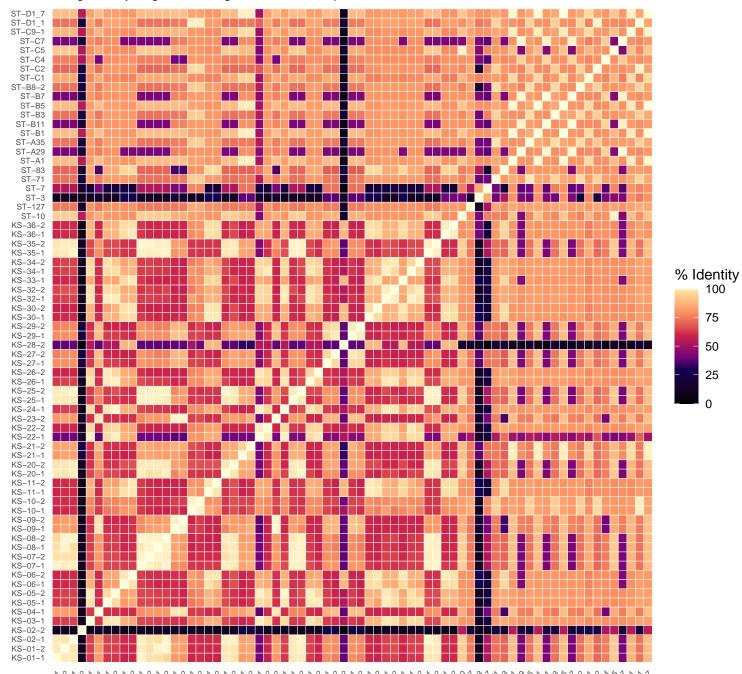
Pairwise Aligned (local) Sequence Identity Heatmap

Weighted by alignment length / shortest sequence



75

50

25