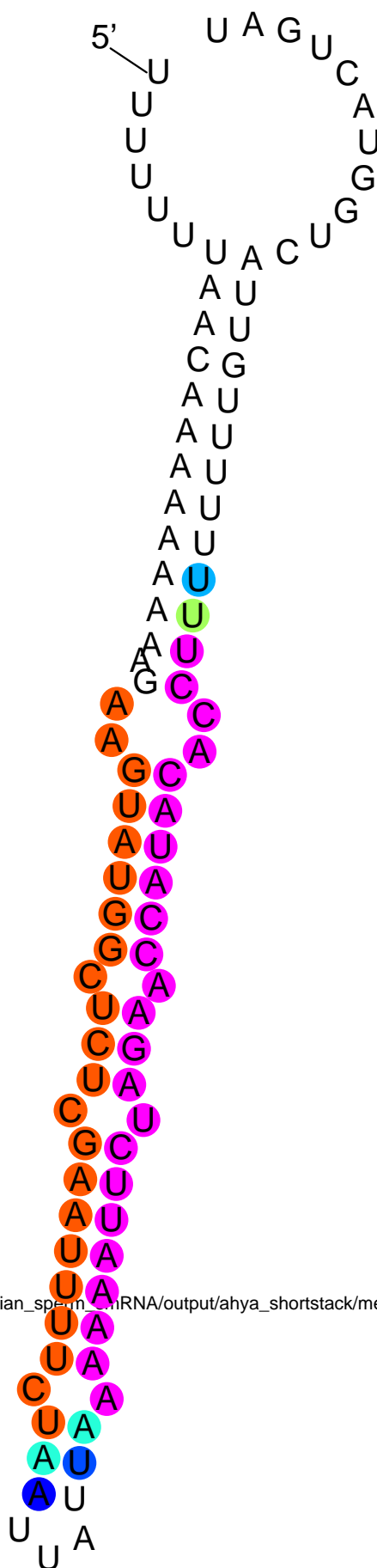


Depth of Coverage



Genome: Ahyacinthus.chrsV1.fasta

Alignments: /work/pi_hputnam_uri_edu/jillashey/cnidarian_spe-miRNA/output/ahya_shortstack/merged_alignments.bam

Location: chr3:26150297-26150391 minus

Name: Cluster_3705