

Additional file 8

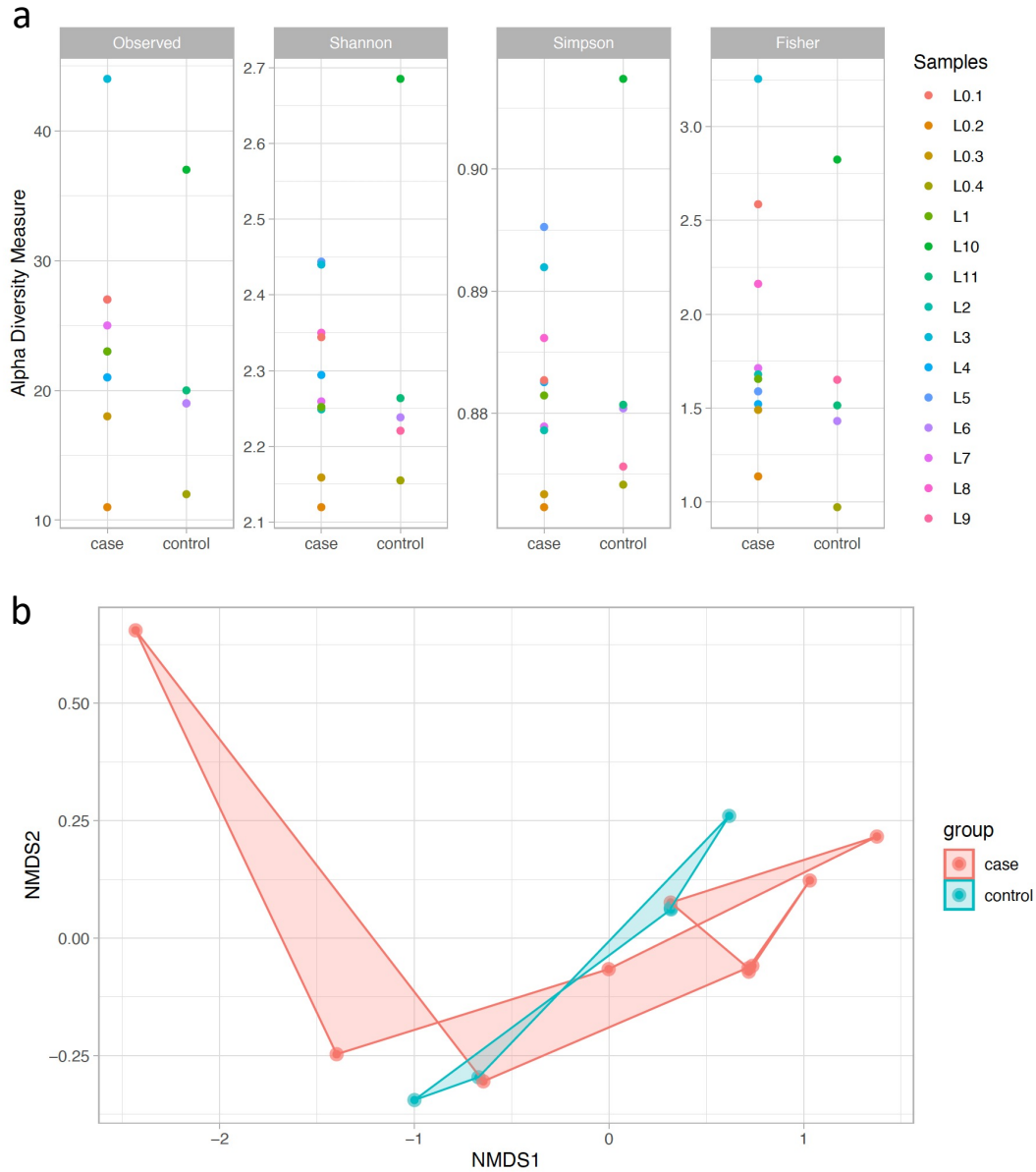


Fig: Alpha and beta diversity of the metatranscriptomic sequence libraries obtained from animals with clinical bovine respiratory disease (cases) and without clinical disease (controls). **Fig(a):** The alpha diversity computed using different methods, no significant differences in the alpha diversity were observed between cases and control. **Fig(b).** NMDS plot using Bray-Curtis dissimilarities between samples in cases and controls. There were no significant differences in the beta diversities between cases and controls.