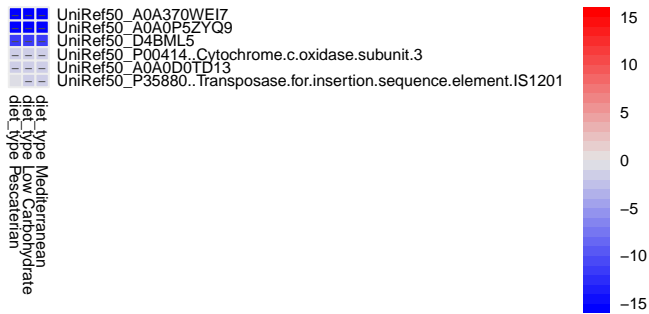
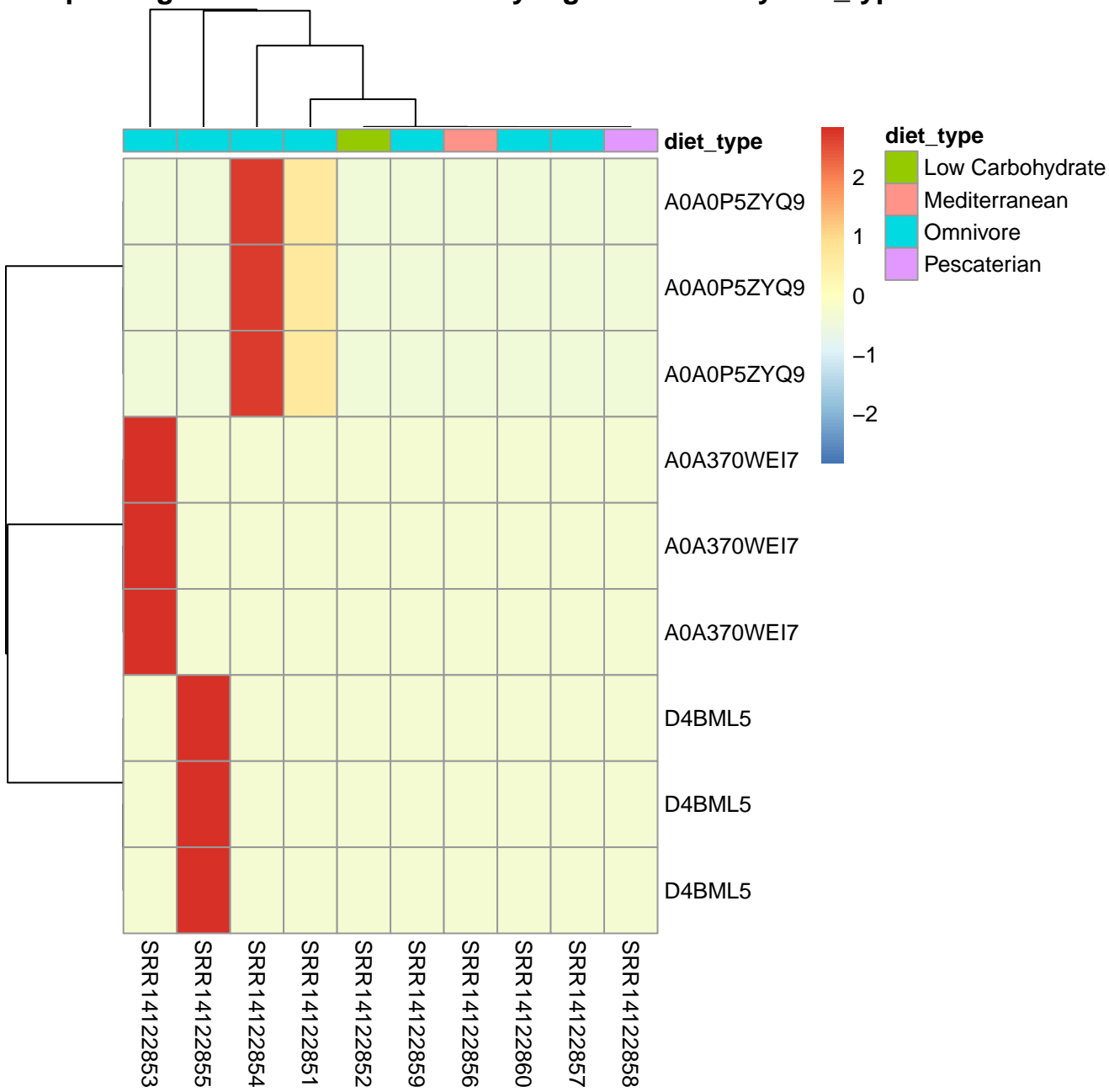


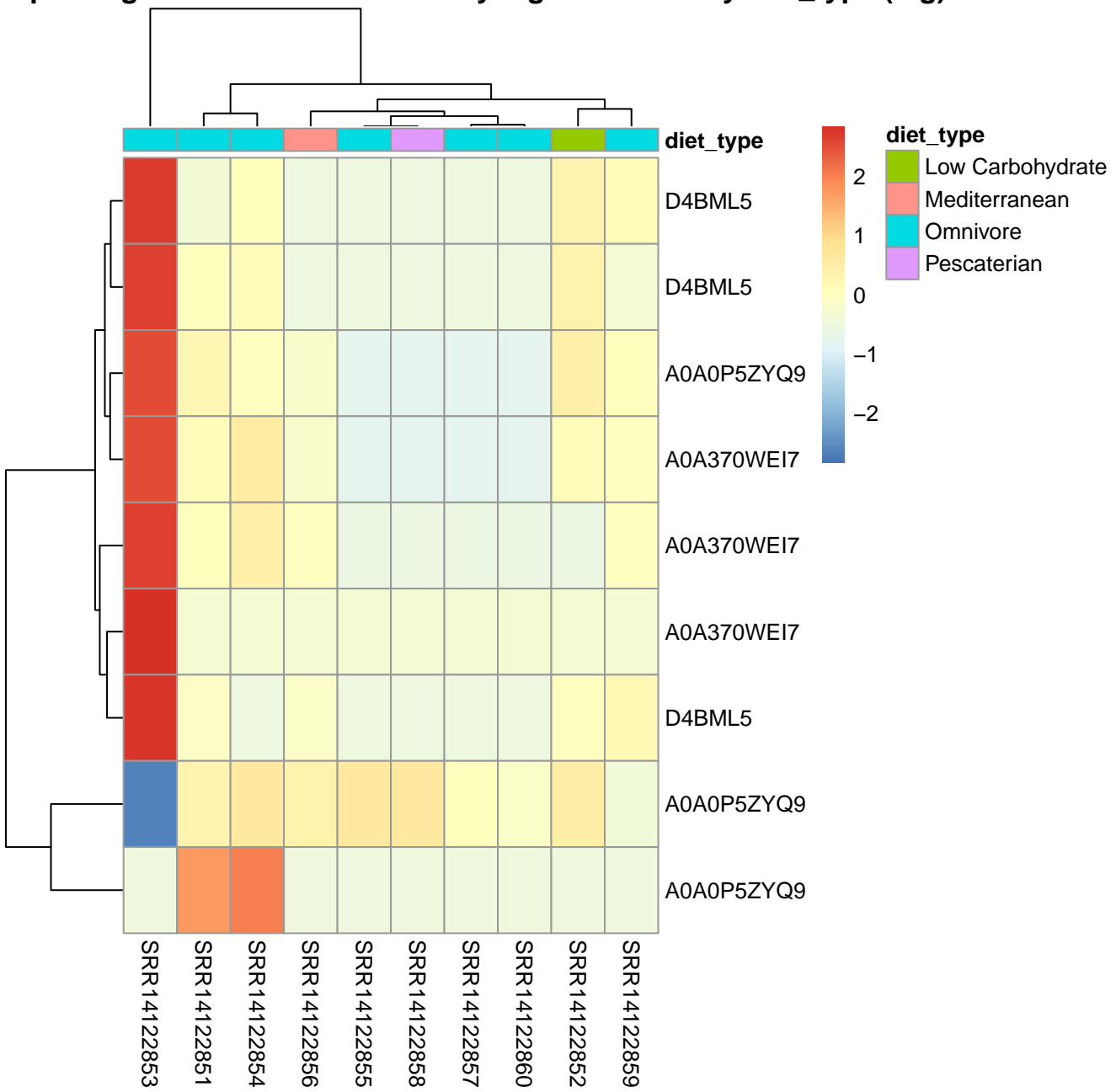
Significant associations (−log(qval)*sign(coeff))



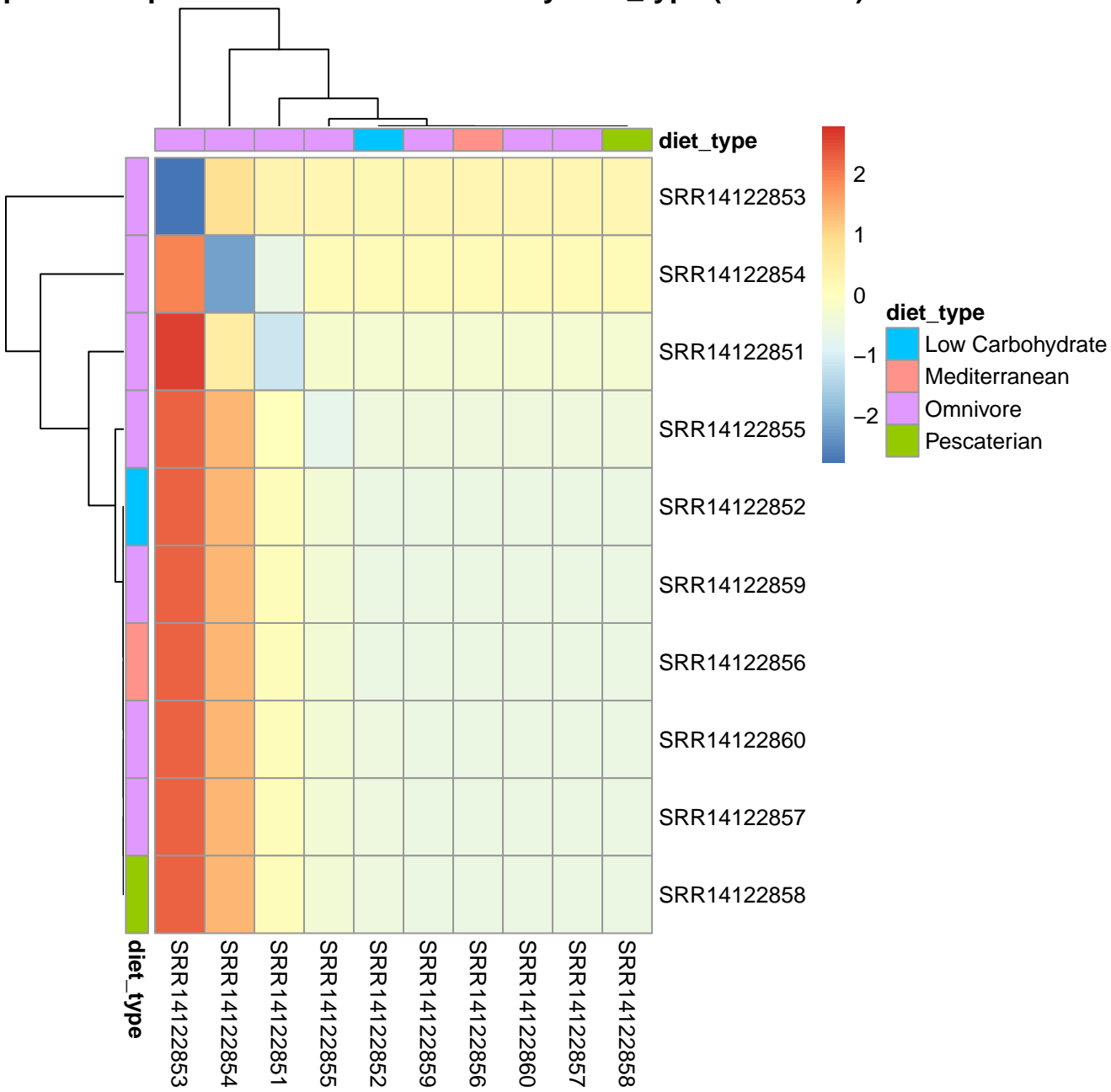
of top 50 significant feature counts by logFC colored by diet_type



top 50 significant feature counts by logFC colored by diet_type (log)



sample to sample count distance colored by diet_type (euclidean)



le to sample log count distance colored by diet_type (euclidean)

