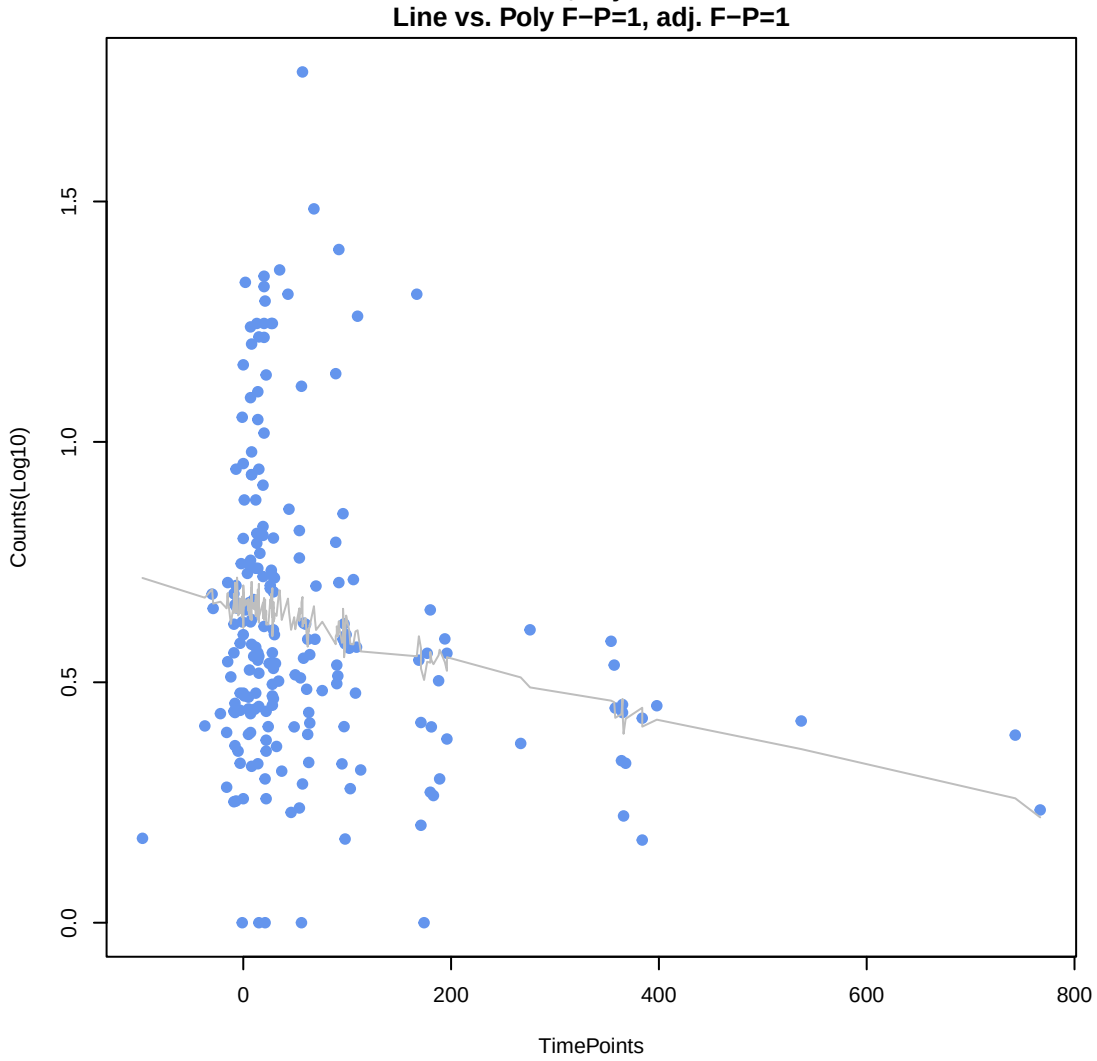
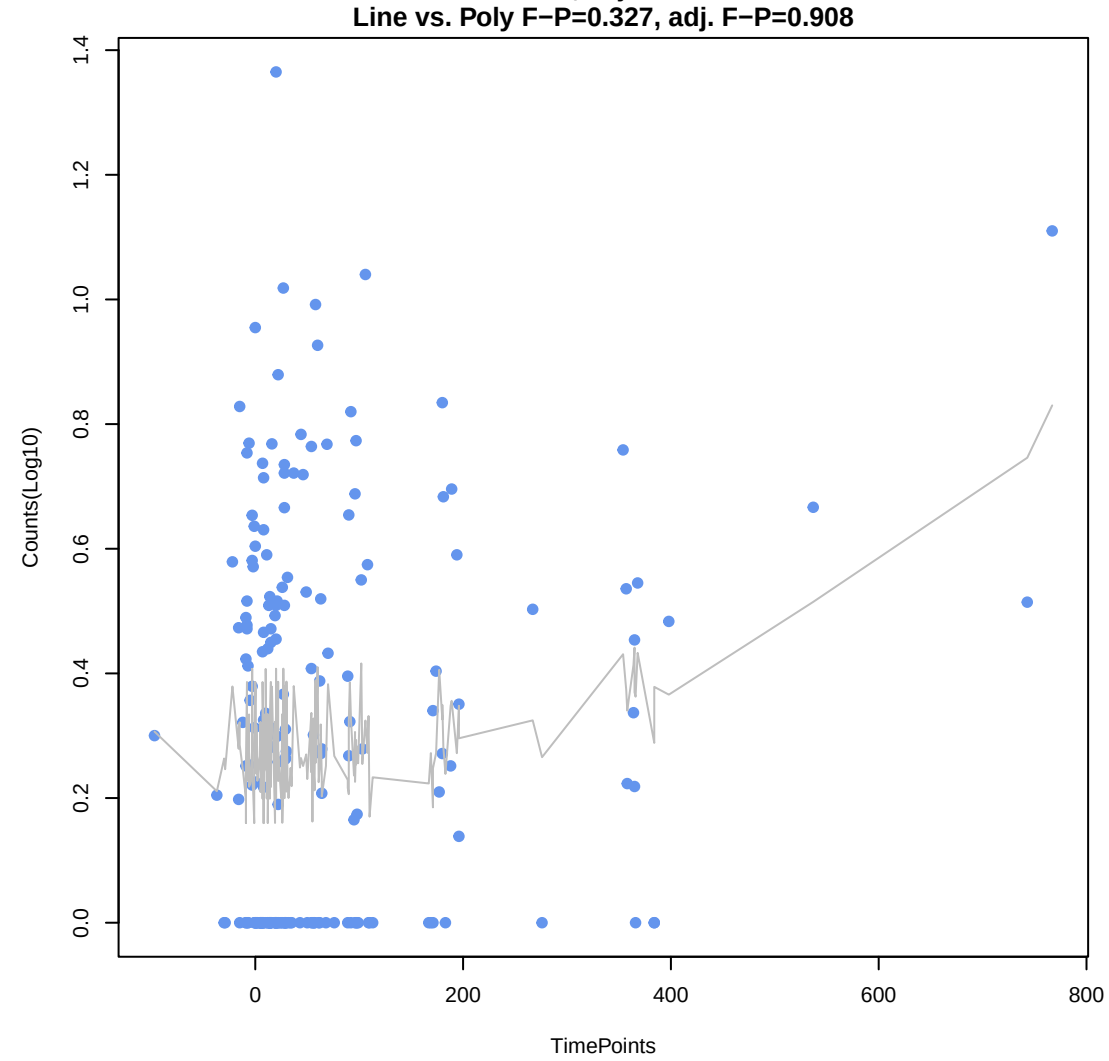
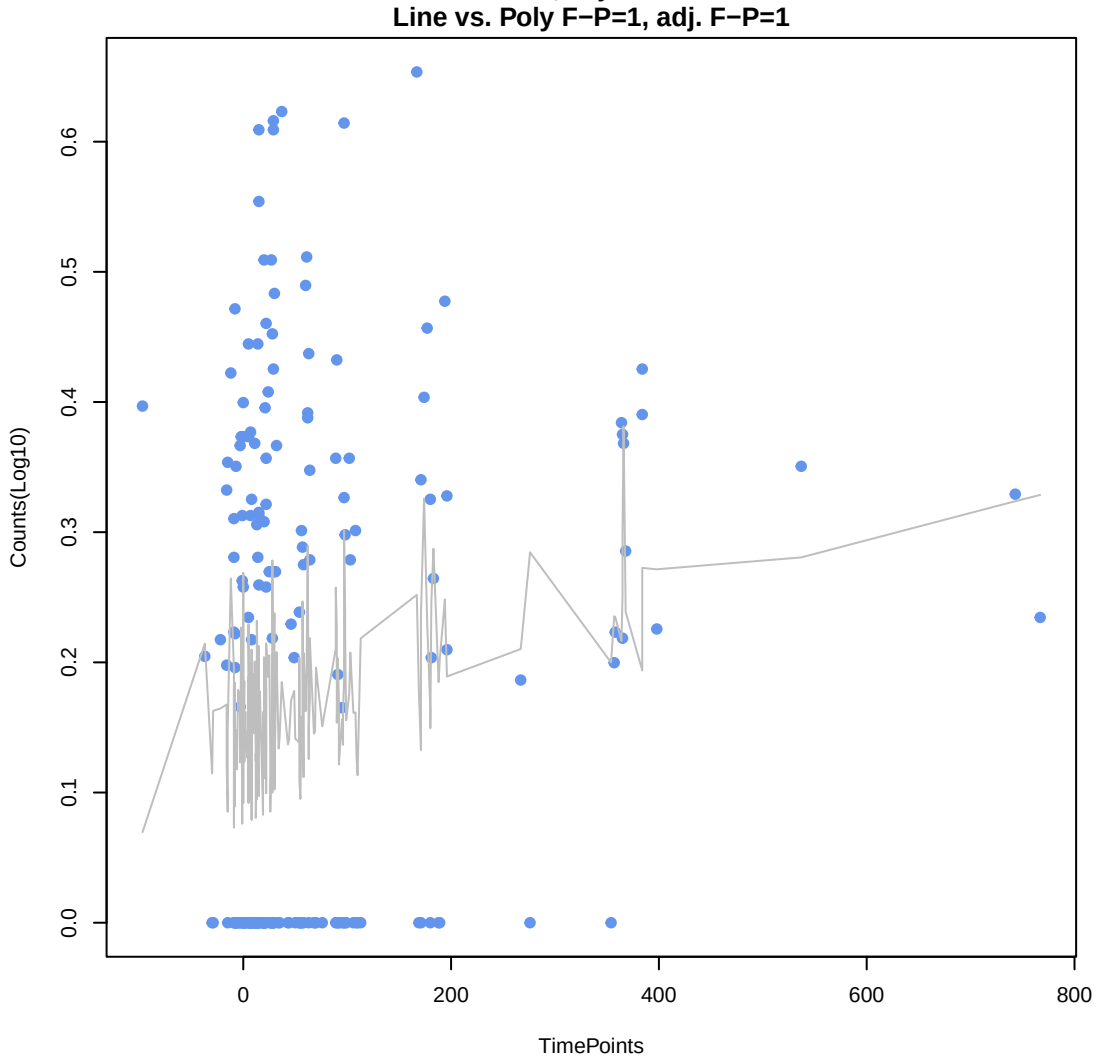
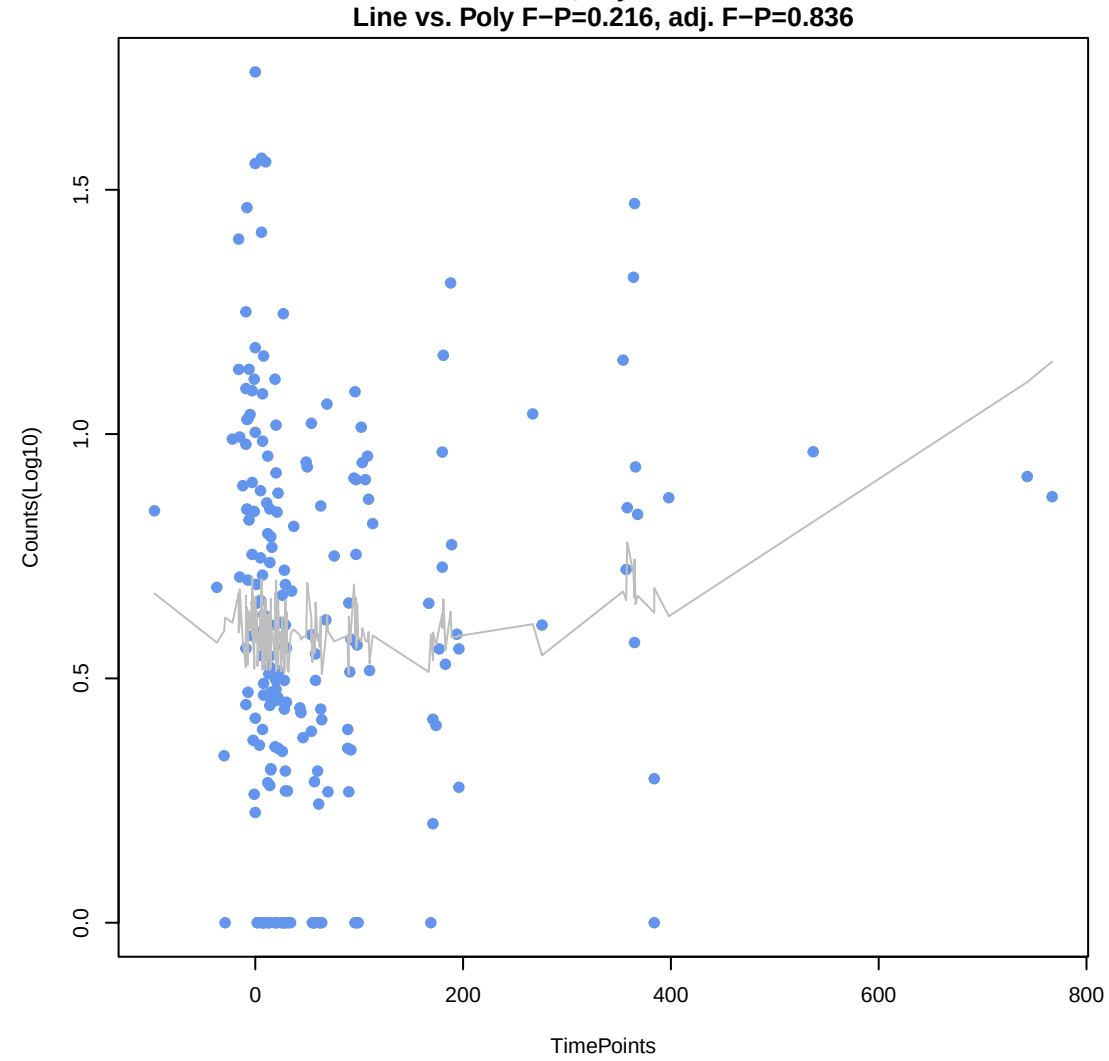
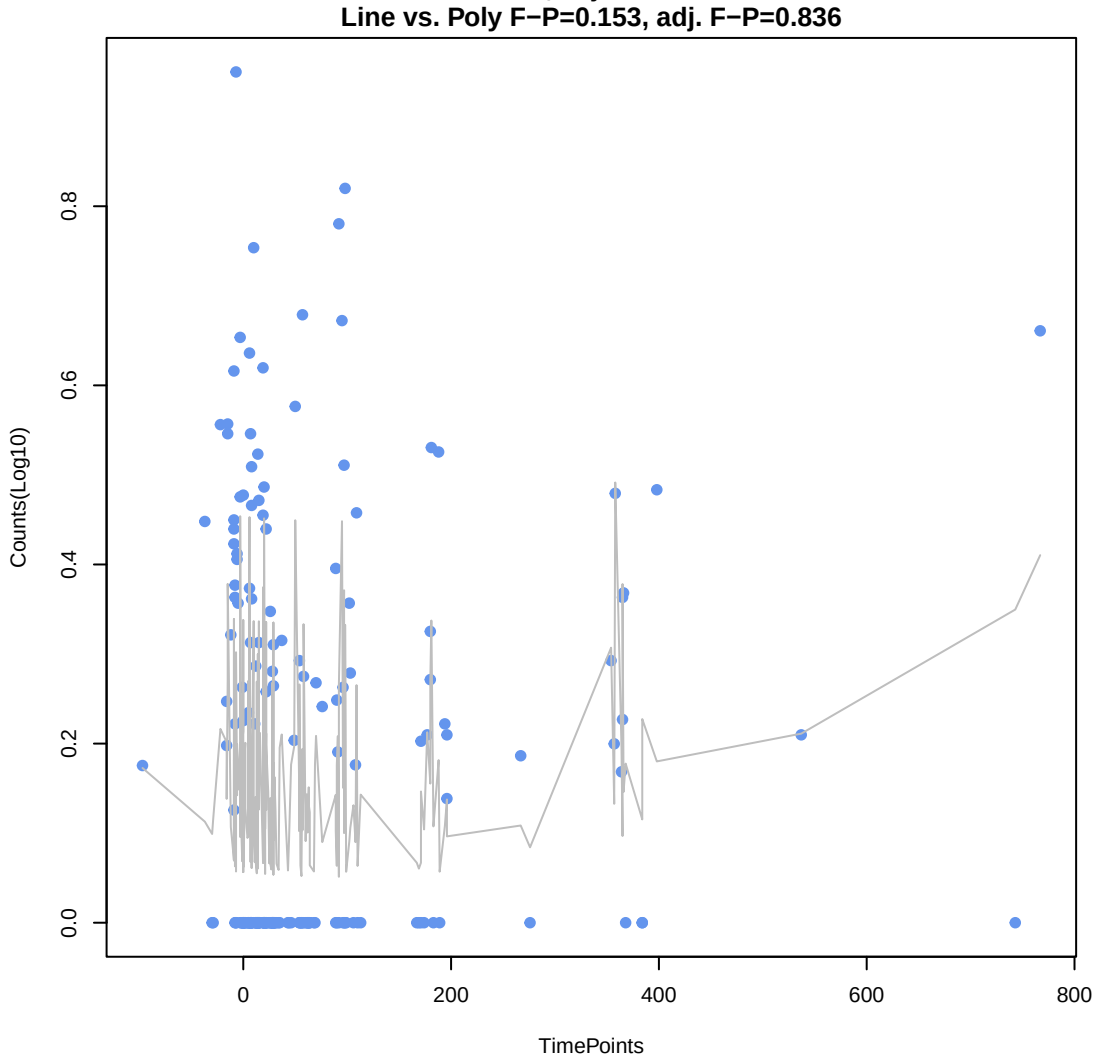
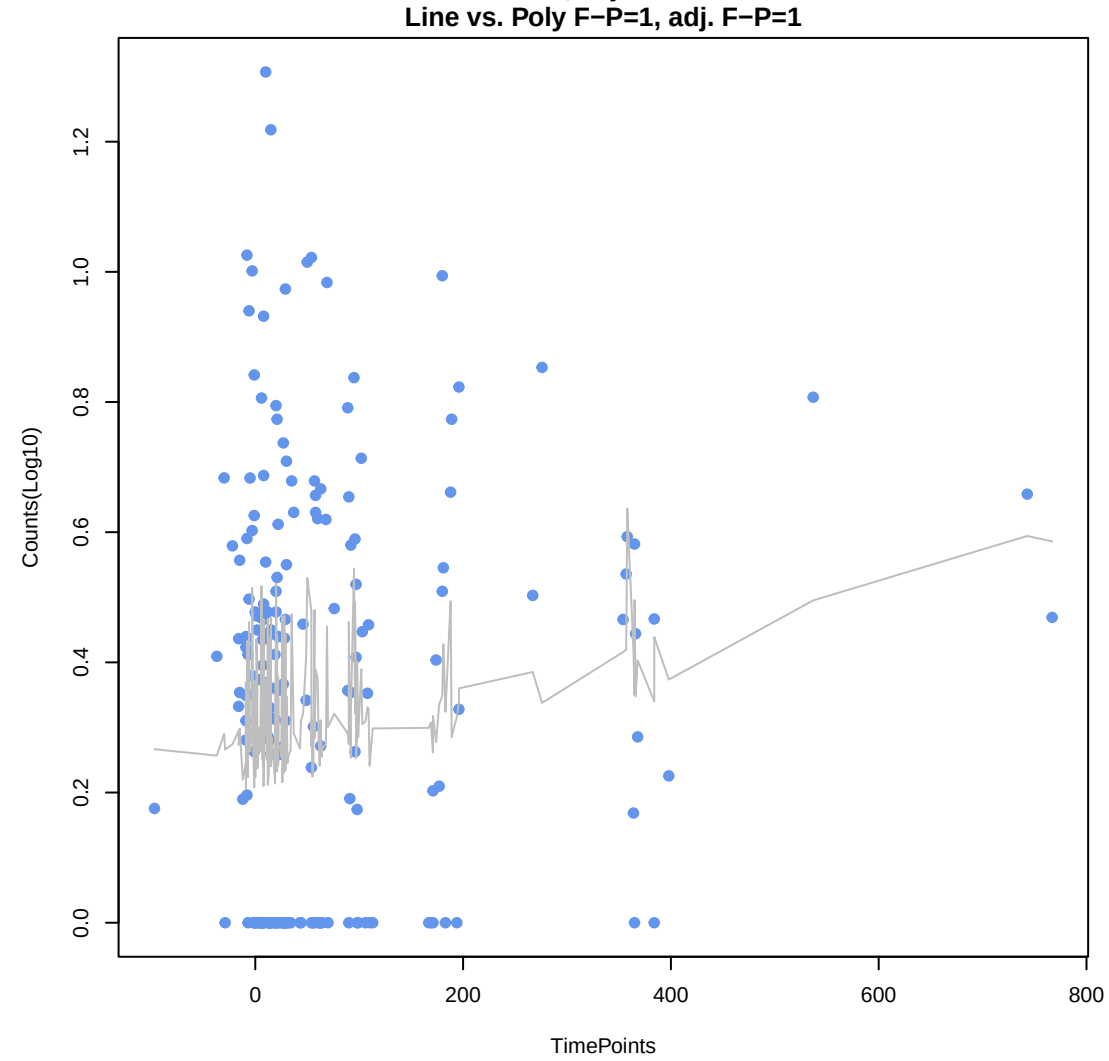

Erm 23S ribosomal RNA methyltransferase
ANOVA P=0.00493, adj. ANOVA−P=0.123
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints

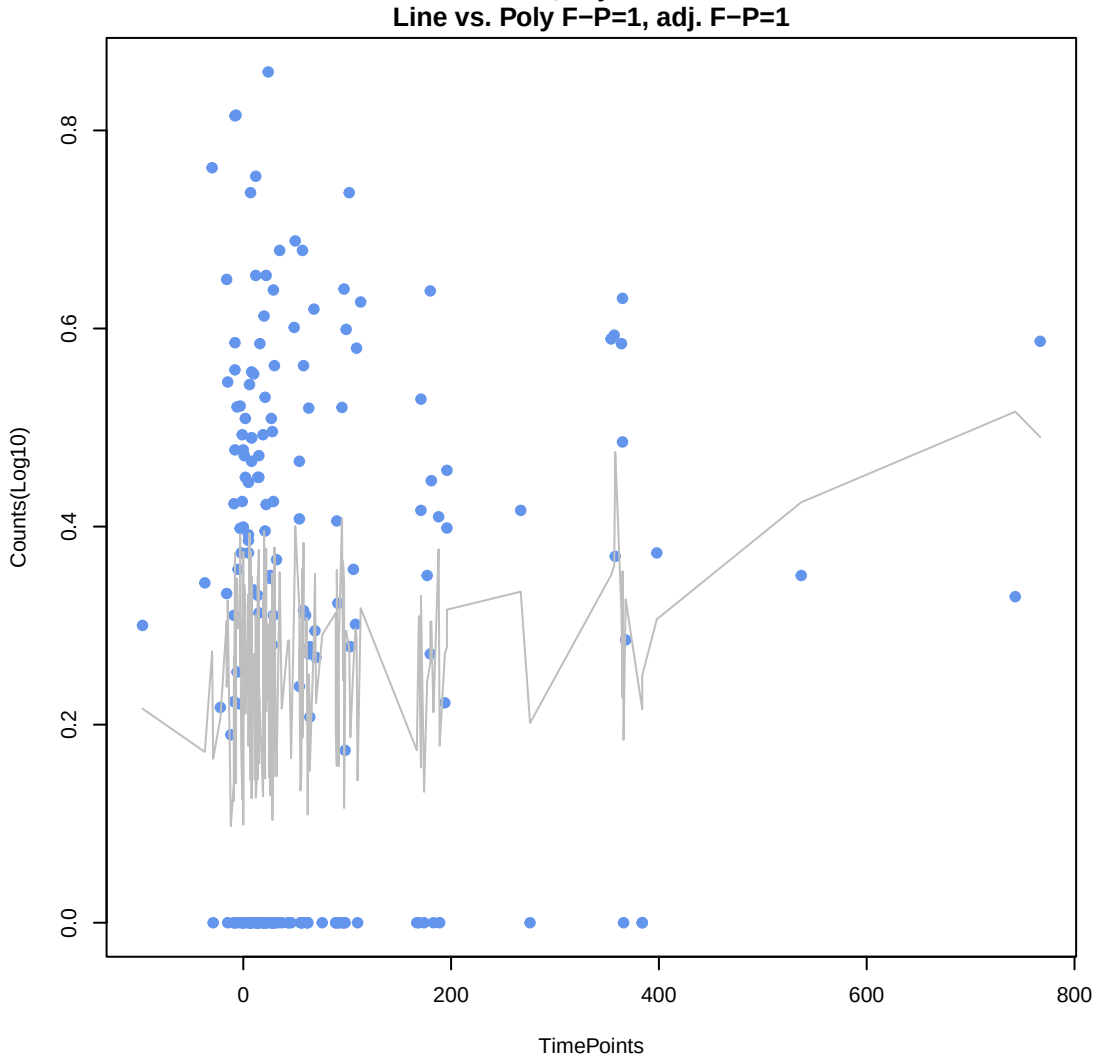

tetracycline−resistant ribosomal protection protein
ANOVA P=0.0212, adj. ANOVA−P=0.191
Line vs. Poly F−P=0.327, adj. F−P=0.908
Counts(Log10)
TimePoints

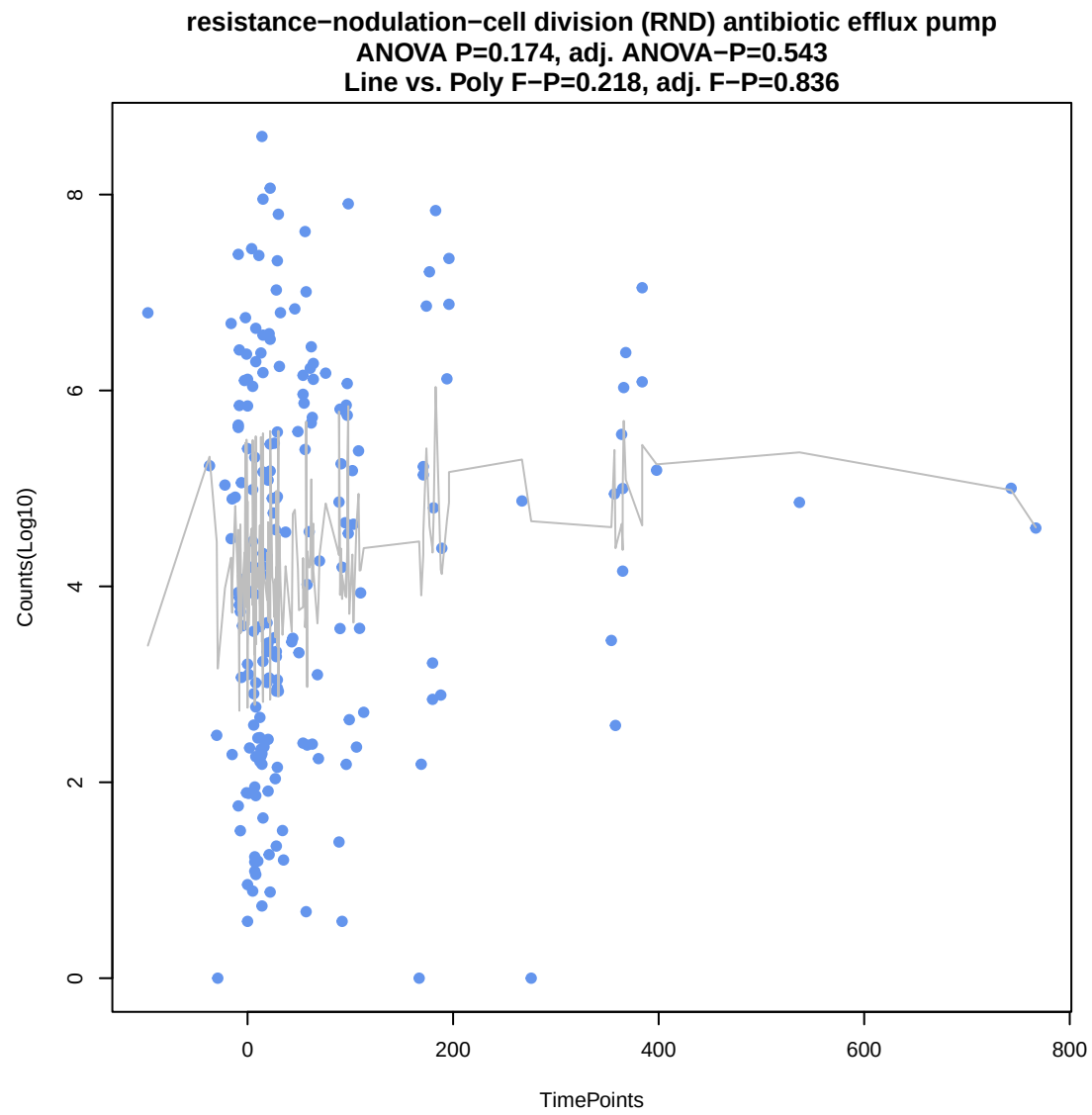

undecaprenyl pyrophosphate related proteins
ANOVA P=0.023, adj. ANOVA−P=0.191
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints

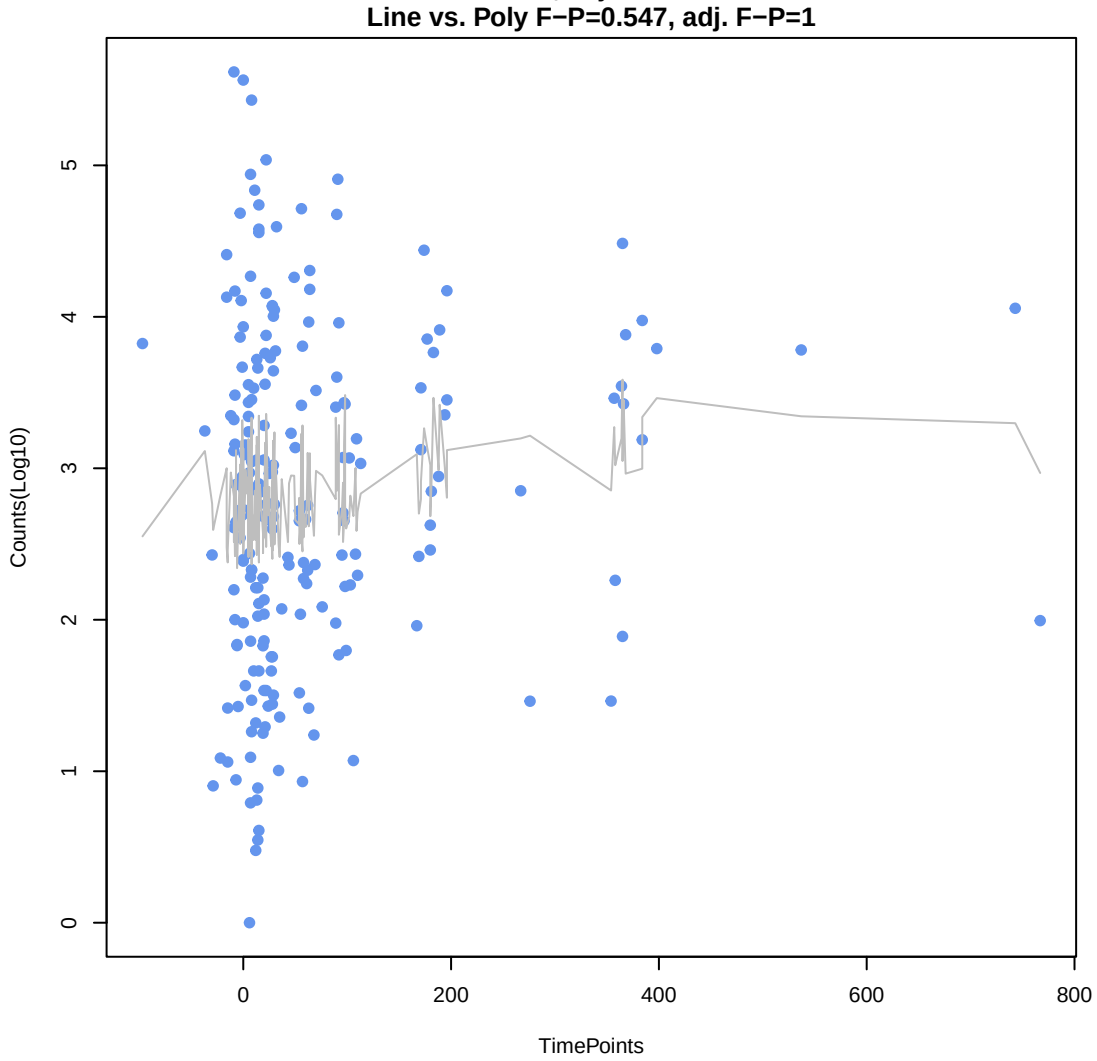

BlaB beta−lactamase
ANOVA P=0.0989, adj. ANOVA−P=0.516
Line vs. Poly F−P=0.216, adj. F−P=0.836
Counts(Log10)
TimePoints

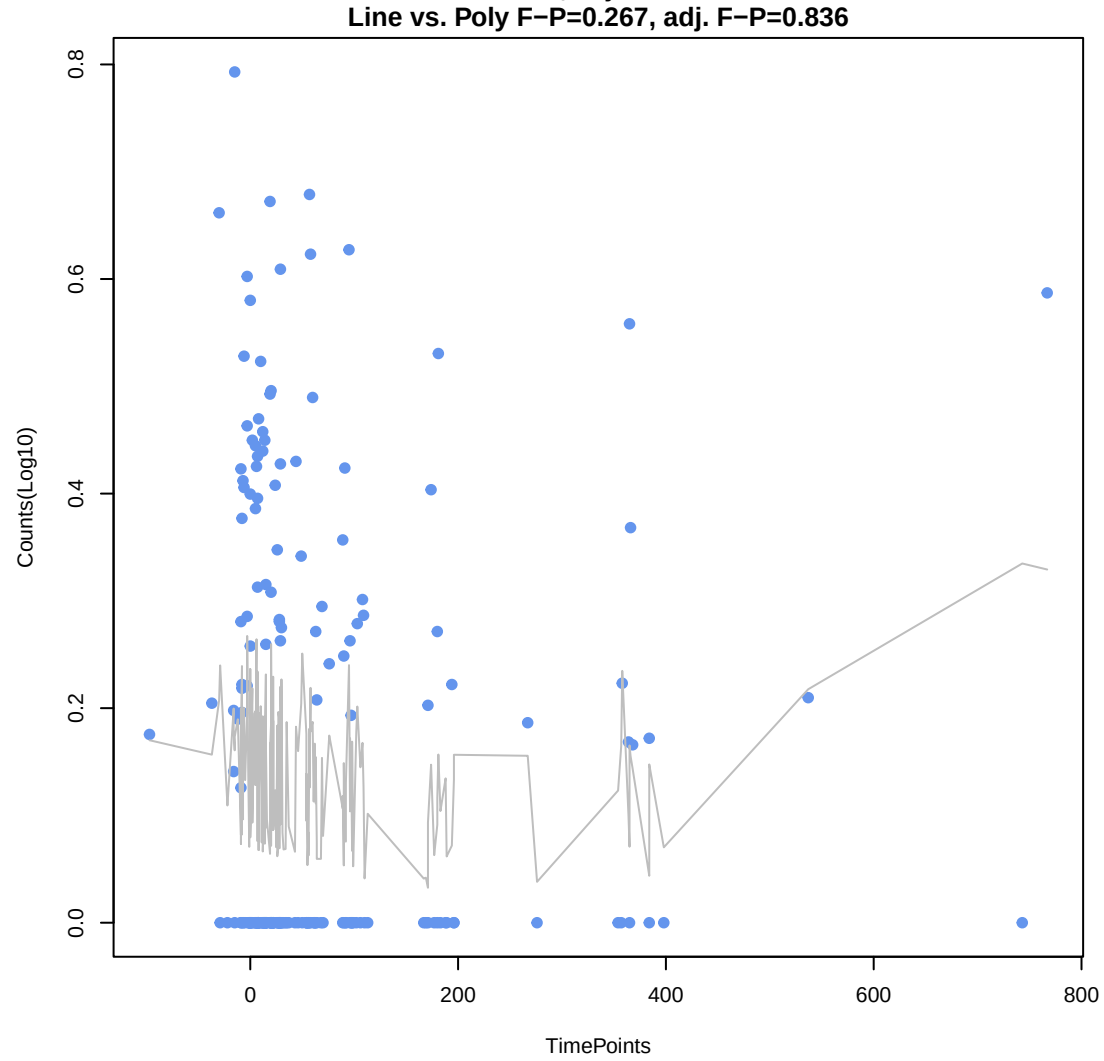

APH(6)
ANOVA P=0.12, adj. ANOVA−P=0.516
Line vs. Poly F−P=0.153, adj. F−P=0.836
Counts(Log10)
TimePoints

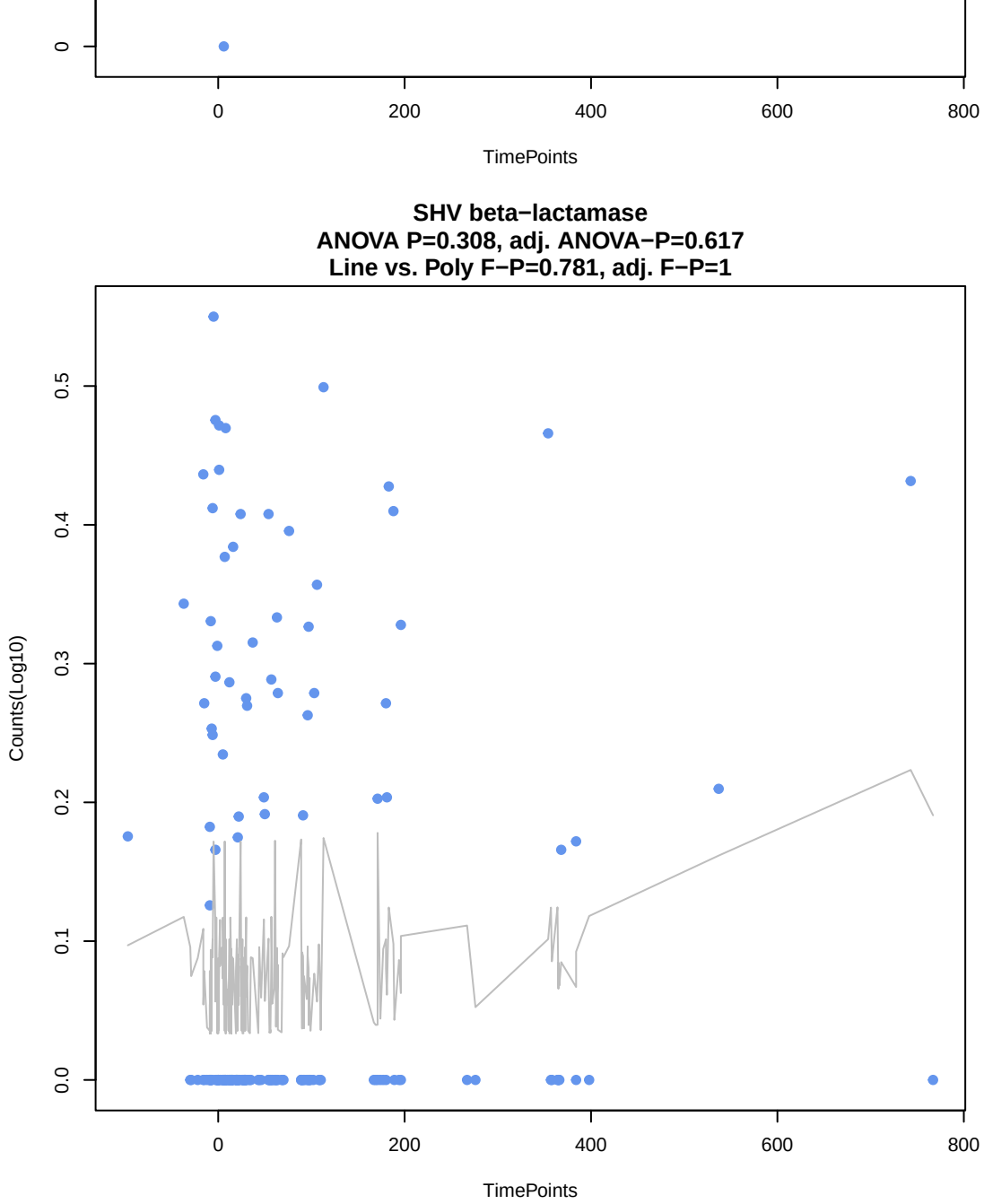

non−erm 23S ribosomal RNA methyltransferase (G748)
ANOVA P=0.124, adj. ANOVA−P=0.516
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints

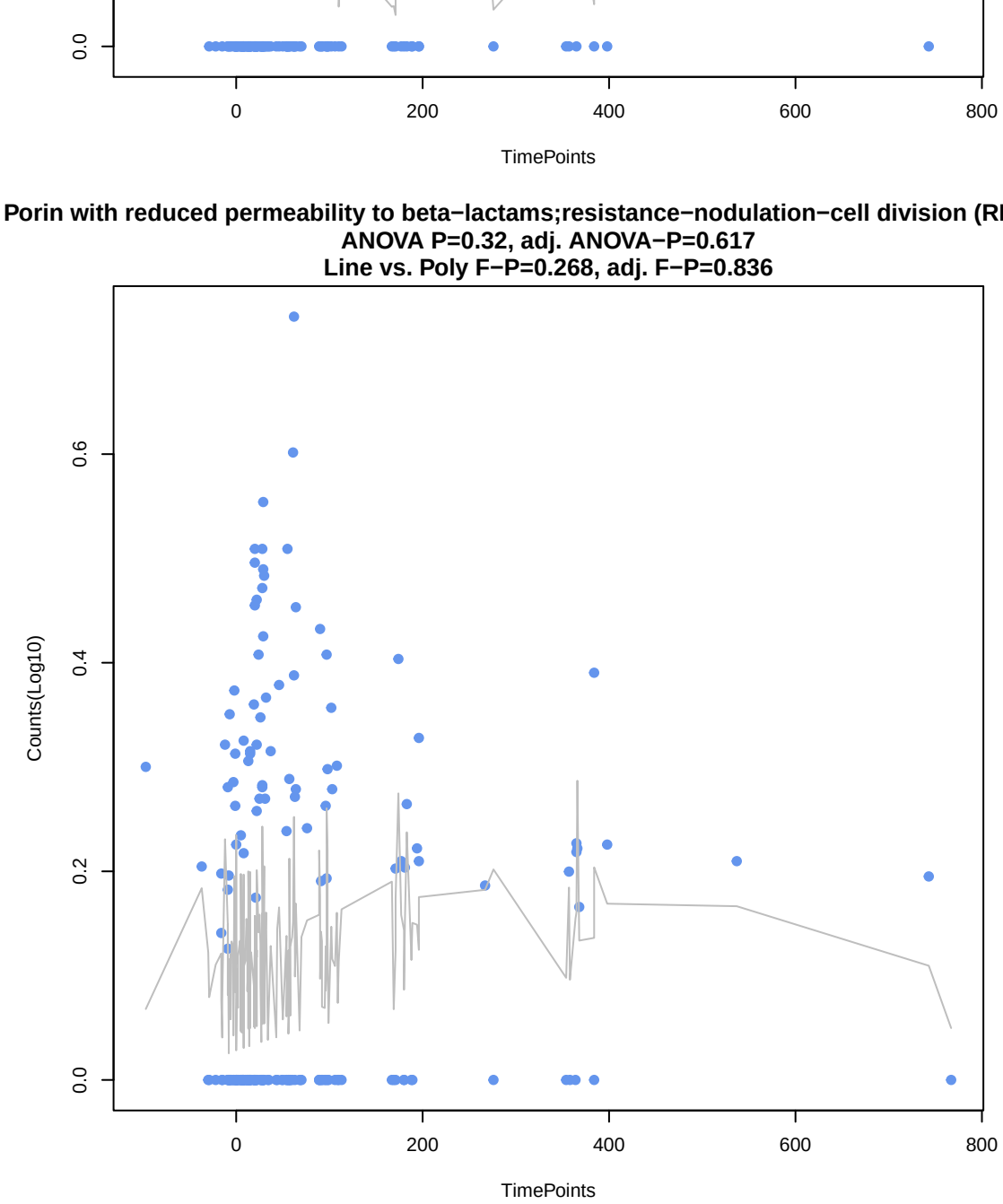

ANA beta−lactamase
ANOVA P=0.169, adj. ANOVA−P=0.543
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints


resistance−nodulation−cell division (RND) antibiotic efflux pump
ANOVA P=0.174, adj. ANOVA−P=0.543
Line vs. Poly F−P=0.218, adj. F−P=0.836
Counts(Log10)
TimePoints

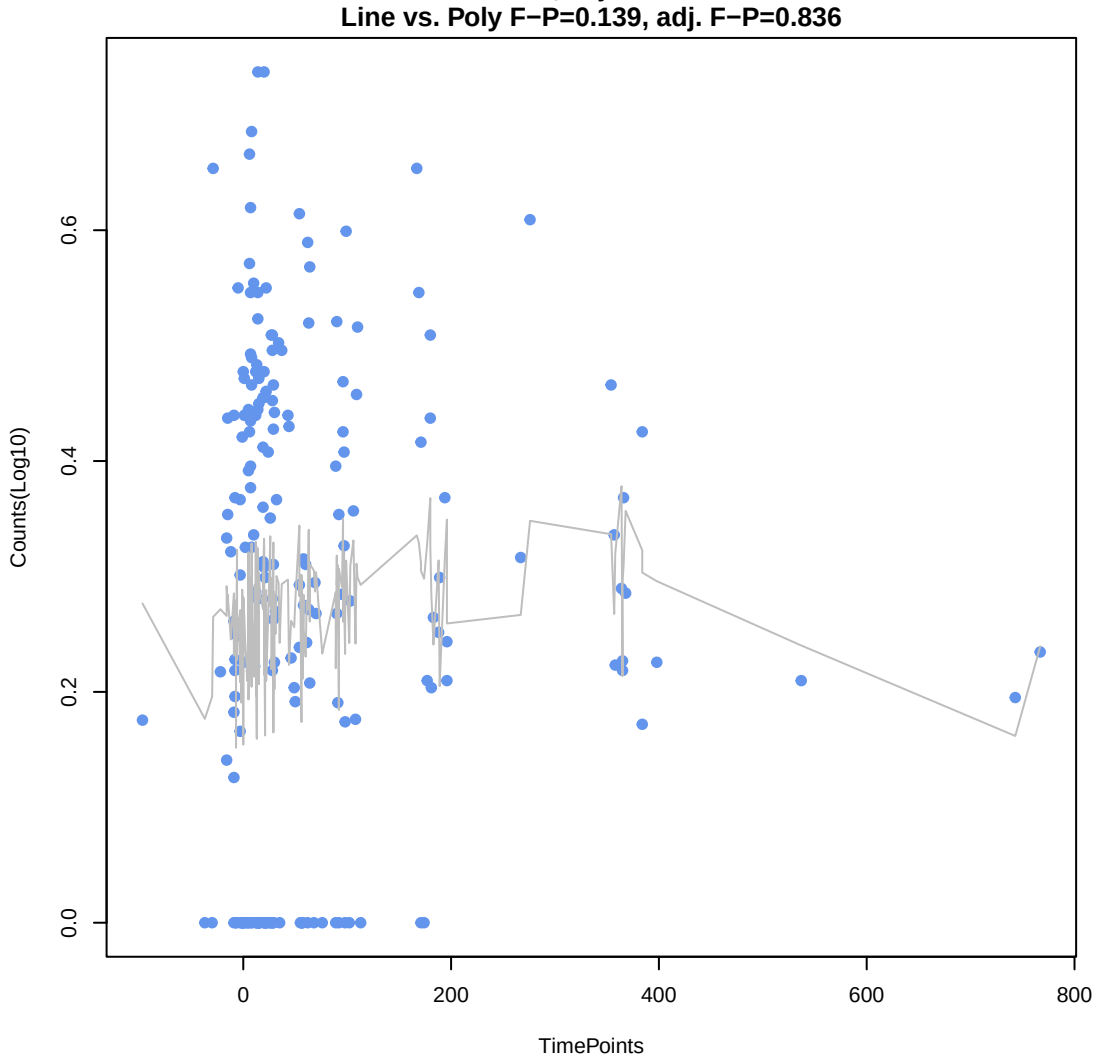

major facilitator superfamily (MFS) antibiotic efflux pump
ANOVA P=0.207, adj. ANOVA−P=0.574
Line vs. Poly F−P=0.547, adj. F−P=1
Counts(Log10)
TimePoints


PDC beta−lactamase
ANOVA P=0.295, adj. ANOVA−P=0.617
Line vs. Poly F−P=0.267, adj. F−P=0.836
Counts(Log10)
TimePoints

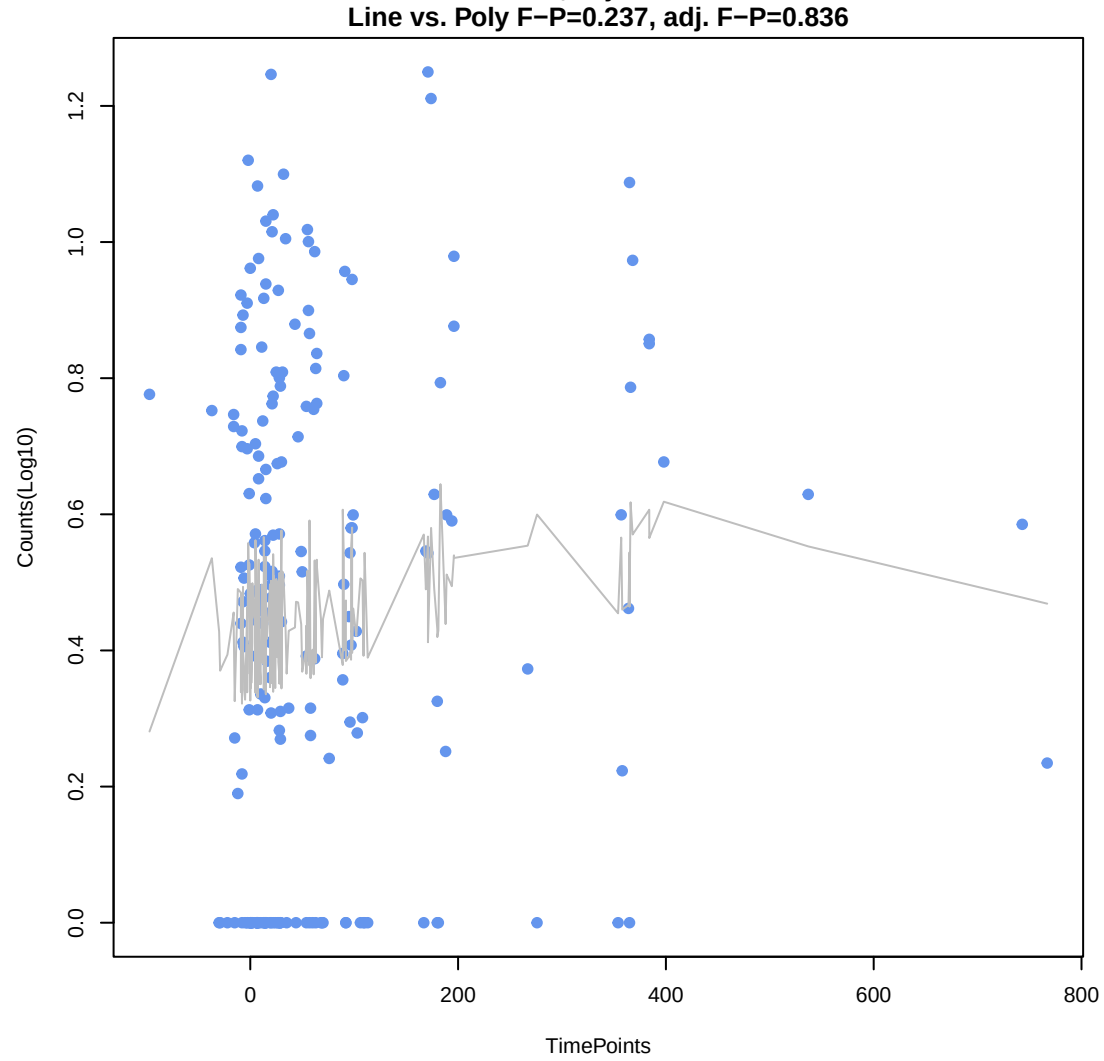

SHV beta−lactamase
ANOVA P=0.308, adj. ANOVA−P=0.617
Line vs. Poly F−P=0.781, adj. F−P=1
Counts(Log10)
TimePoints


al Porin with reduced permeability to beta−lactams;resistance−nodulation−cell division (RN
ANOVA P=0.32, adj. ANOVA−P=0.617
Line vs. Poly F−P=0.268, adj. F−P=0.836
Counts(Log10)
TimePoints

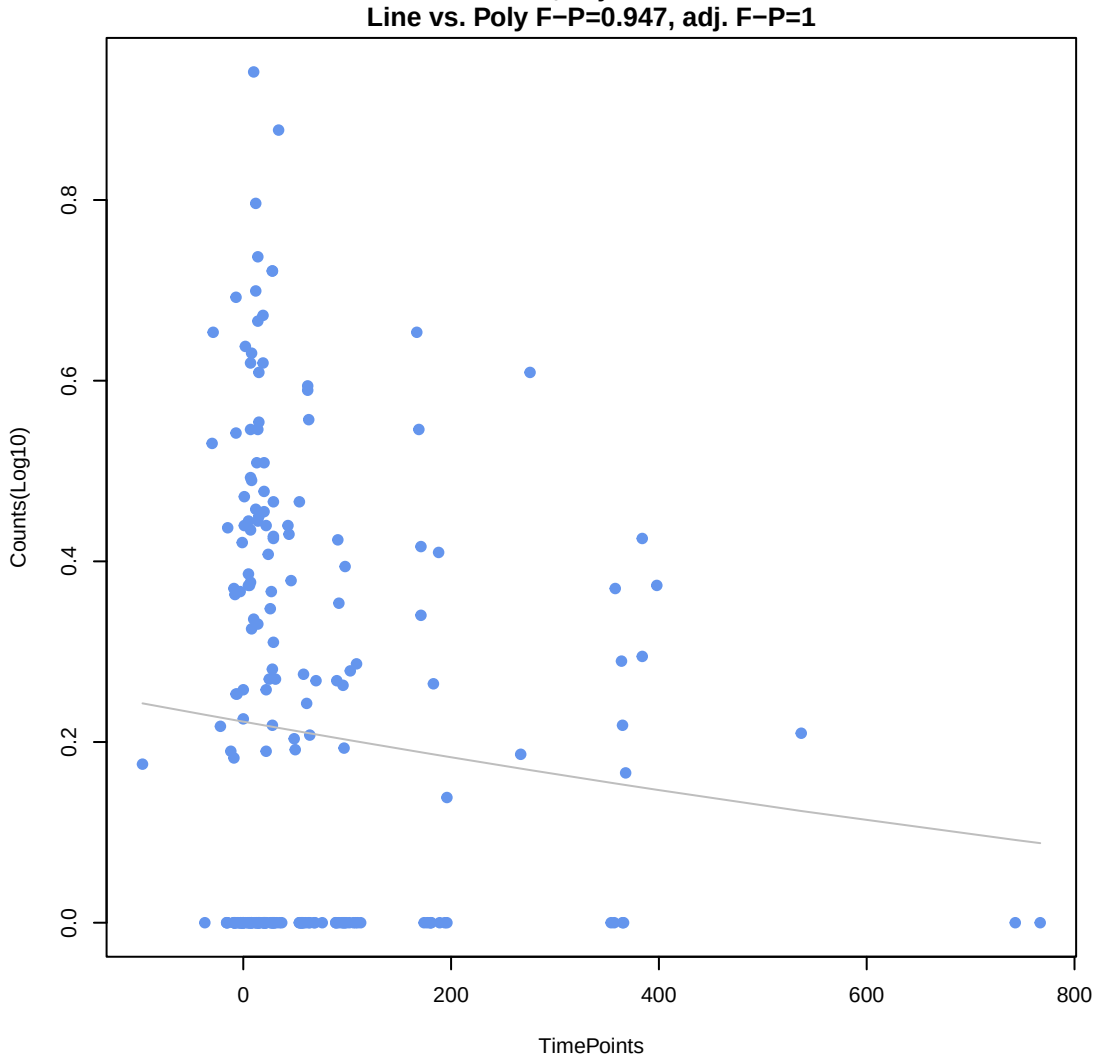
streptothricin acetyltransferase (SAT)
ANOVA P=0.343, adj. ANOVA−P=0.617
Line vs. Poly F−P=0.139, adj. F−P=0.836



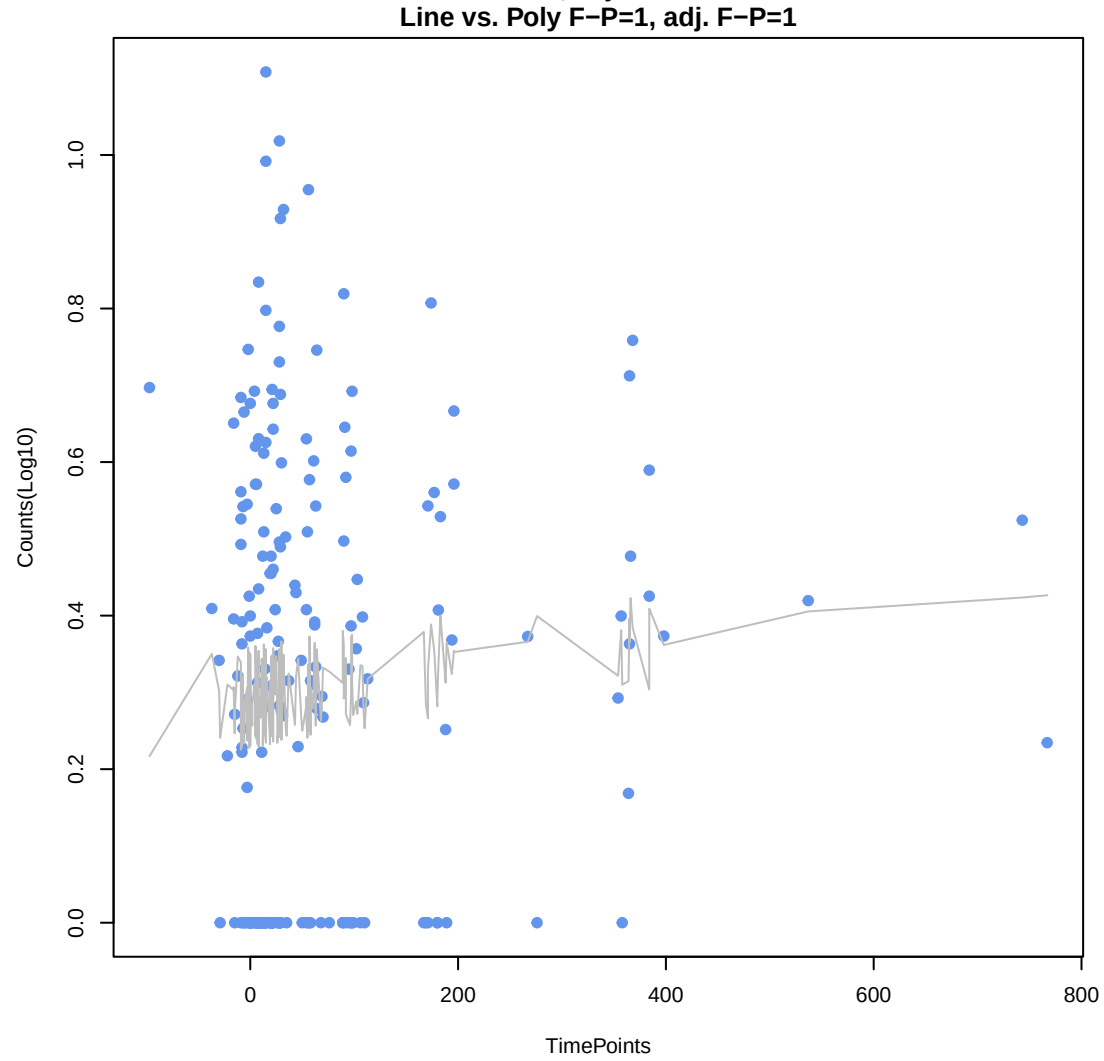
itator superfamily (MFS) antibiotic efflux pump;resistance−nodulation−cell division (RND) a
ANOVA P=0.346, adj. ANOVA−P=0.617
Line vs. Poly F−P=0.237, adj. F−P=0.836



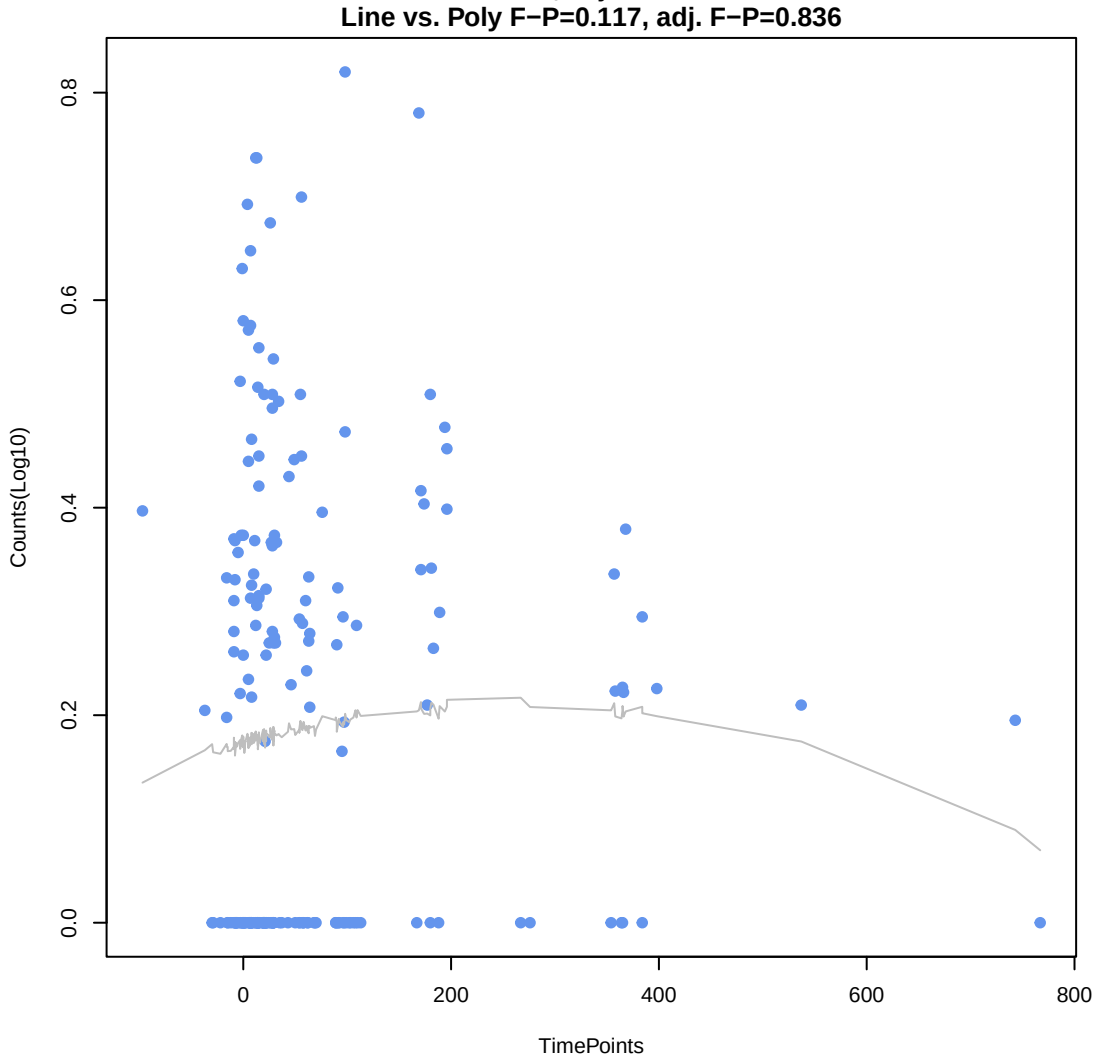
msr−type ABC−F protein
ANOVA P=0.415, adj. ANOVA−P=0.692
Line vs. Poly F−P=0.947, adj. F−P=1



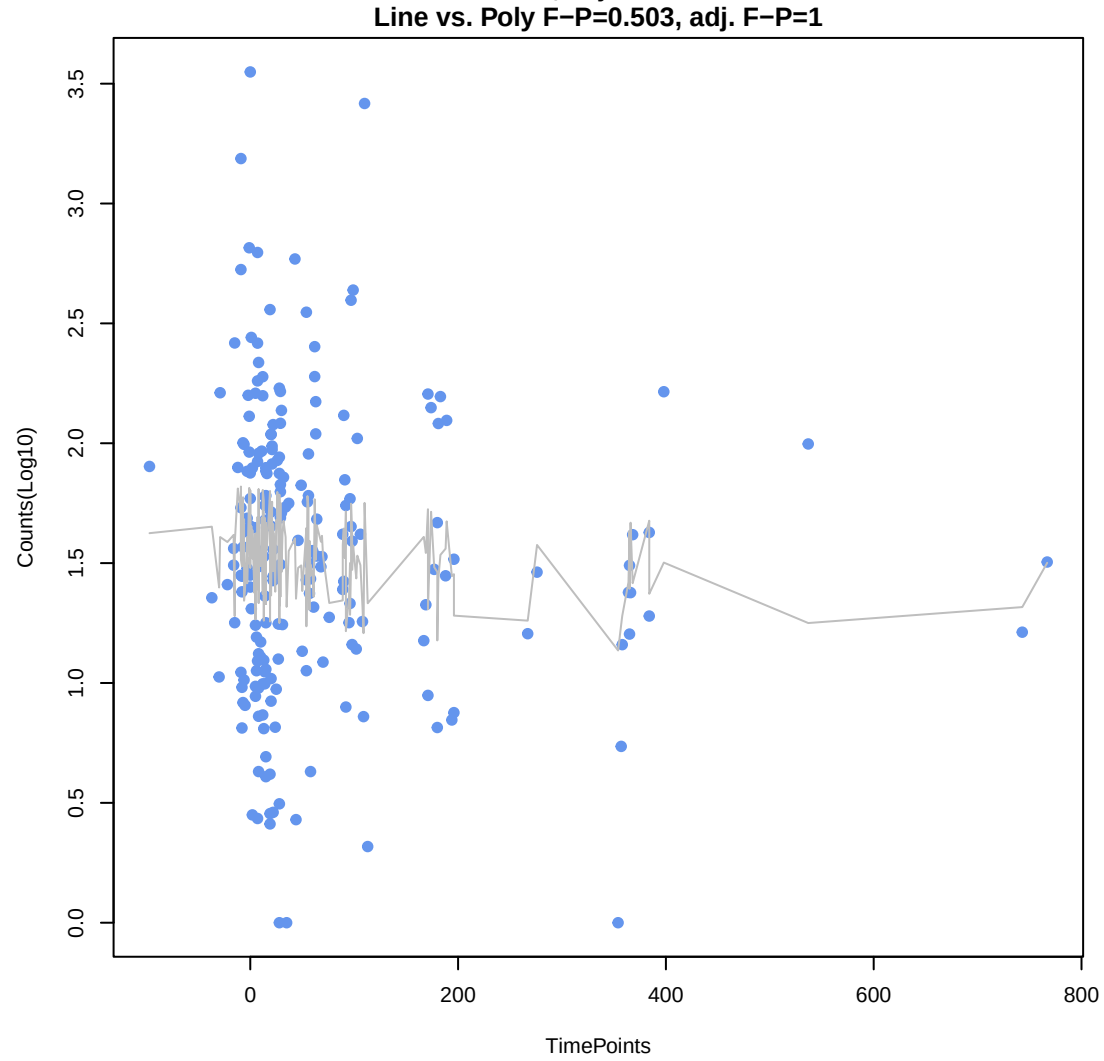
pmr phosphoethanolamine transferase
ANOVA P=0.525, adj. ANOVA−P=0.817
Line vs. Poly F−P=1, adj. F−P=1

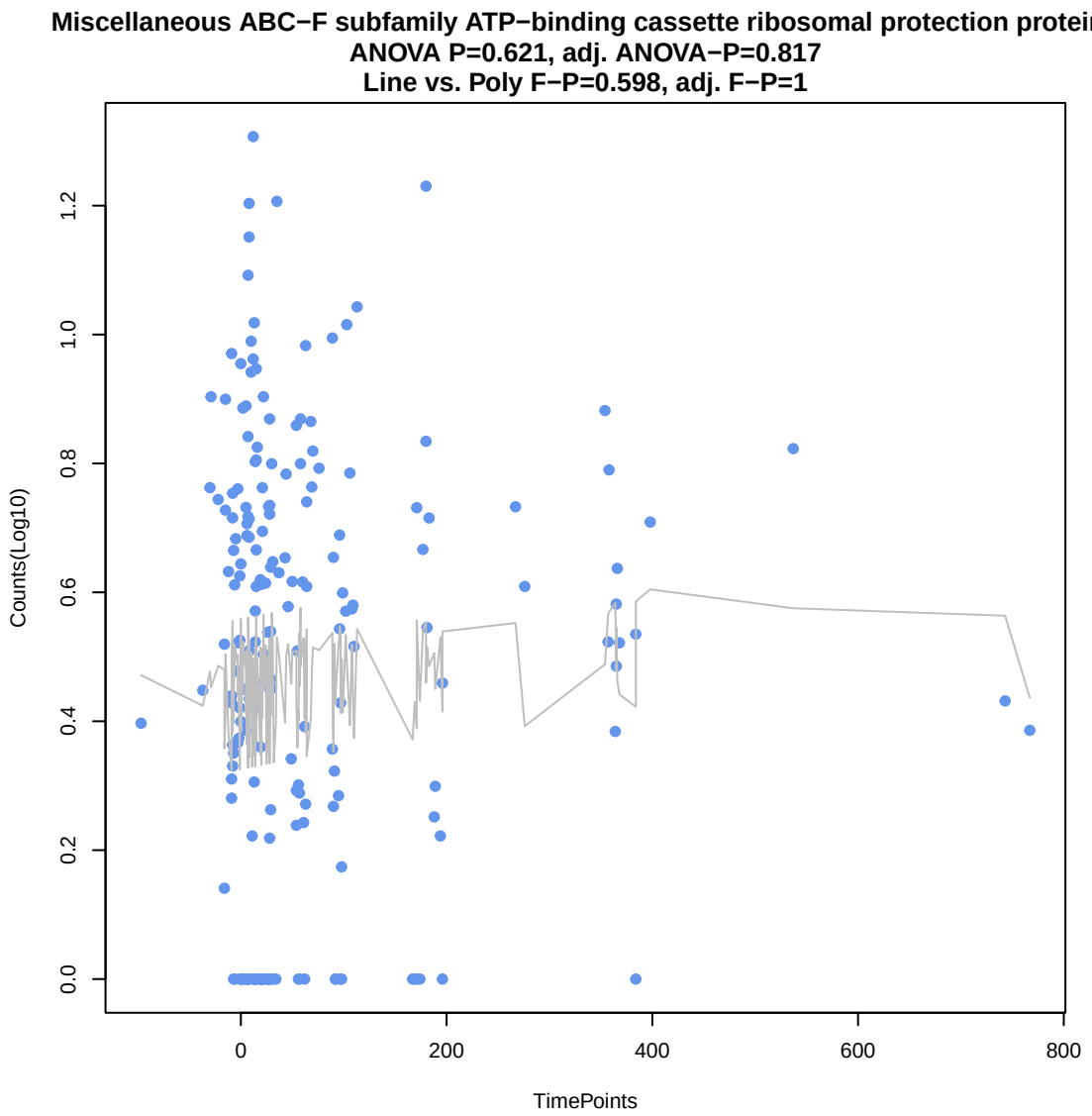


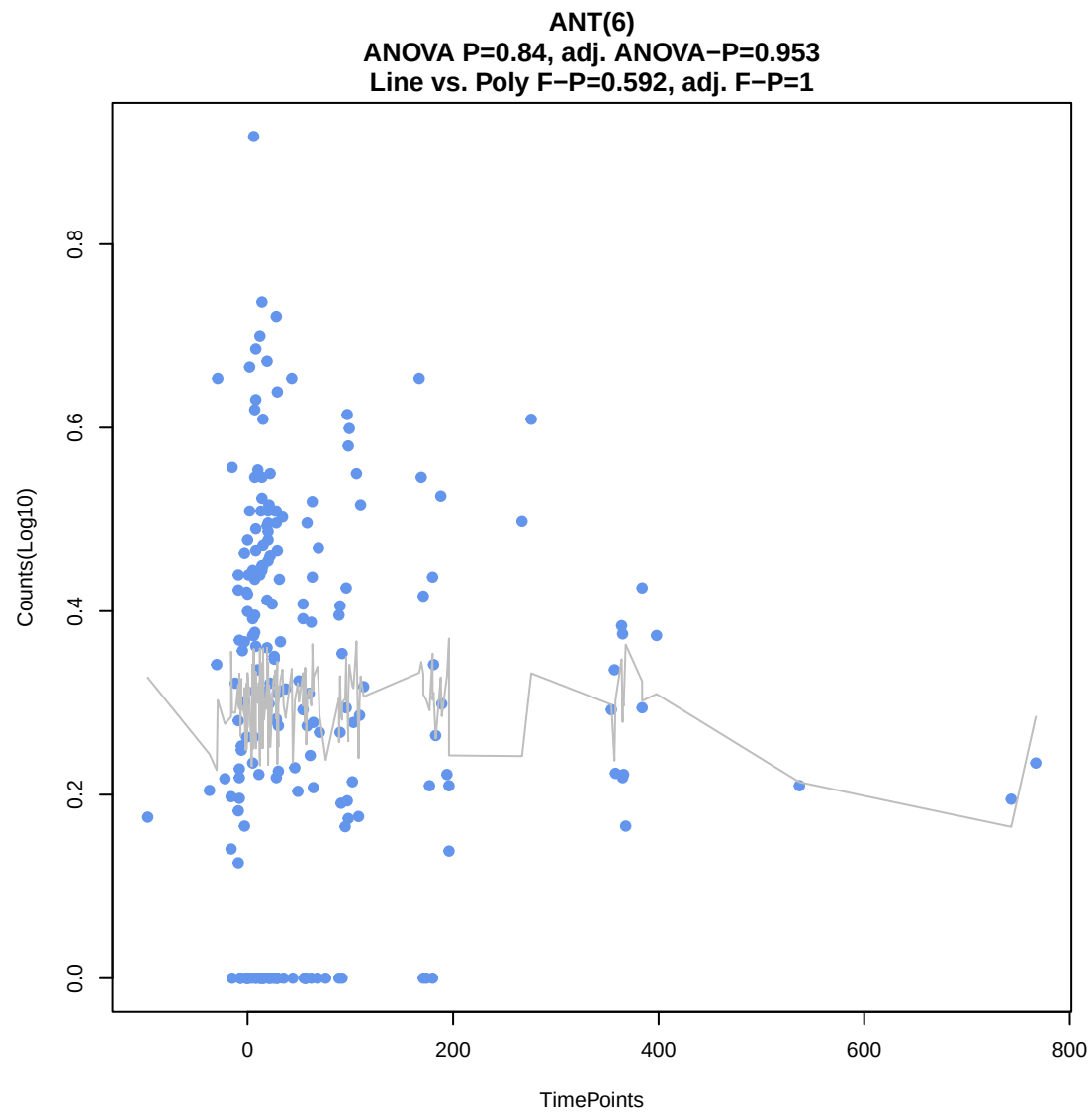
ic efflux pump;major facilitator superfamily (MFS) antibiotic efflux pump;resistance−nodula
ANOVA P=0.572, adj. ANOVA−P=0.817
Line vs. Poly F−P=0.117, adj. F−P=0.836

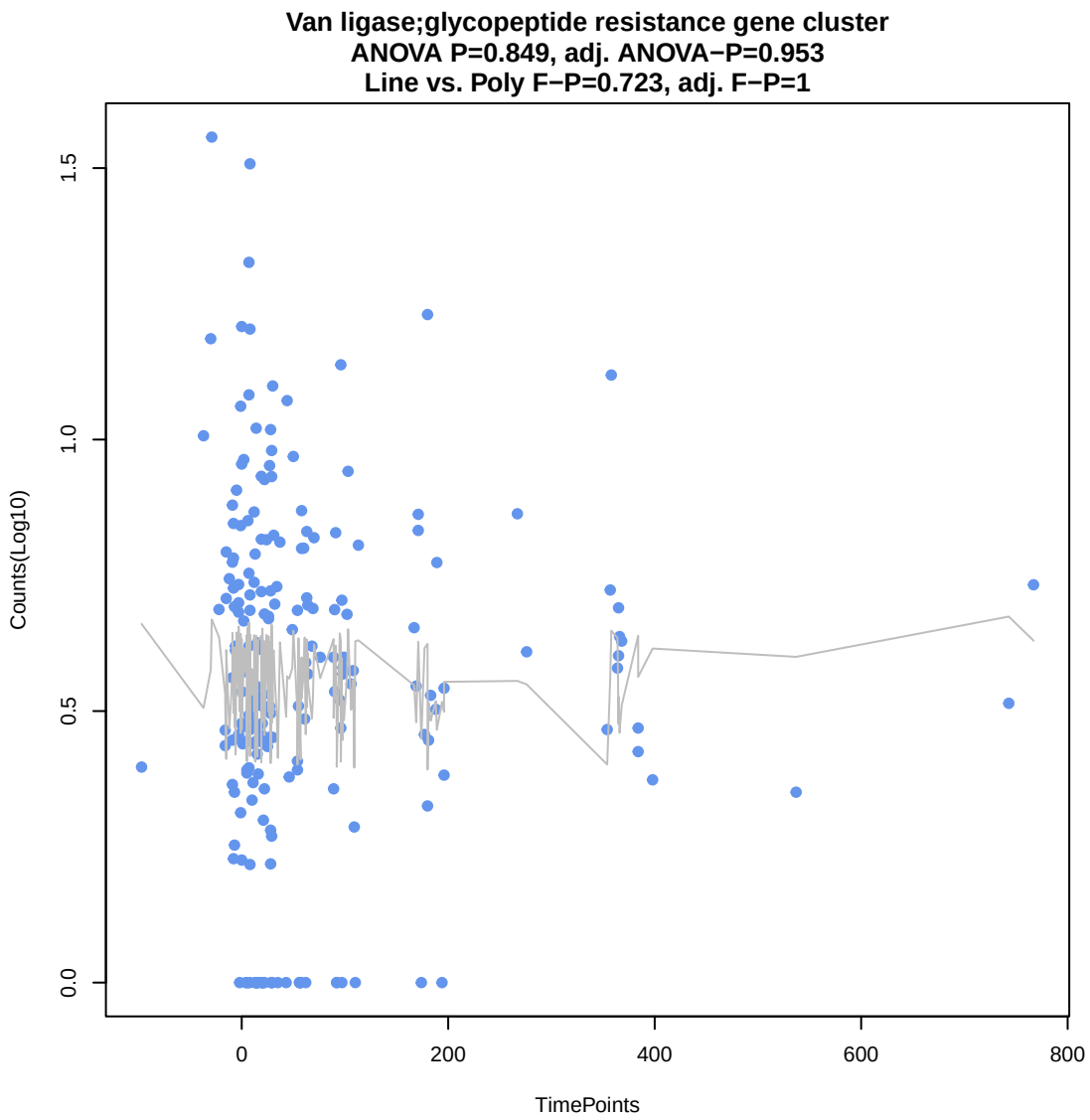


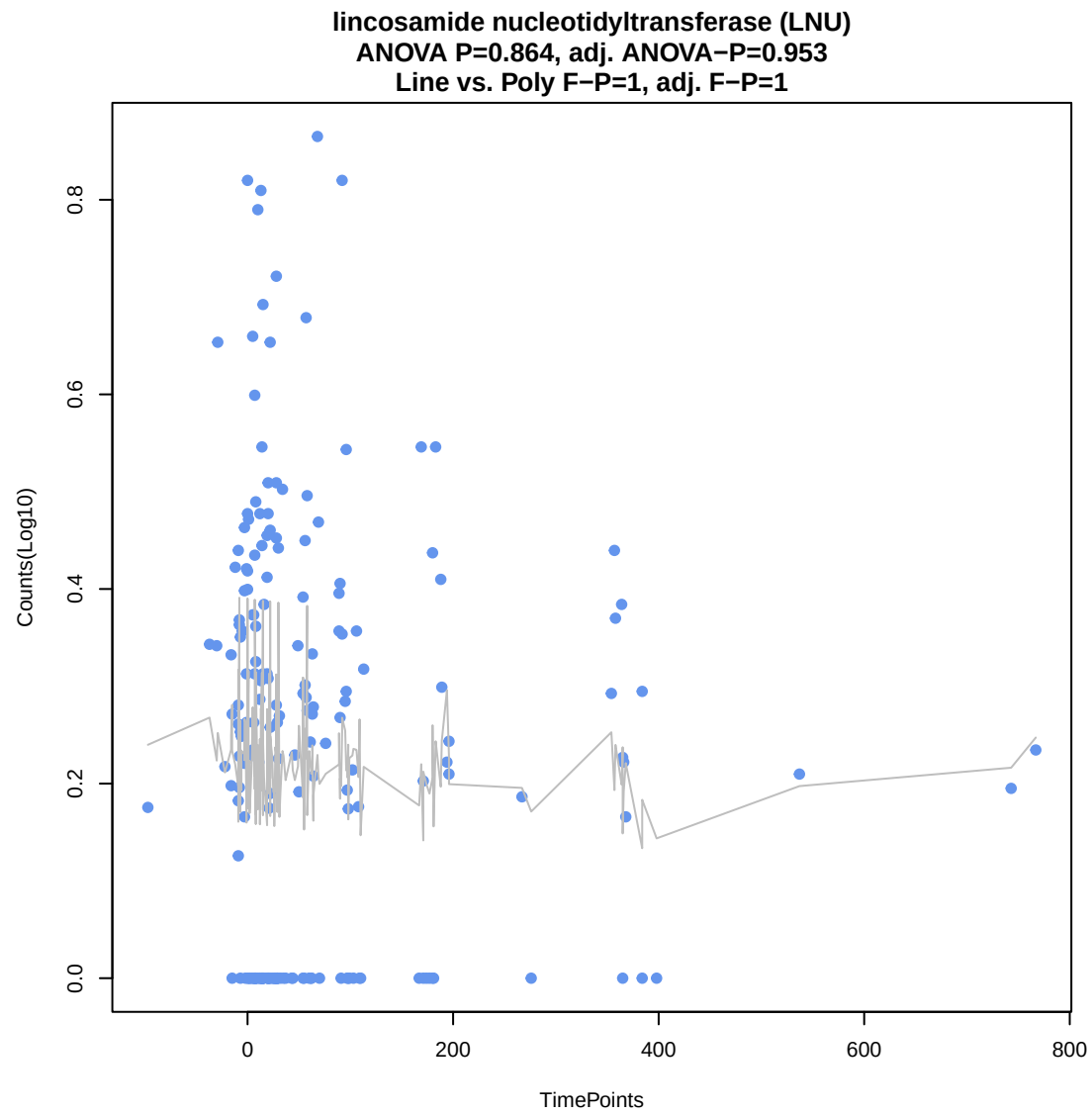
ATP−binding cassette (ABC) antibiotic efflux pump
ANOVA P=0.62, adj. ANOVA−P=0.817
Line vs. Poly F−P=0.503, adj. F−P=1

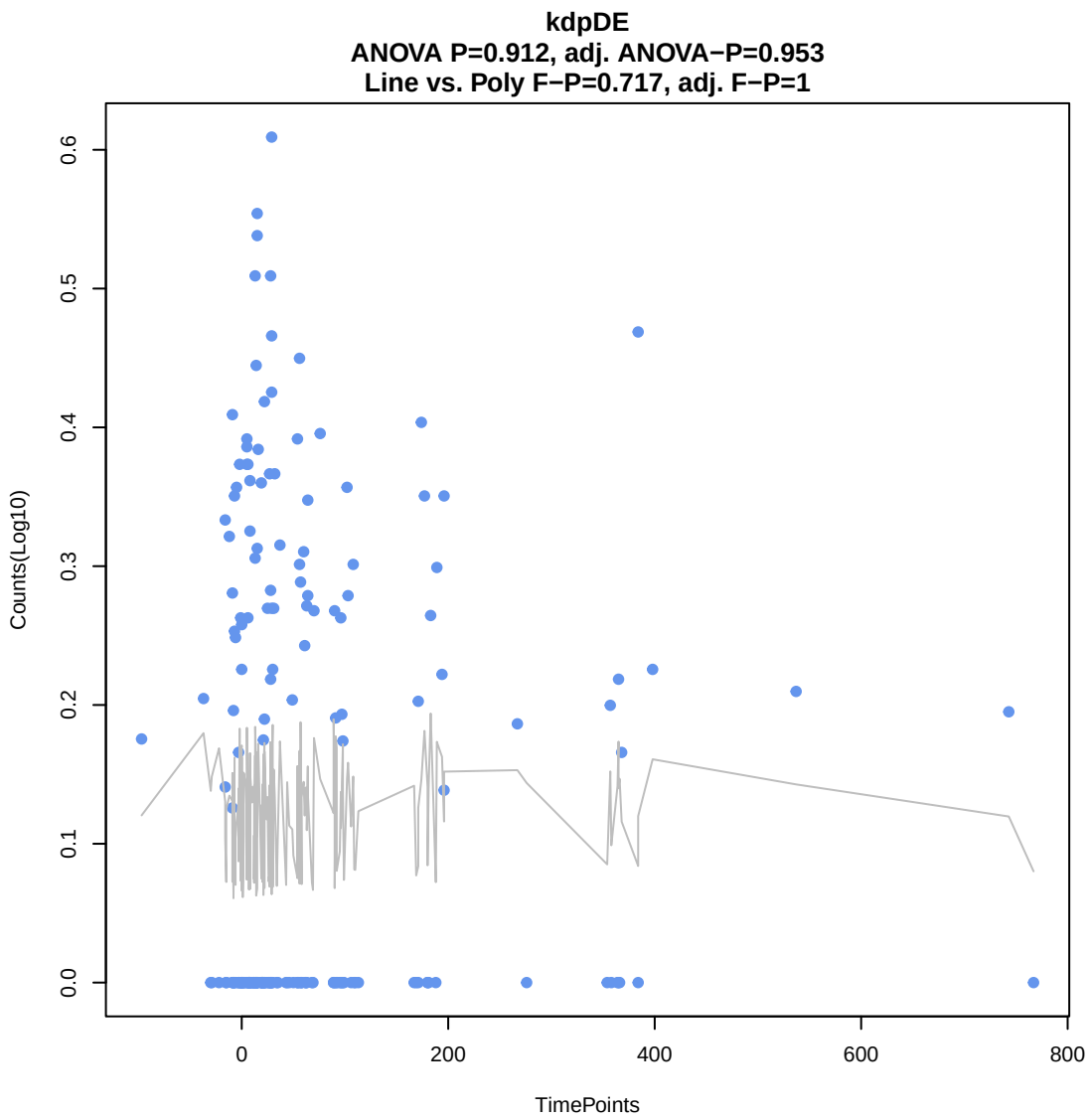


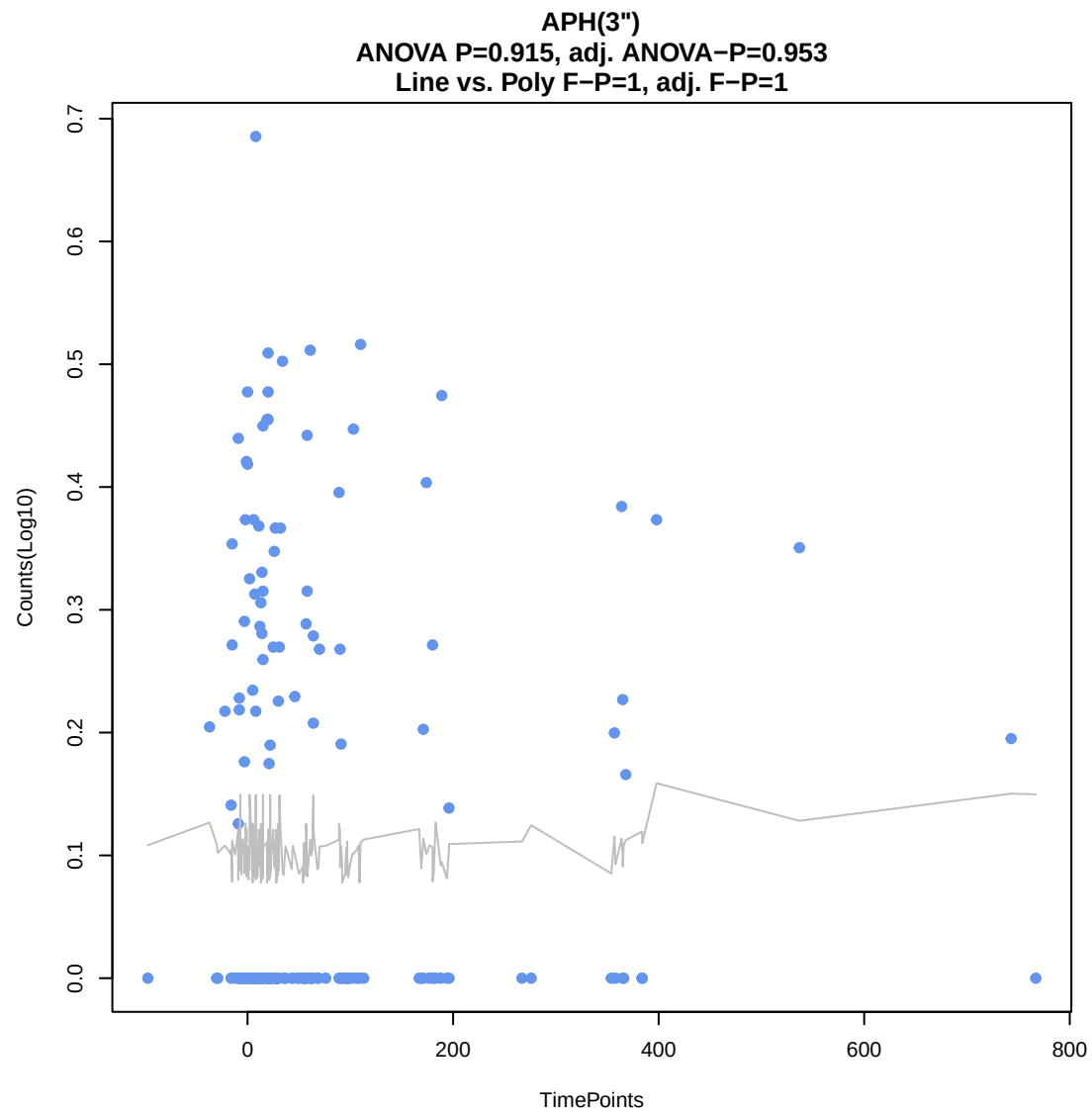

Miscellaneous ABC−F subfamily ATP−binding cassette ribosomal protection protei
ANOVA P=0.621, adj. ANOVA−P=0.817
Line vs. Poly F−P=0.598, adj. F−P=1
Counts(Log10)
TimePoints

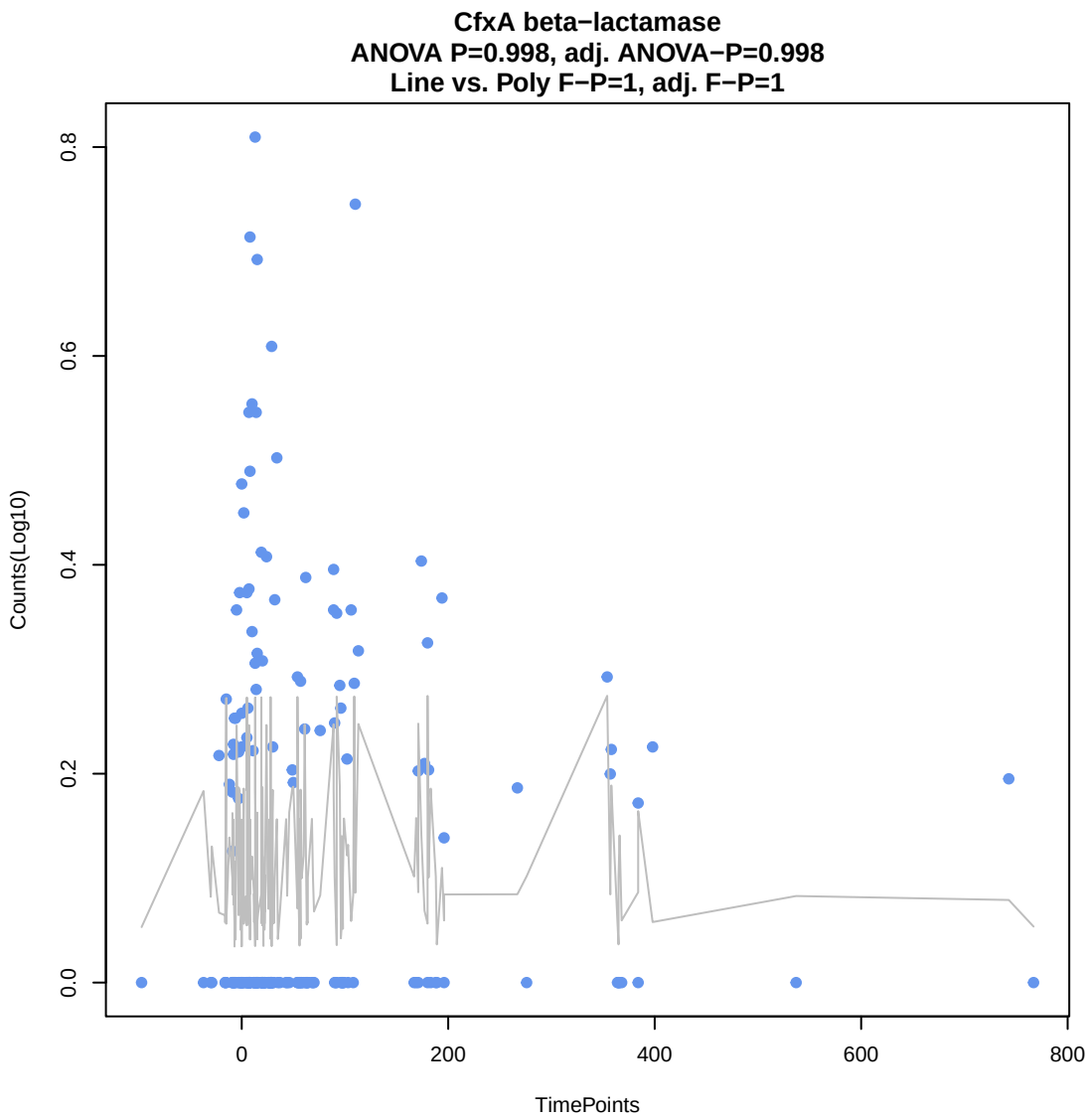

ANT(6)
ANOVA P=0.84, adj. ANOVA−P=0.953
Line vs. Poly F−P=0.592, adj. F−P=1
Counts(Log10)
TimePoints


Van ligase;glycopeptide resistance gene cluster
ANOVA P=0.849, adj. ANOVA−P=0.953
Line vs. Poly F−P=0.723, adj. F−P=1
Counts(Log10)
TimePoints


lincosamide nucleotidyltransferase (LNU)
ANOVA P=0.864, adj. ANOVA−P=0.953
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints


kdpDE
ANOVA P=0.912, adj. ANOVA−P=0.953
Line vs. Poly F−P=0.717, adj. F−P=1
Counts(Log10)
TimePoints


APH(3")
ANOVA P=0.915, adj. ANOVA−P=0.953
Line vs. Poly F−P=1, adj. F−P=1
Counts(Log10)
TimePoints


CfxA beta−lactamase
ANOVA P=0.998, adj. ANOVA−P=0.998
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