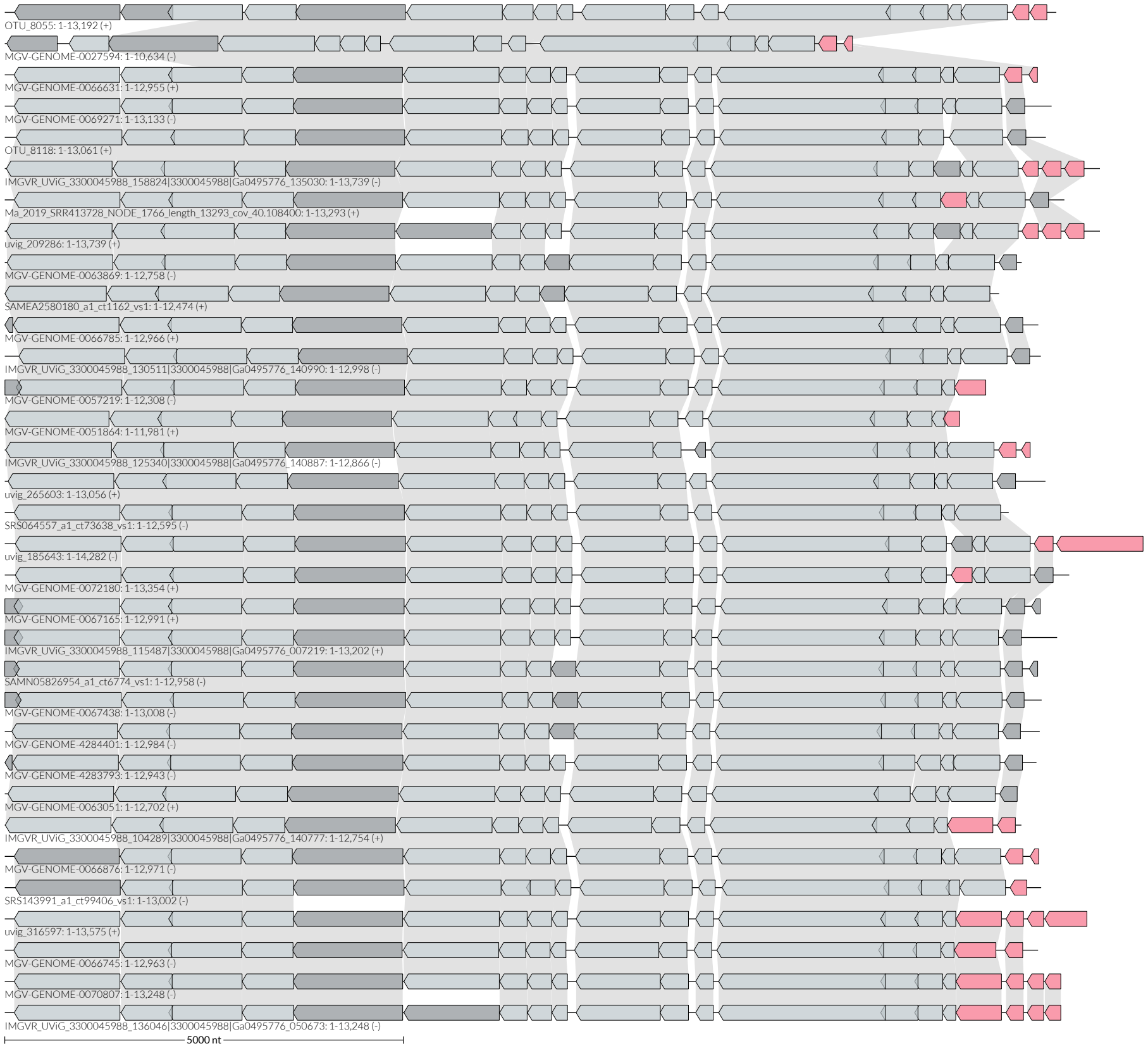
OTU_8055: 1-13,192 (+)
MGV-GENOME-0027594: 1-10,634 (-)
MGV-GENOME-0066631: 1-12,955 (+)
MGV-GENOME-0069271: 1-13,133 (-)
OTU_8118: 1-13,061 (+)
IMGVR_UViG_3300045988_158824|3300045988|Ga0495776_135030: 1-13,739 (-)
Ma_2019_SRR413728_NODE_1766_length_13293_cov_40.108400: 1-13,293 (+)
uvig_209286: 1-13,739 (+)
MGV-GENOME-0063869: 1-12,758 (-)
SAMEA2580180_a1_ct1162_vs1: 1-12,474 (+)
MGV-GENOME-0066785: 1-12,966 (+)
IMGVR_UViG_3300045988_130511|3300045988|Ga0495776_140990: 1-12,998 (-)
MGV-GENOME-0057219: 1-12,308 (-)
MGV-GENOME-0051864: 1-11,981 (+)
IMGVR_UViG_3300045988_125340|3300045988|Ga0495776_140887: 1-12,866 (-)
uvig_265603: 1-13,056 (+)
SRS064557_a1_ct73638_vs1: 1-12,595 (-)
uvig_185643: 1-14,282 (-)
MGV-GENOME-0072180: 1-13,354 (+)
MGV-GENOME-0067165: 1-12,991 (+)
IMGVR_UViG_3300045988_115487|3300045988|Ga0495776_007219: 1-13,202 (+)
SAMN05826954_a1_ct6774_vs1: 1-12,958 (-)
MGV-GENOME-0067438: 1-13,008 (-)
MGV-GENOME-4284401: 1-12,984 (-)
MGV-GENOME-4283793: 1-12,943 (-)
MGV-GENOME-0063051: 1-12,702 (+)
IMGVR_UViG_3300045988_104289|3300045988|Ga0495776_140777: 1-12,754 (+)
MGV-GENOME-0066876: 1-12,971 (-)
SRS143991_a1_ct99406_vs1: 1-13,002 (-)
uvig_316597: 1-13,575 (+)
MGV-GENOME-0066745: 1-12,963 (-)
MGV-GENOME-0070807: 1-13,248 (-)
IMGVR_UViG_3300045988_136046|3300045988|Ga0495776_050673: 1-13,248 (-)
5000 nt