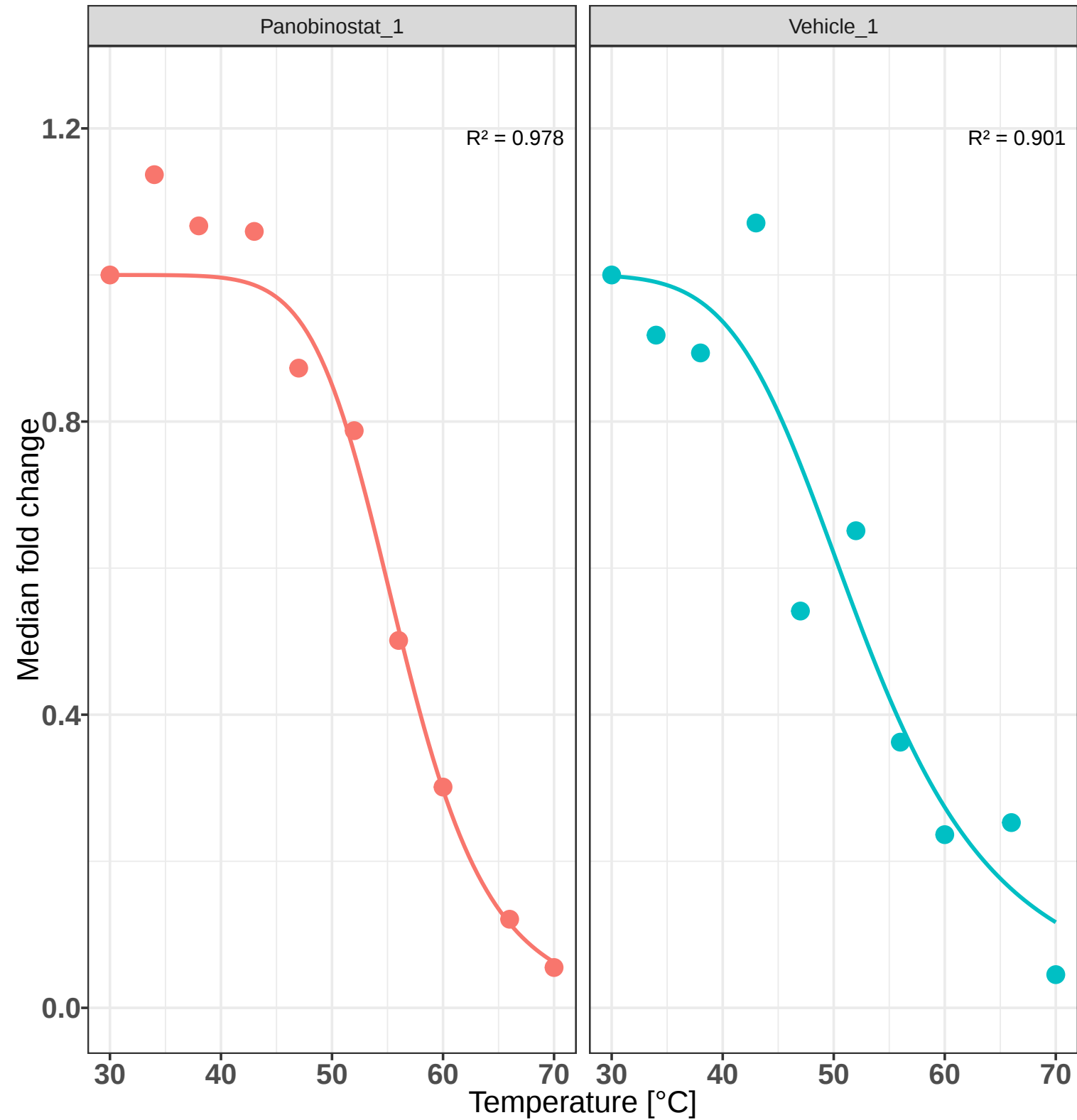
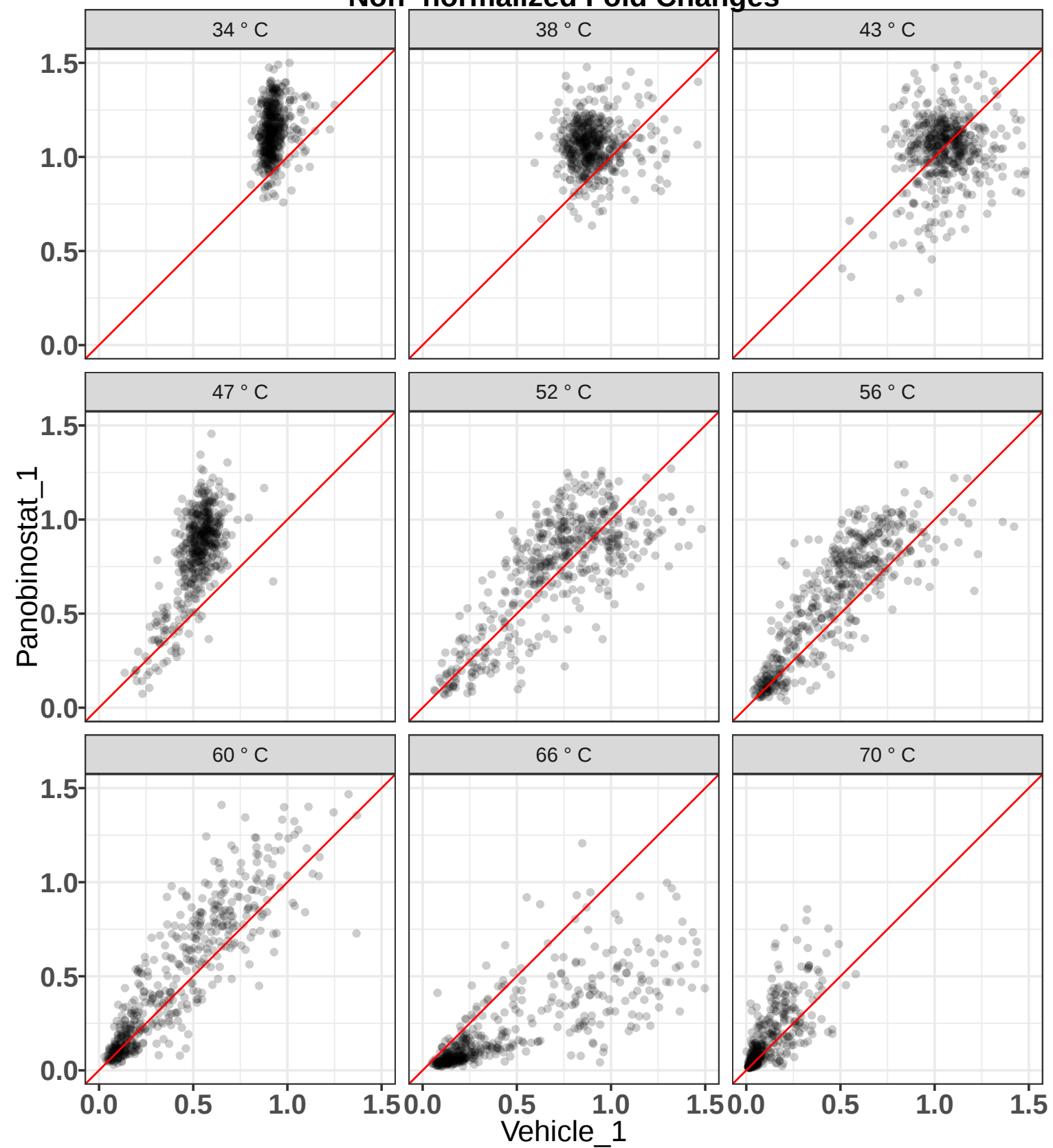


Normalization curves

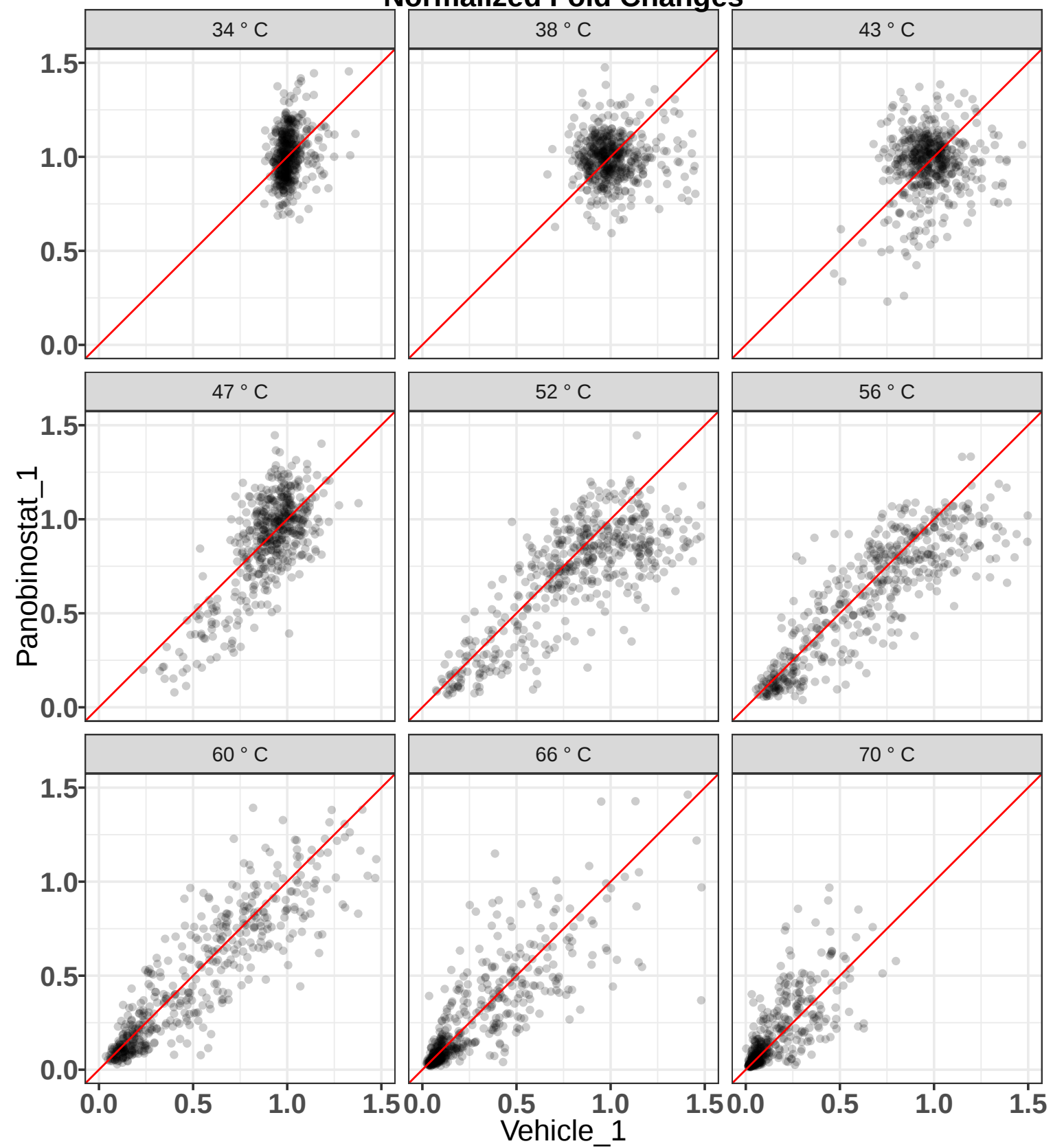
based on 62 proteins



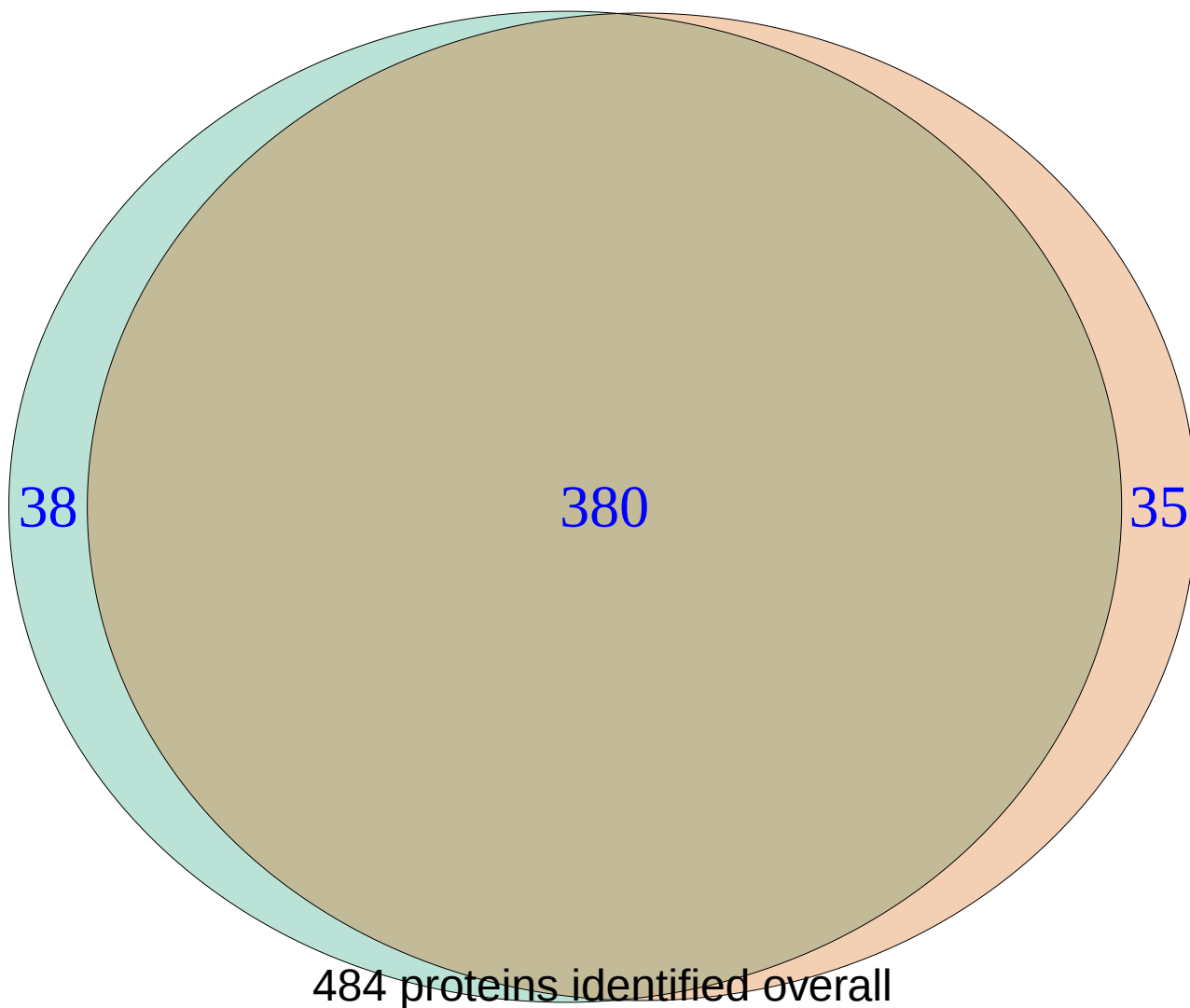
Non-normalized Fold Changes



Normalized Fold Changes



Panobinostat_1
Vehicle_1



Panobinostat_1

418

69

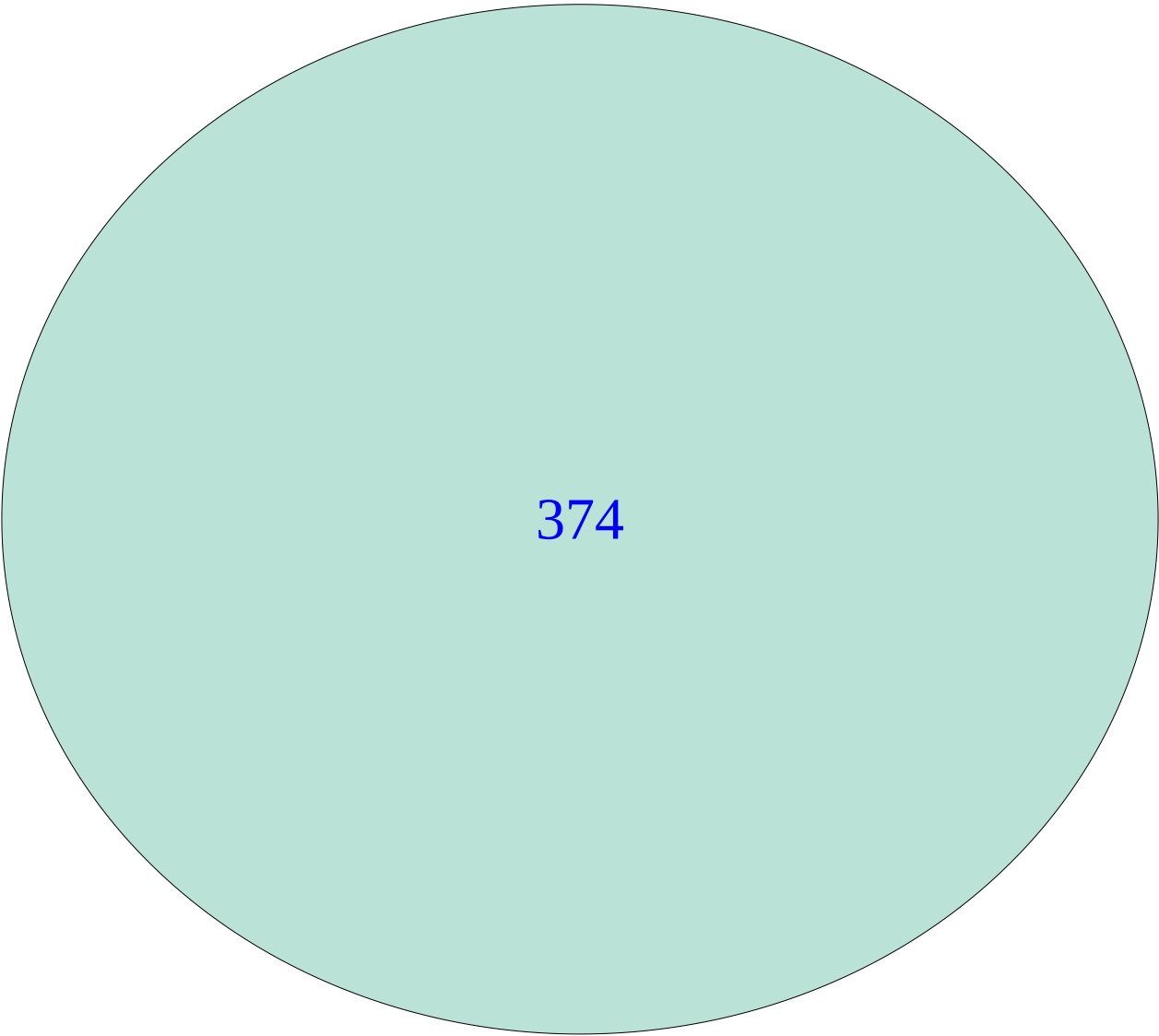
484

R2 >= 0.8

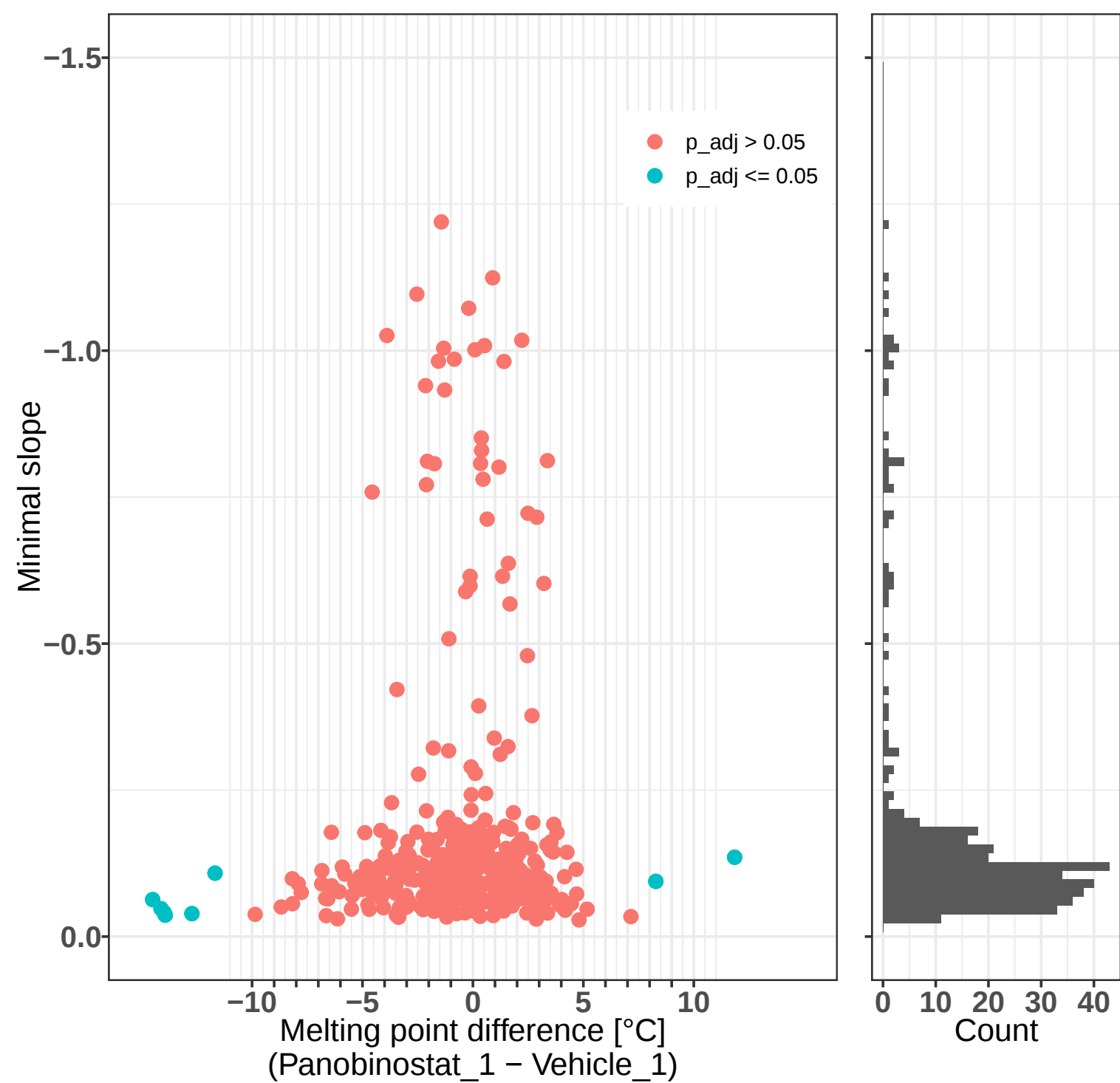
R2 < 0.8

Total

Panobinostat_1_vs_Vehicle_1



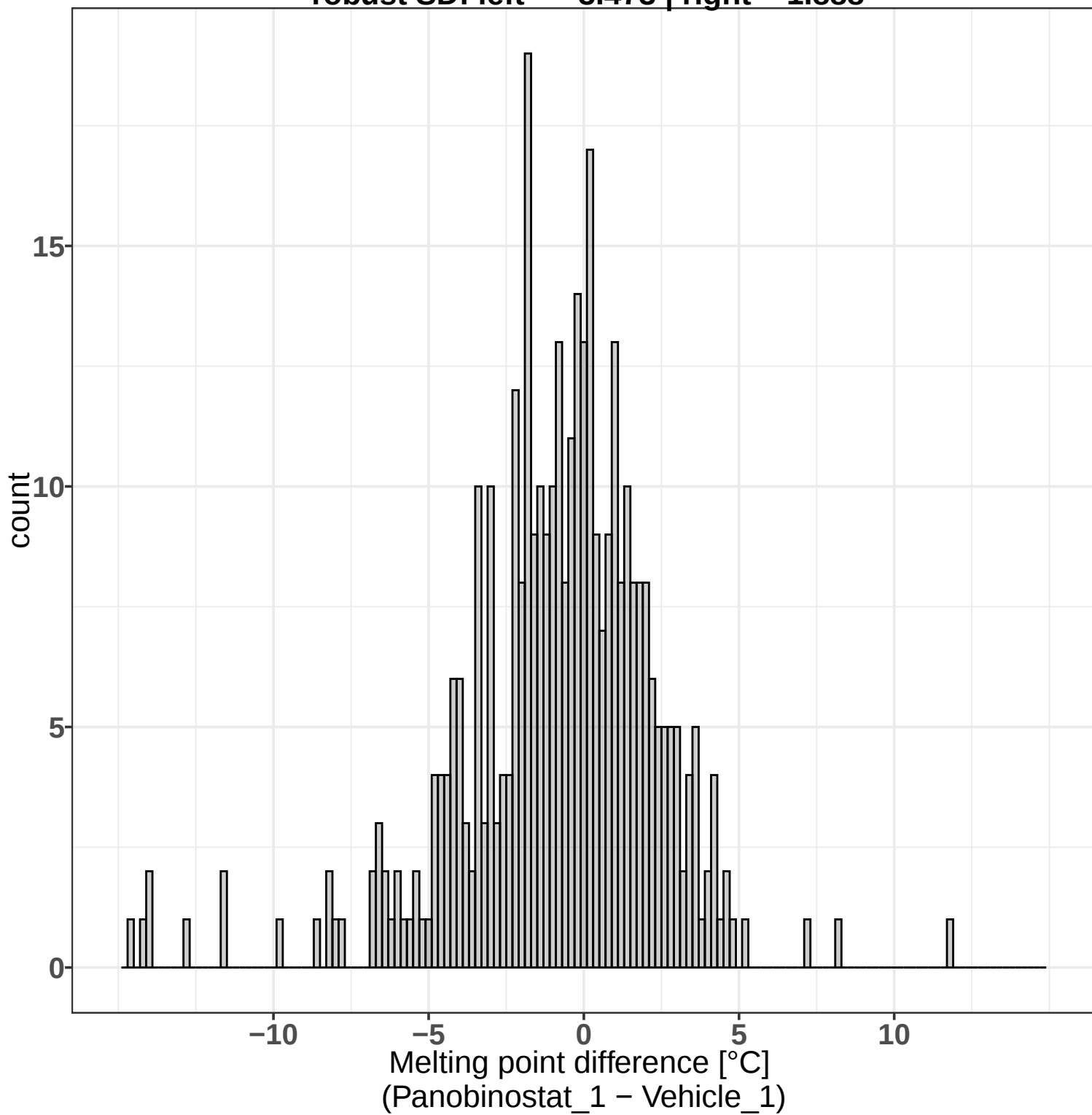
nobinostat_1_vs_Vehicle_1	374	110	484
	passed_filter	filtered_out	Total



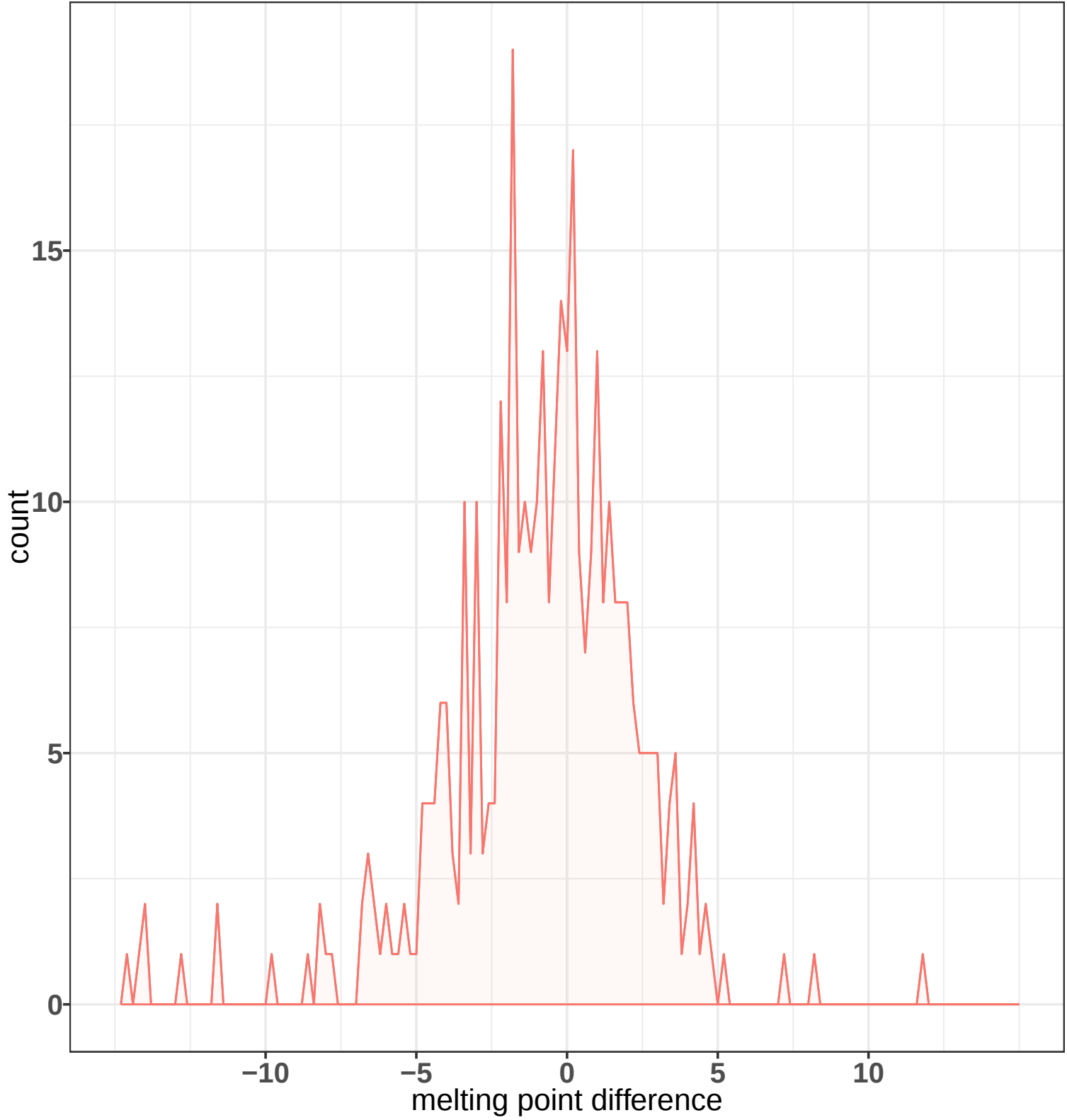
n = 371

Panobinostat_1 vs. Vehicle_1

**median = -0.567 ,
robust SD: left = -3.473 | right = 1.888**



n = 380 , R_sq >= 0.8



comparison  Panobinostat_1 - Vehicle_1