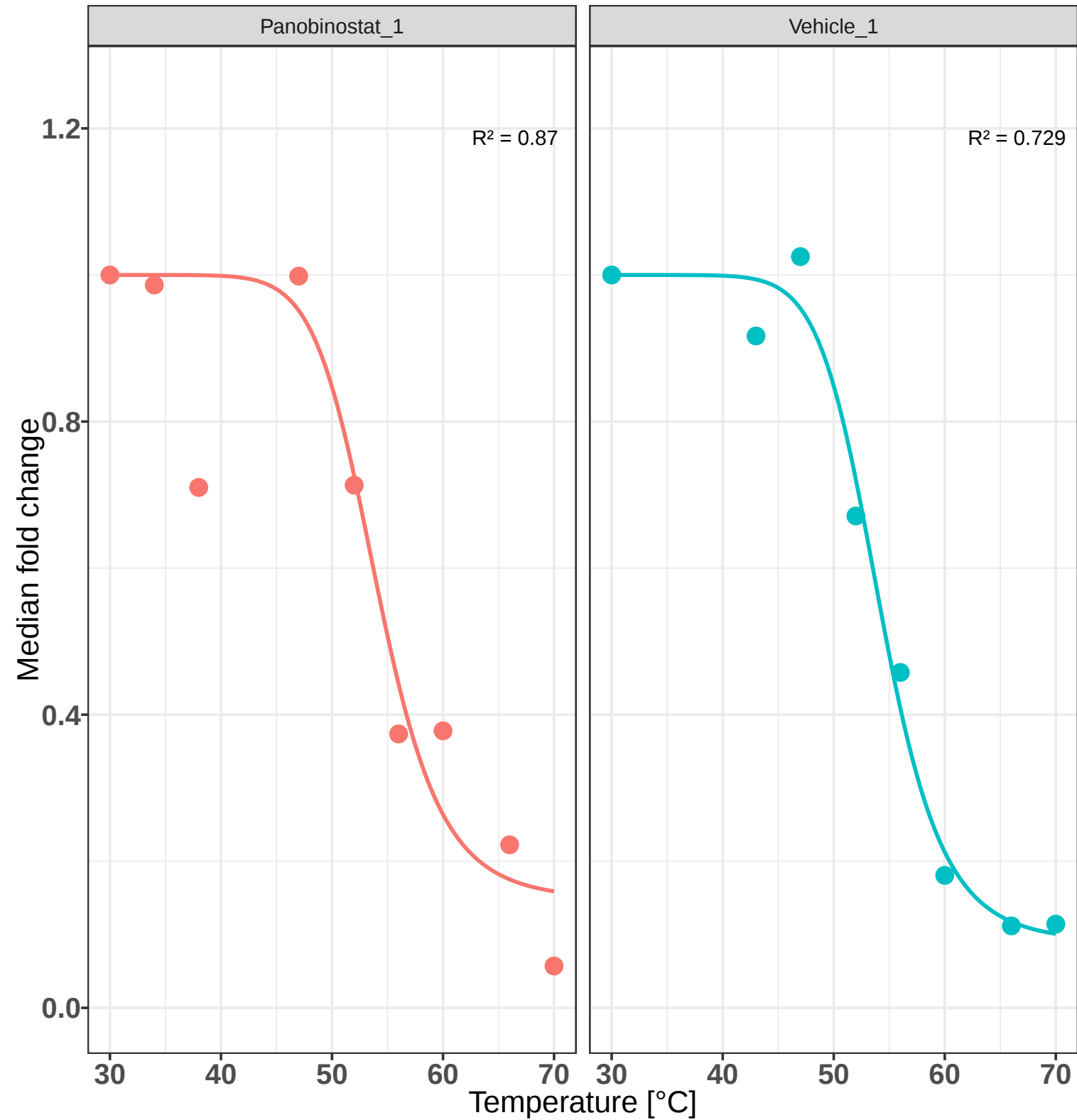
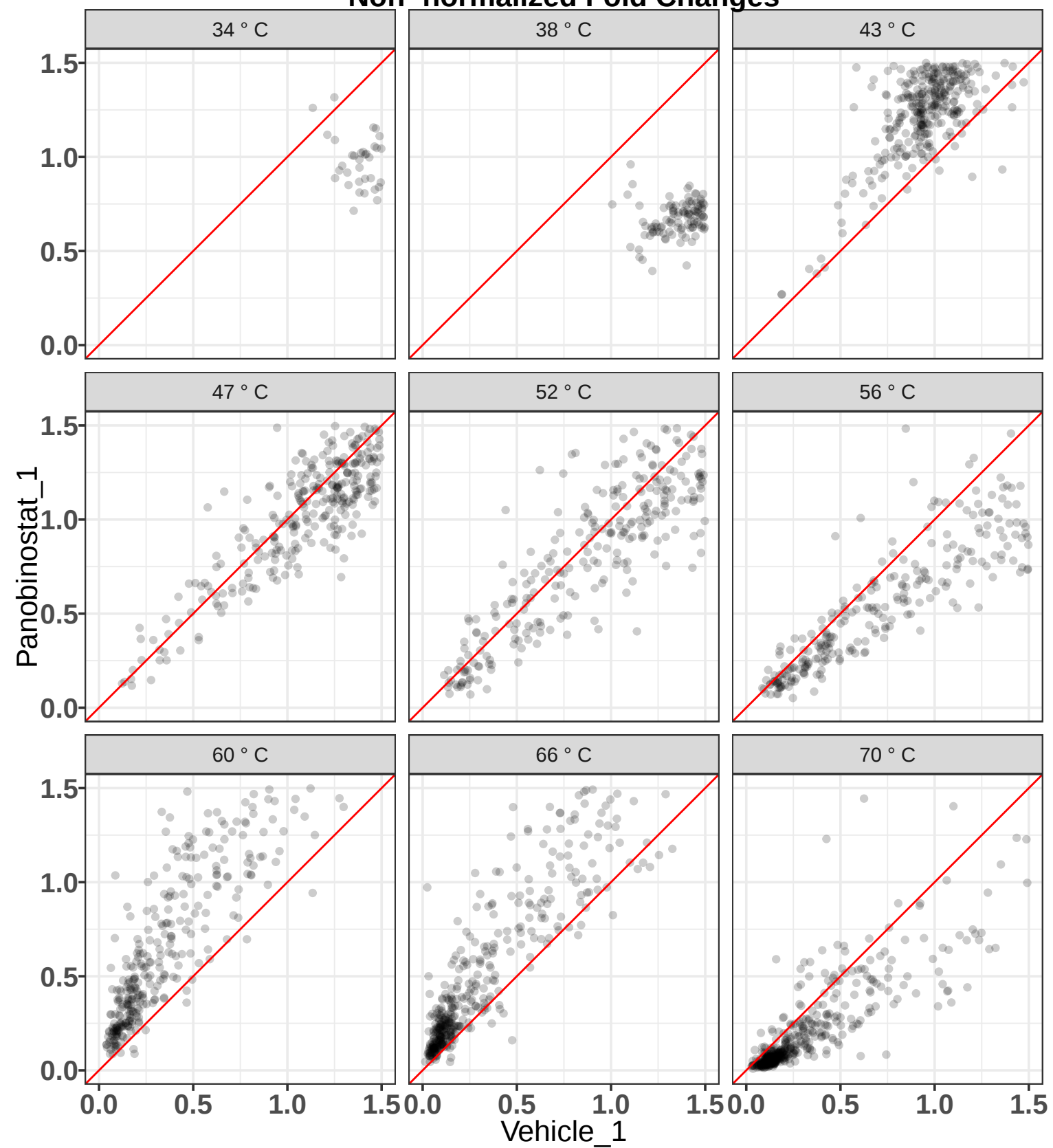


Normalization curves

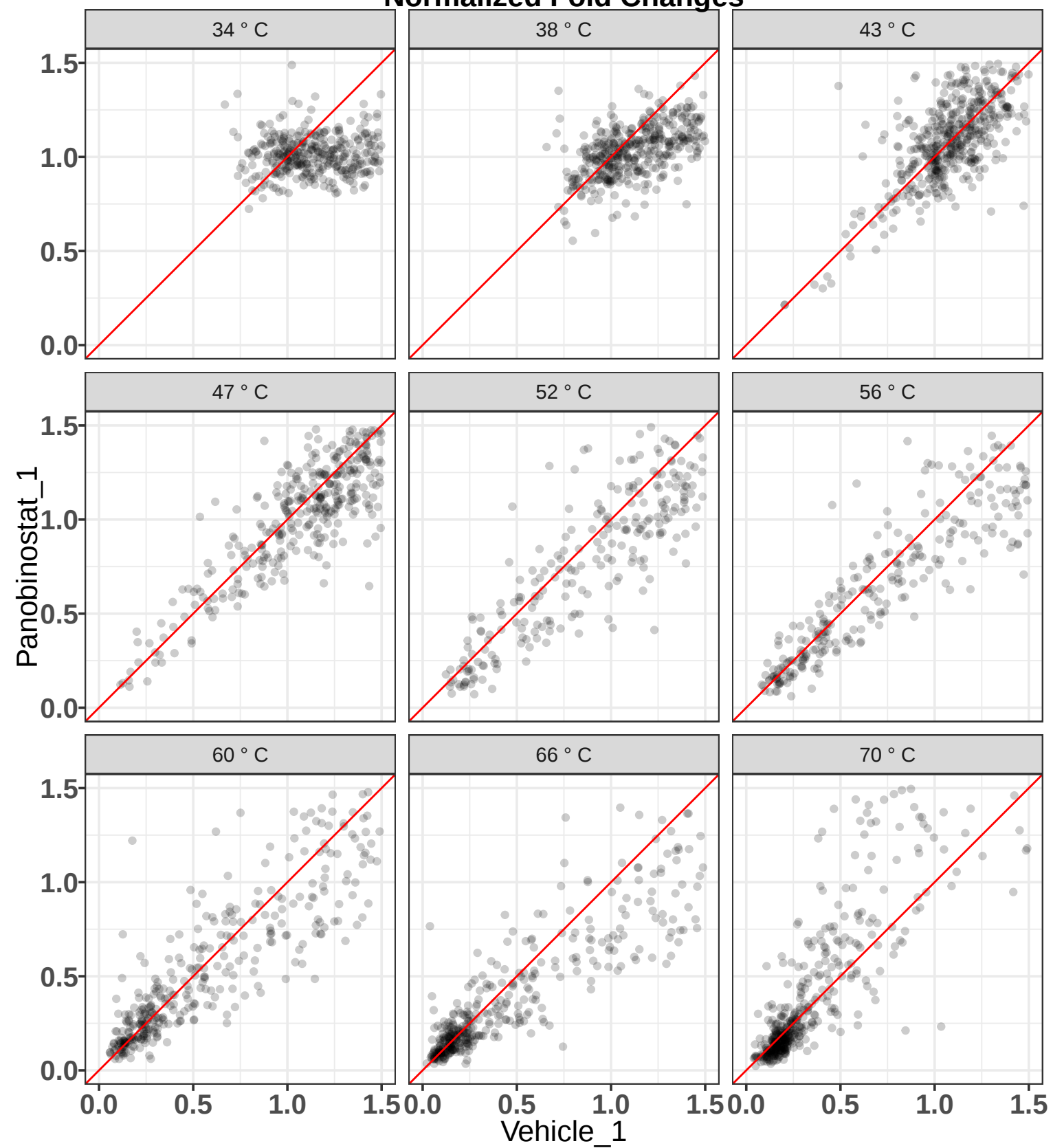
based on 39 proteins

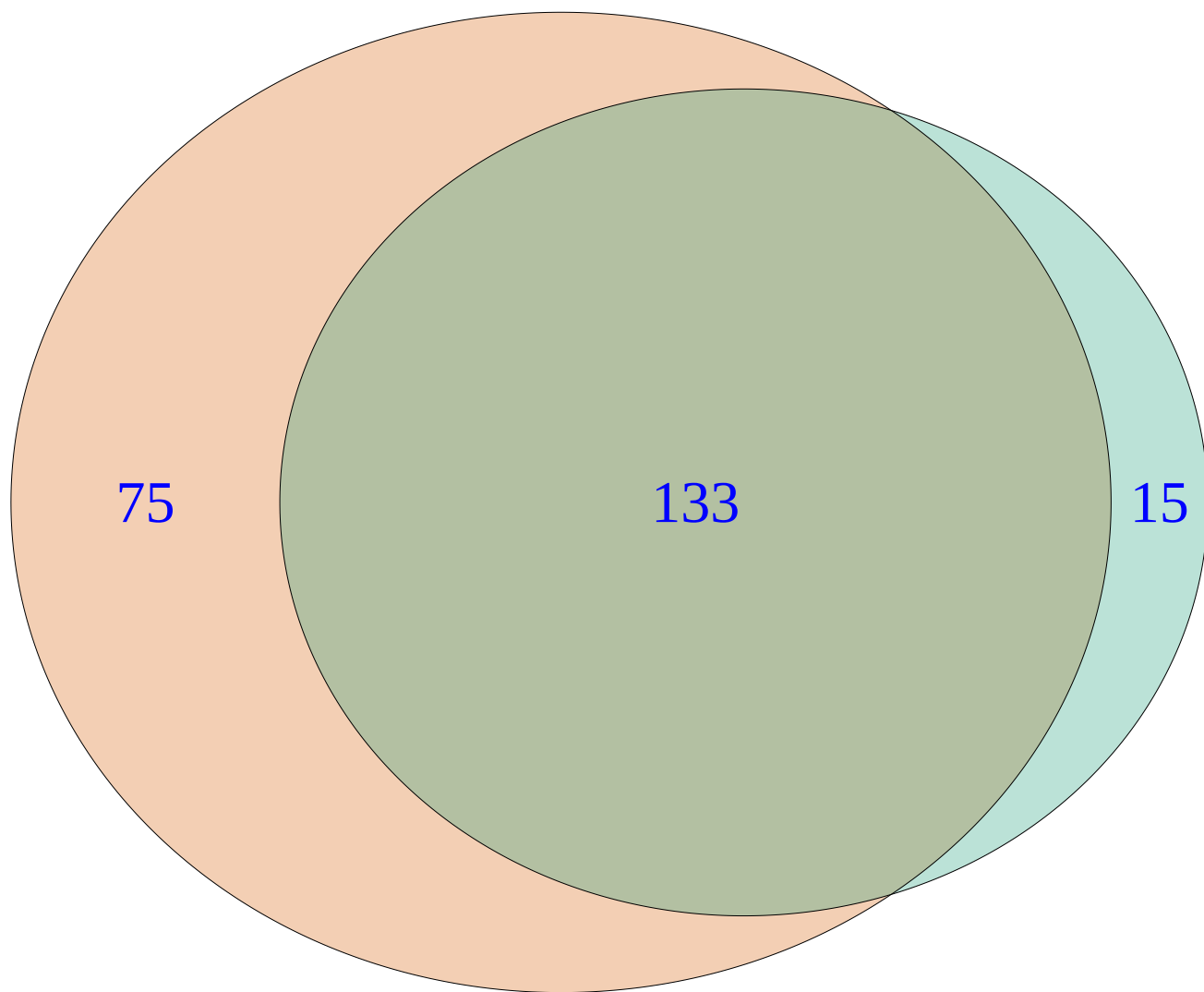


Non-normalized Fold Changes



Normalized Fold Changes





Vehicle_1

484 proteins identified overall

Panobinostat_1

Vehicle_1

208

236

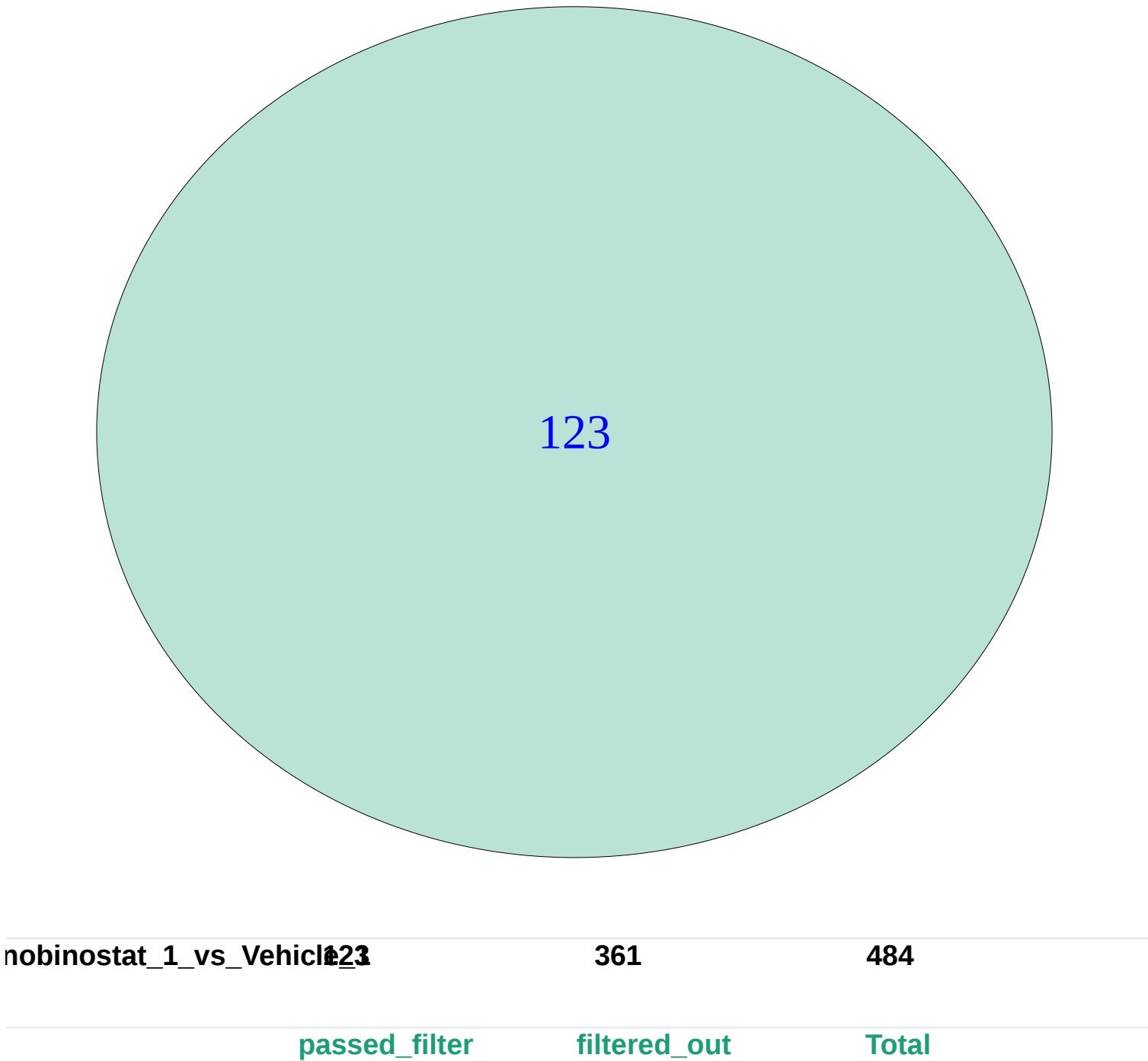
484

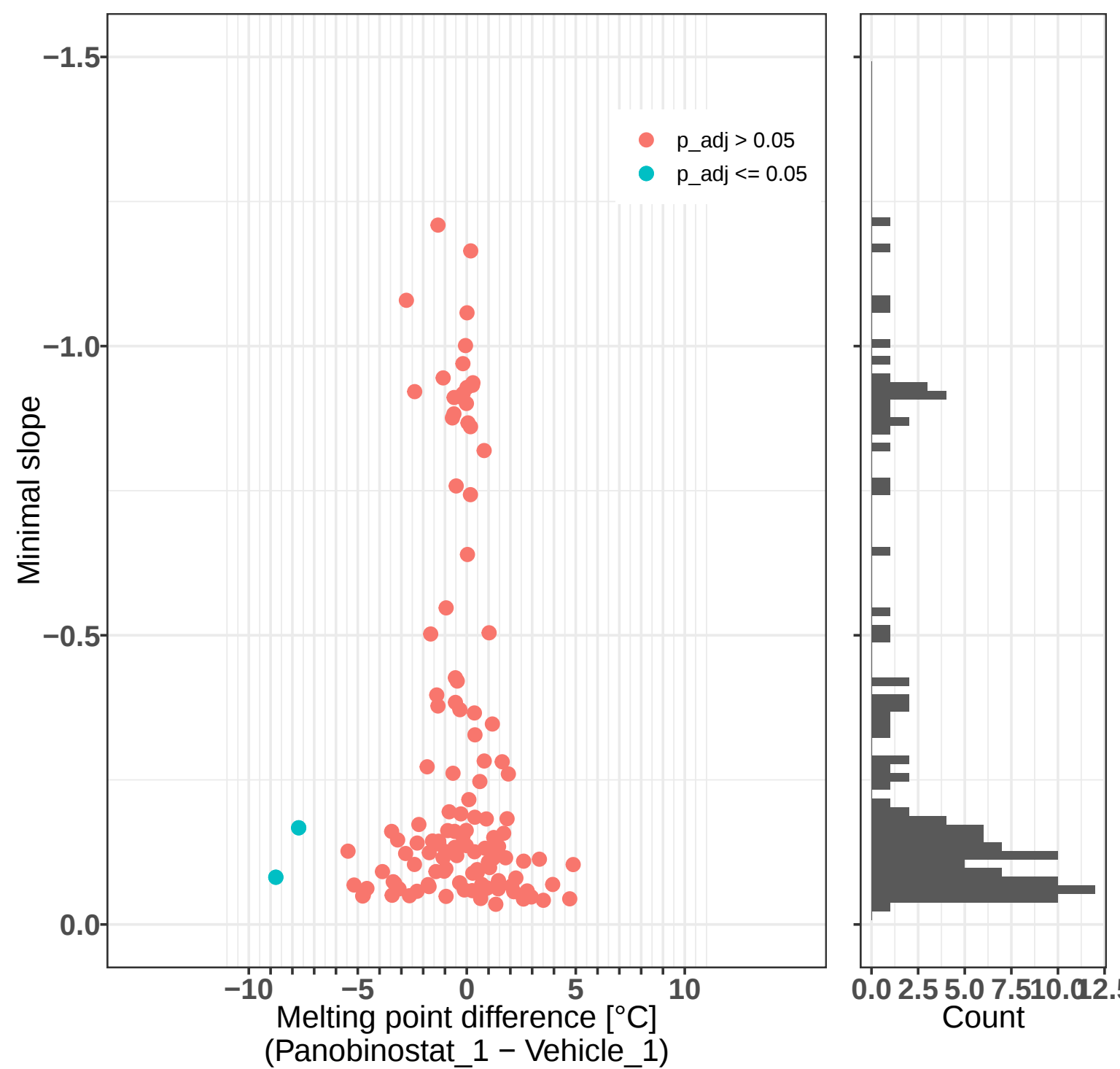
R2 >= 0.8

R2 < 0.8

Total

Panobinostat_1_vs_Vehicle_1

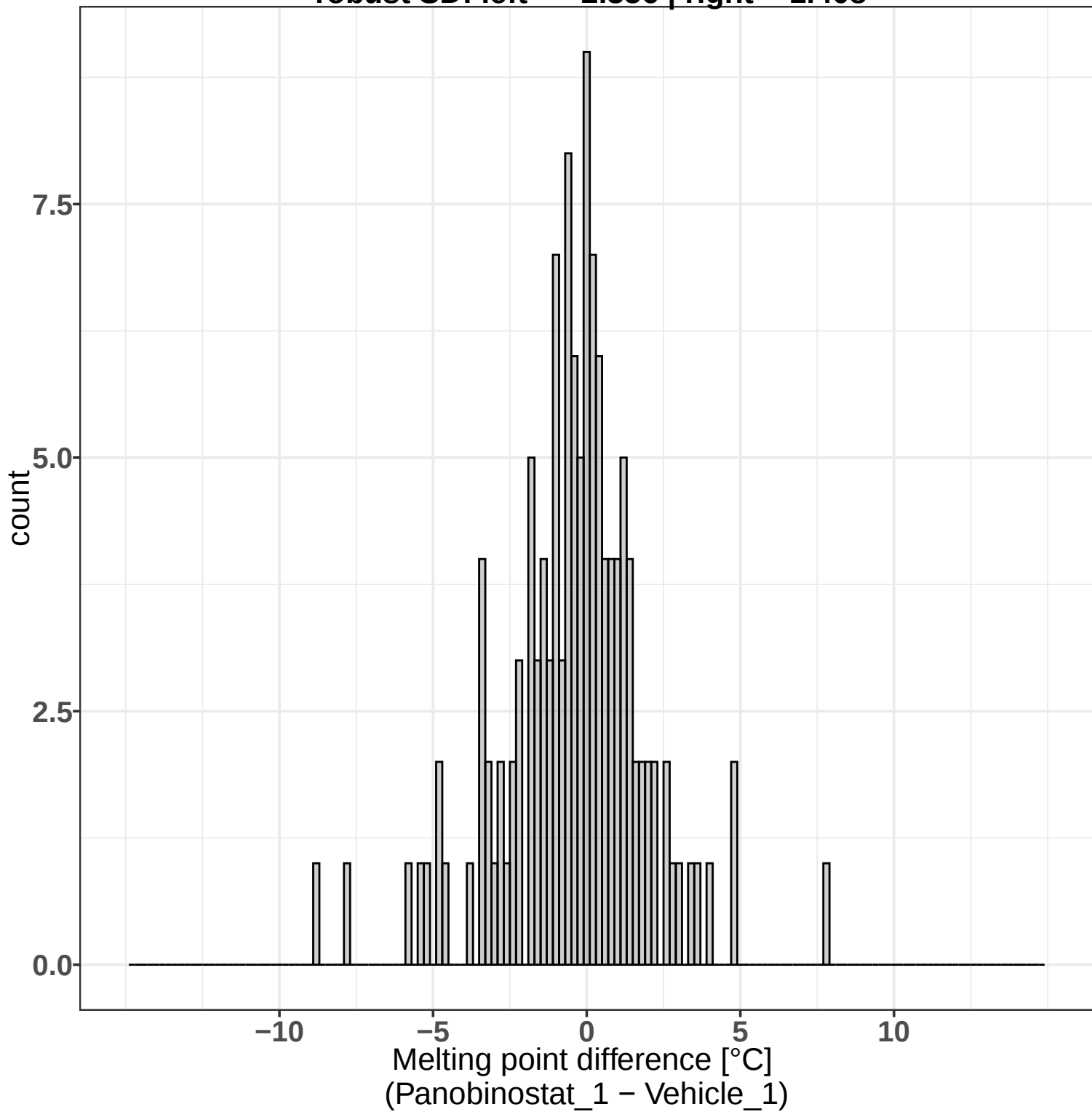




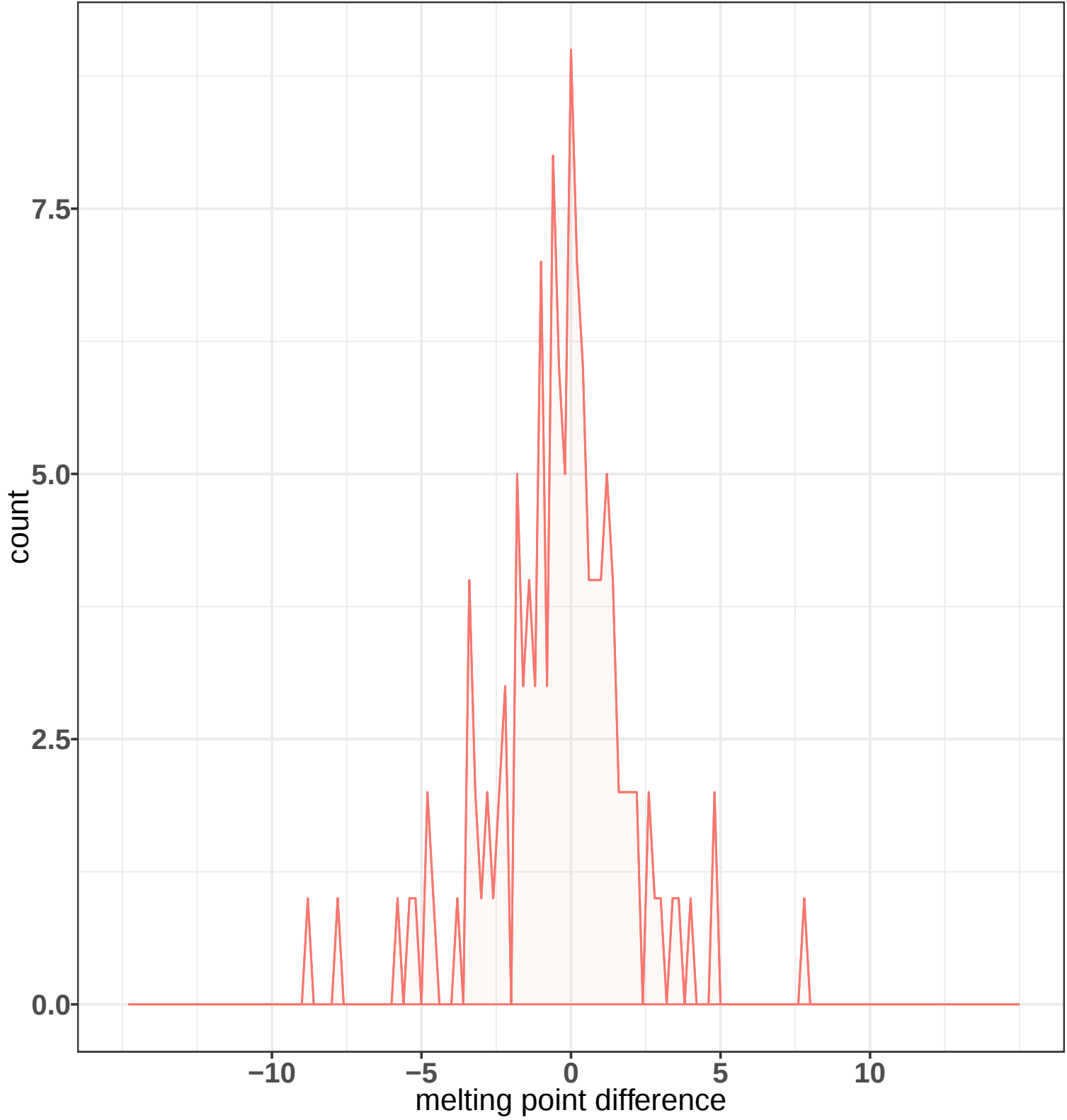
n = 122

Panobinostat_1 vs. Vehicle_1

median = -0.178 ,
robust SD: left = -2.356 | right = 1.408



n = 133 , R_sq >= 0.8



comparison Panobinostat_1 - Vehicle_1