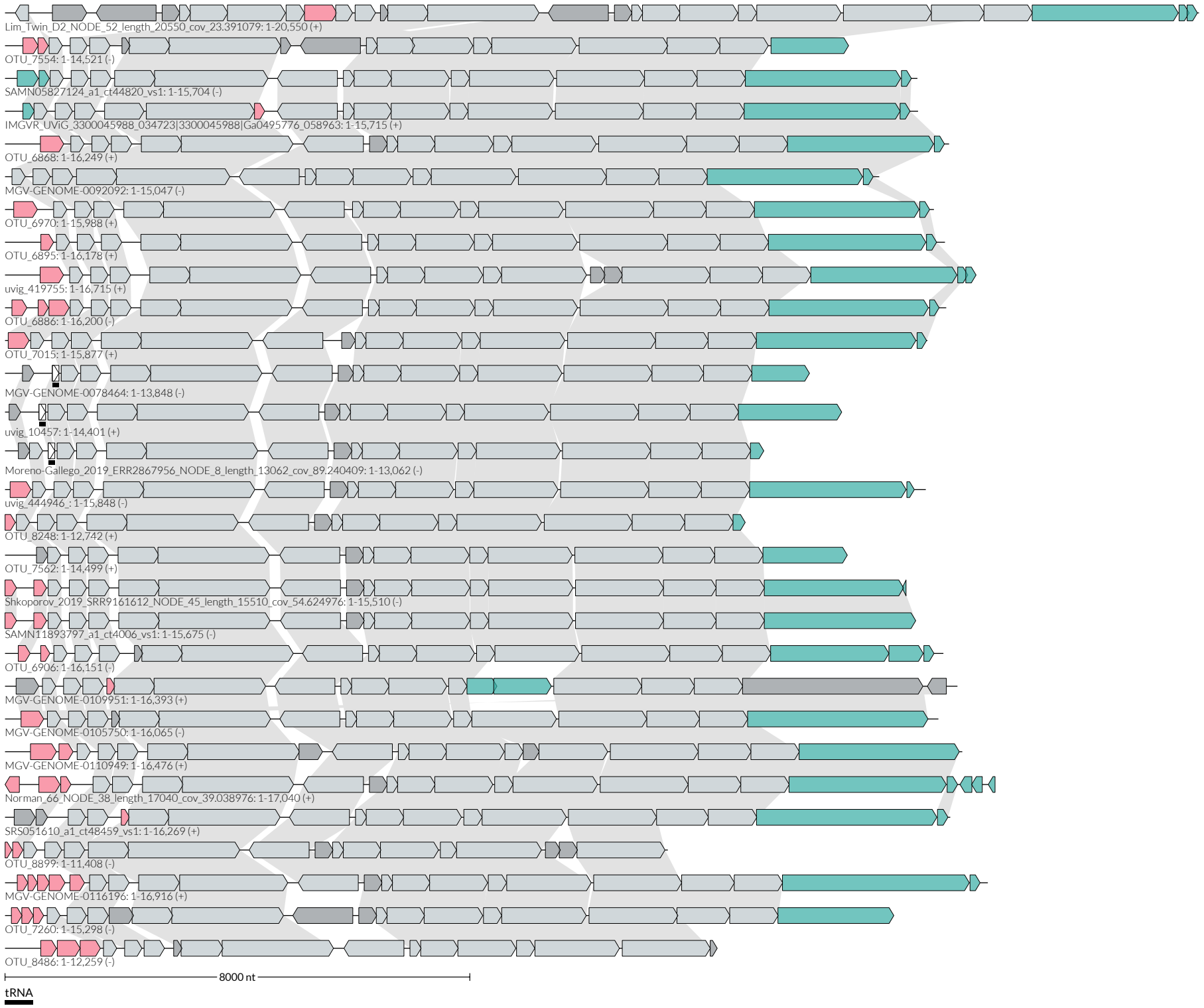
Lim_Twin_D2_NODE_52_length_20550_cov_23.391079: 1-20,550 (+)
OTU_7554: 1-14,521 (-)
SAMN05827124_a1_ct44820_vs1: 1-15,704 (-)
IMGVR_UViG_3300045988_034723|3300045988|Ga0495776_058963: 1-15,715 (+)
OTU_6868: 1-16,249 (+)
MGV-GENOME-0092092: 1-15,047 (-)
OTU_6970: 1-15,988 (+)
OTU_6895: 1-16,178 (+)
uvig_419755: 1-16,715 (+)
OTU_6886: 1-16,200 (-)
OTU_7015: 1-15,877 (+)
MGV-GENOME-0078464: 1-13,848 (-)
uvig_10457: 1-14,401 (+)
Moreno-Gallego_2019_ERR2867956_NODE_8_length_13062_cov_89.240409: 1-13,062 (-)
uvig_444946_: 1-15,848 (-)
OTU_8248: 1-12,742 (+)
OTU_7562: 1-14,499 (+)
Shkoporov_2019_SRR9161612_NODE_45_length_15510_cov_54.624976: 1-15,510 (-)
SAMN11893797_a1_ct4006_vs1: 1-15,675 (-)
OTU_6906: 1-16,151 (-)
MGV-GENOME-0109951: 1-16,393 (+)
MGV-GENOME-0105750: 1-16,065 (-)
MGV-GENOME-0110949: 1-16,476 (+)
Norman_66_NODE_38_length_17040_cov_39.038976: 1-17,040 (+)
SRS051610_a1_ct48459_vs1: 1-16,269 (+)
OTU_8899: 1-11,408 (-)
MGV-GENOME-0116196: 1-16,916 (+)
OTU_7260: 1-15,298 (-)
OTU_8486: 1-12,259 (-)
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