

b

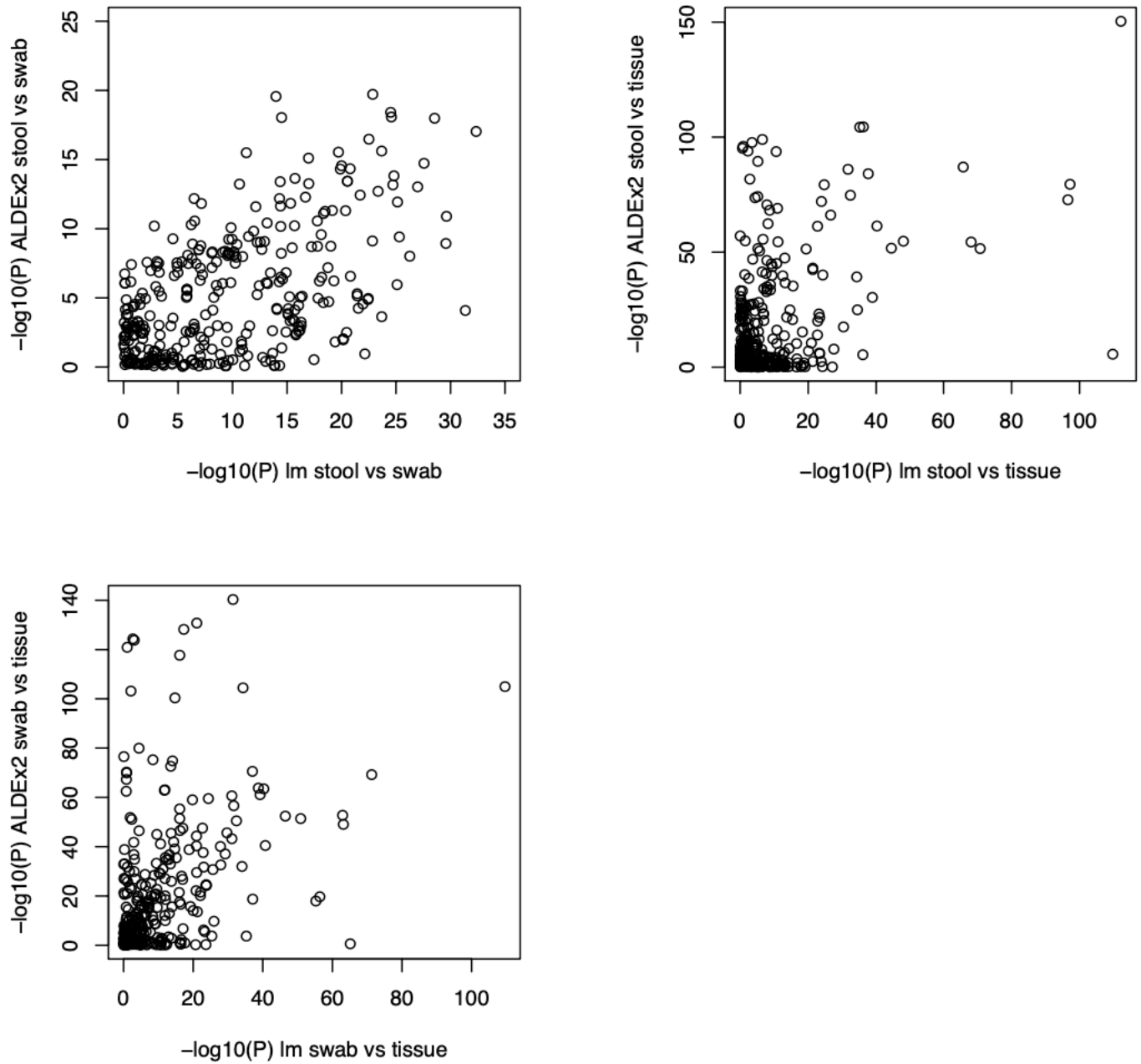


Fig. S3. Comparison of P values from ALDEx2 and mixed effects linear models for each pair of sample types for genus (a) and pathway abundance (b).