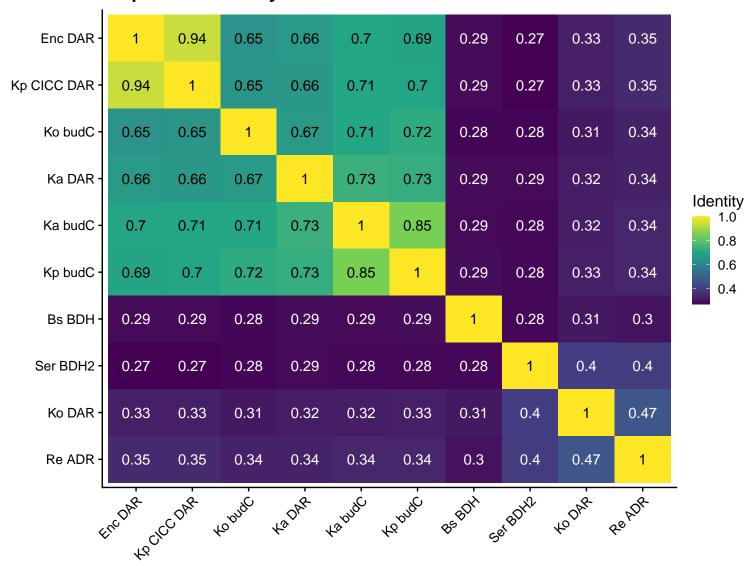
Sequence Similarity Matrix



1.0

0.8 0.6

0.4