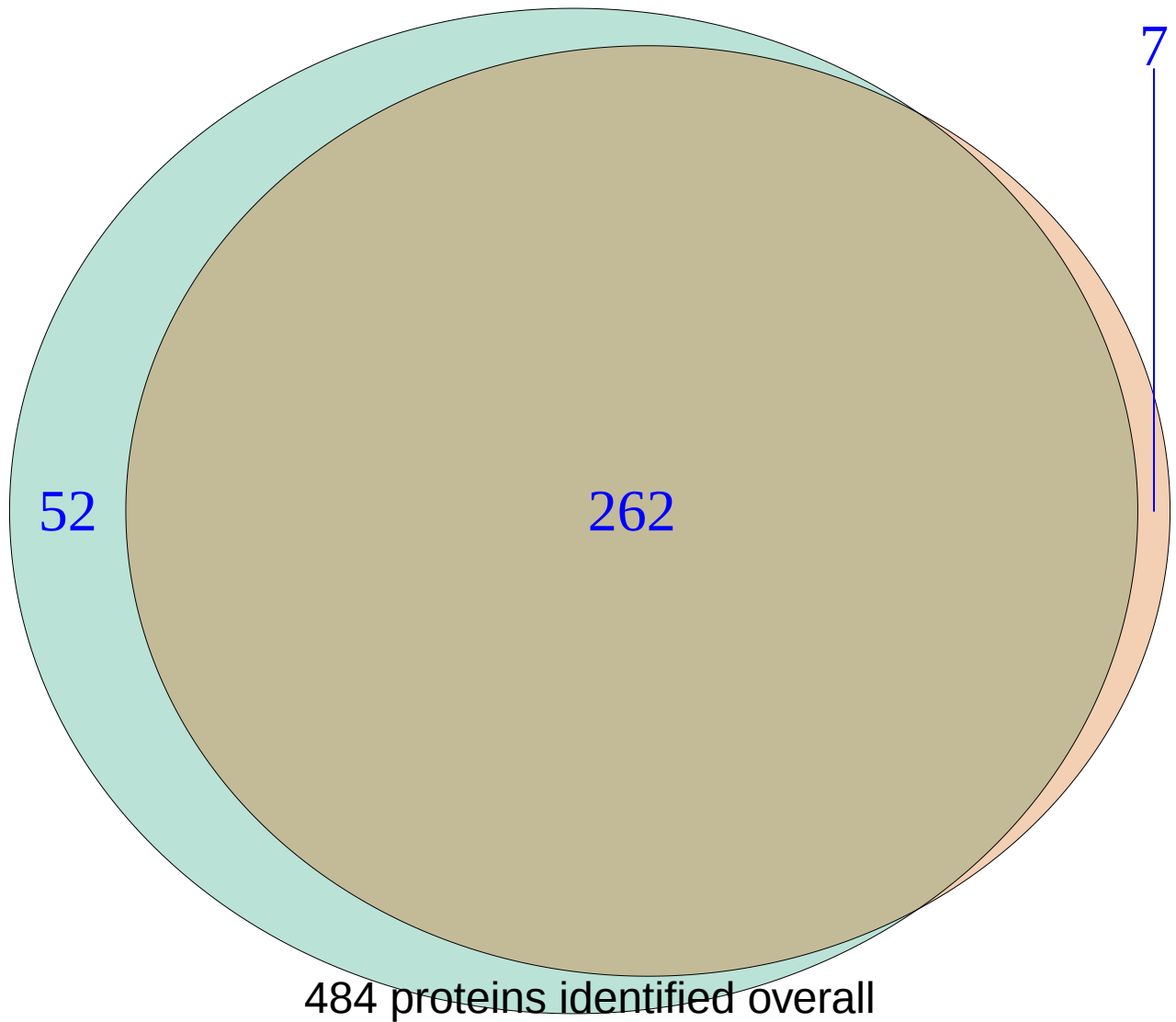
Non-normalized Fold Changes



Normalized Fold Changes



Panobinostat_1
Vehicle_1



Panobinostat_1

Vehicle_1

Intersection

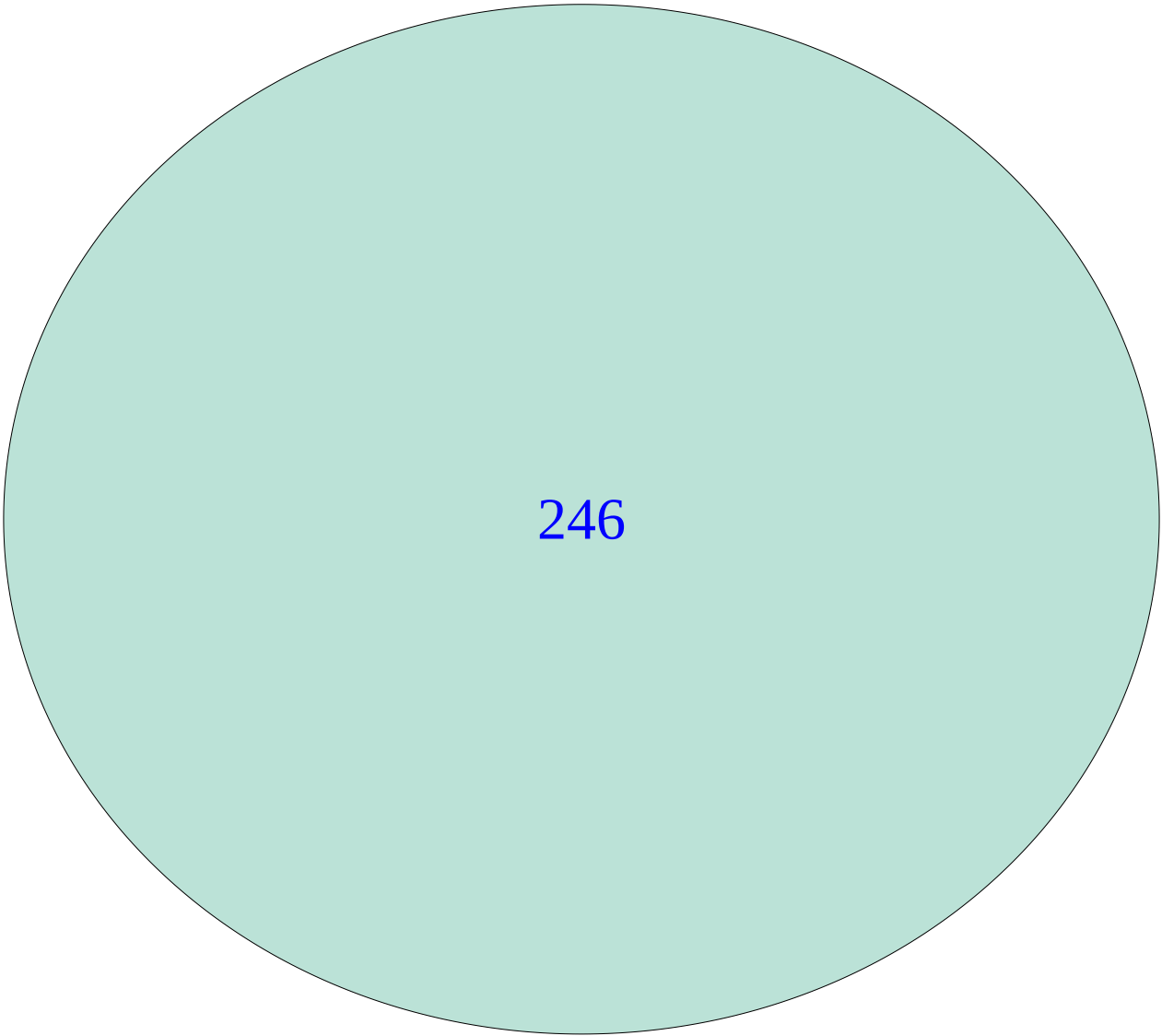
Total

R2 >= 0.8

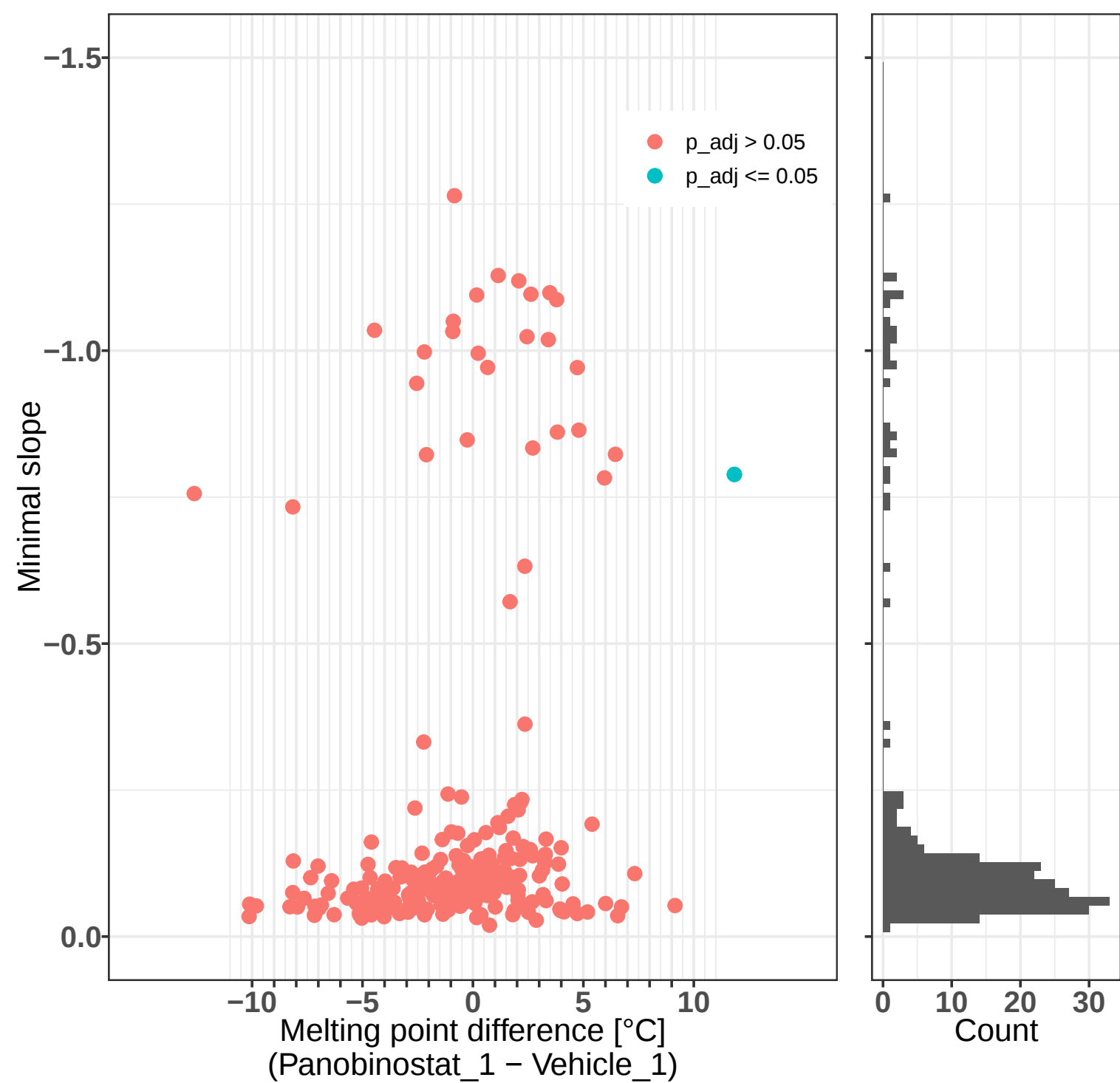
R2 < 0.8

Total

Panobinostat_1_vs_Vehicle_1



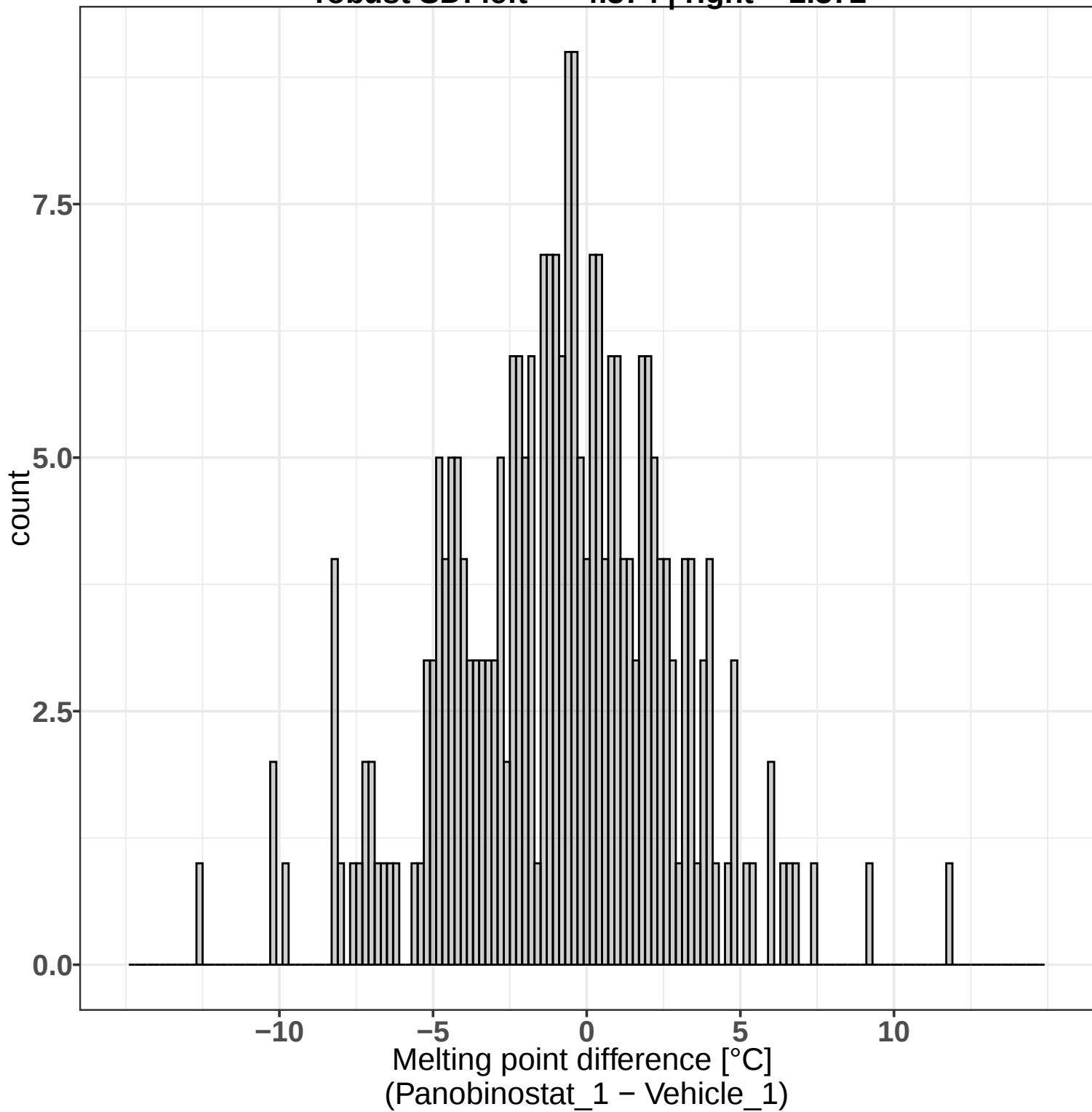
nobinostat_1_vs_Vehicle_1	246	238	484
	passed_filter	filtered_out	Total



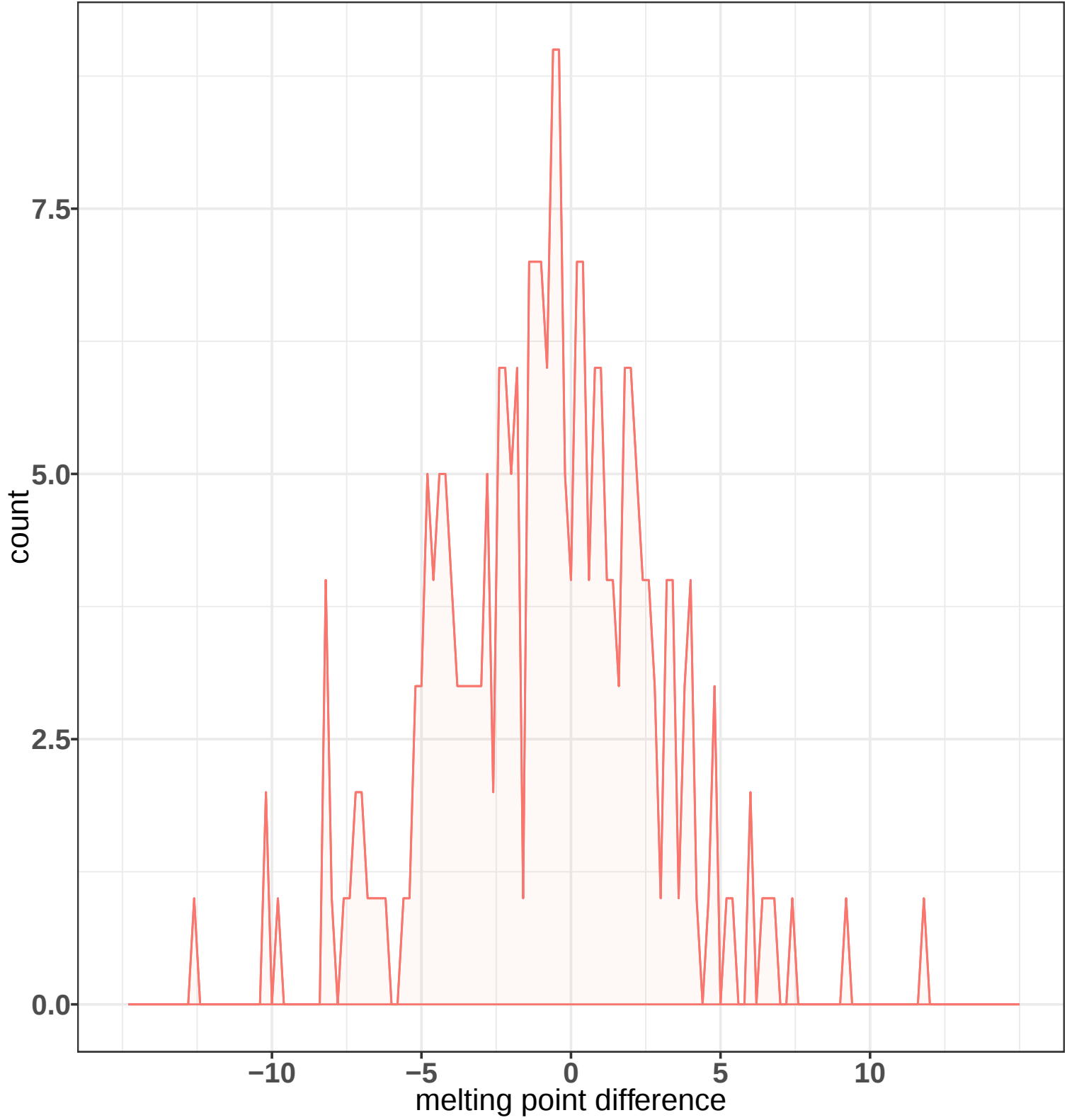
n = 245

Panobinostat_1 vs. Vehicle_1

median = -0.686 ,
robust SD: left = -4.374 | right = 2.372



n = 262 , R_sq >= 0.8



comparison Panobinostat_1 - Vehicle_1