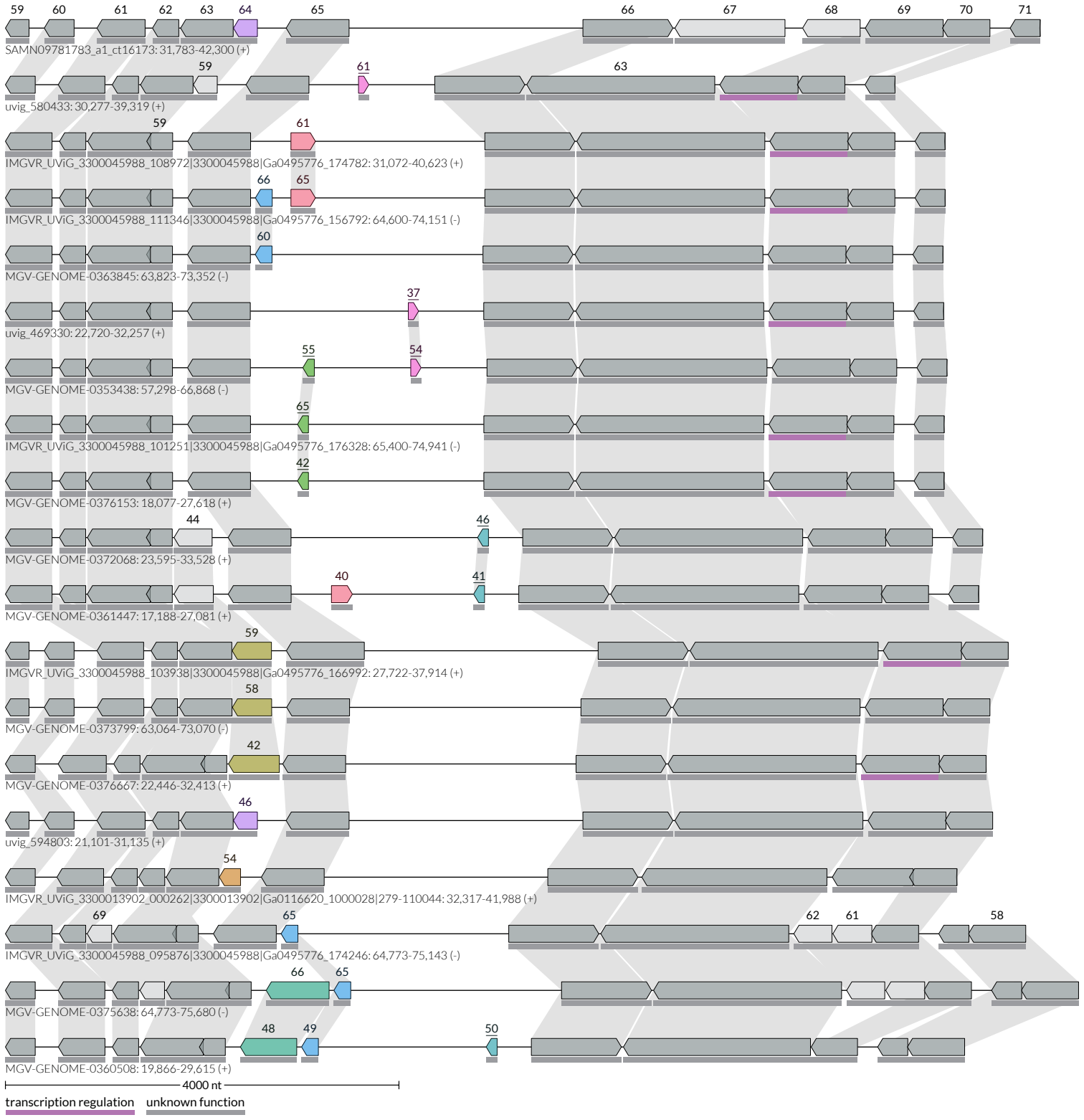
SAMN09781783_a1_ct16173: 31,783-42,300 (+)
uvig_580433: 30,277-39,319 (+)
IMGVR_UViG_3300045988_108972|3300045988|Ga0495776_174782: 31,072-40,623 (+)
IMGVR_UViG_3300045988_111346|3300045988|Ga0495776_156792: 64,600-74,151 (-)
MGV-GENOME-0363845: 63,823-73,352 (-)
uvig_469330: 22,720-32,257 (+)
MGV-GENOME-0353438: 57,298-66,868 (-)
IMGVR_UViG_3300045988_101251|3300045988|Ga0495776_176328: 65,400-74,941 (-)
MGV-GENOME-0376153: 18,077-27,618 (+)
MGV-GENOME-0372068: 23,595-33,528 (+)
MGV-GENOME-0361447: 17,188-27,081 (+)
IMGVR_UViG_3300045988_103938|3300045988|Ga0495776_166992: 27,722-37,914 (+)
MGV-GENOME-0373799: 63,064-73,070 (-)
MGV-GENOME-0376667: 22,446-32,413 (+)
uvig_594803: 21,101-31,135 (+)
IMGVR_UViG_3300013902_000262|3300013902|Ga0116620_1000028|279-110044: 32,317-41,988 (+)
IMGVR_UViG_3300045988_095876|3300045988|Ga0495776_174246: 64,773-75,143 (-)
MGV-GENOME-0375638: 64,773-75,680 (-)
MGV-GENOME-0360508: 19,866-29,615 (+)
4000 nt
transcription regulation
unknown function