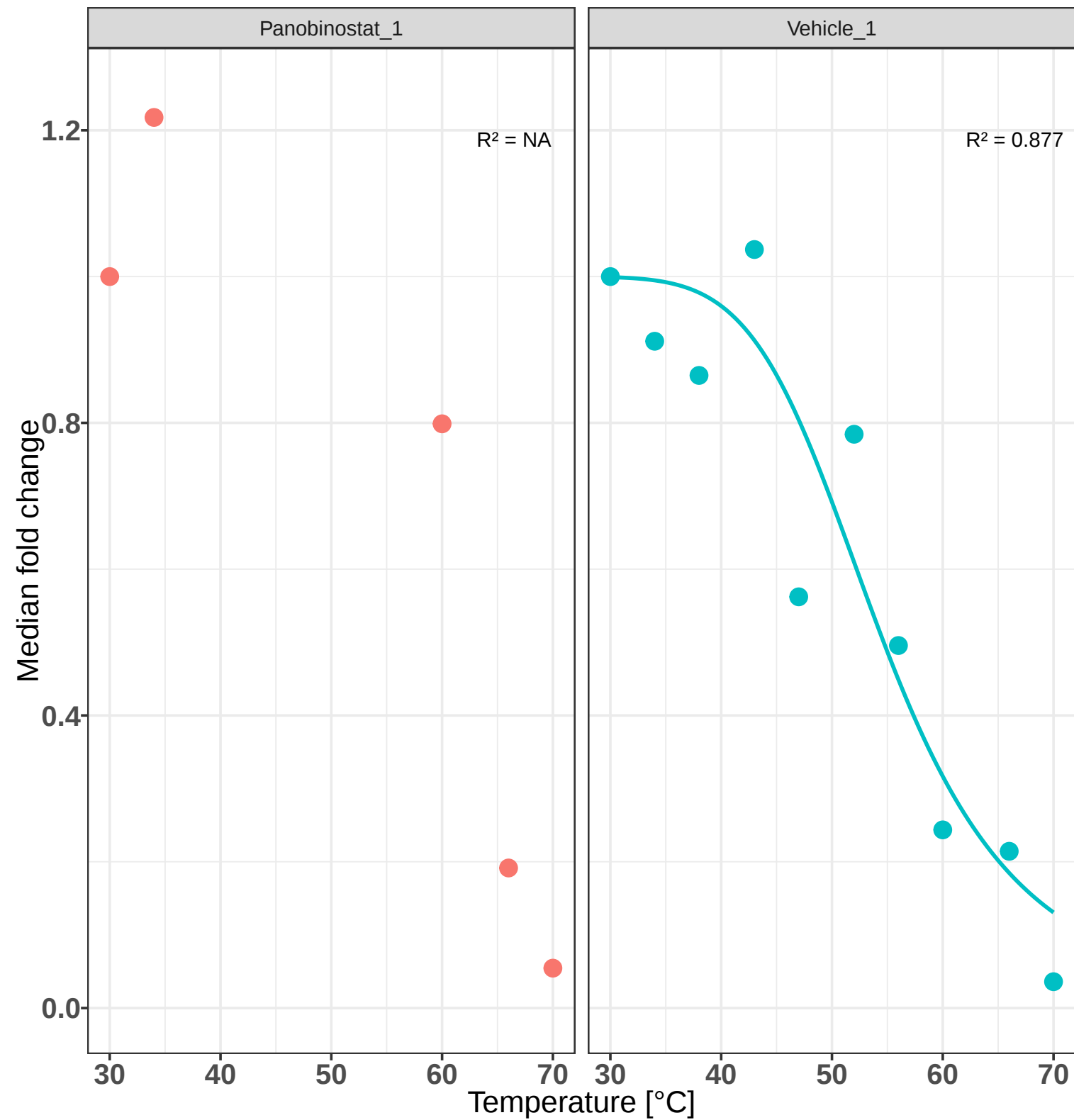
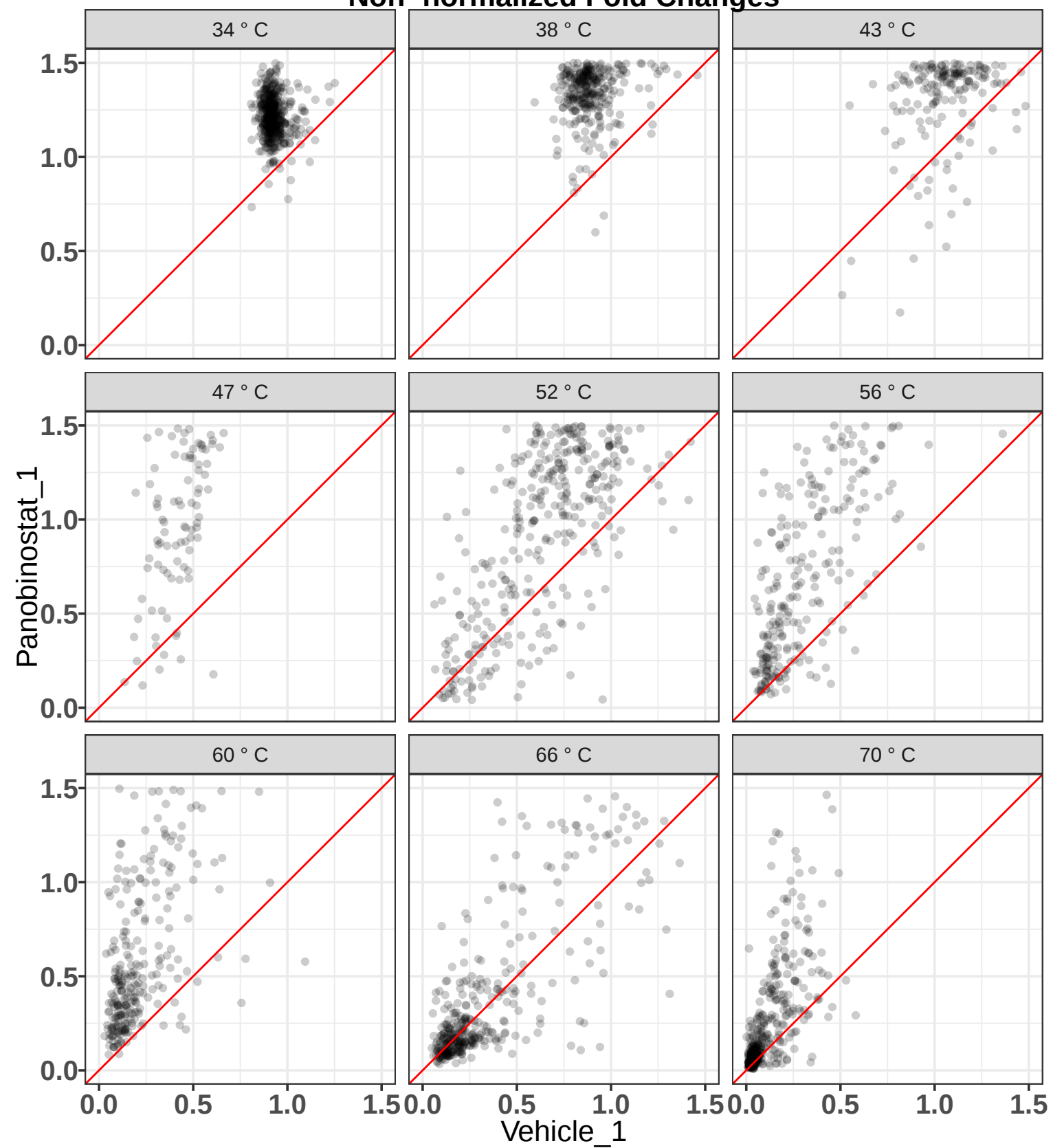


Normalization curves

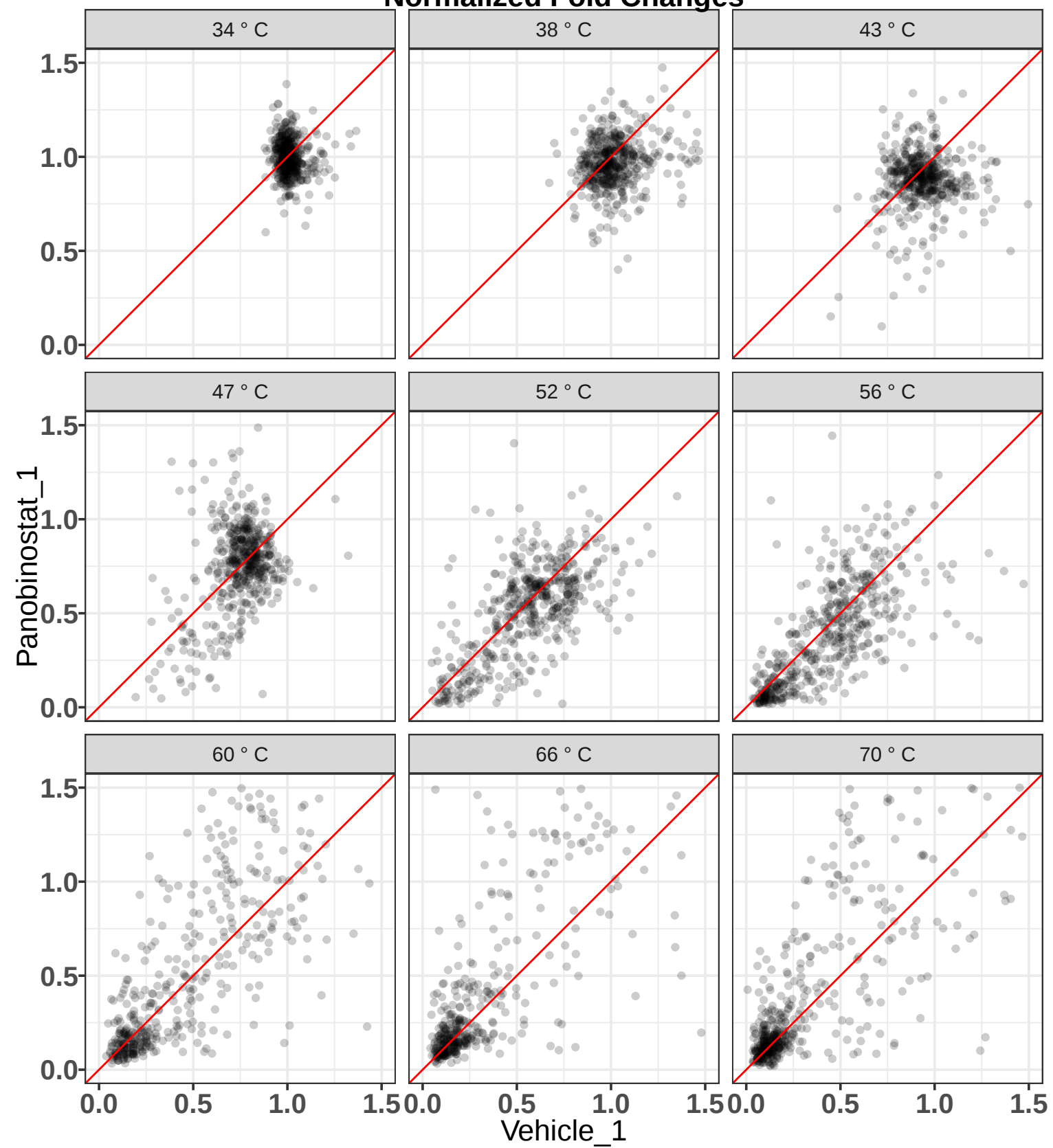
based on 37 proteins



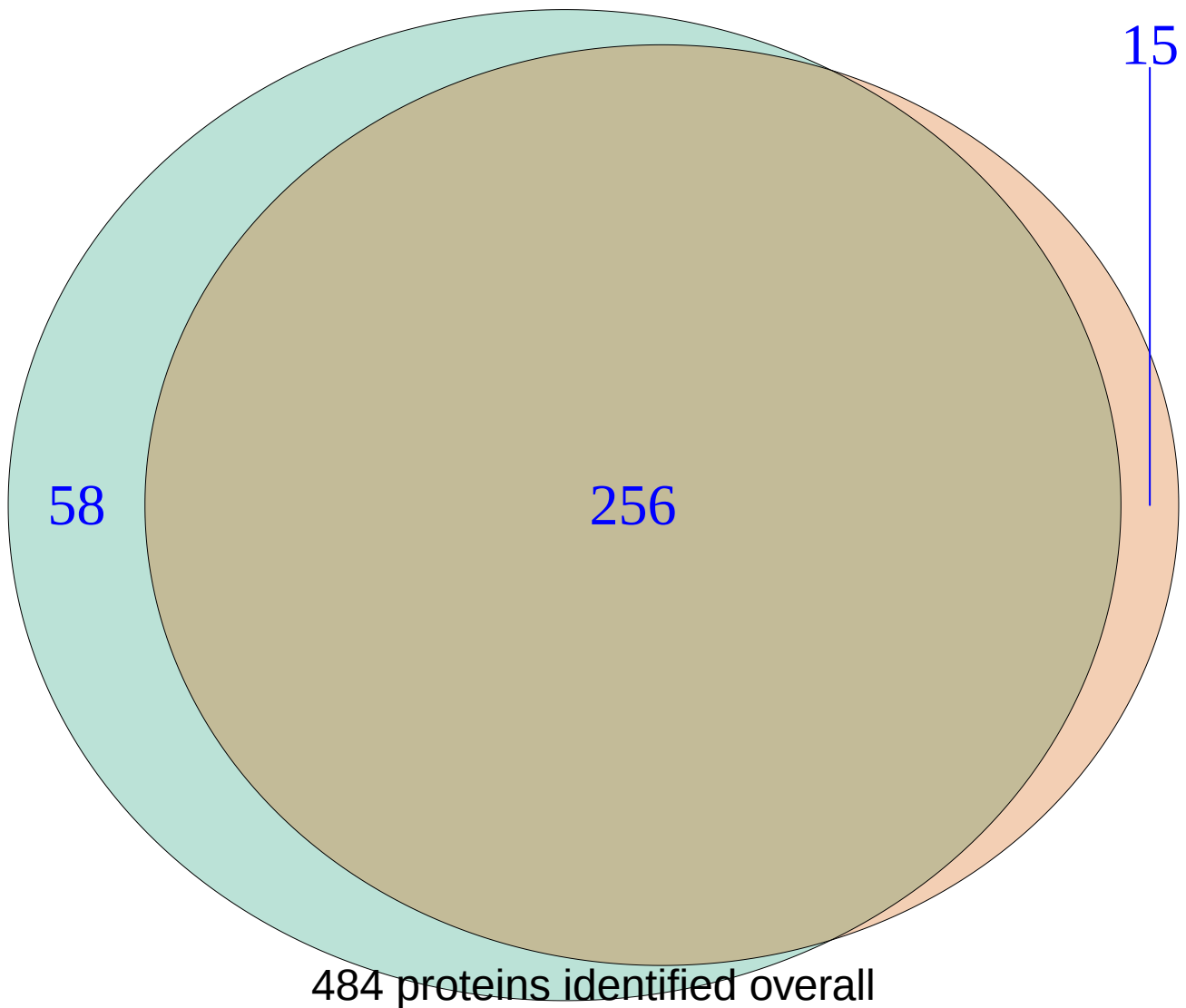
Non-normalized Fold Changes



Normalized Fold Changes



Panobinostat_1
Vehicle_1



Panobinostat_1

Vehicle_1

256

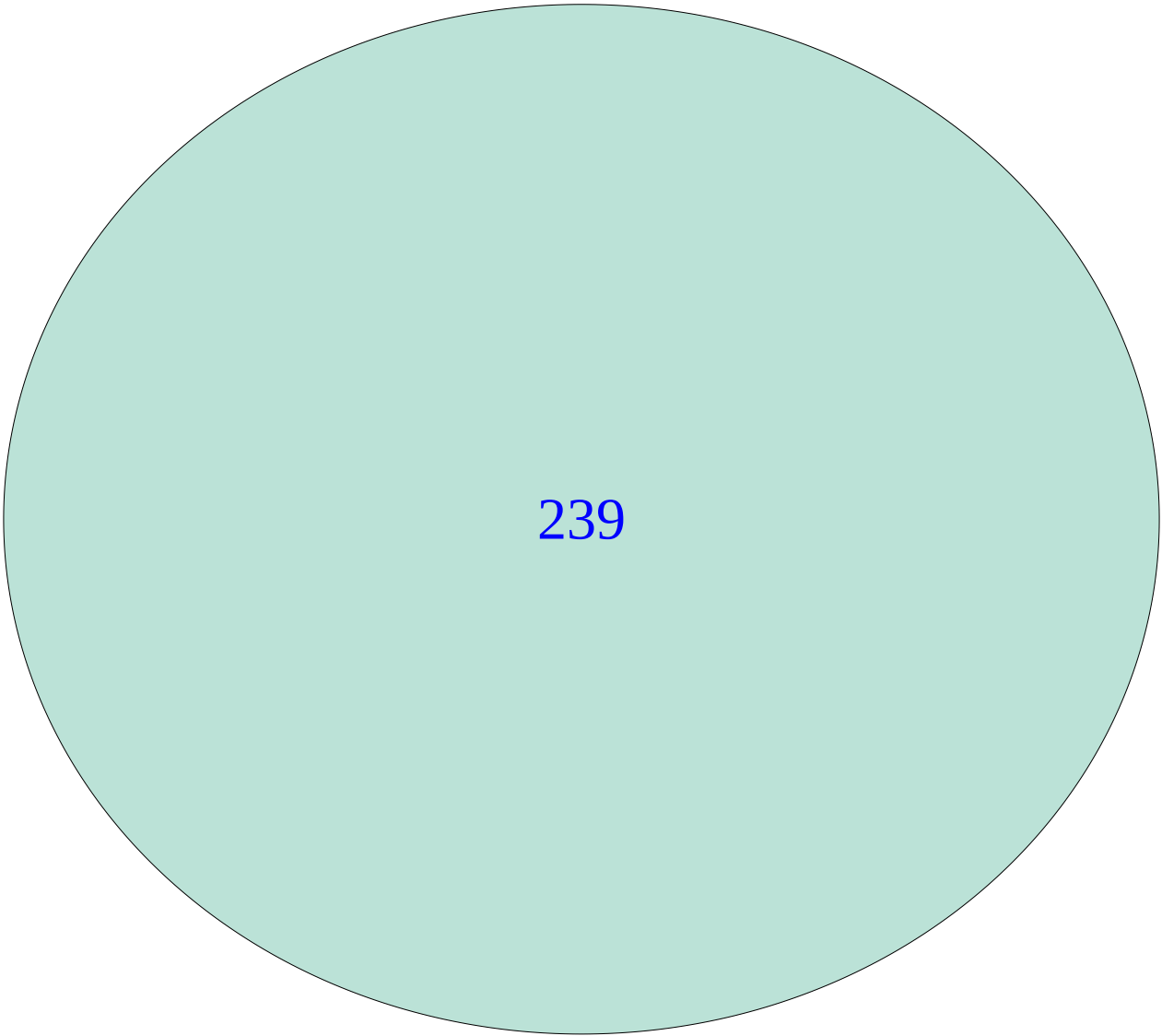
484

R2 >= 0.8

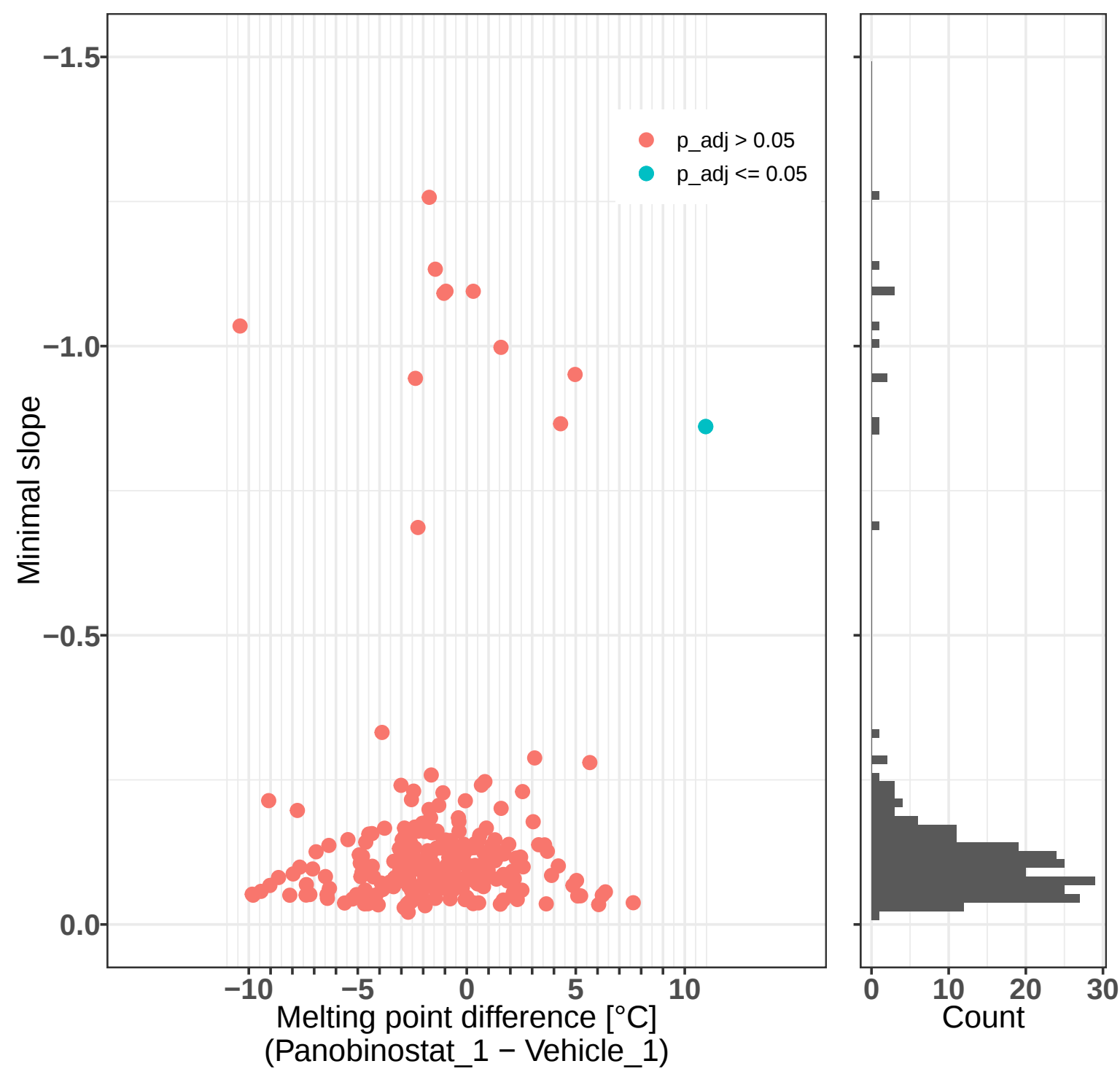
R2 < 0.8

Total

Panobinostat_1_vs_Vehicle_1



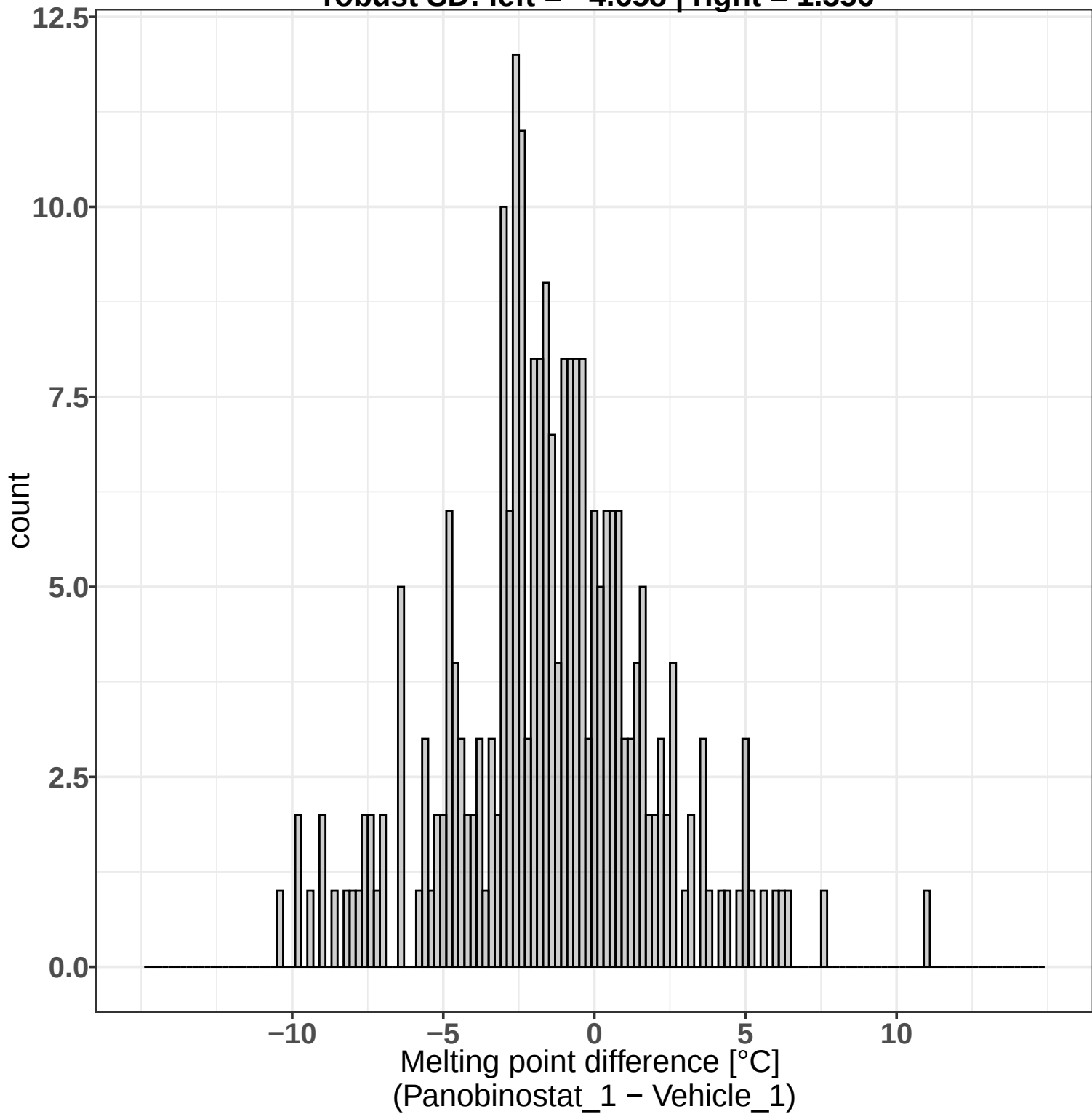
nobinostat_1_vs_Vehicle_1	239	245	484
	passed_filter	filtered_out	Total



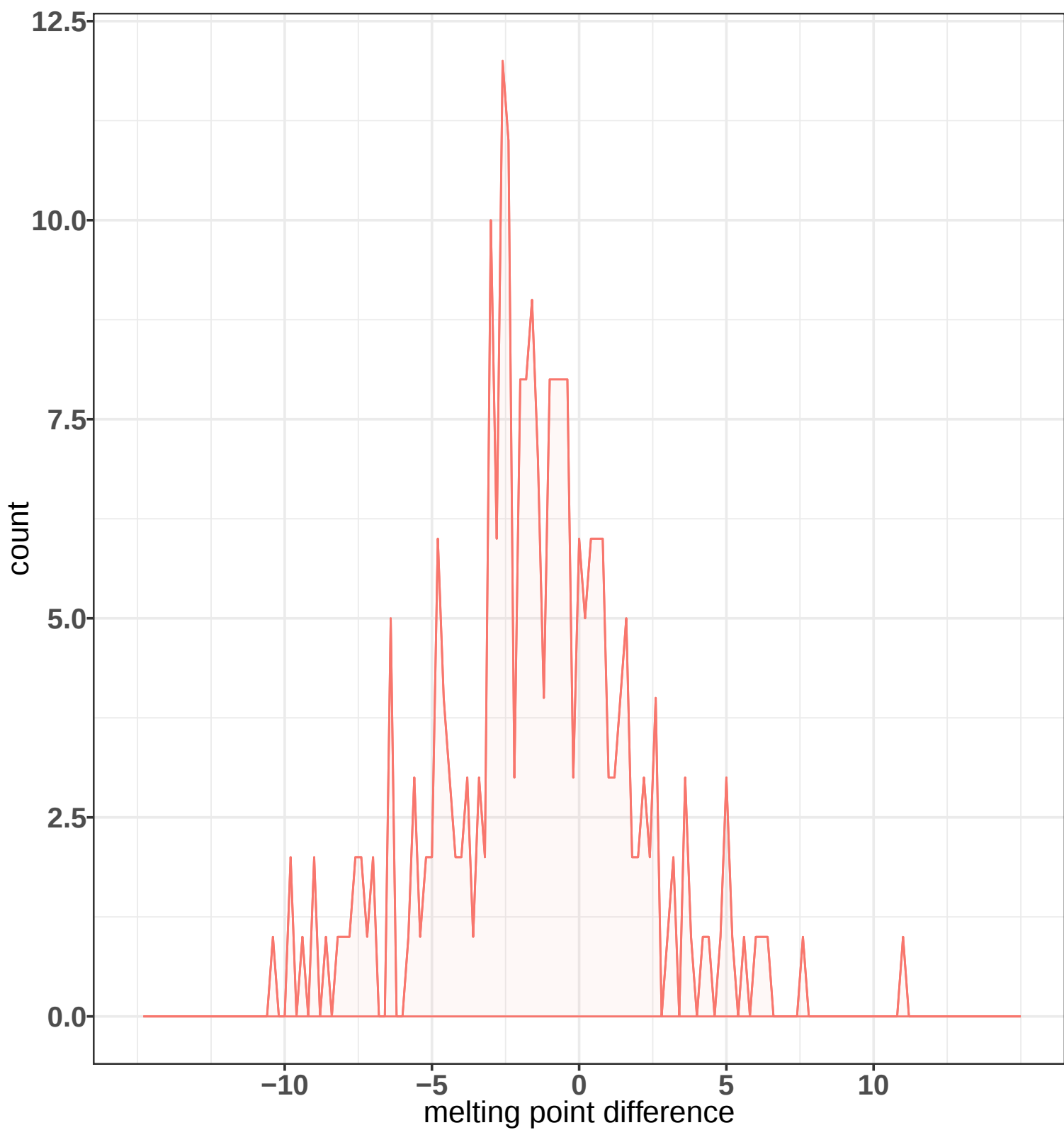
n = 239

Panobinostat_1 vs. Vehicle_1

**median = -1.563 ,
robust SD: left = -4.658 | right = 1.356**



n = 256 , R_sq >= 0.8



comparison Panobinostat_1 - Vehicle_1