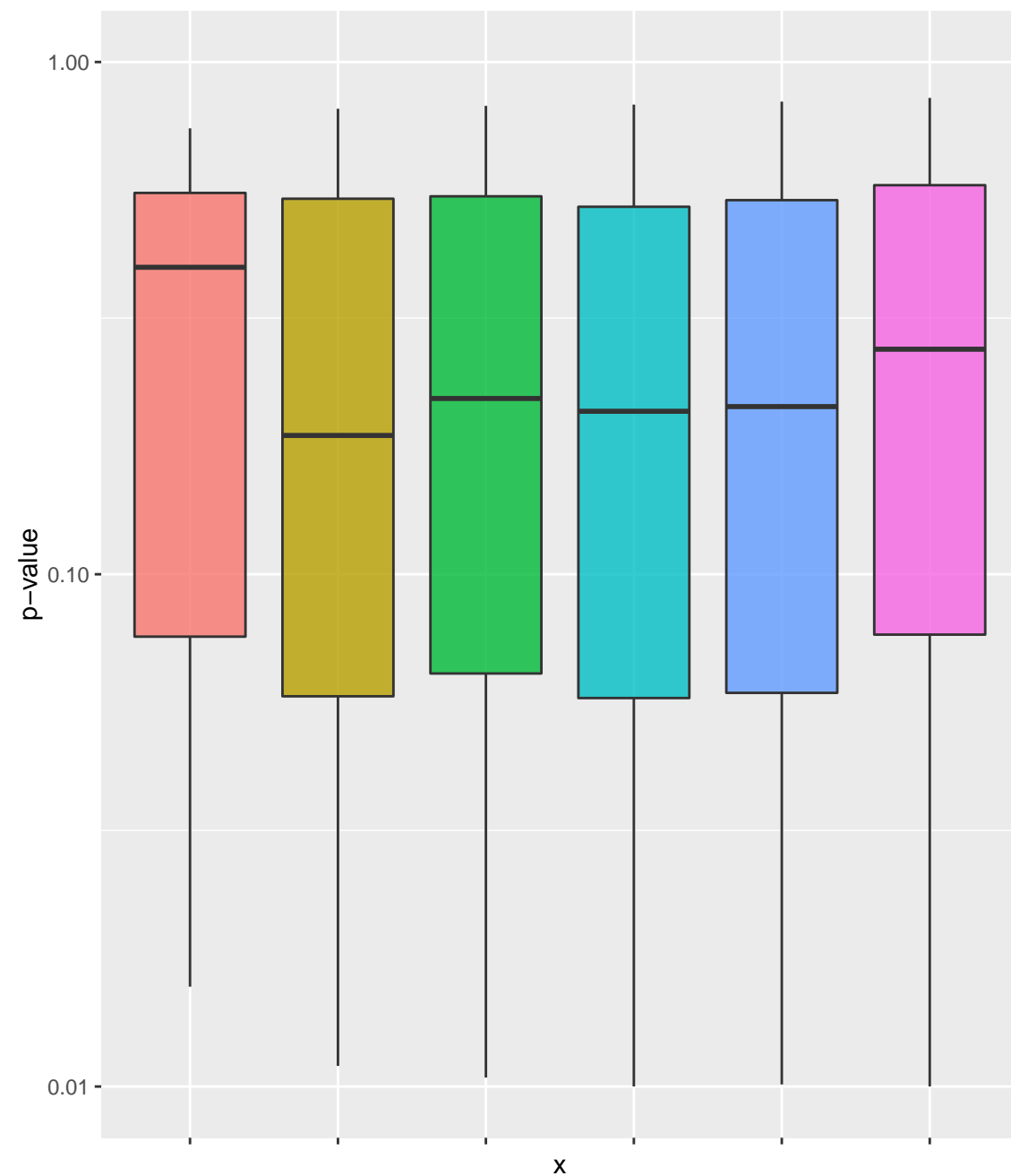
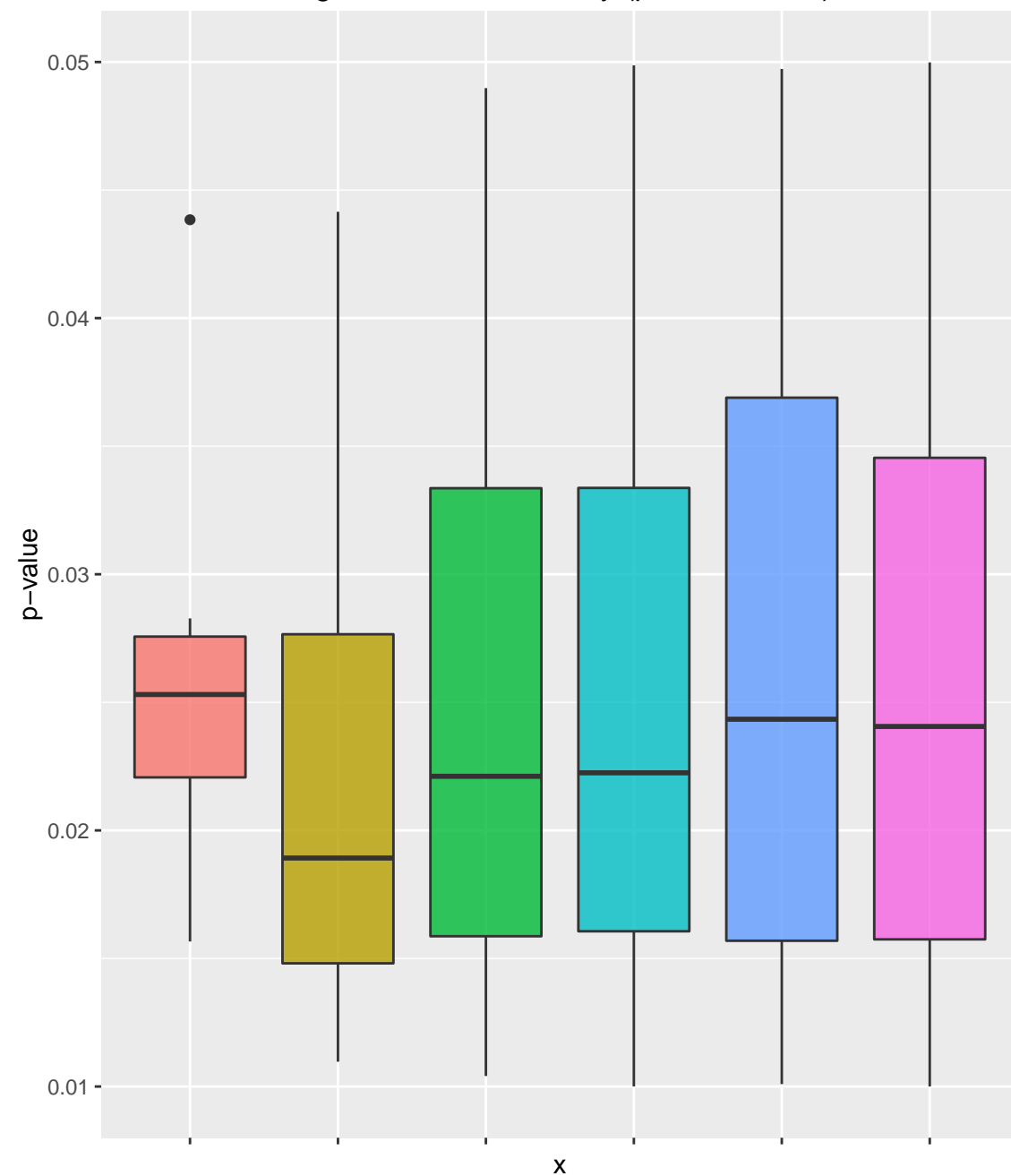


All features



fun.vals go.vals eff.vals path.vals mini.genes.vals other.genes.vals

Significant features only (p-value<0.05)



fun.vals go.vals eff.vals path.vals mini.genes.vals other.genes.vals