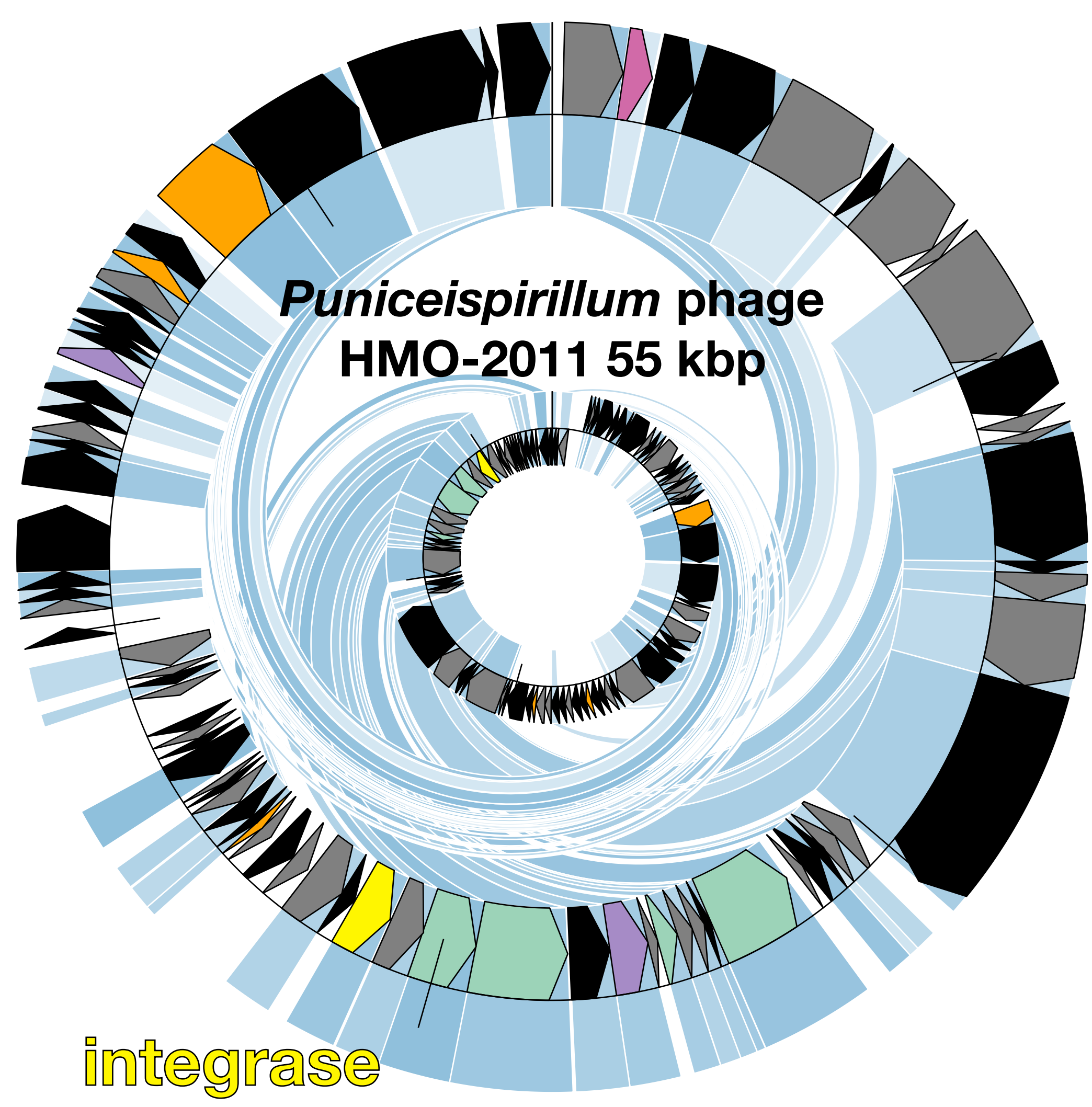
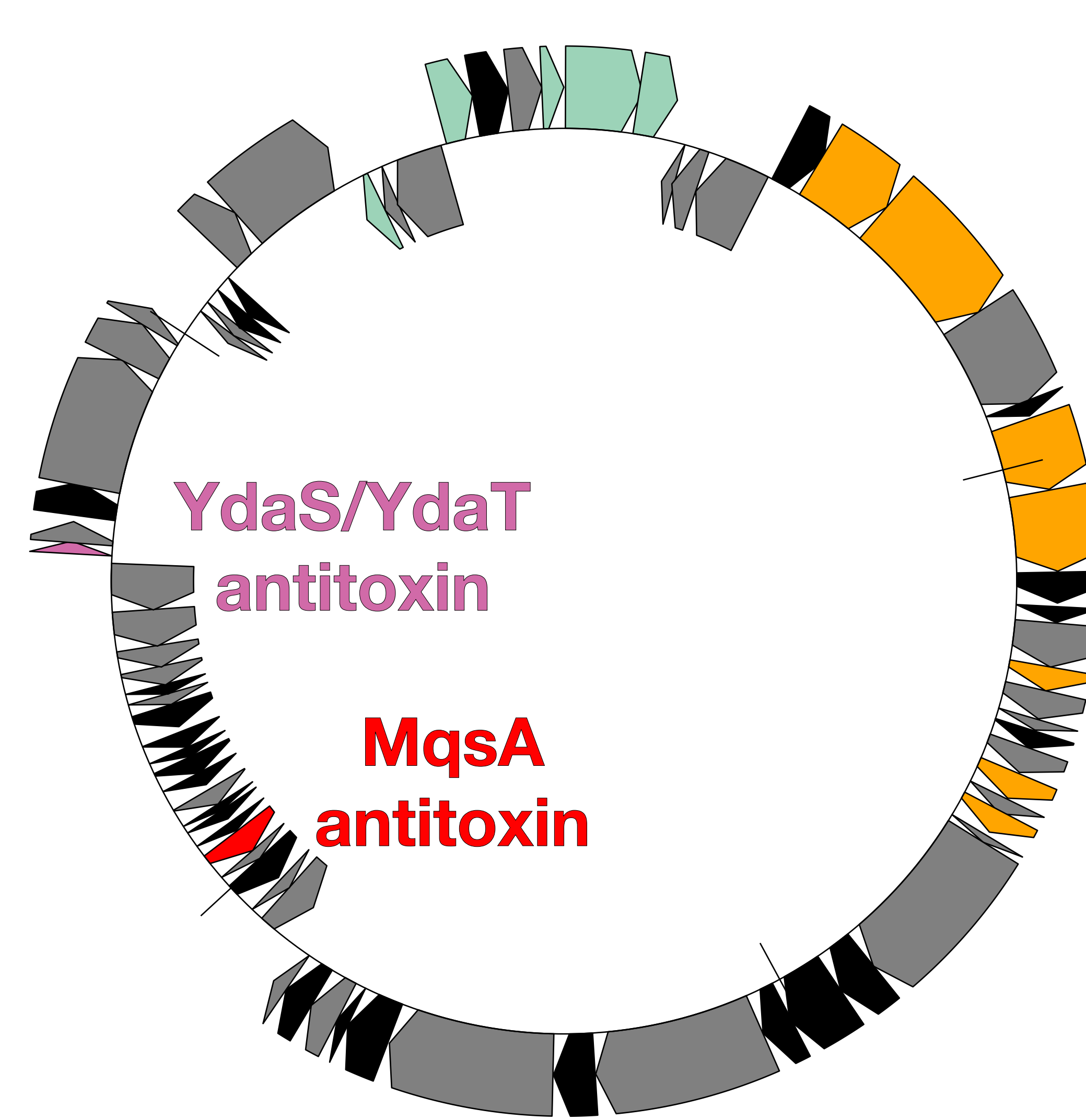


39571. putative SAR116 temperate phage  
complete genome 55kbp



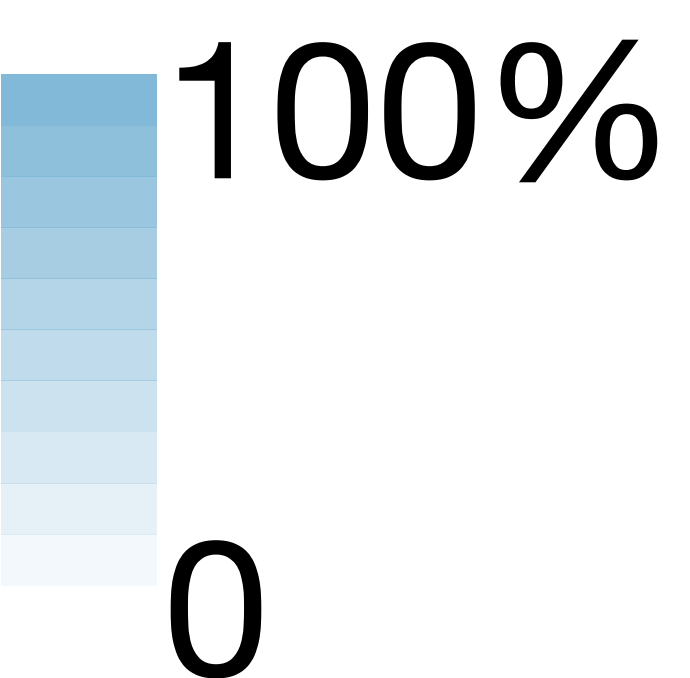
16956. putative *Marinomonas* phage  
complete genome 47kbp



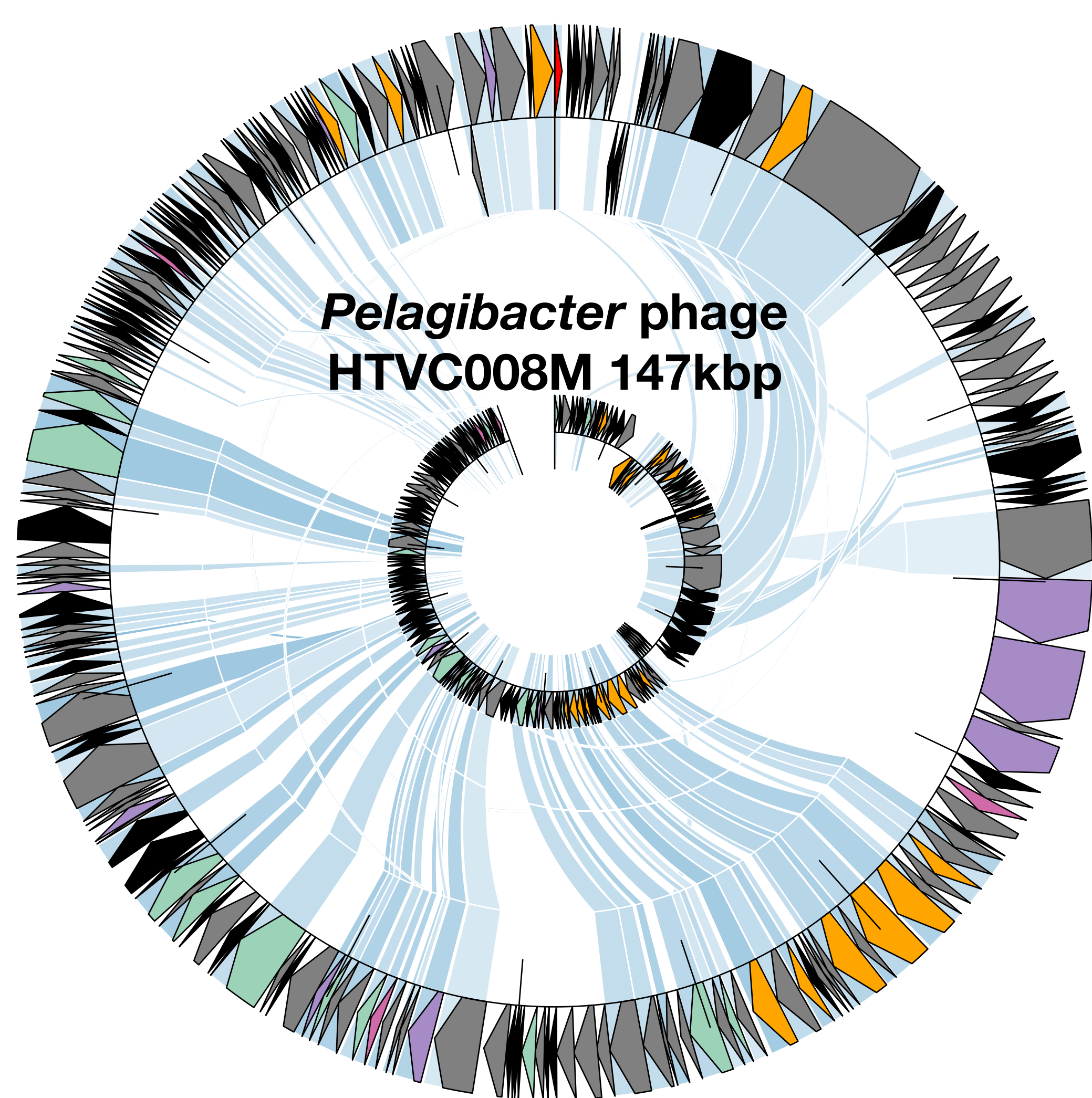
protein

structural  
regulatory  
replication  
lysogeny  
AMG  
other  
hypothetical  
novel viral genes

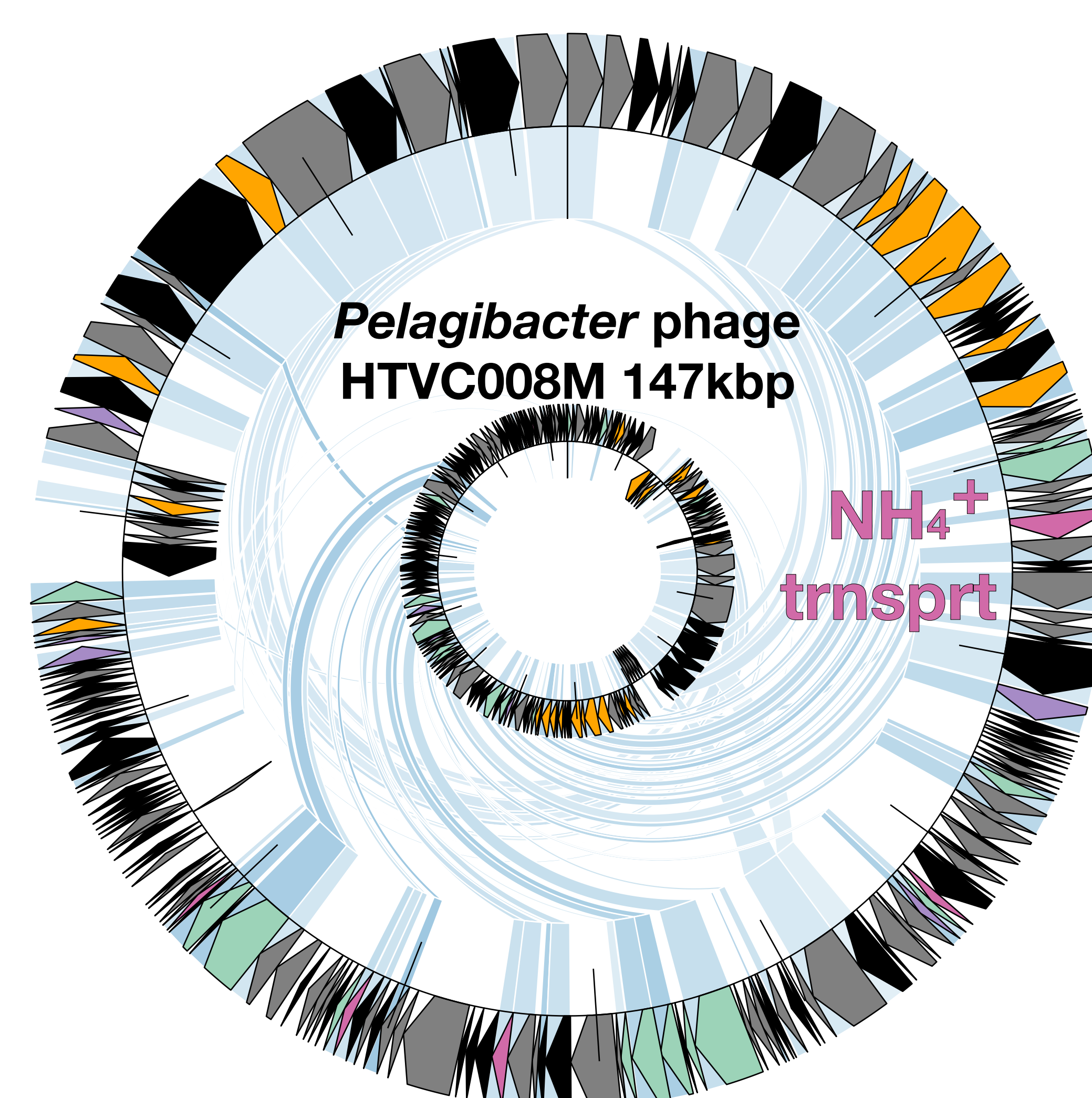
AAI



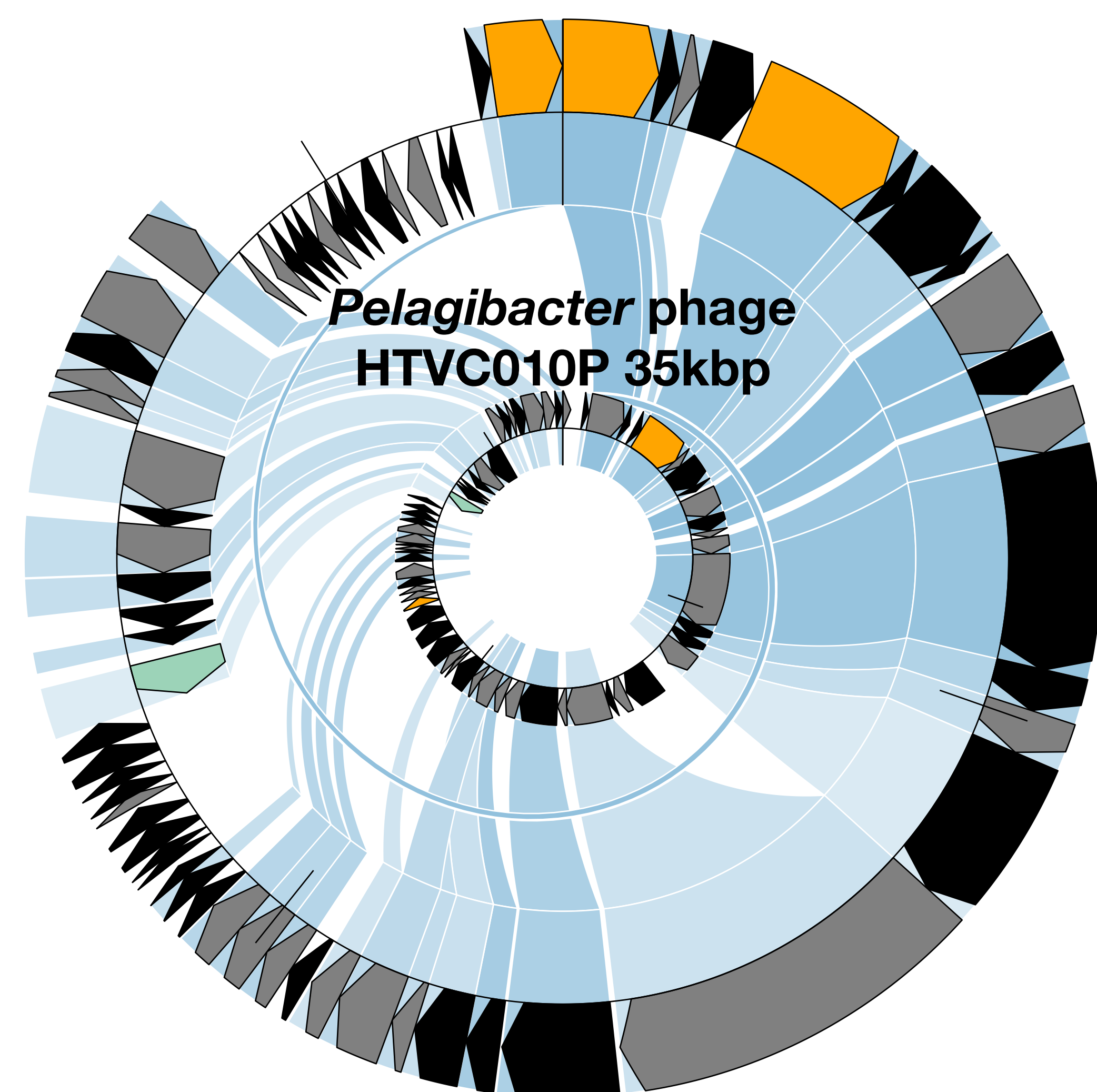
30093. putative SAR11 temperate phage  
complete genome 156kbp  
most abundant temperate phage in both fractions



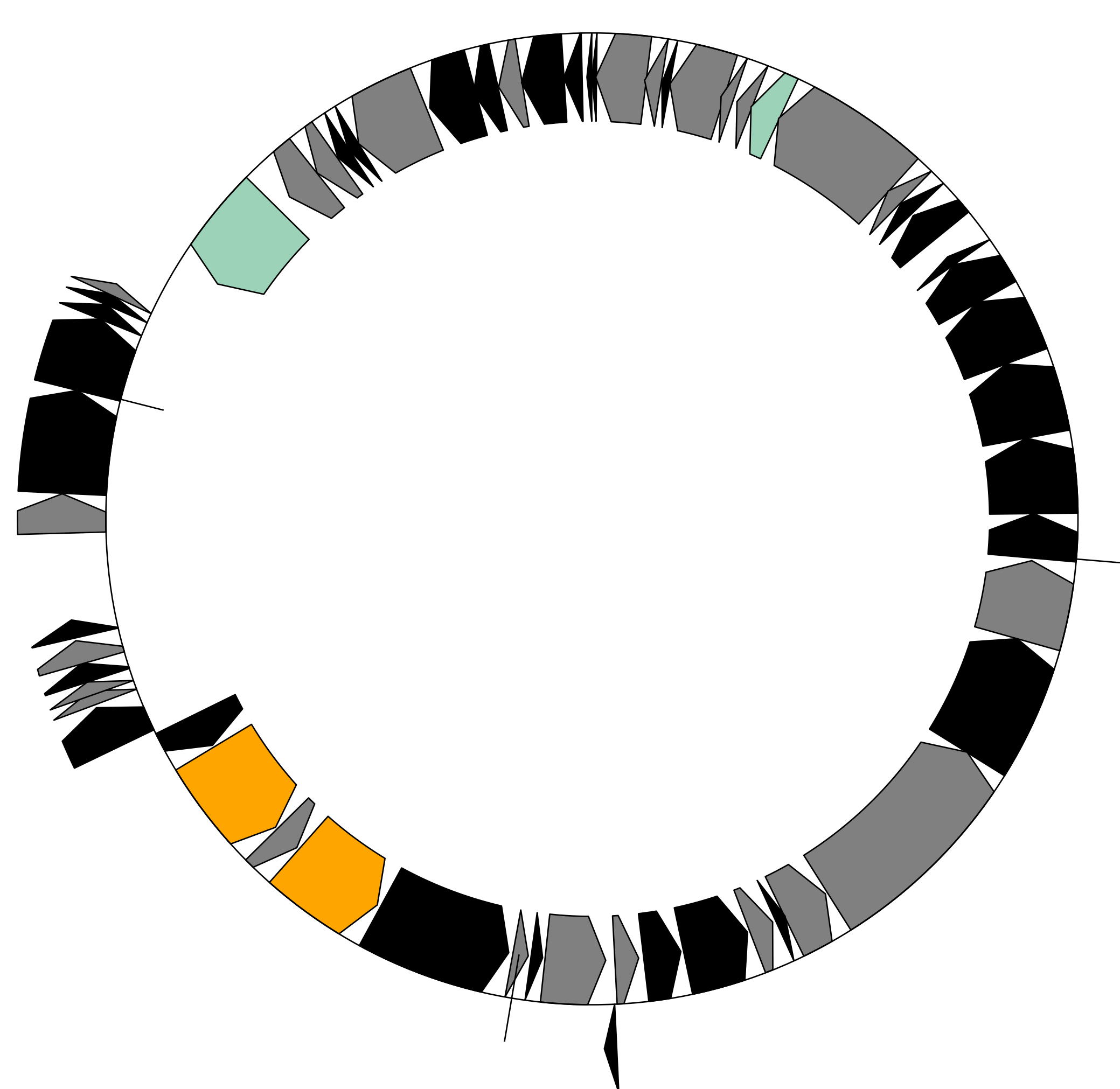
52708. putative SAR11 phage  
complete genome 143kbp



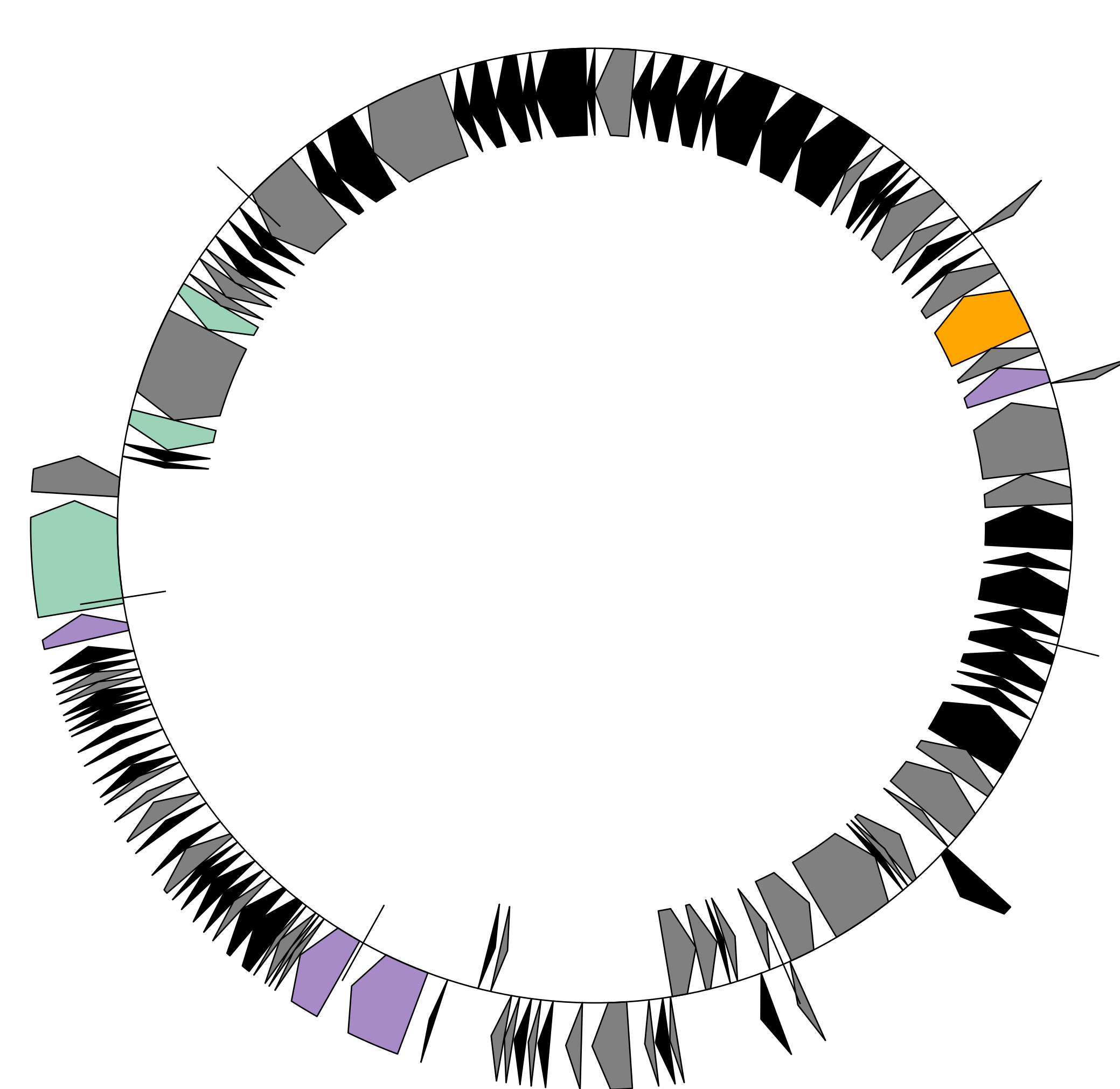
47393. putative SAR11 phage  
complete genome 33kbp



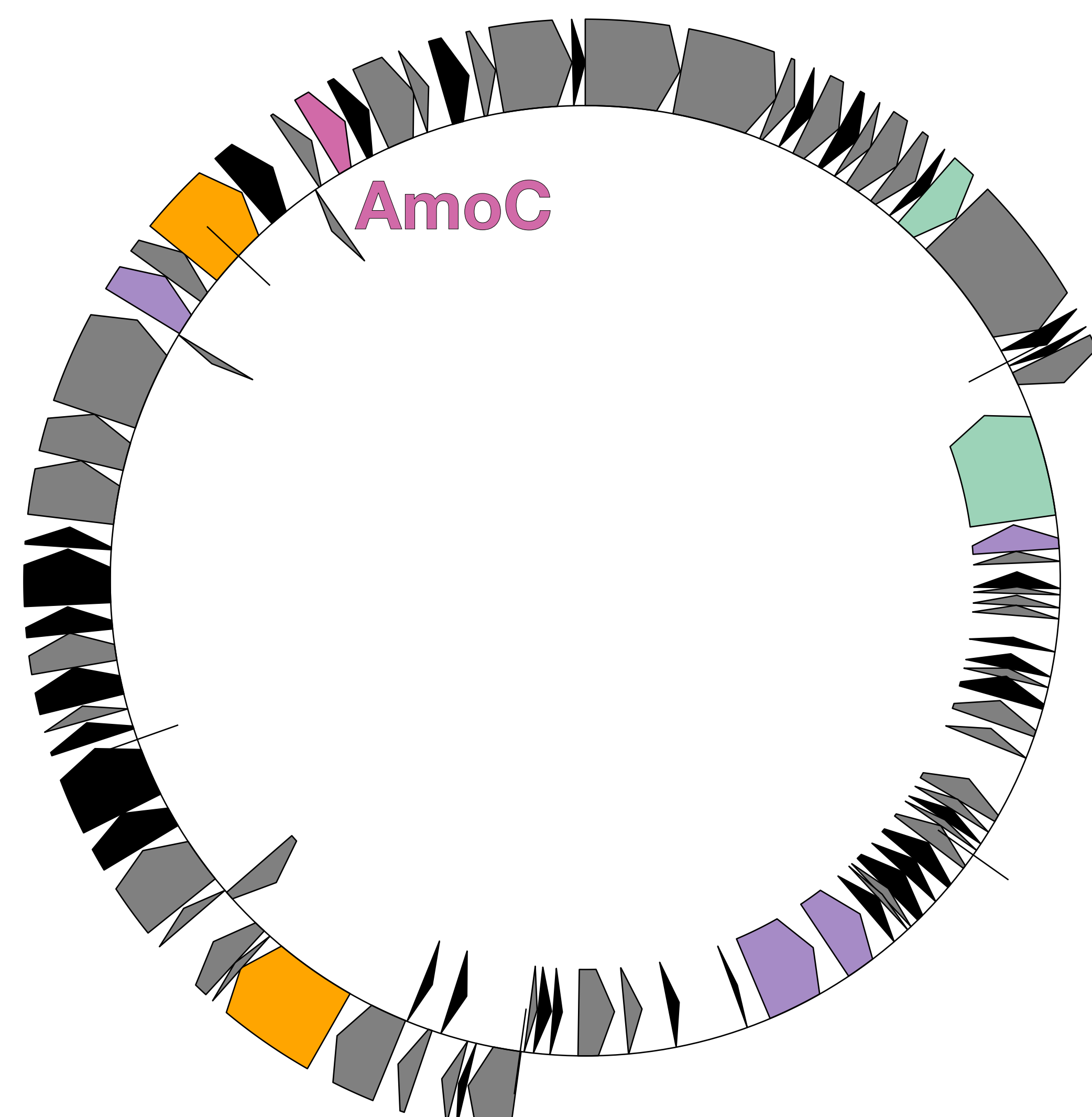
11148. putative archaeal virus  
complete genome 38kbp  
surface



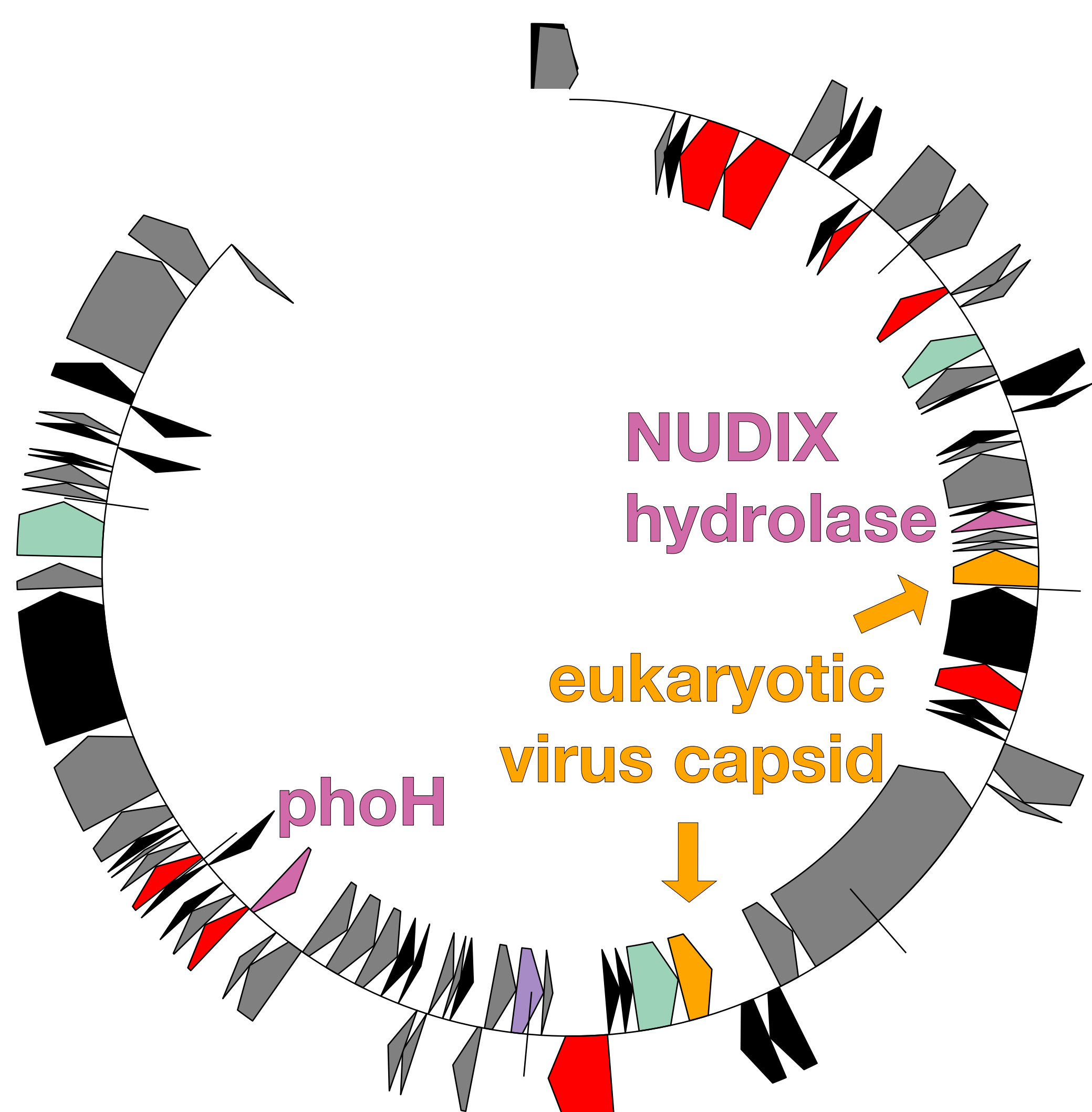
18781. putative Thaumarchaeal virus  
complete genome 69kbp  
below DCM



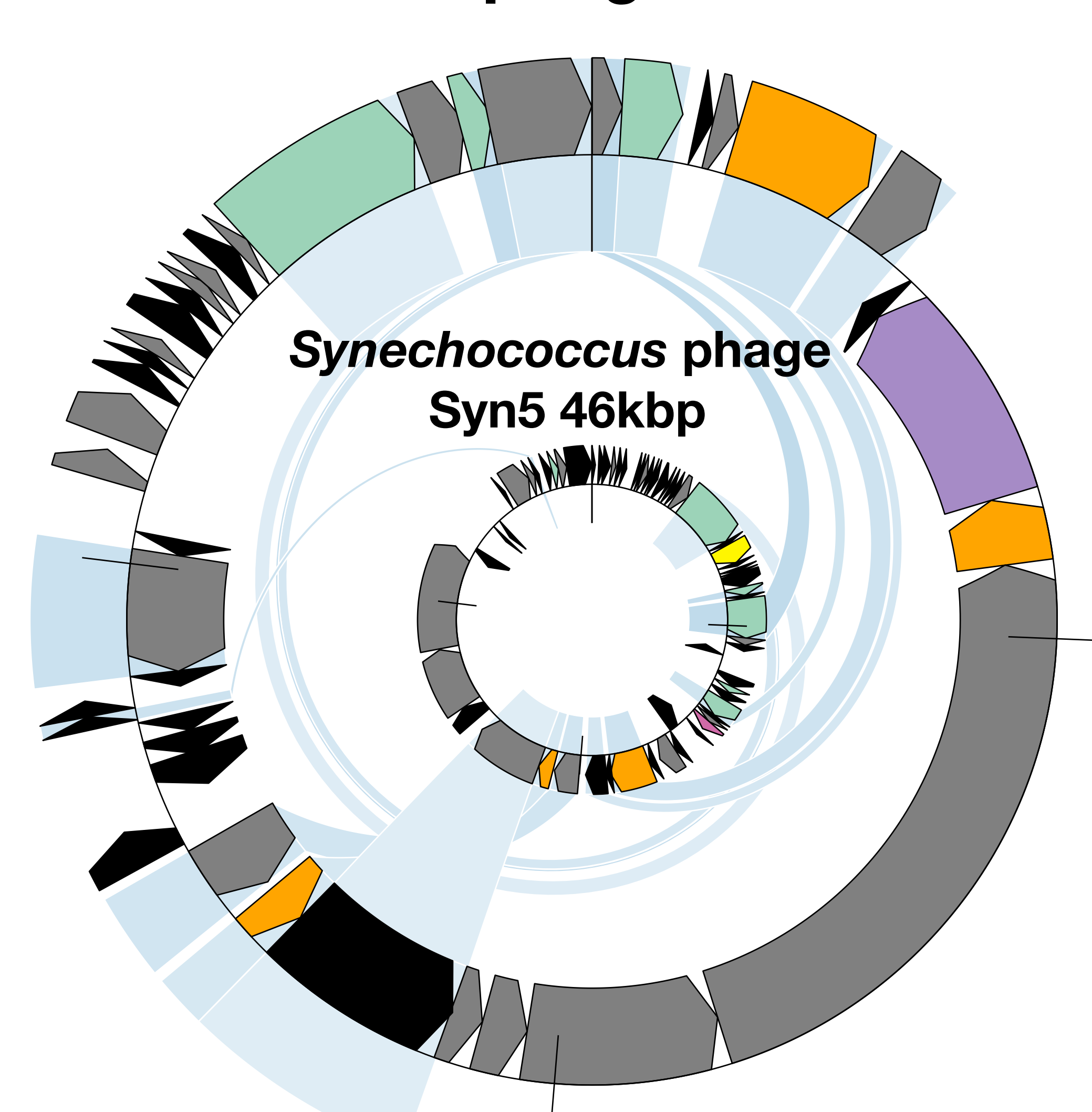
35794. putative Thaumarchaeal virus  
genomic fragment 57kbp  
below DCM



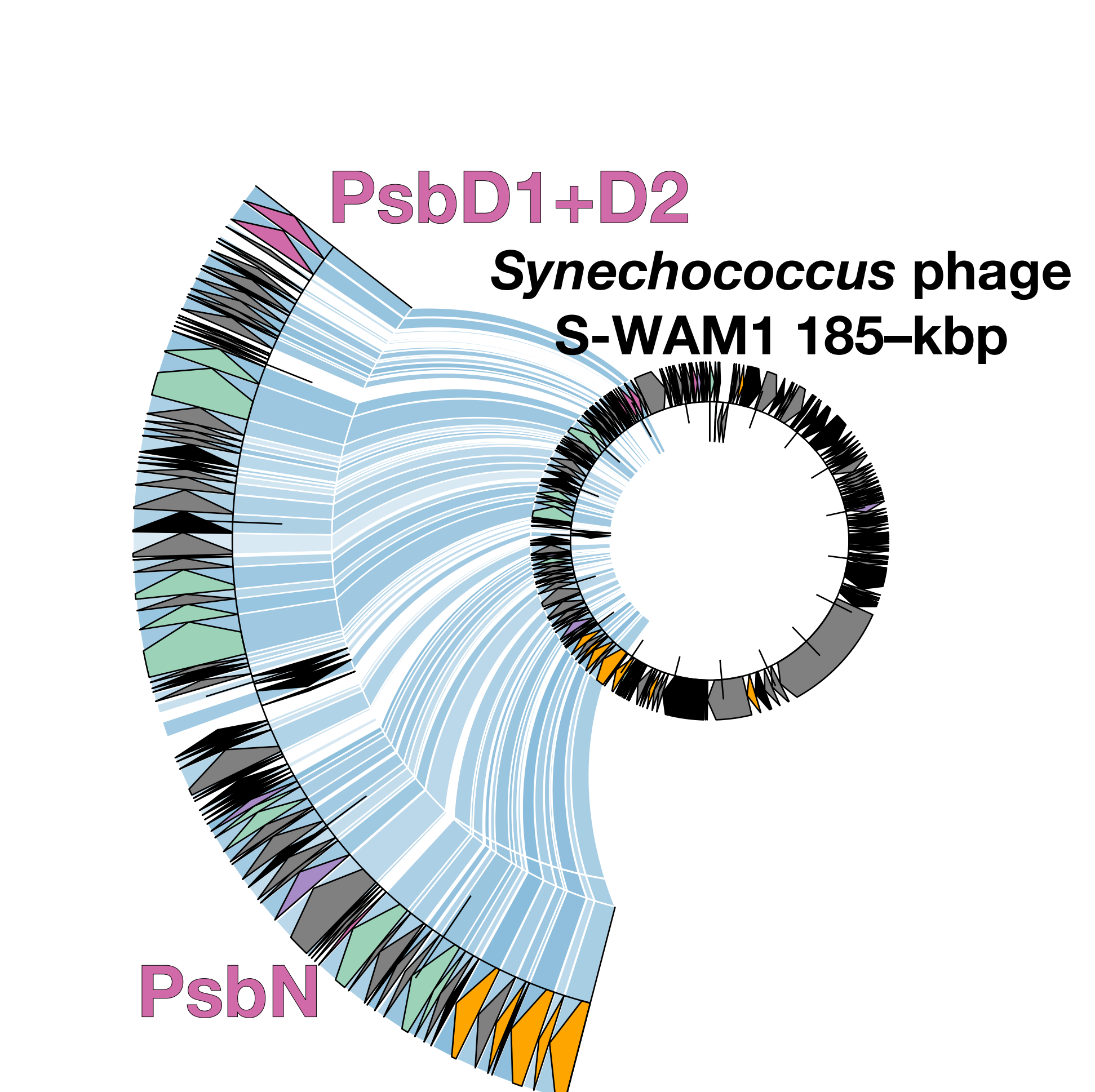
56434. putative *Bathycoccus* virus  
genomic fragment 68kbp



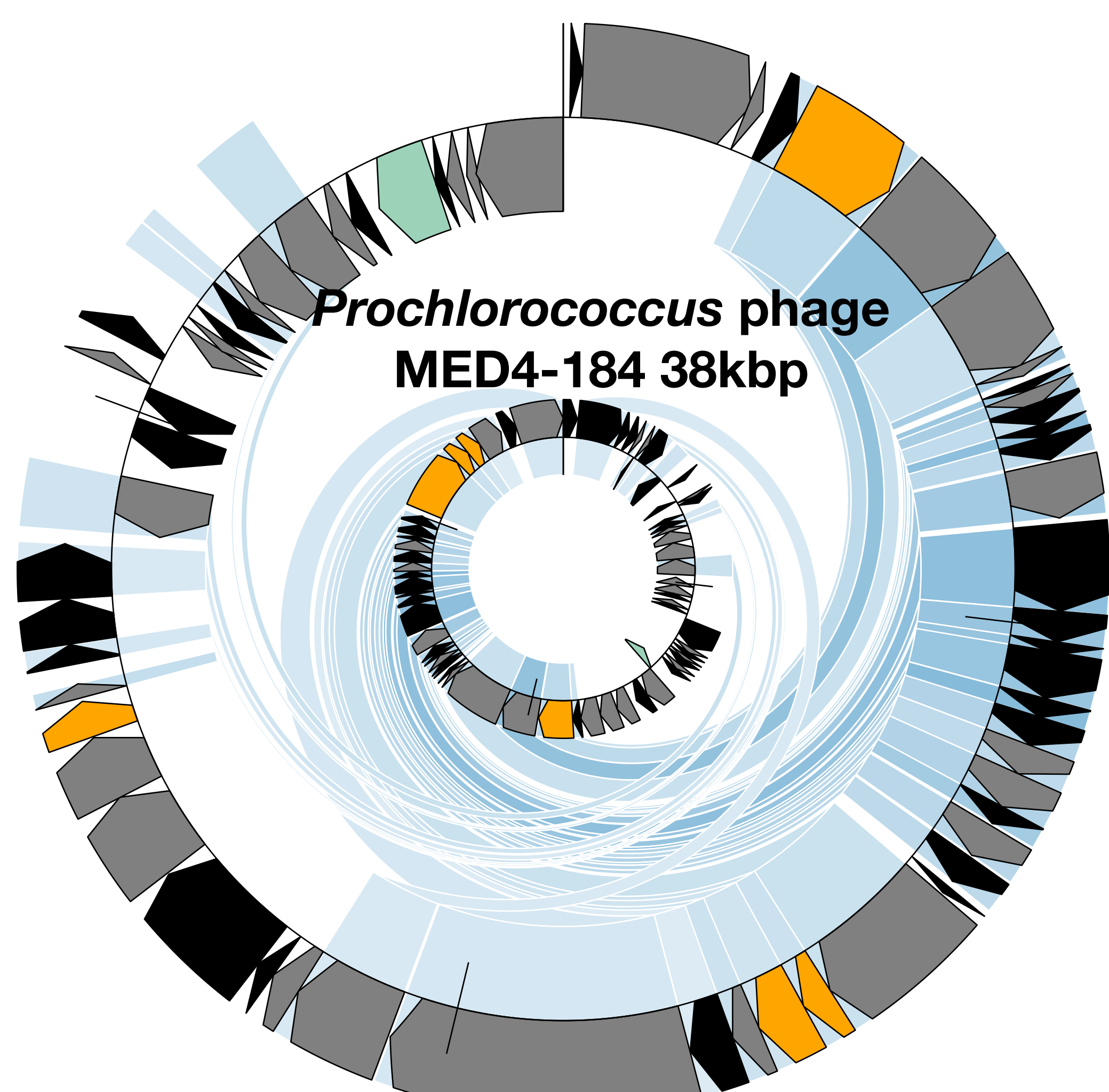
5712. putative *Synechococcus* phage  
complete genome 39kbp  
most abundant phage in cell fraction



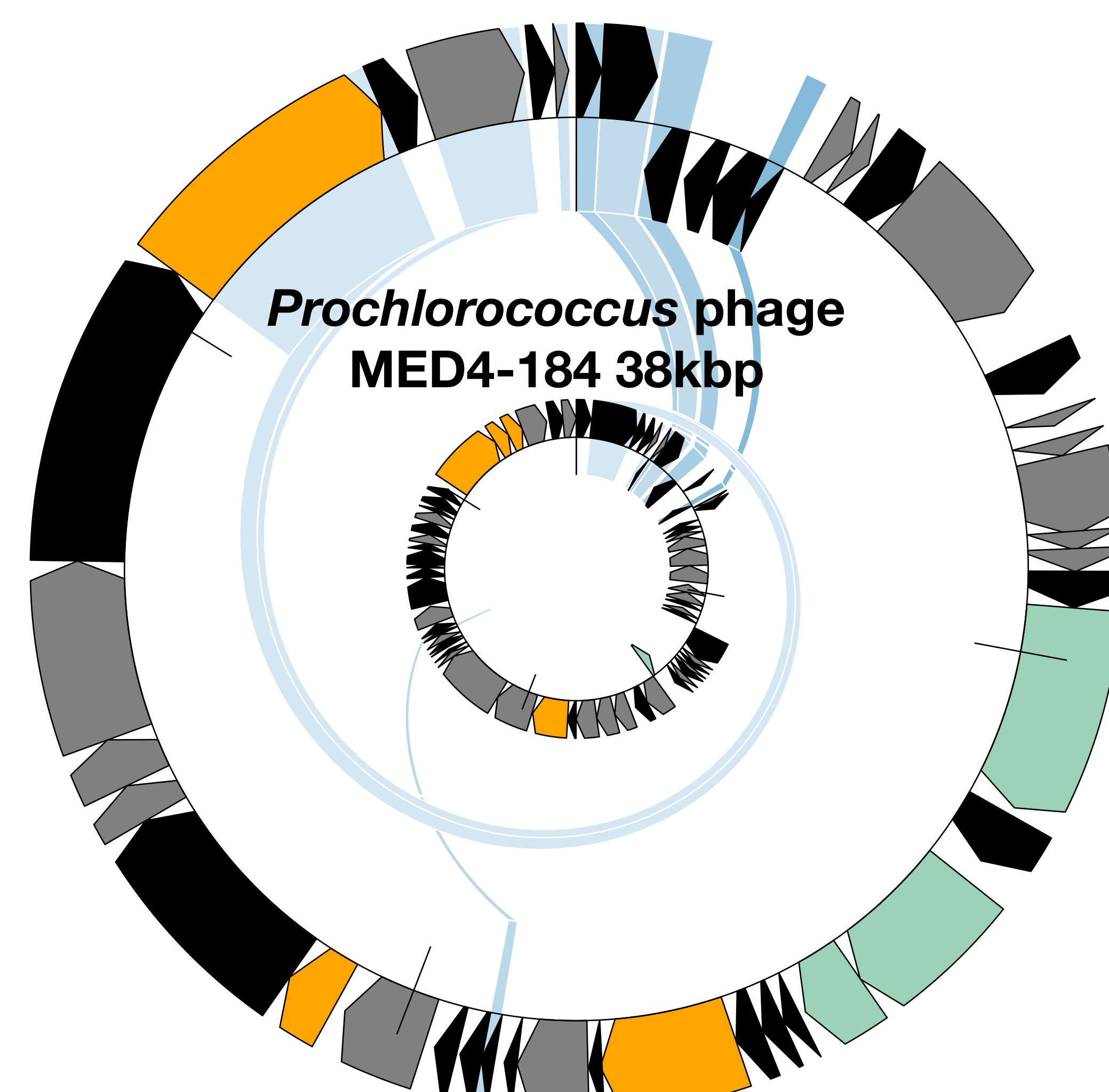
30150. putative *Synechococcus* phage  
genomic fragment 58kbp



47103. putative *Prochlorococcus* phage  
complete genome 36kbp  
most abundant phage in viral fraction



24463. putative *Prochlorococcus* phage  
complete genome 37kbp



41447. putative *Prochlorococcus* phage  
complete genome 199kbp

