IMGVR_UViG_3300045988_151304|3300045988|Ga0495776_135162|126-13844: 1-13,719 (+)
IMGVR_UViG_3300045988_158842|3300045988|Ga0495776_156939|1942-15485: 1-13,544 (+)
IMGVR_UViG_3300045988_154506|3300045988|Ga0495776_099120|1390-15260: 1-13,871 (+)
IMGVR_UViG_3300045988_147884|3300045988|Ga0495776_040748|5231-19290: 1-14,060 (+)
IMGVR_UViG_3300045988_154549|3300045988|Ga0495776_103773|4746-17700: 1-12,955 (+)
IMGVR_UViG_3300007055_000003|3300007055|Ga0102702_100535|2568-15304: 1-12,737 (+)
IMGVR_UViG_3300045988_157687|3300045988|Ga0495776_040805|6644-19317: 1-12,674 (+)
MGV-GENOME-0118041: 1-15,489 (-)
MGV-GENOME-0108107: 1-15,485 (+)
MGV-GENOME-0094654: 1-15,236 (+)
MGV-GENOME-0092936: 1-15,108 (-)
MGV-GENOME-0095518: 1-15,299 (+)
IMGVR_UViG_7000000385_000045|7000000385|SRS017433_Baylor_scaffold_2198: 1-13,783 (+)
MGV-GENOME-0076483: 1-13,686 (+)
Han_2018_ERR1398205_NODE_2095_length_12836_cov_13.284798: 1-12,836 (+)
MGV-GENOME-0140071: 1-19,290 (+)
MGV-GENOME-0137424: 1-18,993 (-)
MGV-GENOME-4343381: 1-17,700 (+)
MGV-GENOME-0150631: 1-20,576 (+)
IMGVR_UViG_3300029678_000037|3300029678|Ga0245195_101811|1600-14916: 1-13,317 (-)
MGV-GENOME-0051814: 1-11,979 (+)
MGV-GENOME-0182355: 1-19,509 (-)
IMGVR_UViG_3300045988_137351|3300045988|Ga0495776_155701|1-18120: 1-18,120 (-)
MGV-GENOME-0155921: 1-19,860 (+)
IMGVR_UViG_3300045988_118083|3300045988|Ga0495776_067331|1-17920: 1-17,920 (-)
MGV-GENOME-0106299: 1-16,110 (-)
IMGVR_UViG_3300045988_150017|3300045988|Ga0495776_108060|4758-20576: 1-15,819 (+)
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