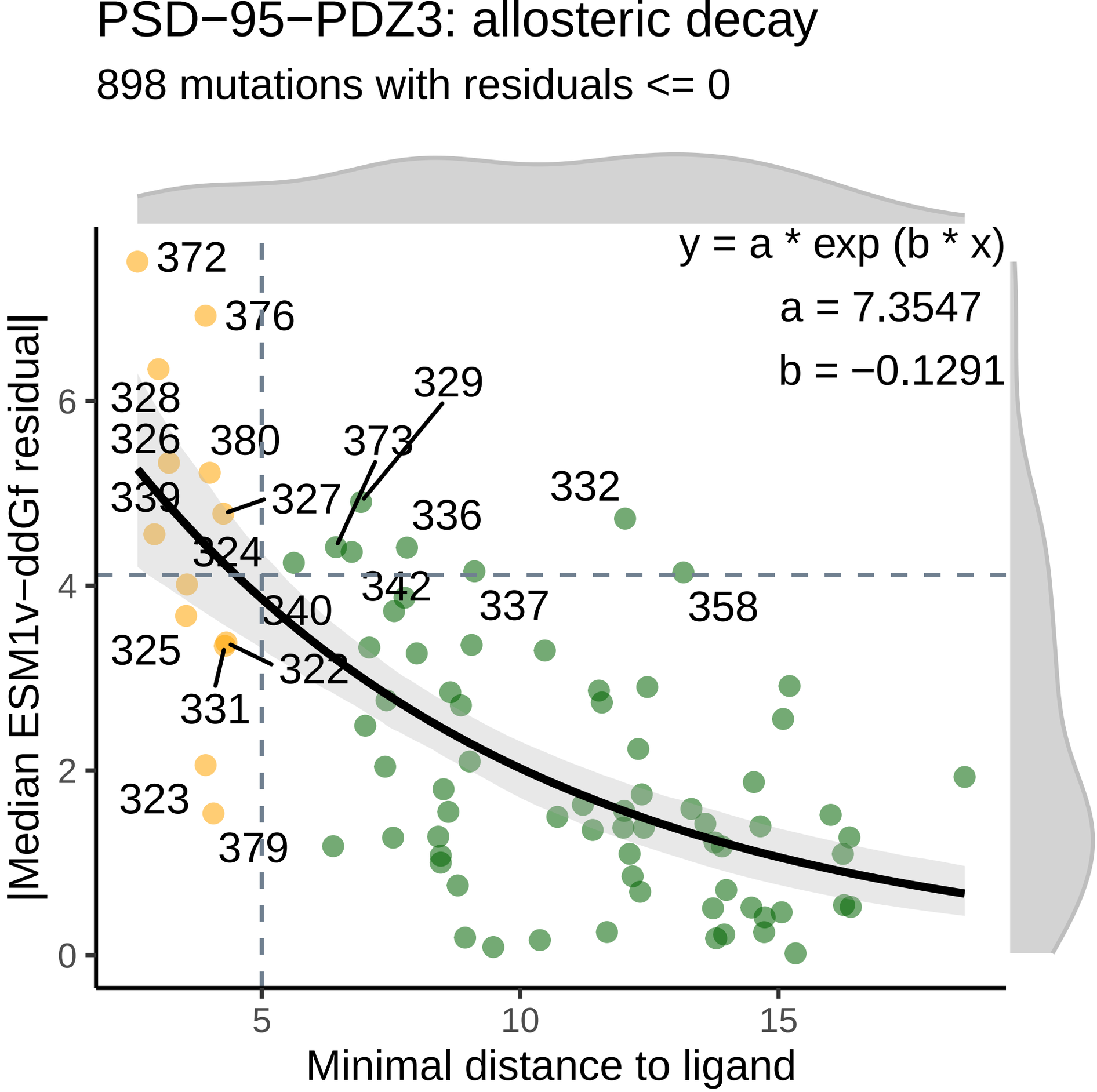
PSD−95−PDZ3: allosteric decay
898 mutations with residuals <= 0
y = a * exp (b * x)
a = 7.3547
b = −0.1291
372
376
328
326
380
329
373
339
327
332
336
324
342
340
337
358
325
322
331
323
379
|Median ESM1v−ddGf residual|
0
2
4
6
5
10
15
Minimal distance to ligand