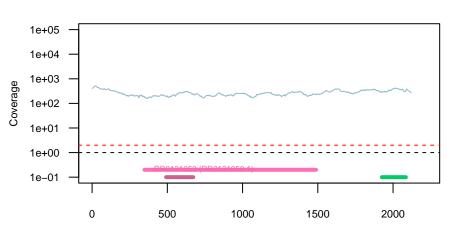
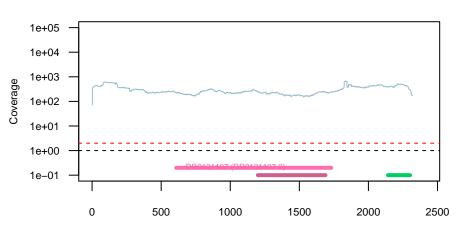
Apla_XP_022090421.1, Apla_XM_022234729.1 (1st domain pair)



Transcript (bp)

arch: HTH_Tnp_Tc5 PHD

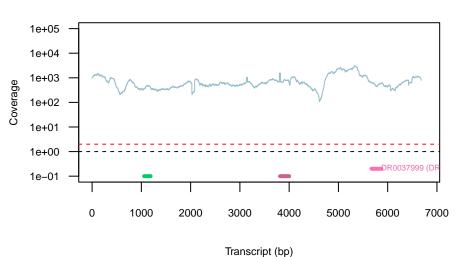
Apla_XP_022090455.1, Apla_XM_022234763.1 (1st domain pair)



Transcript (bp)

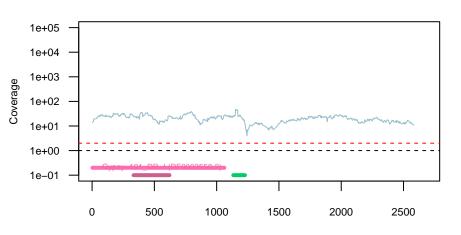
arch: HTH_Tnp_Tc5 DDE_1 PHD

Apla_XP_022104216.1, Apla_XM_022248524.1 (1st domain pair)



arch: CHDNT PHD PHD Chromo SNF2_N Helicase_C DUF1087 DUF1086 CHDCT2

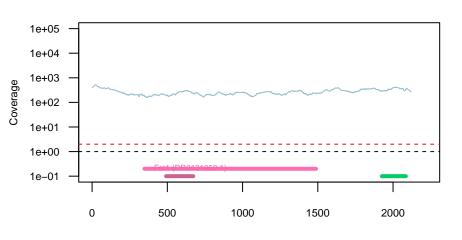
Apla_XP_022111688.1, Apla_XM_022255996.1 (1st domain pair)



Transcript (bp)

arch: Integrase_H2C2 rve PHD TatD_DNase

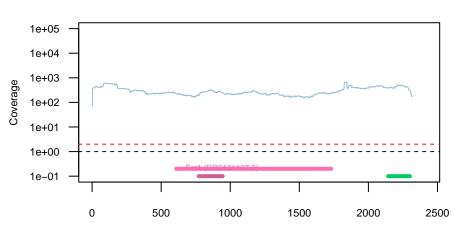
Apla_XP_022090421.1, Apla_XM_022234729.1 (2nd domain pair)



Transcript (bp)

arch: HTH_Tnp_Tc5 PHD

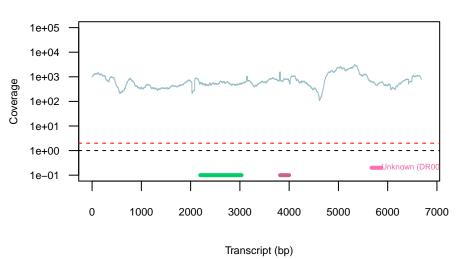
Apla_XP_022090455.1, Apla_XM_022234763.1 (2nd domain pair)



Transcript (bp)

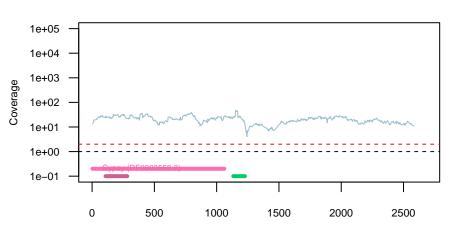
arch: HTH_Tnp_Tc5 DDE_1 PHD

Apla_XP_022104216.1, Apla_XM_022248524.1 (2nd domain pair)



arch: CHDNT PHD PHD Chromo SNF2_N Helicase_C DUF1087 DUF1086 CHDCT2

Apla_XP_022111688.1, Apla_XM_022255996.1 (2nd domain pair)



Transcript (bp)

arch: Integrase_H2C2 rve PHD TatD_DNase