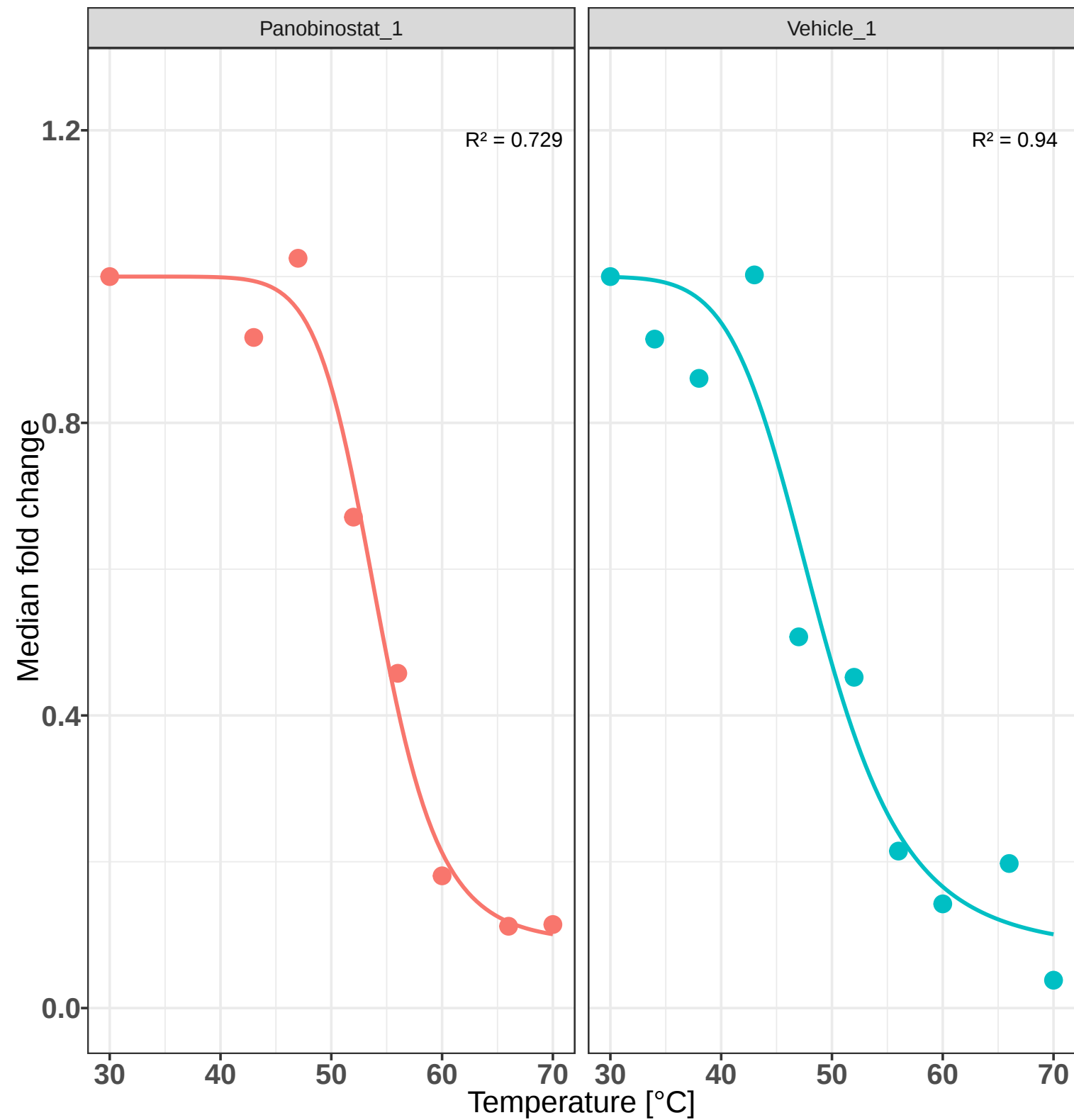
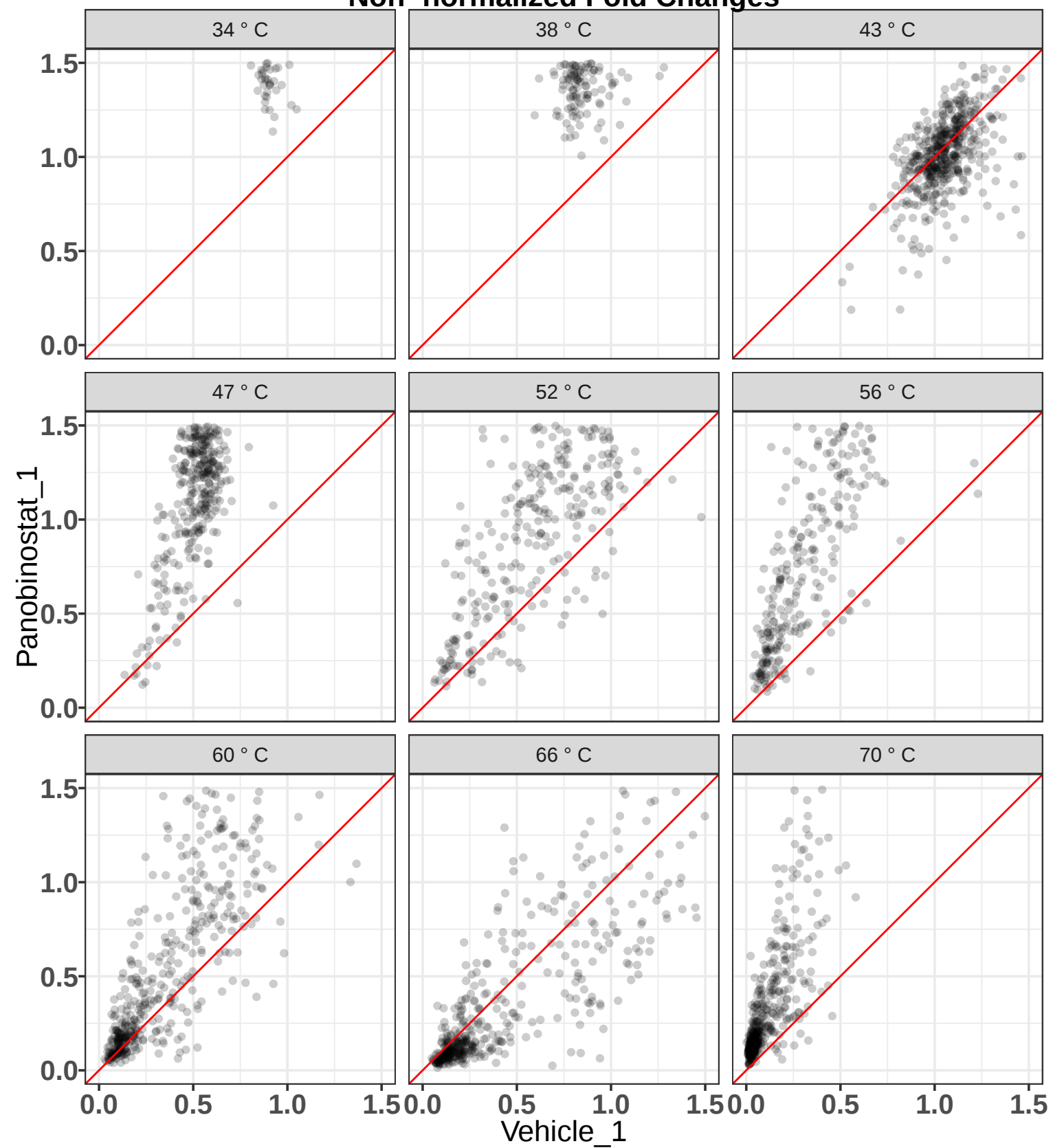


Normalization curves

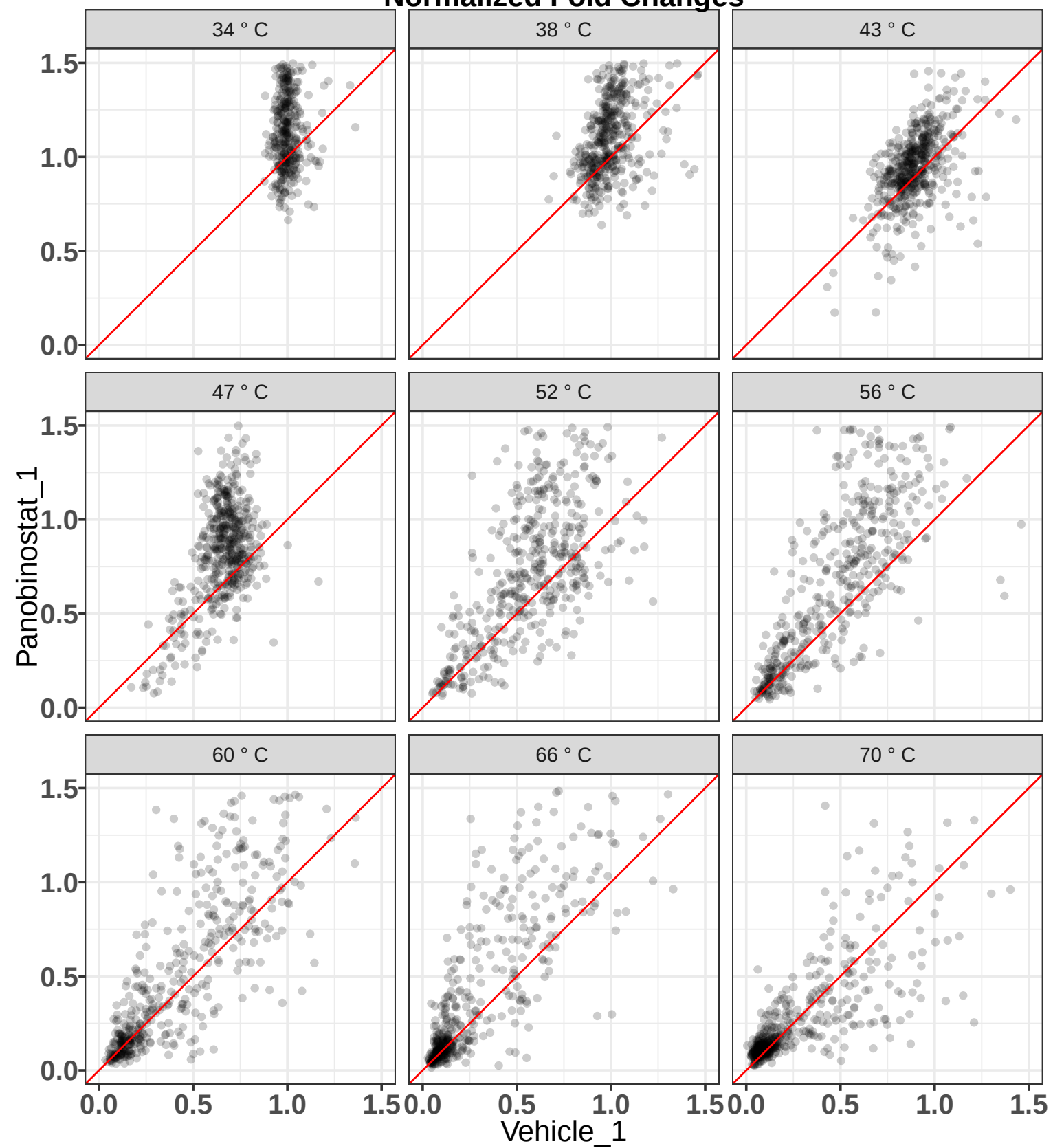
based on 39 proteins



Non-normalized Fold Changes

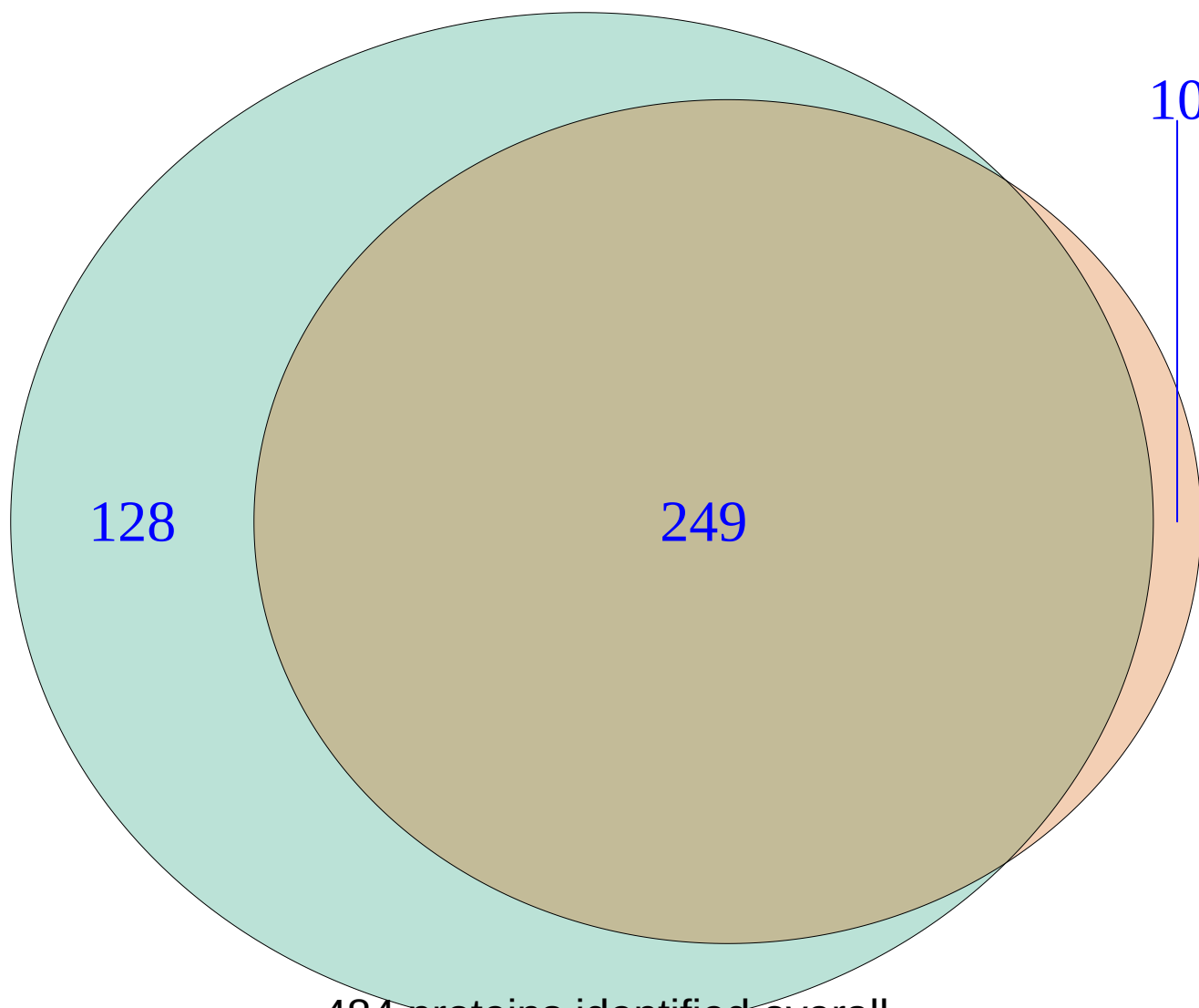


Normalized Fold Changes



Vehicle_1

Panobinostat_1



484 proteins identified overall

Vehicle_1

Panobinostat_1

249

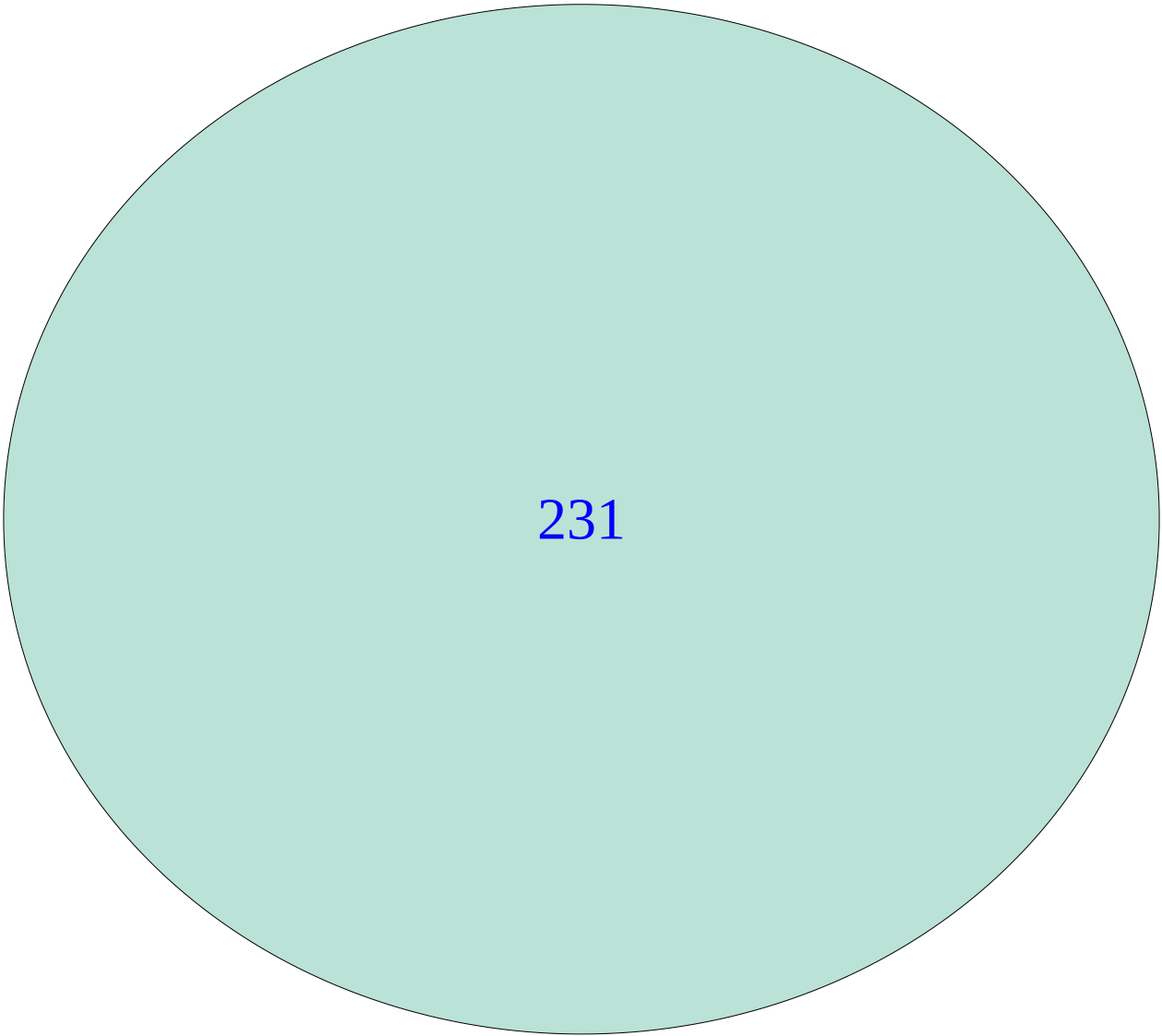
10

$R^2 \geq 0.8$

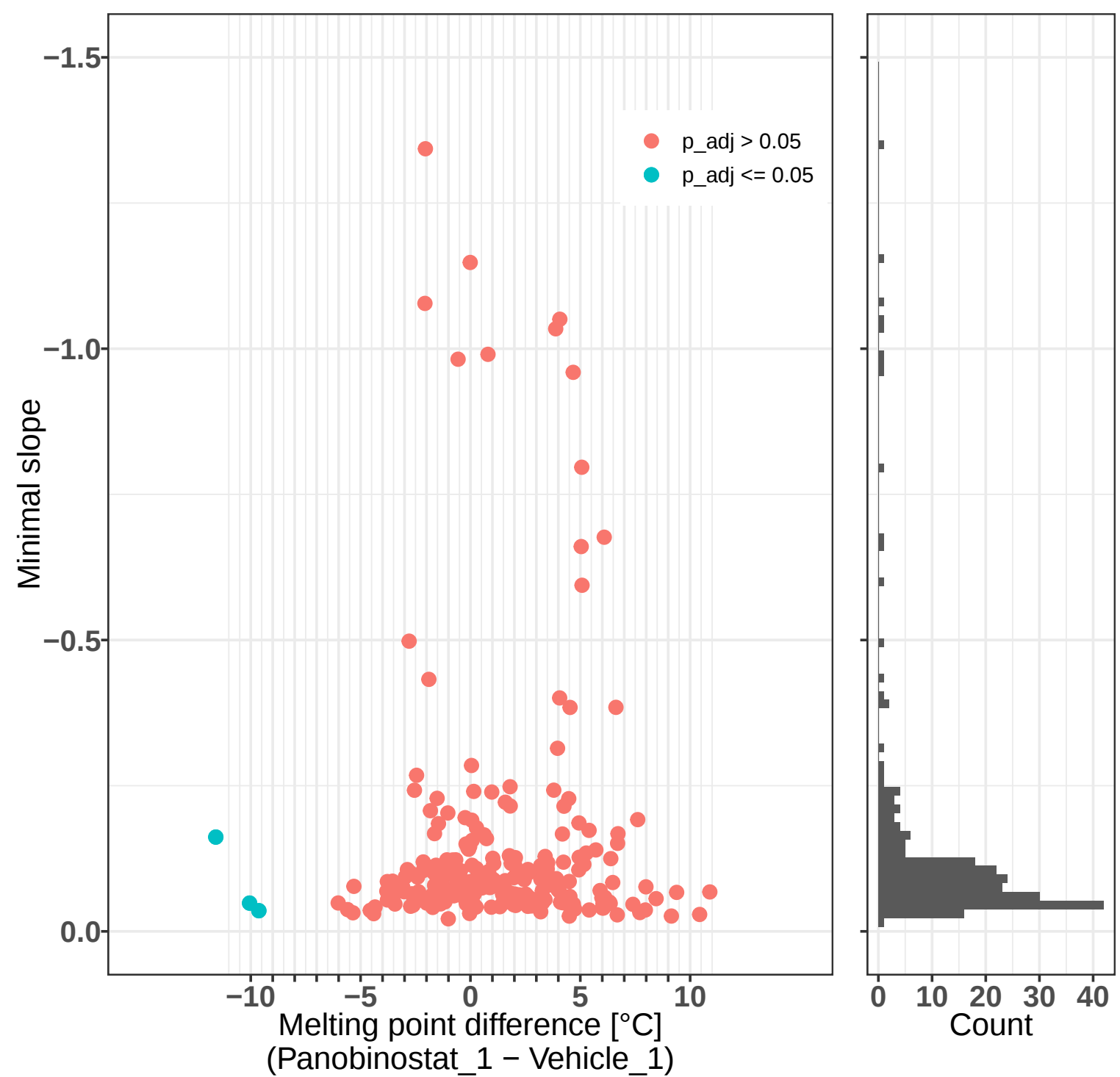
$R^2 < 0.8$

Total

Panobinostat_1_vs_Vehicle_1



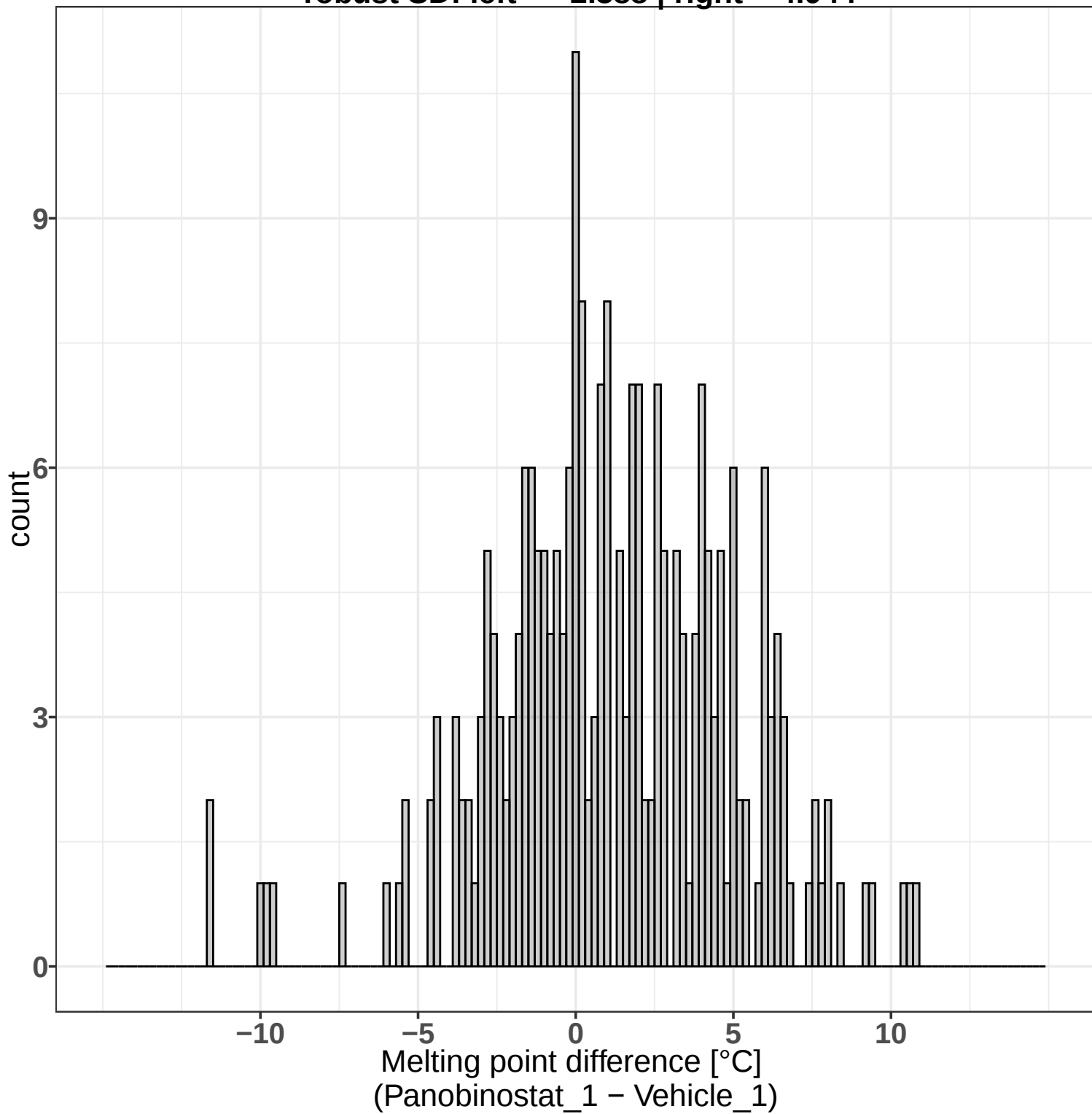
nobinostat_1_vs_Vehicle_1	231	253	484
passed_filter	filtered_out	Total	



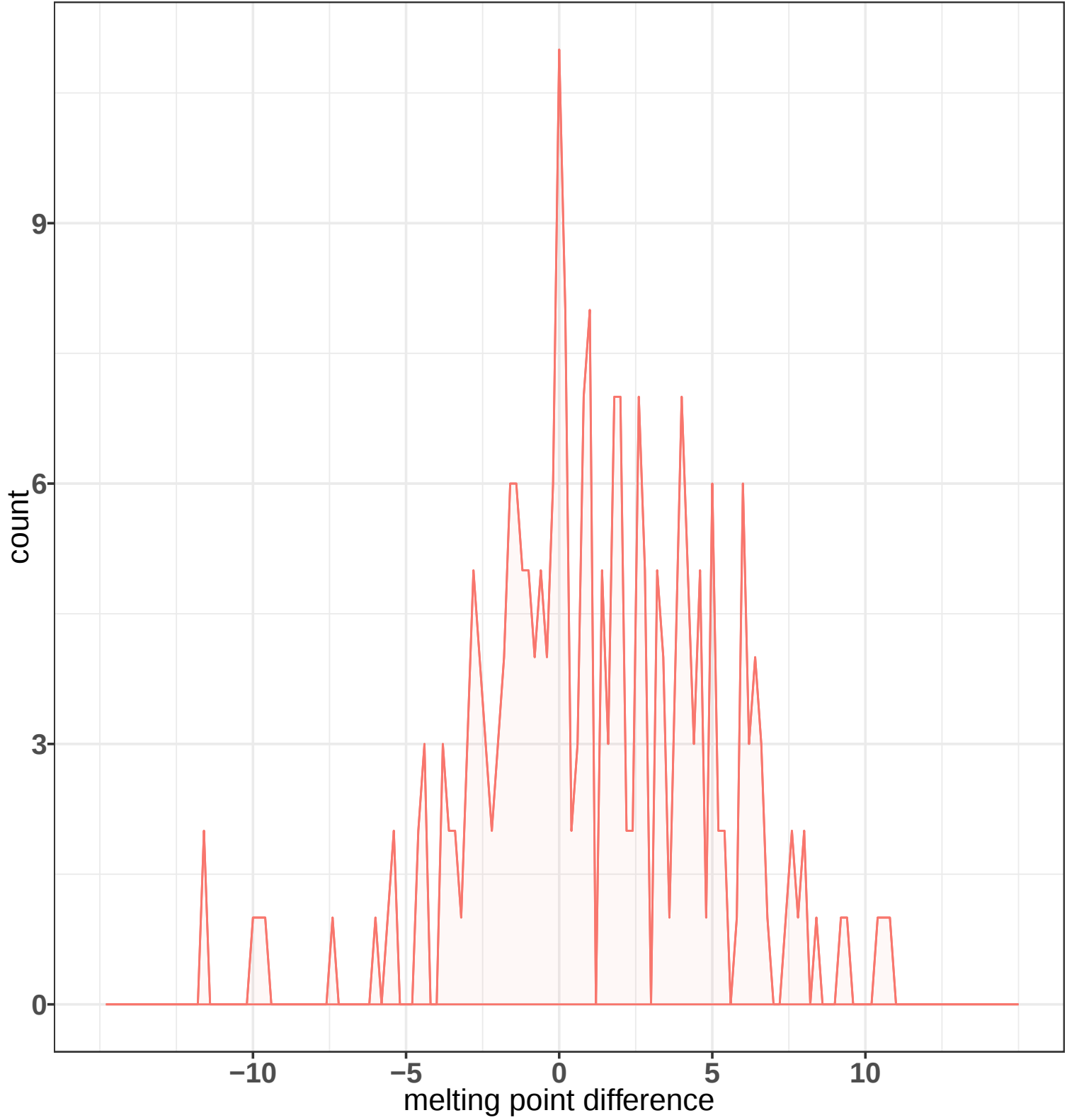
n = 231

Panobinostat_1 vs. Vehicle_1

**median = 0.919 ,
robust SD: left = -2.388 | right = 4.944**



n = 249 , R_sq >= 0.8



comparison Panobinostat_1 - Vehicle_1