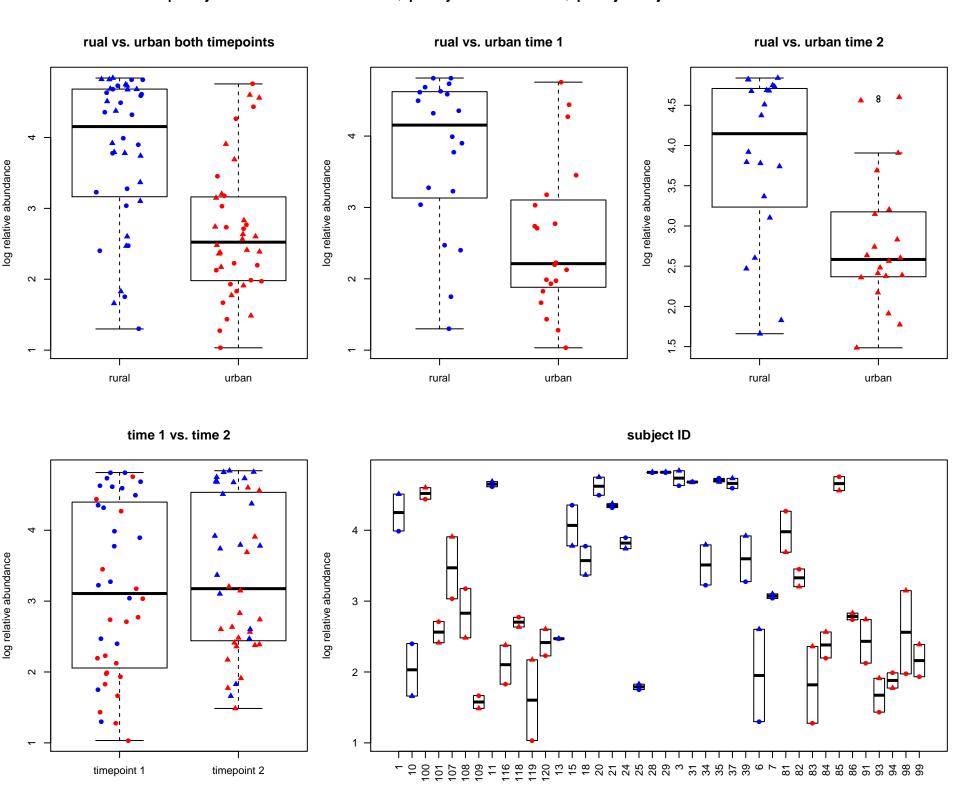
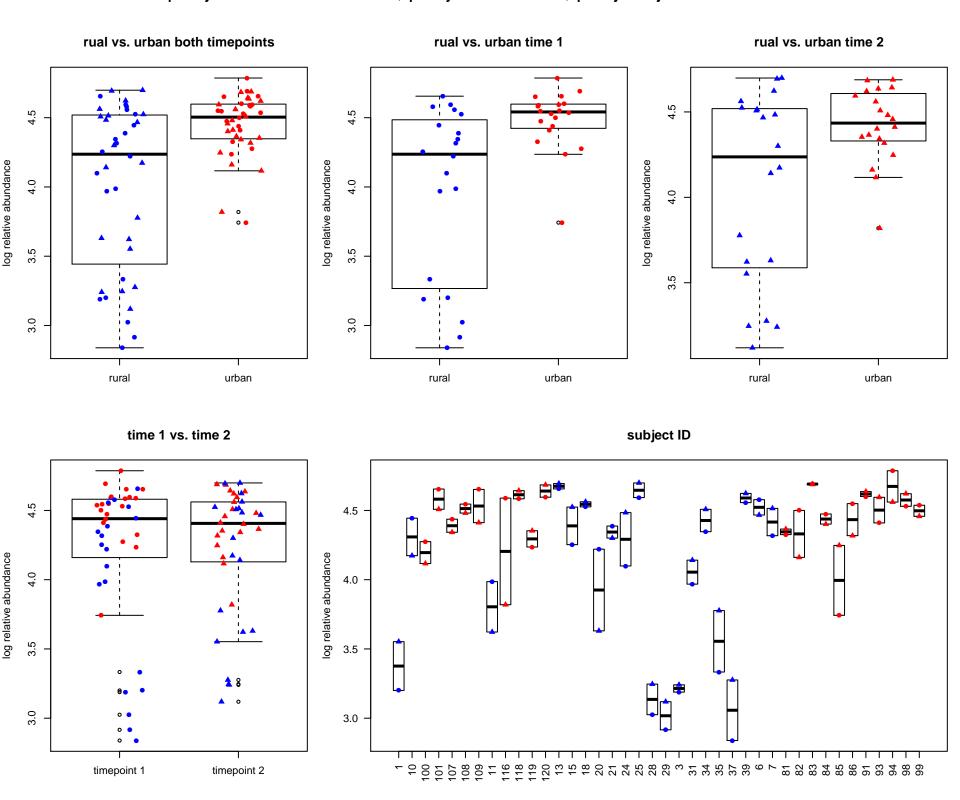
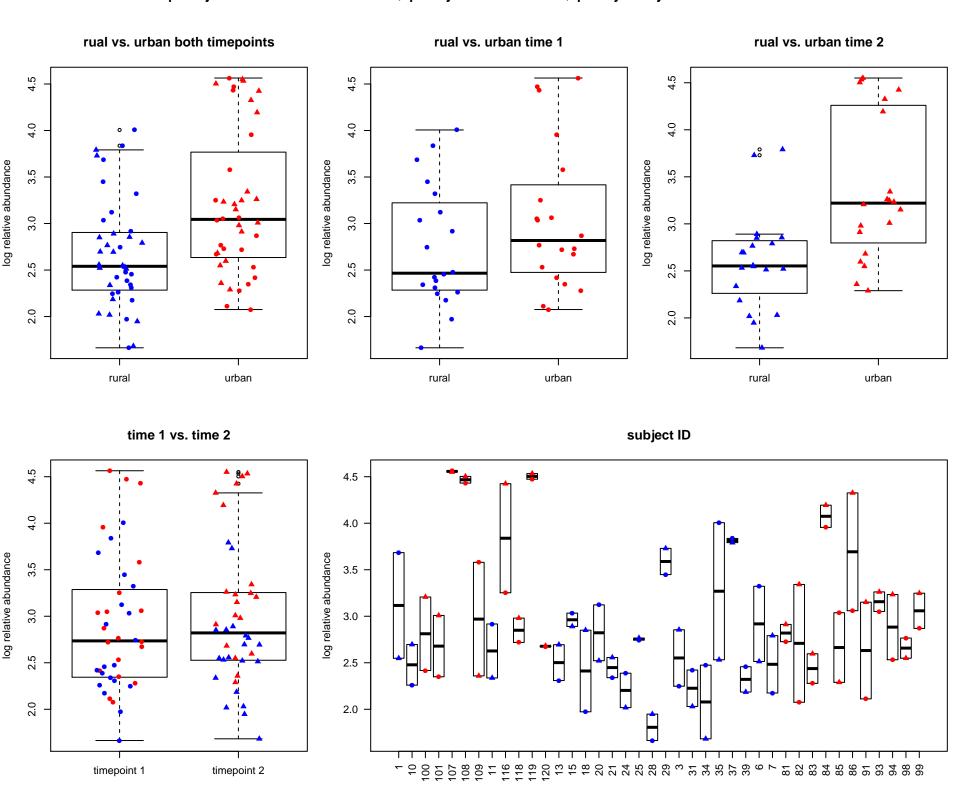
16S family: Prevotellaceae pAdjRuralUrban=0.0279; pAdjTime=0.241; pAdjSubject=9.21e-13



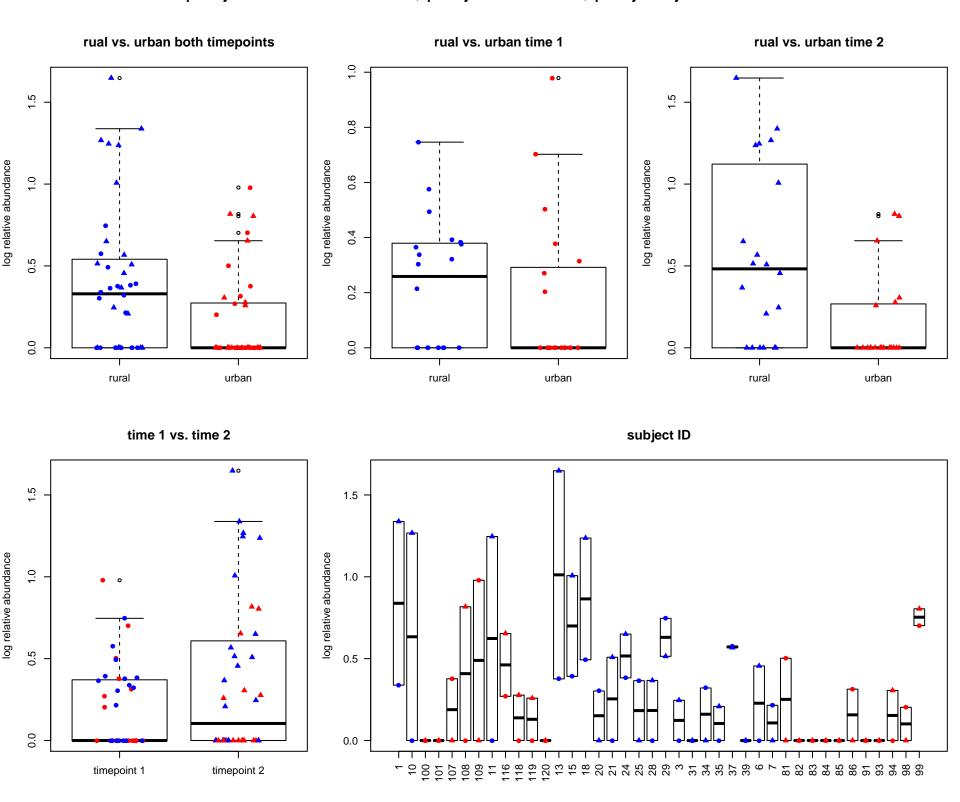
16S family: Bacteroidaceae pAdjRuralUrban=0.0534; pAdjTime=0.916; pAdjSubject=2.69e-10



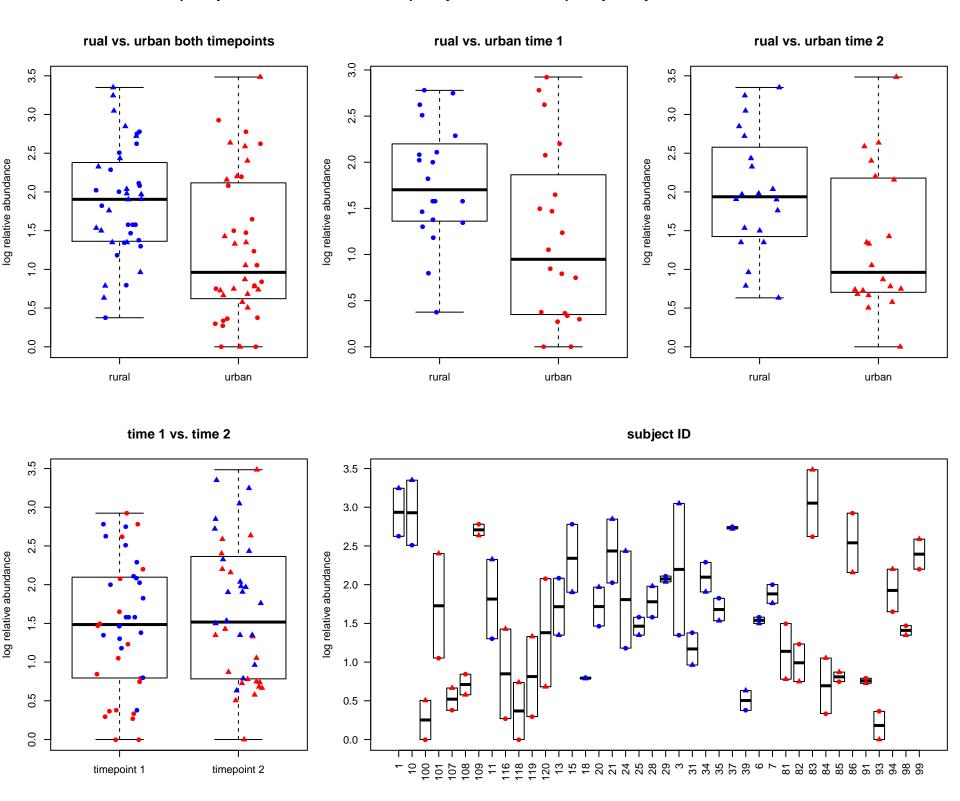
16S family: Enterobacteriaceae pAdjRuralUrban=0.0961; pAdjTime=0.773; pAdjSubject=0.000507



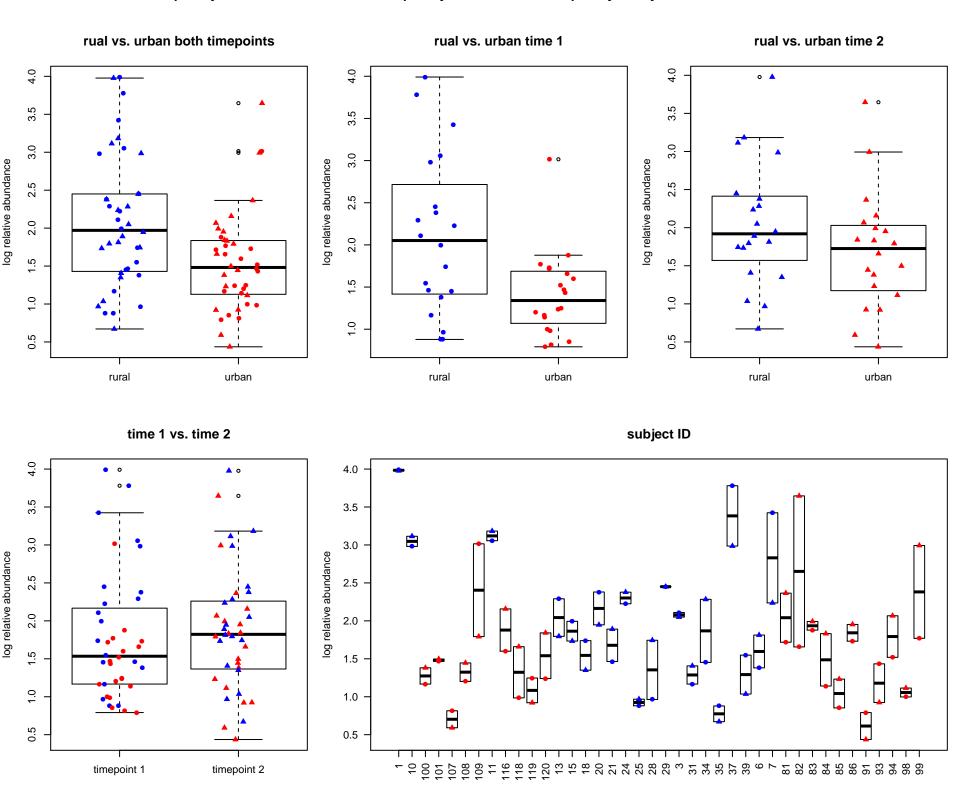
16S family: Campylobacteraceae pAdjRuralUrban=0.0961; pAdjTime=0.241; pAdjSubject=0.605



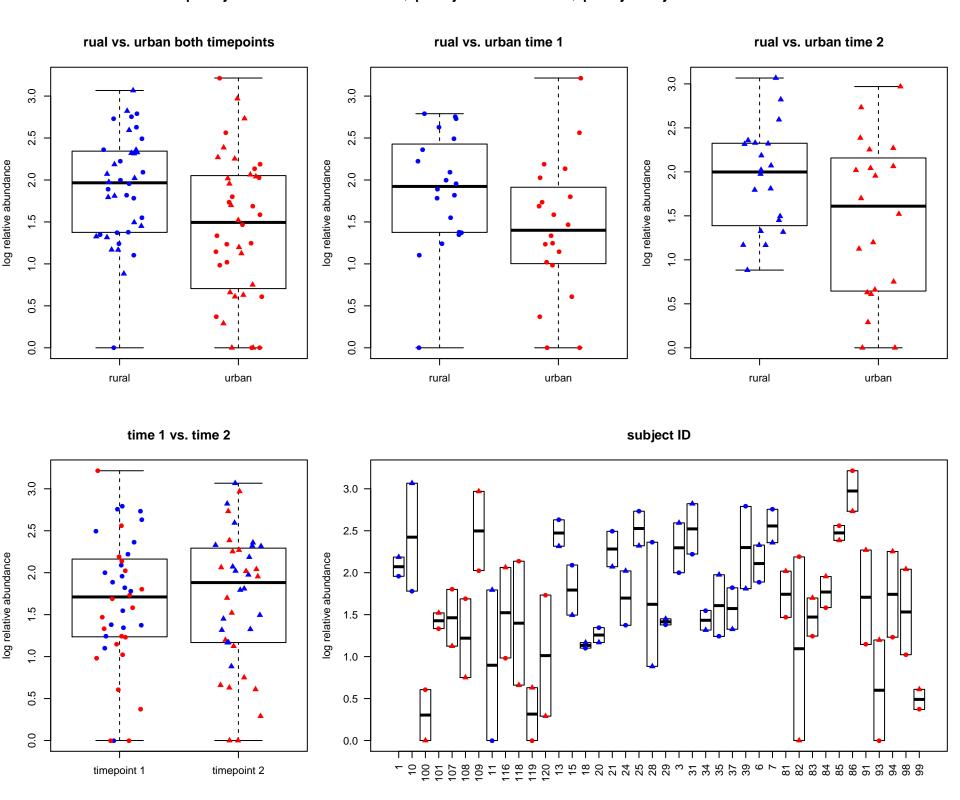
16S family: Pasteurellaceae pAdjRuralUrban=0.107; pAdjTime=0.35; pAdjSubject=1.07e-05



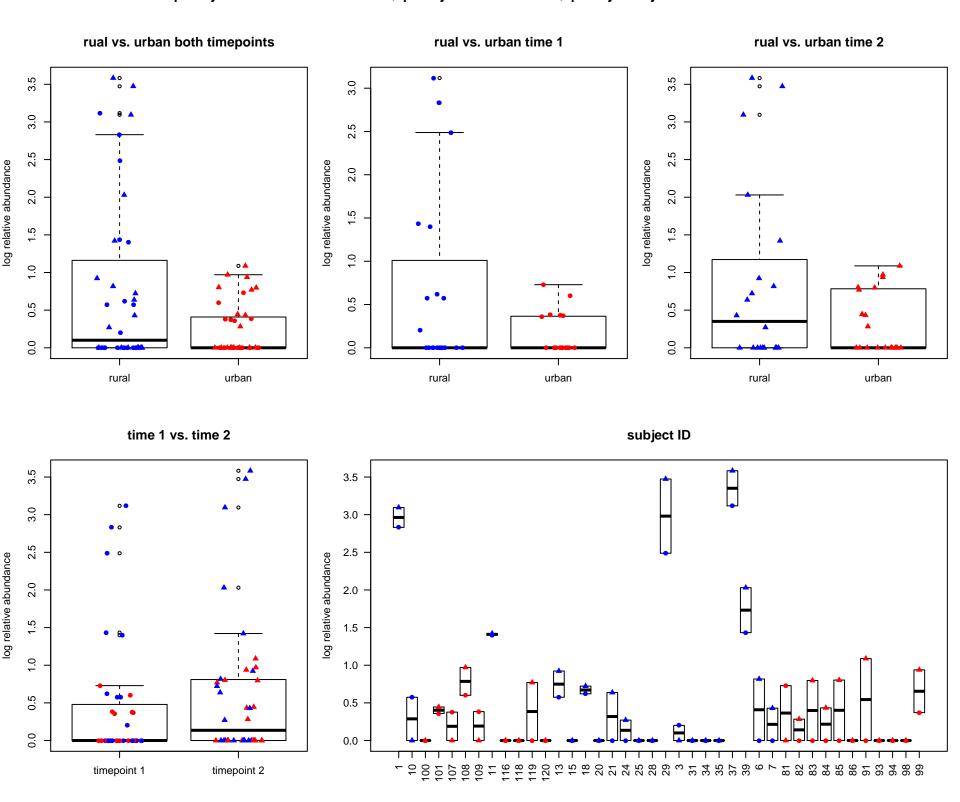
16S family: Streptococcaceae pAdjRuralUrban=0.146; pAdjTime=0.632; pAdjSubject=1.64e-06



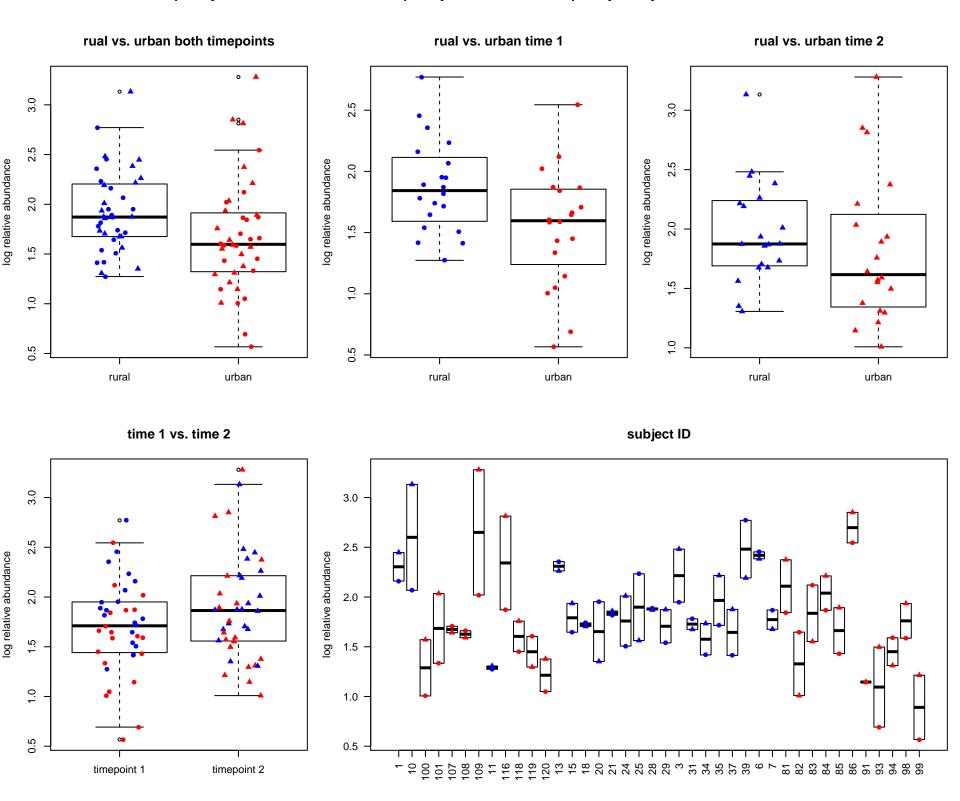
16S family: Clostridiaceae.1 pAdjRuralUrban=0.146; pAdjTime=0.909; pAdjSubject=0.0468



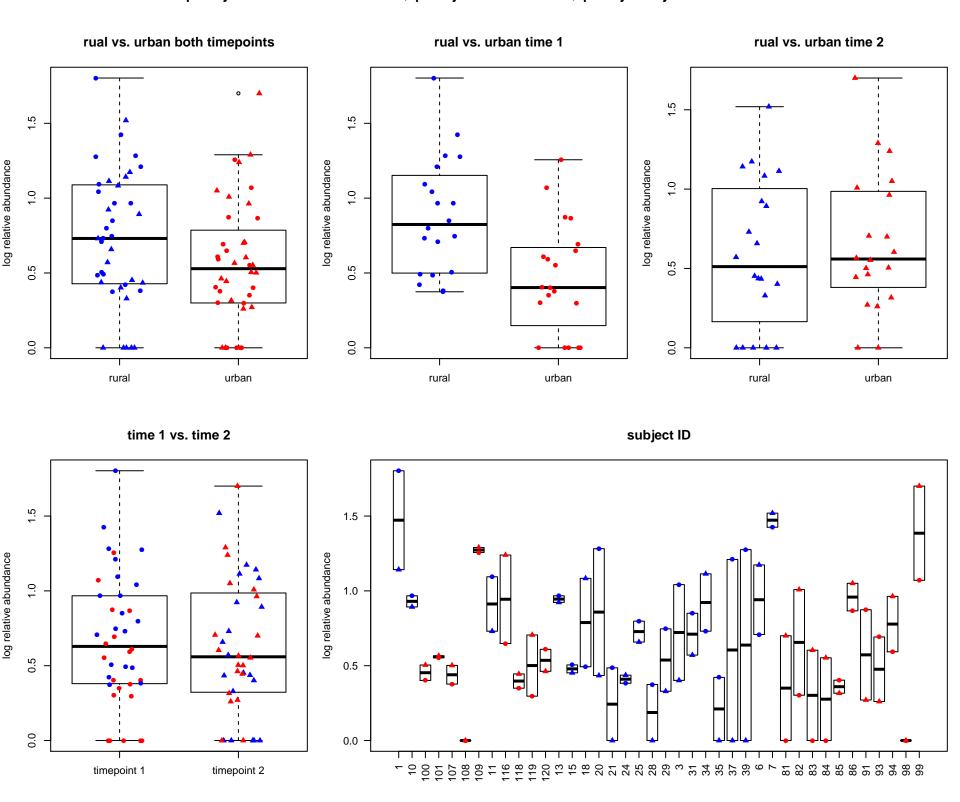
16S family: Succinivibrionaceae pAdjRuralUrban=0.218; pAdjTime=0.166; pAdjSubject=6.33e-13



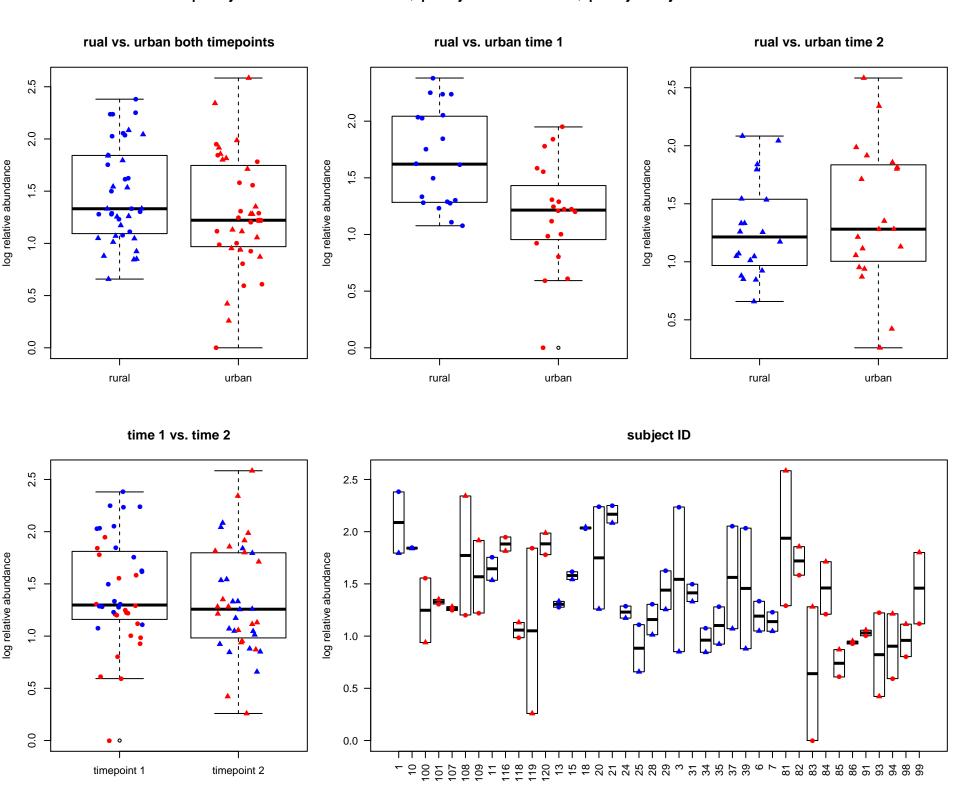
16S family: Peptostreptococcaceae pAdjRuralUrban=0.315; pAdjTime=0.174; pAdjSubject=0.000582



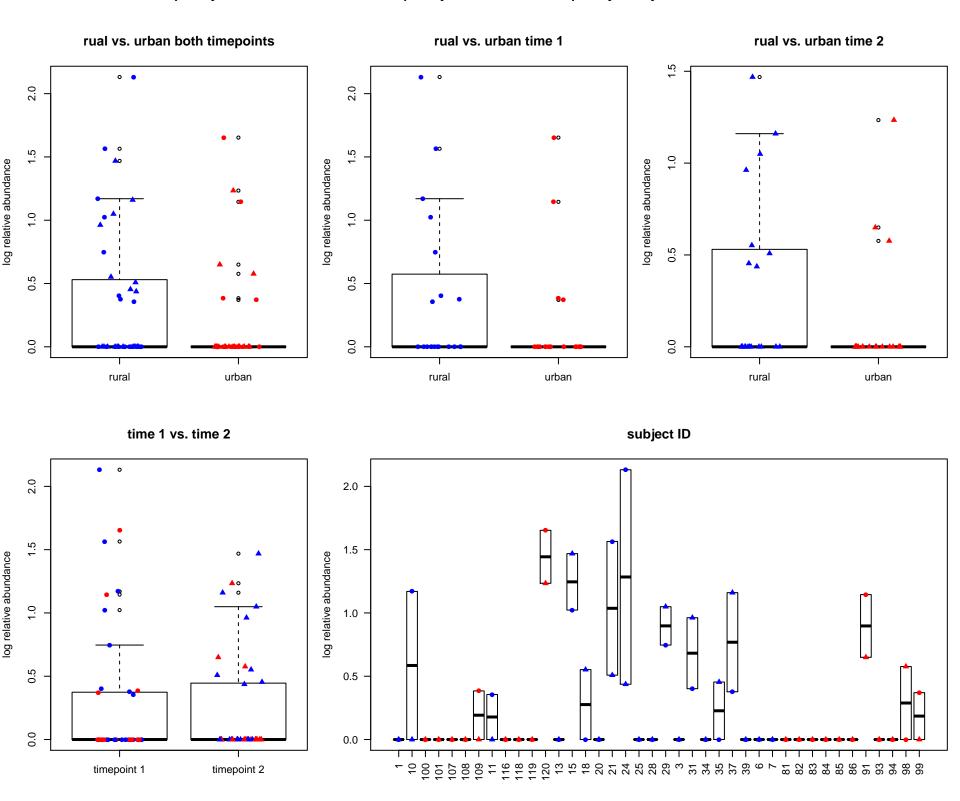
16S family: Actinomycetaceae pAdjRuralUrban=0.322; pAdjTime=0.827; pAdjSubject=0.0364



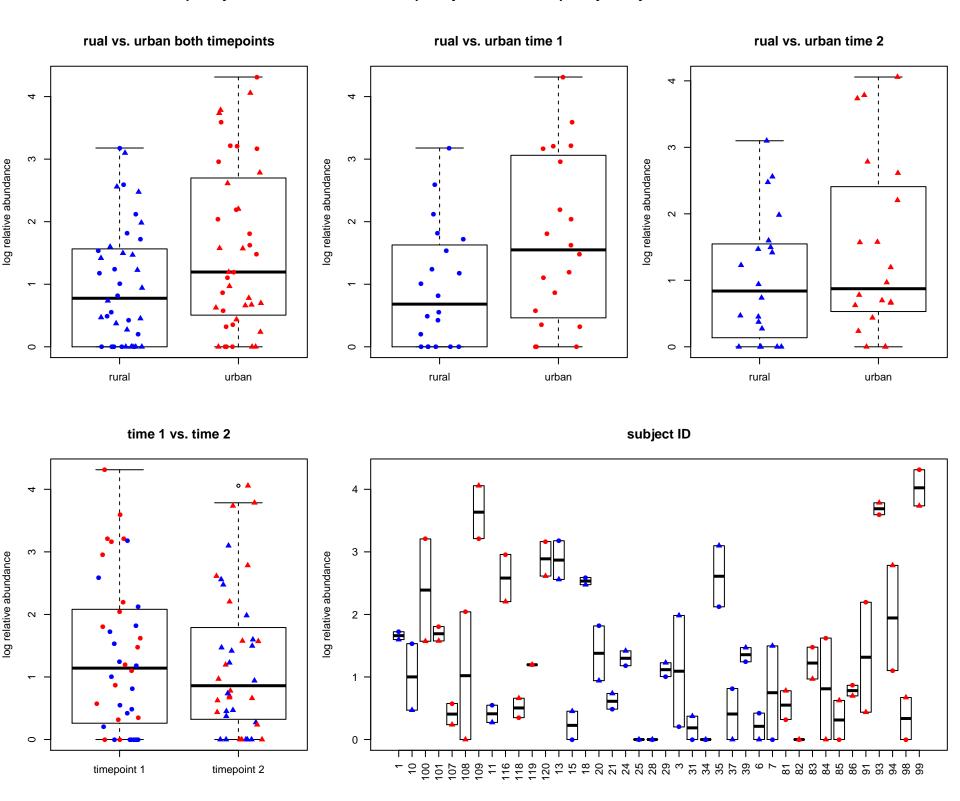
16S family: Coriobacteriaceae pAdjRuralUrban=0.322; pAdjTime=0.773; pAdjSubject=0.278



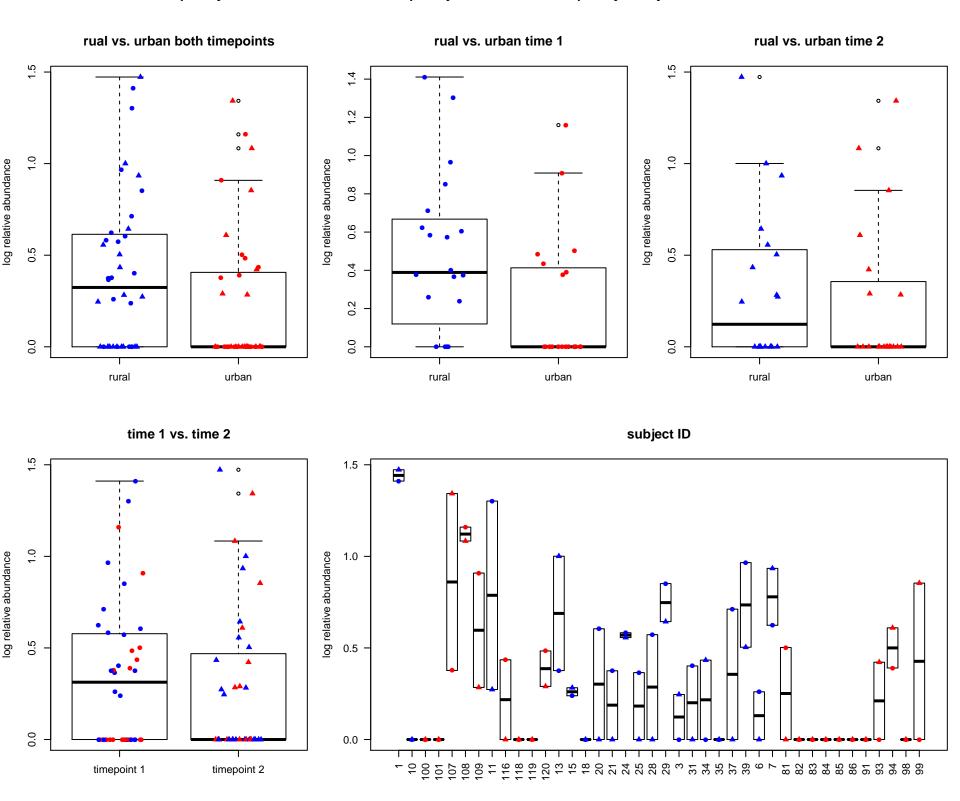
16S family: Oxalobacteraceae pAdjRuralUrban=0.322; pAdjTime=0.794; pAdjSubject=0.000507



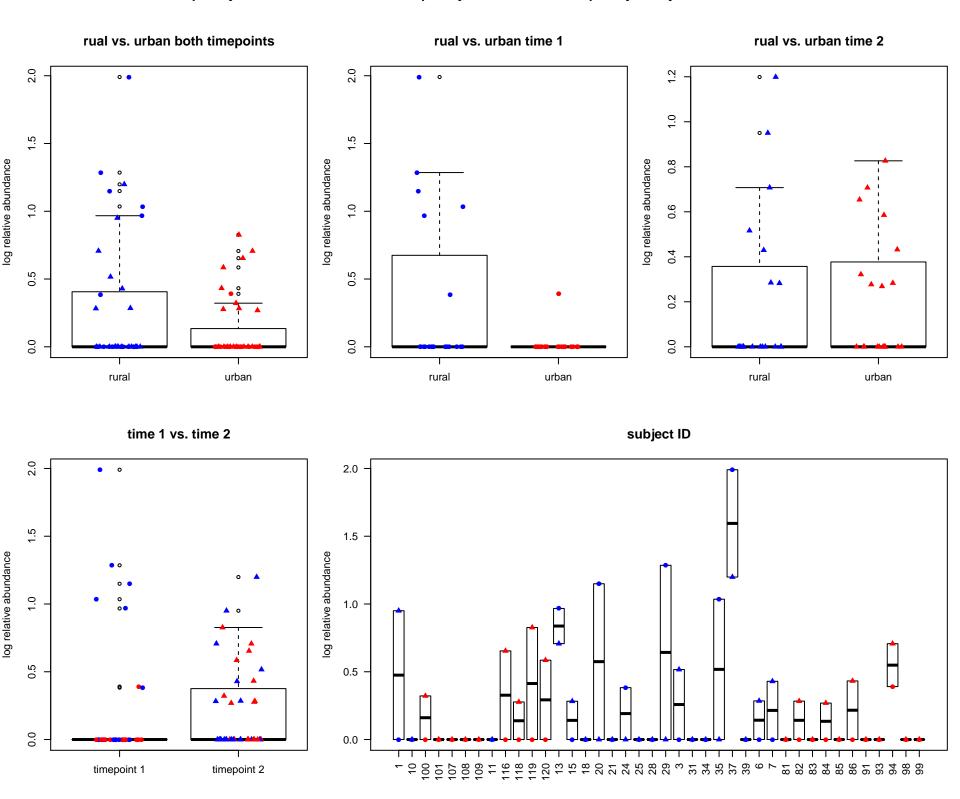
16S family: Fusobacteriaceae pAdjRuralUrban=0.322; pAdjTime=0.8; pAdjSubject=3.01e-07



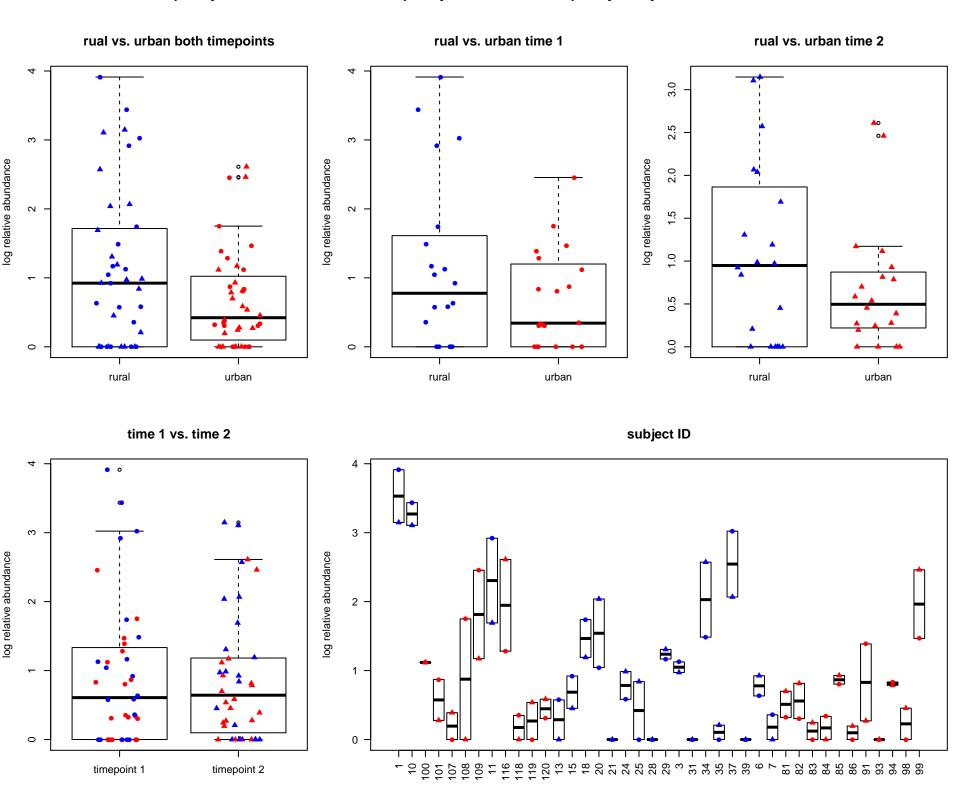
16S family: Clostridiales_Incertae.Sedis.XIII pAdjRuralUrban=0.322; pAdjTime=0.683; pAdjSubject=0.00158



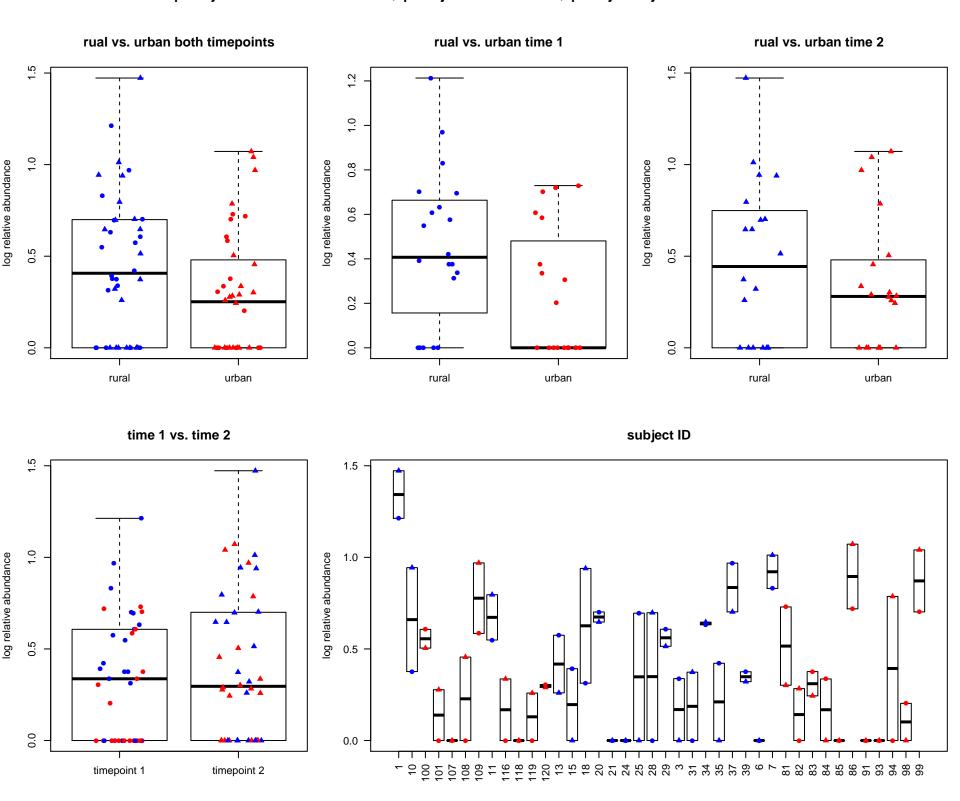
16S family: Enterococcaceae pAdjRuralUrban=0.322; pAdjTime=0.831; pAdjSubject=0.099



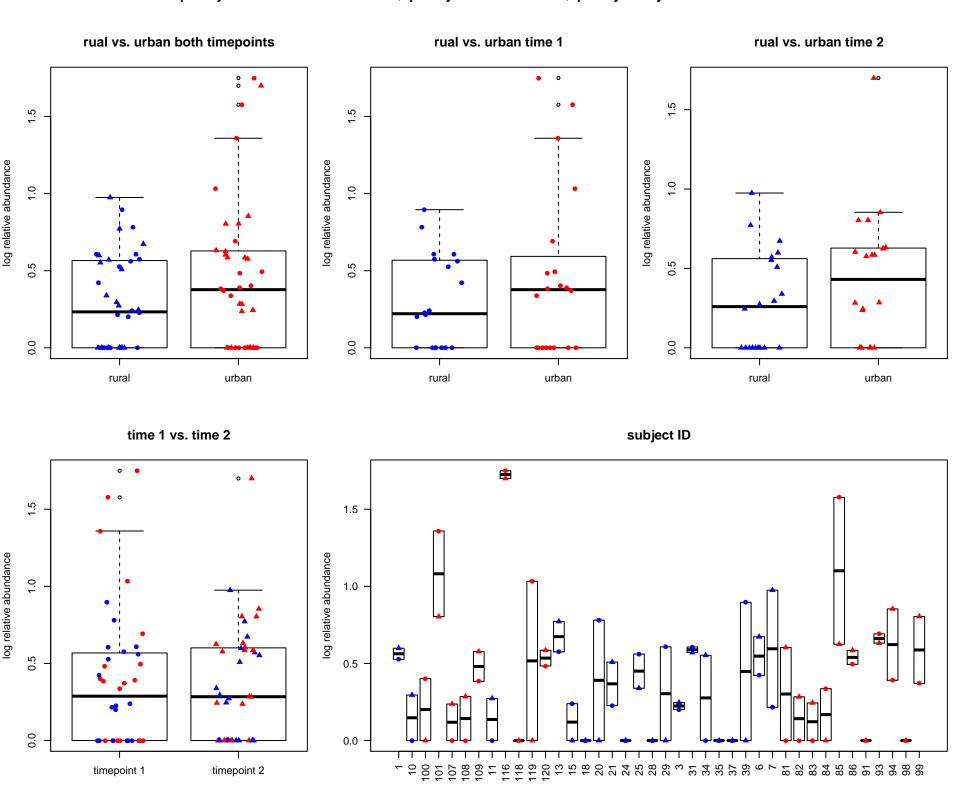
16S family: Lactobacillaceae pAdjRuralUrban=0.322; pAdjTime=0.909; pAdjSubject=6.91e-08



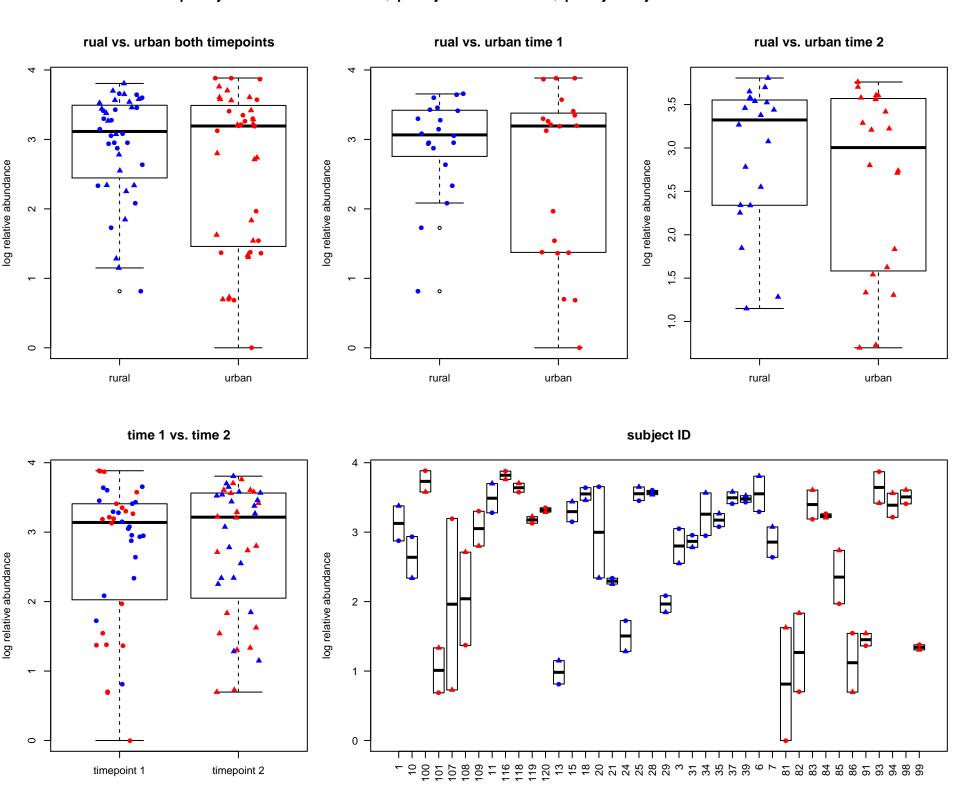
16S family: Bacillales_Incertae.Sedis.XI pAdjRuralUrban=0.322; pAdjTime=0.606; pAdjSubject=0.000194



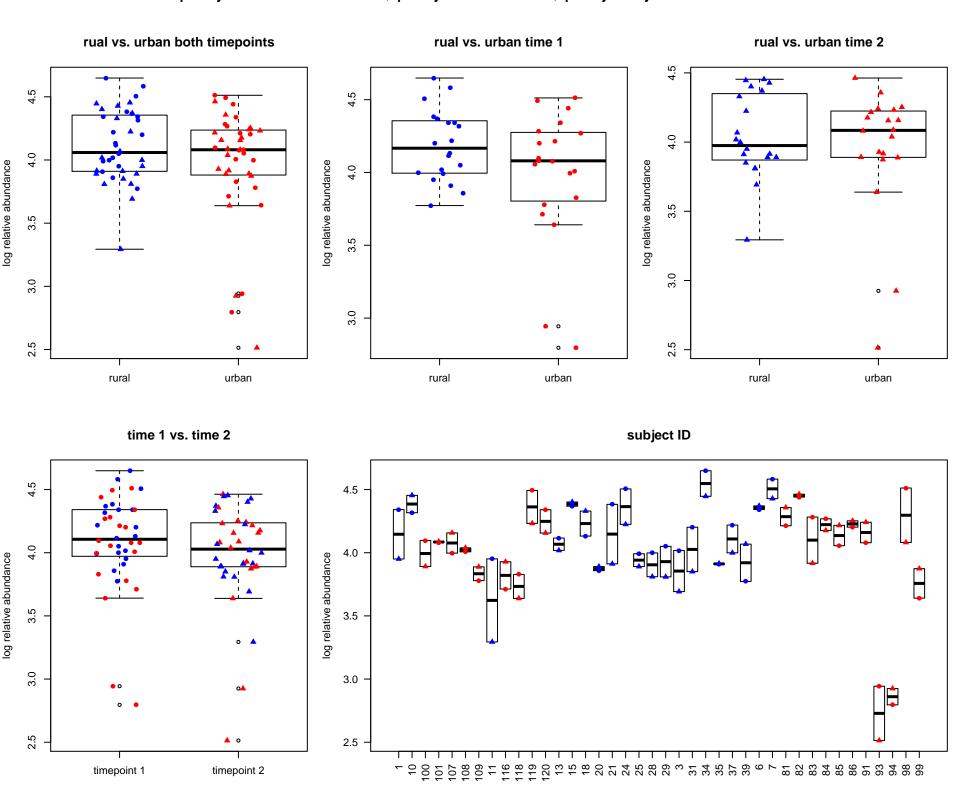
16S family: Eubacteriaceae pAdjRuralUrban=0.355; pAdjTime=0.916; pAdjSubject=0.00208



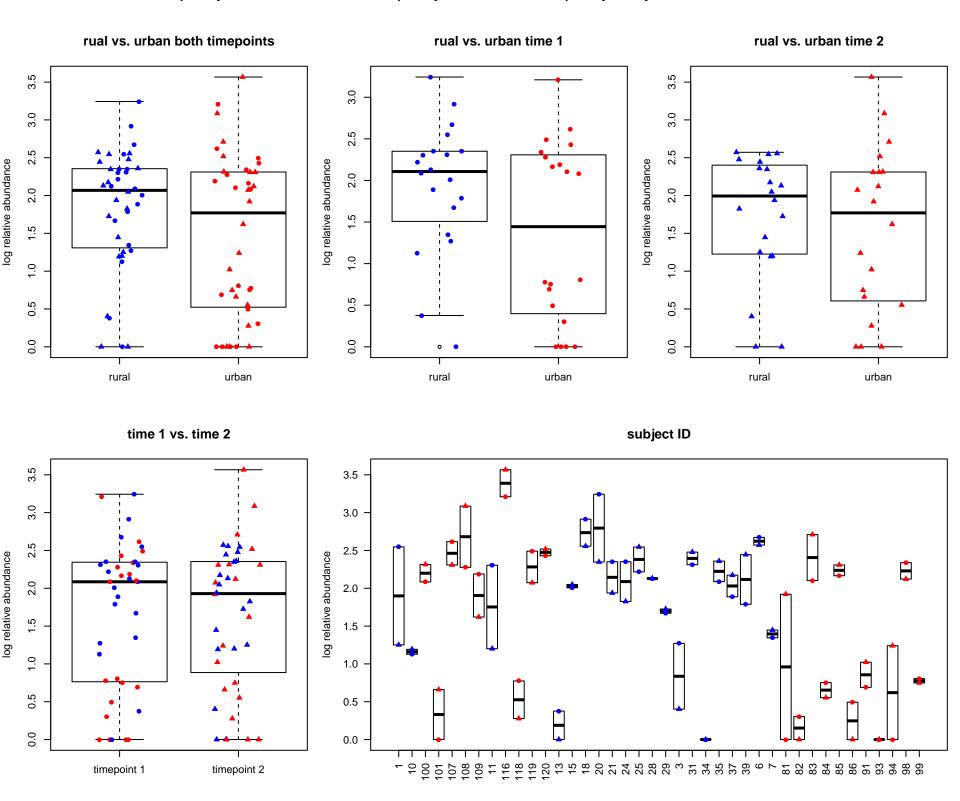
16S family: Acidaminococcaceae pAdjRuralUrban=0.43; pAdjTime=0.831; pAdjSubject=6.91e-08



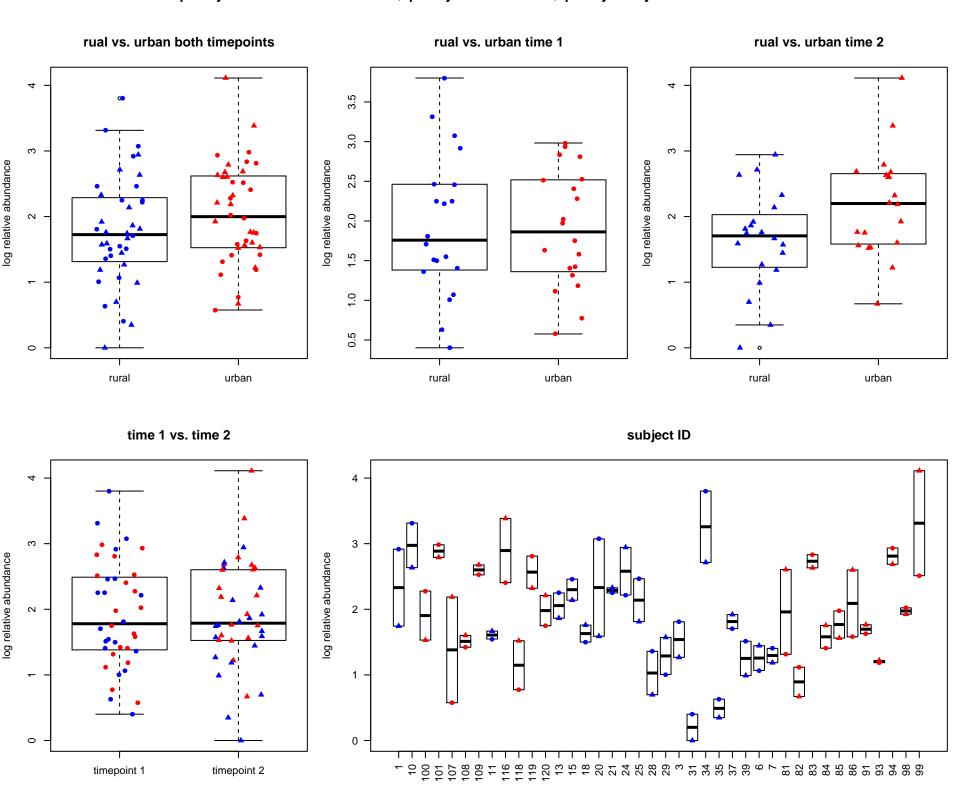
16S family: Ruminococcaceae pAdjRuralUrban=0.43; pAdjTime=0.174; pAdjSubject=5.23e-10



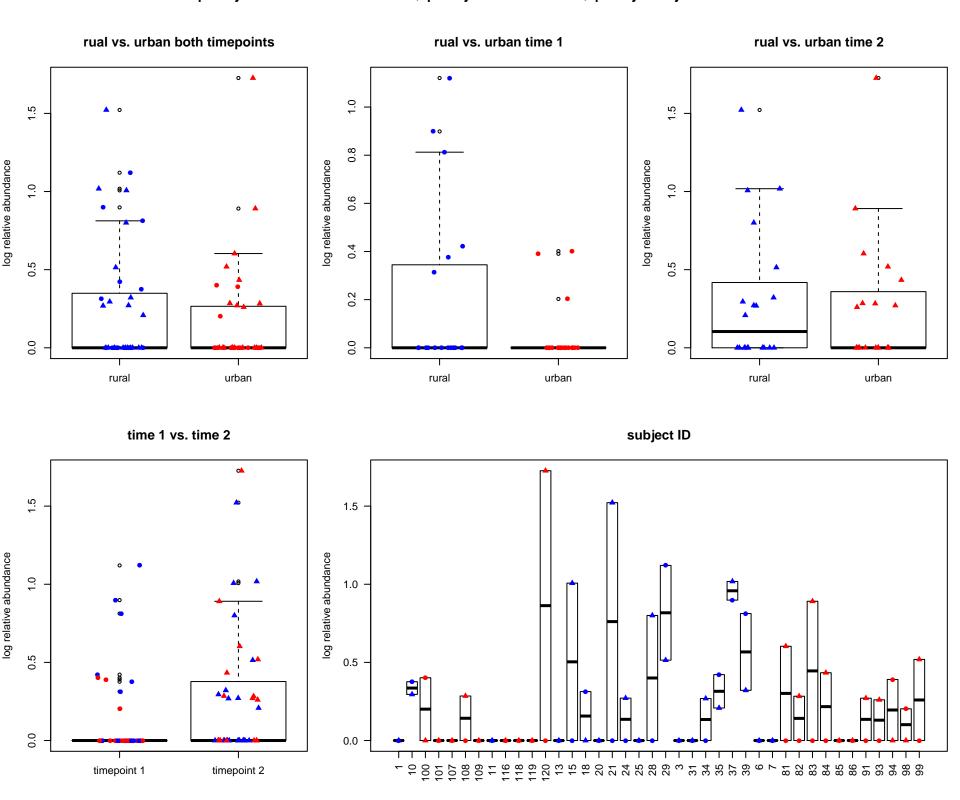
16S family: Desulfovibrionaceae pAdjRuralUrban=0.43; pAdjTime=0.916; pAdjSubject=1.35e-09



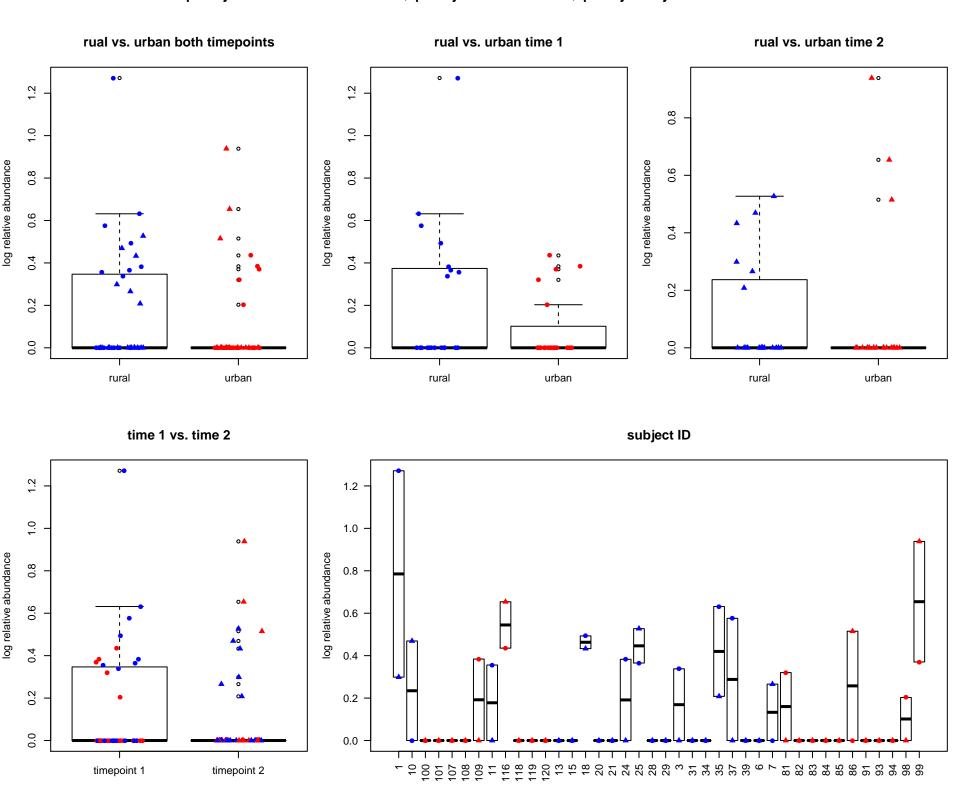
16S family: Bifidobacteriaceae pAdjRuralUrban=0.457; pAdjTime=0.96; pAdjSubject=3.96e-05



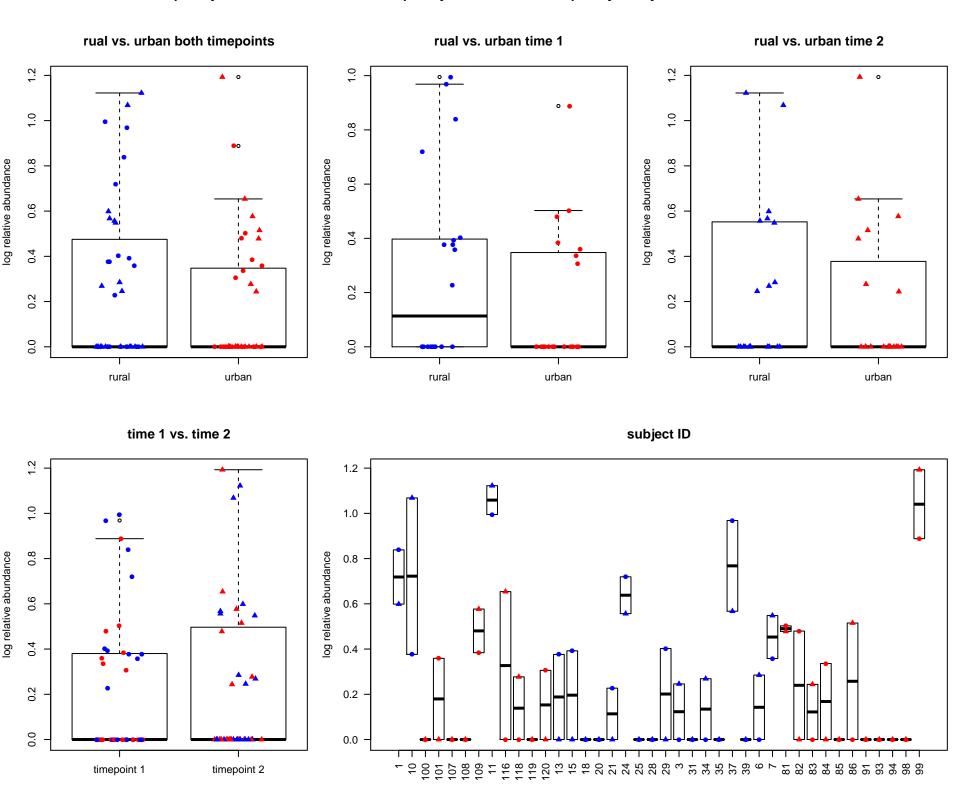
16S family: Comamonadaceae pAdjRuralUrban=0.457; pAdjTime=0.241; pAdjSubject=0.59



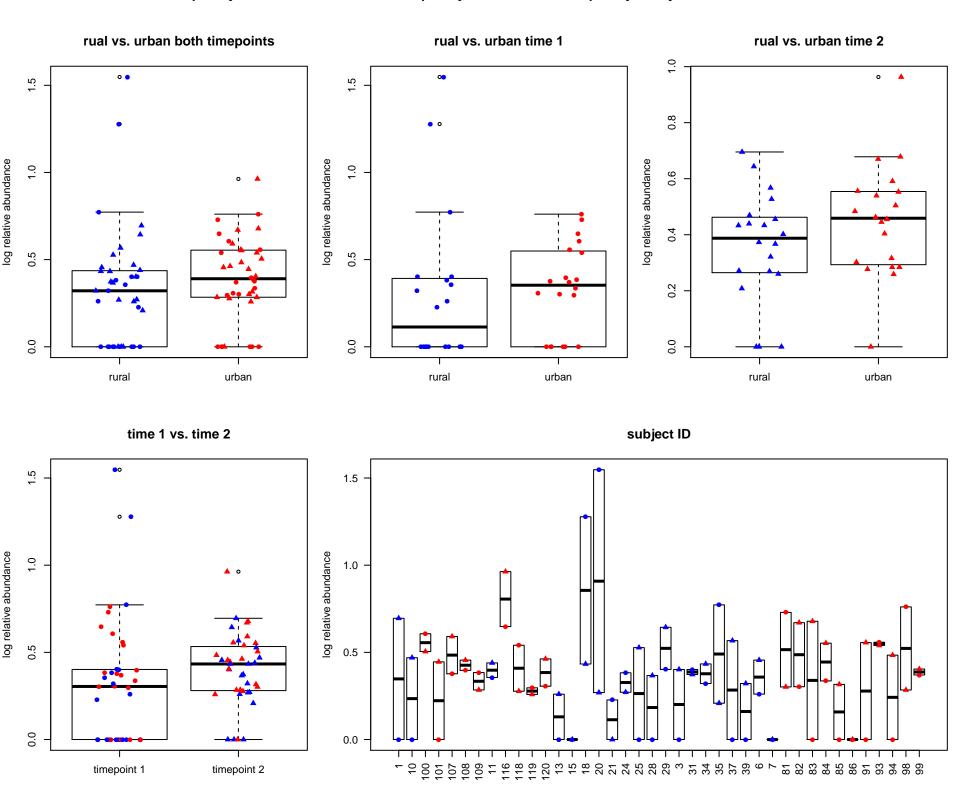
16S family: Corynebacteriaceae pAdjRuralUrban=0.479; pAdjTime=0.683; pAdjSubject=0.0255



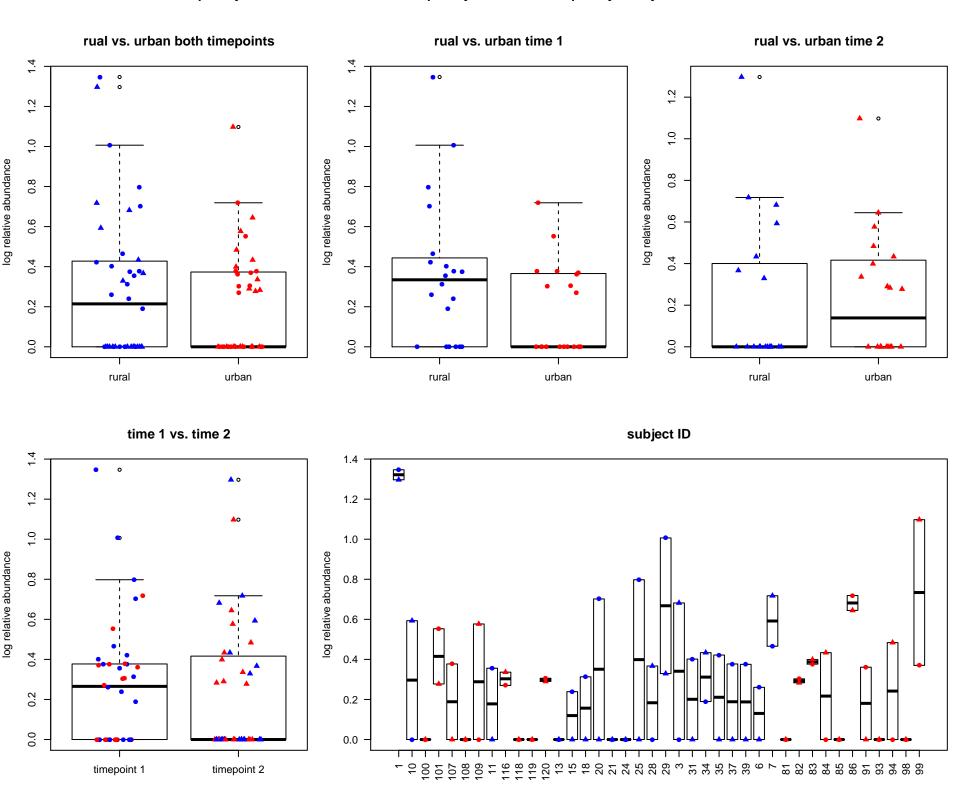
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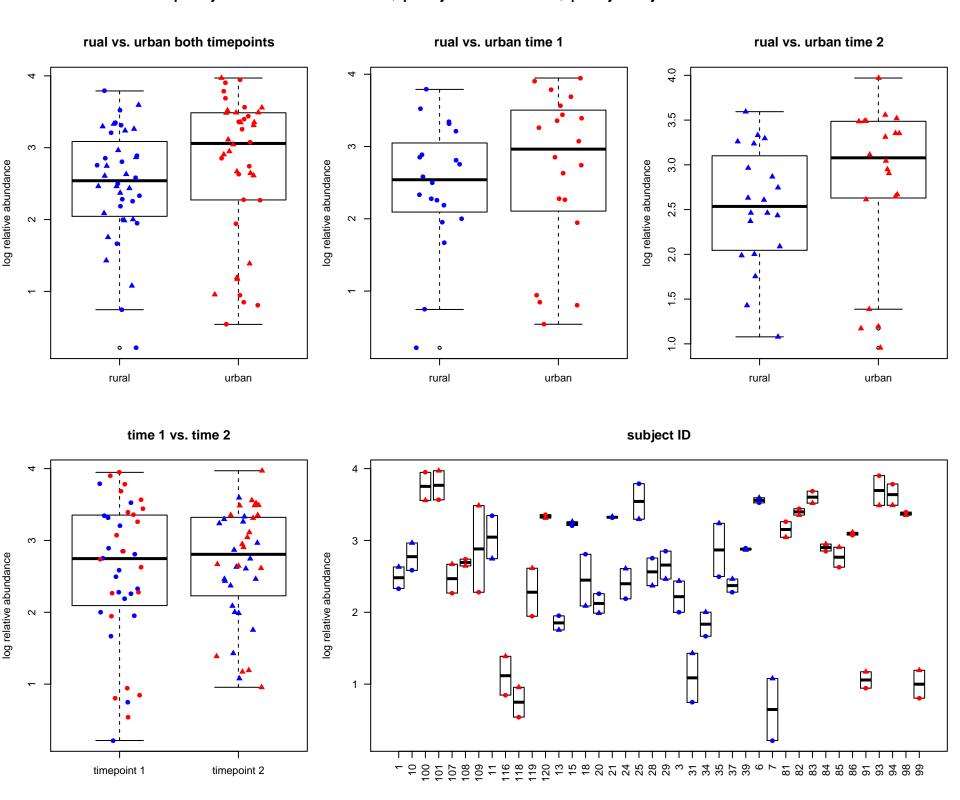


16S family: Shewanellaceae pAdjRuralUrban=0.508; pAdjTime=0.483; pAdjSubject=0.918

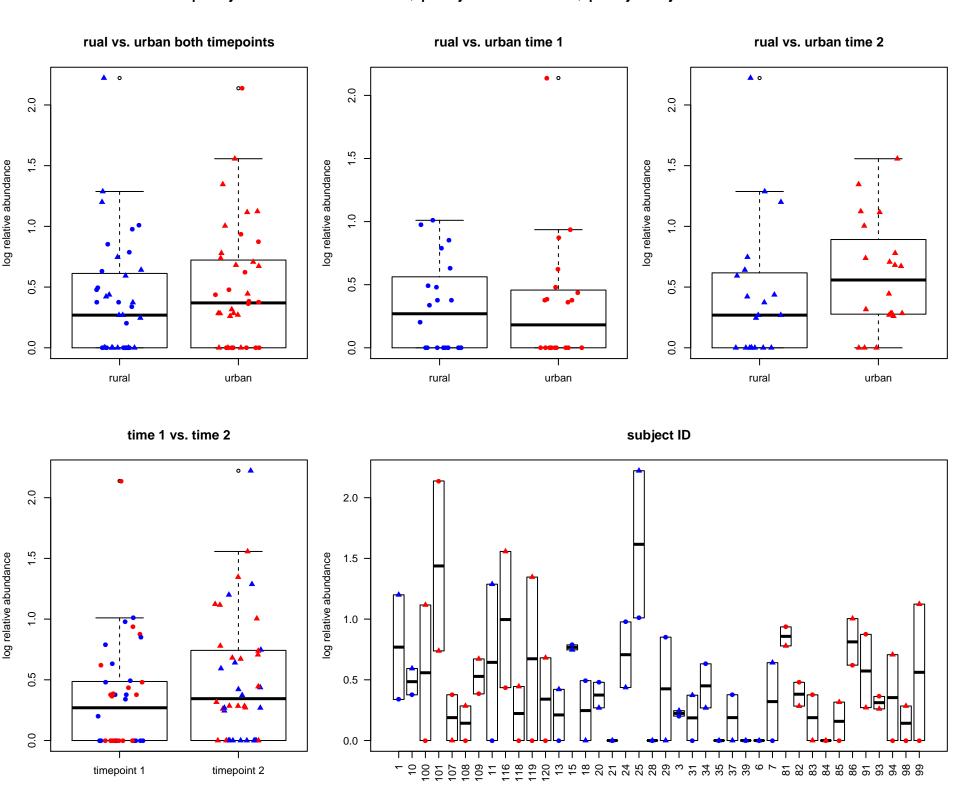


16S family: Carnobacteriaceae pAdjRuralUrban=0.508; pAdjTime=0.8; pAdjSubject=0.0333

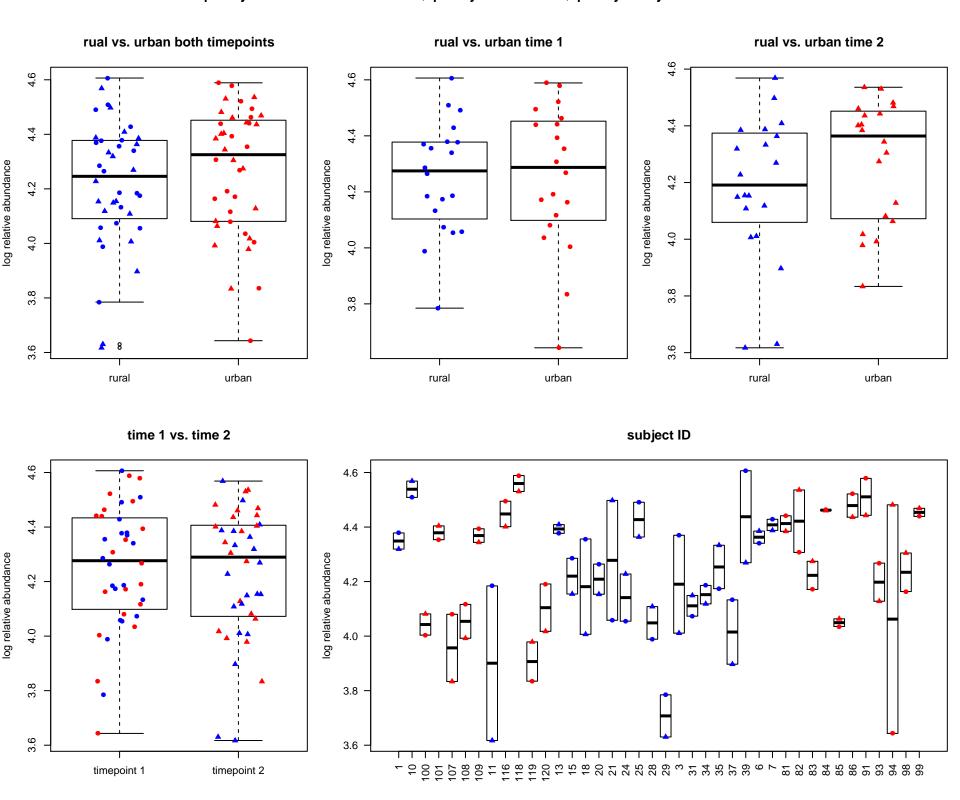




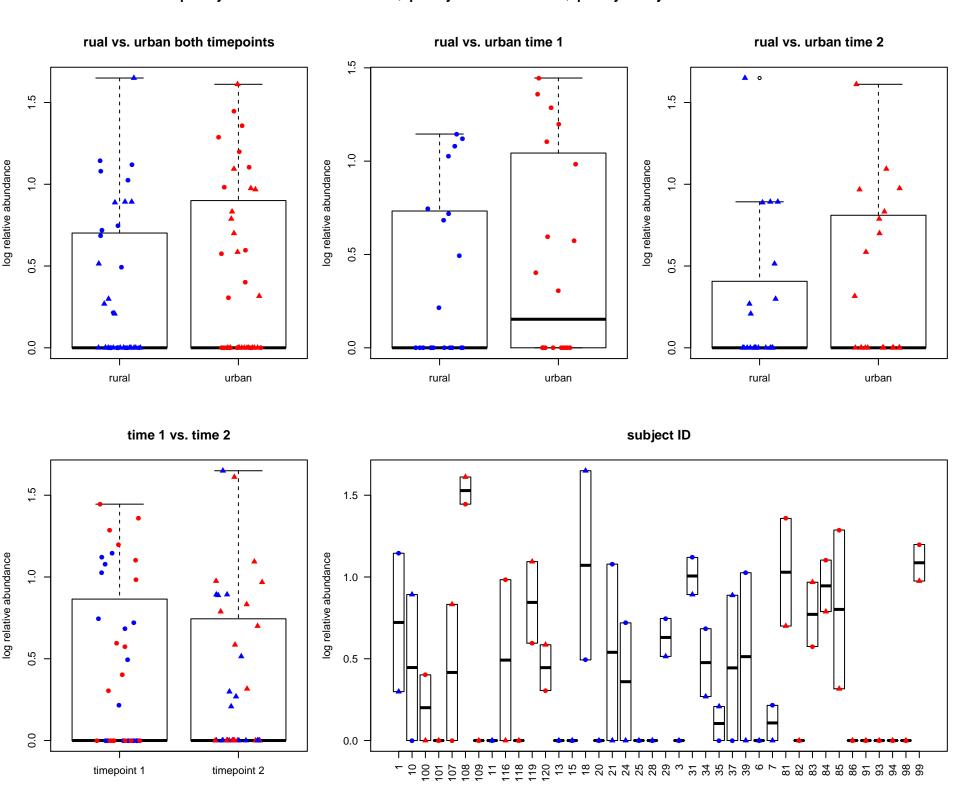
16S family: Chloroplast pAdjRuralUrban=0.576; pAdjTime=0.313; pAdjSubject=0.285



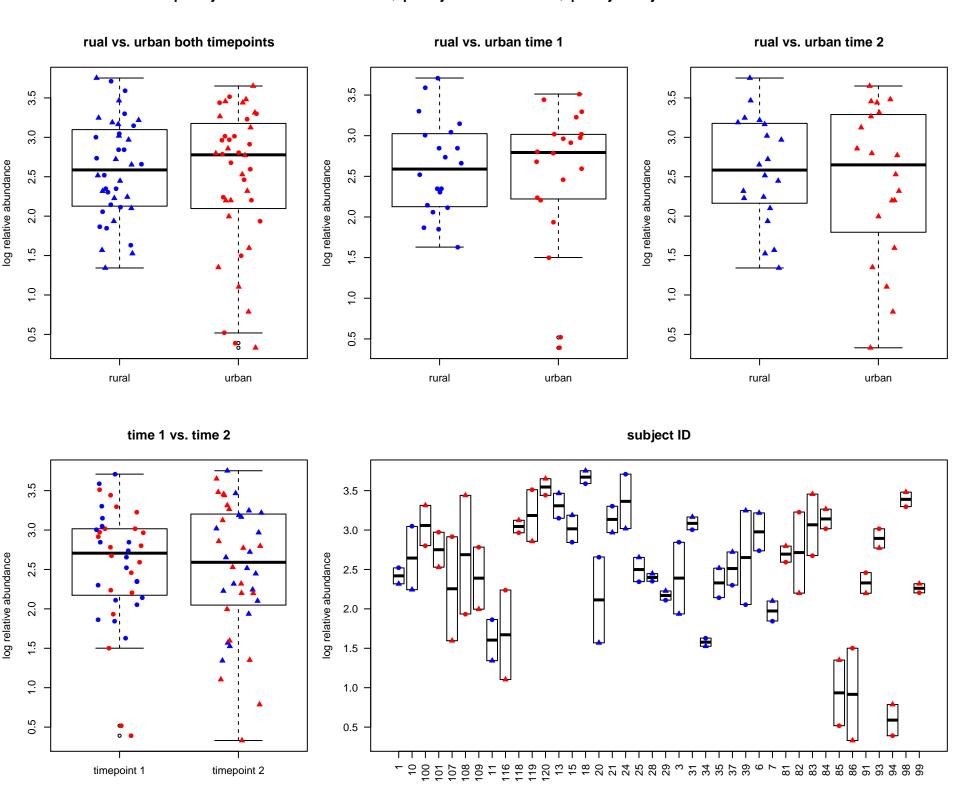
16S family: Lachnospiraceae pAdjRuralUrban=0.576; pAdjTime=0.8; pAdjSubject=0.001



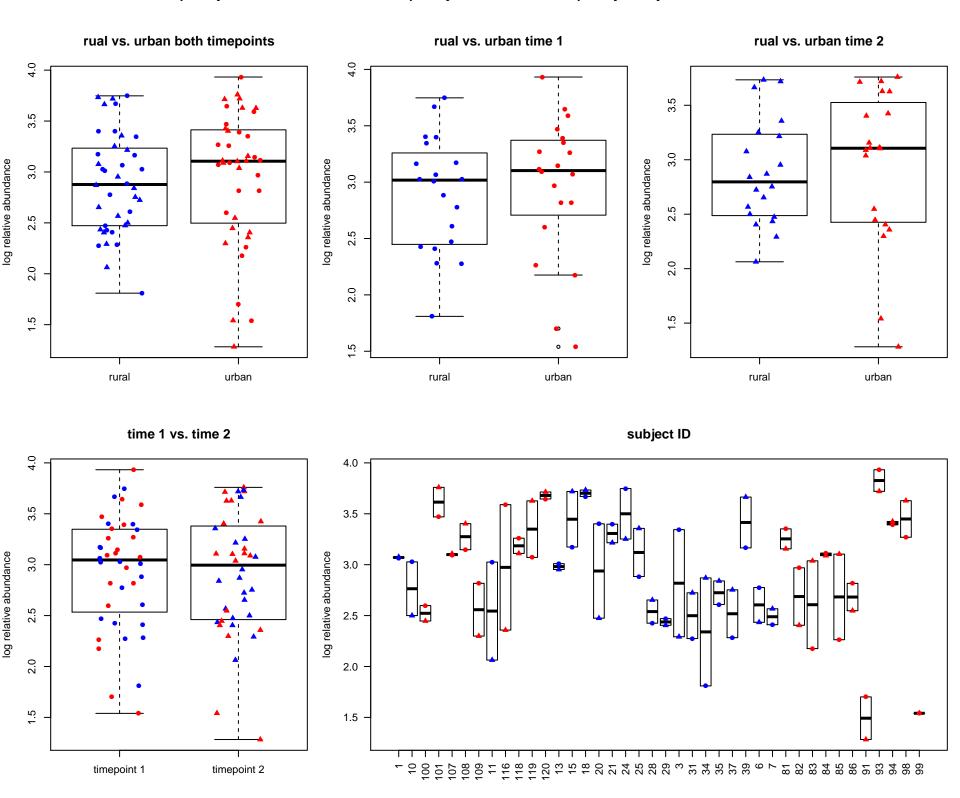
16S family: Synergistaceae pAdjRuralUrban=0.576; pAdjTime=0.766; pAdjSubject=0.00433



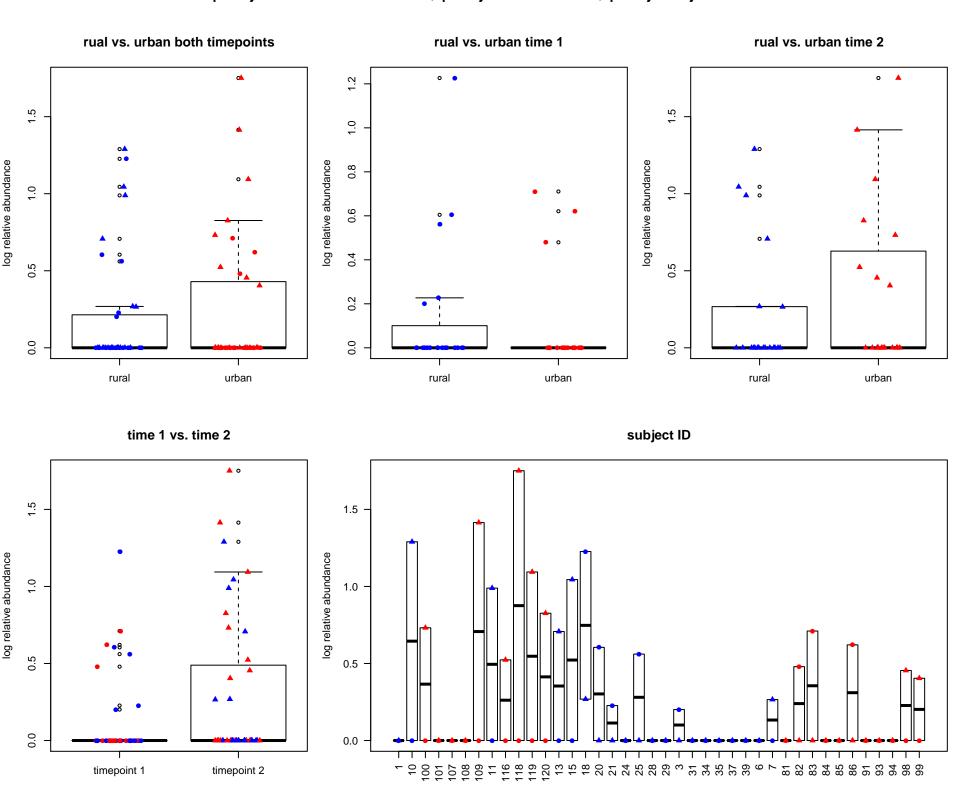
16S family: Rikenellaceae pAdjRuralUrban=0.765; pAdjTime=0.827; pAdjSubject=1.07e-05



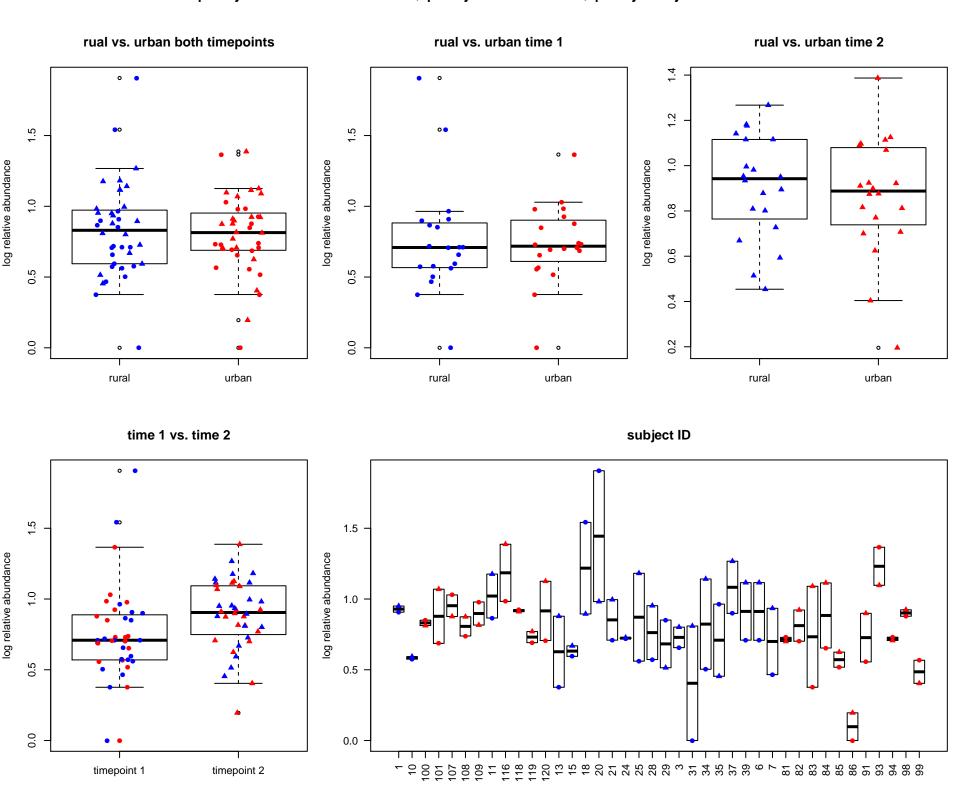
16S family: Porphyromonadaceae pAdjRuralUrban=0.774; pAdjTime=0.916; pAdjSubject=5.1e-05



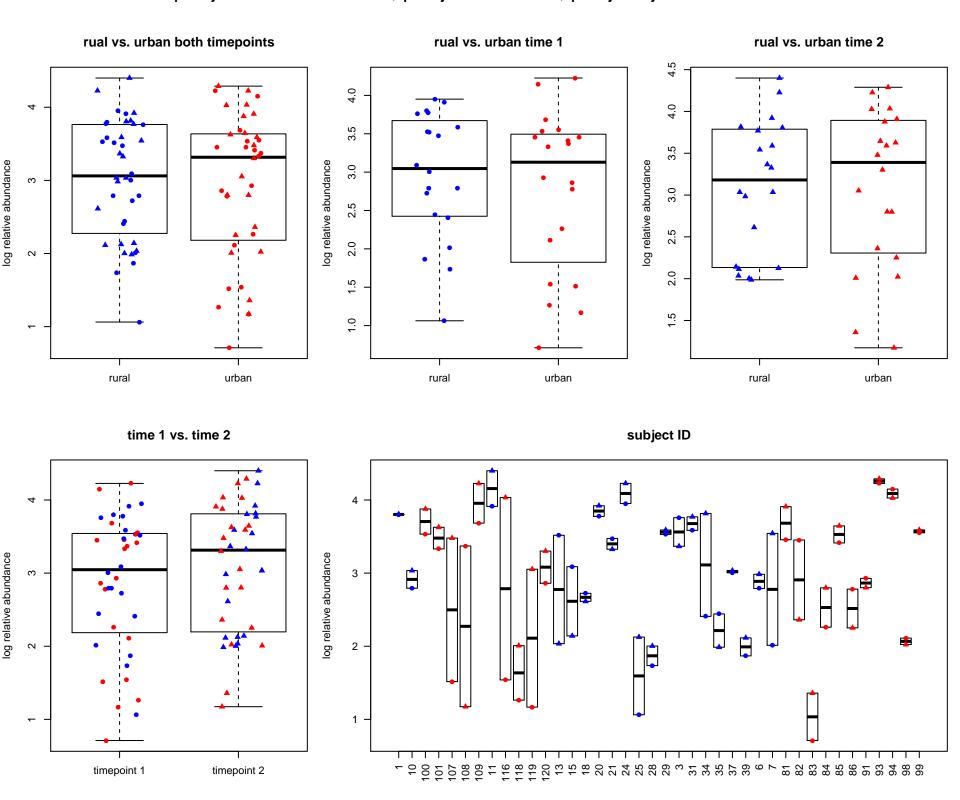
16S family: Flavobacteriaceae pAdjRuralUrban=0.774; pAdjTime=0.245; pAdjSubject=1



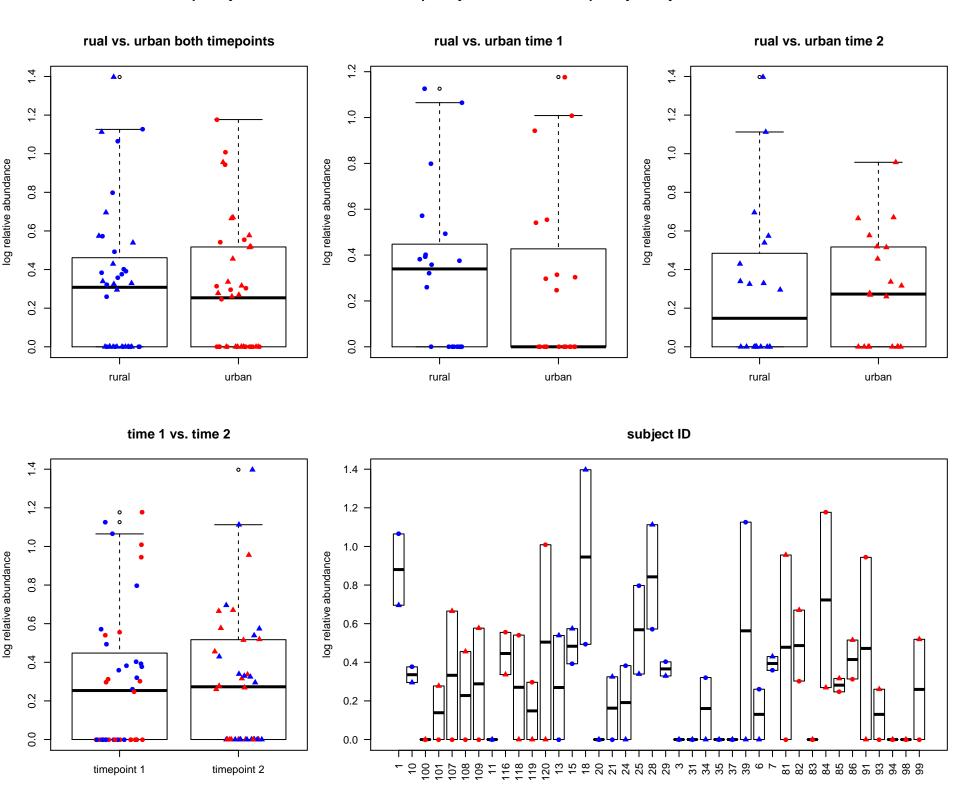
16S family: Halomonadaceae pAdjRuralUrban=0.774; pAdjTime=0.174; pAdjSubject=0.12



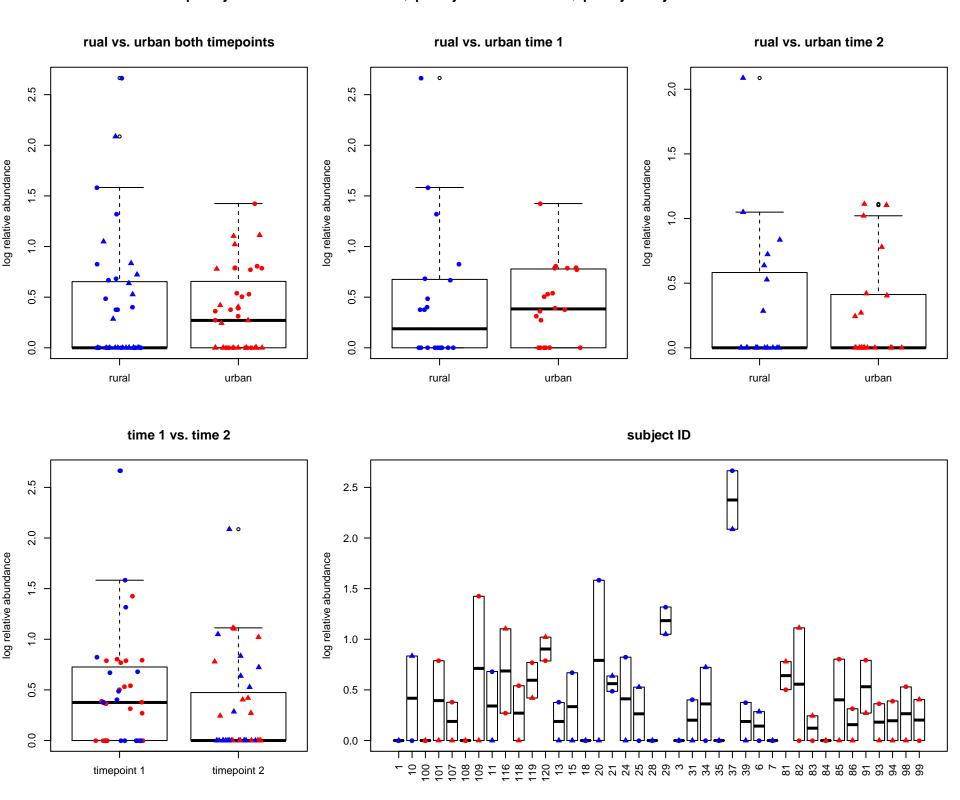
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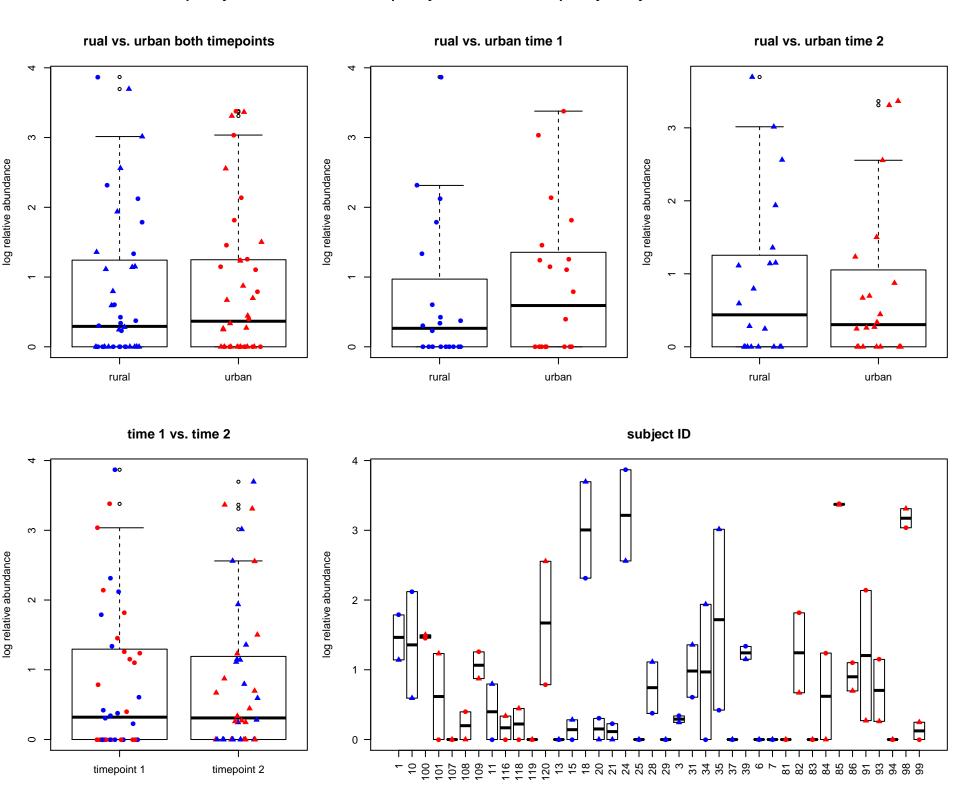
16S family: Clostridiales_Incertae.Sedis.XI pAdjRuralUrban=0.774; pAdjTime=0.979; pAdjSubject=0.627



16S family: Leuconostocaceae pAdjRuralUrban=0.826; pAdjTime=0.313; pAdjSubject=0.0131



16S family: Verrucomicrobiaceae pAdjRuralUrban=0.9; pAdjTime=0.902; pAdjSubject=1.07e-05



16S family: Erysipelotrichaceae pAdjRuralUrban=0.908; pAdjTime=0.8; pAdjSubject=0.0674

