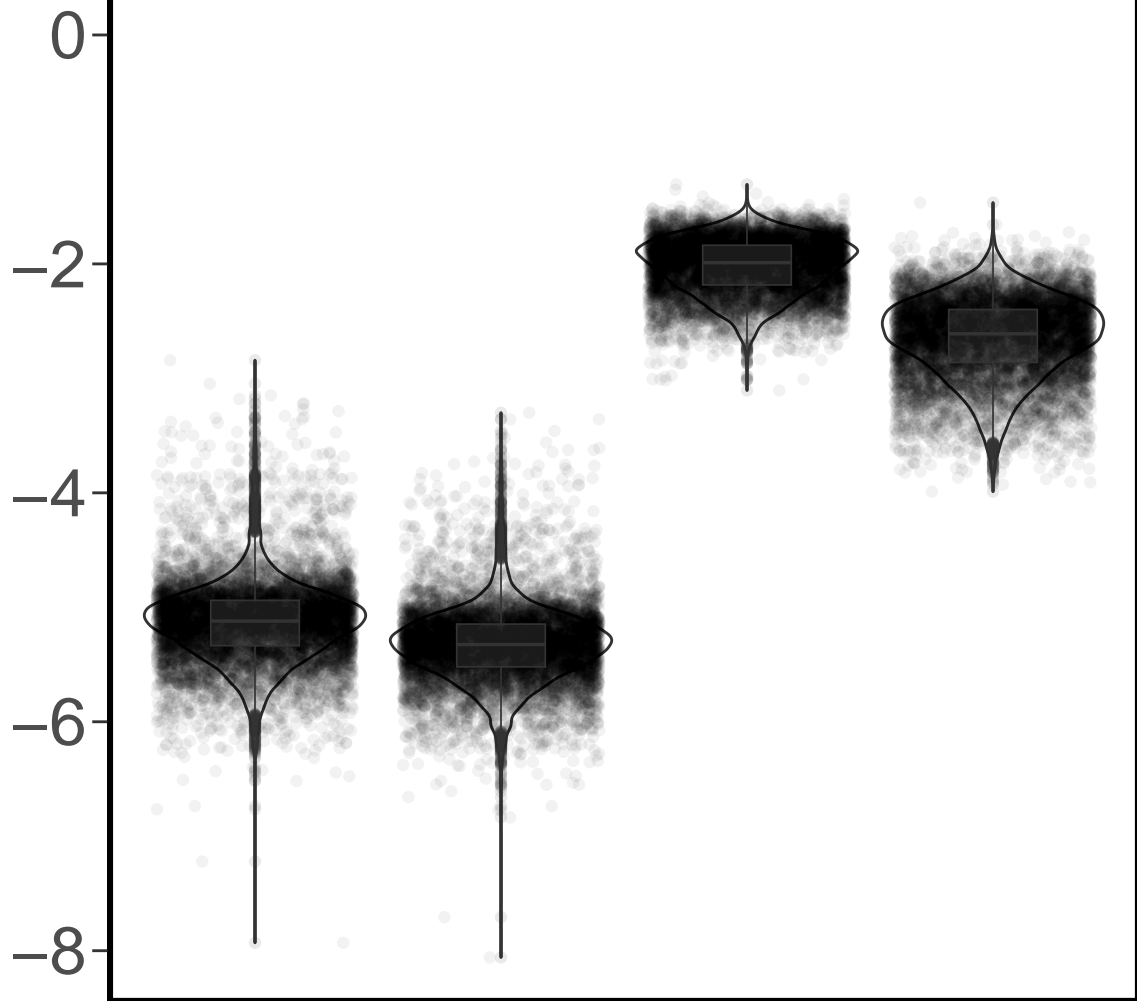


**log<sub>10</sub> raw values – all genes**



B\_base\_mean

B\_expr\_mean

P\_base\_mean

P\_expr\_mean