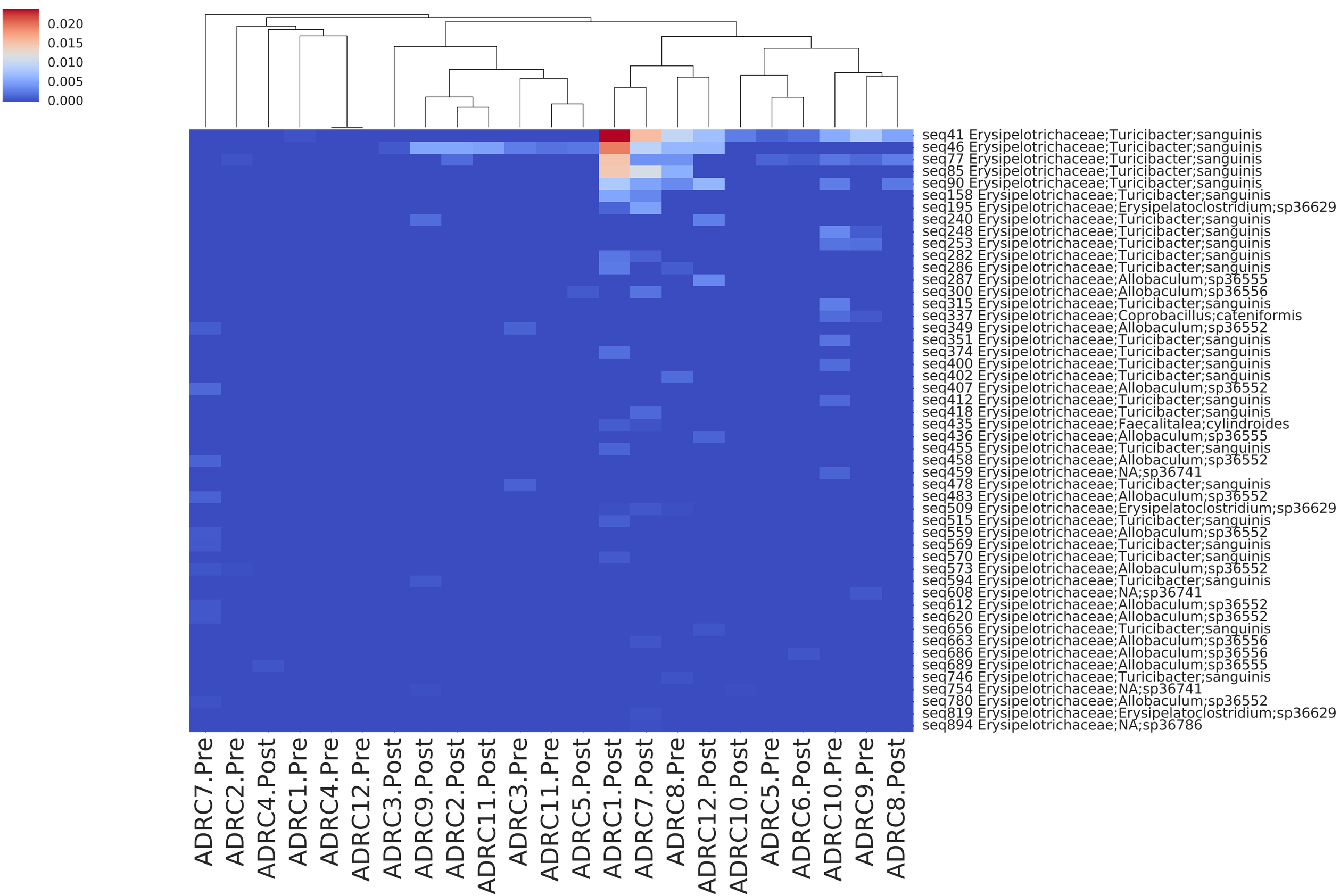
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0.015
0.010
0.005
0.000
seq41 Erysipelotrichaceae;Turicibacter;sanguinis
seq46 Erysipelotrichaceae;Turicibacter;sanguinis
seq77 Erysipelotrichaceae;Turicibacter;sanguinis
seq85 Erysipelotrichaceae;Turicibacter;sanguinis
seq90 Erysipelotrichaceae;Turicibacter;sanguinis
seq158 Erysipelotrichaceae;Turicibacter;sanguinis
seq195 Erysipelotrichaceae;Erysipelatoclostridium;sp36629
seq240 Erysipelotrichaceae;Turicibacter;sanguinis
seq248 Erysipelotrichaceae;Turicibacter;sanguinis
seq253 Erysipelotrichaceae;Turicibacter;sanguinis
seq282 Erysipelotrichaceae;Turicibacter;sanguinis
seq286 Erysipelotrichaceae;Turicibacter;sanguinis
seq287 Erysipelotrichaceae;Allobaculum;sp36555
seq300 Erysipelotrichaceae;Allobaculum;sp36556
seq315 Erysipelotrichaceae;Turicibacter;sanguinis
seq337 Erysipelotrichaceae;Coprobacillus;cateniformis
seq349 Erysipelotrichaceae;Allobaculum;sp36552
seq351 Erysipelotrichaceae;Turicibacter;sanguinis
seq374 Erysipelotrichaceae;Turicibacter;sanguinis
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seq412 Erysipelotrichaceae;Turicibacter;sanguinis
seq418 Erysipelotrichaceae;Turicibacter;sanguinis
seq435 Erysipelotrichaceae;Faecalitalea;cylindroides
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seq509 Erysipelotrichaceae;Erysipelatoclostridium;sp36629
seq515 Erysipelotrichaceae;Turicibacter;sanguinis
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seq746 Erysipelotrichaceae;Turicibacter;sanguinis
seq754 Erysipelotrichaceae;NA;sp36741
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seq894 Erysipelotrichaceae;NA;sp36786
Taxon
ADRC7.Pre
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ADRC4.Post
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