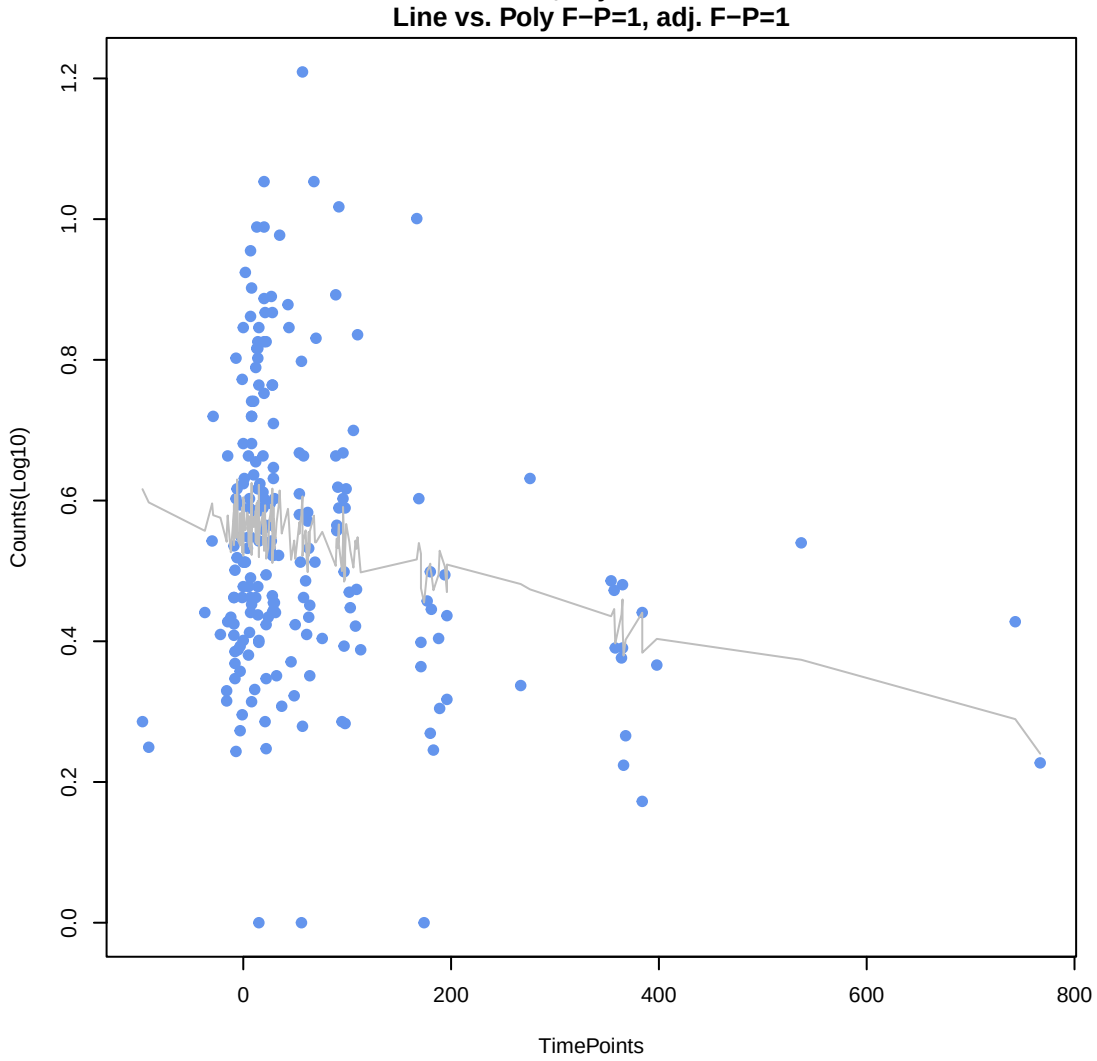
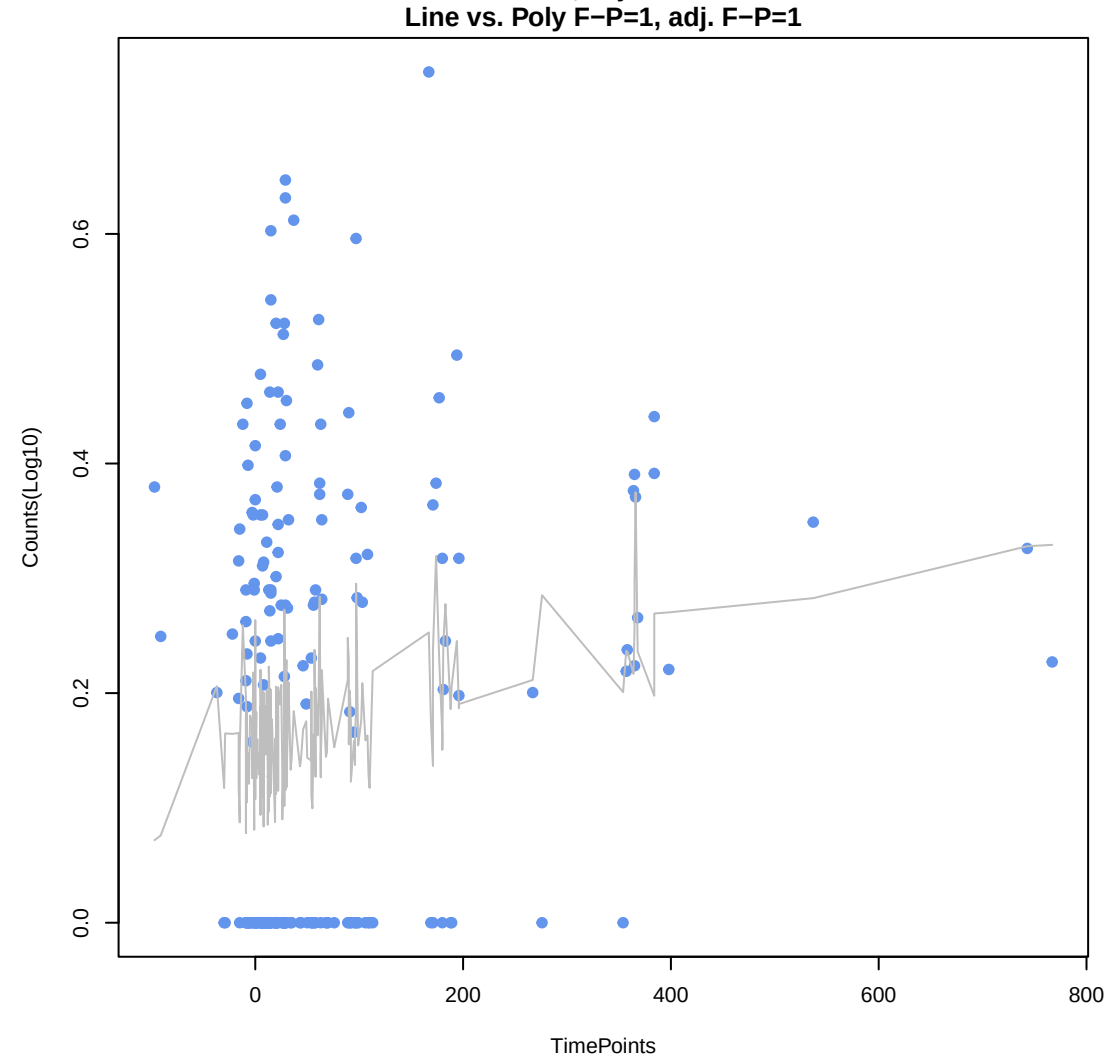
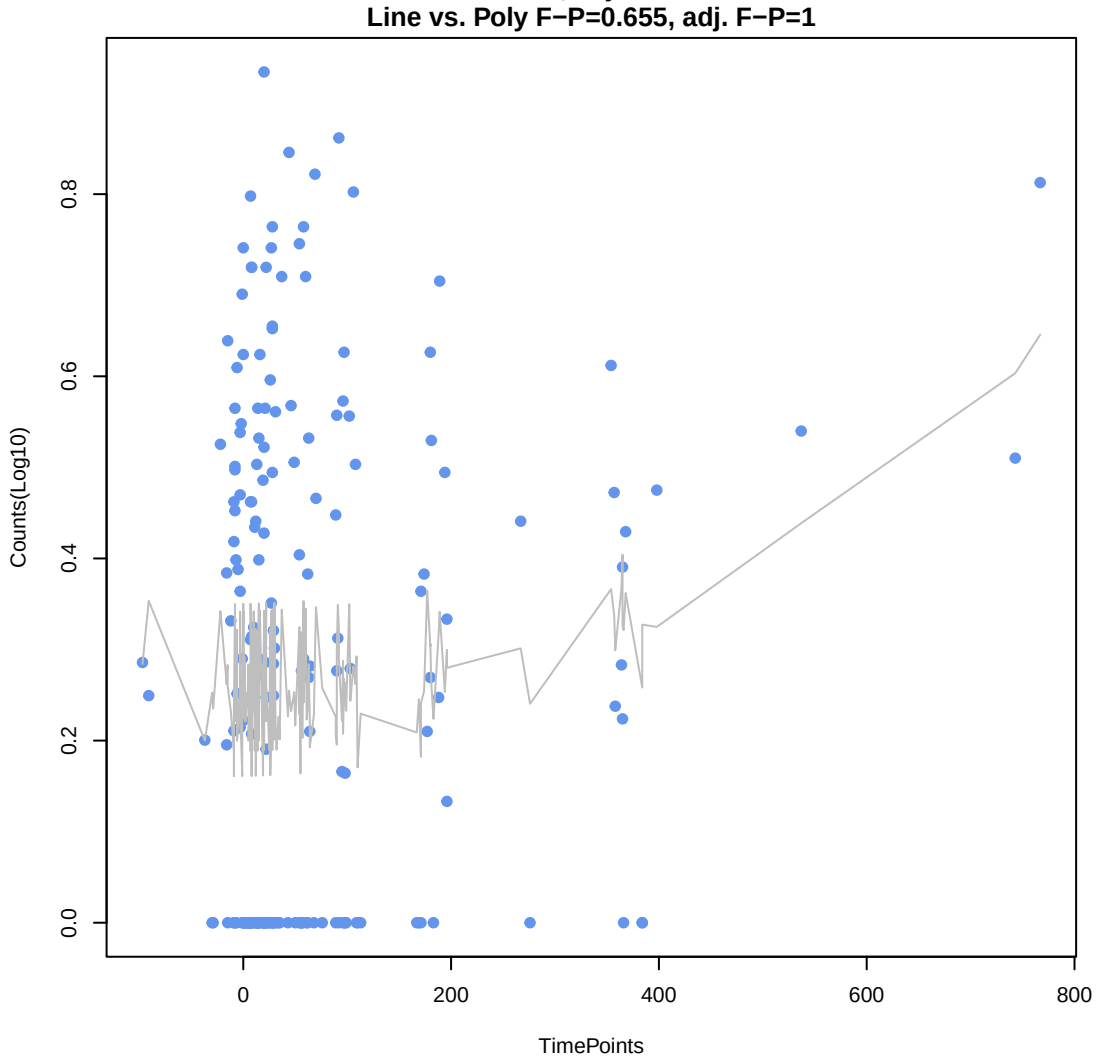
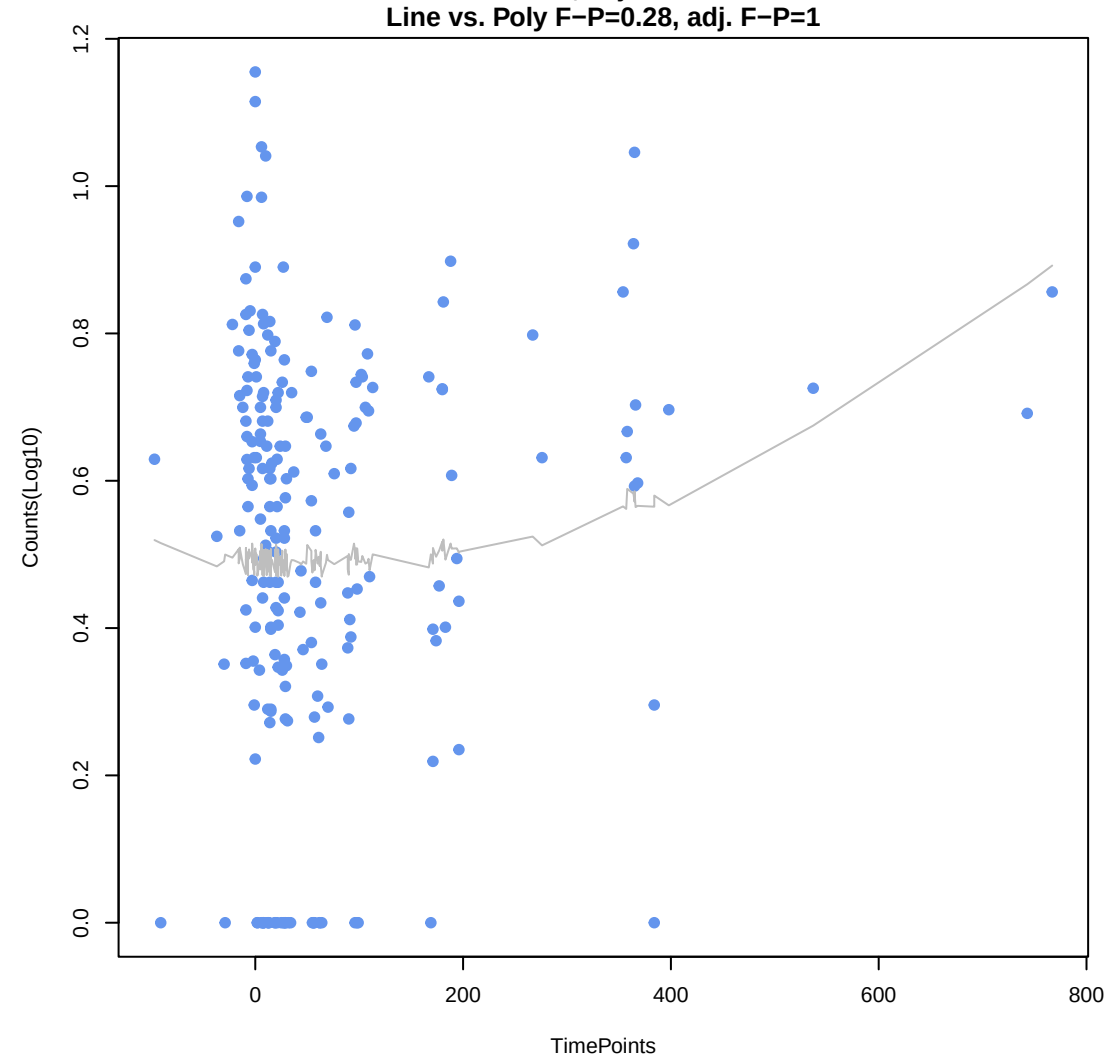
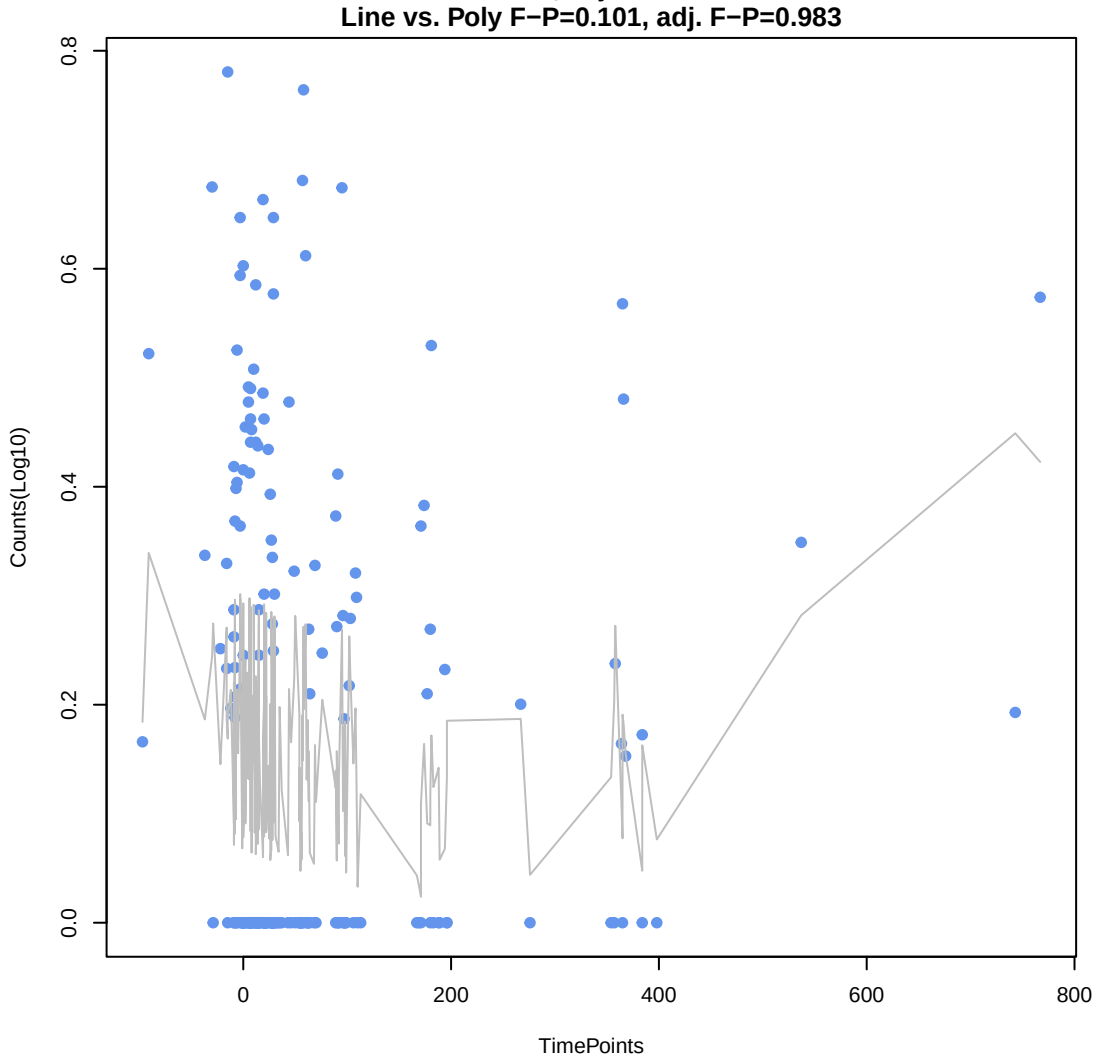
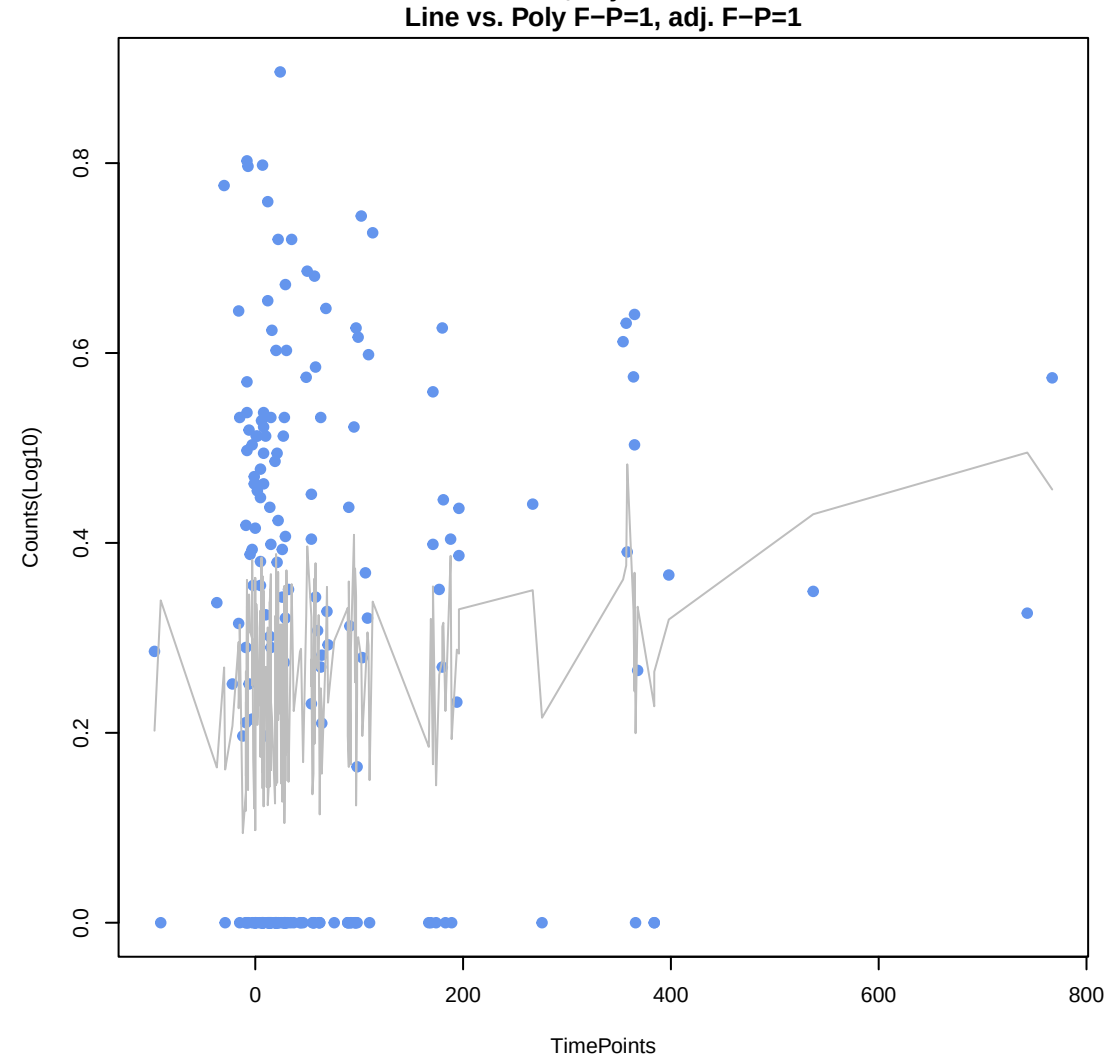

Erm 23S ribosomal RNA methyltransferase
ANOVA P=0.0037, adj. ANOVA−P=0.0962
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints

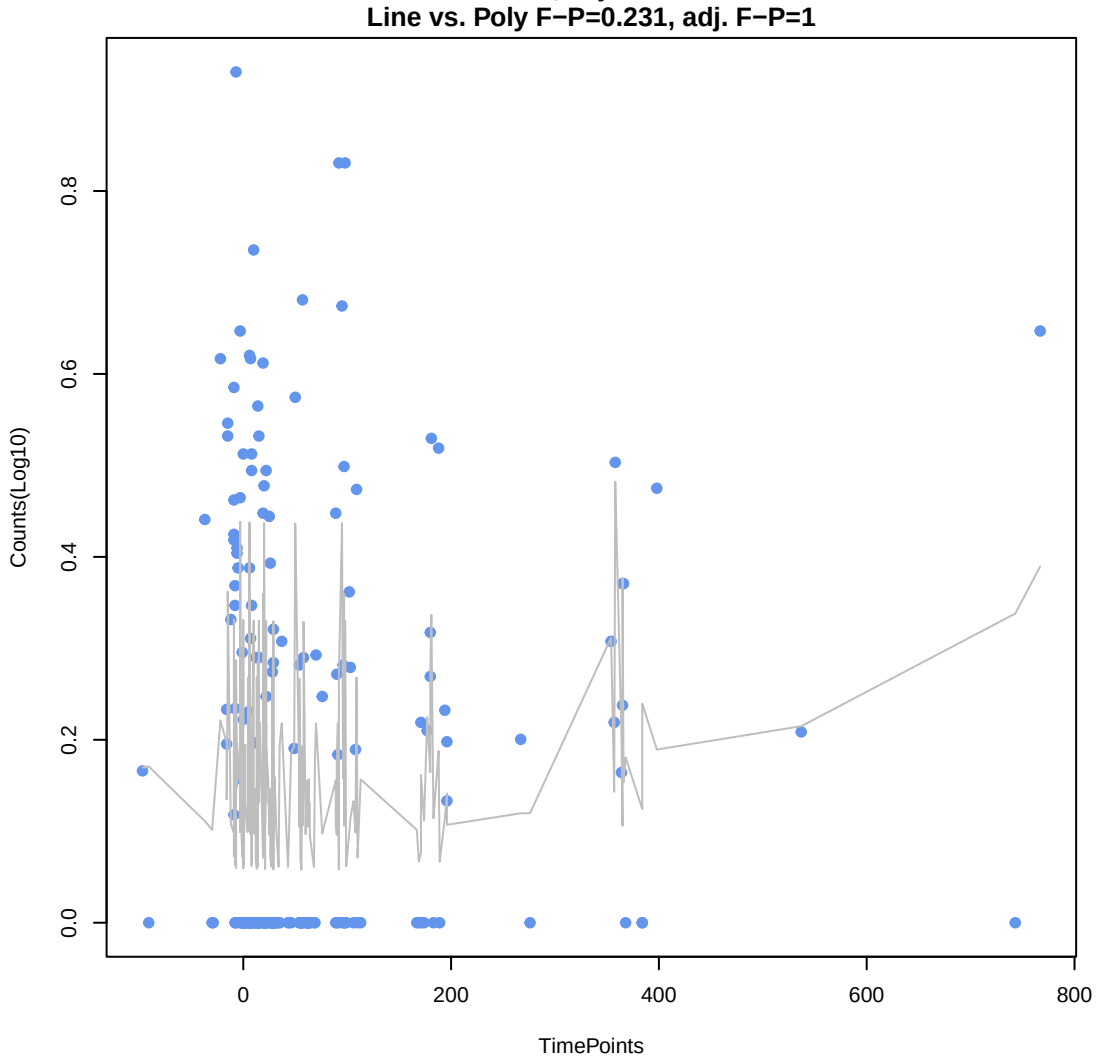

undecaprenyl pyrophosphate related proteins
ANOVA P=0.0246, adj. ANOVA−P=0.319
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints

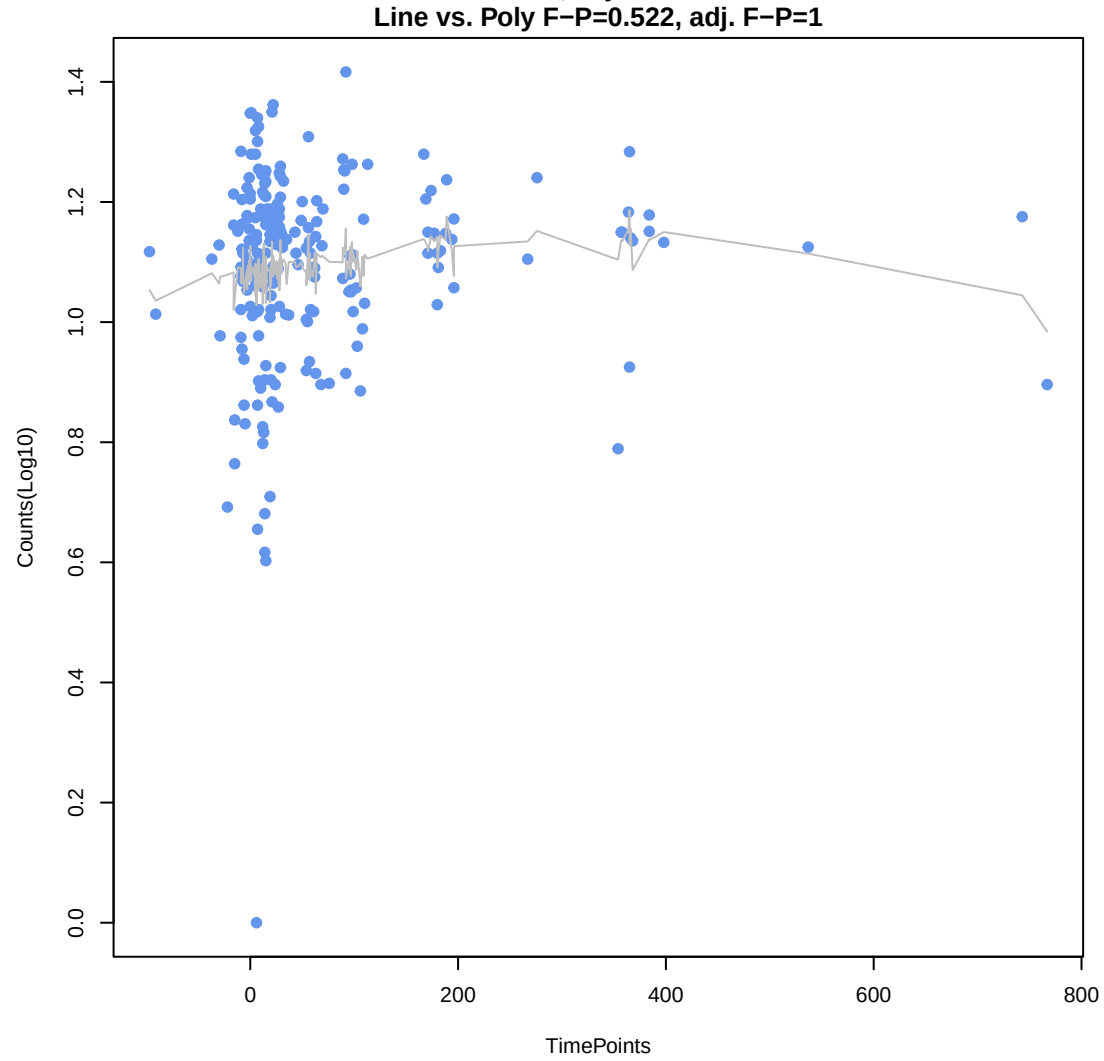

tetracycline−resistant ribosomal protection protein
ANOVA P=0.0714, adj. ANOVA−P=0.554
Line vs. Poly F−P=0.655, adj. F−P=1
Counts(Log10)
TimePoints

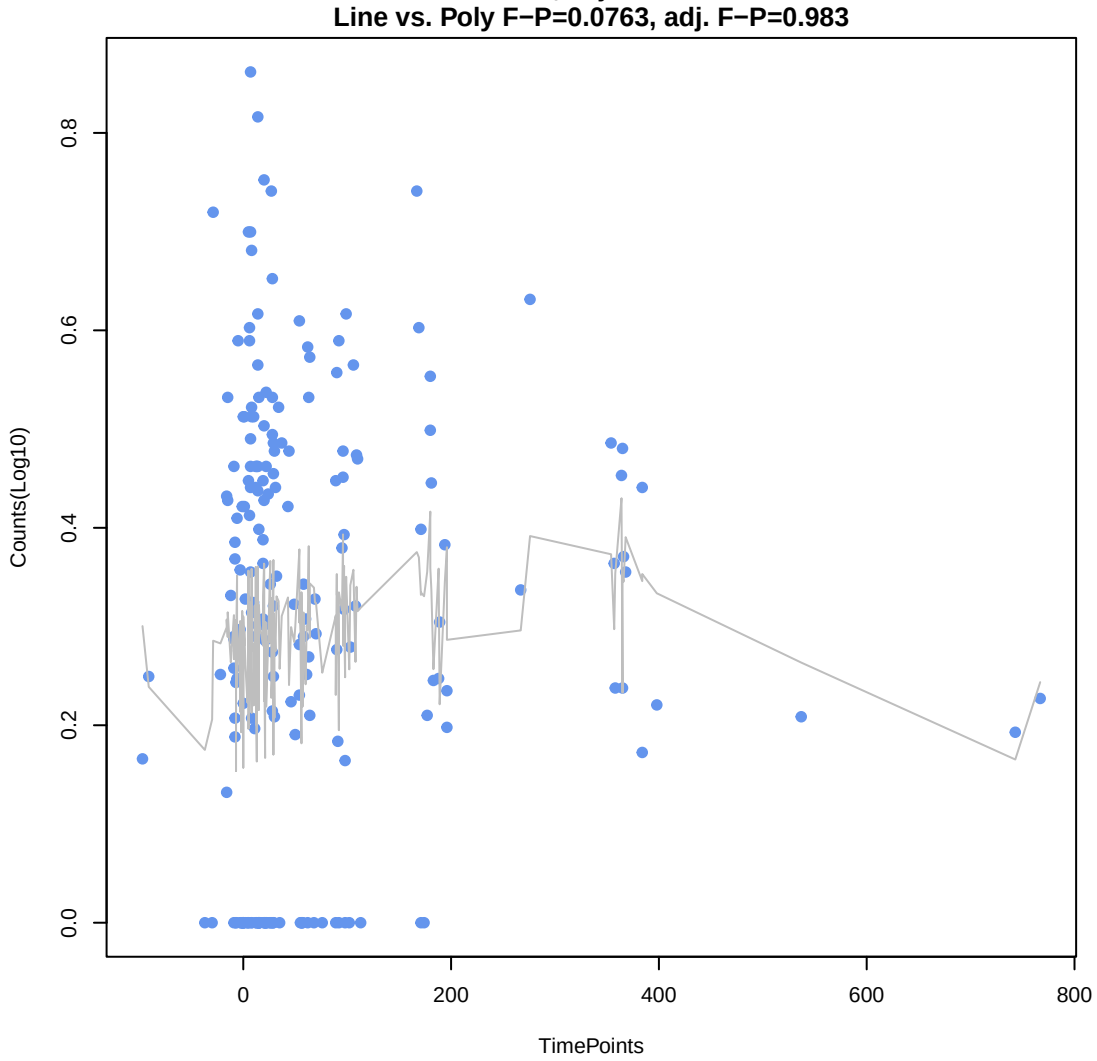

BlaB beta−lactamase
ANOVA P=0.0852, adj. ANOVA−P=0.554
Line vs. Poly F−P=0.28, adj. F−P=1
Counts(Log10)
TimePoints

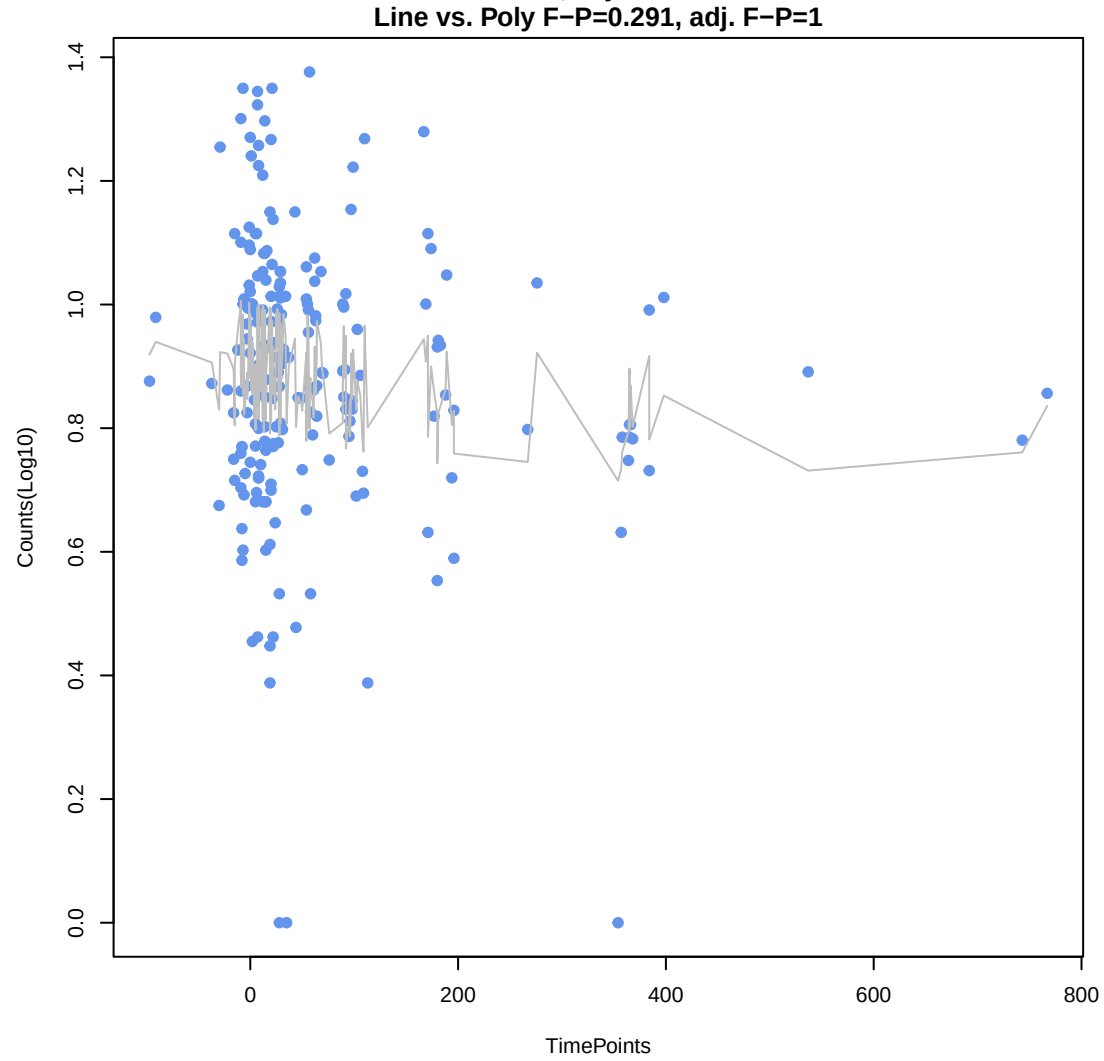

PDC beta−lactamase
ANOVA P=0.115, adj. ANOVA−P=0.596
Line vs. Poly F−P=0.101, adj. F−P=0.983
Counts(Log10)
TimePoints

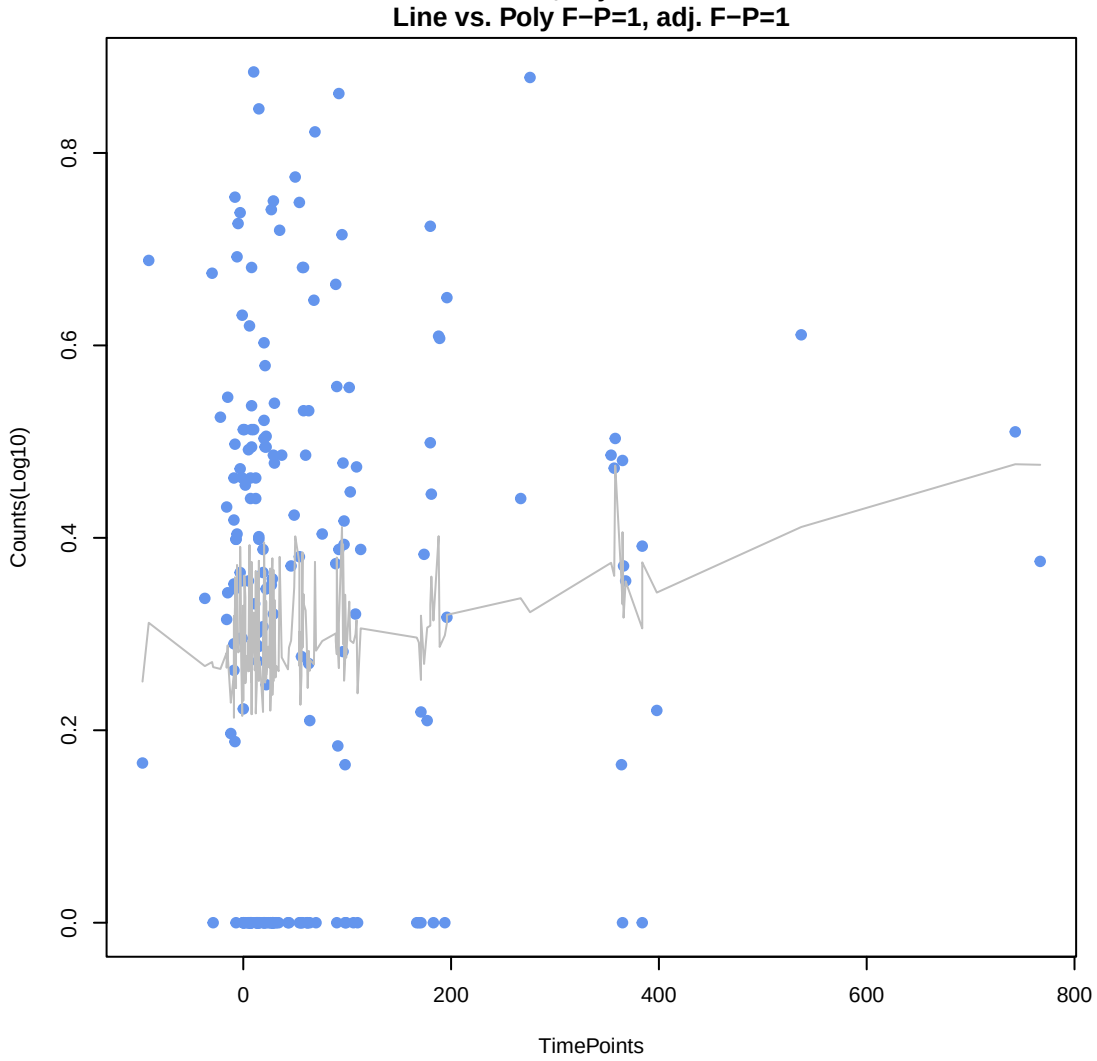

ANA beta−lactamase
ANOVA P=0.145, adj. ANOVA−P=0.605
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints

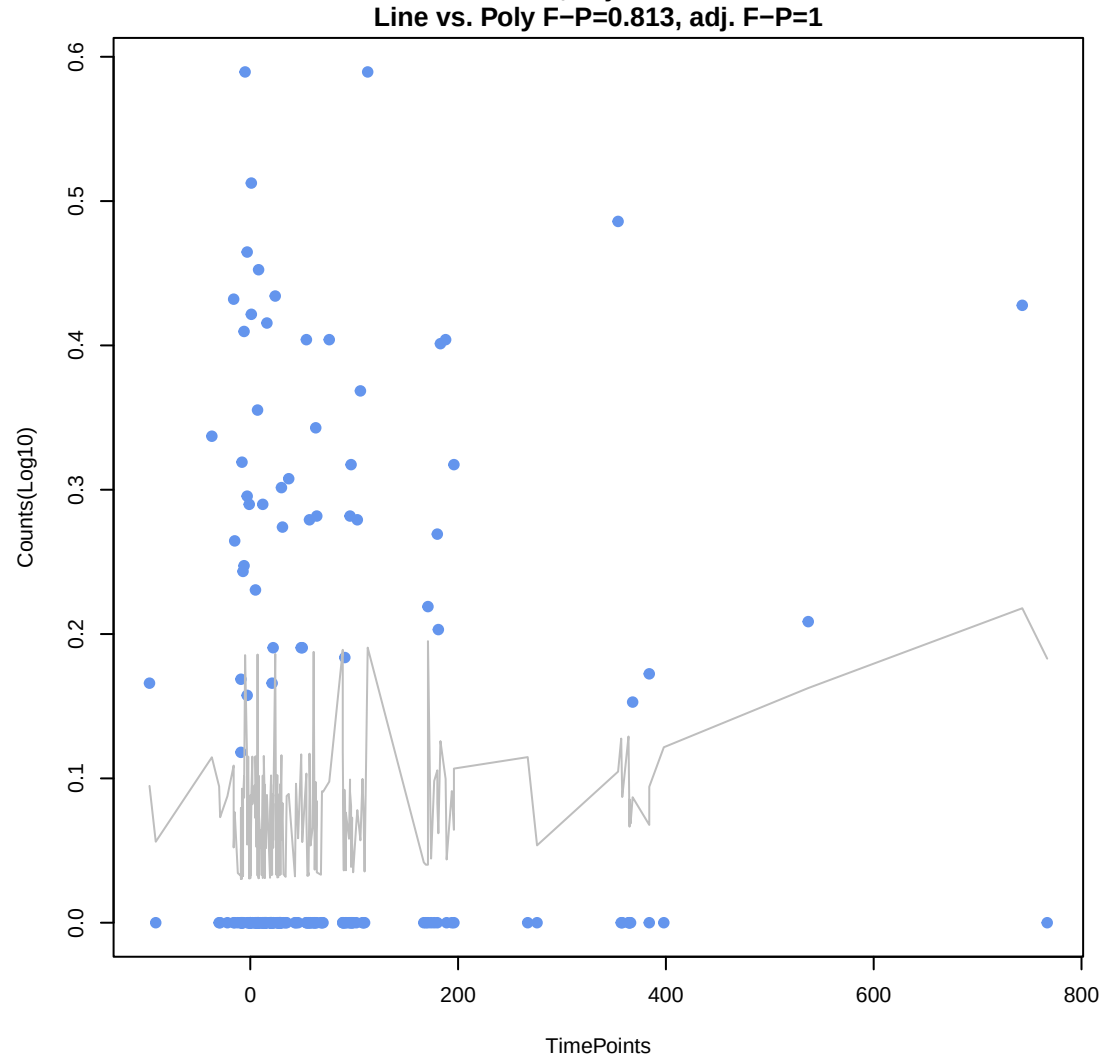

APH(6)
ANOVA P=0.164, adj. ANOVA−P=0.605
Line vs. Poly F−P=0.231, adj. F−P=1
Counts(Log10)
TimePoints


major facilitator superfamily (MFS) antibiotic efflux pump
ANOVA P=0.204, adj. ANOVA−P=0.605
Line vs. Poly F−P=0.522, adj. F−P=1
Counts(Log10)
TimePoints

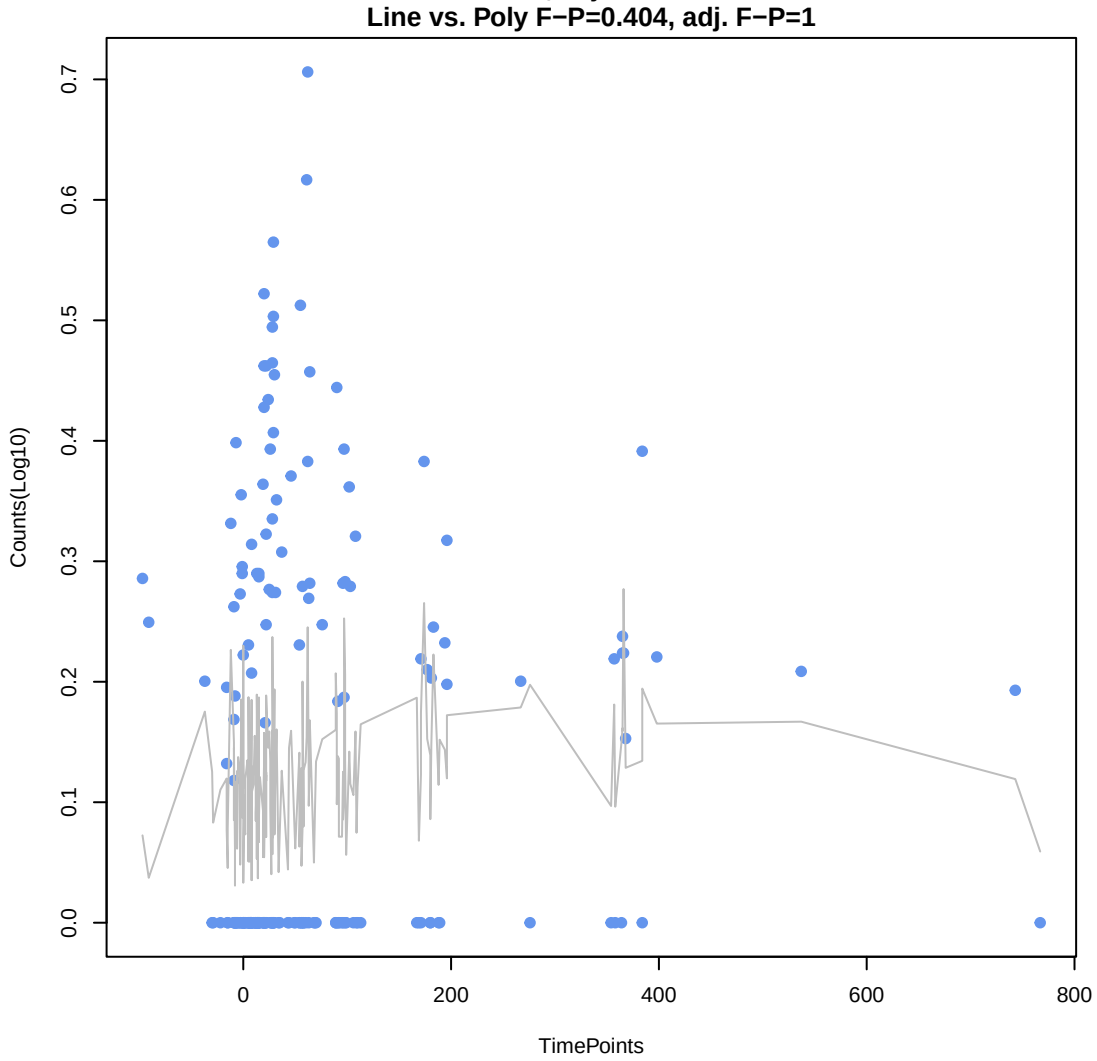

streptothricin acetyltransferase (SAT)
ANOVA P=0.218, adj. ANOVA−P=0.605
Line vs. Poly F−P=0.0763, adj. F−P=0.983
Counts(Log10)
TimePoints


ATP−binding cassette (ABC) antibiotic efflux pump
ANOVA P=0.246, adj. ANOVA−P=0.605
Line vs. Poly F−P=0.291, adj. F−P=1
Counts(Log10)
TimePoints

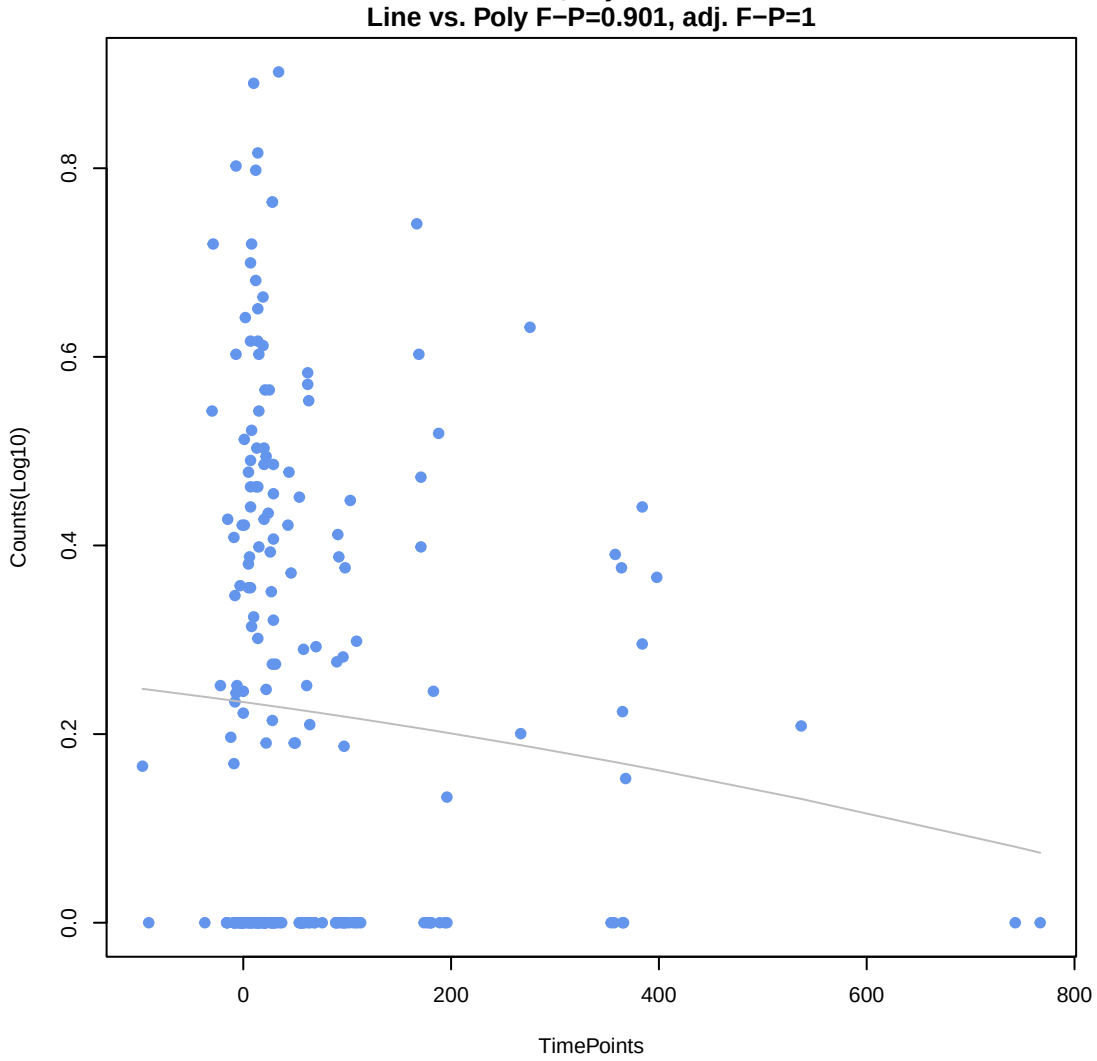

non−erm 23S ribosomal RNA methyltransferase (G748)
ANOVA P=0.256, adj. ANOVA−P=0.605
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints


SHV beta−lactamase
ANOVA P=0.317, adj. ANOVA−P=0.686
Line vs. Poly F−P=0.813, adj. F−P=1
Counts(Log10)
TimePoints

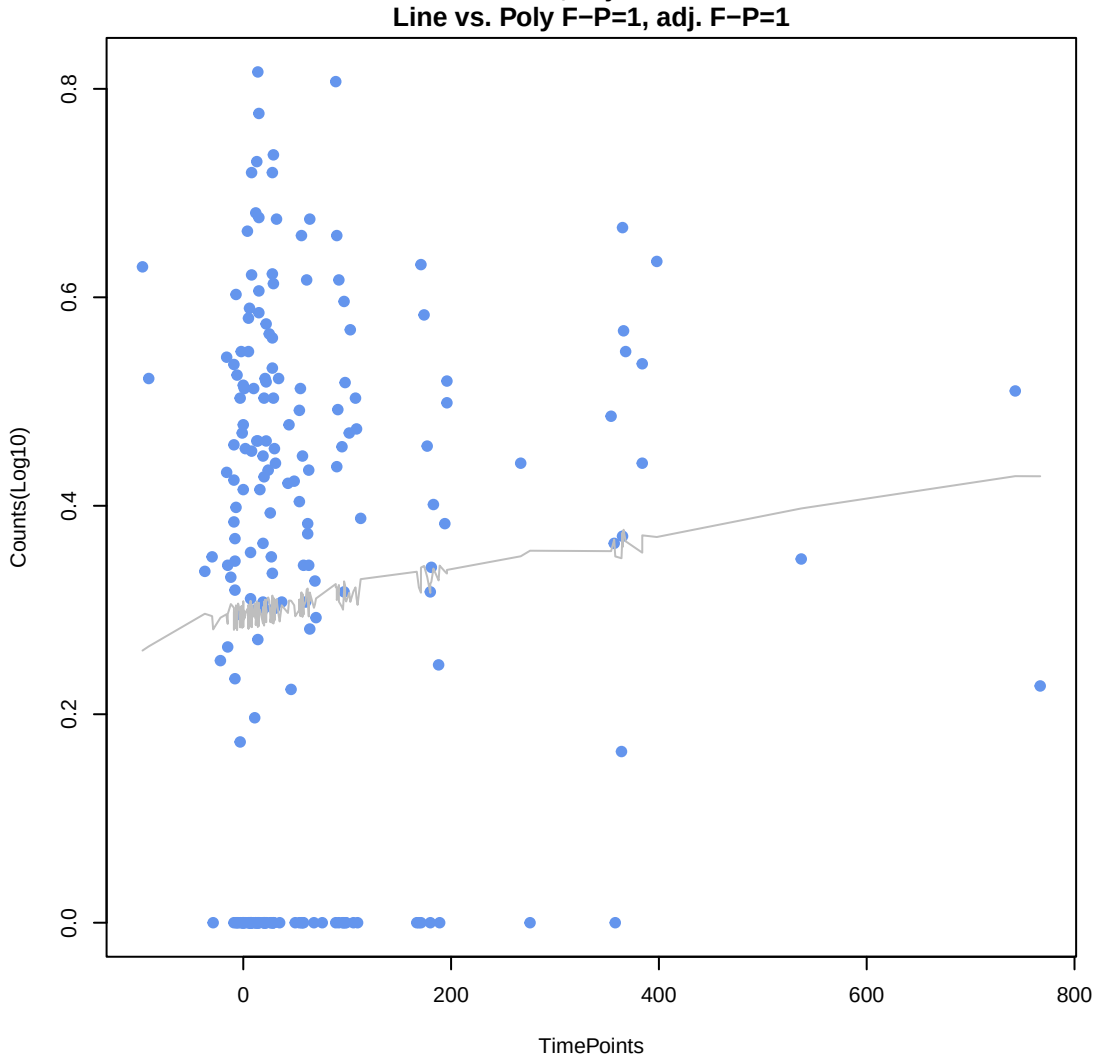
al Porin with reduced permeability to beta−lactams;resistance−nodulation−cell division (RN
ANOVA P=0.4, adj. ANOVA−P=0.77
Line vs. Poly F−P=0.404, adj. F−P=1



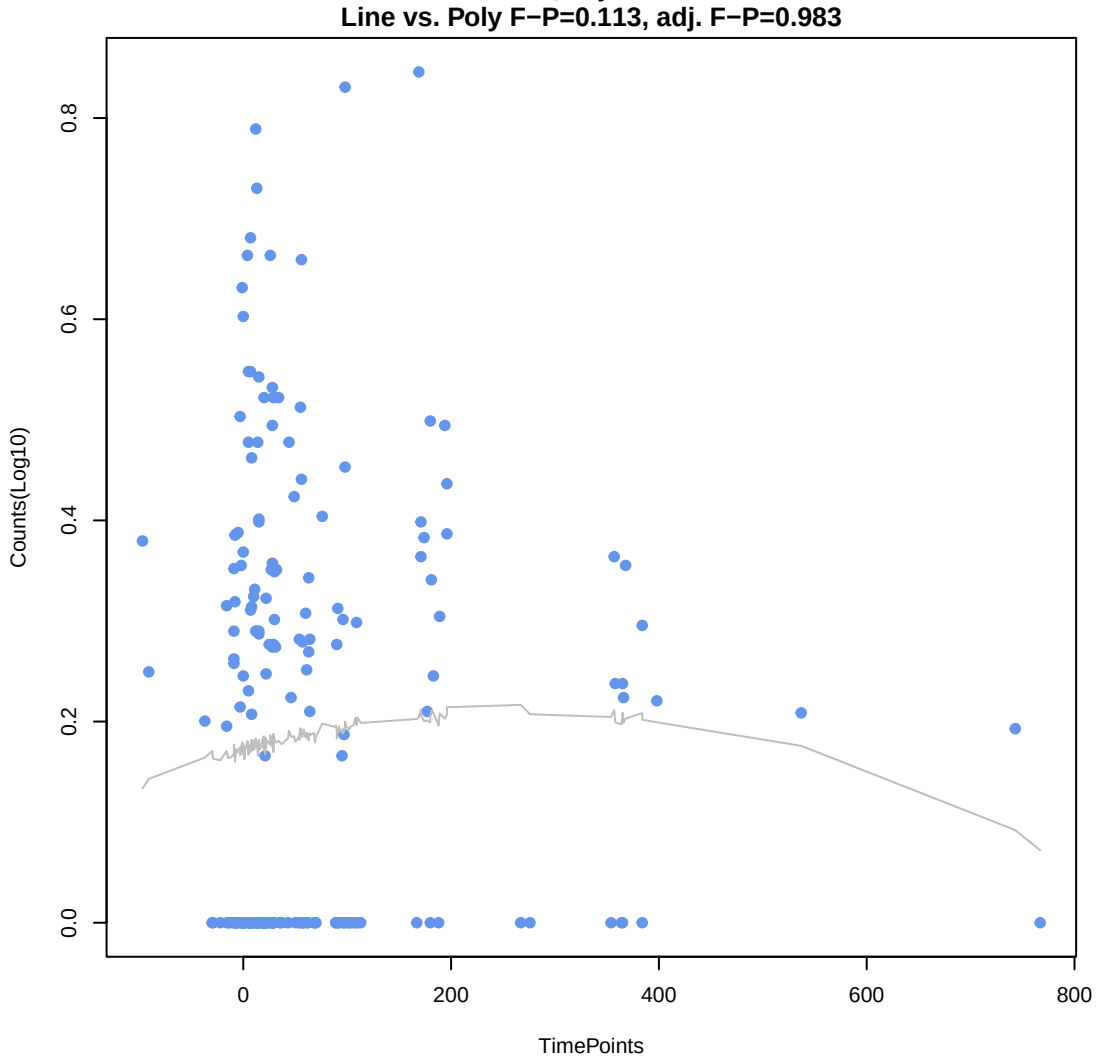

msr−type ABC−F protein
ANOVA P=0.434, adj. ANOVA−P=0.77
Line vs. Poly F−P=0.901, adj. F−P=1



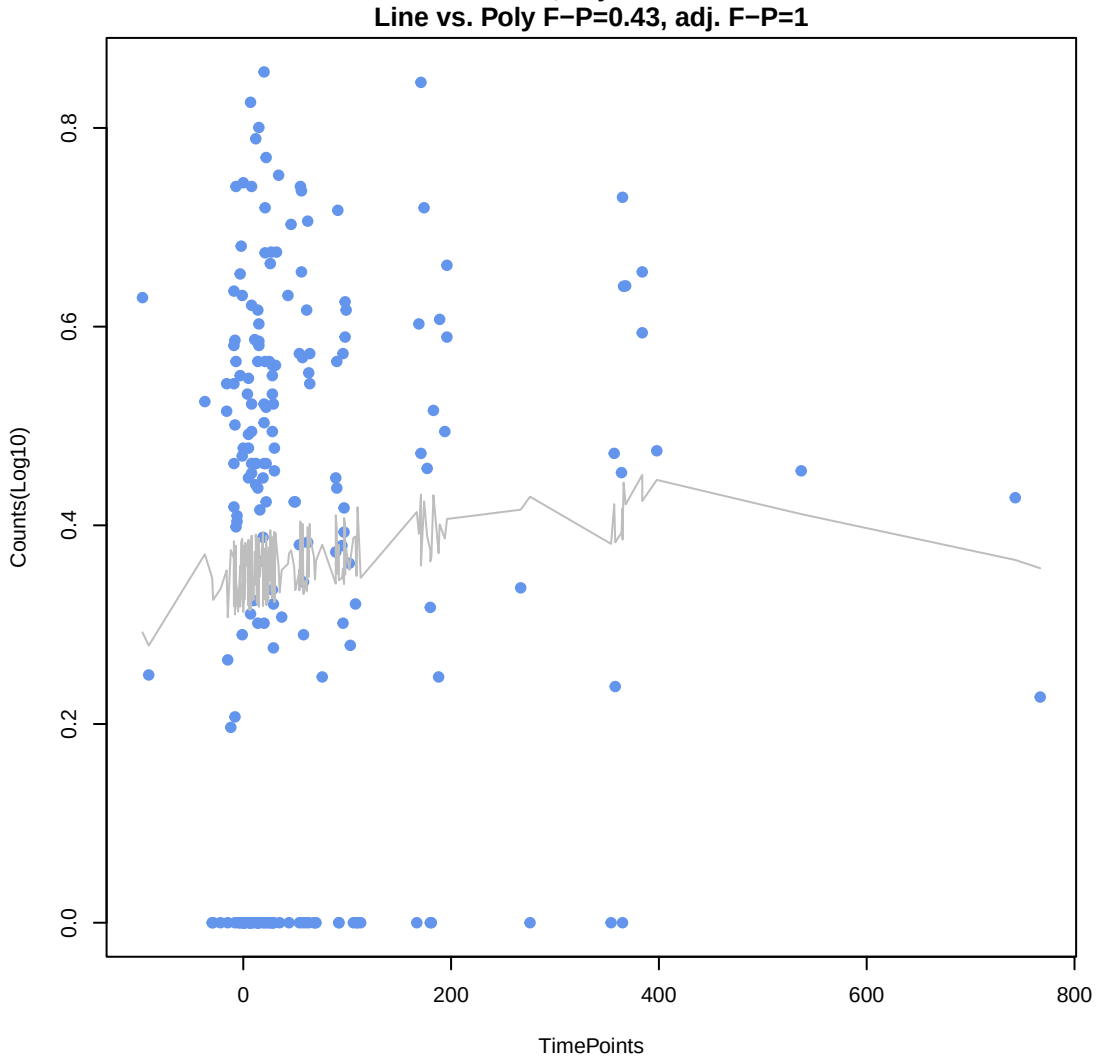

pmr phosphoethanolamine transferase
ANOVA P=0.444, adj. ANOVA−P=0.77
Line vs. Poly F−P=1, adj. F−P=1



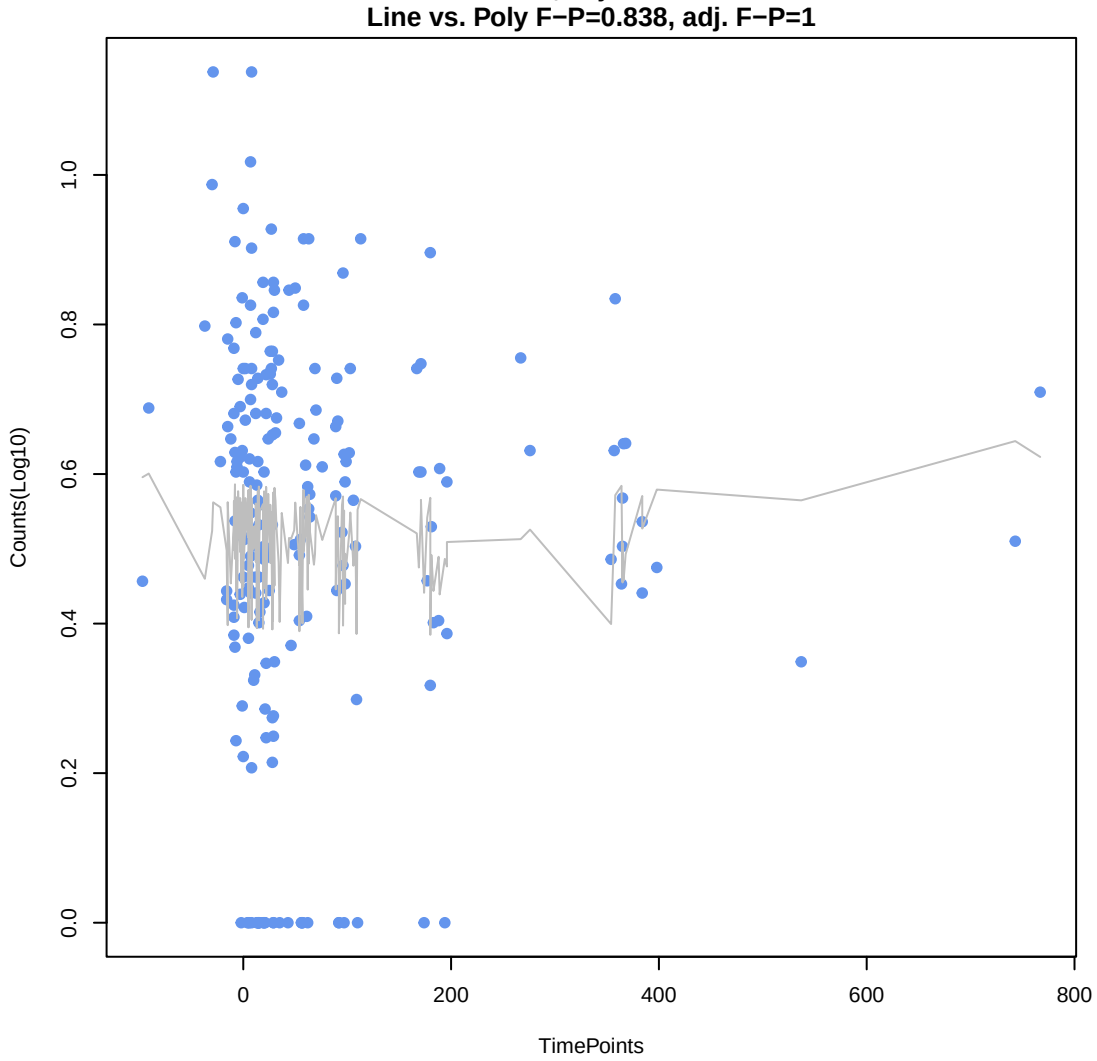

ic efflux pump;major facilitator superfamily (MFS) antibiotic efflux pump;resistance−nodula
ANOVA P=0.562, adj. ANOVA−P=0.878
Line vs. Poly F−P=0.113, adj. F−P=0.983

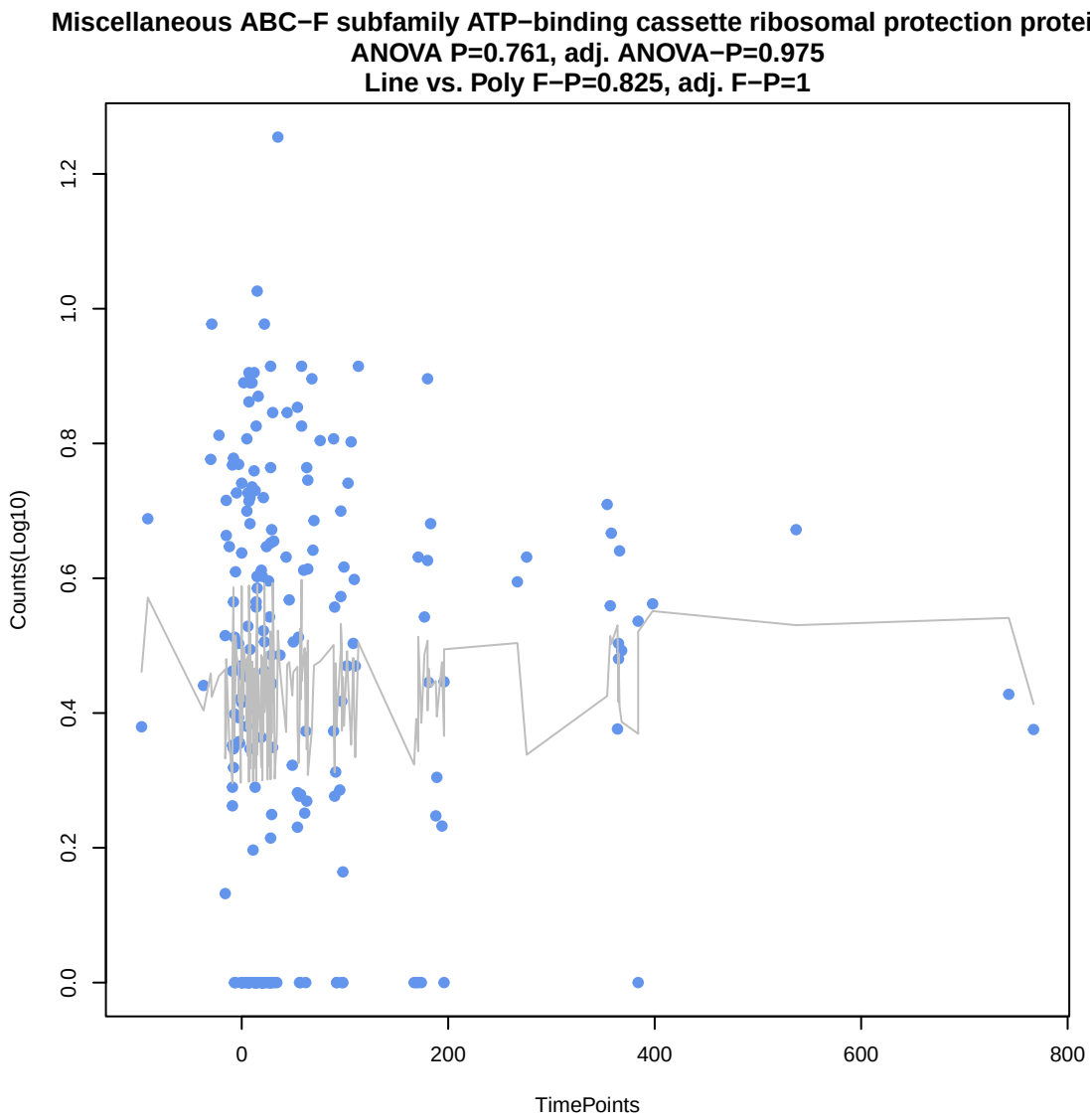


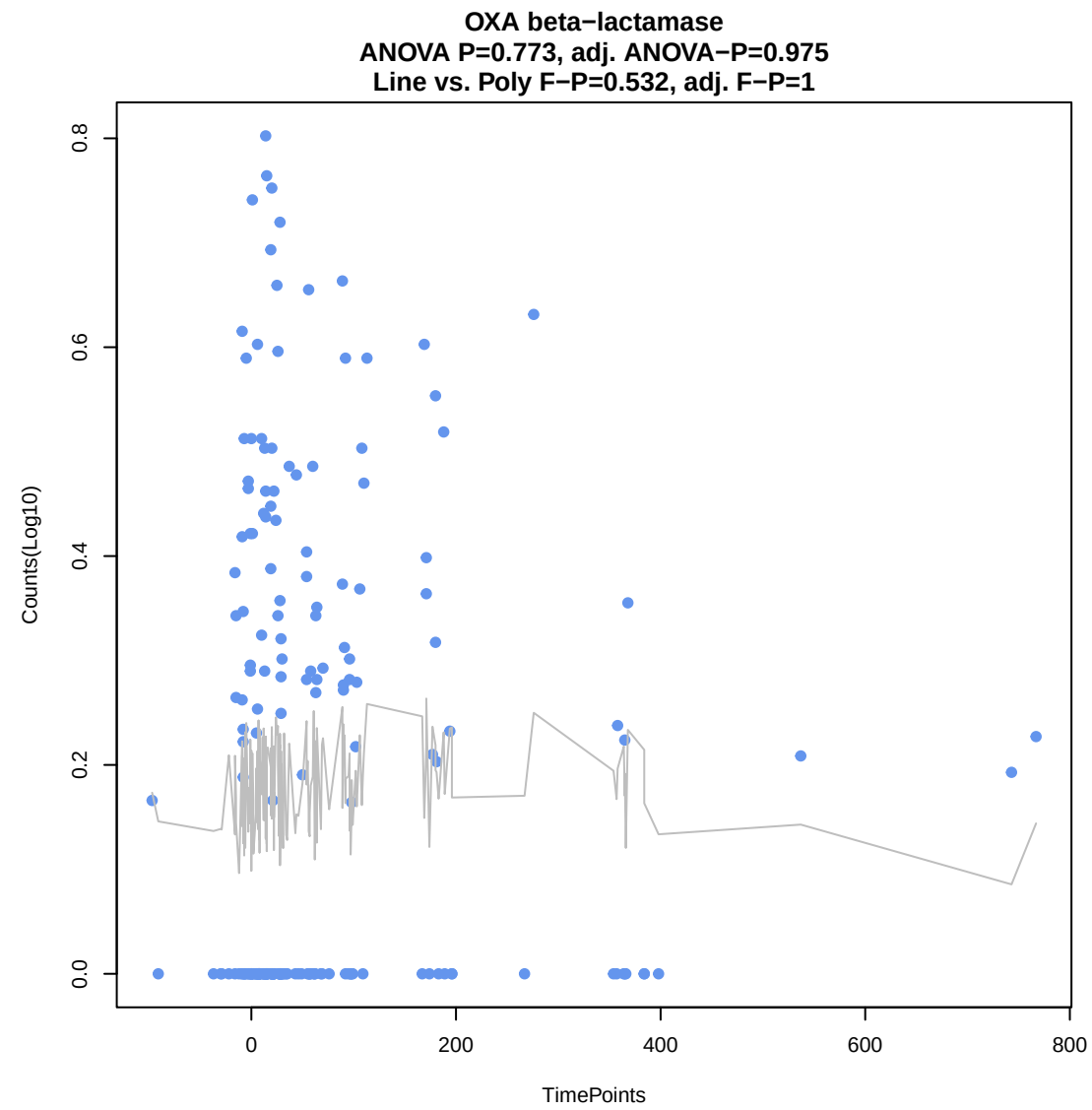

itator superfamily (MFS) antibiotic efflux pump;resistance−nodulation−cell division (RND) a
ANOVA P=0.574, adj. ANOVA−P=0.878
Line vs. Poly F−P=0.43, adj. F−P=1

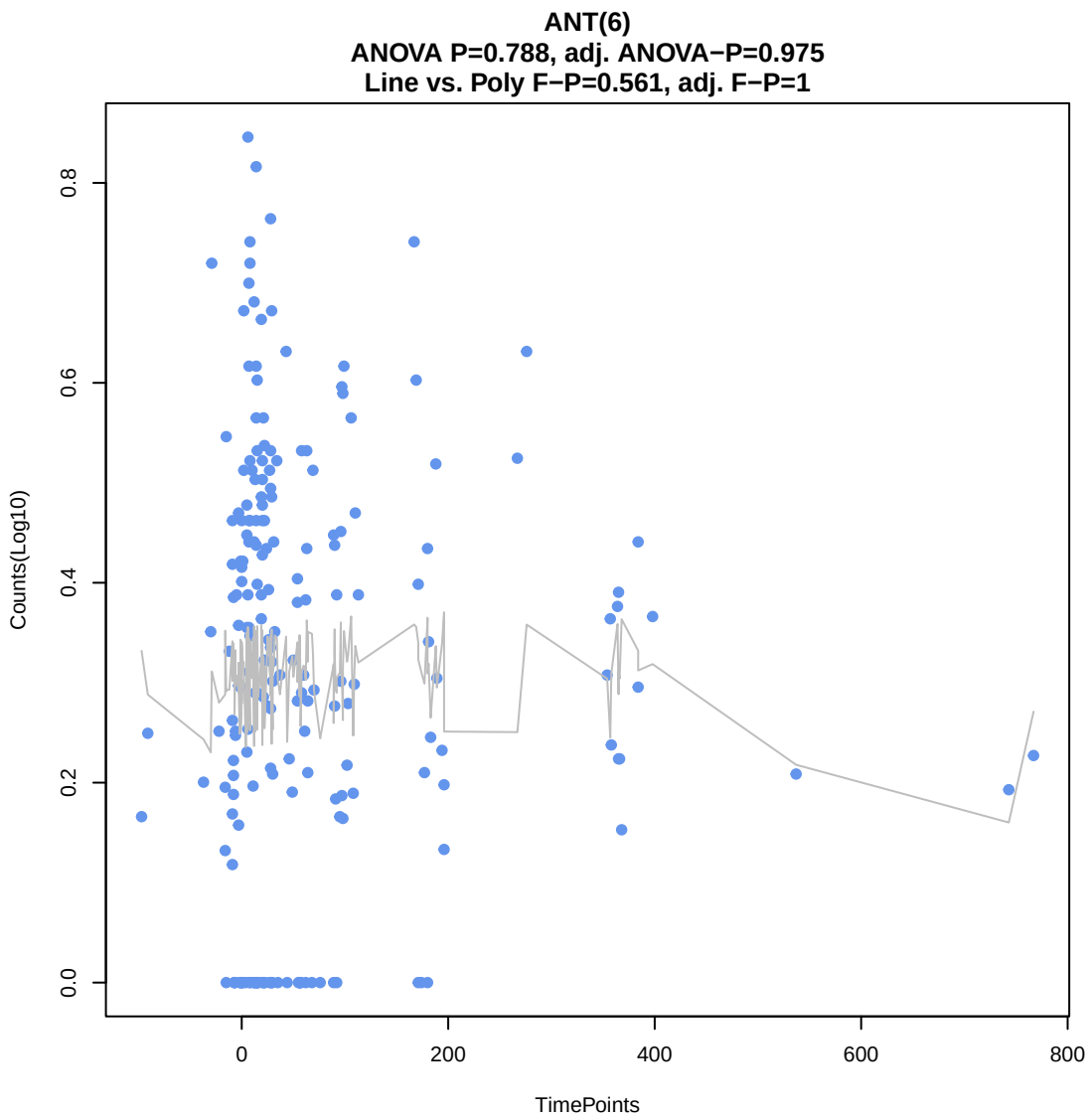


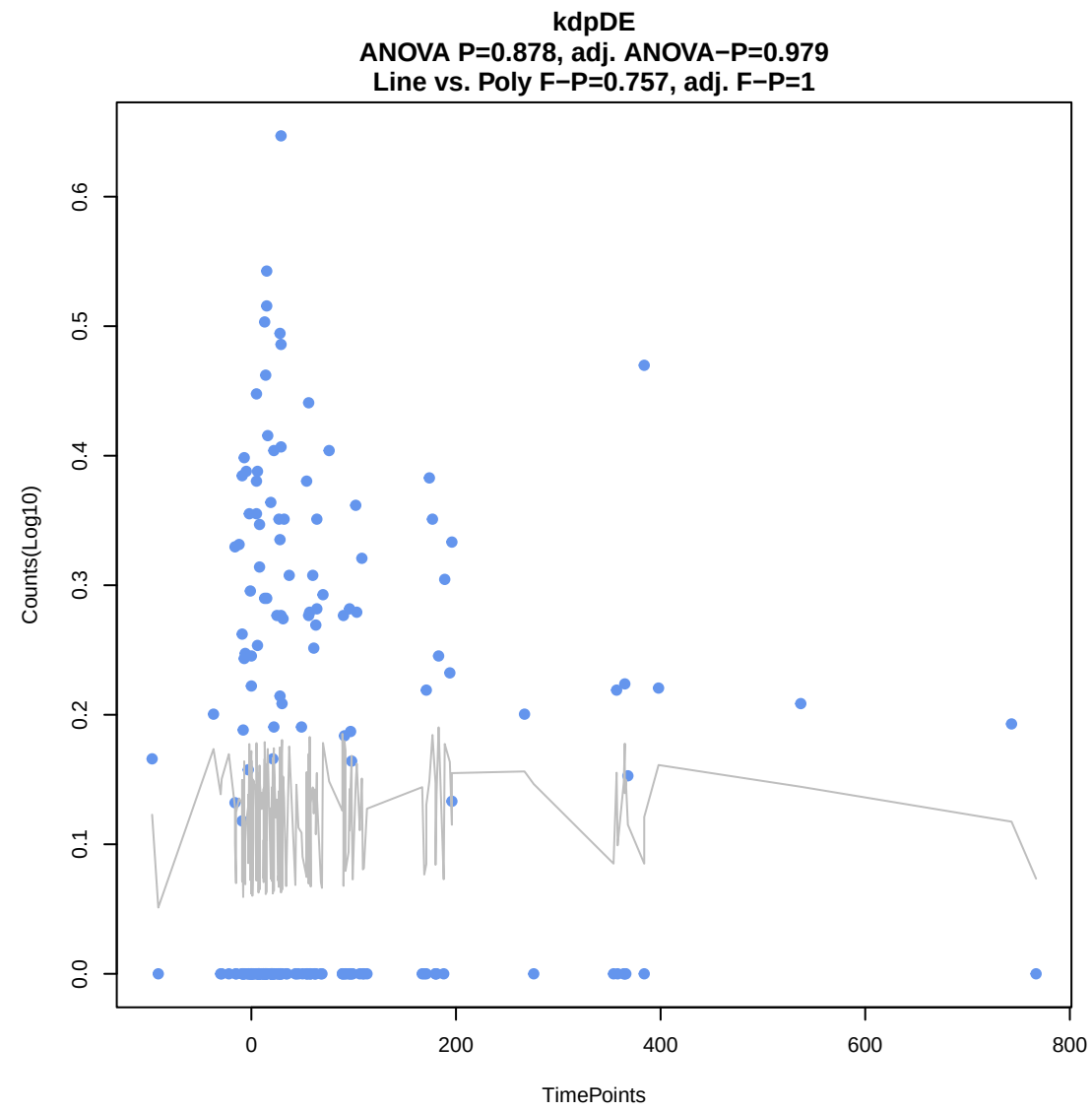

Van ligase;glycopeptide resistance gene cluster
ANOVA P=0.743, adj. ANOVA−P=0.975
Line vs. Poly F−P=0.838, adj. F−P=1

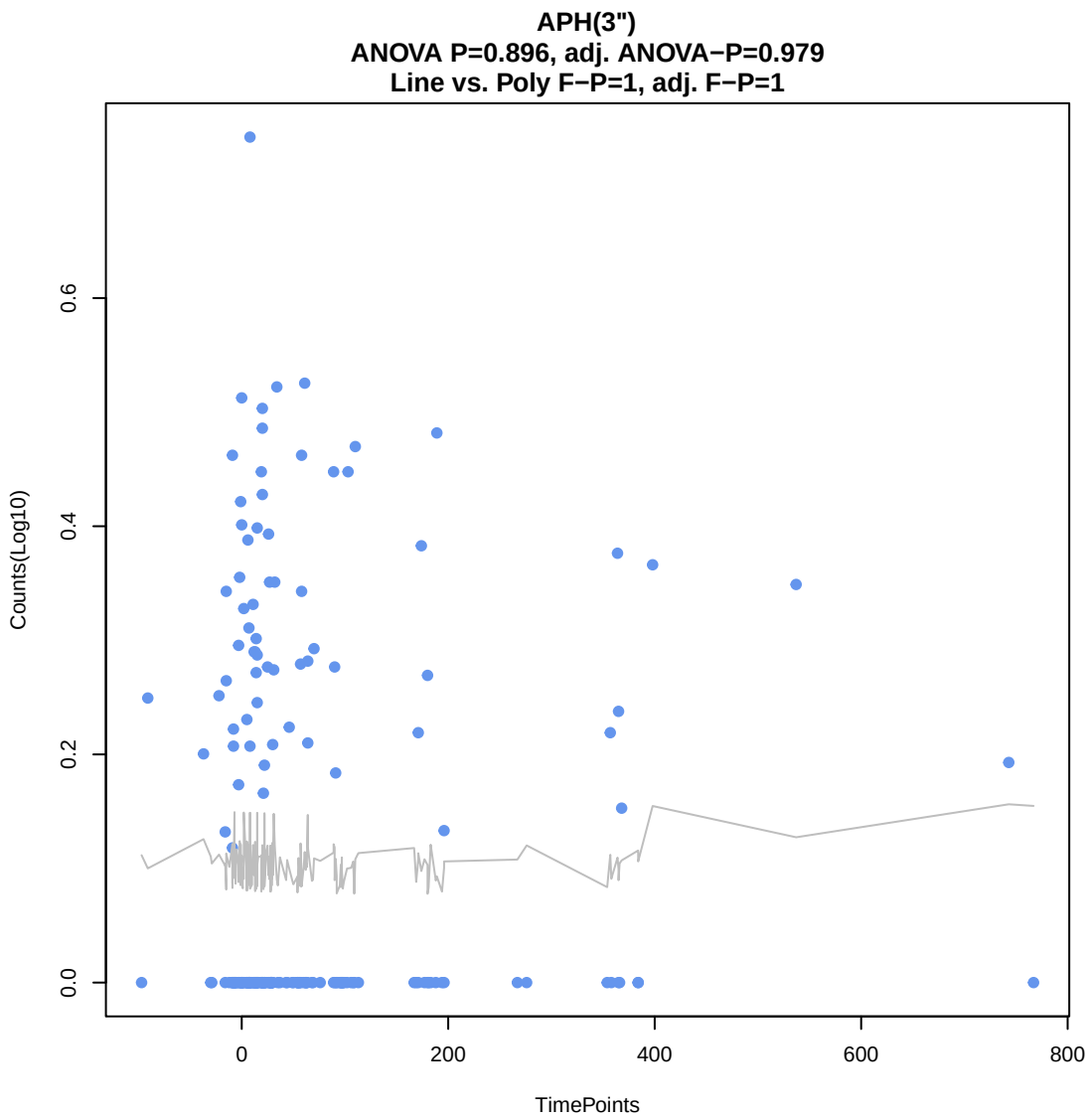


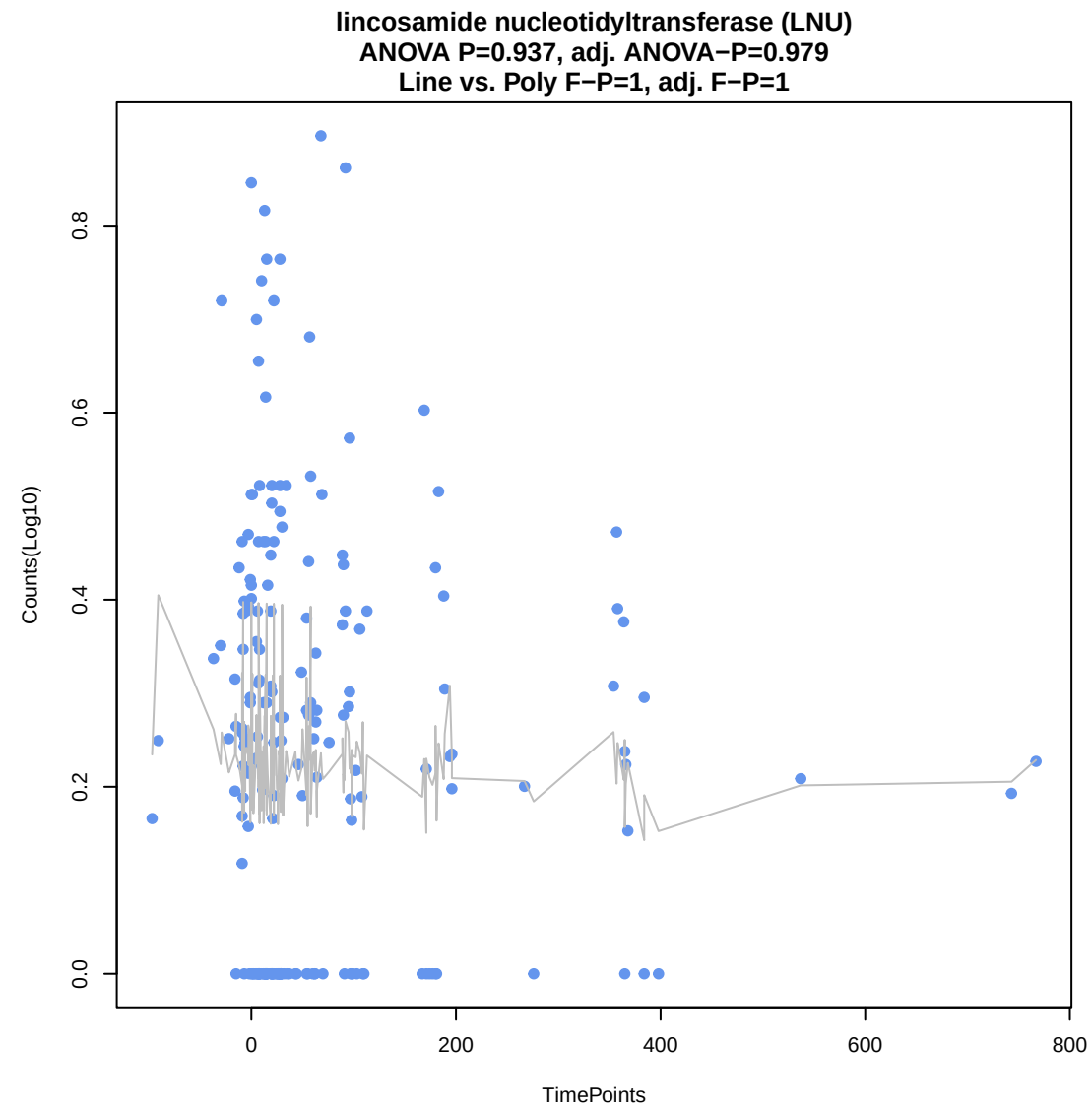

Miscellaneous ABC−F subfamily ATP−binding cassette ribosomal protection protei
ANOVA P=0.761, adj. ANOVA−P=0.975
Line vs. Poly F−P=0.825, adj. F−P=1
Counts(Log10)
TimePoints

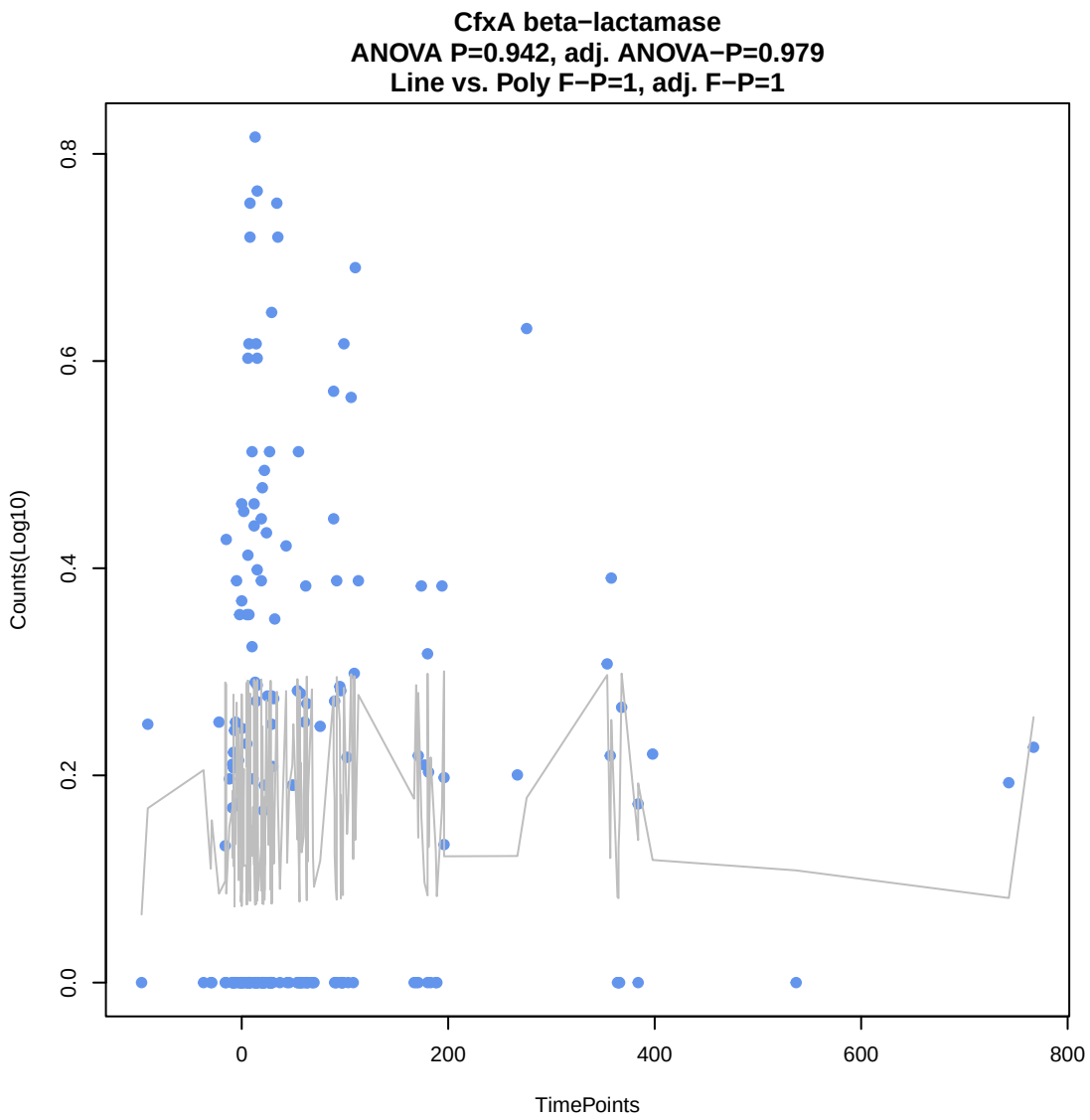

OXA beta−lactamase
ANOVA P=0.773, adj. ANOVA−P=0.975
Line vs. Poly F−P=0.532, adj. F−P=1
Counts(Log10)
TimePoints

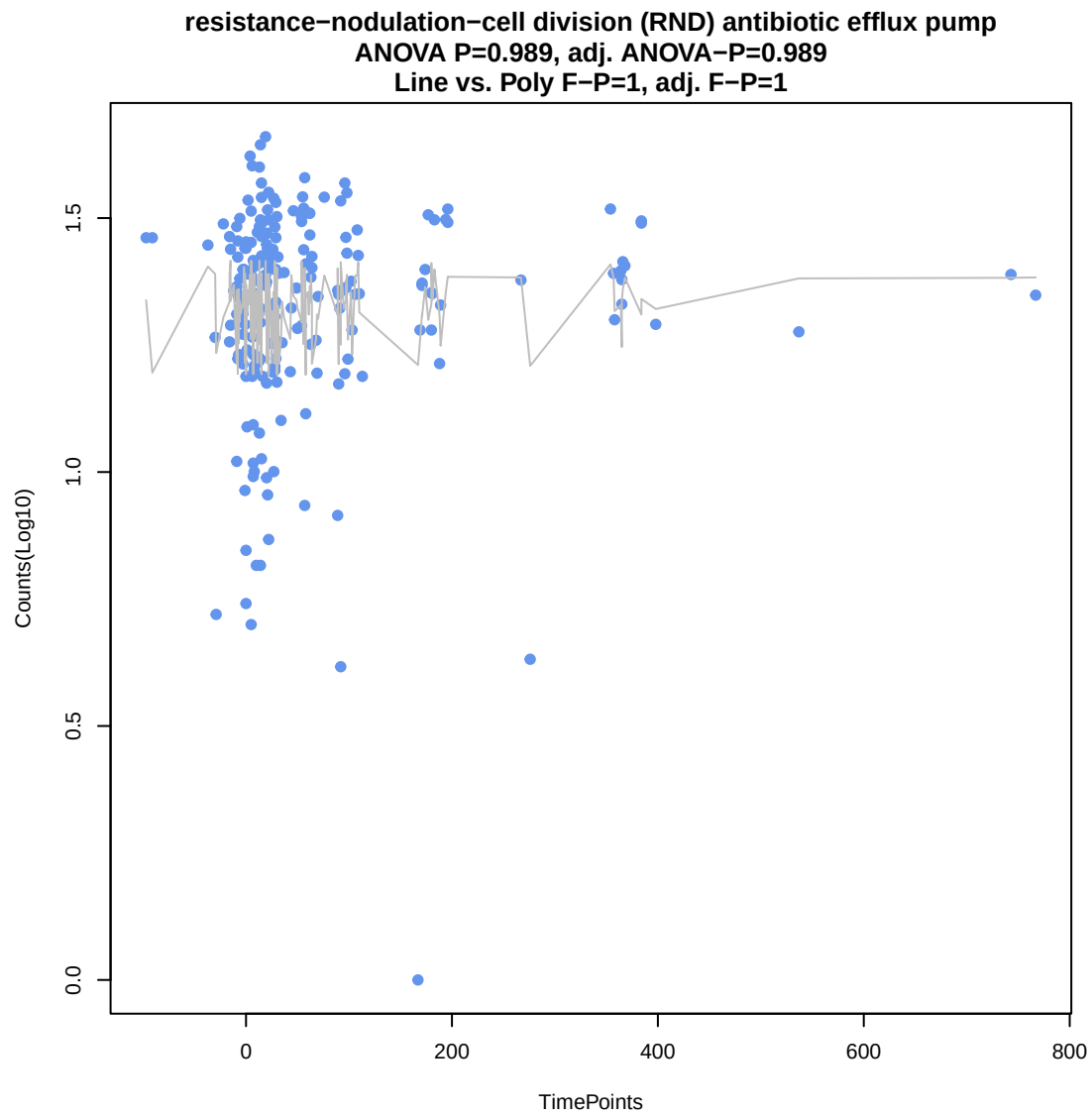

ANT(6)
ANOVA P=0.788, adj. ANOVA−P=0.975
Line vs. Poly F−P=0.561, adj. F−P=1
Counts(Log10)
TimePoints


kdpDE
ANOVA P=0.878, adj. ANOVA−P=0.979
Line vs. Poly F−P=0.757, adj. F−P=1
Counts(Log10)
TimePoints


APH(3")
ANOVA P=0.896, adj. ANOVA−P=0.979
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints


lincosamide nucleotidyltransferase (LNU)
ANOVA P=0.937, adj. ANOVA−P=0.979
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints


CfxA beta−lactamase
ANOVA P=0.942, adj. ANOVA−P=0.979
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
0.0
0.2
0.4
0.6
0.8
0
200
400
600
800
TimePoints


resistance−nodulation−cell division (RND) antibiotic efflux pump
ANOVA P=0.989, adj. ANOVA−P=0.989
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
0.0
0.5
1.0
1.5
0
200
400
600
800
TimePoints