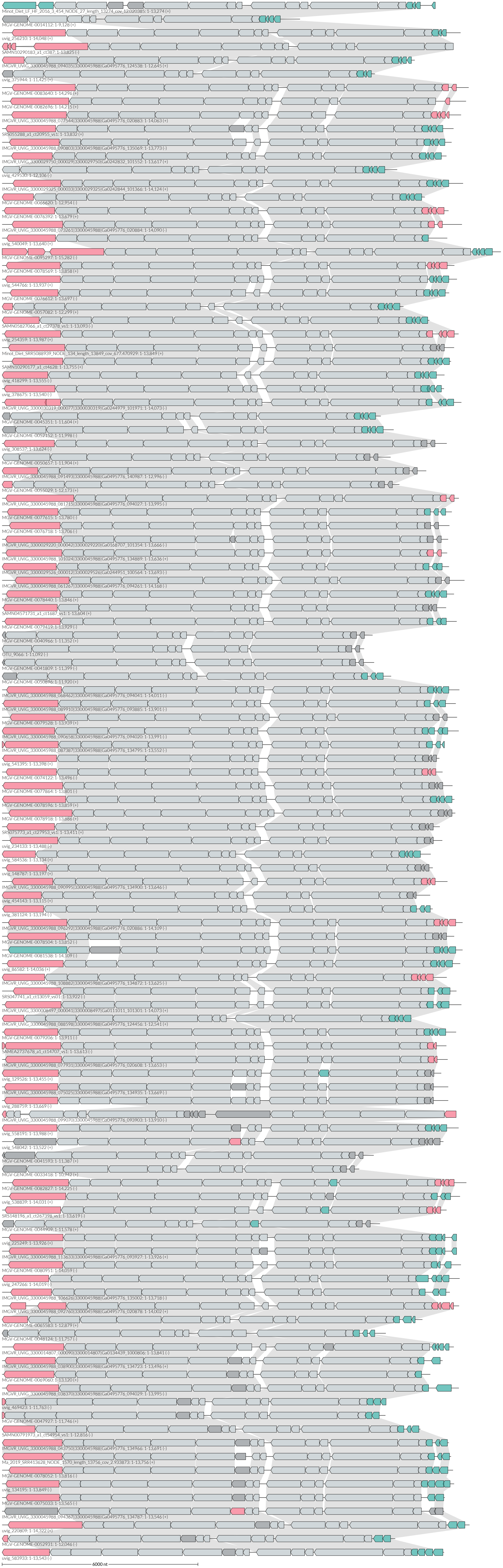
Minot_Diet_LF_HF_2016_3_454_NODE_27_length_13274_cov_12.020385: 1-13,274 (+)
MGV-GENOME-0014112: 1-9,128 (+)
uvig_256210: 1-14,048 (+)
SAMN10290183_a1_ct387: 1-13,825 (-)
IMGVR_UViG_3300045988_094035|3300045988|Ga0495776_124538: 1-12,645 (+)
uvig_375944: 1-11,425 (+)
MGV-GENOME-0083640: 1-14,296 (+)
MGV-GENOME-0082696: 1-14,215 (+)
IMGVR_UViG_3300045988_077544|3300045988|Ga0495776_020883: 1-14,063 (+)
SRS055288_a1_ct20955_vs1: 1-13,832 (+)
IMGVR_UViG_3300045988_090803|3300045988|Ga0495776_135069: 1-13,773 (-)
IMGVR_UViG_3300029750_000029|3300029750|Ga0242832_101552: 1-13,617 (+)
uvig_429530: 1-12,106 (-)
IMGVR_UViG_3300029325_000033|3300029325|Ga0242844_101366: 1-14,124 (+)
MGV-GENOME-0066620: 1-12,954 (-)
MGV-GENOME-0076392: 1-13,679 (+)
IMGVR_UViG_3300045988_073261|3300045988|Ga0495776_020884: 1-14,090 (-)
uvig_540049: 1-13,640 (+)
MGV-GENOME-0095297: 1-15,282 (-)
MGV-GENOME-0078569: 1-13,858 (+)
uvig_544766: 1-13,937 (+)
MGV-GENOME-0076612: 1-13,697 (-)
MGV-GENOME-0057082: 1-12,299 (+)
SAMN05827066_a1_ct27378_vs1: 1-13,093 (-)
uvig_254359: 1-13,987 (+)
Minot_Diet_SRR5088939_NODE_134_length_13849_cov_677.470929: 1-13,849 (+)
SAMN10290177_a1_ct4628: 1-13,755 (+)
uvig_418299: 1-13,555 (-)
uvig_378675: 1-13,540 (-)
IMGVR_UViG_3300030319_000077|3300030319|Ga0244979_101971: 1-14,073 (-)
MGV-GENOME-0045351: 1-11,604 (+)
MGV-GENOME-0052152: 1-11,998 (-)
uvig_308537: 1-13,624 (-)
MGV-GENOME-0050657: 1-11,904 (+)
IMGVR_UViG_3300045988_091493|3300045988|Ga0495776_140987: 1-12,996 (-)
MGV-GENOME-0055029: 1-12,173 (+)
IMGVR_UViG_3300045988_081715|3300045988|Ga0495776_094027: 1-13,995 (-)
MGV-GENOME-0077615: 1-13,780 (-)
MGV-GENOME-0076718: 1-13,706 (-)
IMGVR_UViG_3300029220_000042|3300029220|Ga0168707_101354: 1-13,666 (-)
IMGVR_UViG_3300045988_101024|3300045988|Ga0495776_134889: 1-13,636 (+)
IMGVR_UViG_3300029526_000012|3300029526|Ga0244951_100564: 1-13,693 (-)
IMGVR_UViG_3300045988_061267|3300045988|Ga0495776_094261: 1-14,168 (-)
MGV-GENOME-0078440: 1-13,846 (+)
SAMN04571731_a1_ct1687_vs1: 1-13,604 (+)
MGV-GENOME-0079419: 1-13,929 (-)
MGV-GENOME-0040966: 1-11,352 (+)
OTU_9066: 1-11,092 (-)
MGV-GENOME-0041809: 1-11,399 (-)
MGV-GENOME-0050896: 1-11,920 (+)
IMGVR_UViG_3300045988_068462|3300045988|Ga0495776_094041: 1-14,011 (-)
IMGVR_UViG_3300045988_089910|3300045988|Ga0495776_093885: 1-13,901 (-)
MGV-GENOME-0079528: 1-13,939 (+)
IMGVR_UViG_3300045988_090658|3300045988|Ga0495776_094020: 1-13,991 (-)
IMGVR_UViG_3300045988_087387|3300045988|Ga0495776_134795: 1-13,552 (-)
uvig_541395: 1-13,398 (+)
MGV-GENOME-0074122: 1-13,496 (-)
MGV-GENOME-0077864: 1-13,801 (-)
MGV-GENOME-0078596: 1-13,859 (+)
MGV-GENOME-0078918: 1-13,886 (+)
SRS075773_a1_ct27953_vs1: 1-13,411 (+)
uvig_234133: 1-13,488 (-)
uvig_584536: 1-13,134 (+)
uvig_148787: 1-13,197 (+)
IMGVR_UViG_3300045988_090995|3300045988|Ga0495776_134900: 1-13,646 (-)
uvig_454143: 1-13,115 (+)
uvig_381124: 1-13,194 (-)
IMGVR_UViG_3300045988_096292|3300045988|Ga0495776_020886: 1-14,109 (-)
MGV-GENOME-0078504: 1-13,852 (-)
MGV-GENOME-0081538: 1-14,109 (-)
uvig_86582: 1-14,036 (+)
IMGVR_UViG_3300045988_108882|3300045988|Ga0495776_134872: 1-13,625 (-)
SRS047741_a1_ct13059_vs01: 1-13,922 (-)
IMGVR_UViG_3300008497_000041|3300008497|Ga0111011_101301: 1-14,073 (+)
IMGVR_UViG_3300045988_088598|3300045988|Ga0495776_124456: 1-12,541 (+)
MGV-GENOME-0079206: 1-13,911 (-)
SAMEA2737678_a1_ct14707_vs1: 1-13,613 (-)
IMGVR_UViG_3300045988_077931|3300045988|Ga0495776_020608: 1-13,653 (-)
uvig_129526: 1-13,455 (+)
IMGVR_UViG_3300045988_075025|3300045988|Ga0495776_134935: 1-13,669 (-)
uvig_288759: 1-13,669 (-)
IMGVR_UViG_3300045988_099070|3300045988|Ga0495776_093903: 1-13,910 (-)
uvig_558191: 1-13,988 (+)
uvig_548042: 1-13,522 (+)
MGV-GENOME-0041593: 1-11,387 (+)
MGV-GENOME-0033418: 1-10,942 (+)
MGV-GENOME-0082827: 1-14,225 (-)
uvig_538839: 1-14,031 (+)
SRS148196_a1_ct267398_vs1: 1-13,619 (-)
MGV-GENOME-0044909: 1-11,578 (+)
uvig_225249: 1-13,926 (+)
IMGVR_UViG_3300045988_113633|3300045988|Ga0495776_093927: 1-13,926 (+)
MGV-GENOME-0080951: 1-14,059 (-)
uvig_247266: 1-14,019 (-)
IMGVR_UViG_3300045988_106626|3300045988|Ga0495776_135002: 1-13,718 (-)
IMGVR_UViG_3300045988_092760|3300045988|Ga0495776_020878: 1-14,002 (+)
MGV-GENOME-0065583: 1-12,879 (+)
MGV-GENOME-0048124: 1-11,757 (-)
IMGVR_UViG_3300014807_000090|3300014807|Ga0134439_1000806: 1-13,841 (-)
IMGVR_UViG_3300045988_038900|3300045988|Ga0495776_134723: 1-13,496 (+)
MGV-GENOME-0069060: 1-13,120 (+)
IMGVR_UViG_3300045988_038370|3300045988|Ga0495776_094029: 1-13,995 (-)
uvig_469423: 1-11,763 (-)
MGV-GENOME-0047927: 1-11,746 (+)
SAMN00791973_a1_ct54954_vs1: 1-12,816 (-)
IMGVR_UViG_3300045988_043750|3300045988|Ga0495776_134966: 1-13,691 (-)
Ma_2019_SRR413628_NODE_1570_length_13756_cov_2.933873: 1-13,756 (+)
MGV-GENOME-0078052: 1-13,816 (-)
uvig_134195: 1-13,849 (-)
MGV-GENOME-0075033: 1-13,565 (-)
IMGVR_UViG_3300045988_094367|3300045988|Ga0495776_134787: 1-13,546 (+)
uvig_220809: 1-14,322 (+)
MGV-GENOME-0052931: 1-12,046 (-)
uvig_583933: 1-13,543 (-)
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