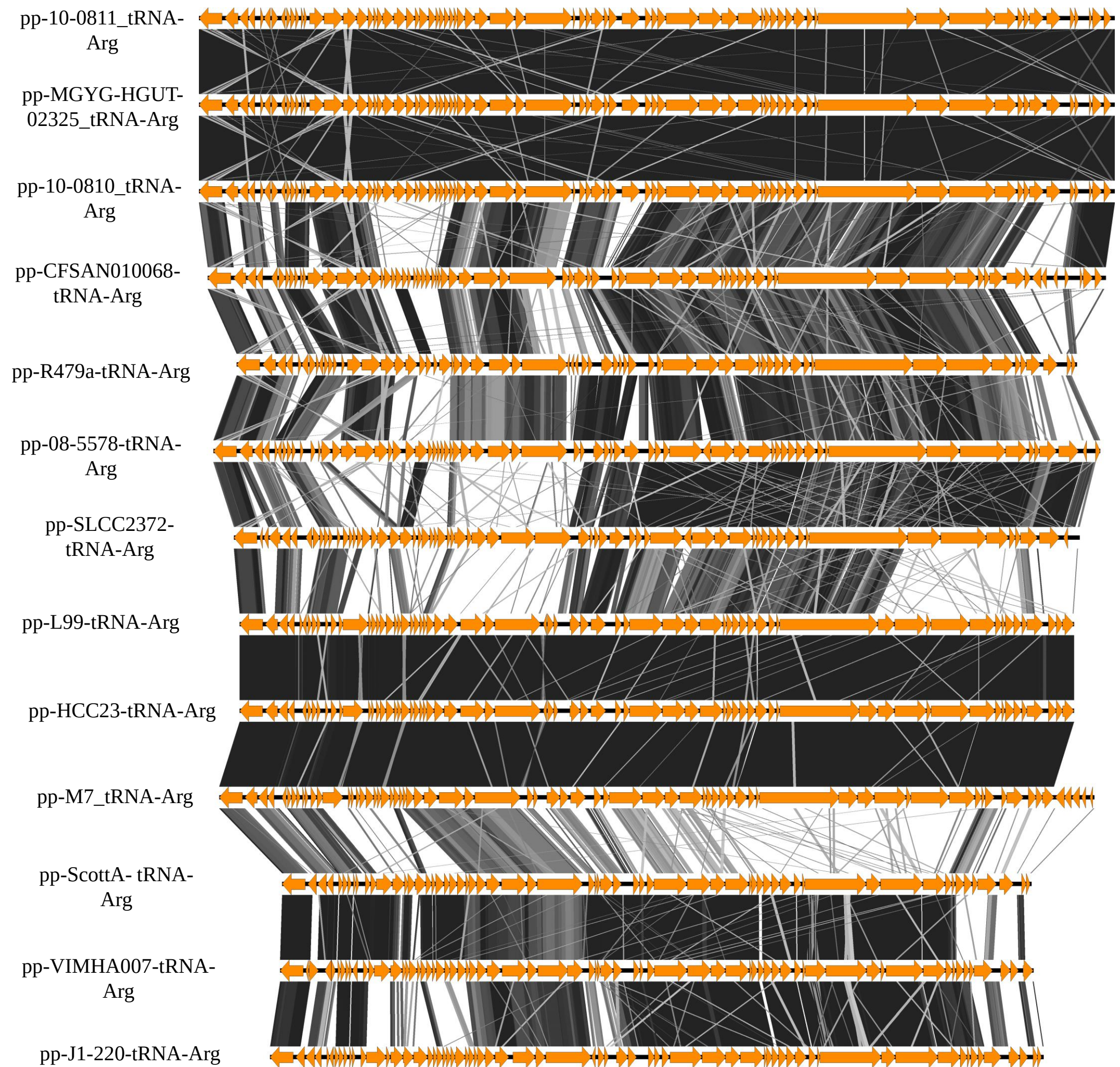
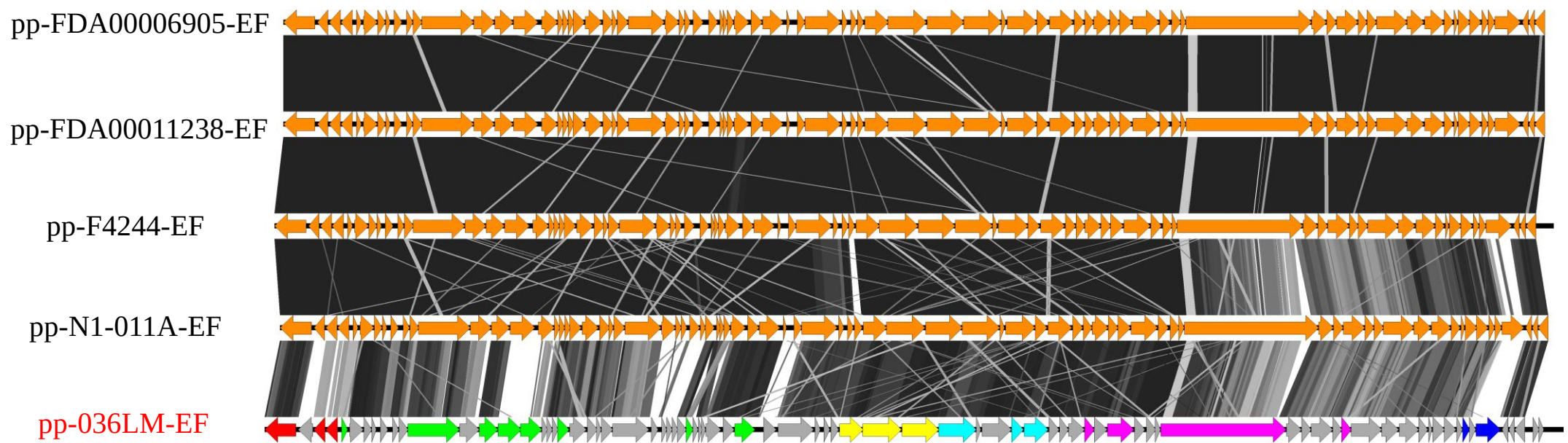


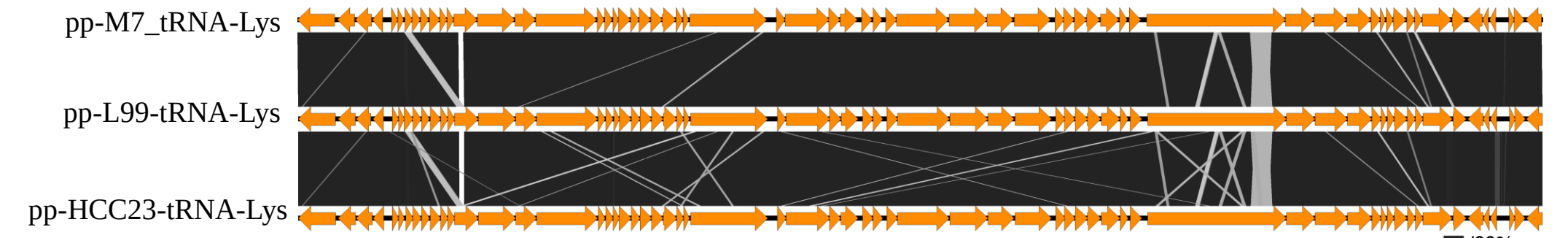
Cluster I



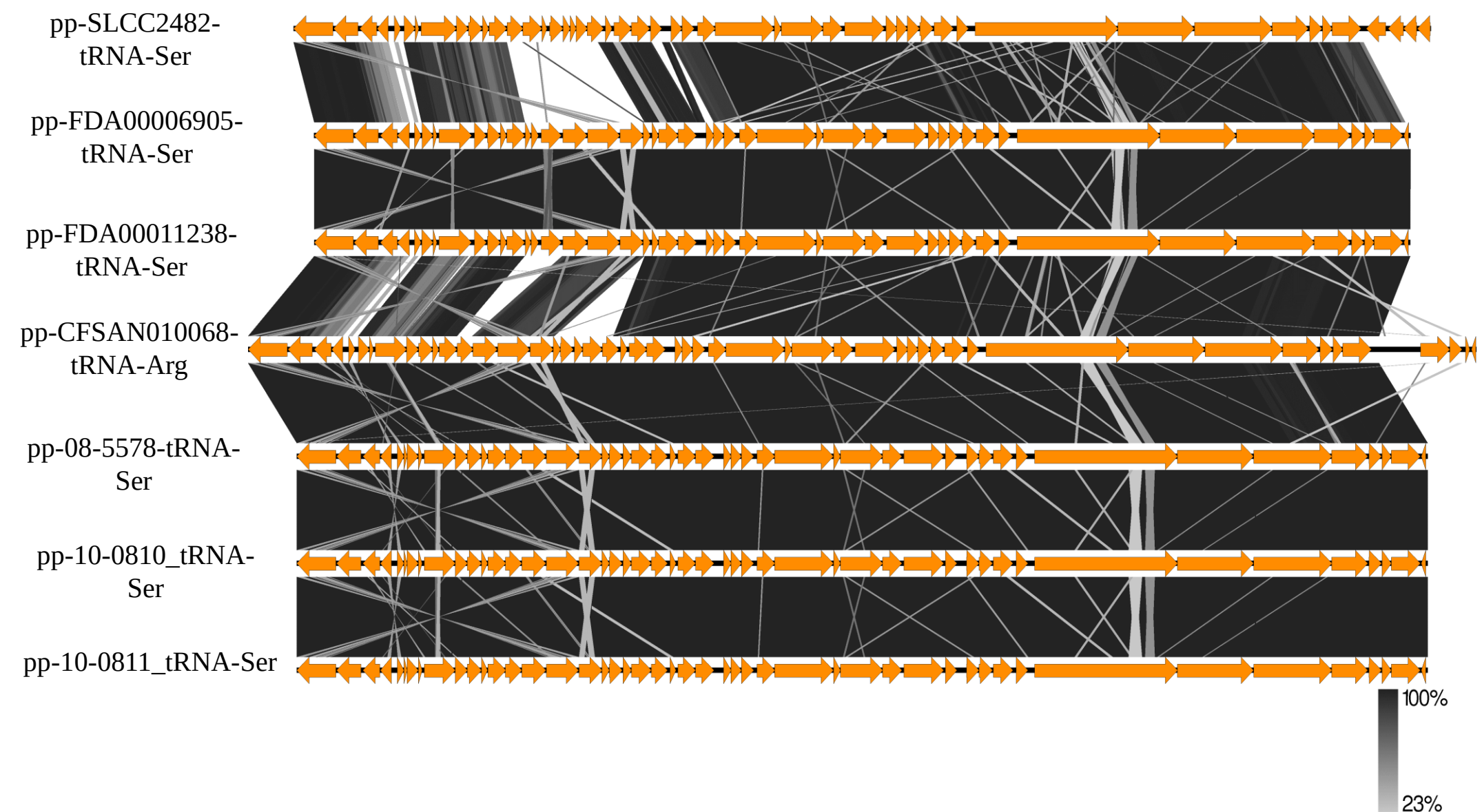
Cluster II



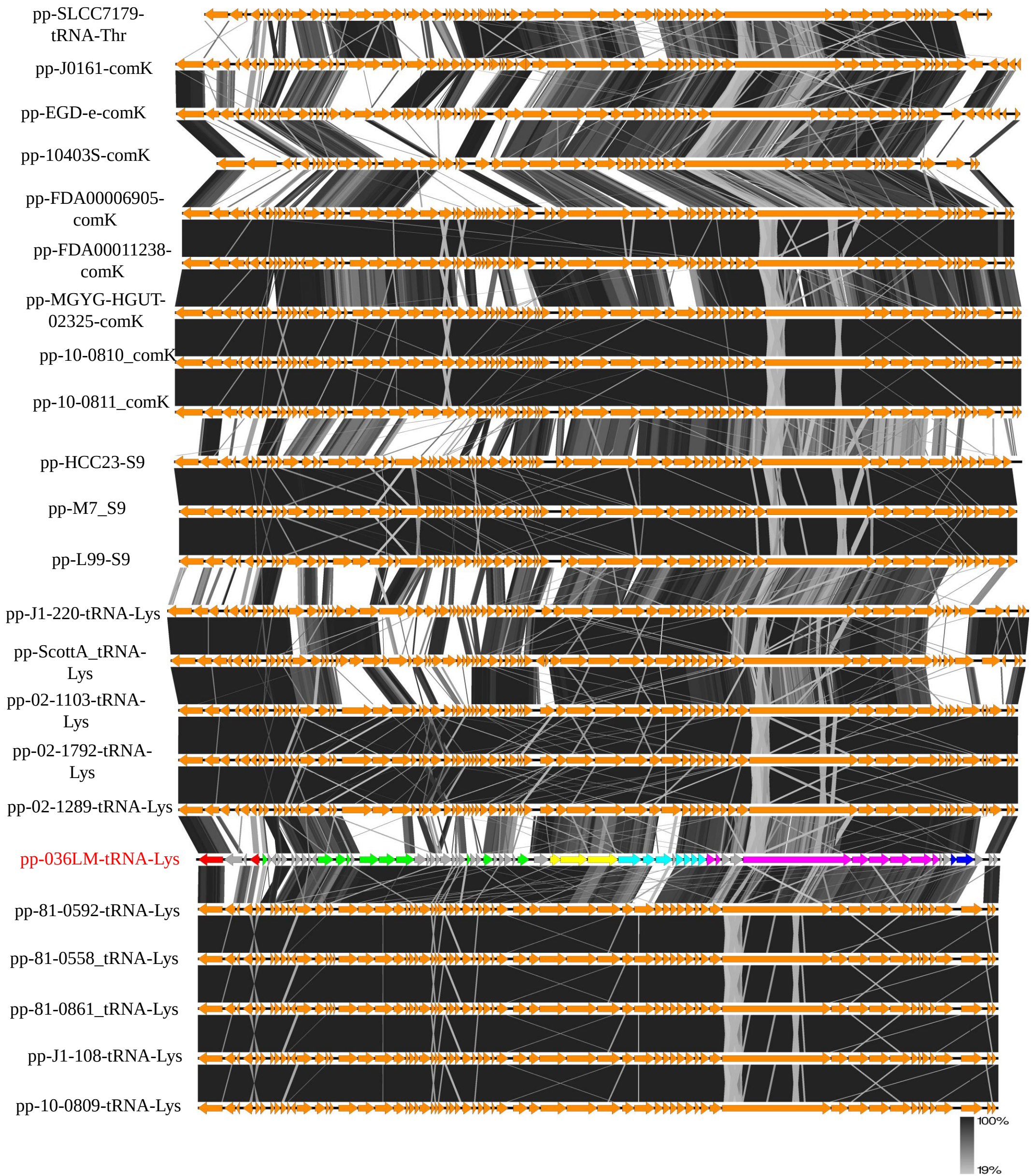
Cluster III



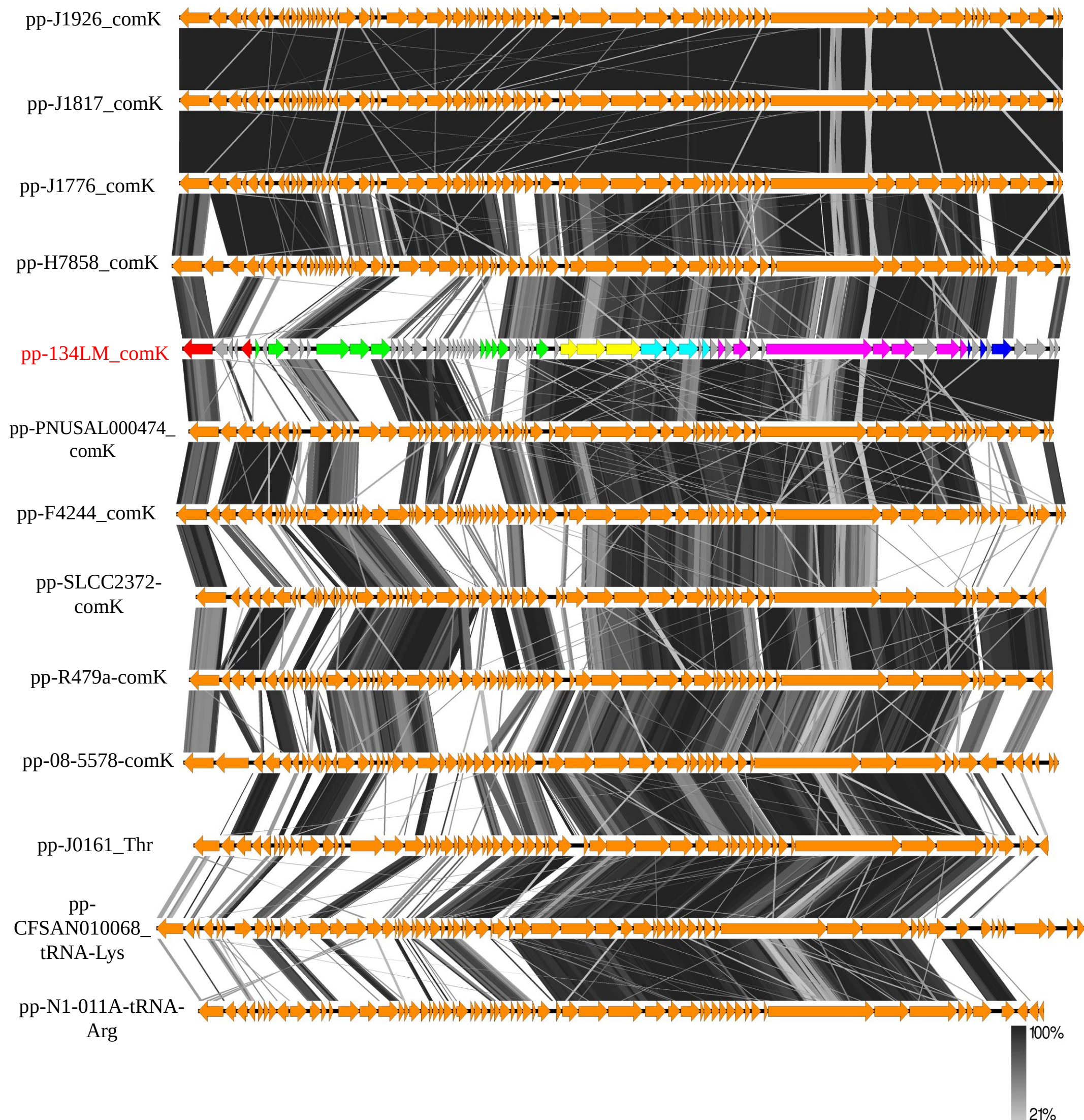
Cluster IV



Cluster V



Cluster VI



Supplement figure S1. Linear genome visualization of the prophage sequences in each of the six identified clusters conducted using Easyfig 2.1. The shade of grey between genomes indicates the level of nucleotide similarity (the darker, the higher similar).