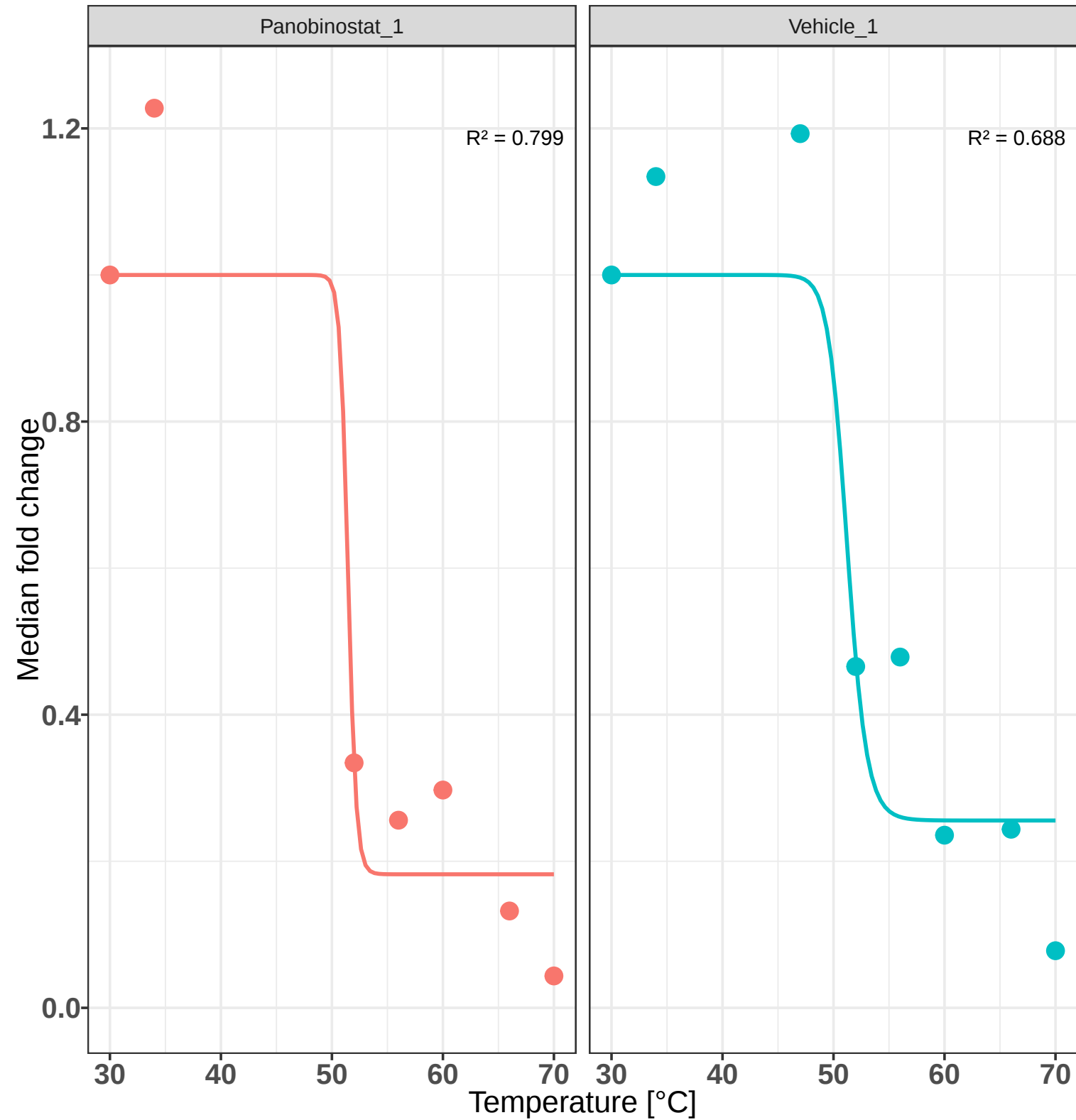
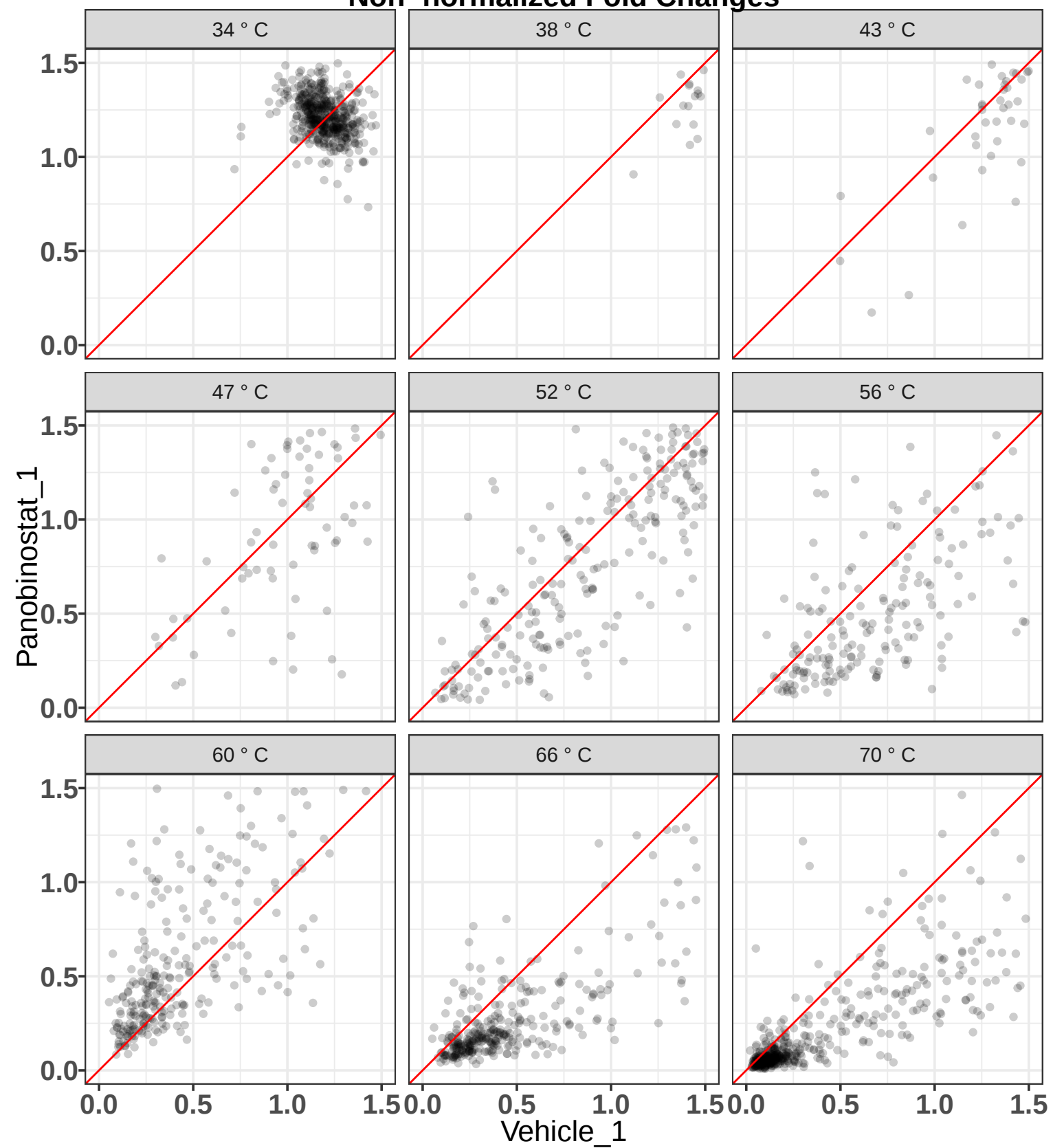


Normalization curves

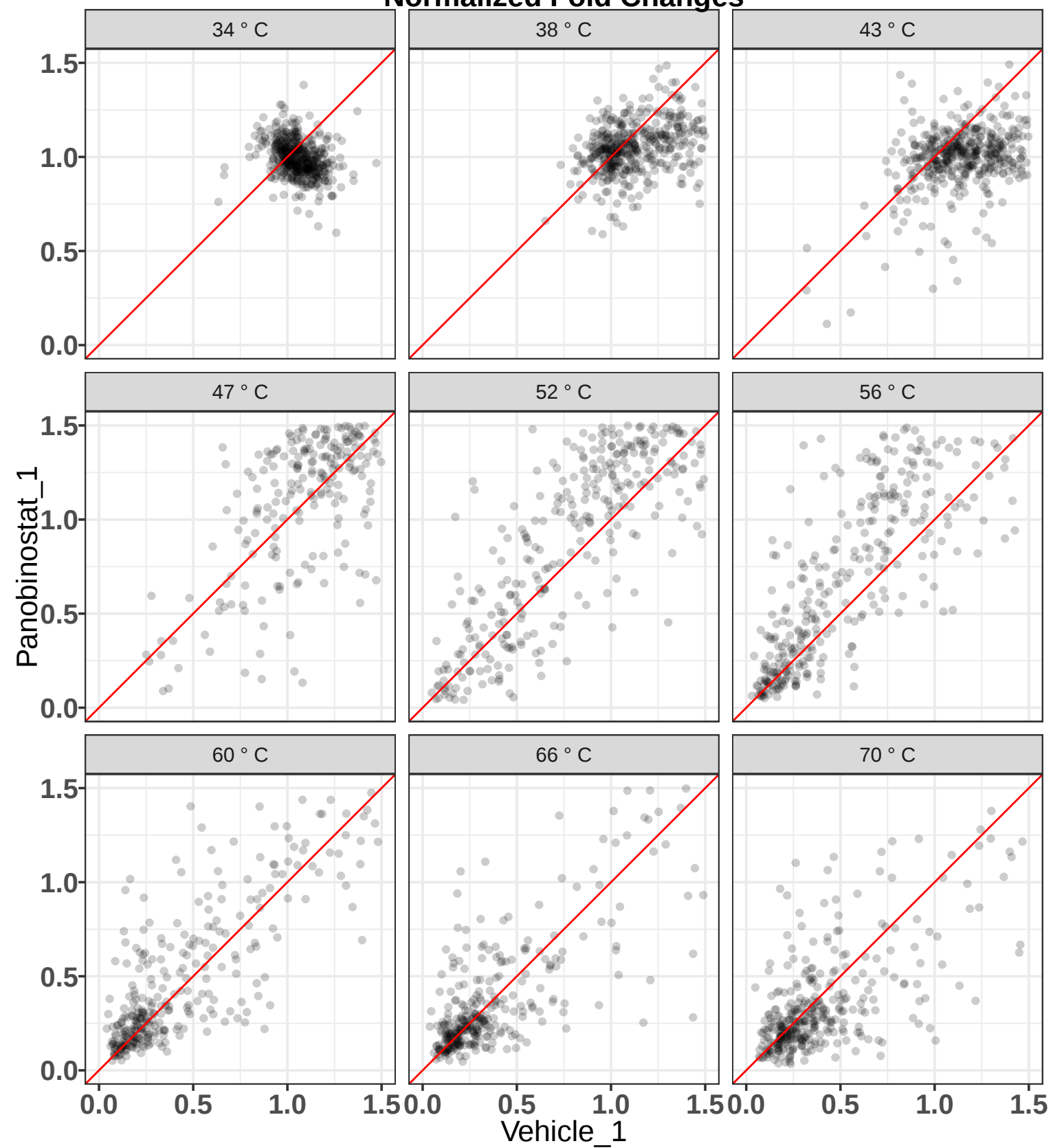
based on 29 proteins



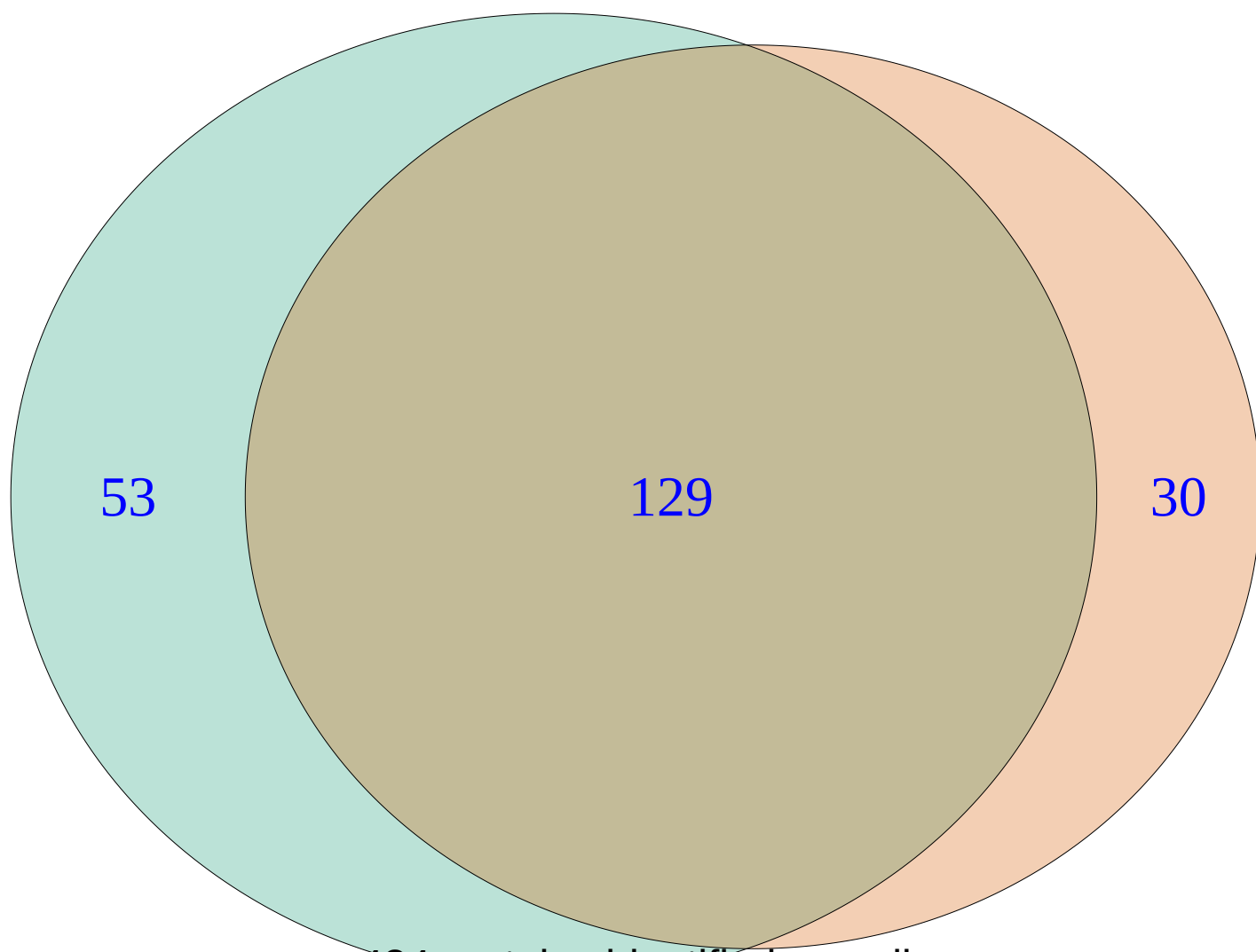
Non-normalized Fold Changes



Normalized Fold Changes



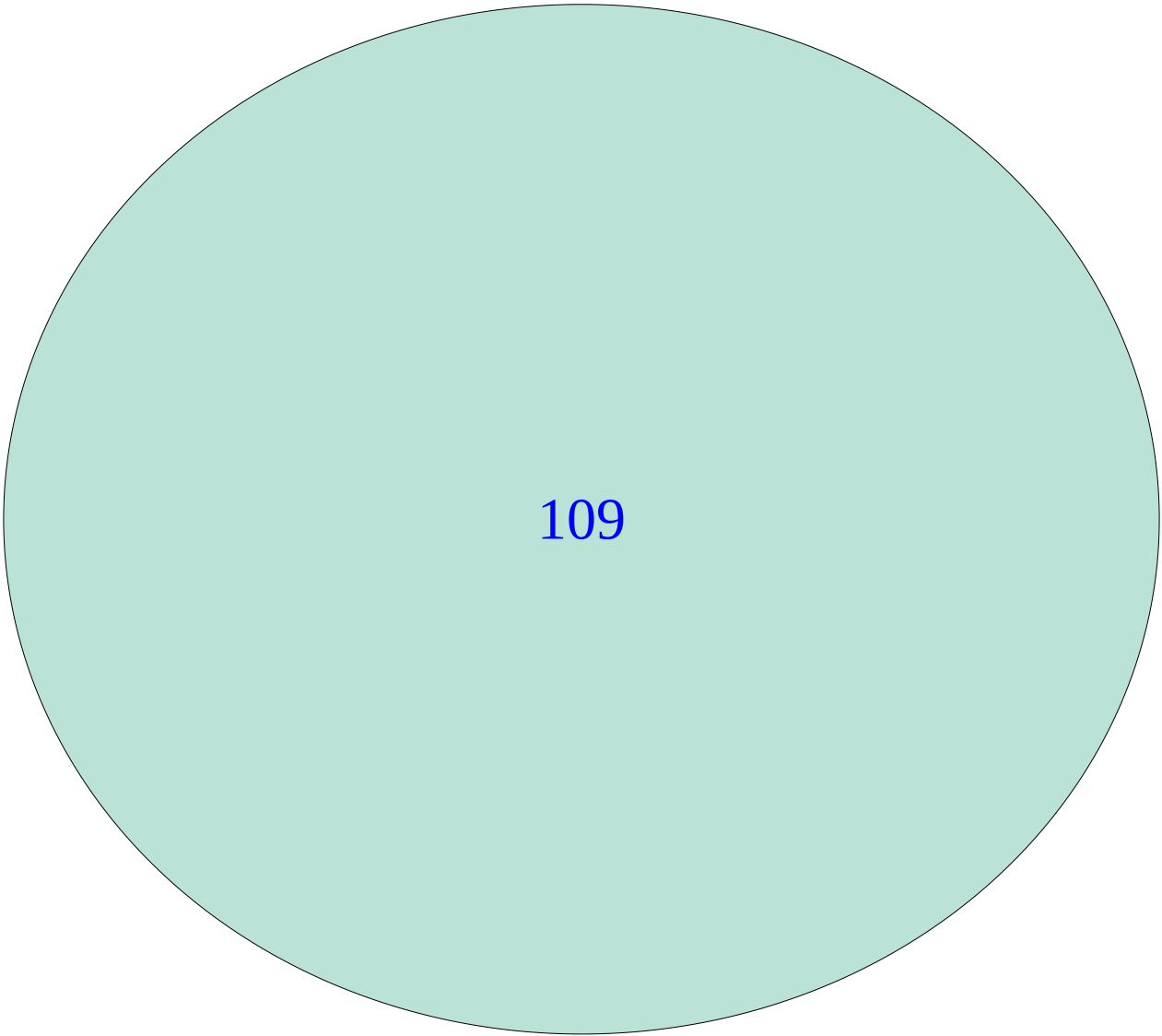
Vehicle_1 Panobinostat_1



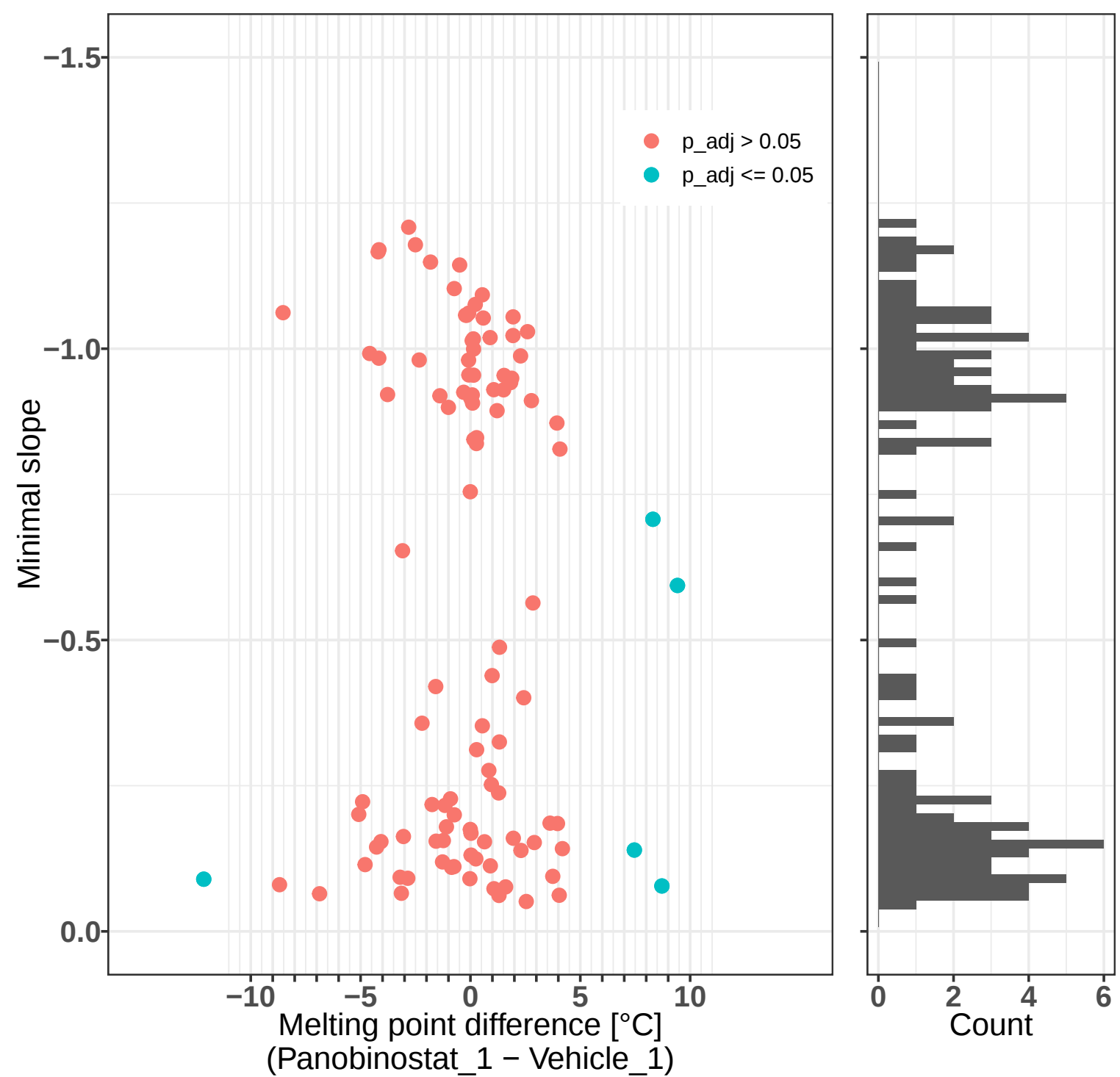
484 proteins identified overall

Panobinostat_1	Vehicle_1	182	302	484
		R2 >= 0.8	R2 < 0.8	Total

Panobinostat_1_vs_Vehicle_1



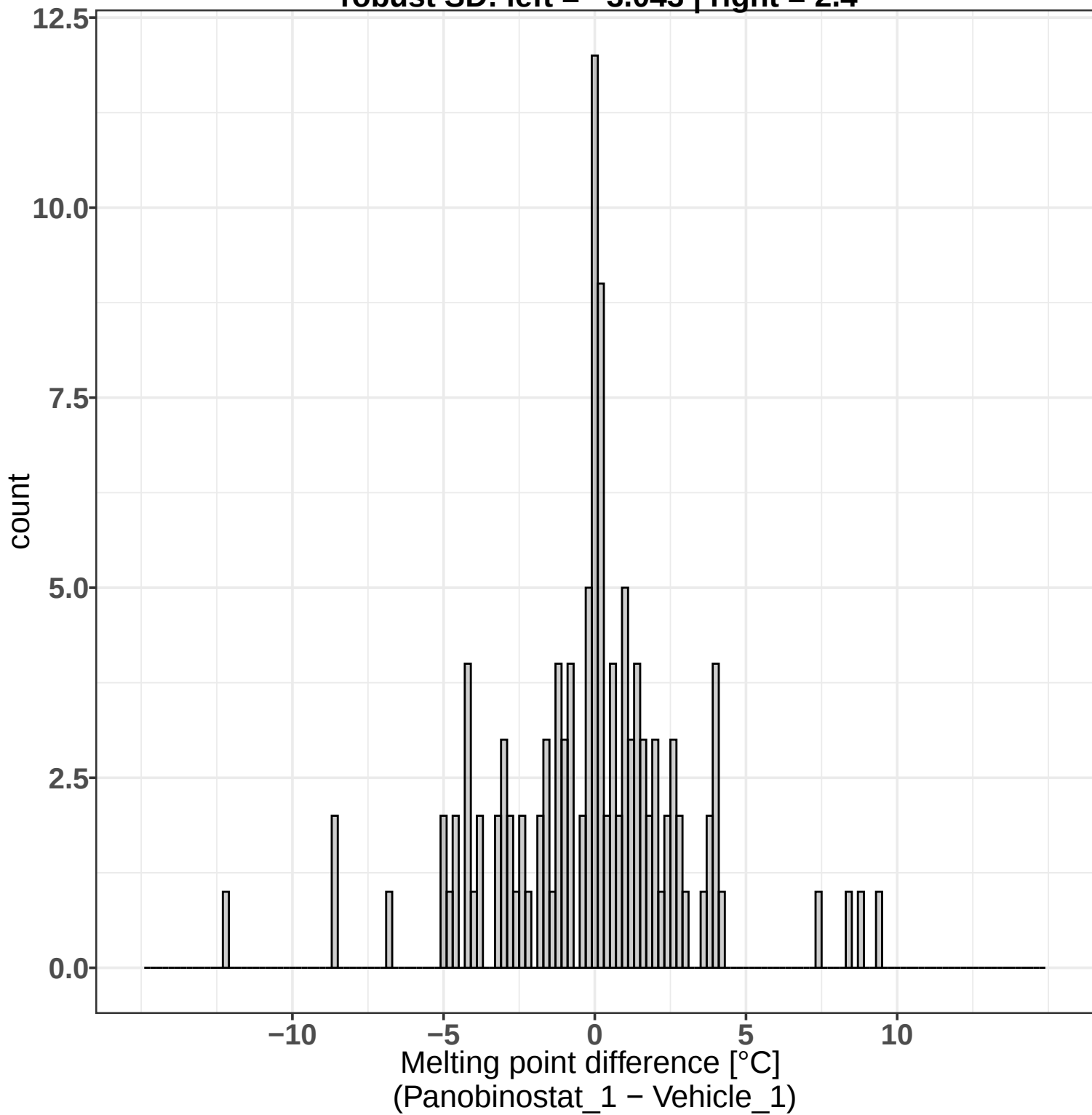
nobinostat_1_vs_Vehicle_1	375	484
passed_filter	filtered_out	Total



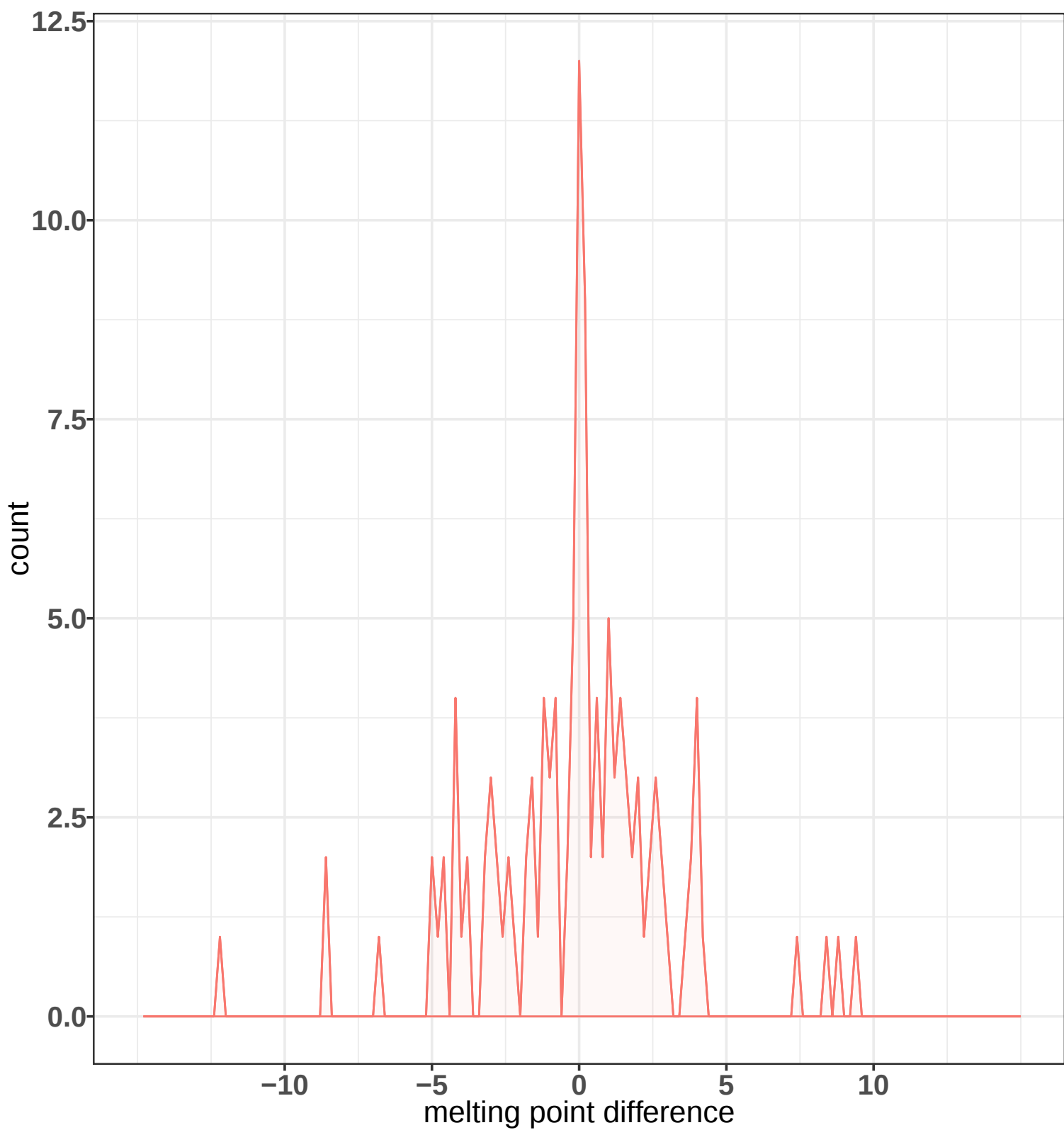
n = 107

Panobinostat_1 vs. Vehicle_1

median = 0.081 ,
robust SD: left = -3.043 | right = 2.4



n = 129 , R_sq >= 0.8



comparison Panobinostat_1 - Vehicle_1