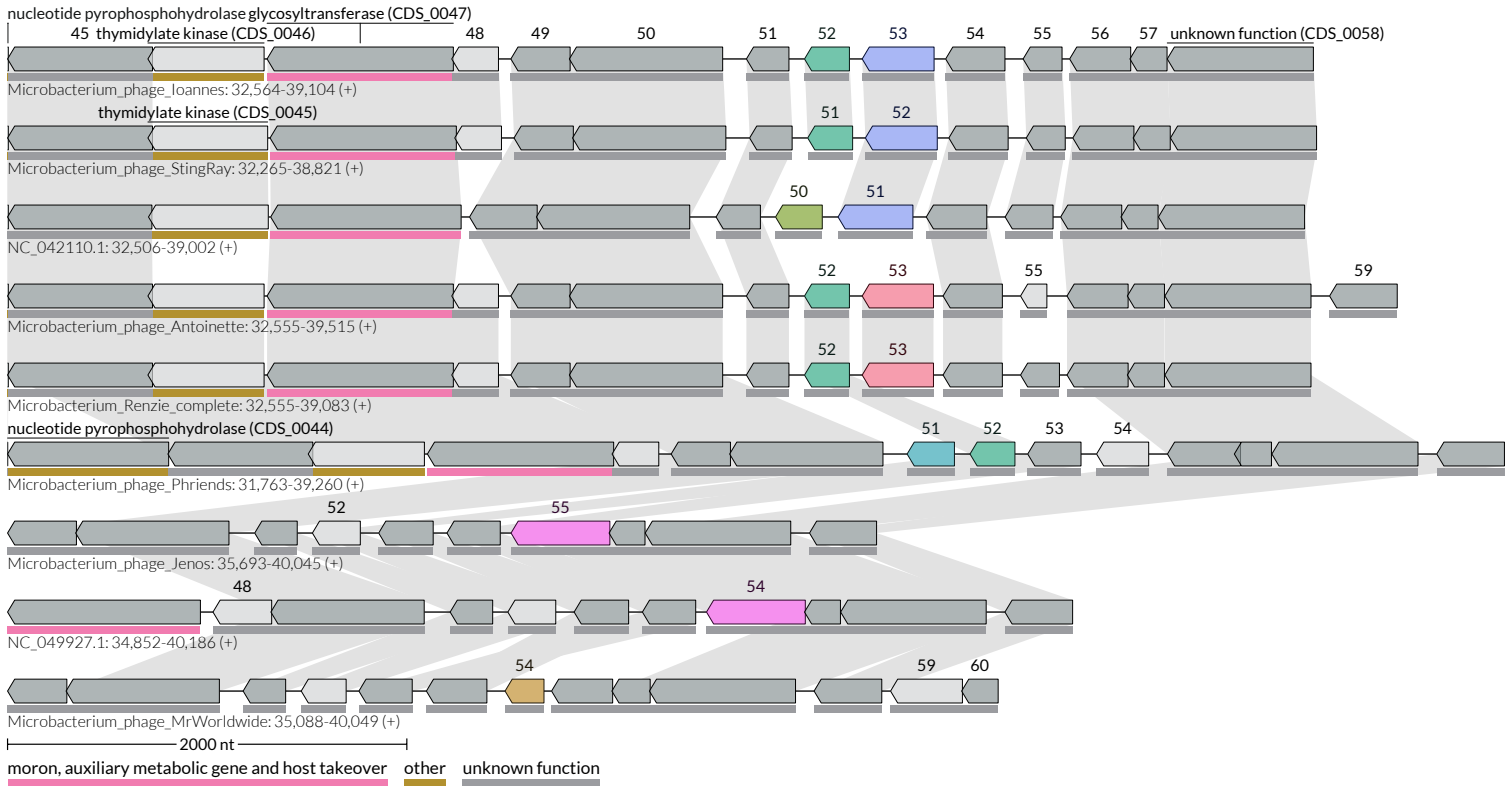
nucleotide pyrophosphohydrolase glycosyltransferase (CDS_0047)
45
thymidylate kinase (CDS_0046)
48
49
50
51
52
53
54
55
56
57
unknown function (CDS_0058)
Microbacterium_phage_Ioannes: 32,564-39,104 (+)
thymidylate kinase (CDS_0045)
51
52
Microbacterium_phage_StingRay: 32,265-38,821 (+)
50
51
NC_042110.1: 32,506-39,002 (+)
52
53
55
59
Microbacterium_phage_Antoinette: 32,555-39,515 (+)
52
53
Microbacterium_Renzie_complete: 32,555-39,083 (+)
nucleotide pyrophosphohydrolase (CDS_0044)
51
52
53
54
Microbacterium_phage_Phriends: 31,763-39,260 (+)
52
55
Microbacterium_phage_Jenos: 35,693-40,045 (+)
48
54
NC_049927.1: 34,852-40,186 (+)
54
59
60
Microbacterium_phage_MrWorldwide: 35,088-40,049 (+)
2000 nt
moron, auxiliary metabolic gene and host takeover
other
unknown function