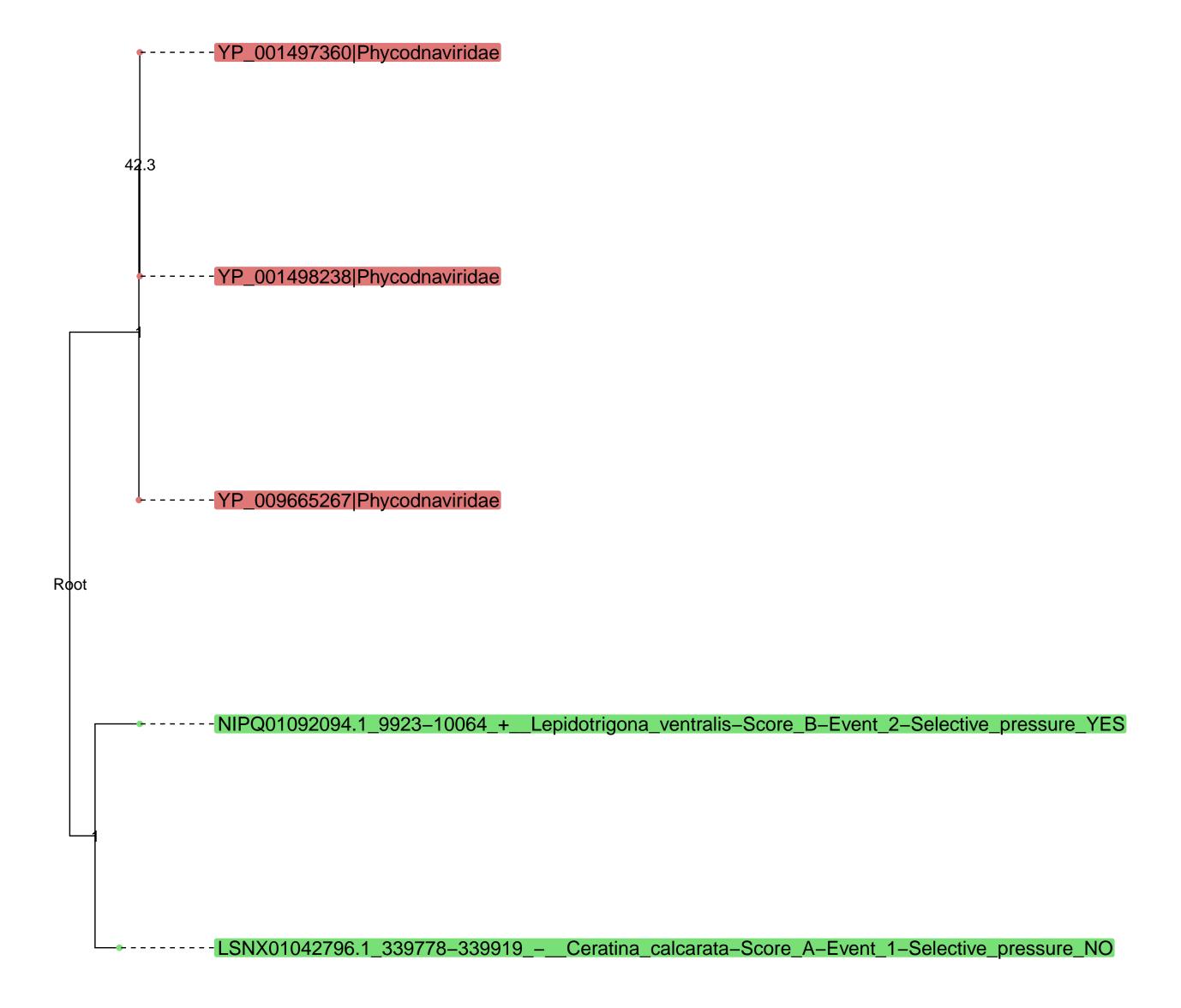


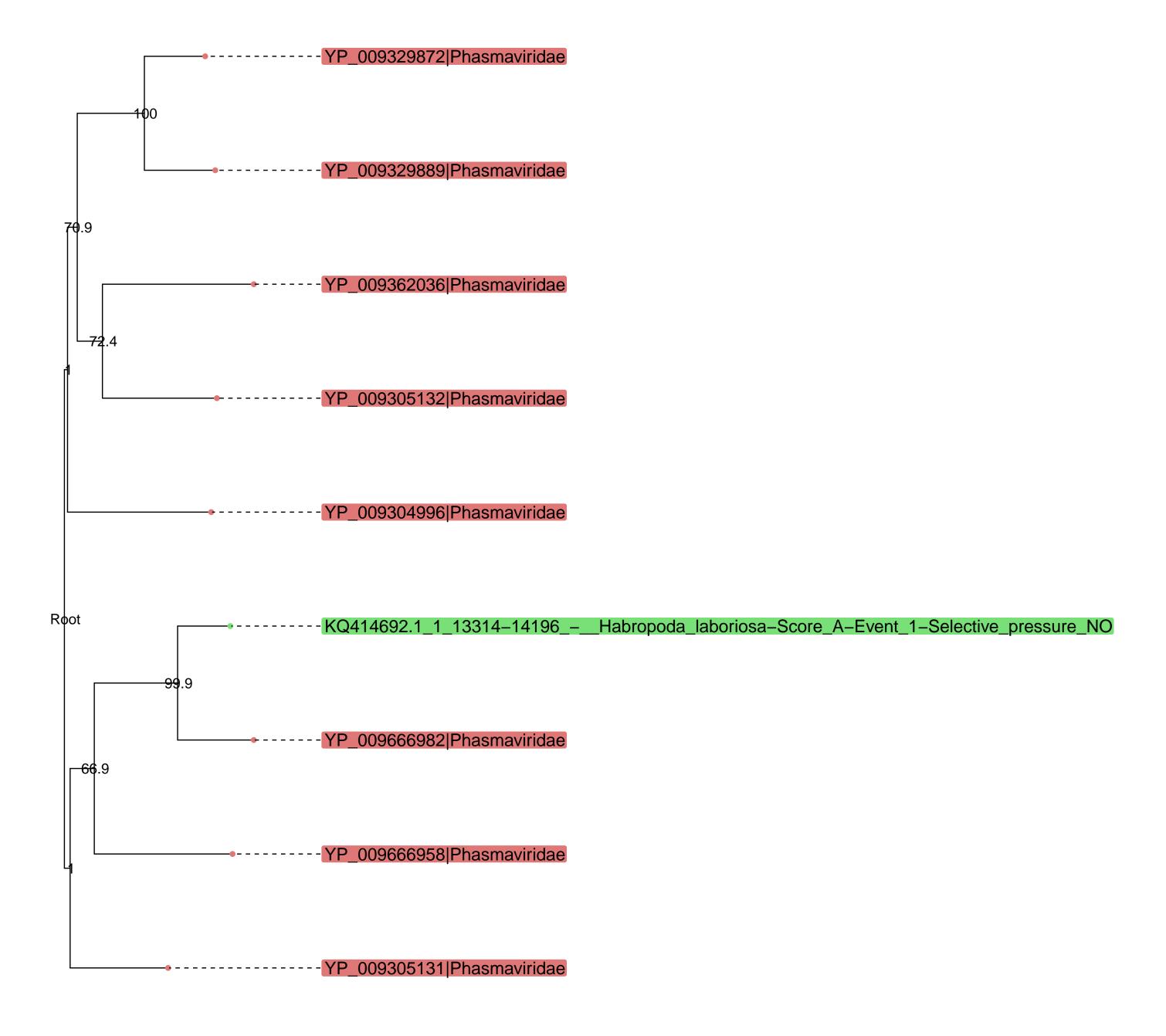
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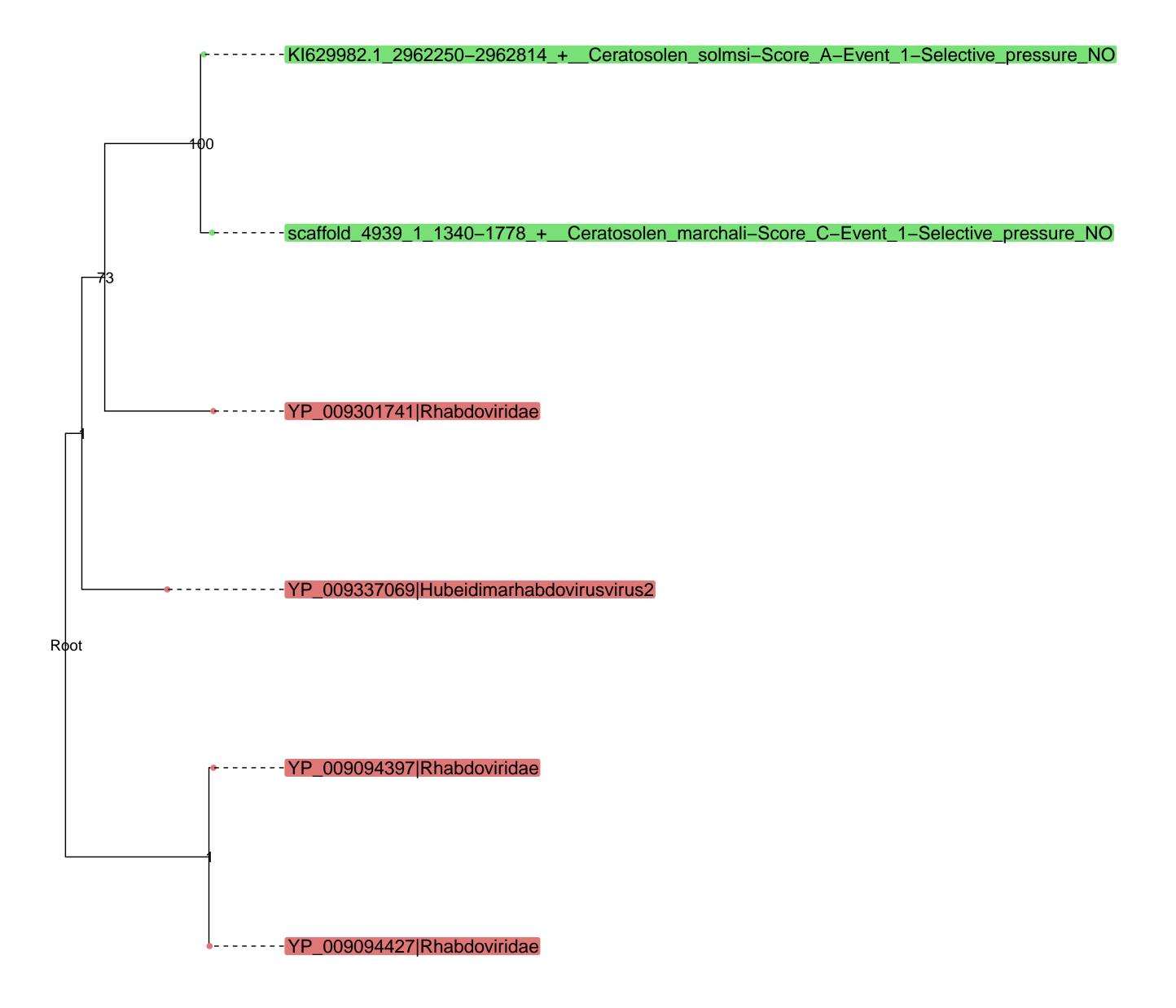
IDBA_scaffold_38286_12-663_+__Leptopilina_boulardi-Score_C-Event_2-Selective_pressure_YES

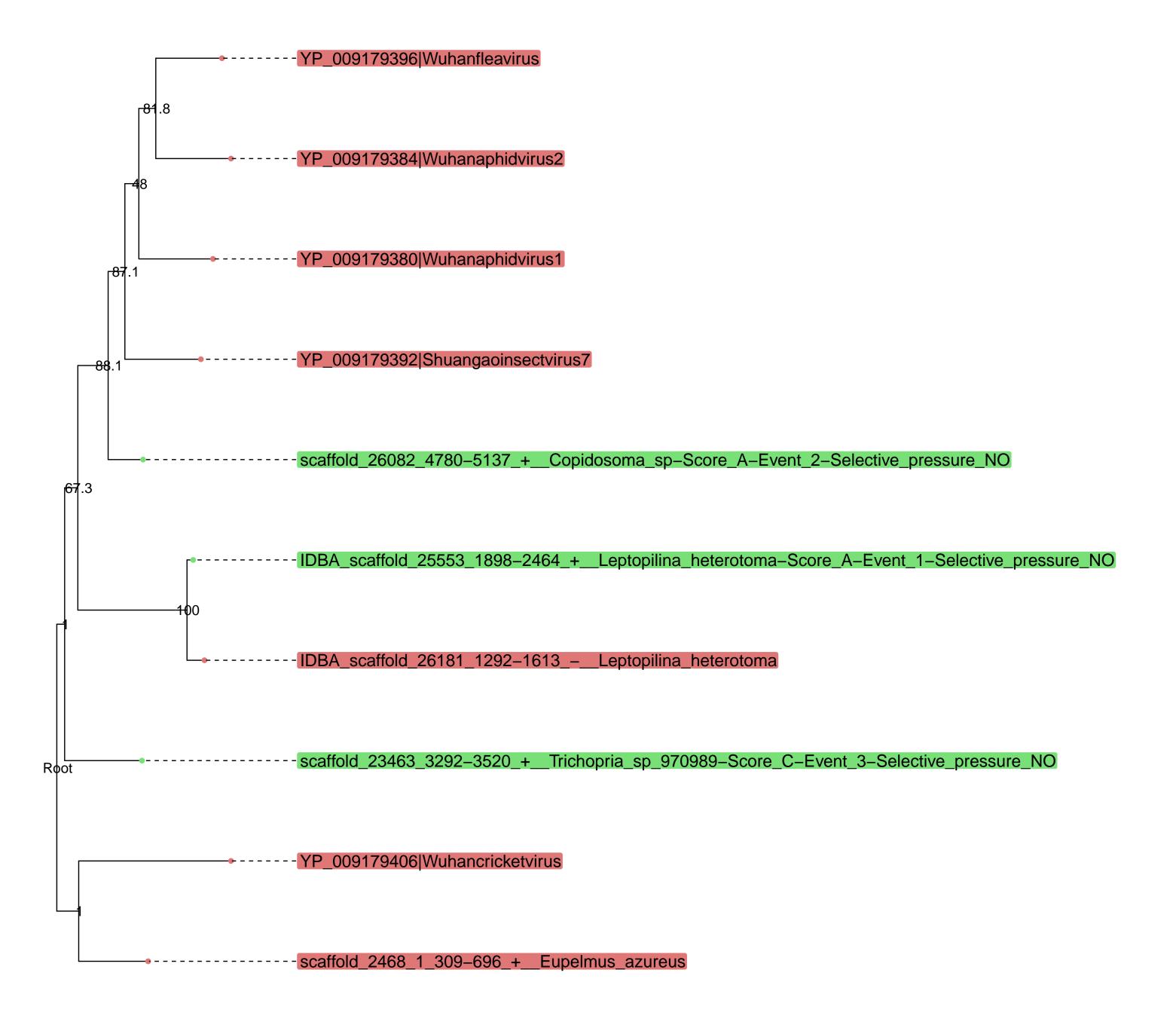
scaffold_4991_9447-10632_+__Ganaspis_brasiliensis-Score_B-Event_4-Selective_pressure_NO

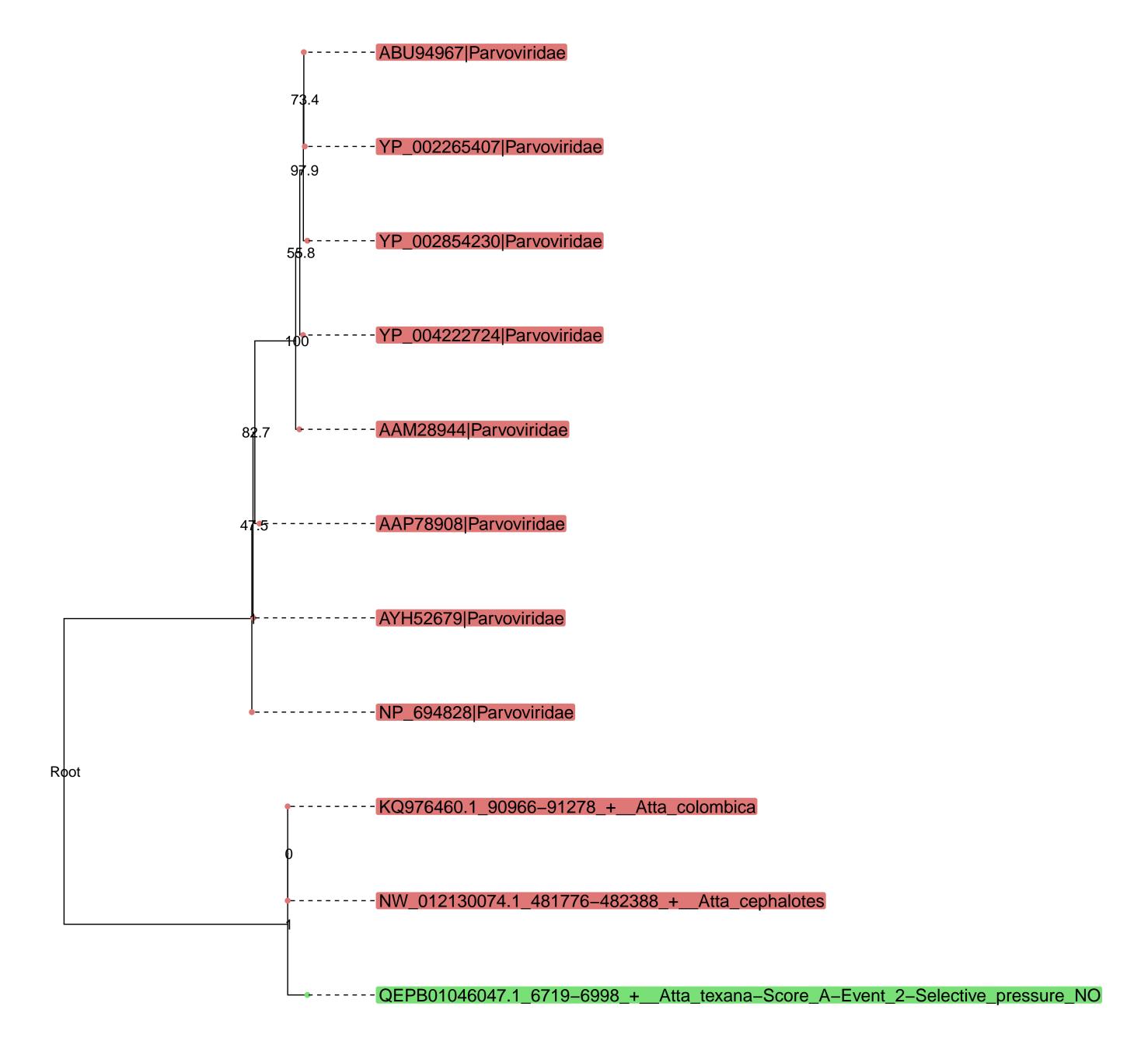
scaffold_2445_0_18385-20359_+__Leptopilina_boulardi-Score_A-Event_1-Selective_pressure_NO

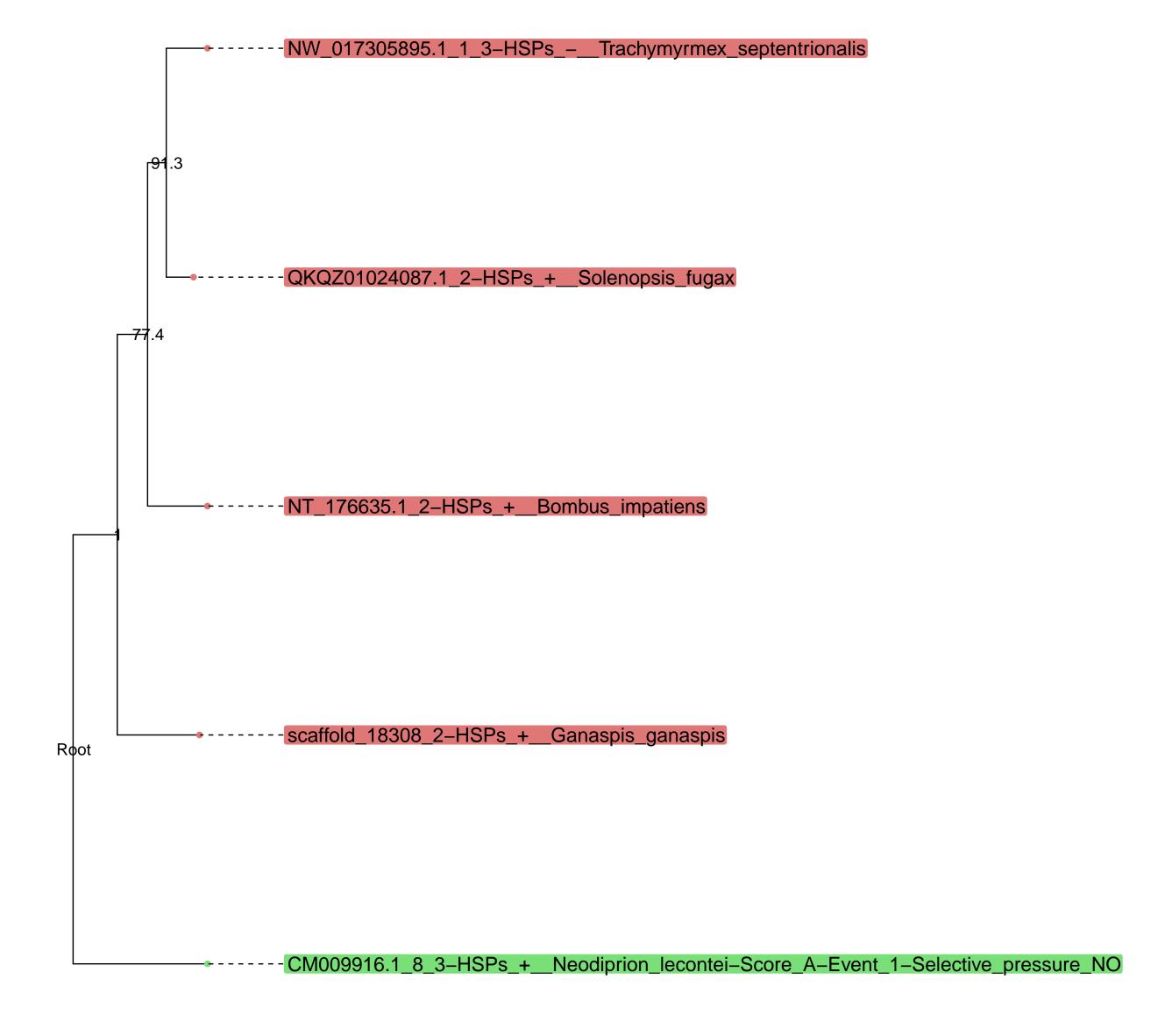






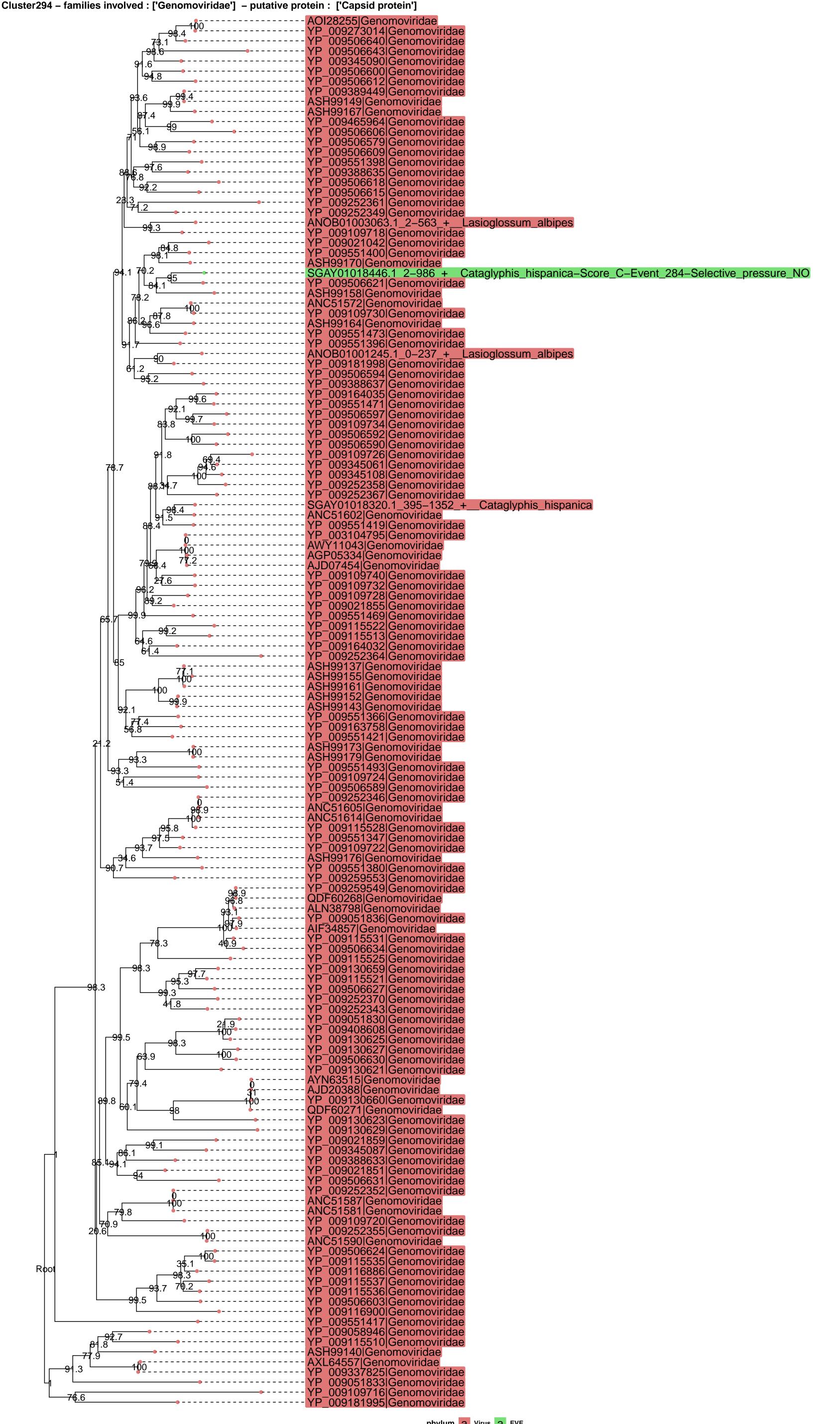


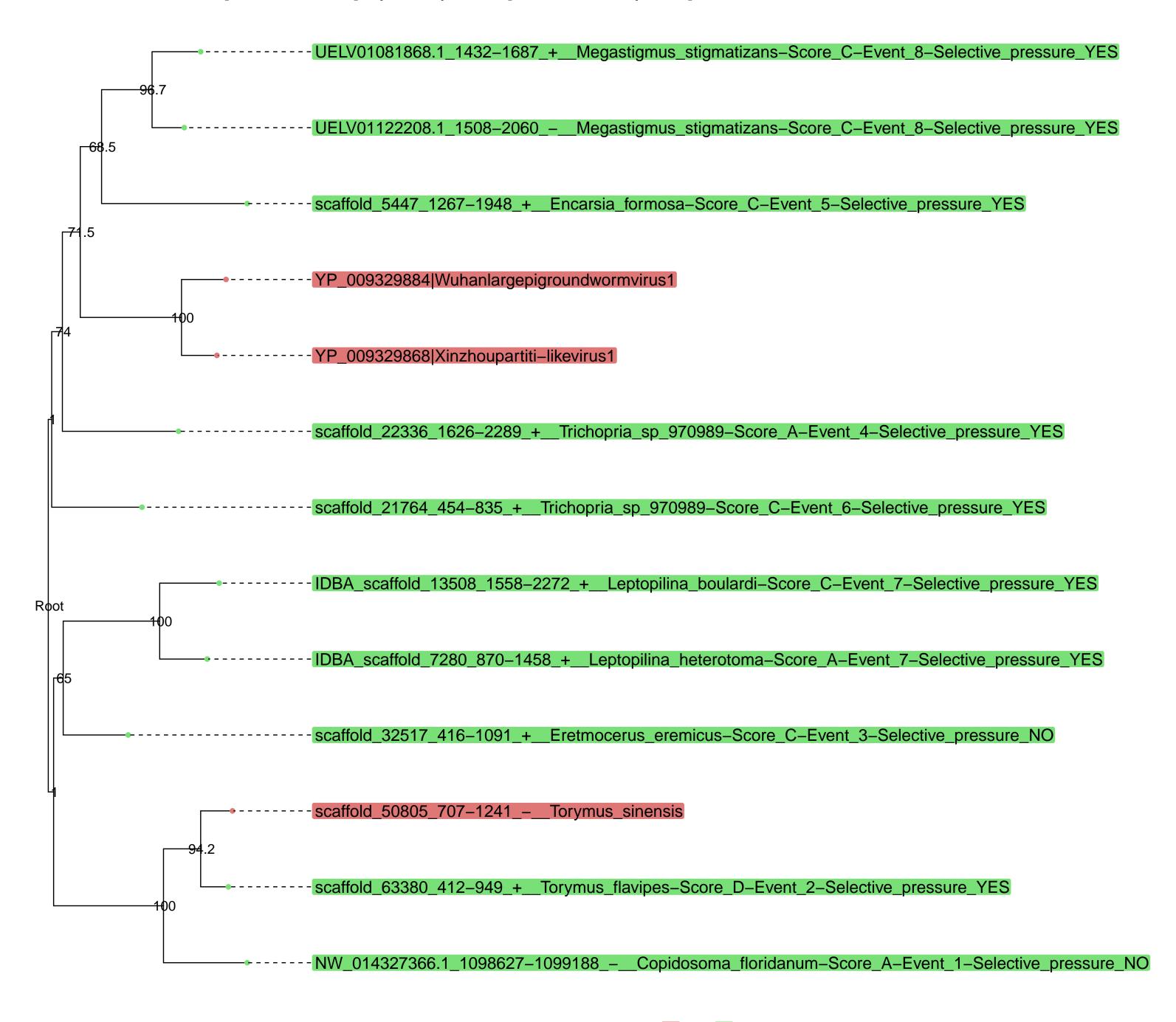


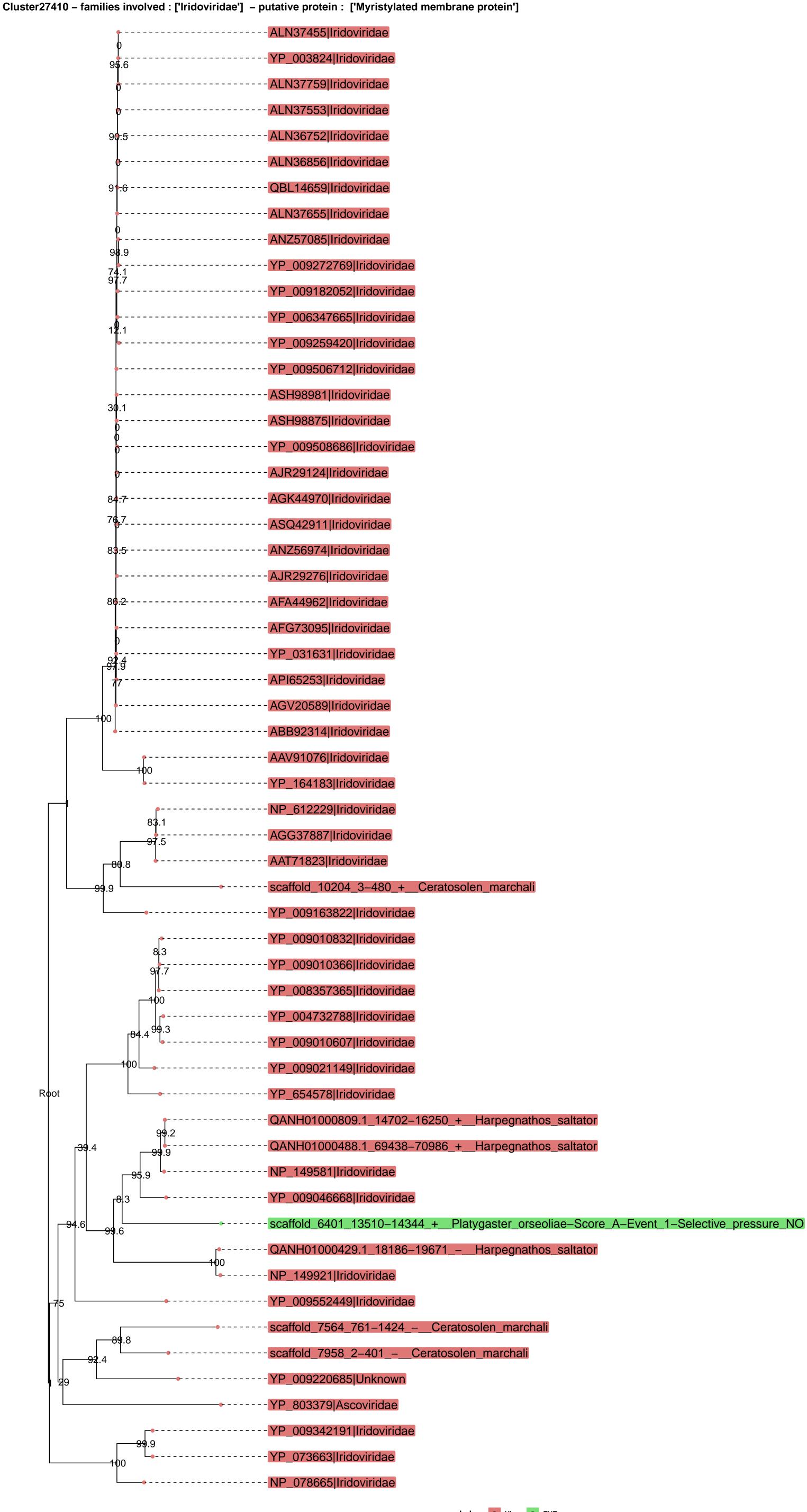


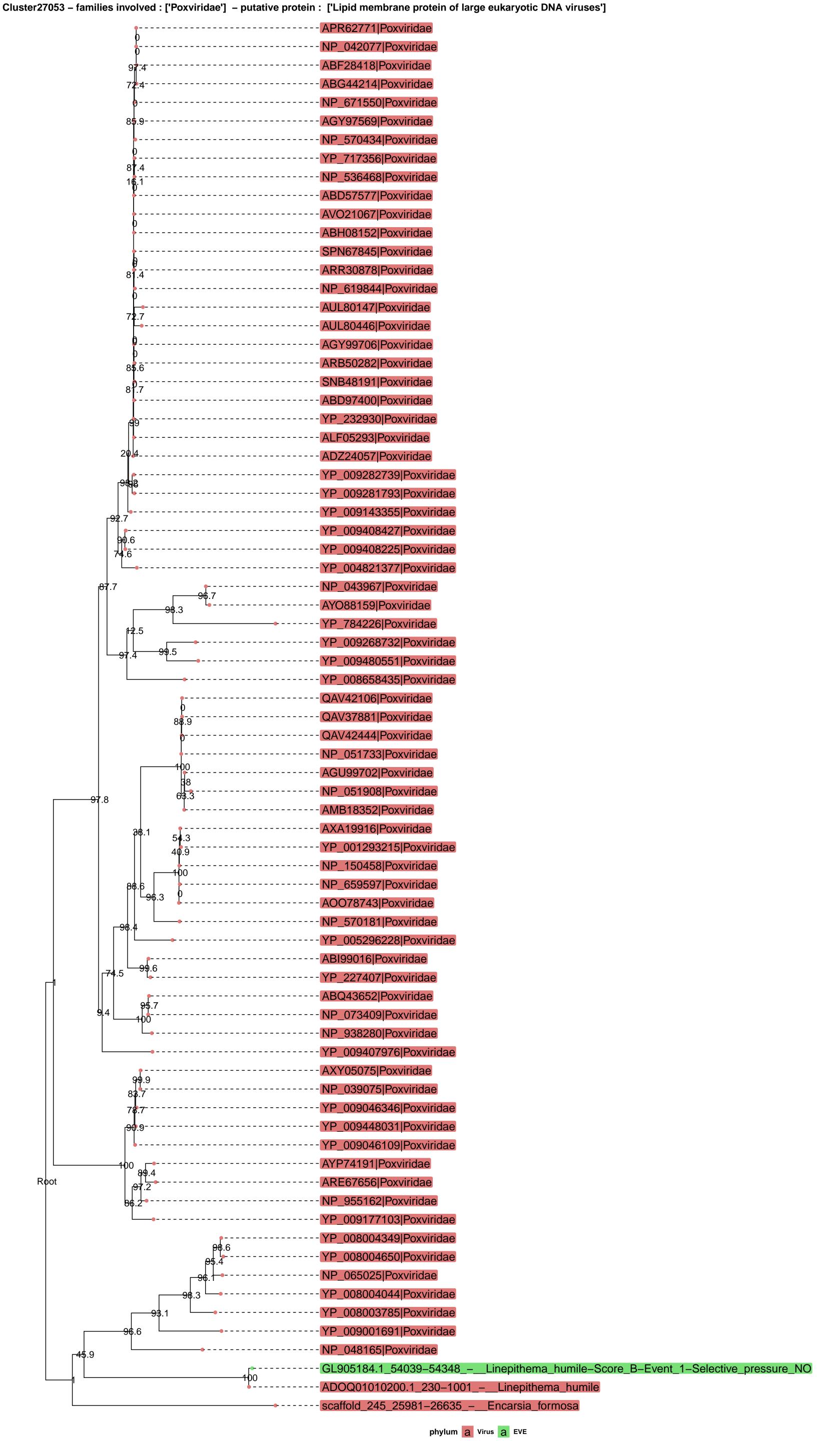


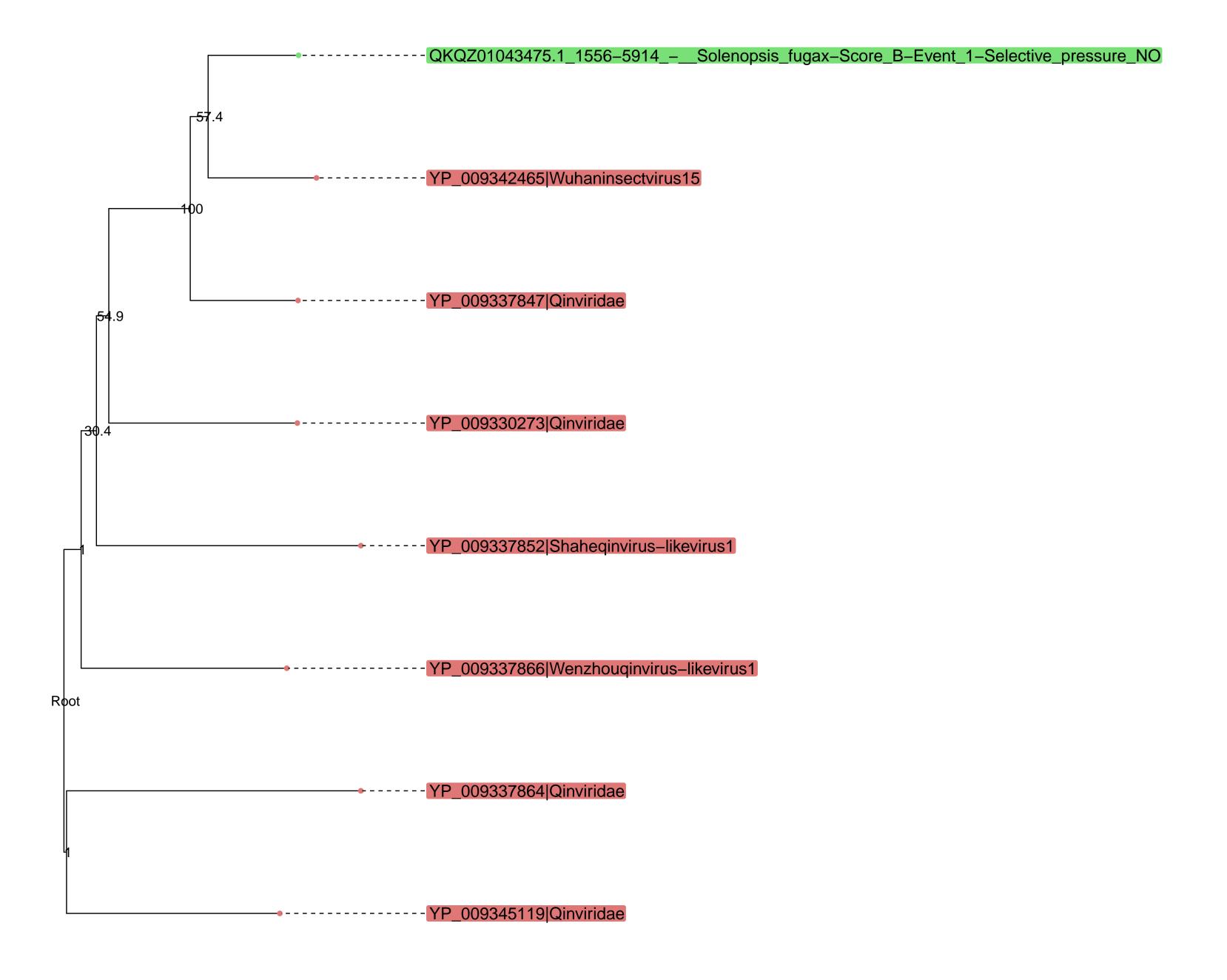


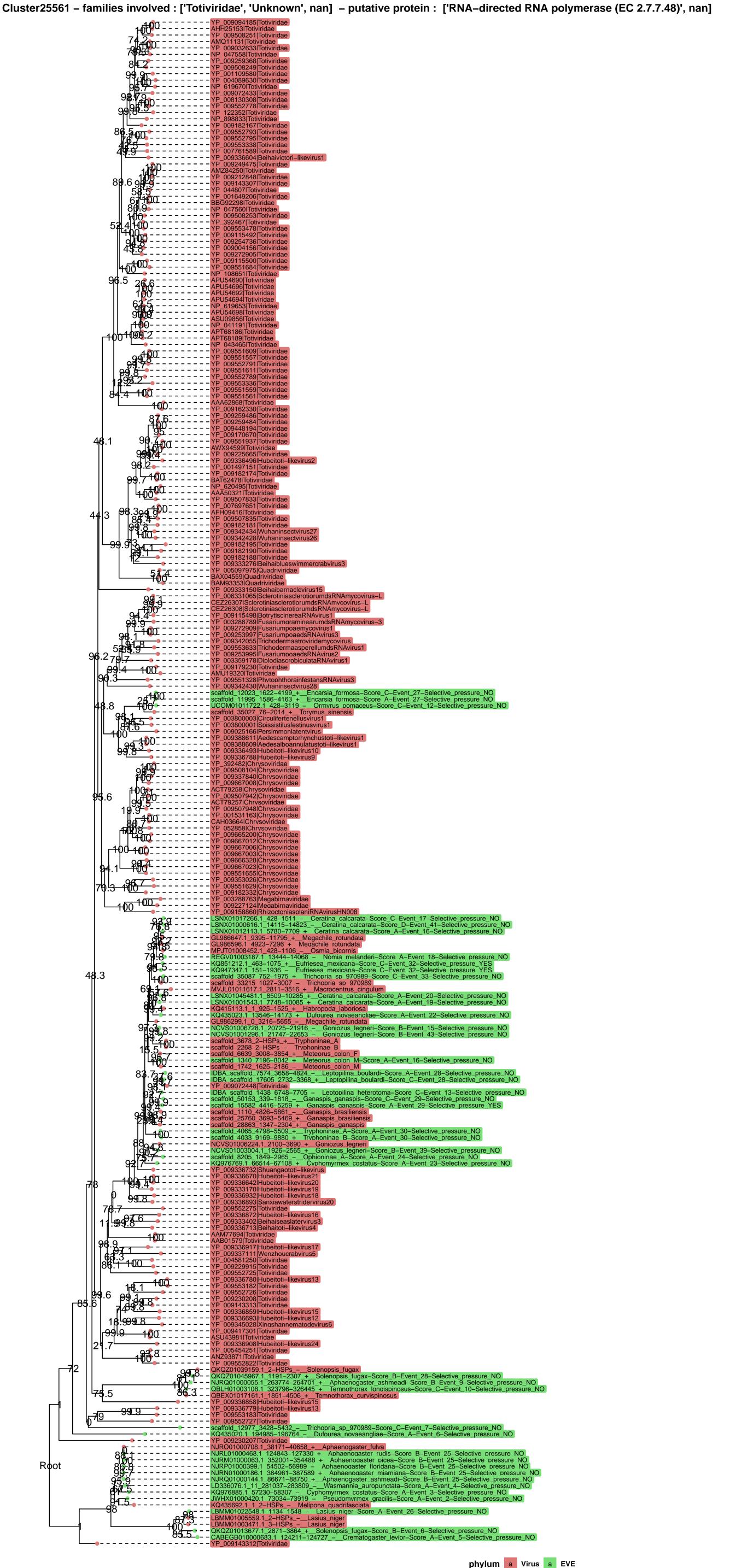












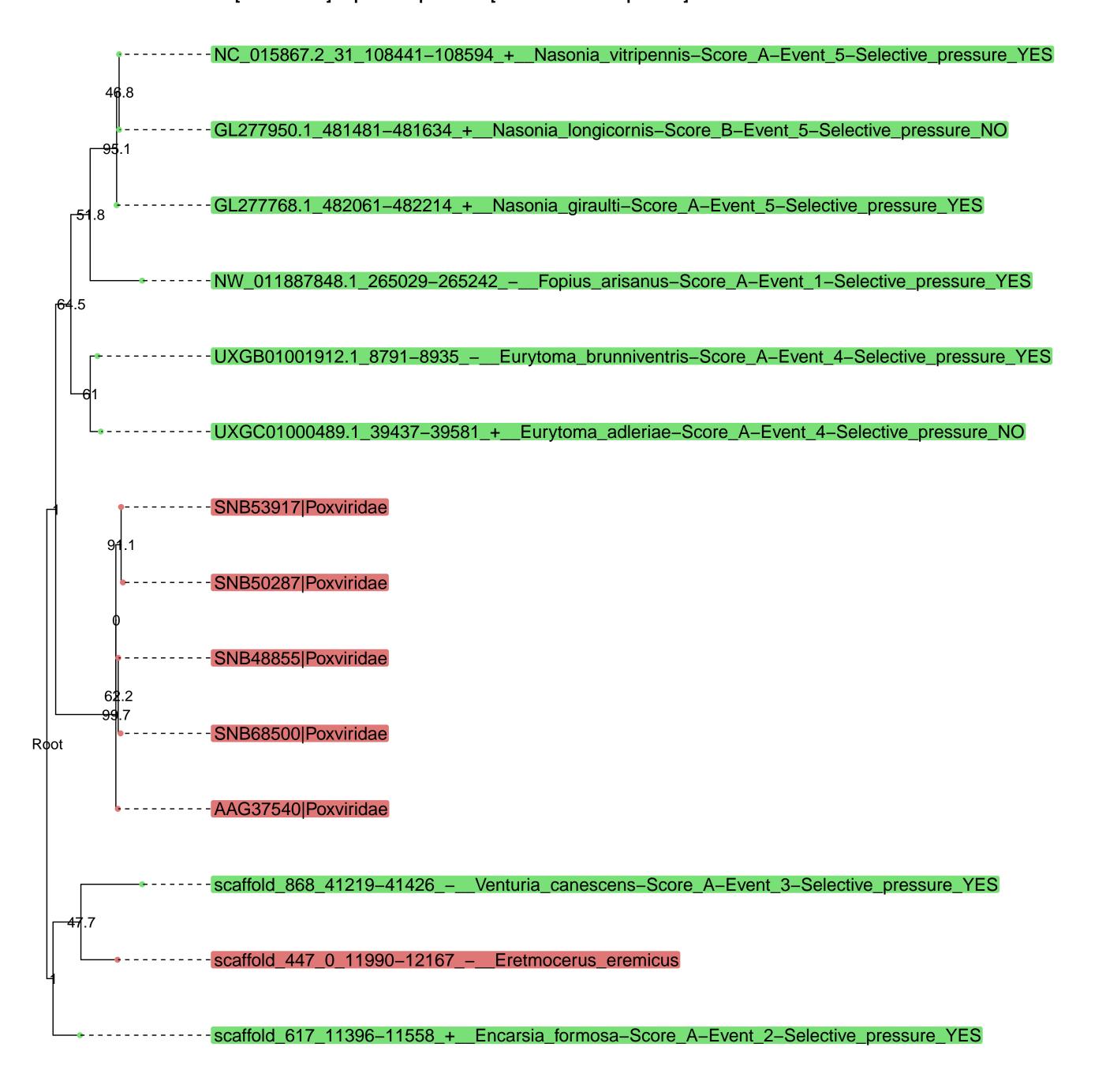
Root

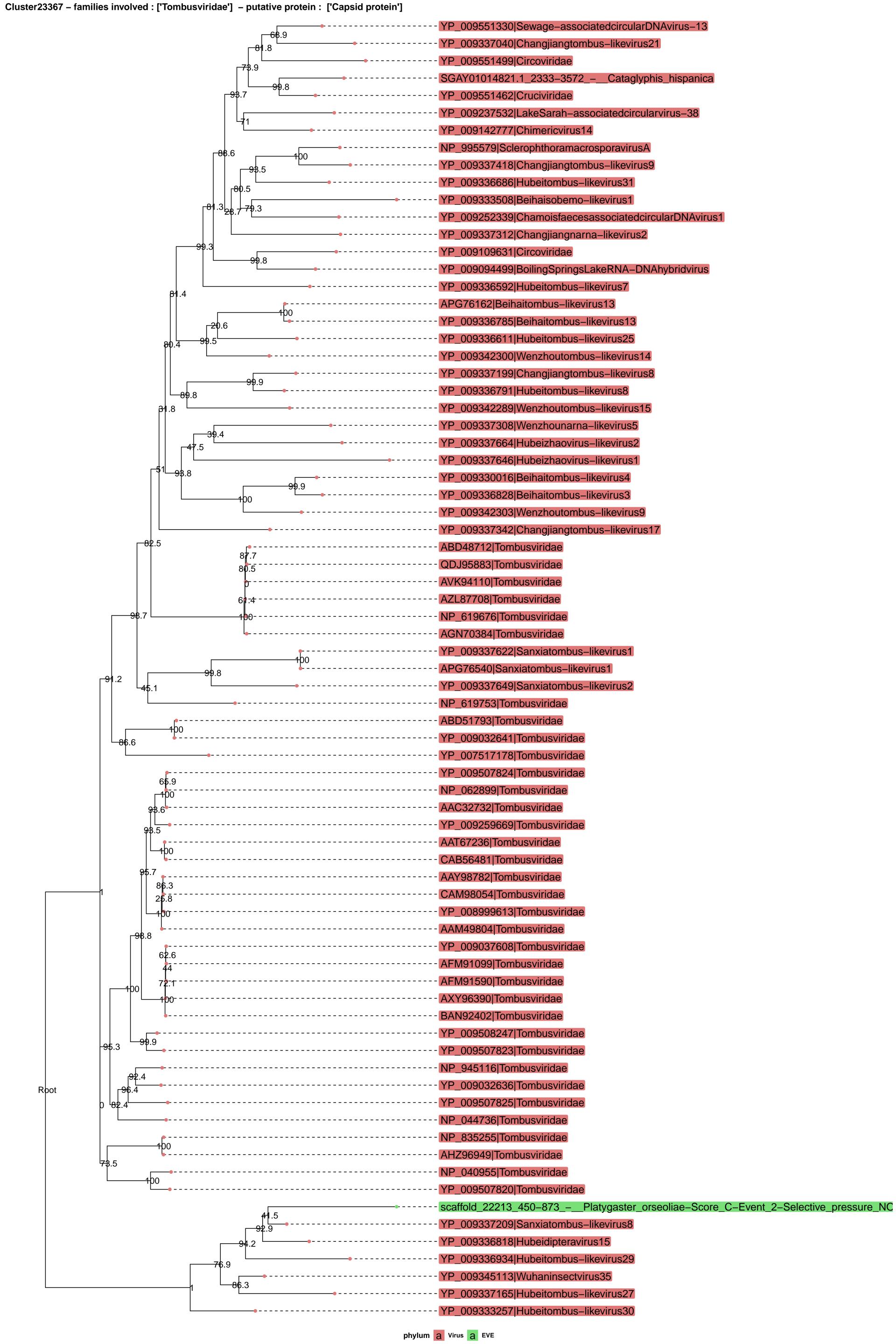
AGO32218|Parvoviridae

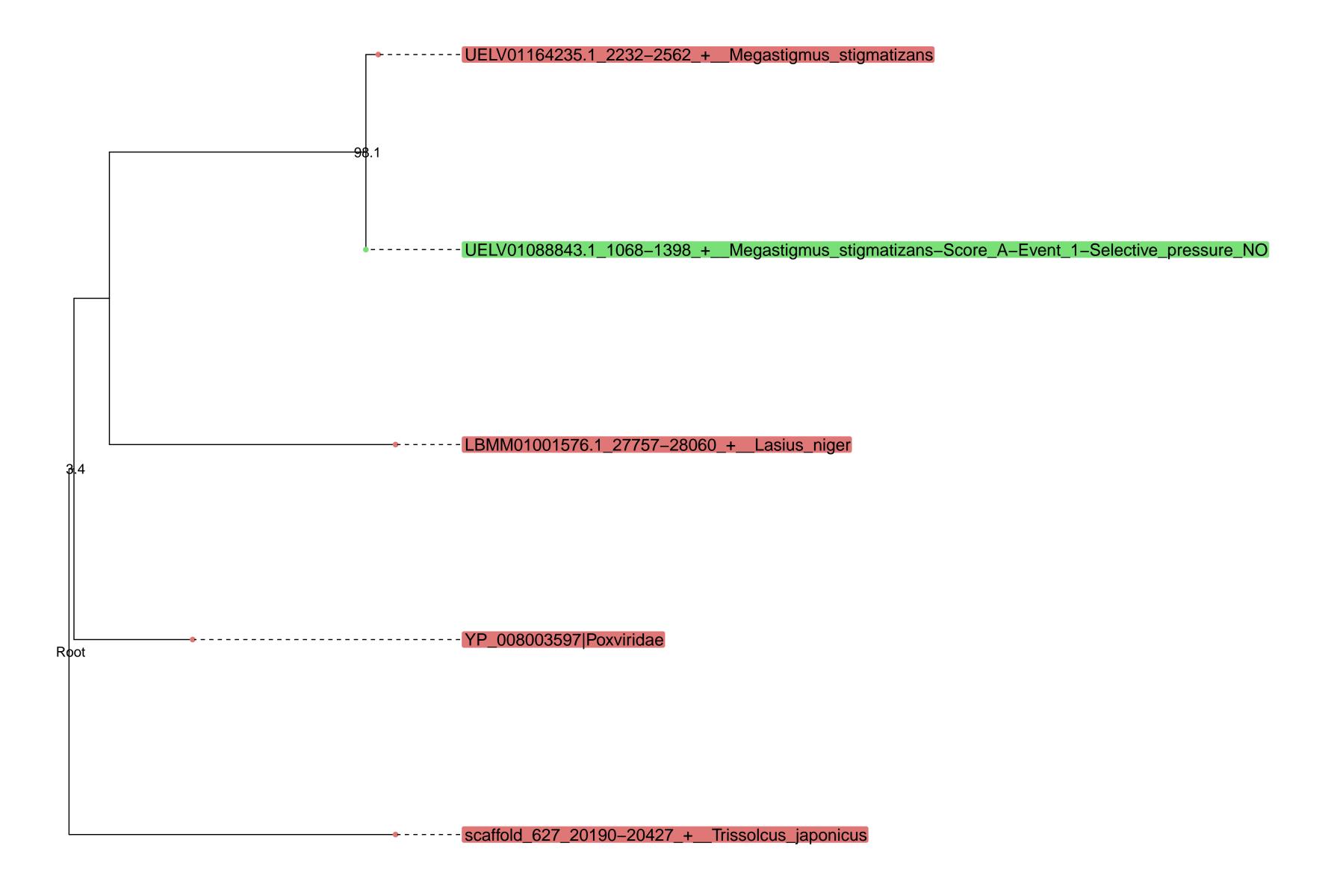
-YP_227599|Parvoviridae

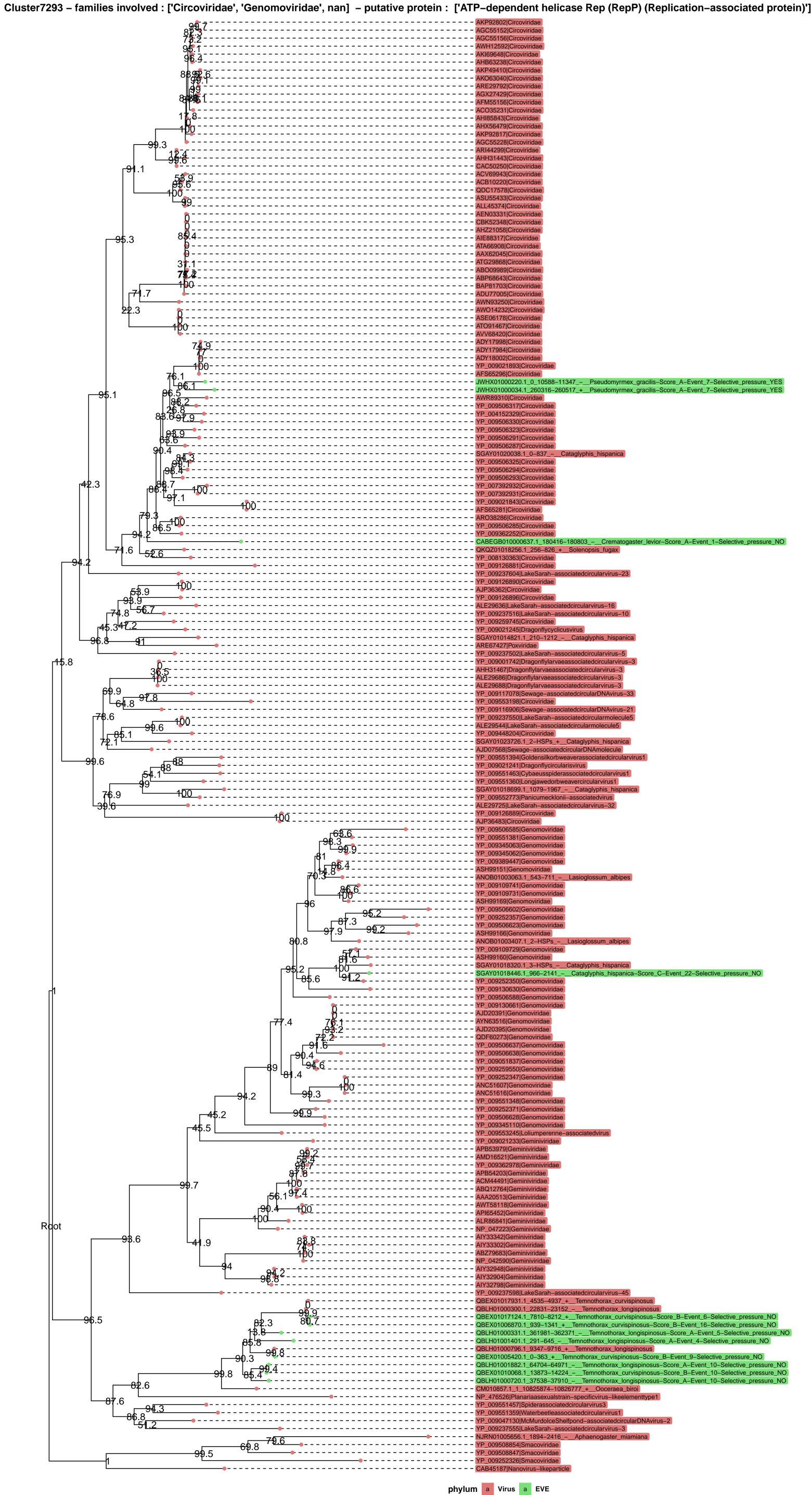


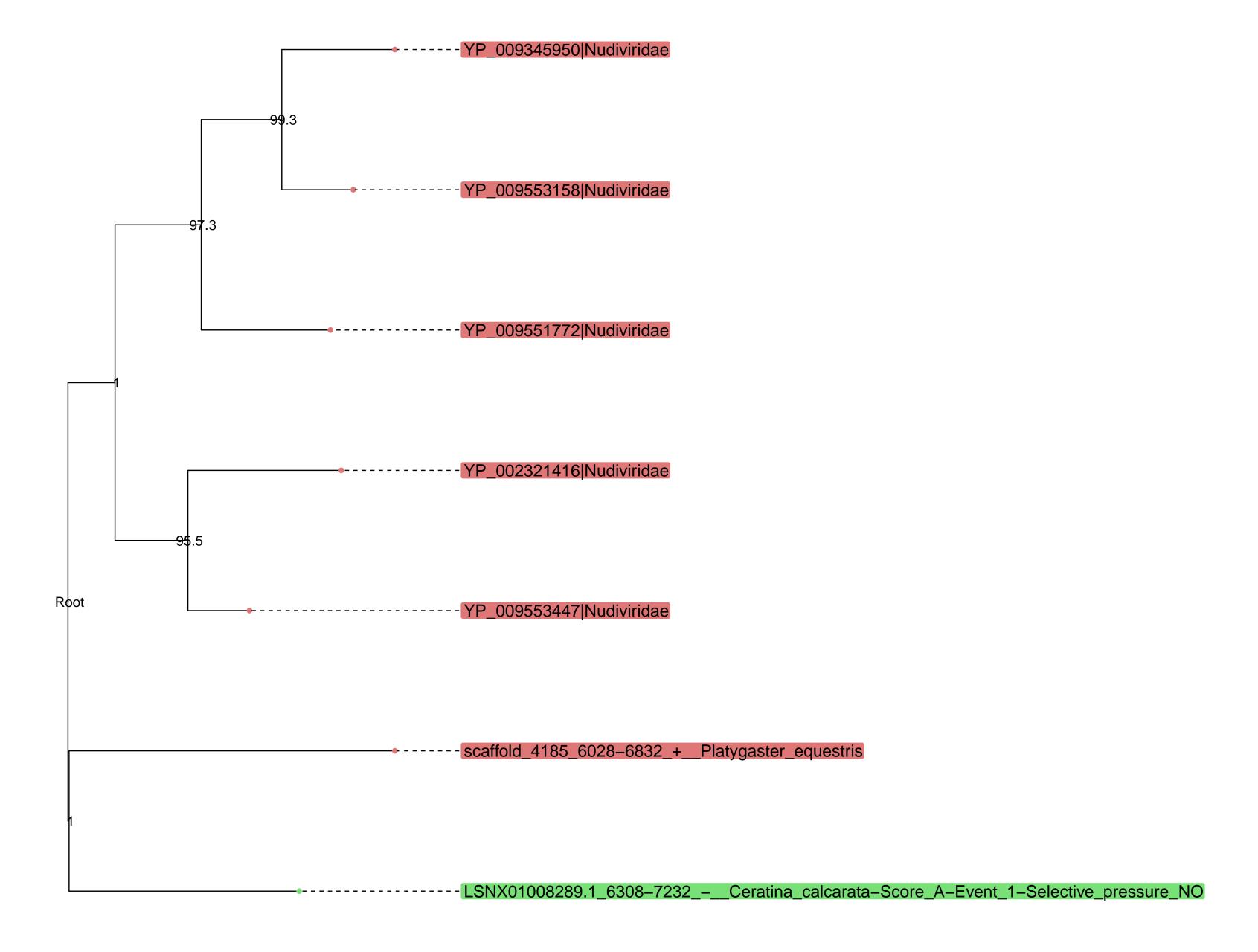
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Cluster23952 – families involved : ['Reoviridae'] – putative protein : ['RNA-directed RNA polymerase (EC 2.7.7.48)']
                                                                   AKV60527|Reoviridae
                                                                   AKV60521|Reoviridae
                                                                   AKV60519|Reoviridae
                                                                   AKV60453|Reoviridae
                                                                   AKV60481|Reoviridae
                                                                   AWL83791|Reoviridae
                                                                   AGJ83471|Reoviridae
                                                                   AYA21648|Reoviridae
                                                                   - AYA21662|Reoviridae
                                                                   AKV60439|Reoviridae
                                                                   AKP24203|Reoviridae
                                                                   ALU65925|Reoviridae
                                                                   ADI49541|Reoviridae
                                                                   AKV60493|Reoviridae
                                                                   AFN71110|Reoviridae
                                                                   ALX36925|Reoviridae
                                                                   ALW82954|Reoviridae
                                                                   AGJ83541|Reoviridae
                                                                   AKV60440|Reoviridae
                                                                   AKM21102|Reoviridae
                                                                   AAA87364|Reoviridae
                                                                   AAA87362|Reoviridae
                                                                   AKM21124|Reoviridae
                                                                   ATY47045|Reoviridae
                                                                   AOX23960|Reoviridae
                                                                   AOG59217|Reoviridae
                                                                   AGJ83481|Reoviridae
                                                                   ALU65935|Reoviridae
                                                                   AOX23970|Reoviridae
                                                                   AHM82384|Reoviridae
                                                                   AOX24020|Reoviridae
                                                                   AFM31250|Reoviridae
                                                                   AOX24060|Reoviridae
                                                                   ADI49561|Reoviridae
                                                                   AOX23930|Reoviridae
                                                                   ATY47042|Reoviridae
                                                                   ATY47021|Reoviridae
                                                                   ATY47046|Reoviridae
                                                                   AKV60550|Reoviridae
                                                                   AKV60447|Reoviridae
                                                                   ANK78265|Reoviridae
                                                                   ALI51254|Reoviridae
                                                                   AFI73116|Reoviridae
                                                                   AKV60463|Reoviridae
                                                                   AKV60444|Reoviridae
                                                                   AKP24150|Reoviridae
                                                                   AKP24140|Reoviridae
                                                                   ATY47722|Reoviridae
                                                                   ATY47749|Reoviridae
                                                                   ATY47766|Reoviridae
                                                                   AGM38080|Reoviridae
                                                                   ATY47711|Reoviridae
                                                                   ATY47712|Reoviridae
                                                                   AIT40287|Reoviridae
                                                                   ATY47720|Reoviridae
                                                                   ATY47717|Reoviridae
                                                                   ATY47724|Reoviridae
                                                                   ATY47760|Reoviridae
                                                                   ATY47762|Reoviridae
                                                                   AGJ83611|Reoviridae
                                                                   AFK91732|Reoviridae
                                                                   AIT40285|Reoviridae
     Root
                                                                   AIT40283|Reoviridae
                                                                   AFK91708|Reoviridae
                                                                   YP_009507687|Reoviridae
                                                                   ADU57374|Reoviridae
                                                                   ADU57376|Reoviridae
                                                                   AXB87376|Reoviridae
                                                                   ADU57375|Reoviridae
                                                                   YP_009158877|Reoviridae
                                                                   AGZ92011|Reoviridae
                                                                    AGZ92013|Reoviridae
                                                                   AIV43180|Reoviridae
                        68.4
                                                                   AGZ92010|Reoviridae
                                                                   AMQ36817|Reoviridae
                                                                   APC23688|Reoviridae
                                                                   scf7180005162336_3490-5239_-_Leptopilina_clavipes-Score_A-Event_1-Selective_pressure_NO
                                                                         phylum a Virus a EVE
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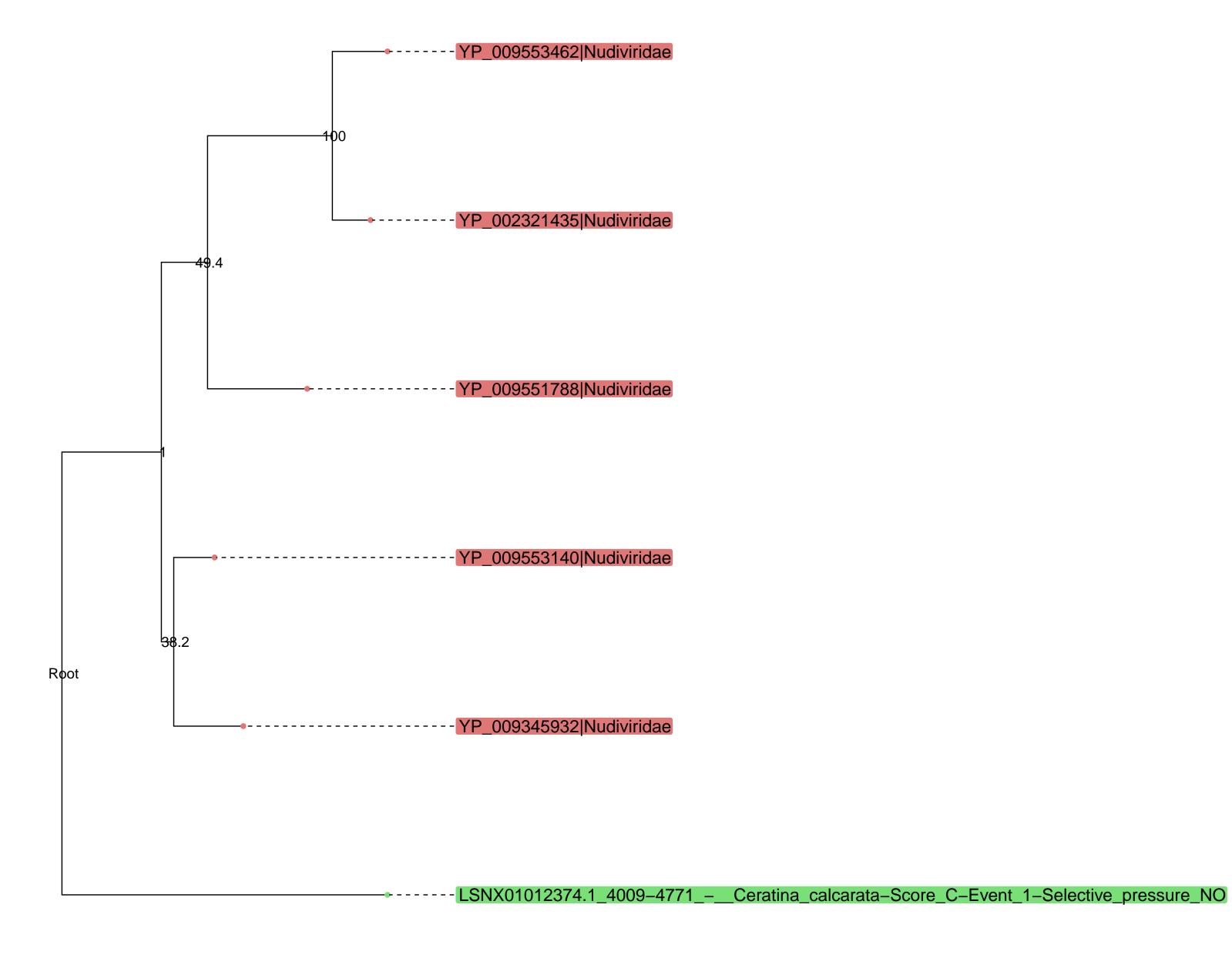


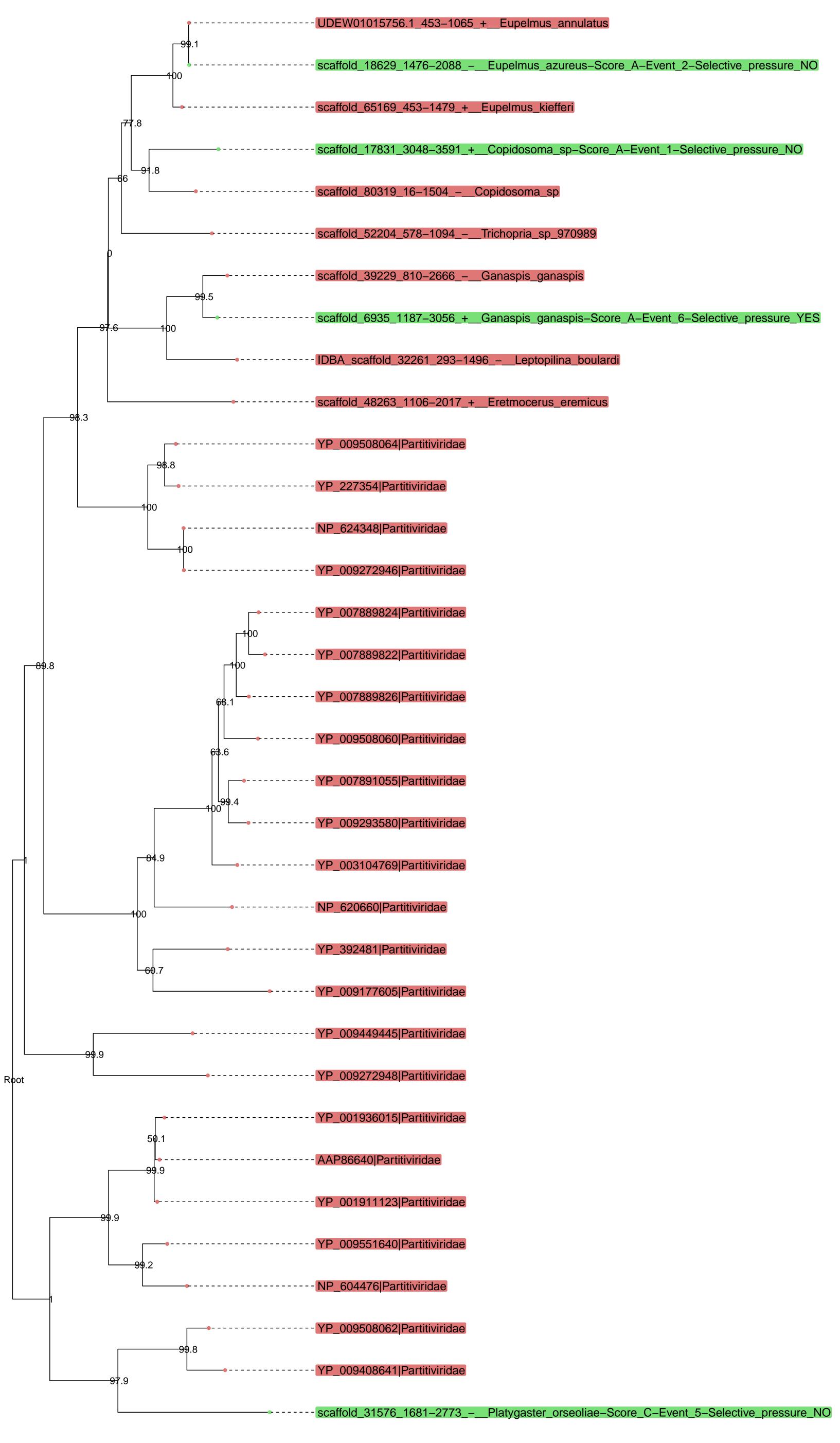


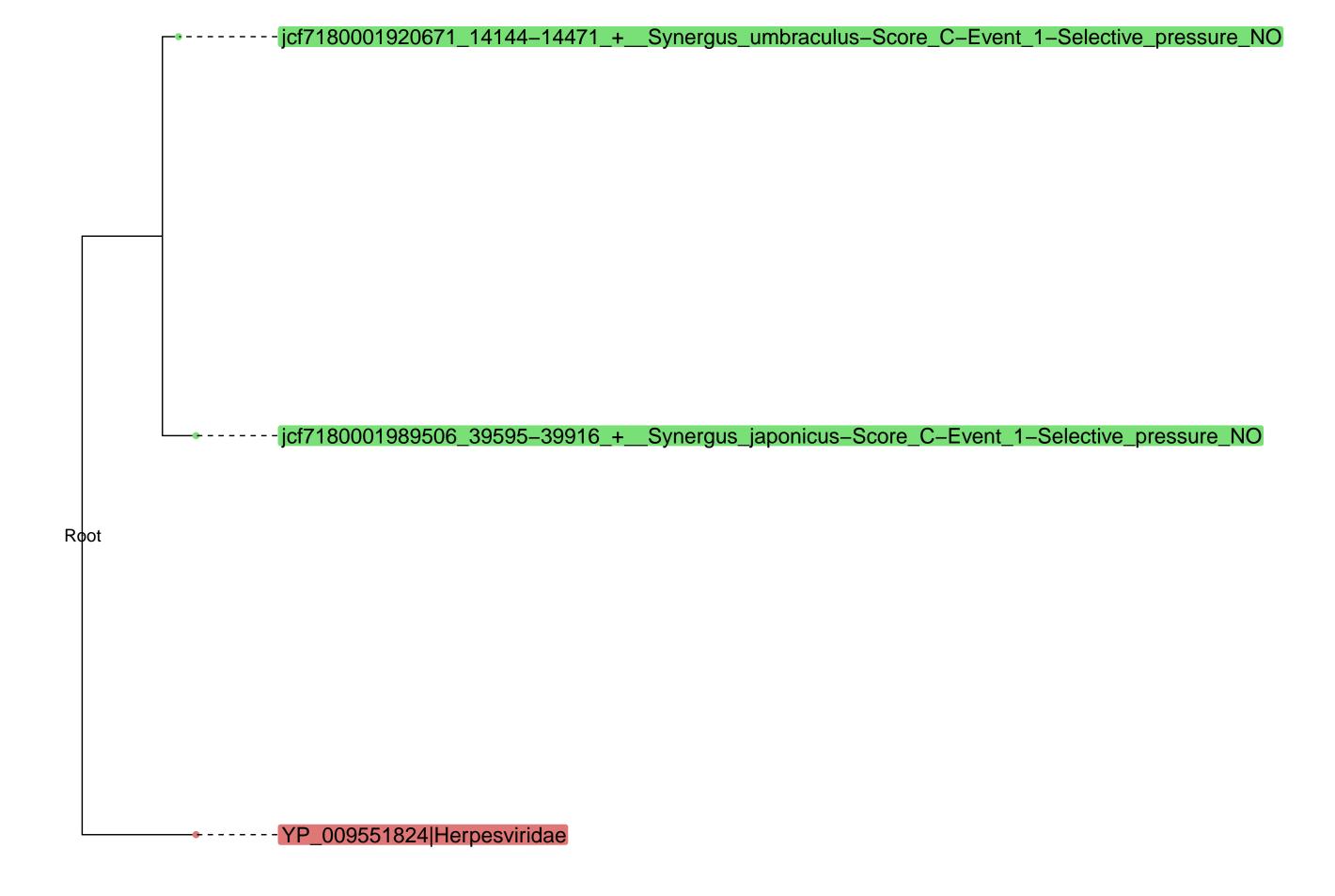


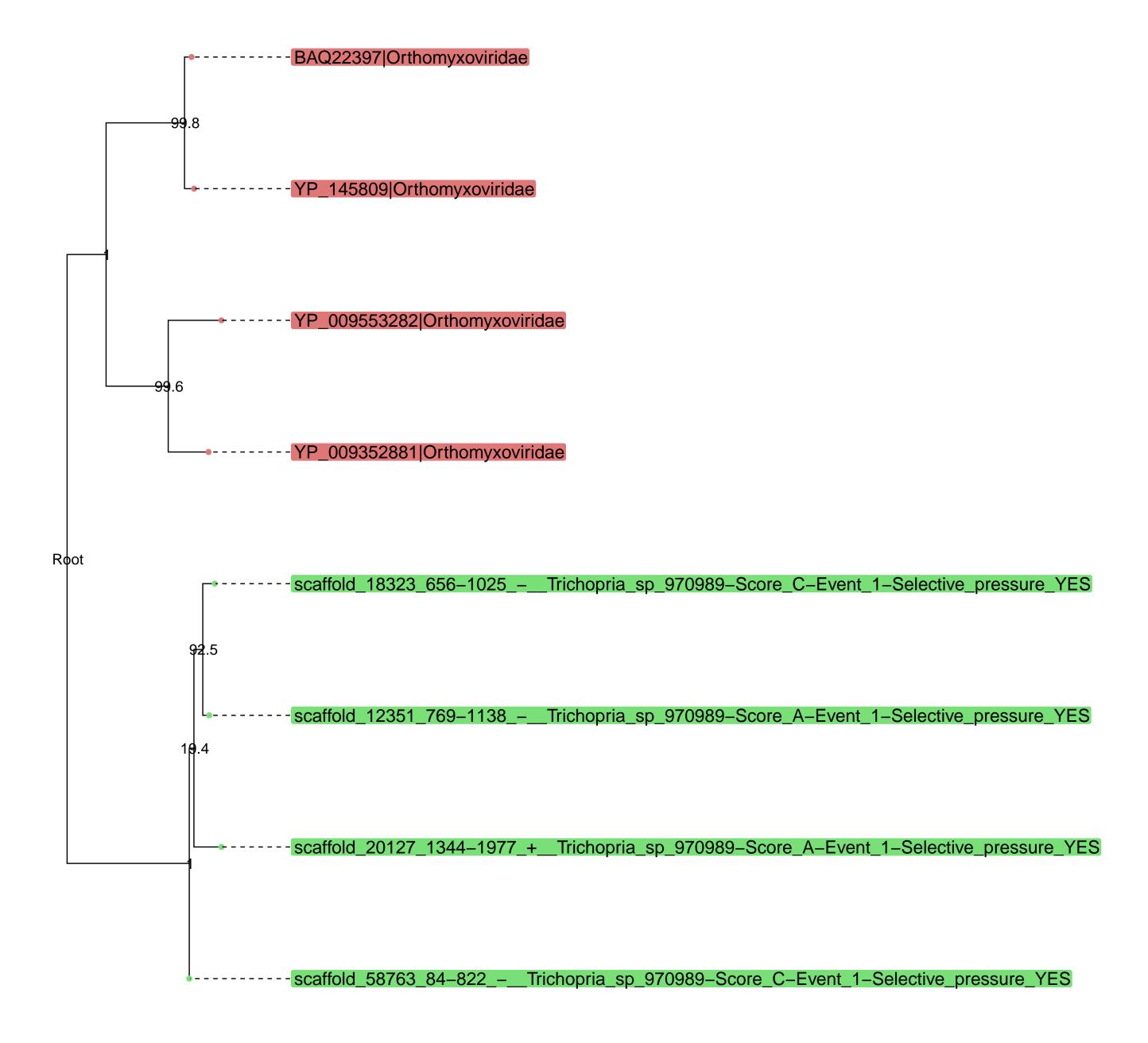


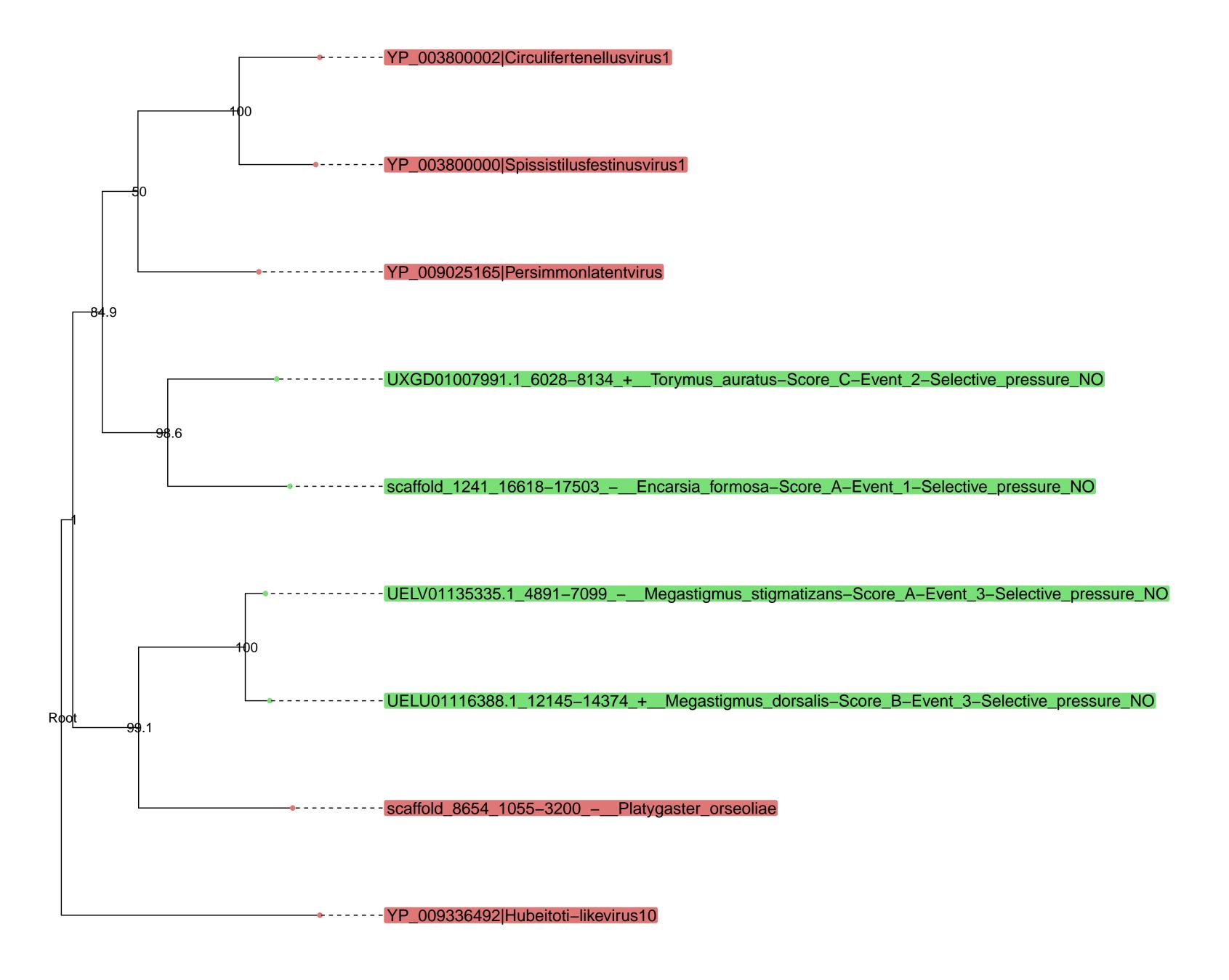


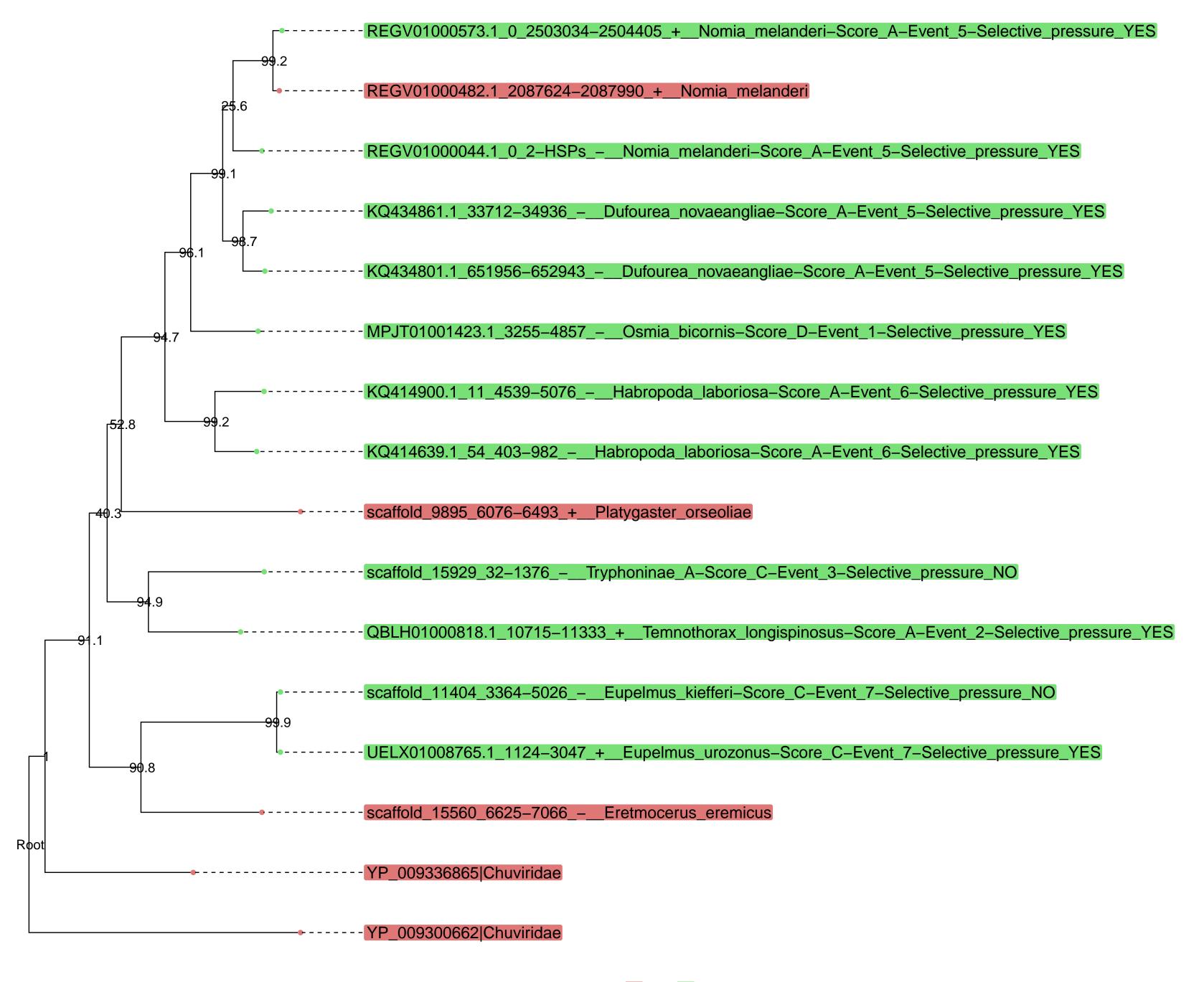


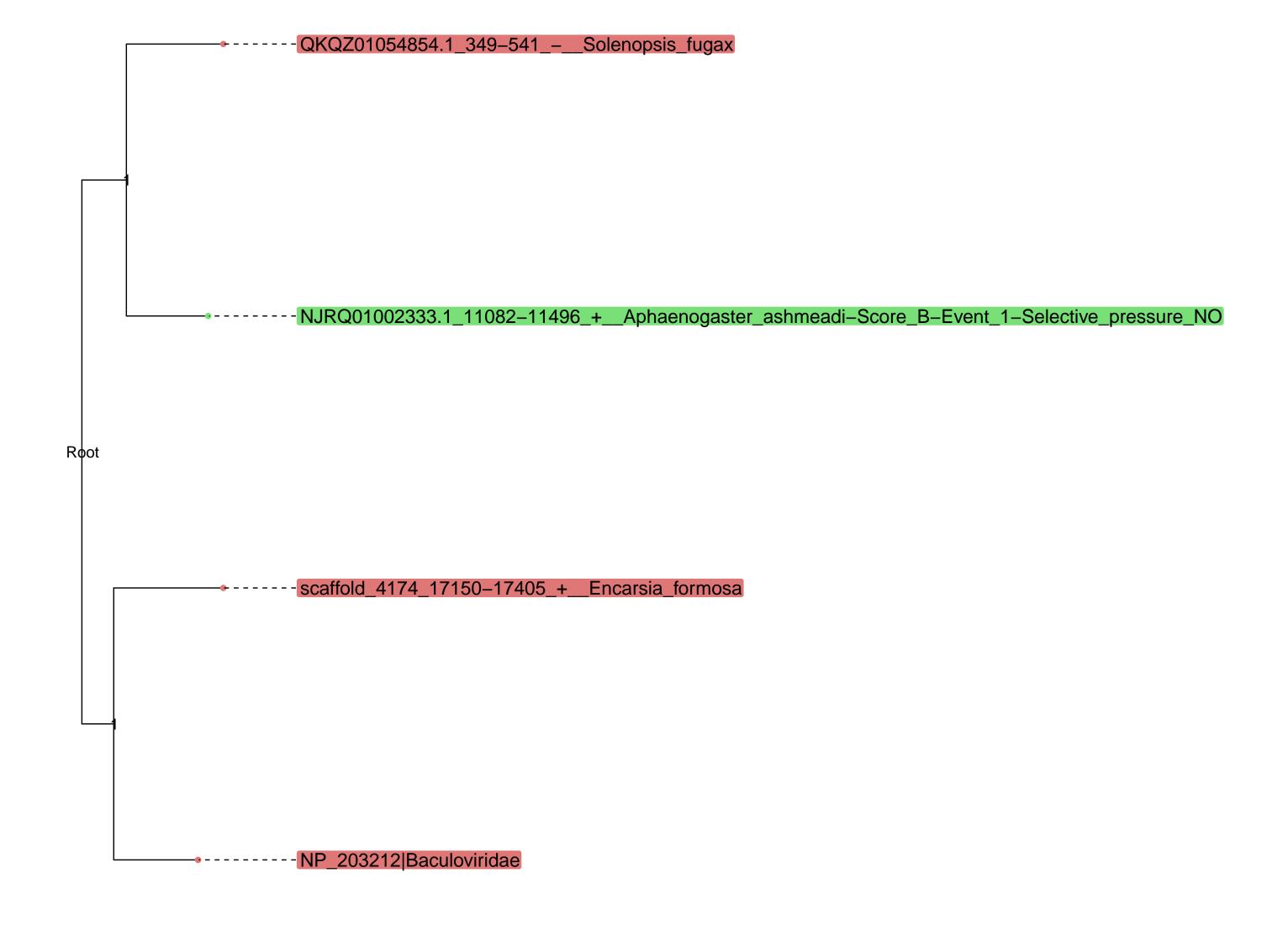


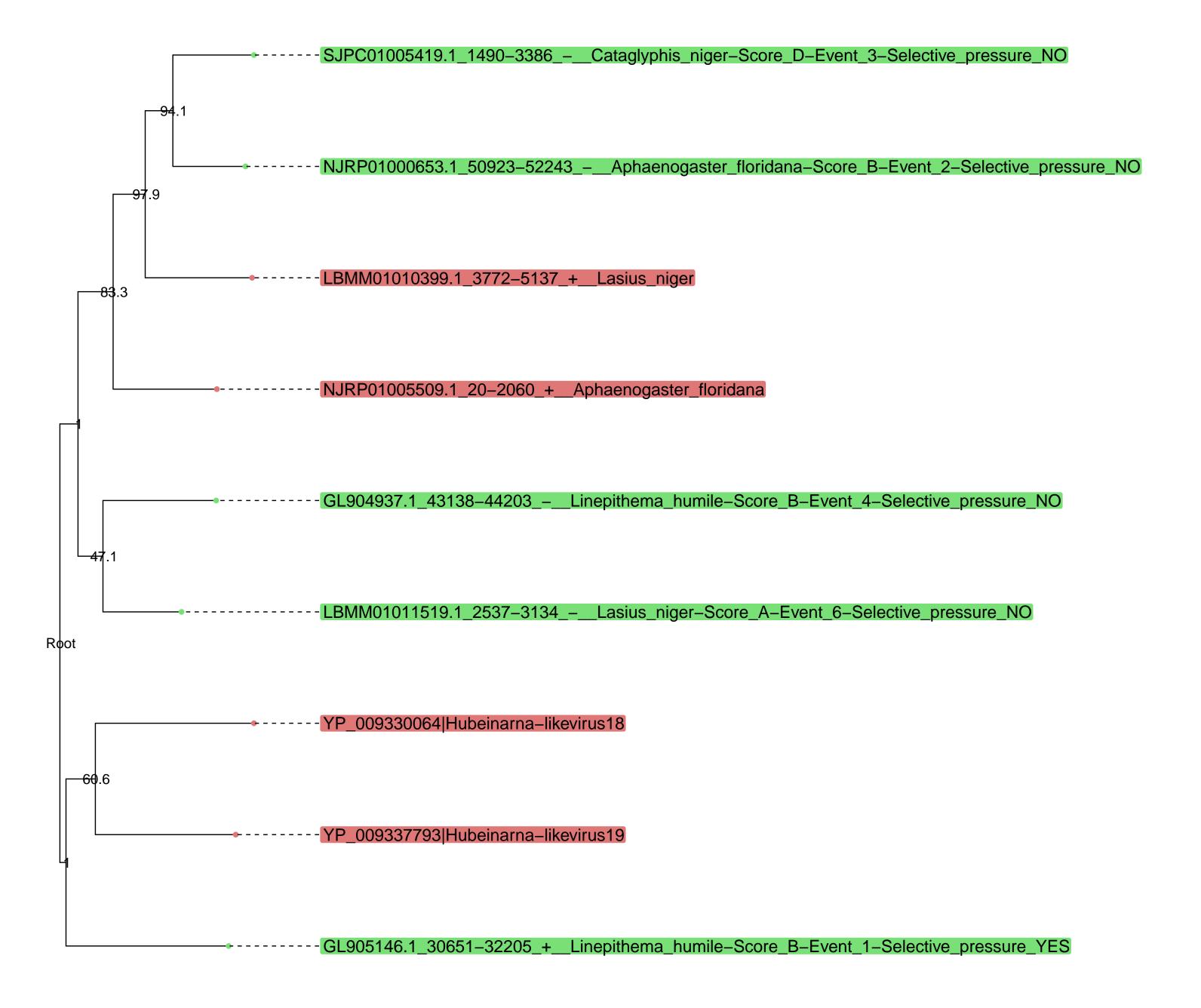


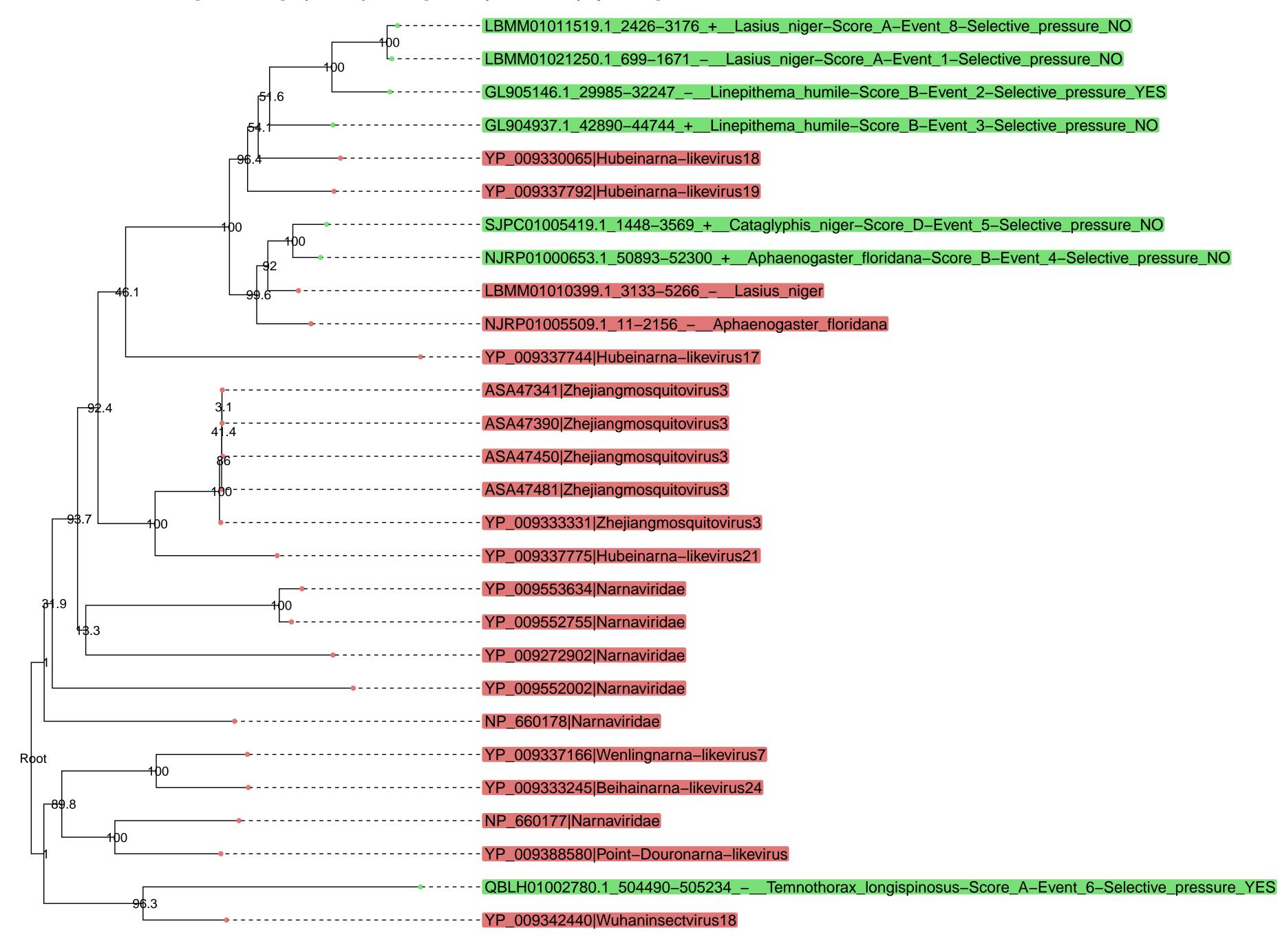


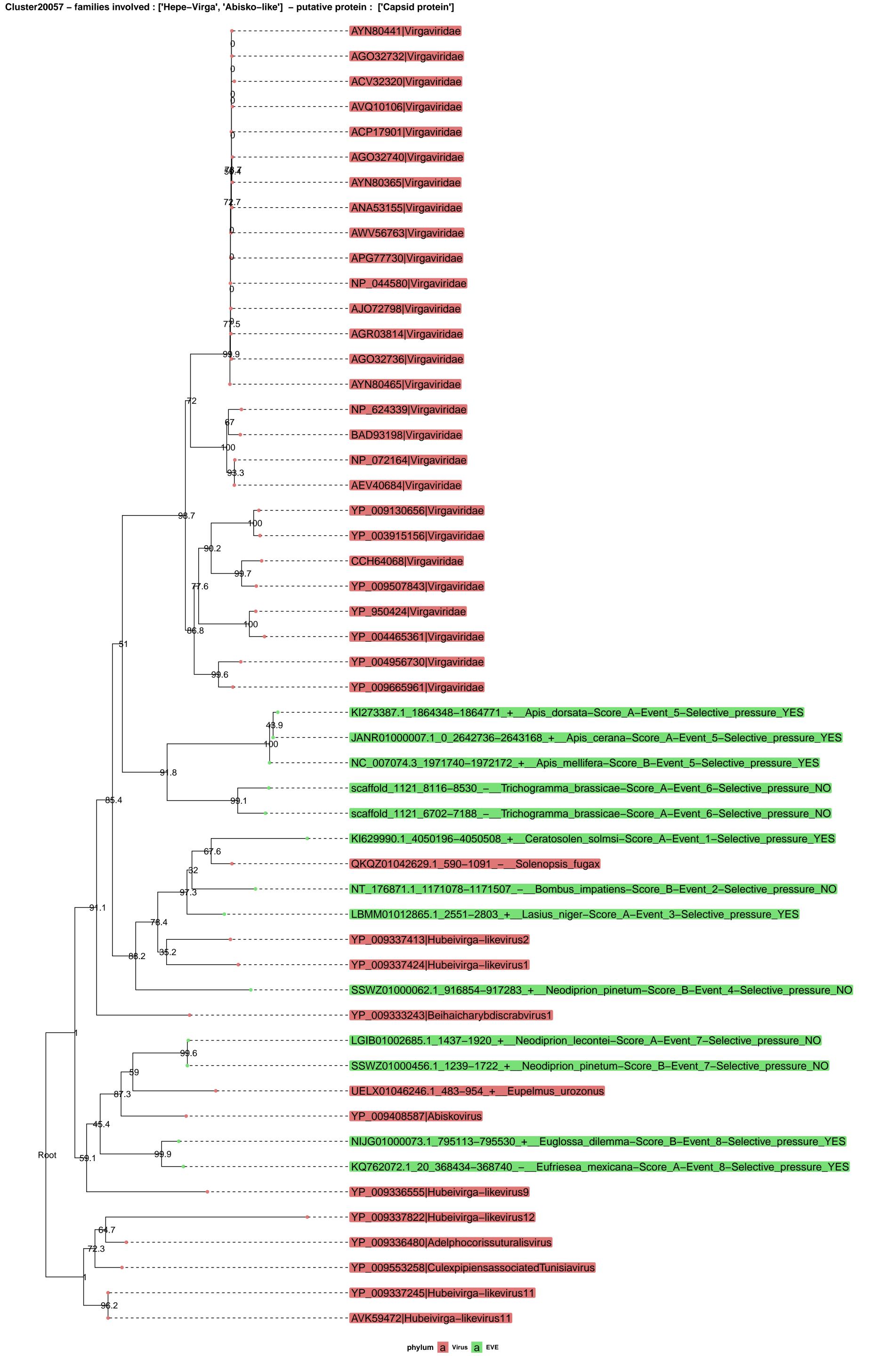


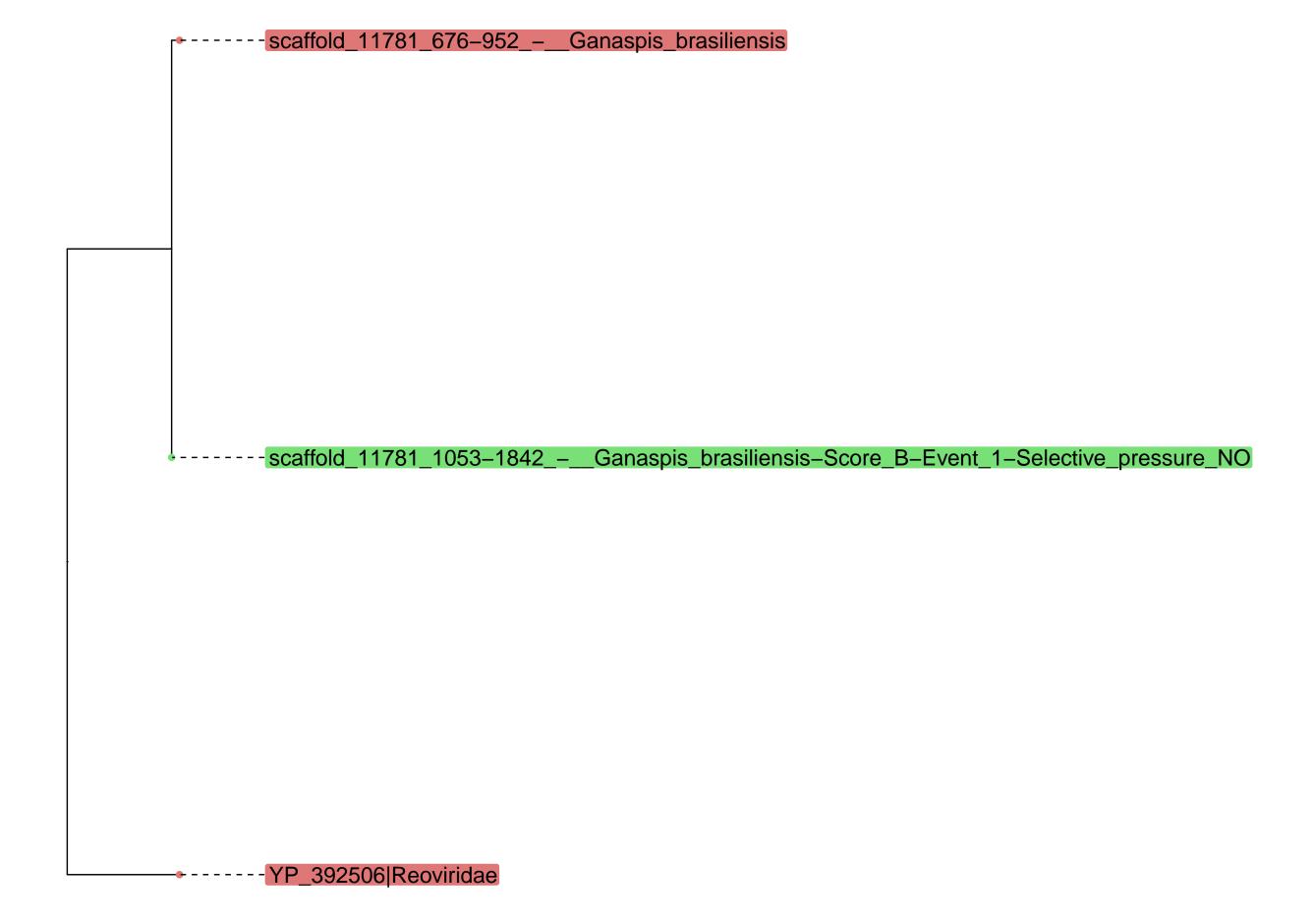


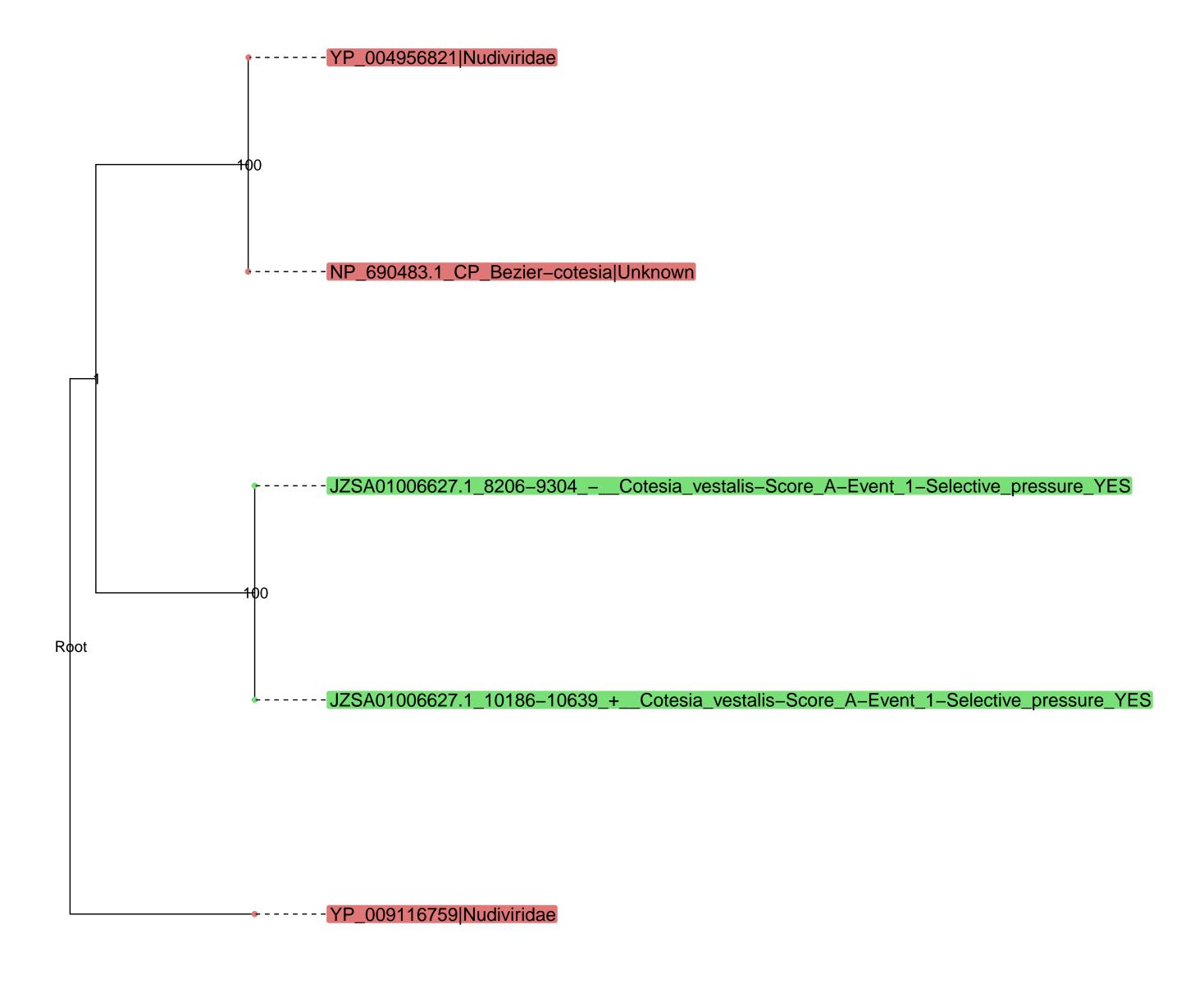


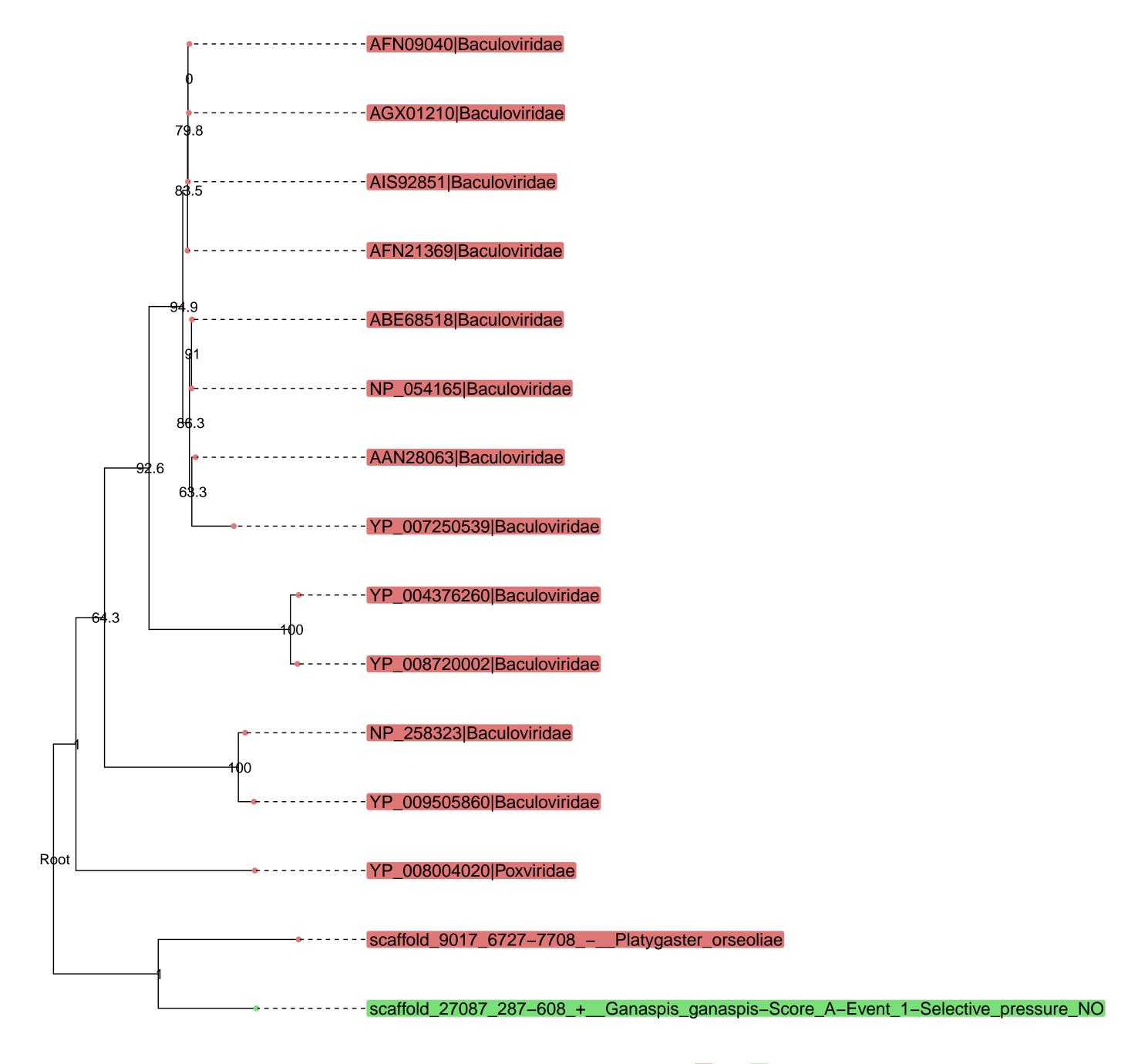


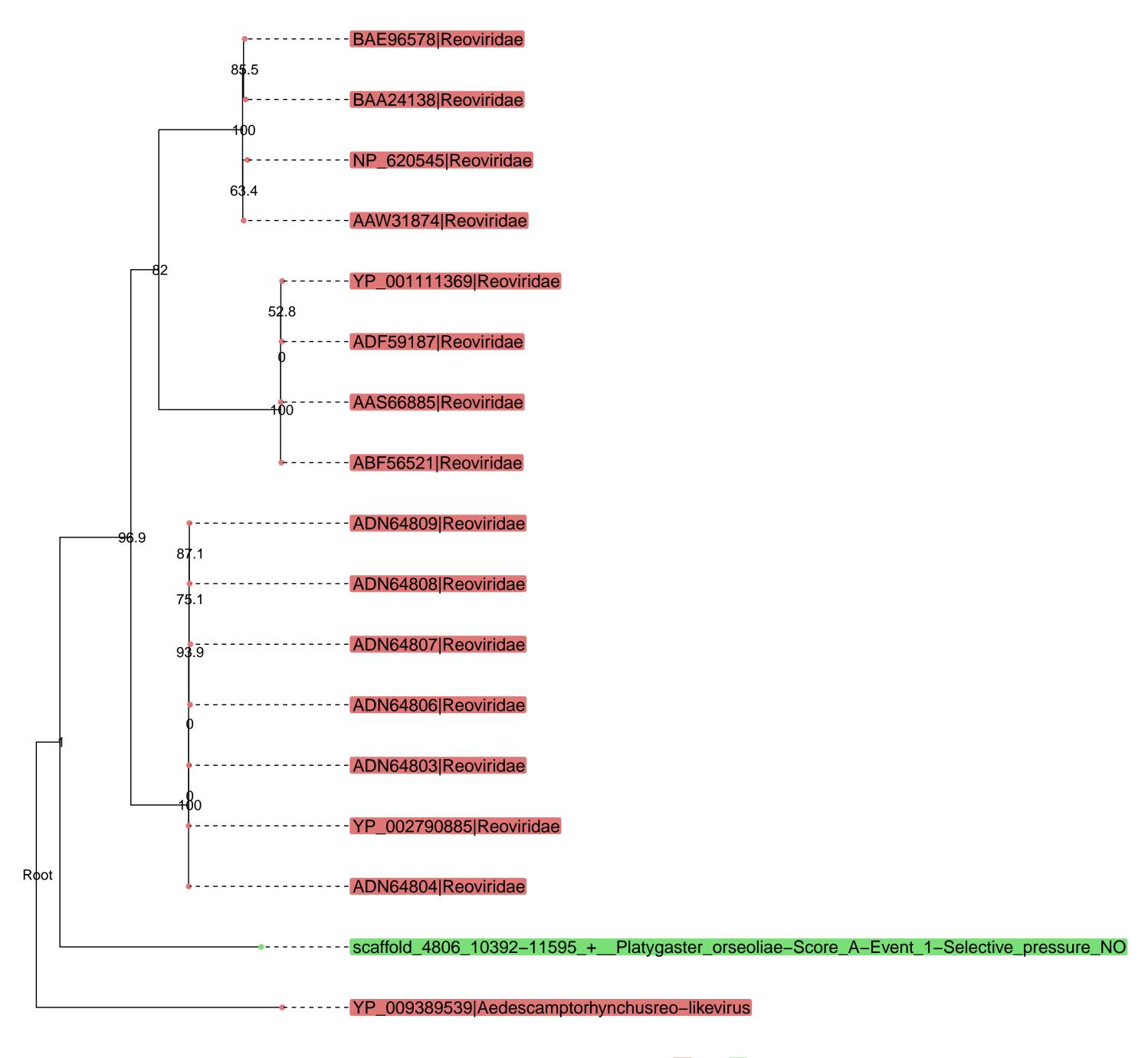




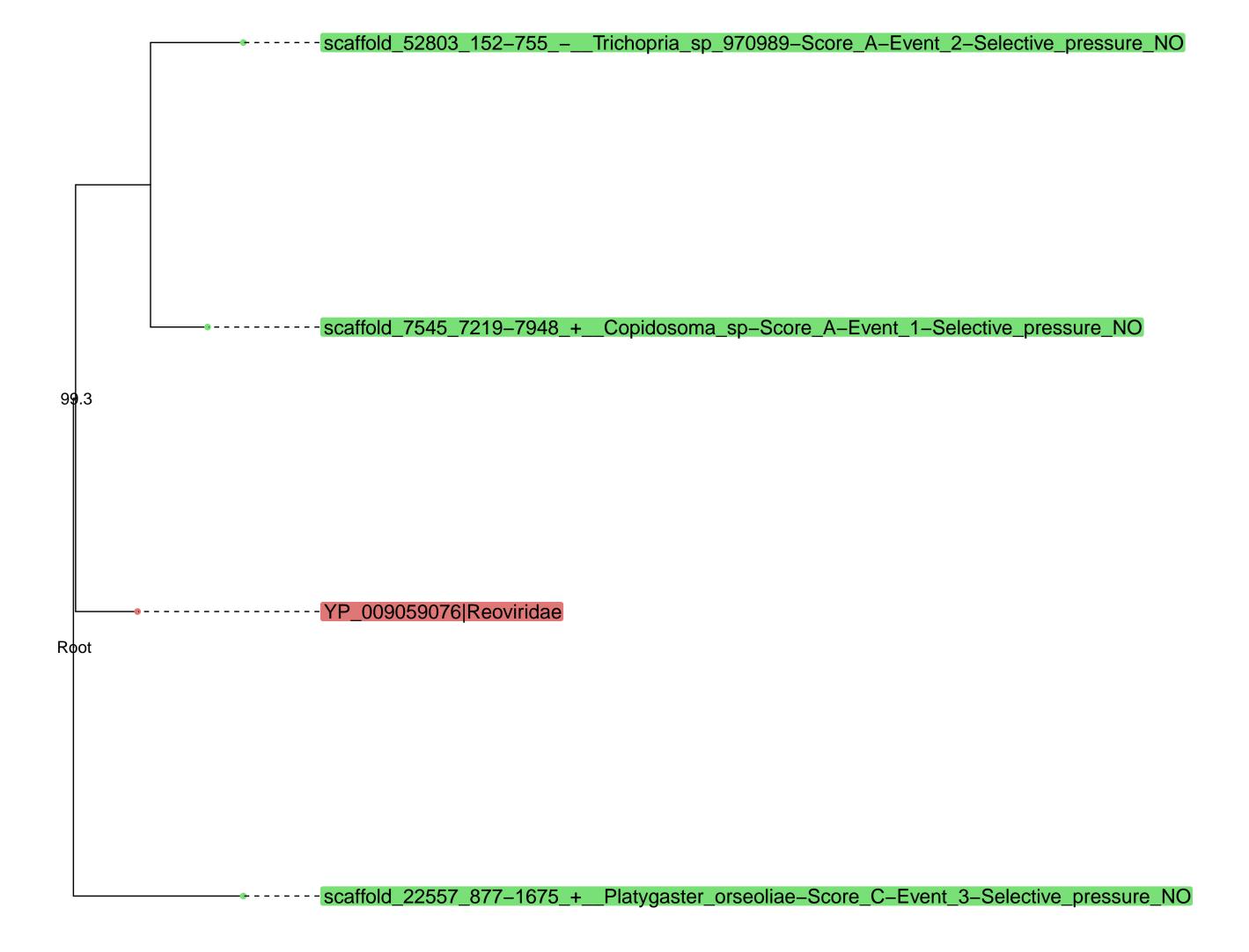


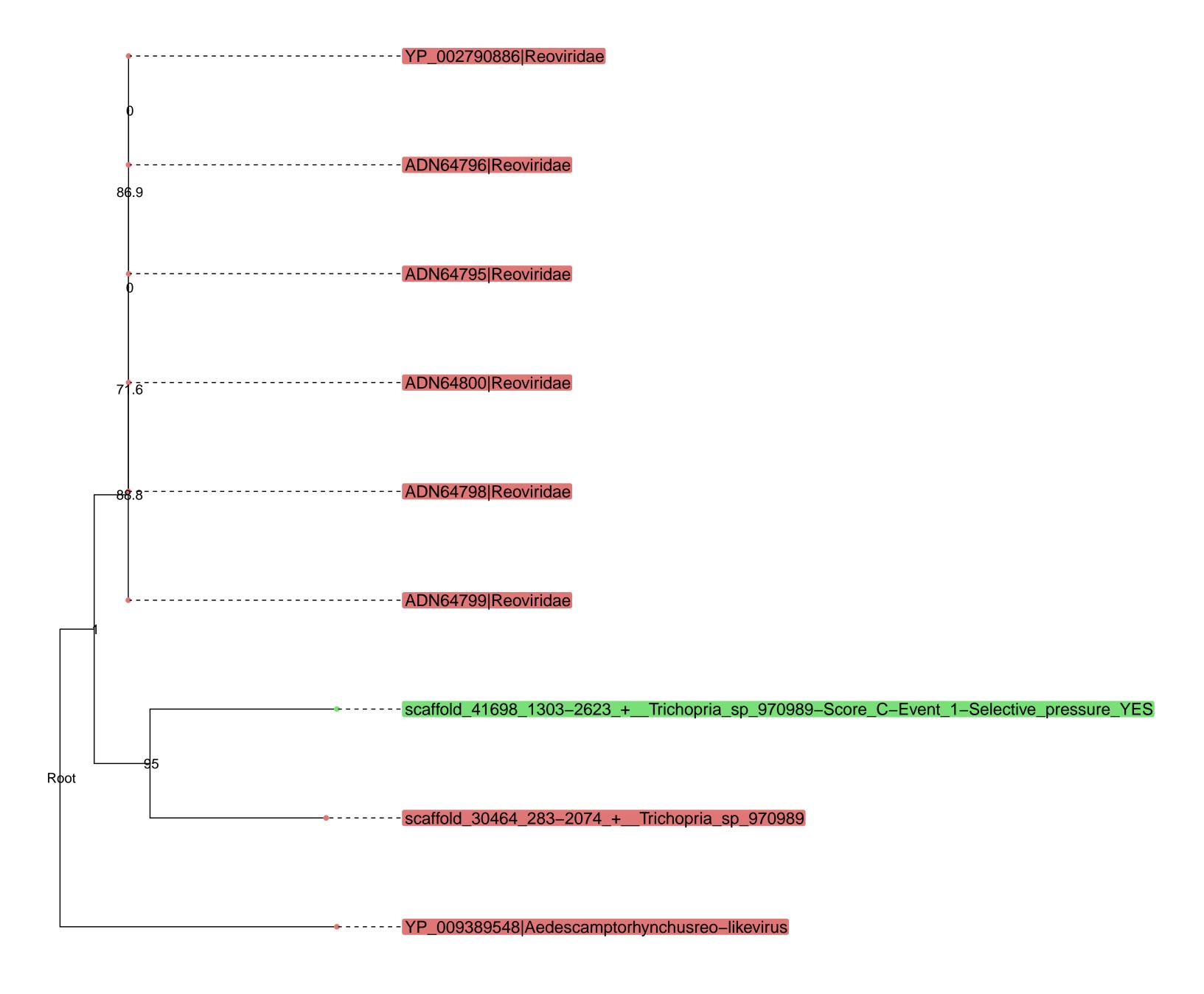


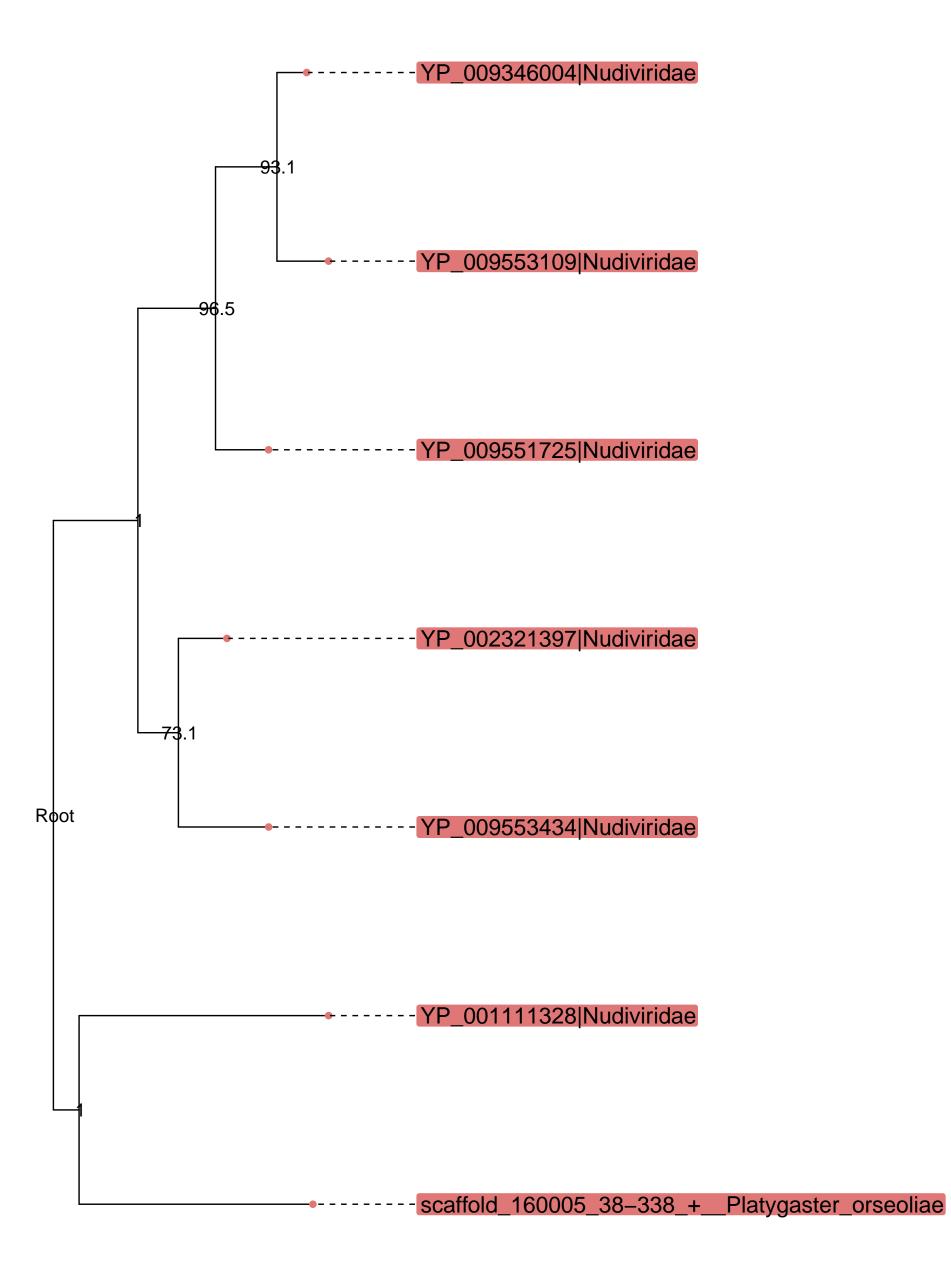


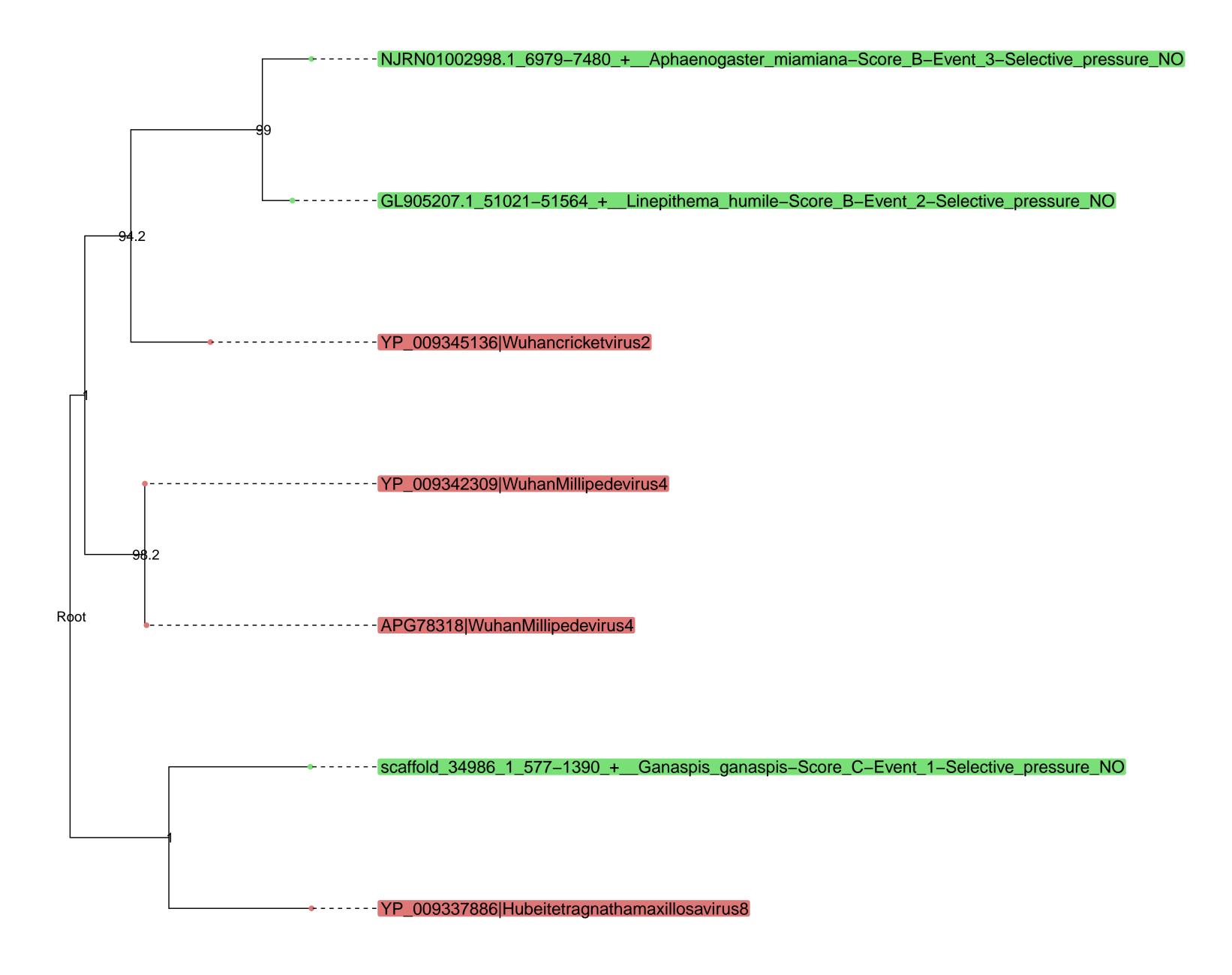


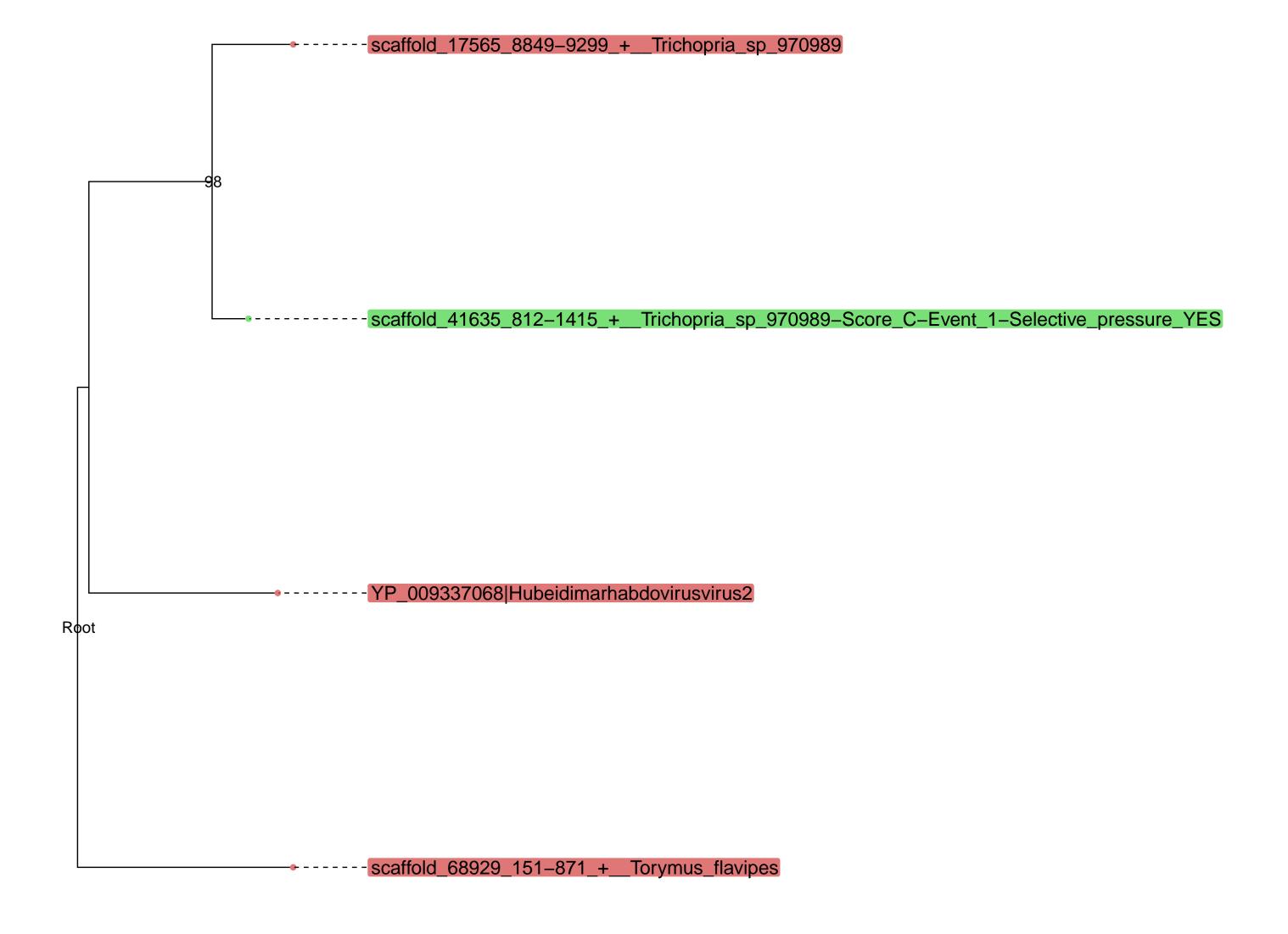


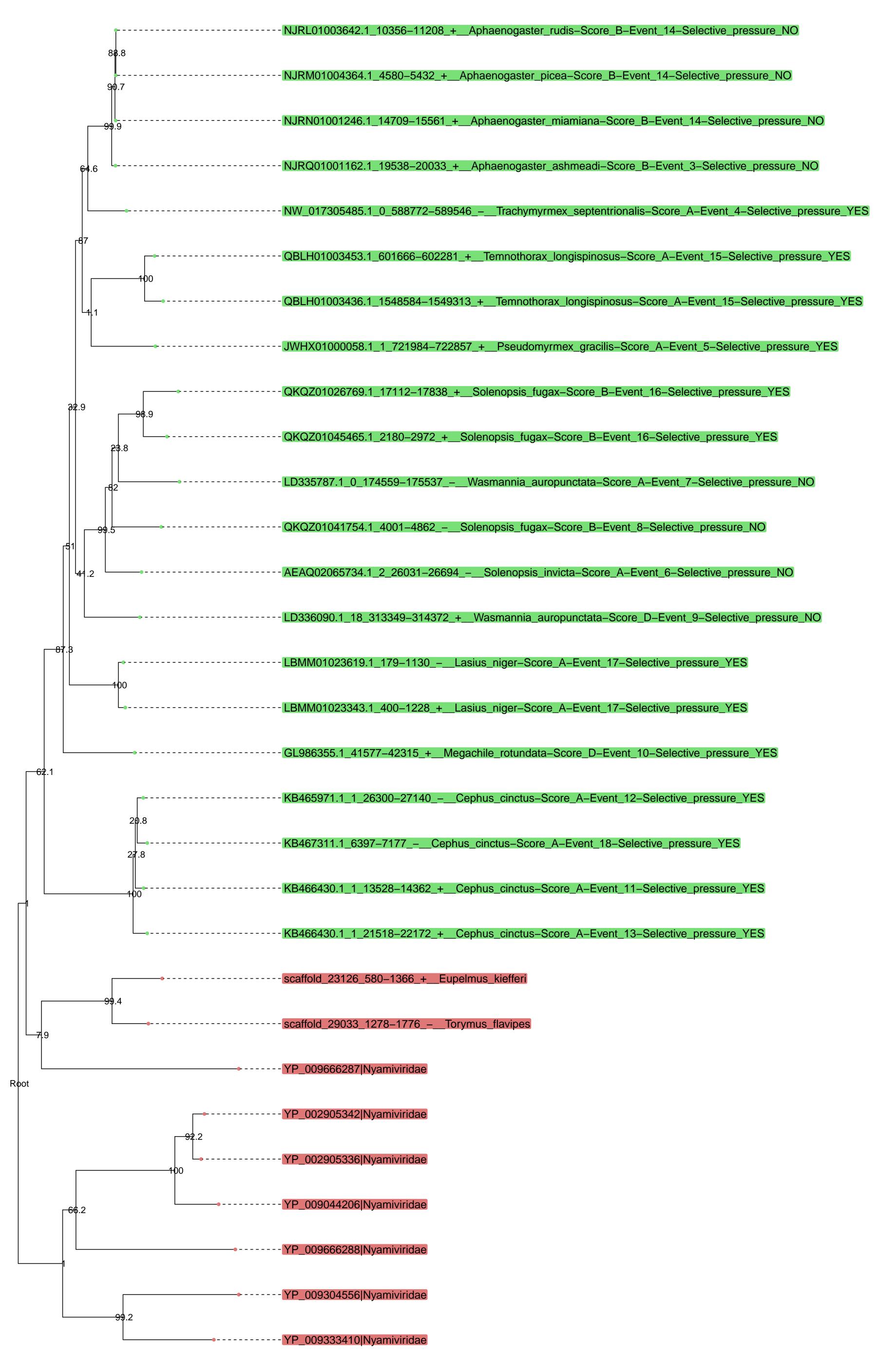


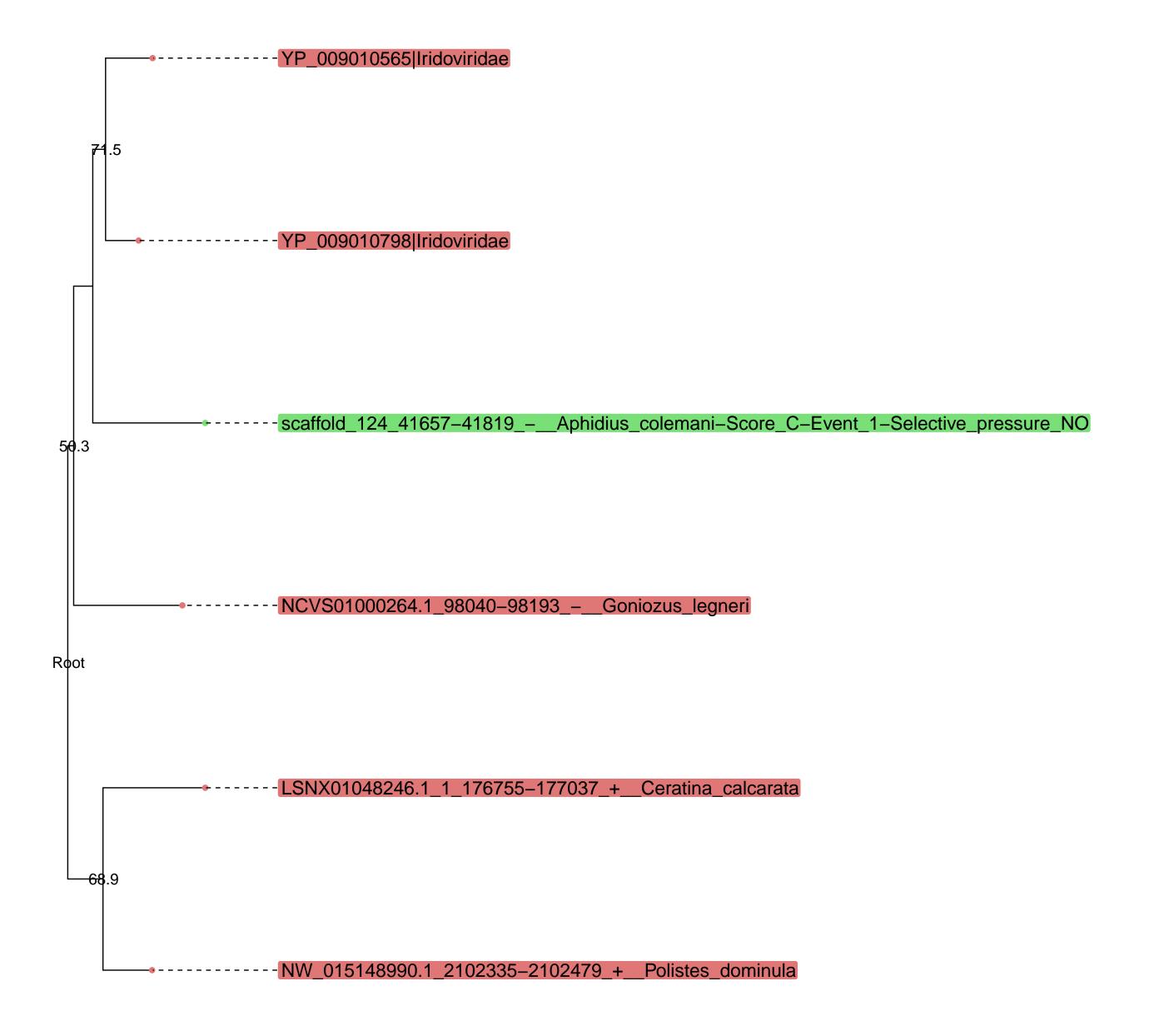




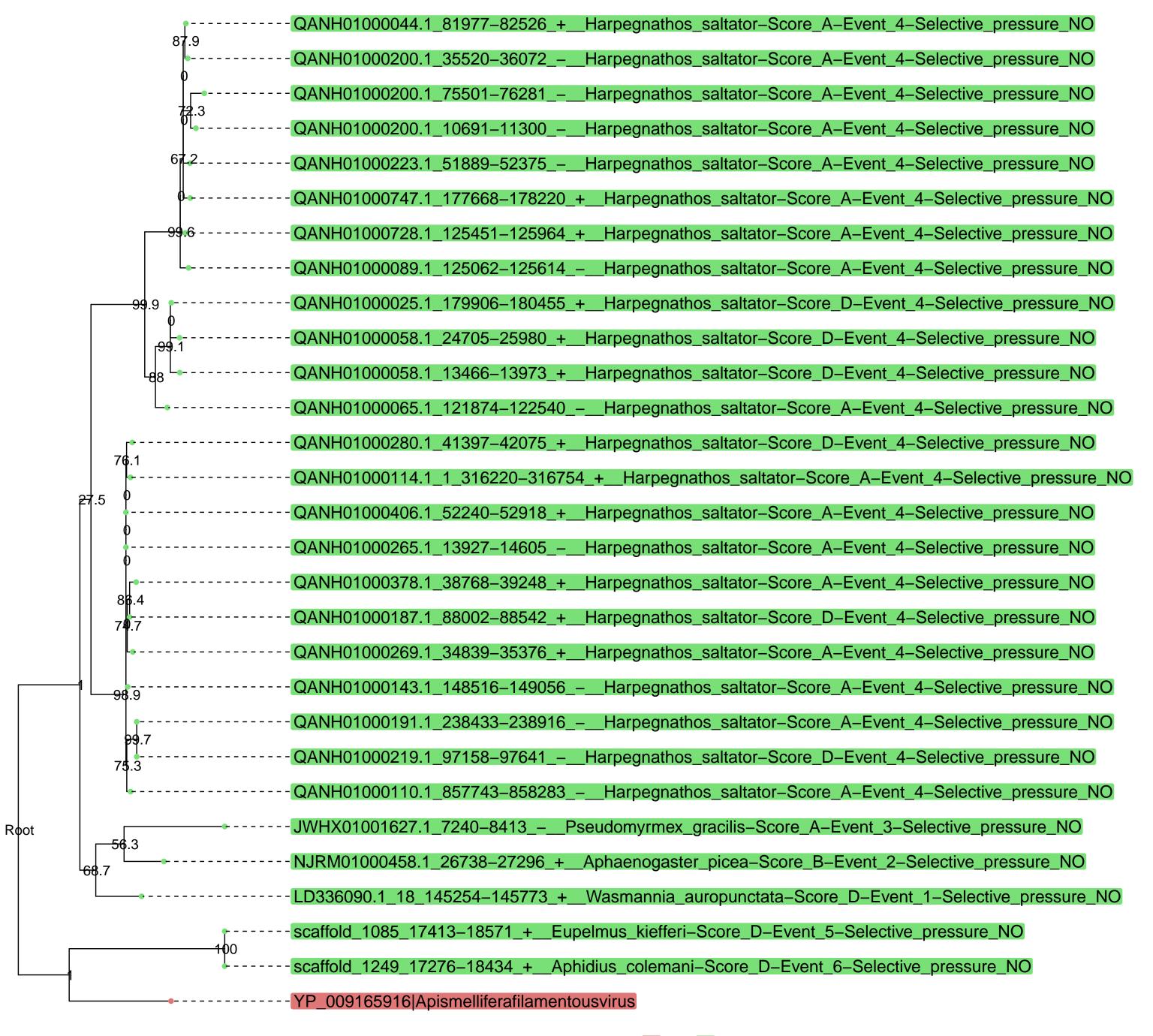




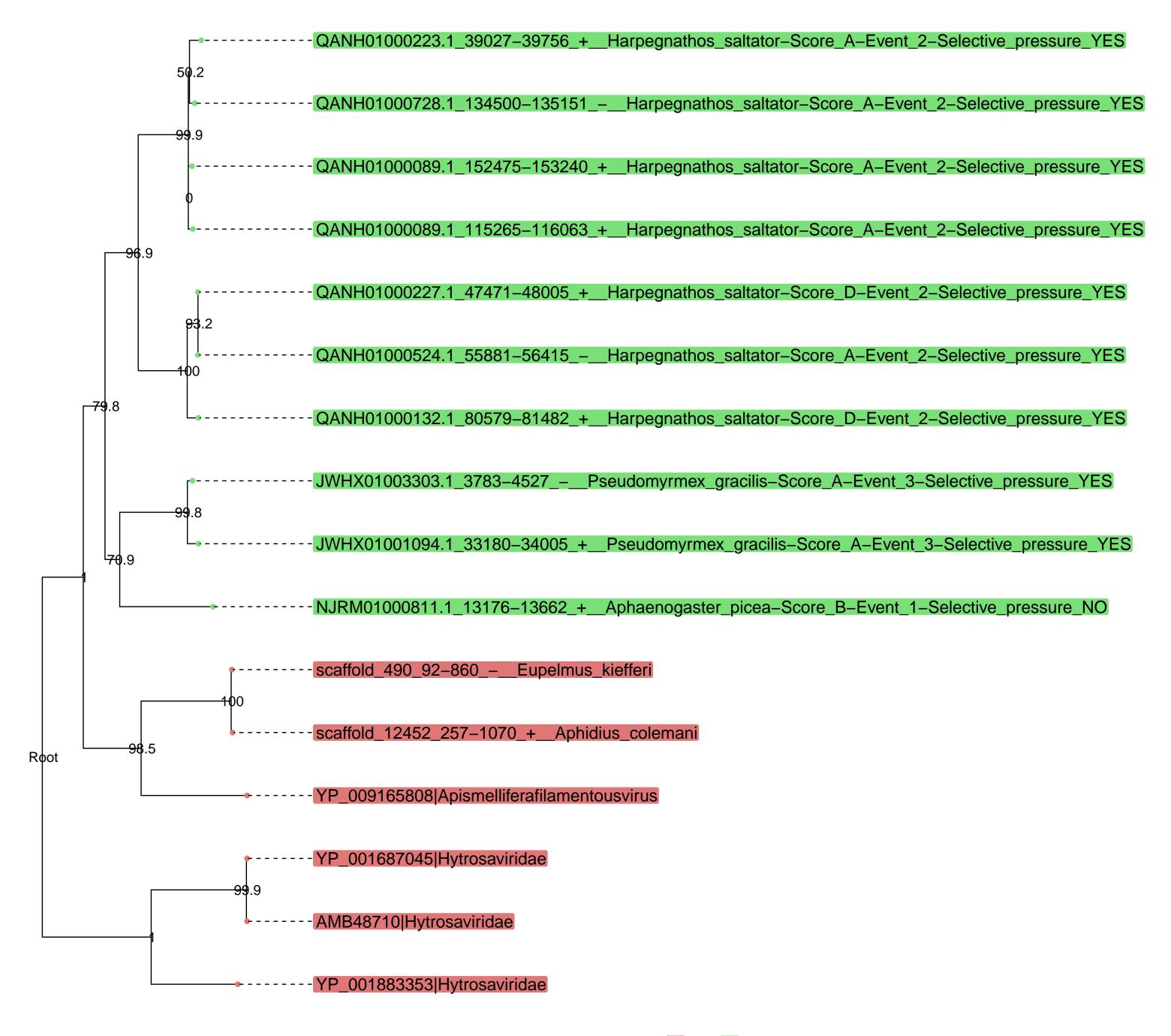


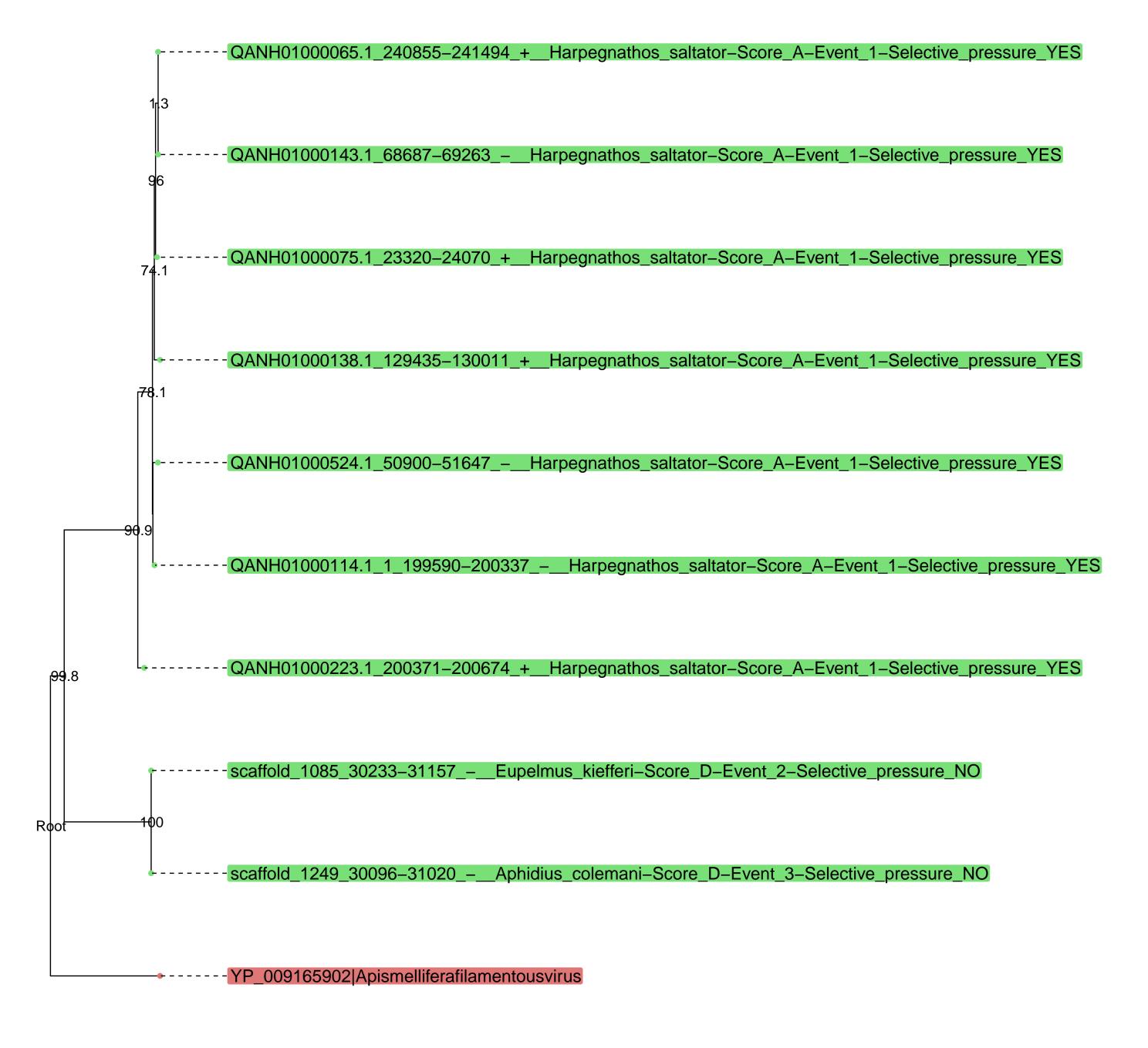


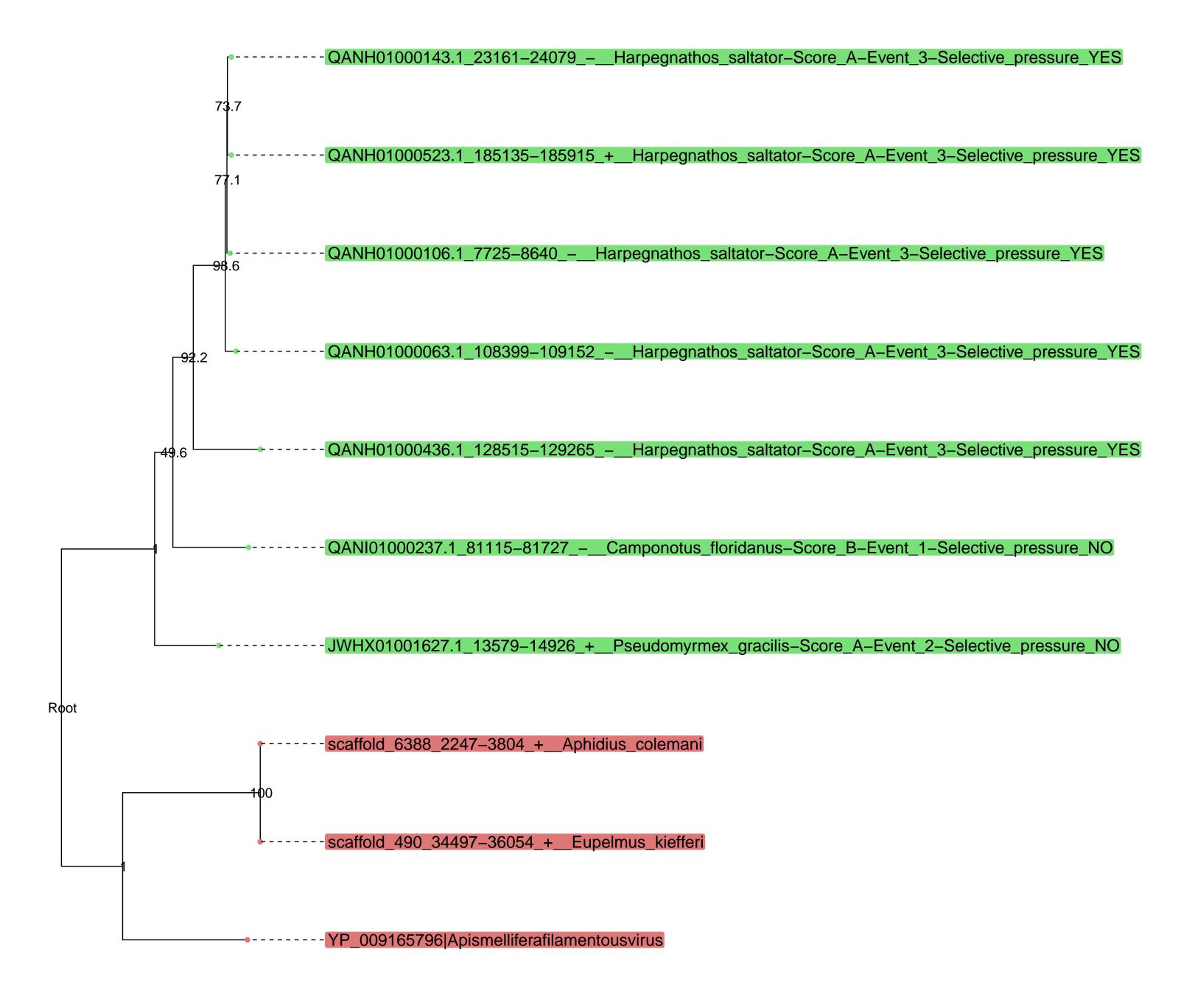
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Cluster17775 – families involved : ['Baculoviridae'] – putative protein : ['p26']
                                                      CDG73170|Baculoviridae
                                                      CDG73033|Baculoviridae
                                                      CDG72881|Baculoviridae
                                                      NP_037889|Baculoviridae
                                                      YP_002332831|Baculoviridae
                                                      - ADV91363|Baculoviridae
                                                      YP_001036423|Baculoviridae
                                                      ACA02687|Baculoviridae
                                                      AYN45077|Baculoviridae
                                                       YP_002268179|Baculoviridae
                                                      YP_009112700|Baculoviridae
                                                      YP_529812|Baculoviridae
                                                      AFP95871|Baculoviridae
                                                      NP_613241|Baculoviridae
                                                      ACH88673|Baculoviridae
                                                      NP_689331|Baculoviridae
                                                      YP_009011212|Baculoviridae
                                                      YP_009049954|Baculoviridae
                                                      YP_009666799|Baculoviridae
                                                      YP_009116933|Baculoviridae
             <del>- 97</del>.6
                                                      AOL57023|Baculoviridae
                                                      AOL56739|Baculoviridae
                                                      QBI90242|Baculoviridae
                                                      YP_308908|Baculoviridae
                                                      YP_249623|Baculoviridae
                                                      QED40546|Baculoviridae
                                                      YP_717557|Baculoviridae
                                                      AKN90978|Baculoviridae
                                                      YP_009001785|Baculoviridae
                                                      -YP_009186701|Baculoviridae
                                                     -YP_009552165|Baculoviridae
               β9.9
                                                      YP_006607871|Baculoviridae
                                                      YP_001650930|Baculoviridae
                                                      YP_002854661|Baculoviridae
                                                      YP_009165631|Baculoviridae
                                                      YP_008378498|Baculoviridae
                                                      NP_848319|Baculoviridae
                                                     - AGR57032|Baculoviridae
               <del>-99</del>.6
                                                      NP_818678|Baculoviridae
                                                      YP_002300548|Baculoviridae
                                                      AJP07579|Baculoviridae
        <del>83</del>.7
                                                      - AJP07309|Baculoviridae
                                                      AJP07715|Baculoviridae
                                                     - AJP07174|Baculoviridae
                                                      NP_075091|Baculoviridae
                                                      QBM78987|Baculoviridae
                                                      AMN15404|Baculoviridae
                                                      AIG63200|Baculoviridae
         <del>38</del>.4
                                                      ALD88584|Baculoviridae
                                                      NW_014463936.1_216974-217865_+__Microplitis_demolitor-Score_A-Event_1-Selective_pressure_YES
                                                     -YP_758317|Baculoviridae
                                                      - AXU41465|Baculoviridae
                                                      AIS92852|Baculoviridae
                                                      ACQ57310|Baculoviridae
                                                      - AGX01339|Baculoviridae
                                                     -NP_047534|Baculoviridae
                                                     - AFN21230|Baculoviridae
                                                     - AFN09041|Baculoviridae
      Root
                                                      AFN21092|Baculoviridae
                                                      ABE68519|Baculoviridae
                                                     NP_054166|Baculoviridae
                                                      - AAN28082|Baculoviridae
                                                     -YP_007250540|Baculoviridae
                                                     -YP_009316042|Baculoviridae
                                                      - AXE72060|Baculoviridae
                           <del>-91</del>.8
                                                      YP_803526|Baculoviridae
                                                      YP_610991|Baculoviridae
                                                     - NP_932739|Baculoviridae
                                                     -YP_473209|Baculoviridae
                                                     - NP_203288|Baculoviridae
                                                     - YP_009666547|Baculoviridae
                                                      YP_009250006|Baculoviridae
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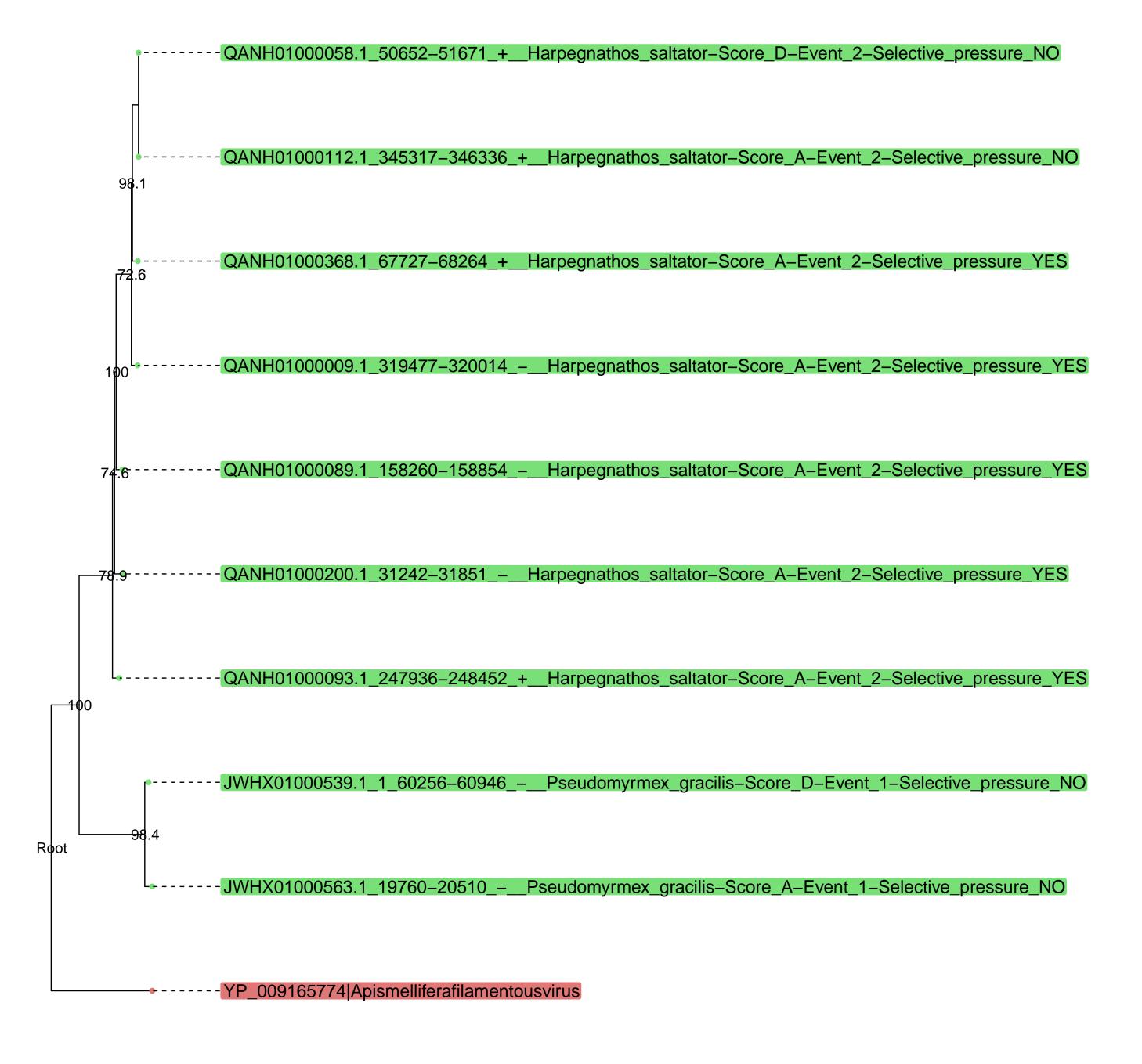


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Cluster3481 – families involved: ['Apis_filamentous-like'] – putative protein: ['Virulence-associated protein E']
                                  QANH01000164.1_550271-550601_-_Harpegnathos_saltator
                                  QANH01000164.1 530657–530972 – Harpegnathos saltator
                                  QANH01000328.1_69710-70253_+__Harpegnathos_saltator
                                 QANH01000755.1_70434-71220_+__Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                  QANH01000526.1_126107-126722_-_Harpegnathos_saltator
                                 QANH01000056.1_25695-26052_+__Harpegnathos_saltator
                                 QANH01000747.1_1535-2228_+__Harpegnathos_saltator
                                 QANH01000580.1_2405-3278_+__Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                 QANH01000254.1_35318-36140_+__Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                 -QANH01000254.1_22880-23702_+__Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                 QANH01000114.1_1_159437-159875_+__Harpegnathos_saltator
                                 QANH01000330.1_430-1093_+__Harpegnathos_saltator
                                  QANH01000621.1_50441-51269_-_Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                  QANH01000621.1_1408-1963_+__Harpegnathos_saltator
                                 QANH01000782.1_128823-129591_+__Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                  QANH01000283.1_77317-78115_+__Harpegnathos_saltator-Score_D-Event_6-Selective_pressure_NO
                                  QANH01000100.1_57761-58301_-__Harpegnathos_saltator
                                  QANH01000178.1_2-HSPs_-_Harpegnathos_saltator
                                 QANH01000283.1_28696-29446_+__Harpegnathos_saltator-Score_D-Event_6-Selective_pressure_NO
                                  QANH01000051.1_50108-50588_-__Harpegnathos_saltator
                                 QANH01000257.1_34366-34930_-__Harpegnathos_saltator
                                  QANH01000526.1_233310-233703_+__Harpegnathos_saltator
                                  QANH01000001.1_631285-631864_+__Harpegnathos_saltator
                                  QANH01000737.1_44589-45435_+__Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                  QANH01000117.1_2-HSPs_-_Harpegnathos_saltator
                                  QANH01000728.1_66849-67653_-_Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                  QANH01000258.1_87195-87711_-_Harpegnathos_saltator
                                 QANH01000142.1_70083-70887_-__Harpegnathos_saltator
                                 QANH01000820.1_8769-9483_-__Harpegnathos_saltator
                                 -QANH01000030.1_209058-209502_-_Harpegnathos_saltator
                                  QANH01000219.1_49347-50139_-_Harpegnathos_saltator-Score_D-Event_6-Selective_pressure_NO
                                  QANH01000686.1_10402-11203_-__Harpegnathos_saltator
                    99.8
80.5_
                                  QANH01000169.1_713935-714739_-__Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                  QANH01000443.1_32924-33614_+__Harpegnathos_saltator
                    75.2
                                  QANH01000015.1_2-HSPs_+__Harpegnathos_saltator
                                  QANH01000473.1_2-HSPs_-_Harpegnathos_saltator
                    88.9
                                  QANH01000431.1_2-HSPs_-_Harpegnathos_saltator
                                  QANH01000093.1_136437-137001_+__Harpegnathos_saltator
                                  QANH01000029.1_73093-73573_-__Harpegnathos_saltator
                                 QANH01000132.1_90225-90957_-_Harpegnathos_saltator-Score_D-Event_6-Selective_pressure_NO
                                 QANH01000015.1_846072-846672_+__Harpegnathos_saltator
                                  QANH01000100.1_75973-76540_-_Harpegnathos_saltator
                   87
                                 QANH01000747.1_79776-80670_-_Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                  QANH01000110.1_557271-557934_+__Harpegnathos_saltator
                                 QANH01000041.1_91644-92214_+__Harpegnathos_saltator
                                  QANH01000762.1_24501-24984_-_Harpegnathos_saltator
                                  QANH01000367.1_2204-2873_+__Harpegnathos_saltator
                                  QANH01000368.1_213270-213705_-_Harpegnathos_saltator
                                  QANH01000124.1_141885-142722_+__Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                   71.9
                                  QANH01000535.1_73744-74437_+__Harpegnathos_saltator
                                 QANH01000041.1_52710-53469_+__Harpegnathos_saltator-Score_D-Event_6-Selective_pressure_NO
                                 QANH01000521.1_34032-34728_-__Harpegnathos_saltator
                                 QANH01000280.1_104787-105339_+__Harpegnathos_saltator
                                  QANH01000130.1_39537-40224_+__Harpegnathos_saltator
                   98.5
                                  QANH01000041.1_2682-3447_+__Harpegnathos_saltator-Score_D-Event_6-Selective_pressure_NO
                                  QANH01000046.1_2130476-2131232_-__Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                  QANH01000028.1_4858-5299_+__Harpegnathos_saltator
                                  QANH01000823.1_2531-3257_+__Harpegnathos_saltator-Score_A-Event_6-Selective_pressure_NO
                                  QANH01000777.1_13889-14324_-_Harpegnathos_saltator
                                 QANH01000403.1_106672-107089_+__Harpegnathos_saltator
                                  QANH01000520.1_24041-24839_-__Harpegnathos_saltator
                 L<del>82</del>157.4
                                  QANH01000376.1_16873-17471_+__Harpegnathos_saltator
                                  KQ467049.1_62228-62642_-_Dinoponera_quadriceps
                                  LBMM01014528.1_922-1558_-__Lasius_niger
                                  NJRQ01001145.1_6535-7201_+__Aphaenogaster_ashmeadi
                                 NJRM01000458.1_42004-42619_-_Aphaenogaster_picea
                                 JWHX01001200.1_29566-30364_+_Pseudomyrmex_gracilis-Score_A-Event_2-Selective_pressure_NO
                                  GL905377.1 226012-226729 - Linepithema humile
     Root
                                  scaffold_25_26715-27438_-__Eupelmus_kiefferi
                                  scaffold 1031 11857-12580 + Aphidius colemani
                                 YP_009165874|Apismelliferafilamentousvirus
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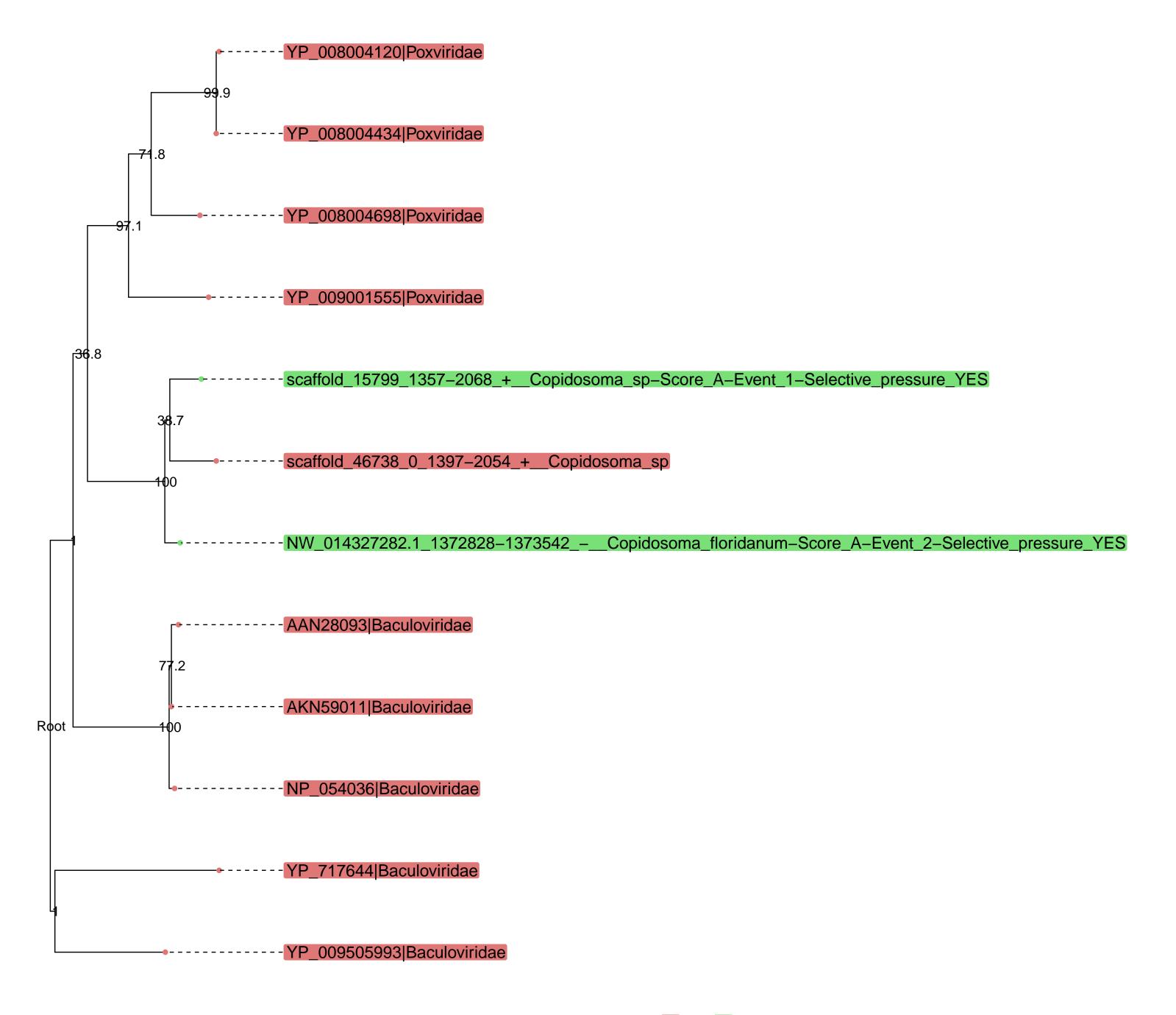




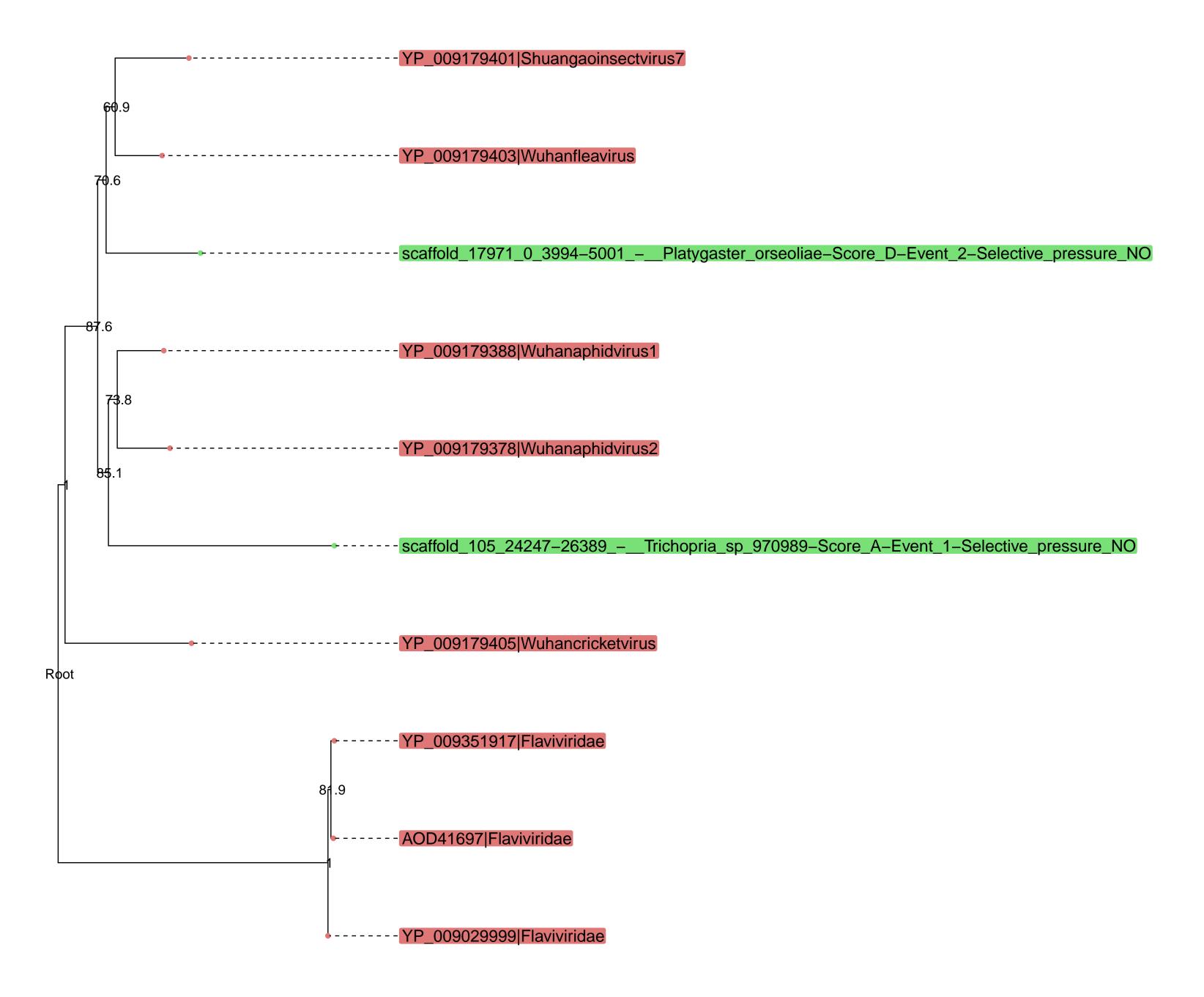


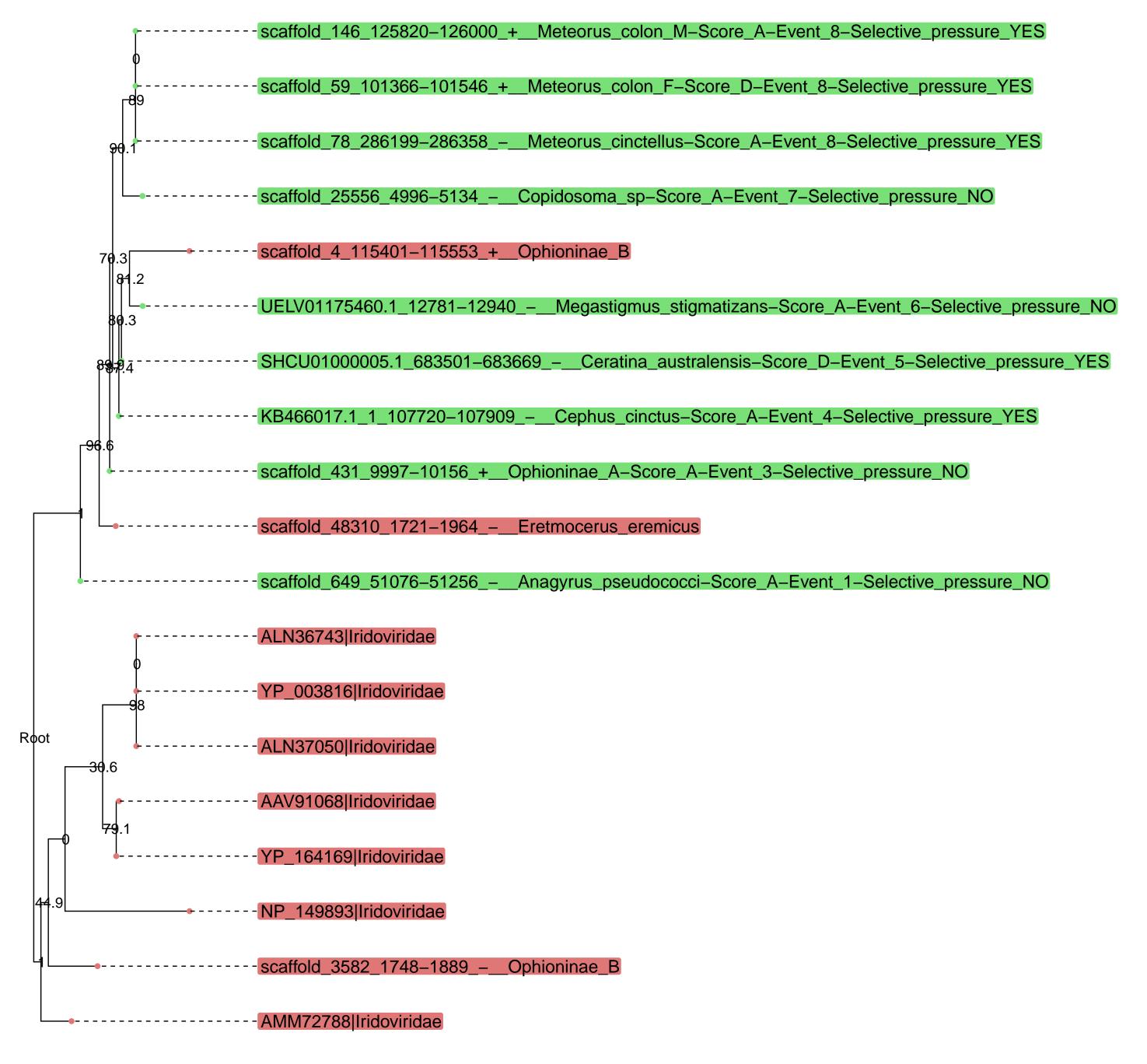


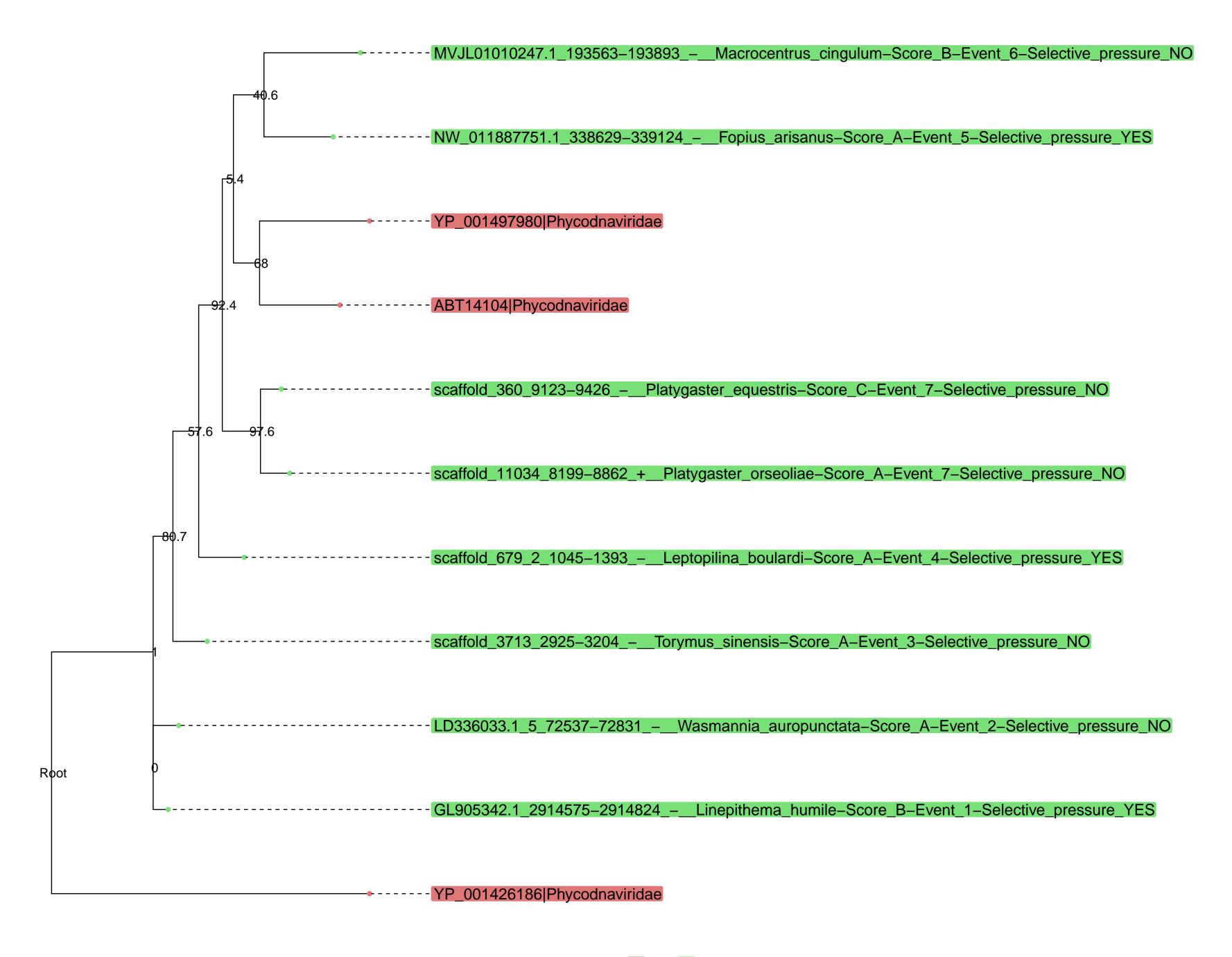
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Cluster17658 – families involved: ['Orthomyxoviridae'] – putative protein: ['RNA-directed RNA polymerase catalytic subunit (EC 2.7.7.48)']
                                                          AGK63053|Orthomyxoviridae
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                                                         ACF41842|Orthomyxoviridae
                                                         AEX92881|Orthomyxoviridae
                                                         BAV59609|Orthomyxoviridae
                                                         AGU93028|Orthomyxoviridae
                                                         NP_040985|Orthomyxoviridae
                                                          YP_308665|Orthomyxoviridae
                                                         NP_859040|Orthomyxoviridae
                                                          YP_009118628|Orthomyxoviridae
                                                          YP_308847|Orthomyxoviridae
                                                          YP_009118471|Orthomyxoviridae
                                                          YP_308851|Orthomyxoviridae
                                                         BAA00002|Orthomyxoviridae
                                     76.7
                                                         NP_056657|Orthomyxoviridae
                                                          AFH57863|Orthomyxoviridae
                                                          AON76701|Orthomyxoviridae
                                                         ALE66294|Orthomyxoviridae
                                                         ALE66283|Orthomyxoviridae
                                                         AON76709|Orthomyxoviridae
                                                          YP_009449556|Orthomyxoviridae
                                                         AMN87905|Orthomyxoviridae
                                                         BBC14928|Orthomyxoviridae
                                                         BAY00630|Orthomyxoviridae
                                                         BAV18003|Orthomyxoviridae
                                                         AMN87906|Orthomyxoviridae
                 <del>-9</del>9.3
                                                          AMN87903|Orthomyxoviridae
                                                         CEE50061|Orthomyxoviridae
                                                          YP_089653|Orthomyxoviridae
                                                          YP_009553280|Orthomyxoviridae
                                             <del>-99</del>.8
                                                           P_009352882|Orthomyxoviridae
                                                          YP_145794|Orthomyxoviridae
                                                          BAQ22394|Orthomyxoviridae
                                                         AEC12828|Orthomyxoviridae
                                                          ABF68025|Orthomyxoviridae
                                                          YP_145804|Orthomyxoviridae
                                                         - ABF68026|Orthomyxoviridae
                                                         - AAK48525|Orthomyxoviridae
                                                         AQT26412|Orthomyxoviridae
                                                         ANM86269|Orthomyxoviridae
                                                         ANM86249|Orthomyxoviridae
                                                         ANM86259|Orthomyxoviridae
                                                         - AQT26406|Orthomyxoviridae
                                                         ABG65760|Orthomyxoviridae
                                                         ACG56669|Orthomyxoviridae
                                                         ADF36506|Orthomyxoviridae
                                                         ACG56670|Orthomyxoviridae
                                                         ABW93483|Orthomyxoviridae
                                                         ABG65756|Orthomyxoviridae
                                                         ACV71155|Orthomyxoviridae
                                                         ABG65757|Orthomyxoviridae
                                                         ACG56671|Orthomyxoviridae
                                                         ACB45380|Orthomyxoviridae
                                                         ADF36496|Orthomyxoviridae
                                                         CAA05486|Orthomyxoviridae
                                                         ABG65767|Orthomyxoviridae
                                                          ABG65768|Orthomyxoviridae
                                                         ACJ37398|Orthomyxoviridae
                                                          ABG65759|Orthomyxoviridae
                                                          ABG65761 | Orthomyxoviridae
                                               72.9
                                                          ABG65764|Orthomyxoviridae
                                                          ADR77506|Orthomyxoviridae
                                                          ABG65758|Orthomyxoviridae
                                                                                       _Eupelmus_annulatus-Score_C-Event_14-Selective_pressure_NO
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     Root
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                                   <del>3$</del>.5
                                                          jcf7180001990402_33441-33723_+__Synergus_japonicus
                                    .3
                                                          cf7180001991336_4266-4698_-_Synergus_japonicus
                                    <del>72</del>.5
                                                          cf7180001923109_17241-17409_+__Synergus_umbraculus
                                                          scaffold_97357_402-627_+__Torymus_flavipes
                                                          scaffold_23685_1-364_+__Torymus_flavipes
                                                          scaffold_127387_253-478_+__Torymus_flavipes
                                                          UELX01013219.1_1885-4180_+__Eupelmus_urozonus
                             6$.9<sup>L</sup>7$.6
                                                          scaffold_19286_229-562_+__Torymus_flavipes
                                                          MPJT01000349.1_20741-21770_-__Osmia_bicornis-Score_A-Event_2-Selective_pressure_NO
                                                          JPHQ01017677.1_3-HSPs_-_Polistes_canadensis
                                                          KB466430.1_1_8450-9329_+__Cephus_cinctus-Score_A-Event_3-Selective_pressure_NO
                                                          MPJT01001439.1_8211-10258_-__Osmia_bicornis-Score_A-Event_8-Selective_pressure_NO
                                                          scaffold_47071_1097-1280_+__Trichopria_sp_970989
                                                          scaffold_104227_348-534_-__Ceratosolen_corneri
                                                          UELU01145086.1_6850-7114_-_Megastigmus_dorsalis
                                                          scaffold_102660_311-1010_+__Ganaspis_brasiliensis
                           [<del>7</del>$.<del>|5.5</del>∳.7
                                                          scaffold_180186_78-462_-_Ganaspis_brasiliensis
                                                          scaffold_382561_9-234_+__Torymus_sinensis
                                                          scaffold_95135_4-919_+__Ganaspis_brasiliensis
                                                          LSNX01007440.1_78-240_-_Ceratina_calcarata
                                                          YP_009508043|Orthomyxoviridae
                                                          YP_009665204|Unknown
                                                          YP_009110686|Orthomyxoviridae
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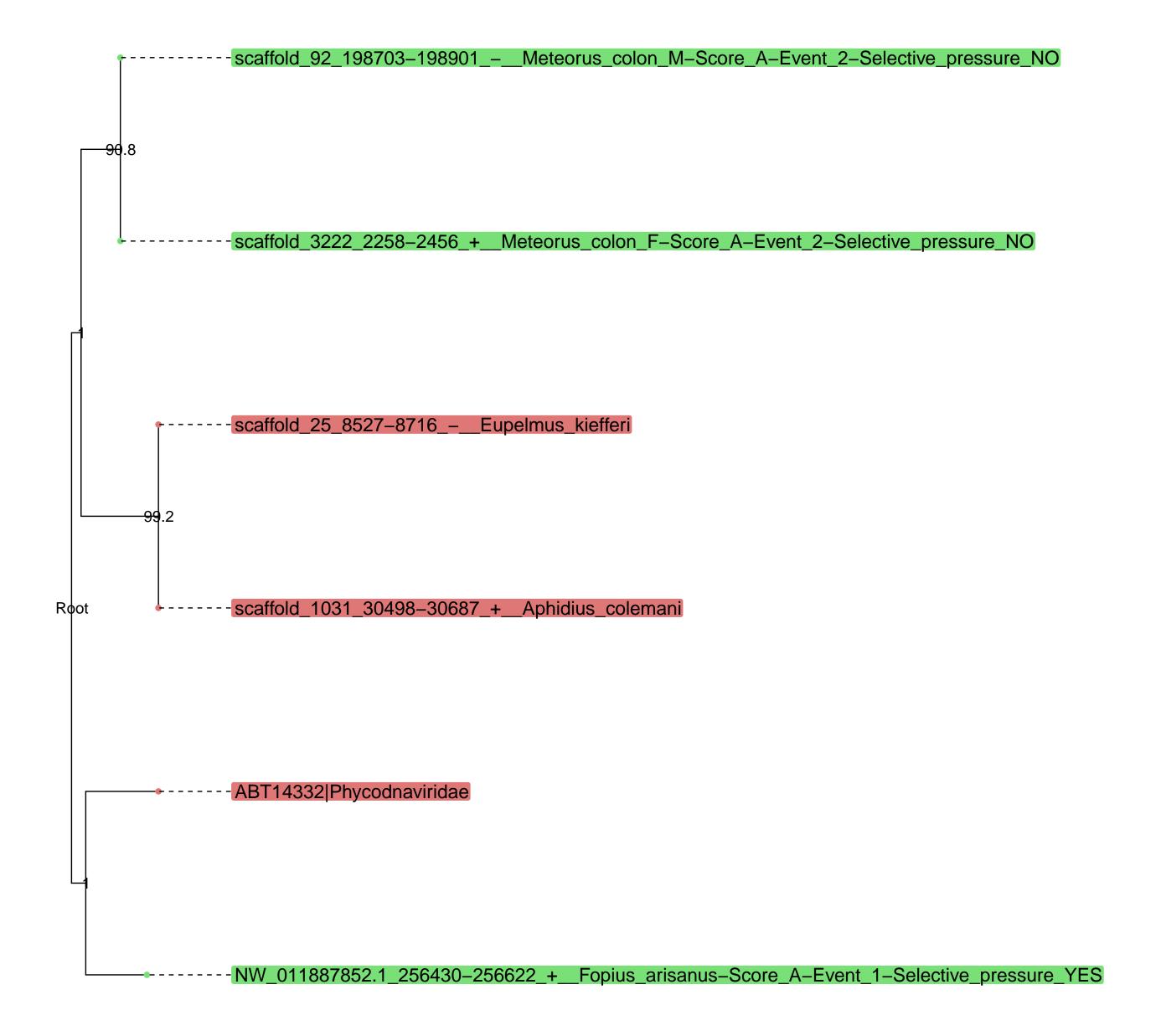


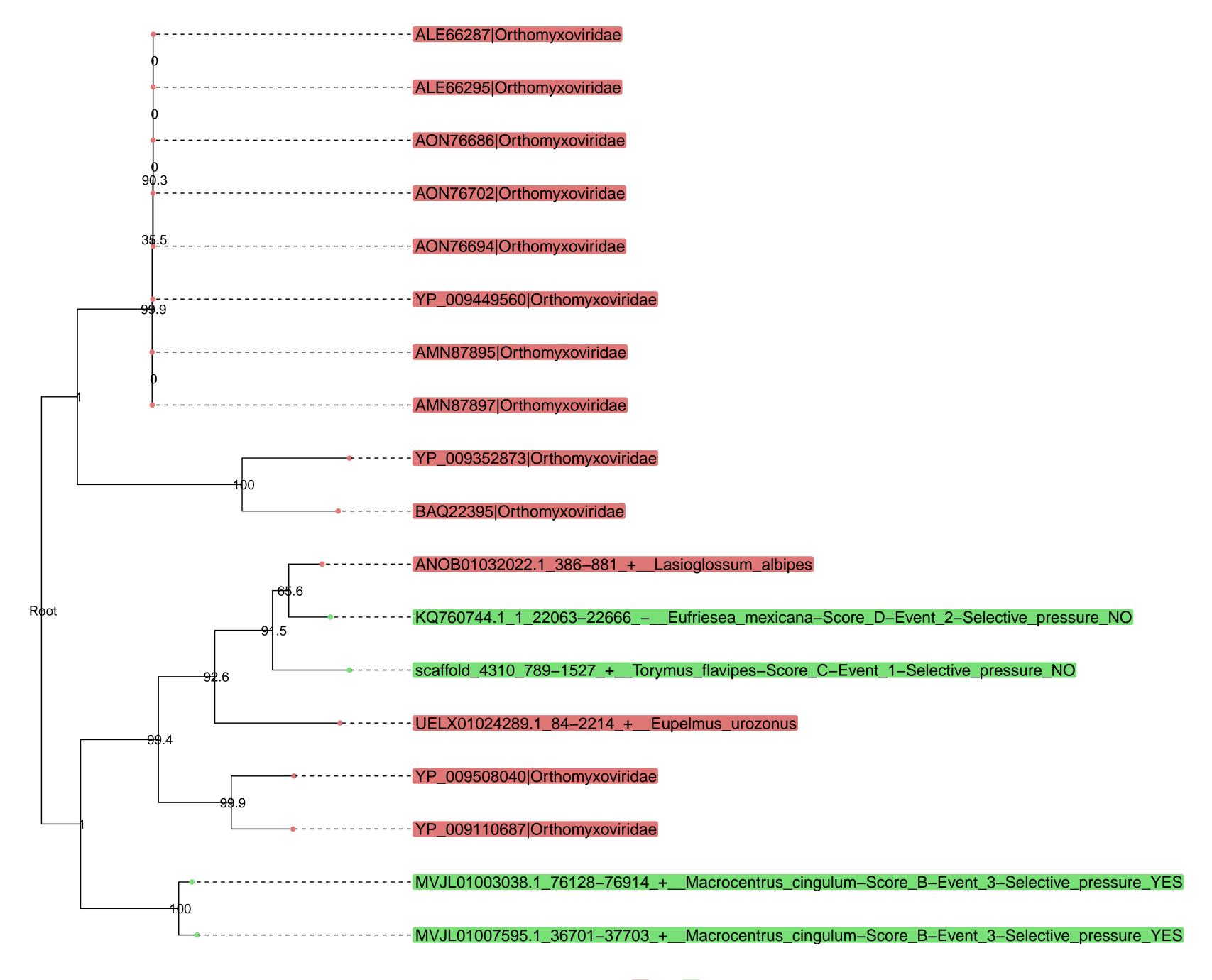
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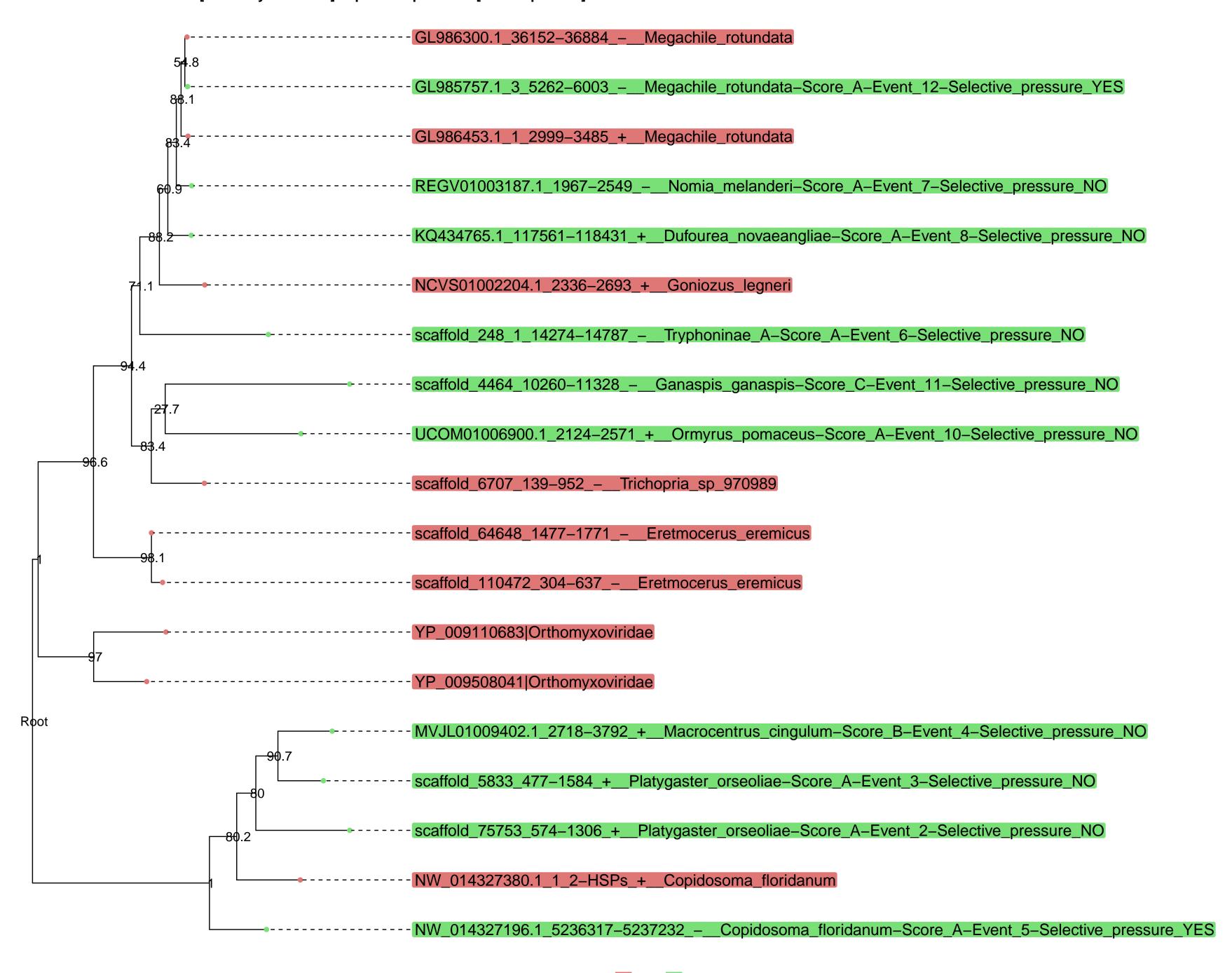


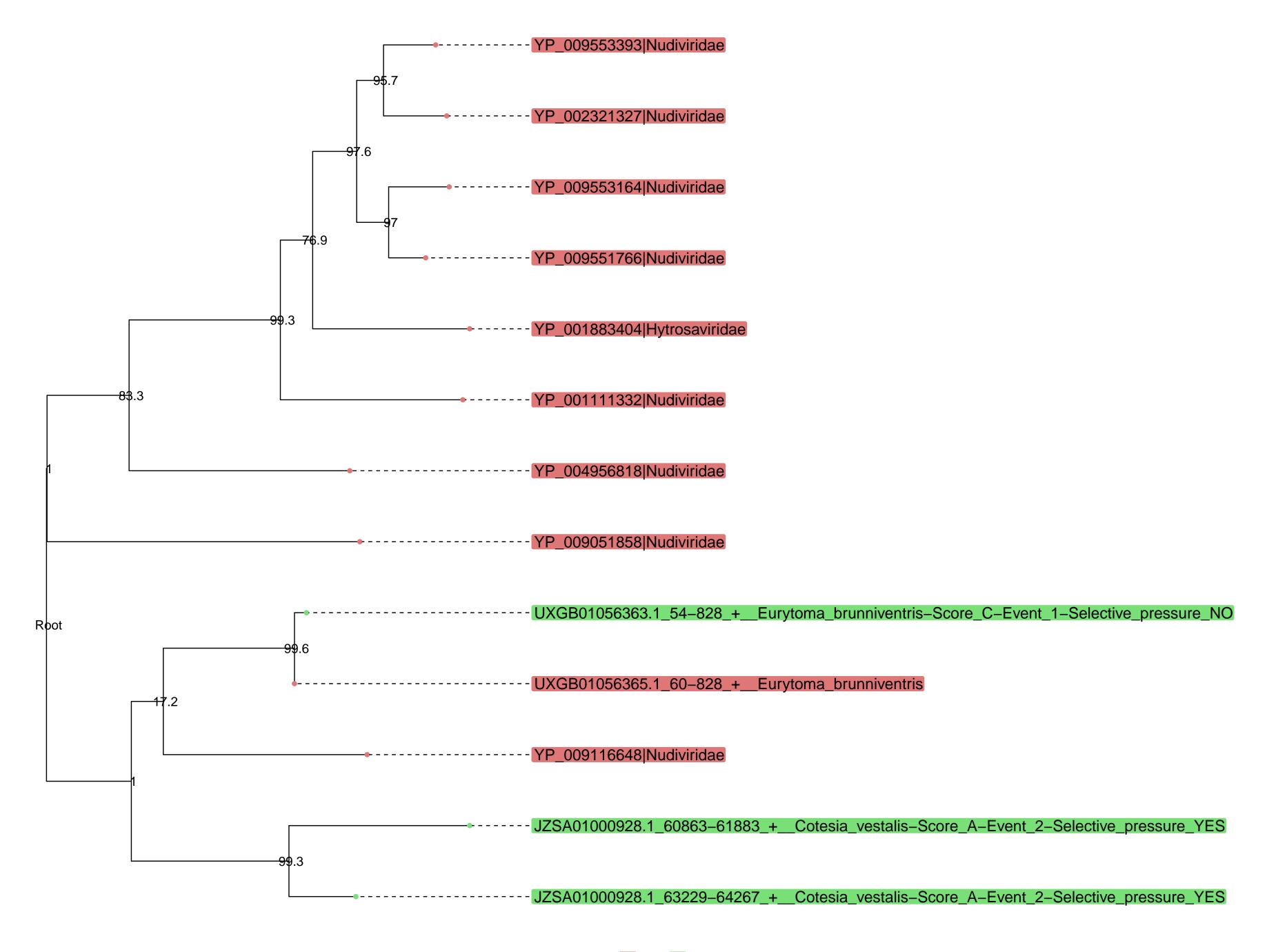


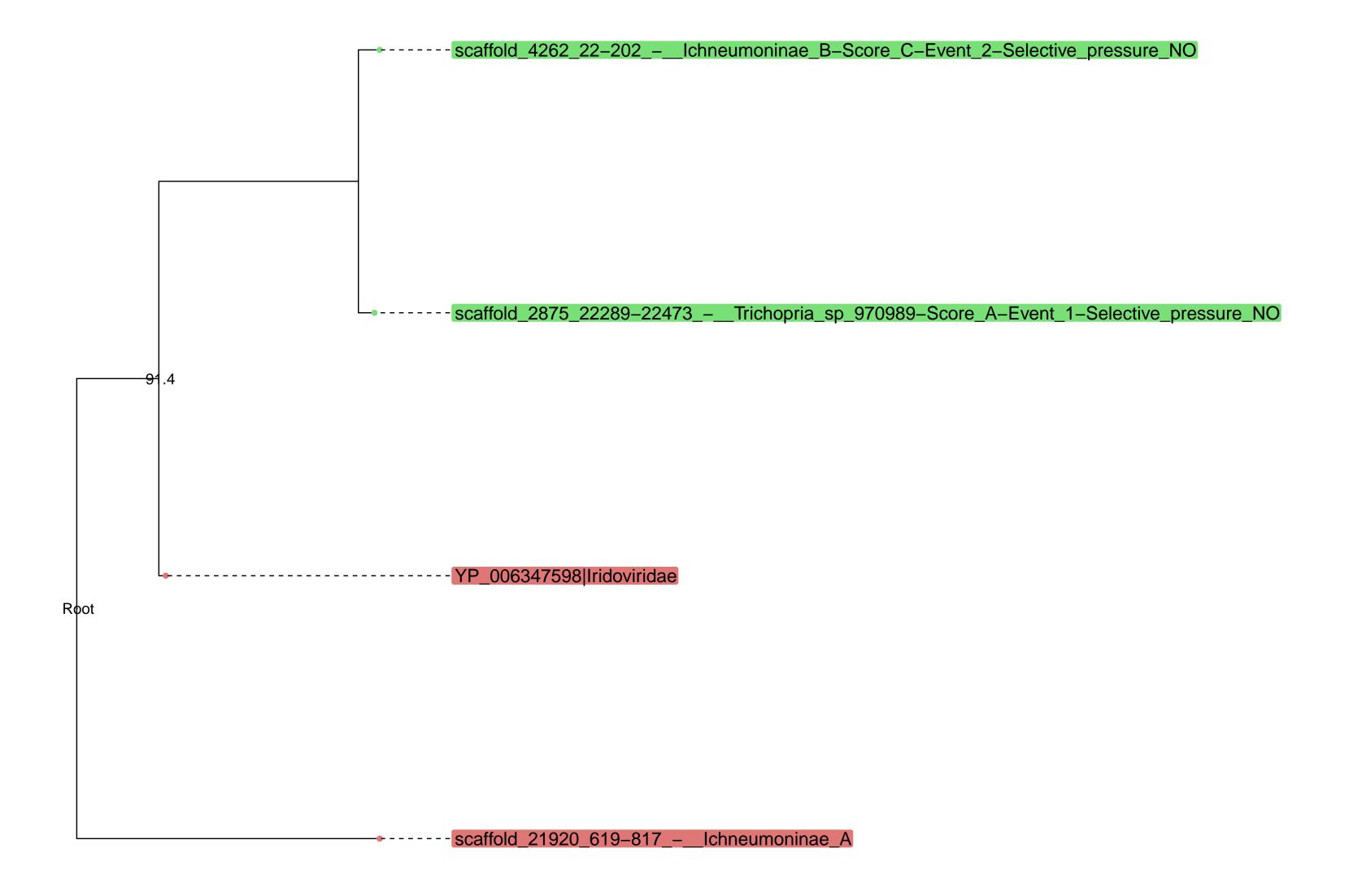


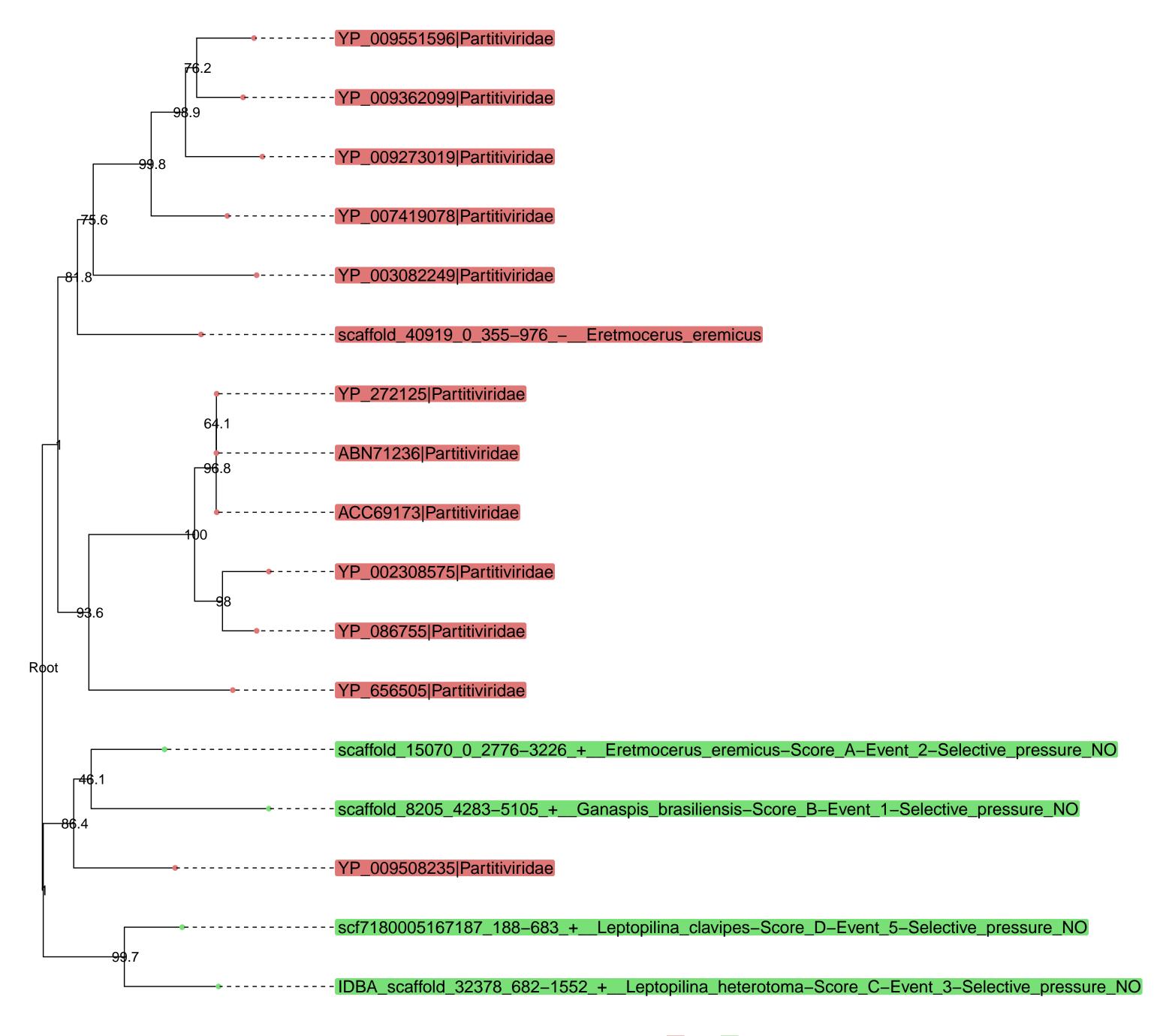


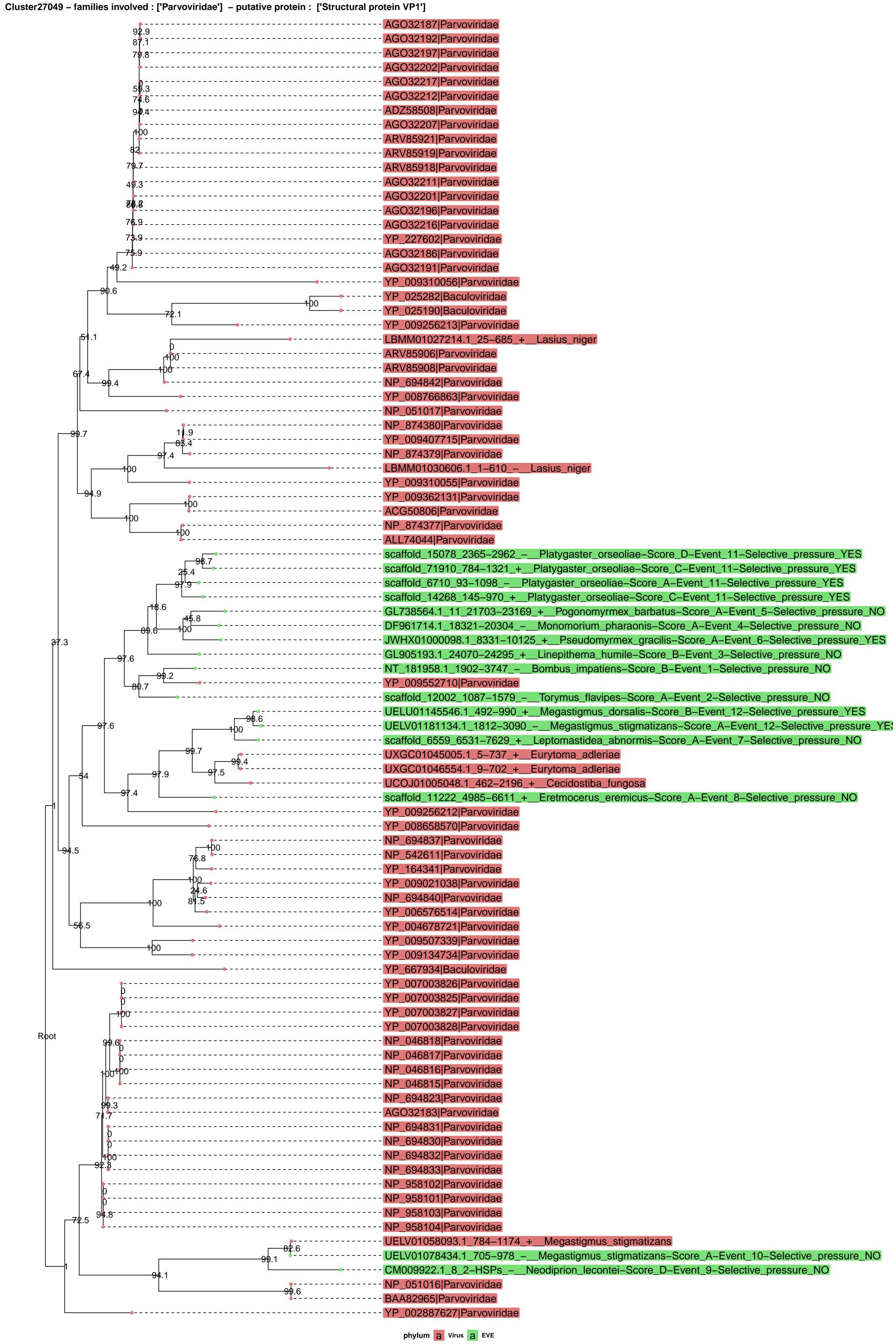


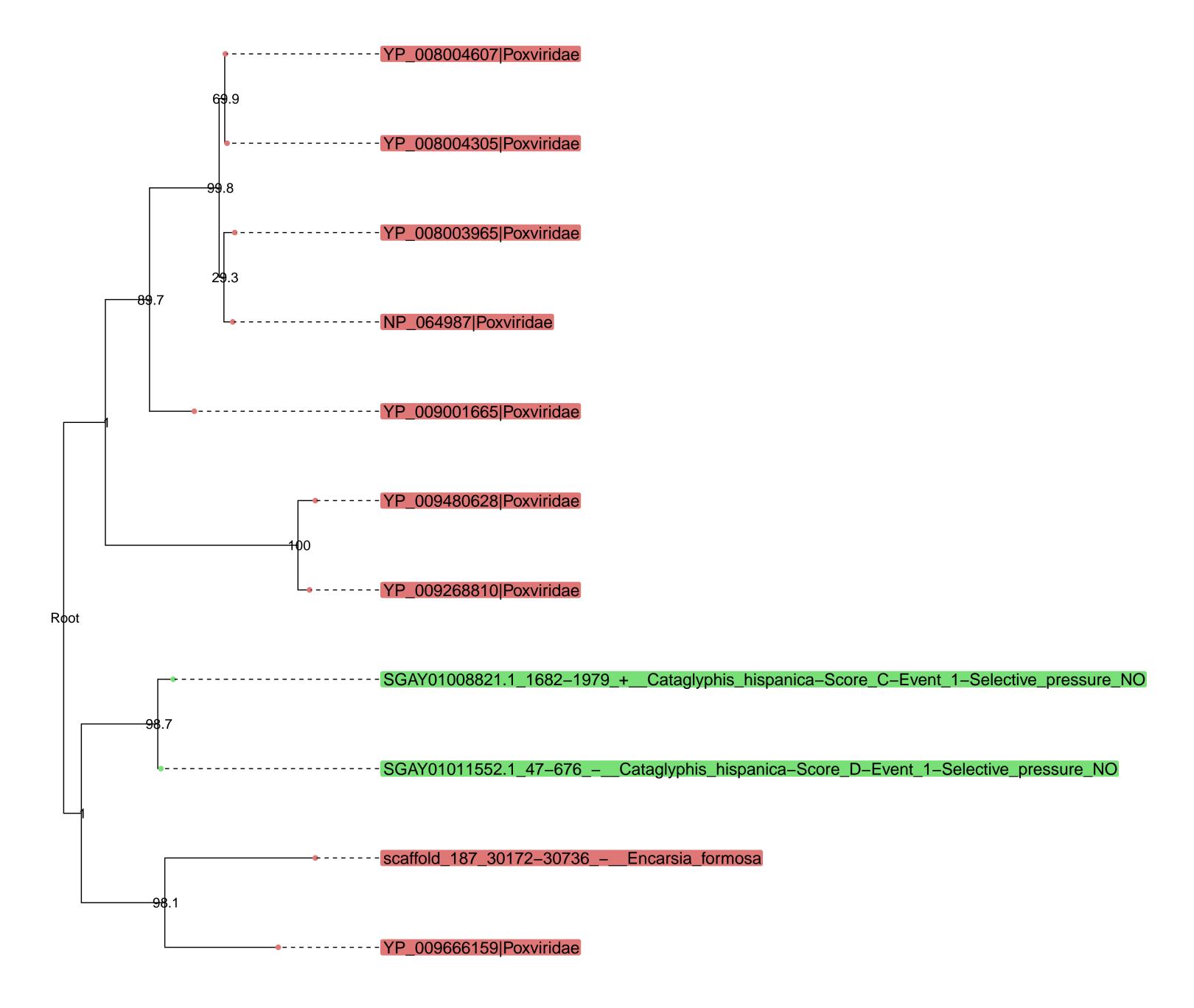


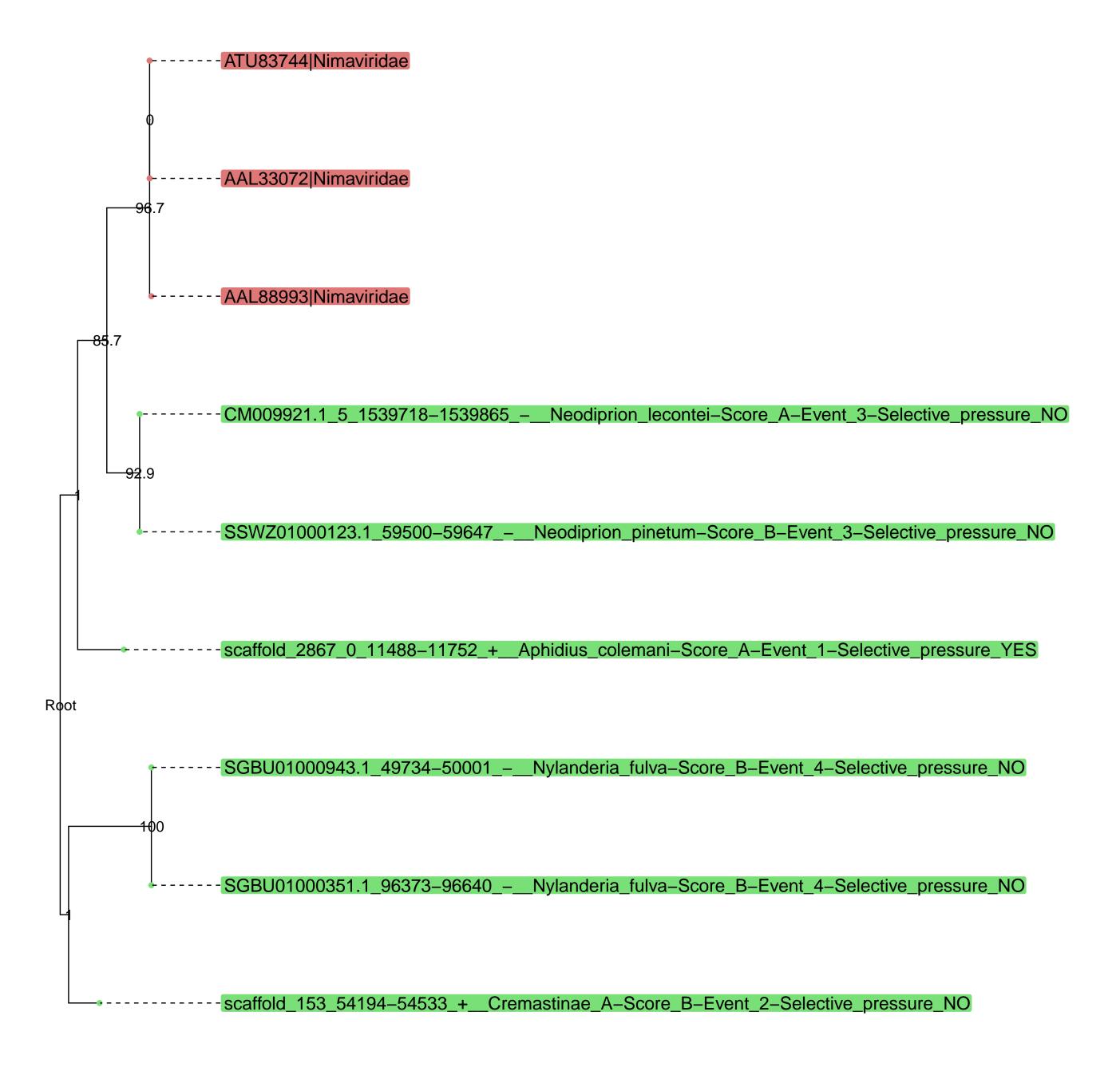


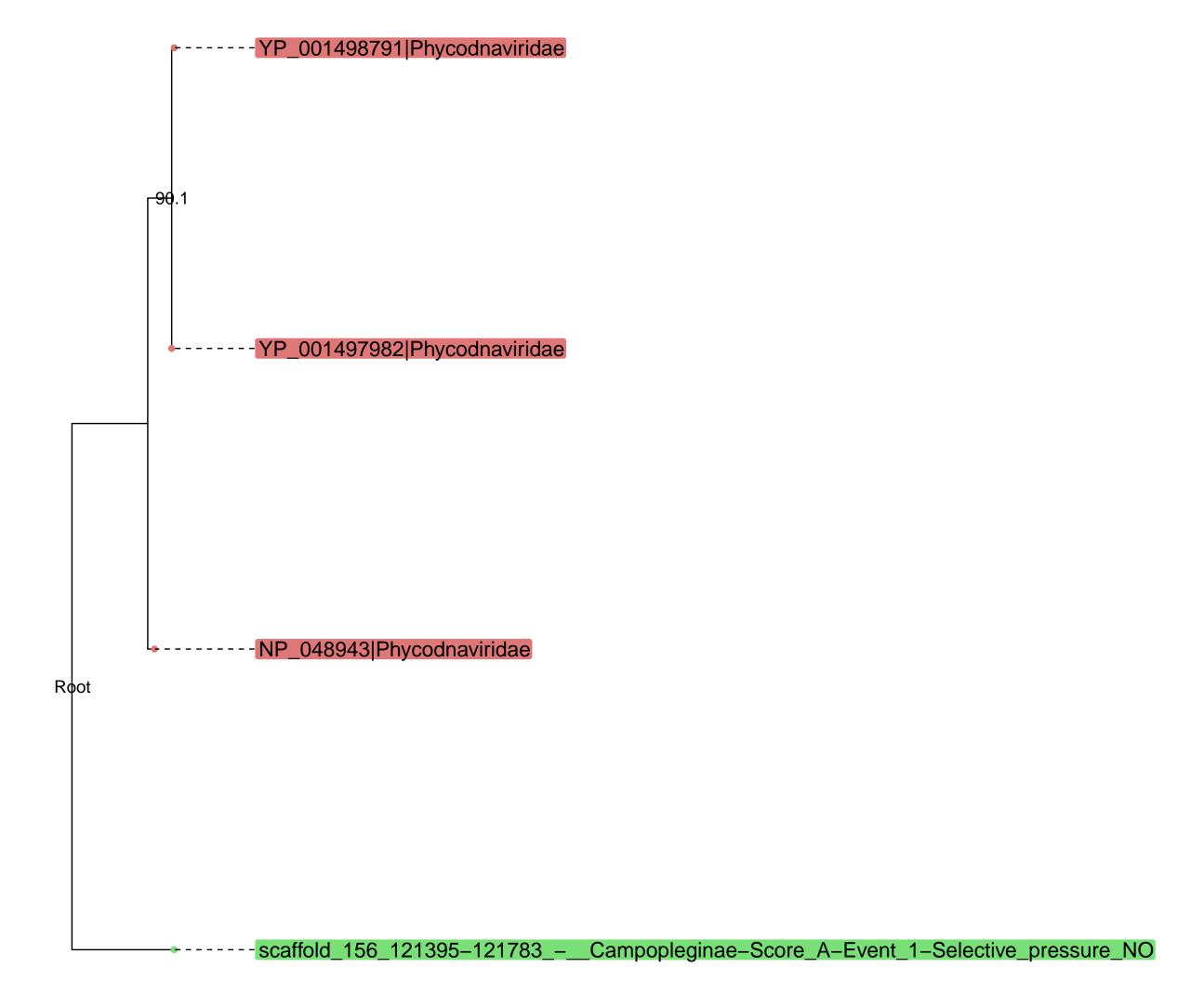


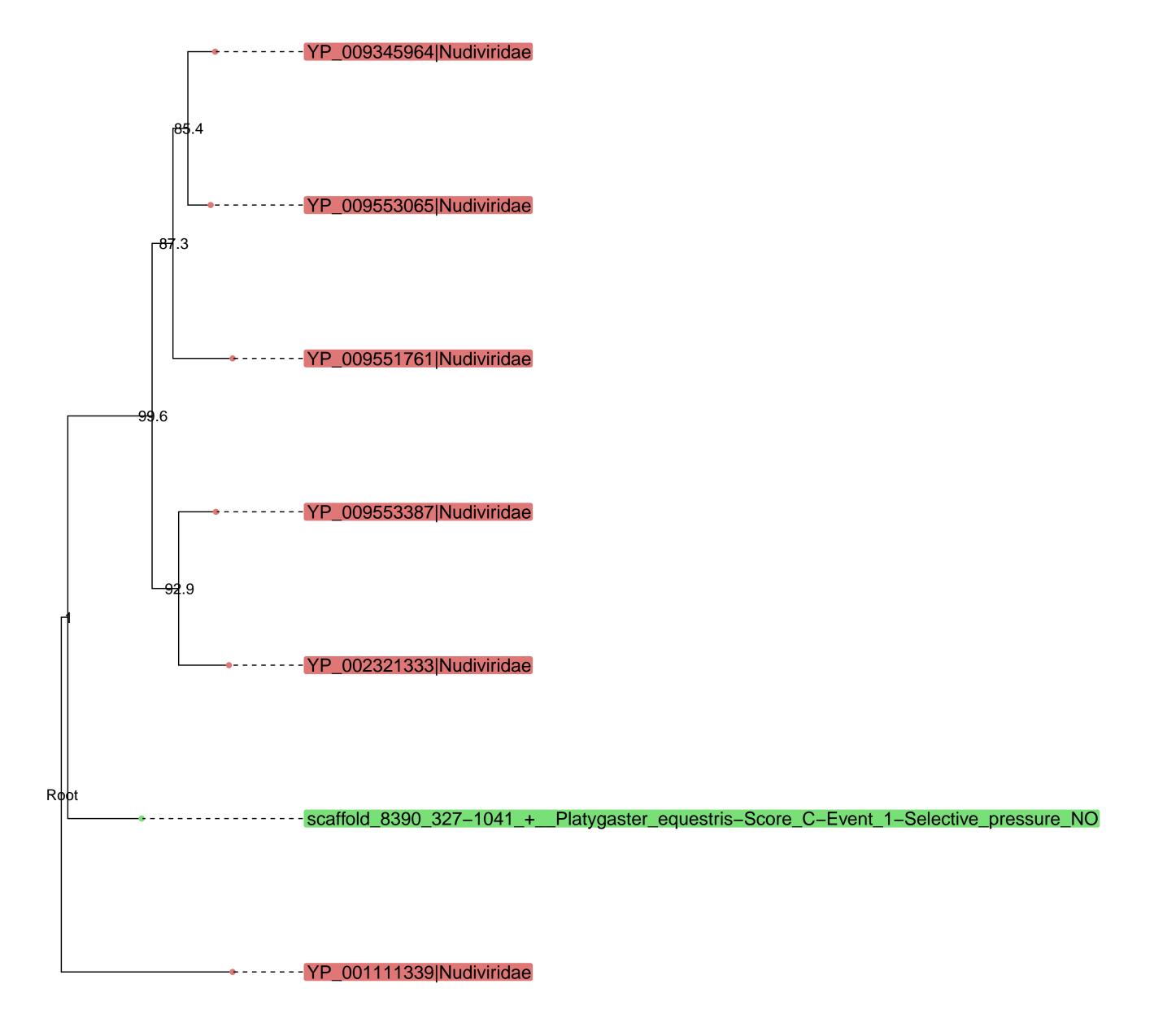


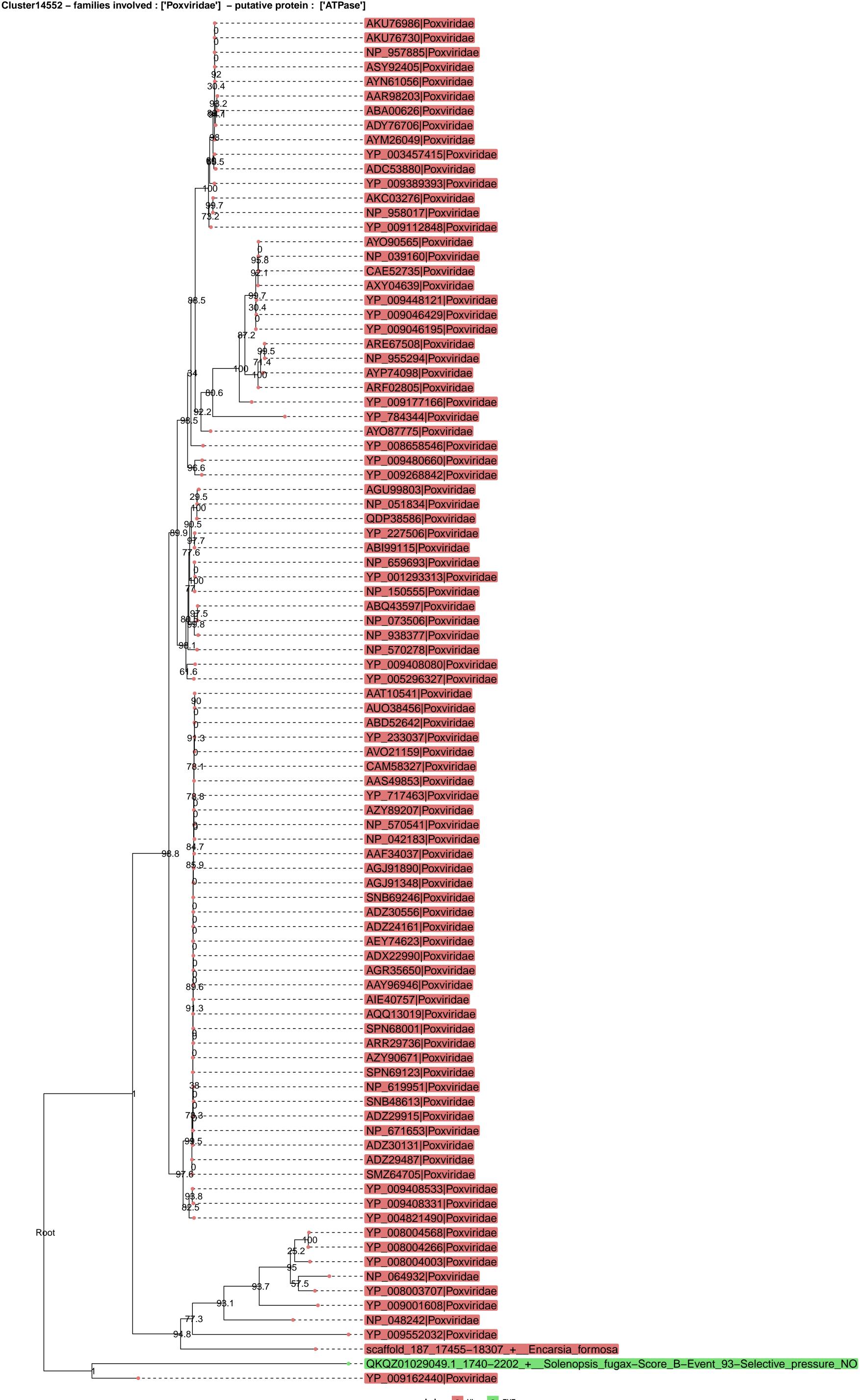


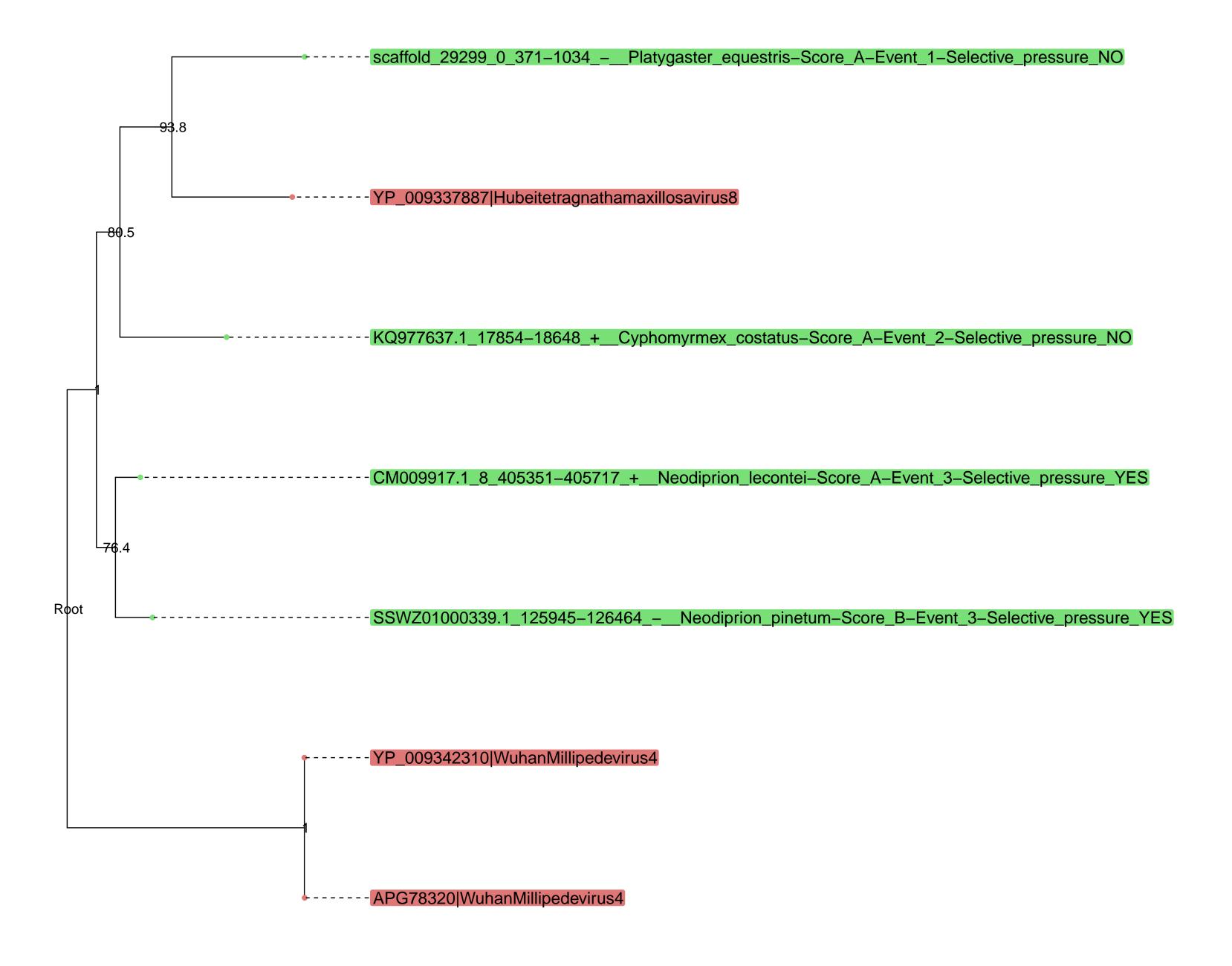


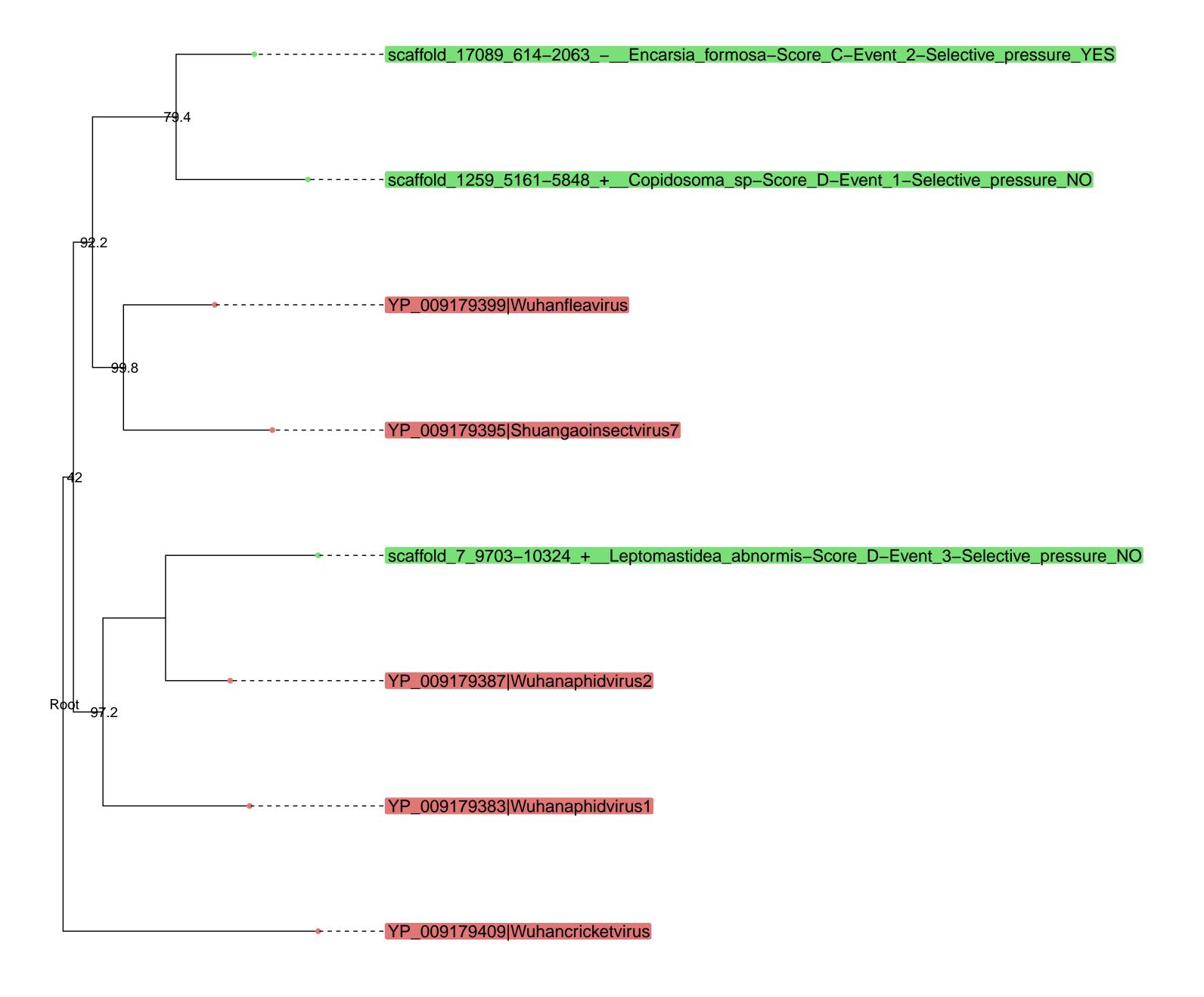


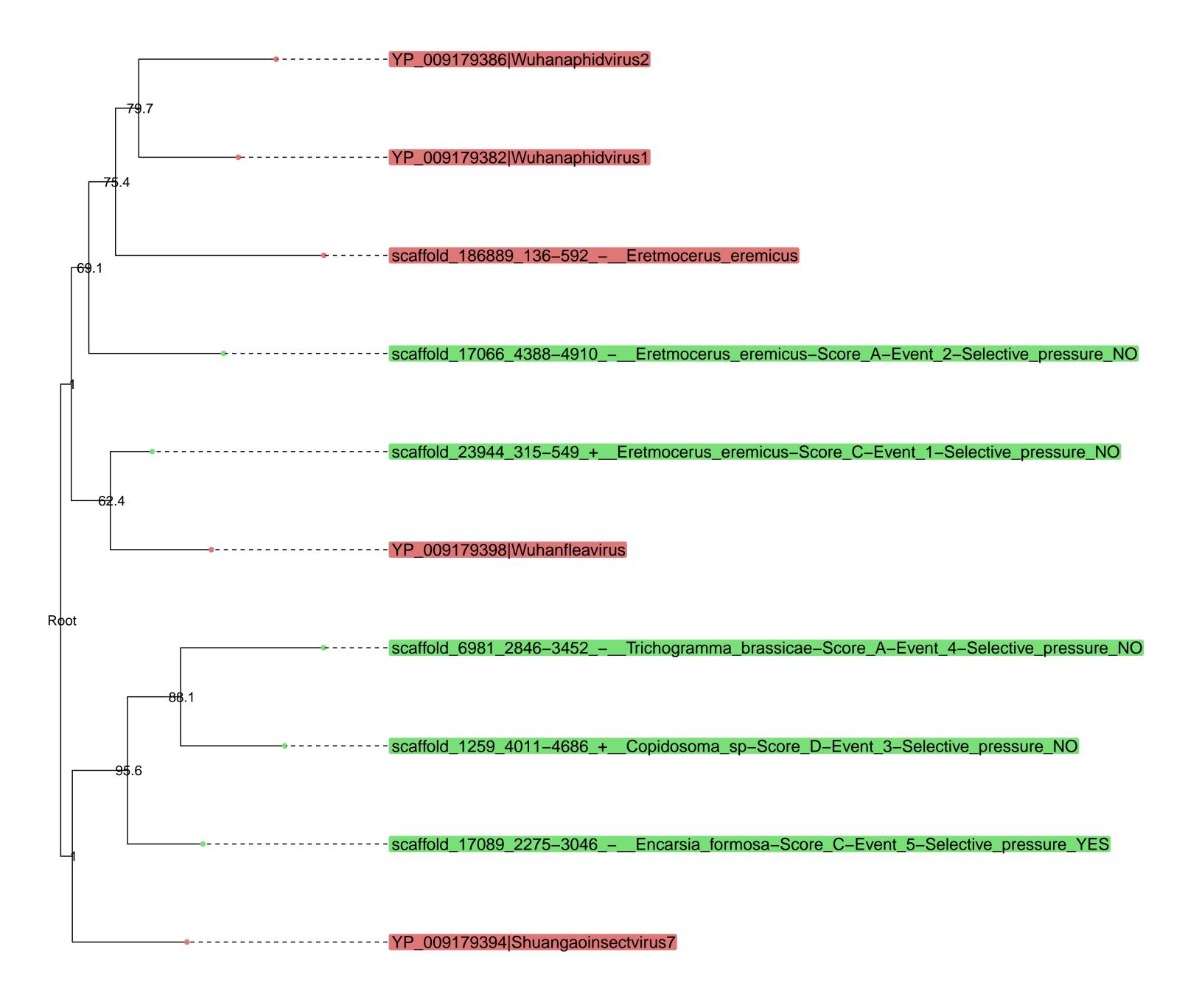




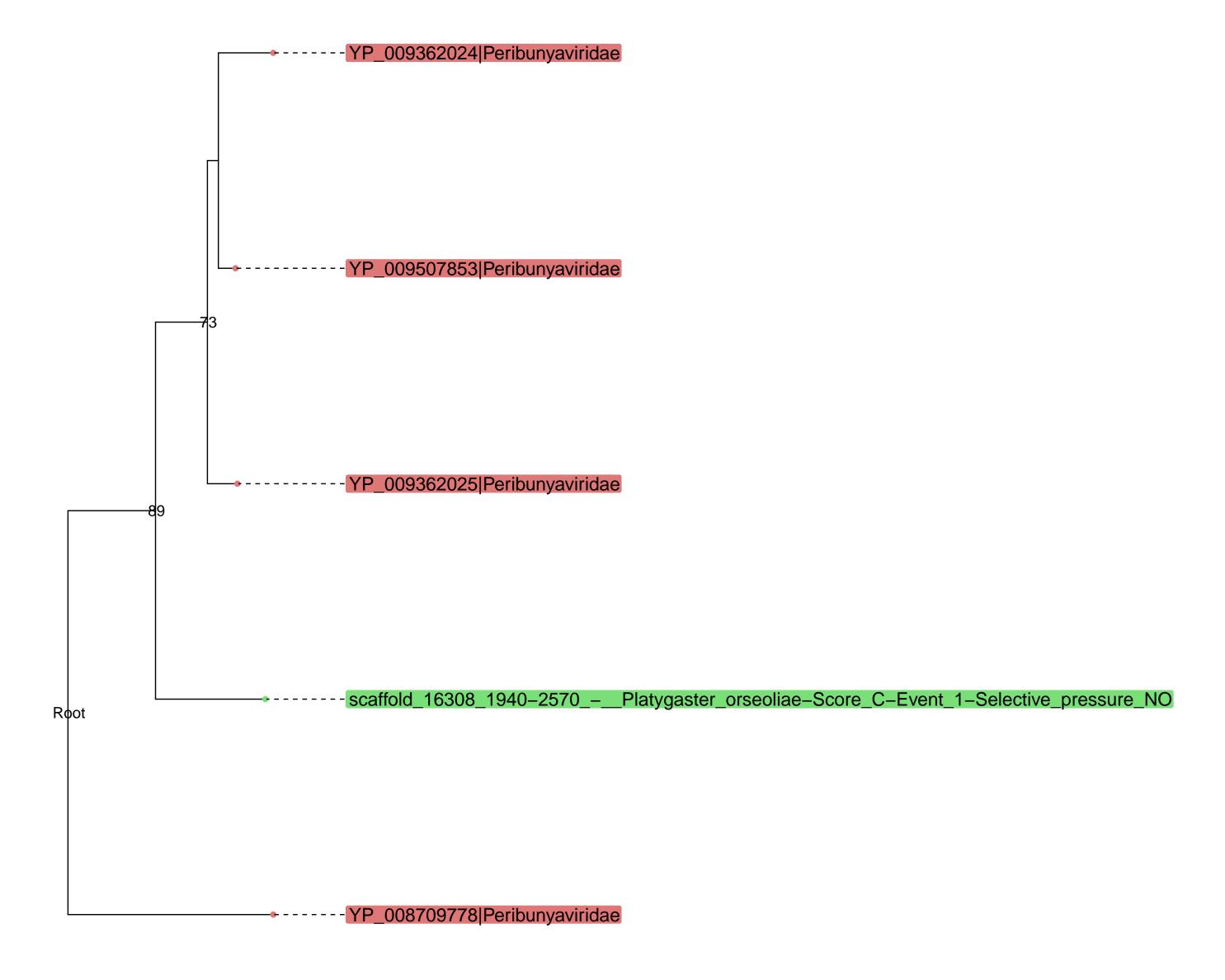


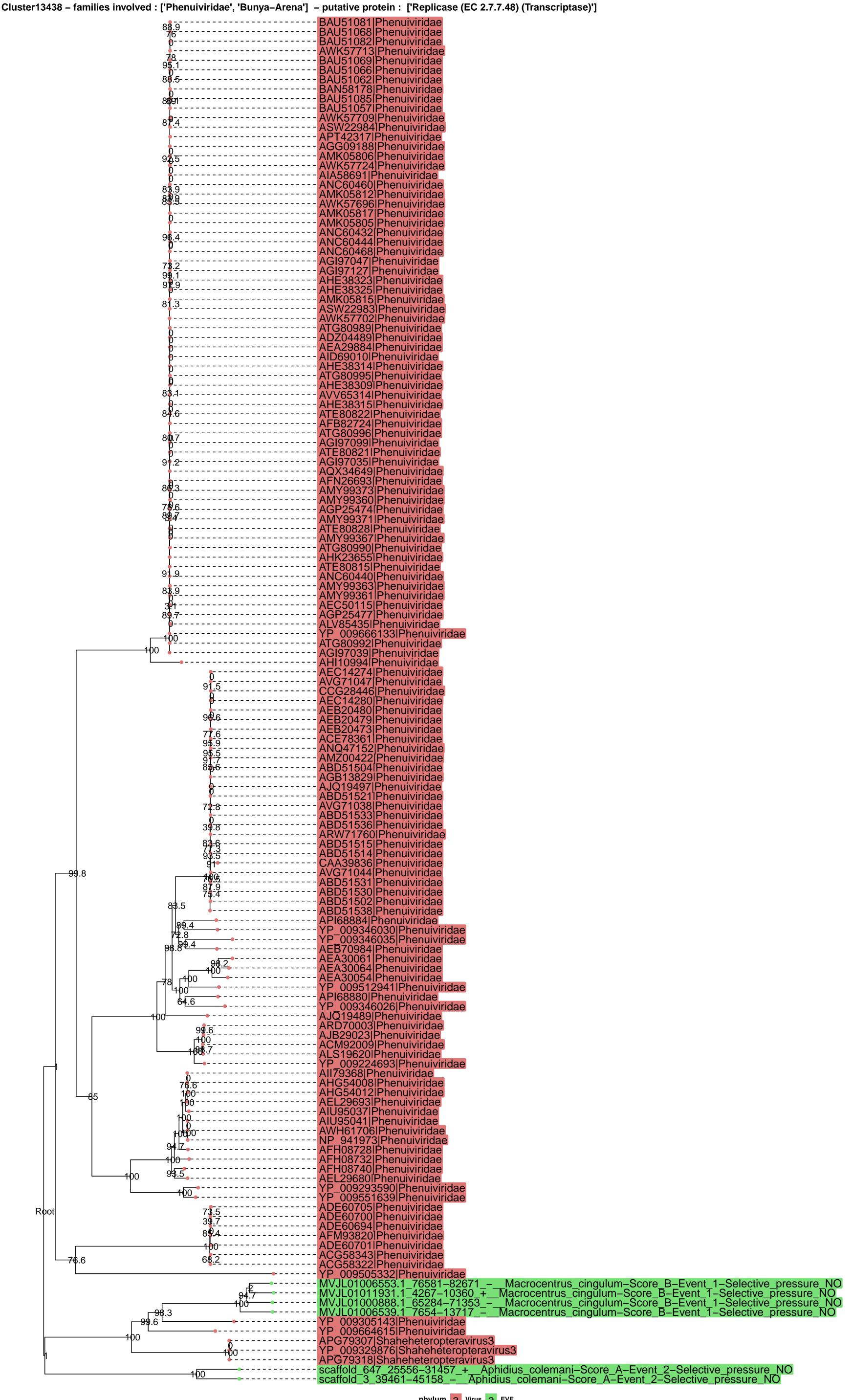


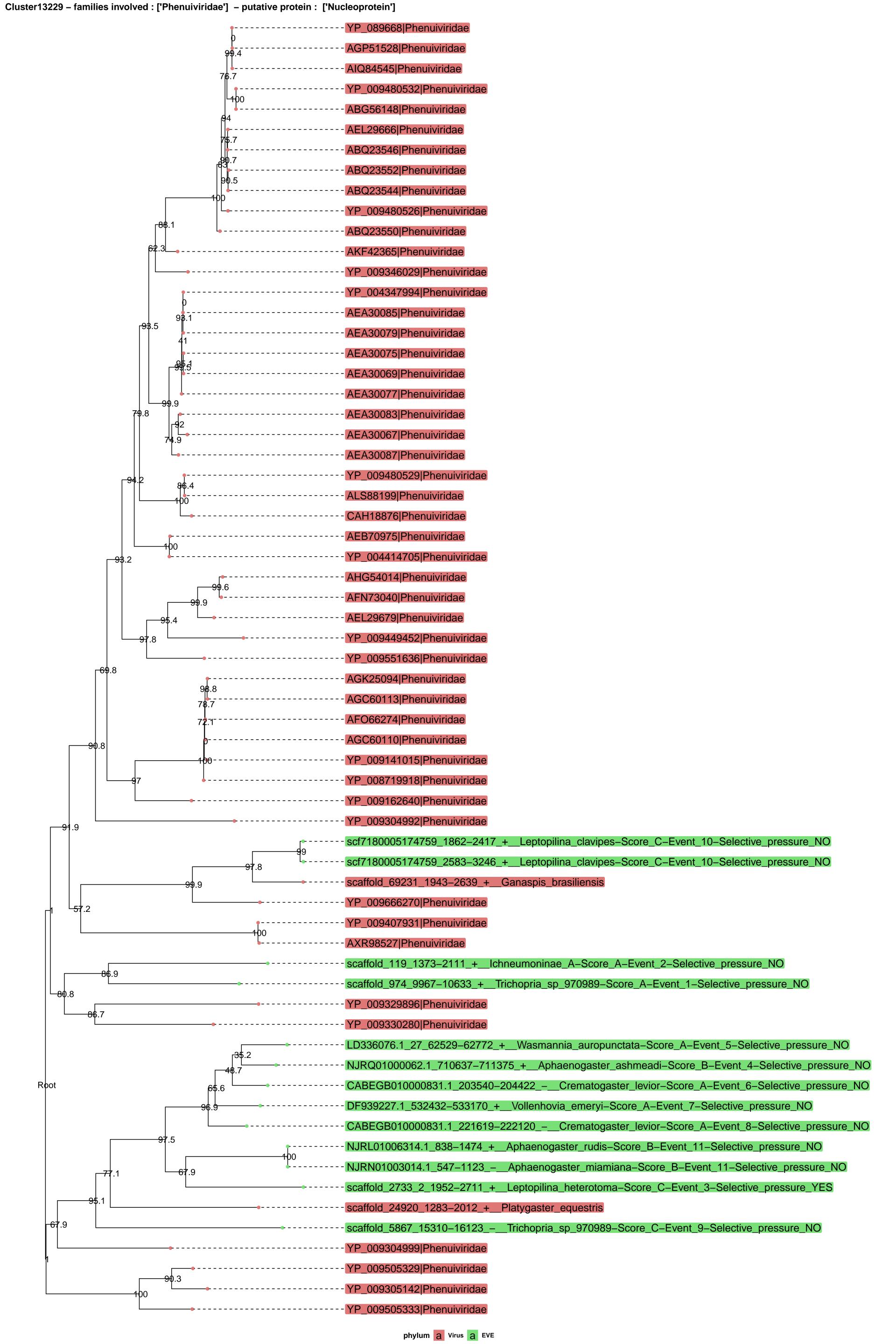


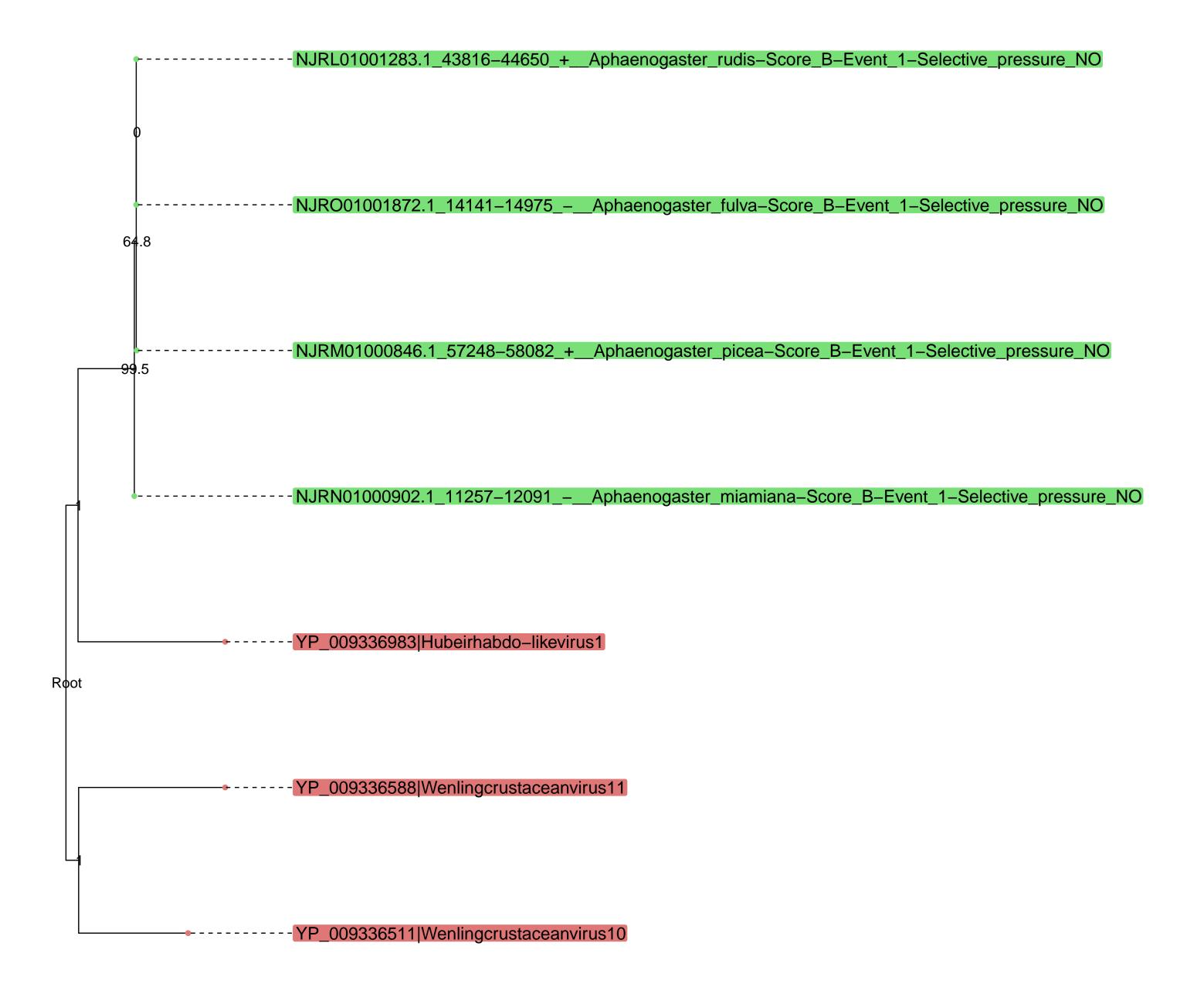


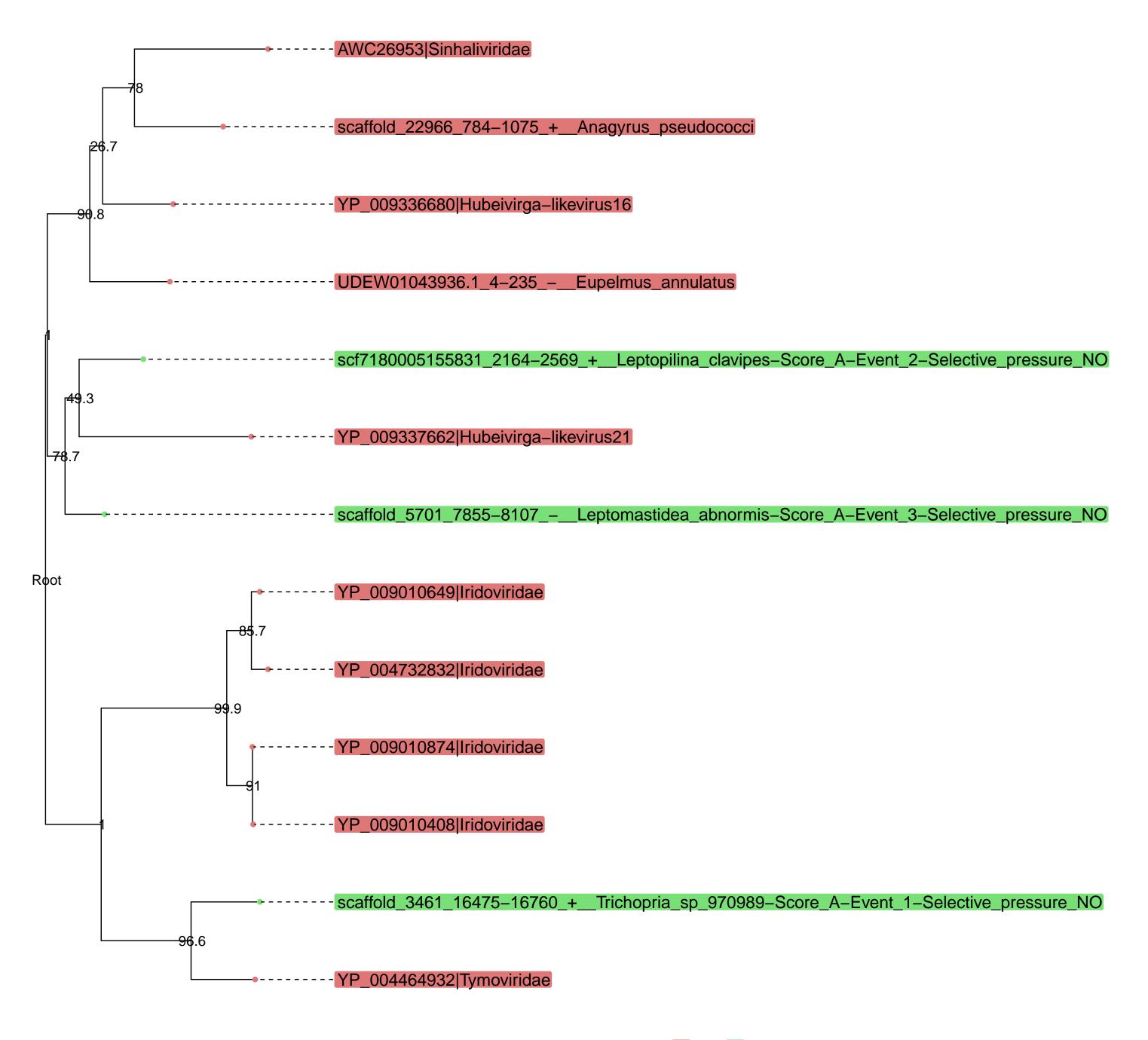
-YP_009116686|Nudiviridae

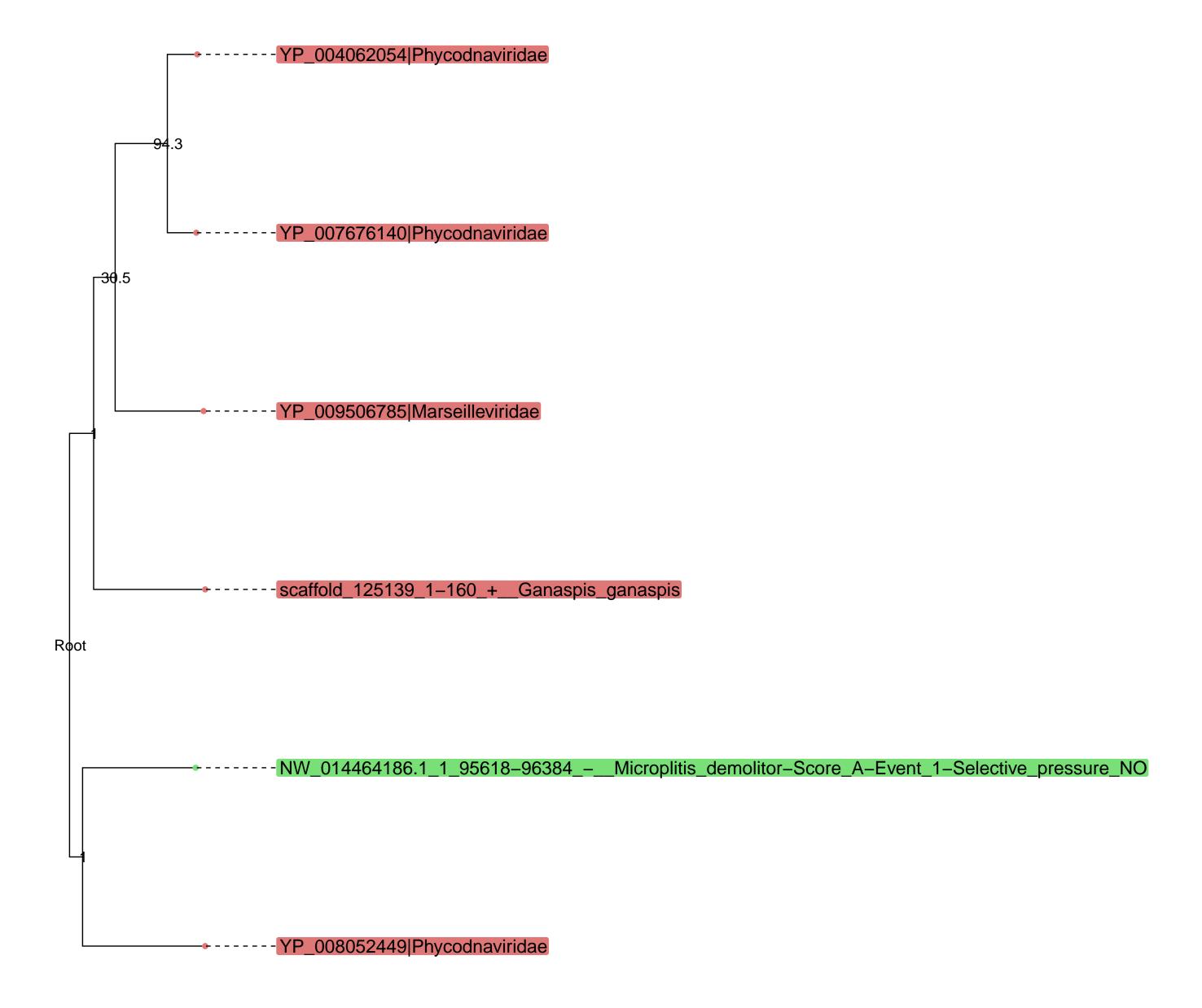


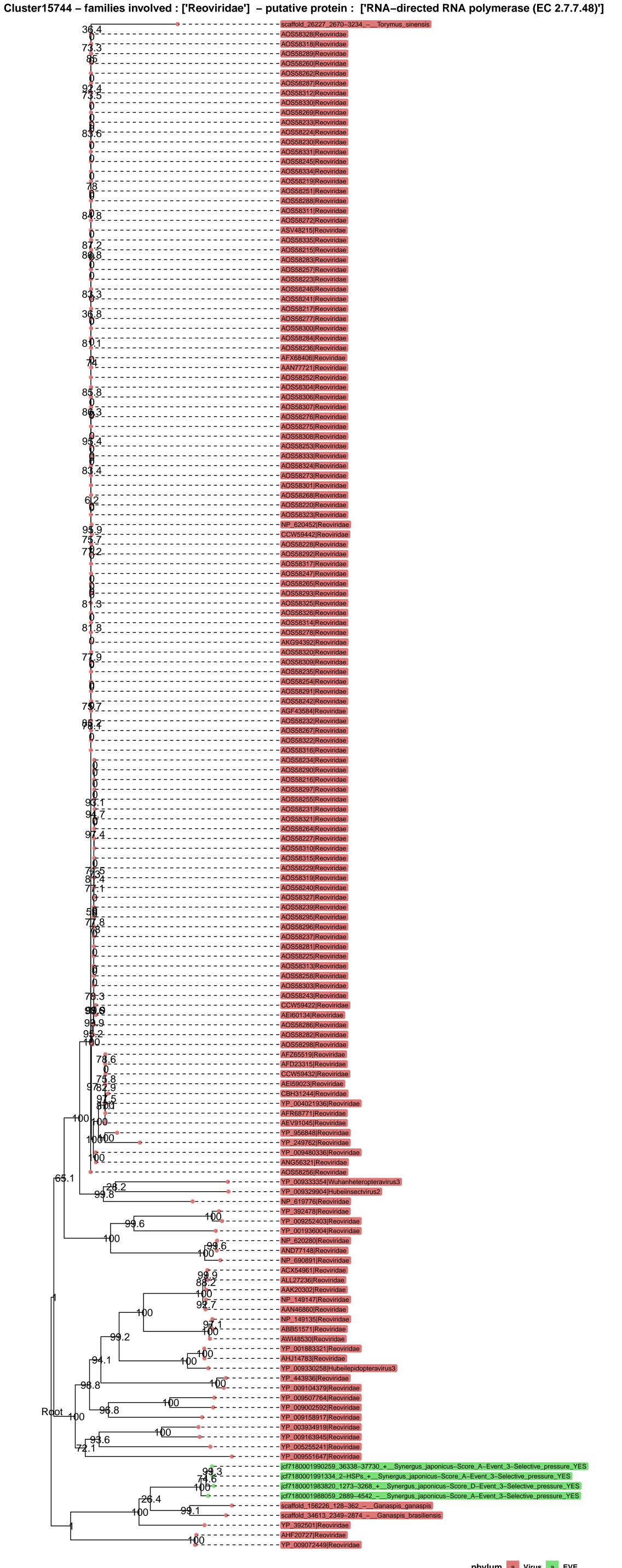




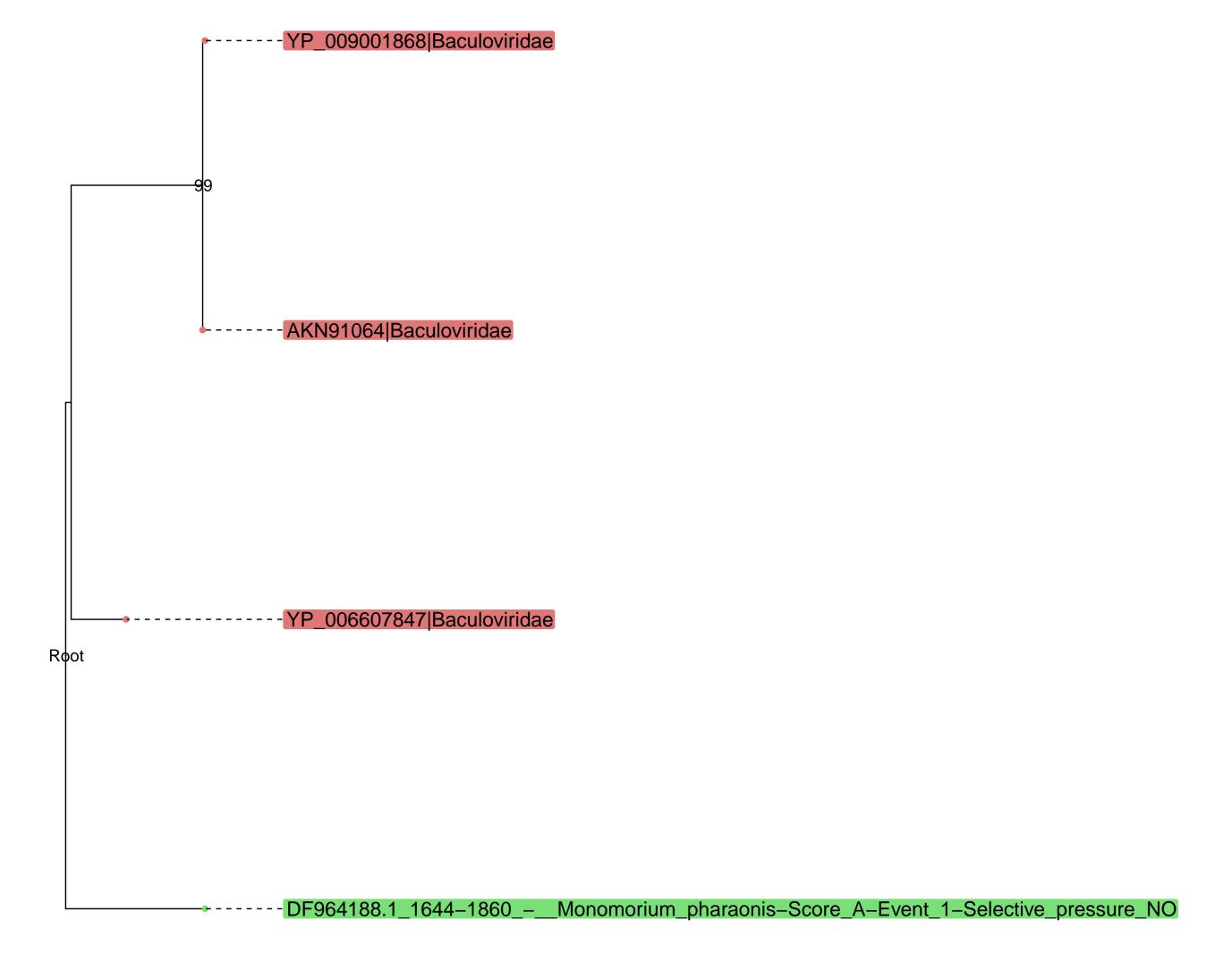


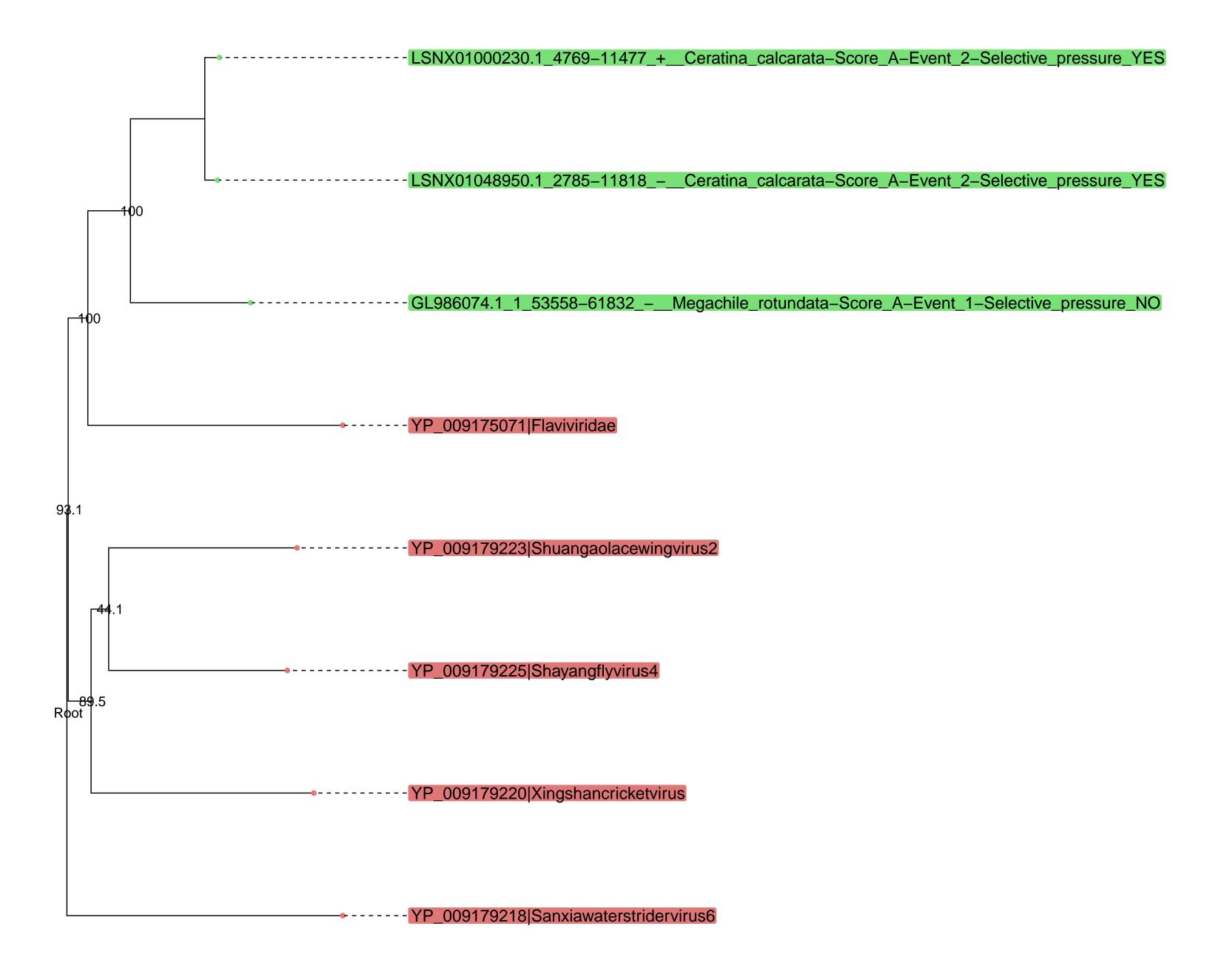


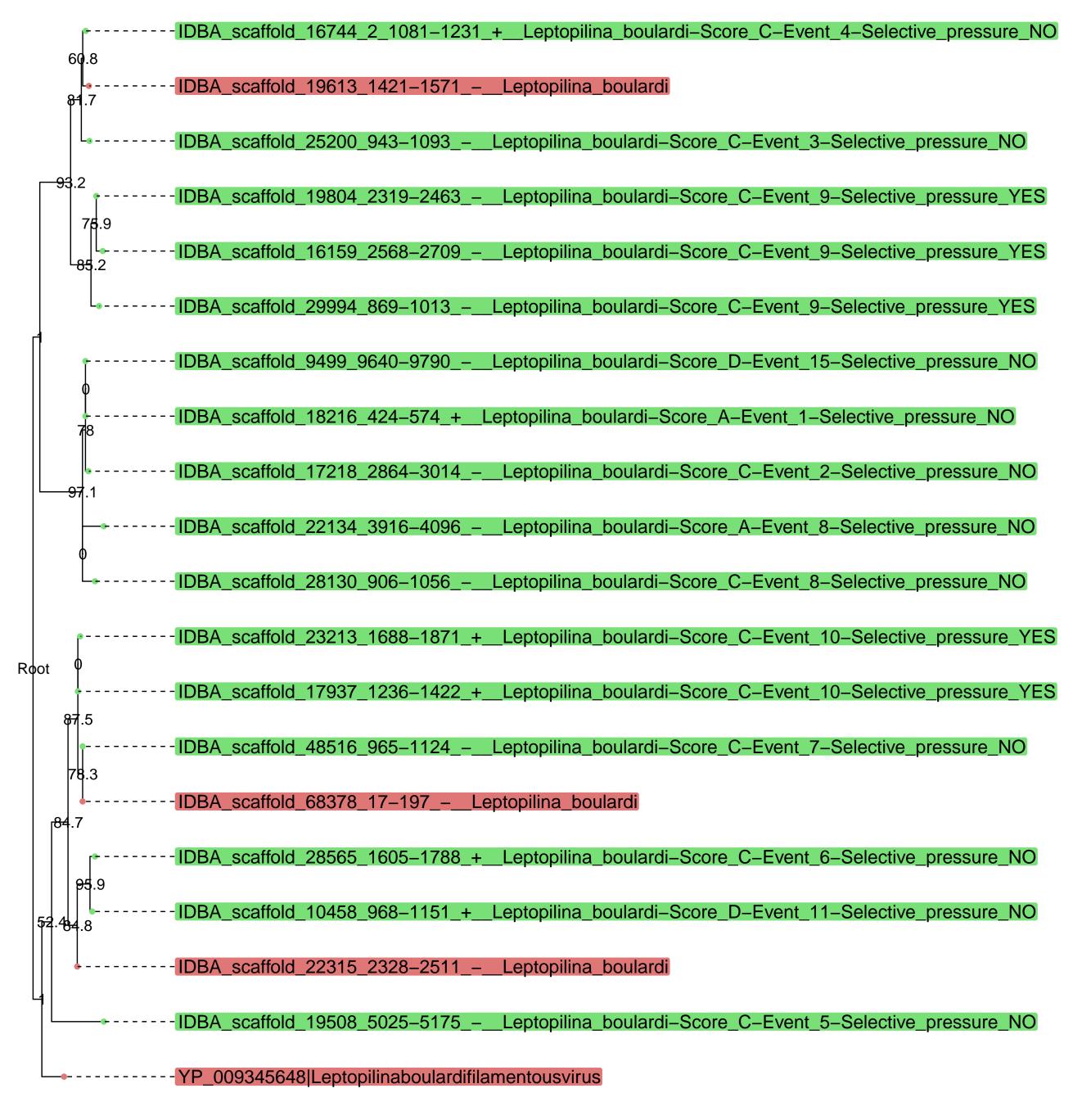


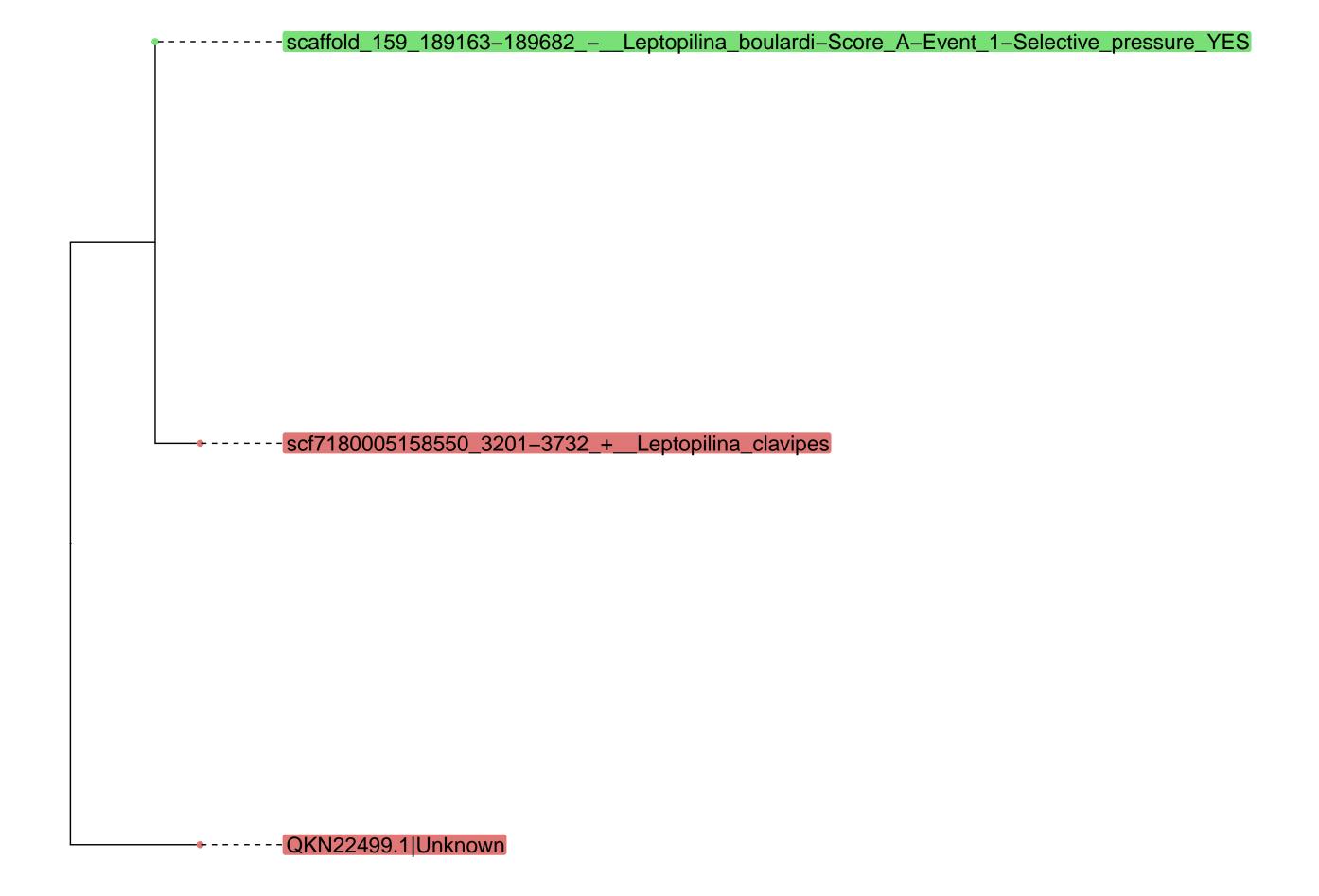


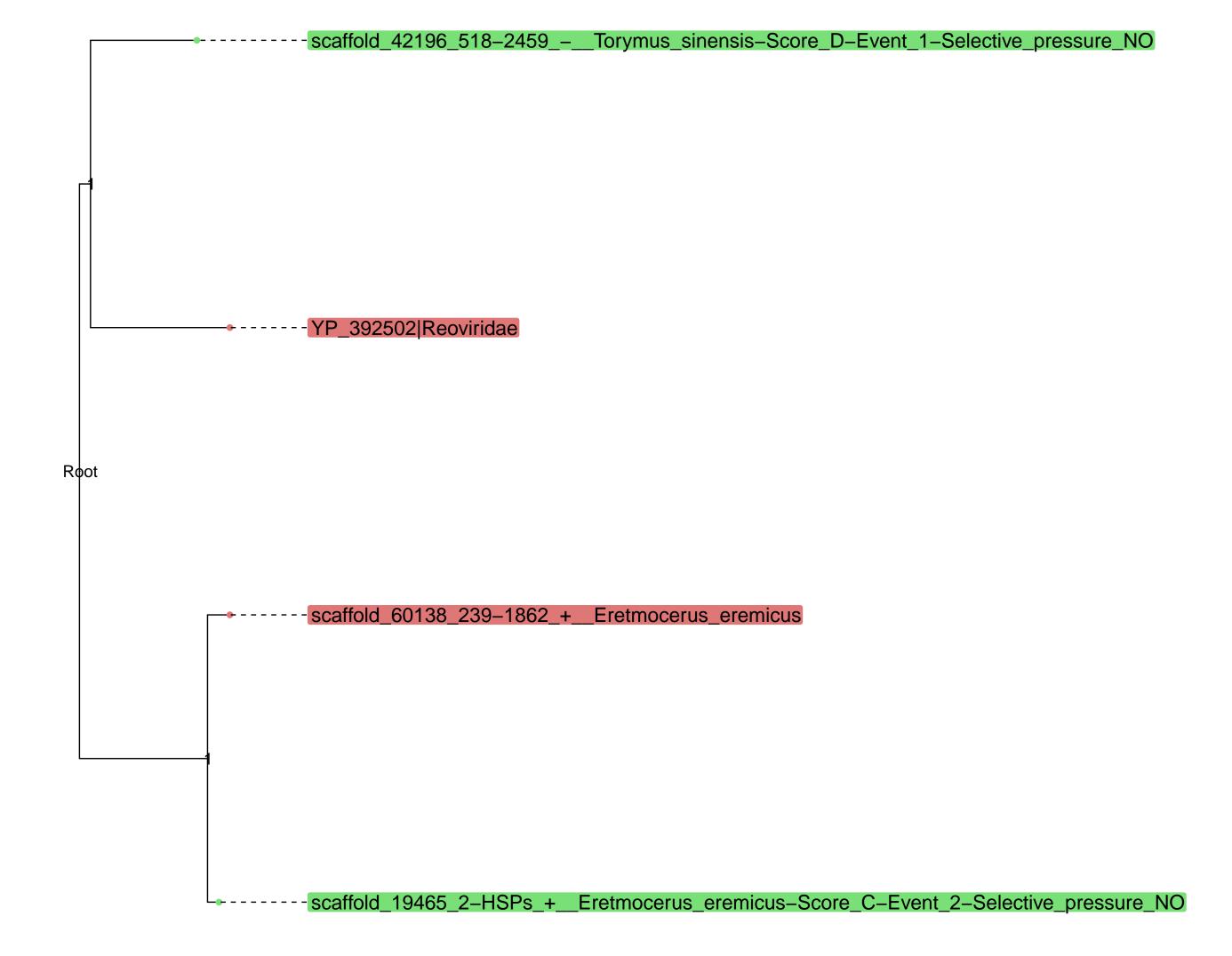


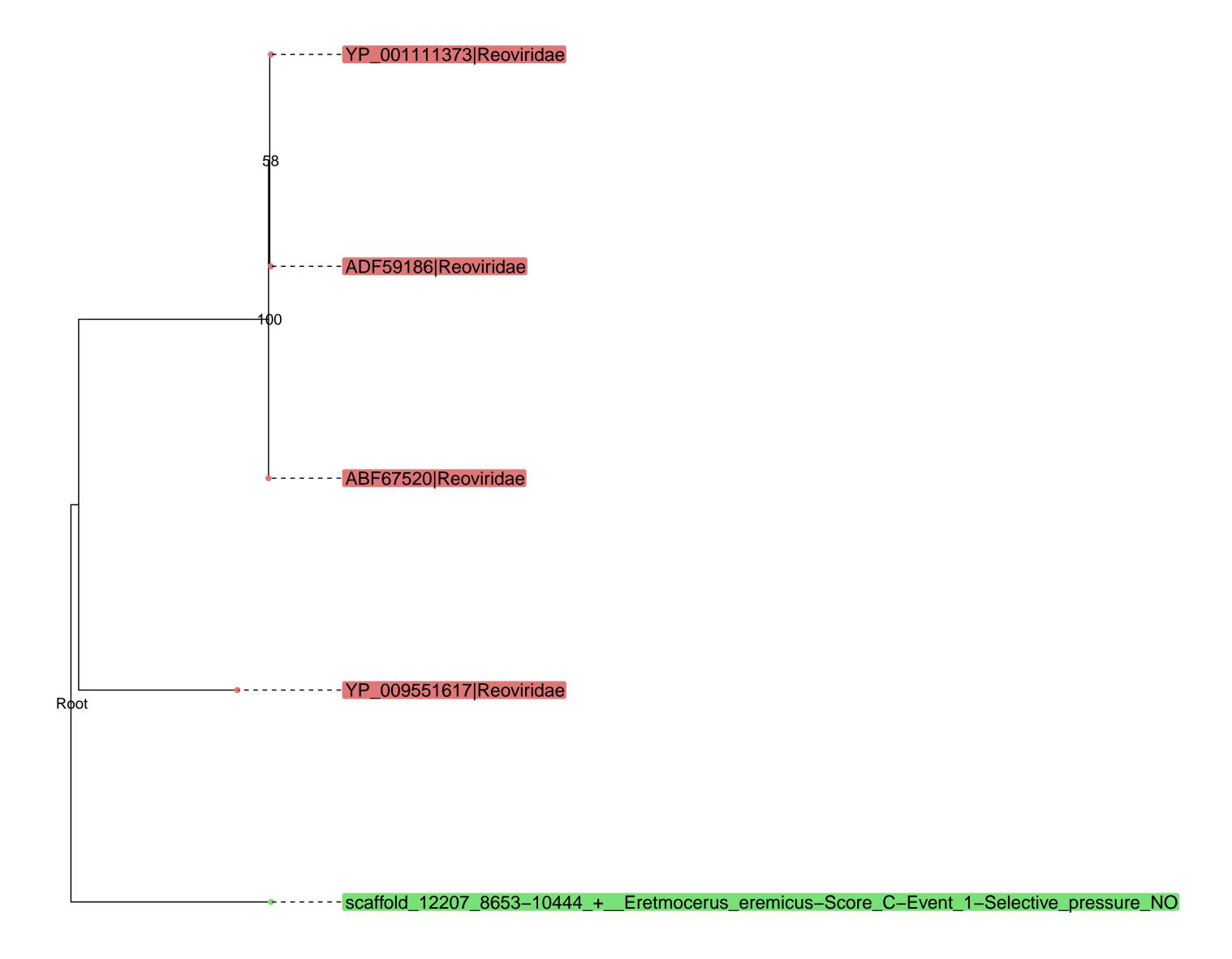


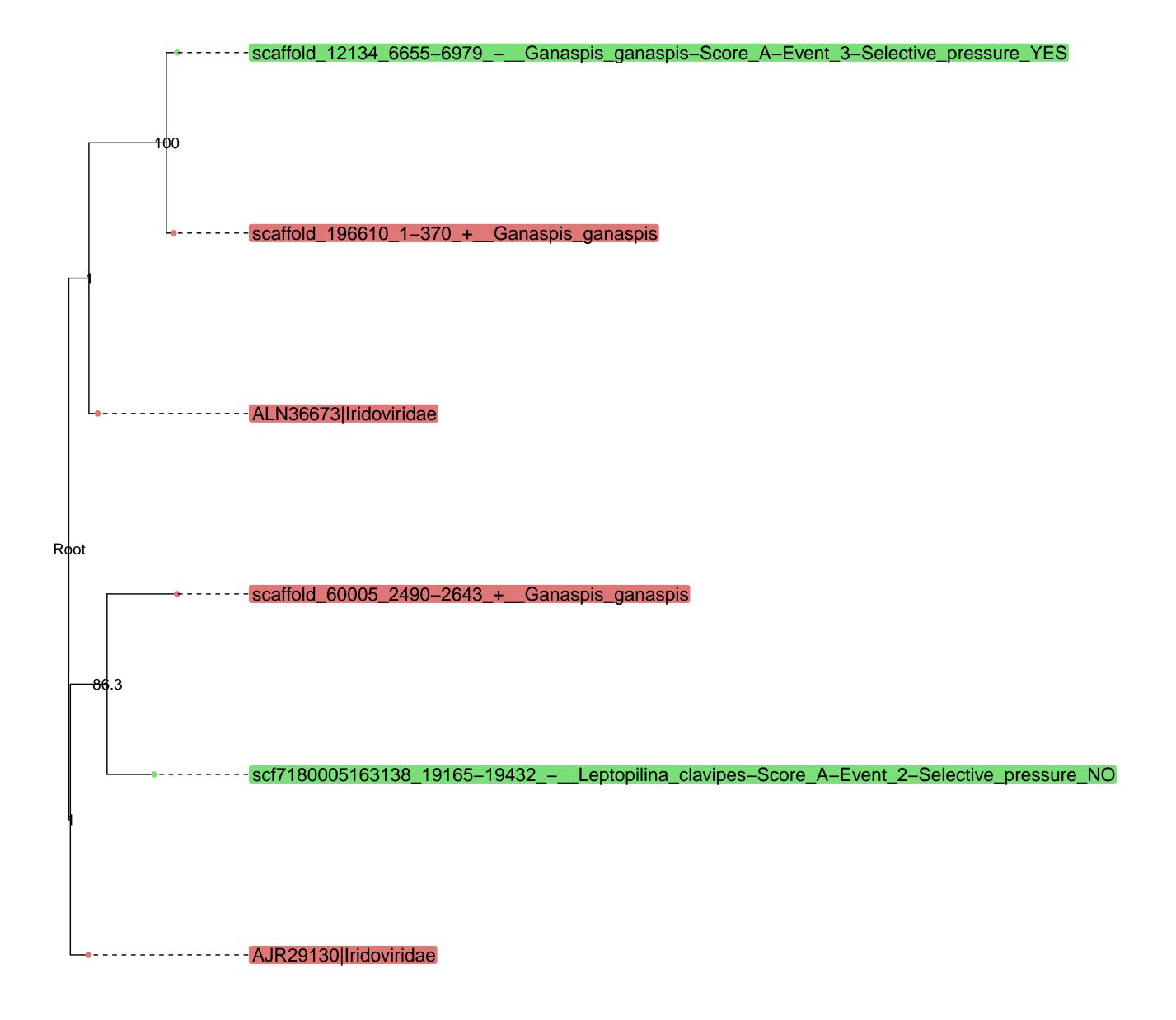


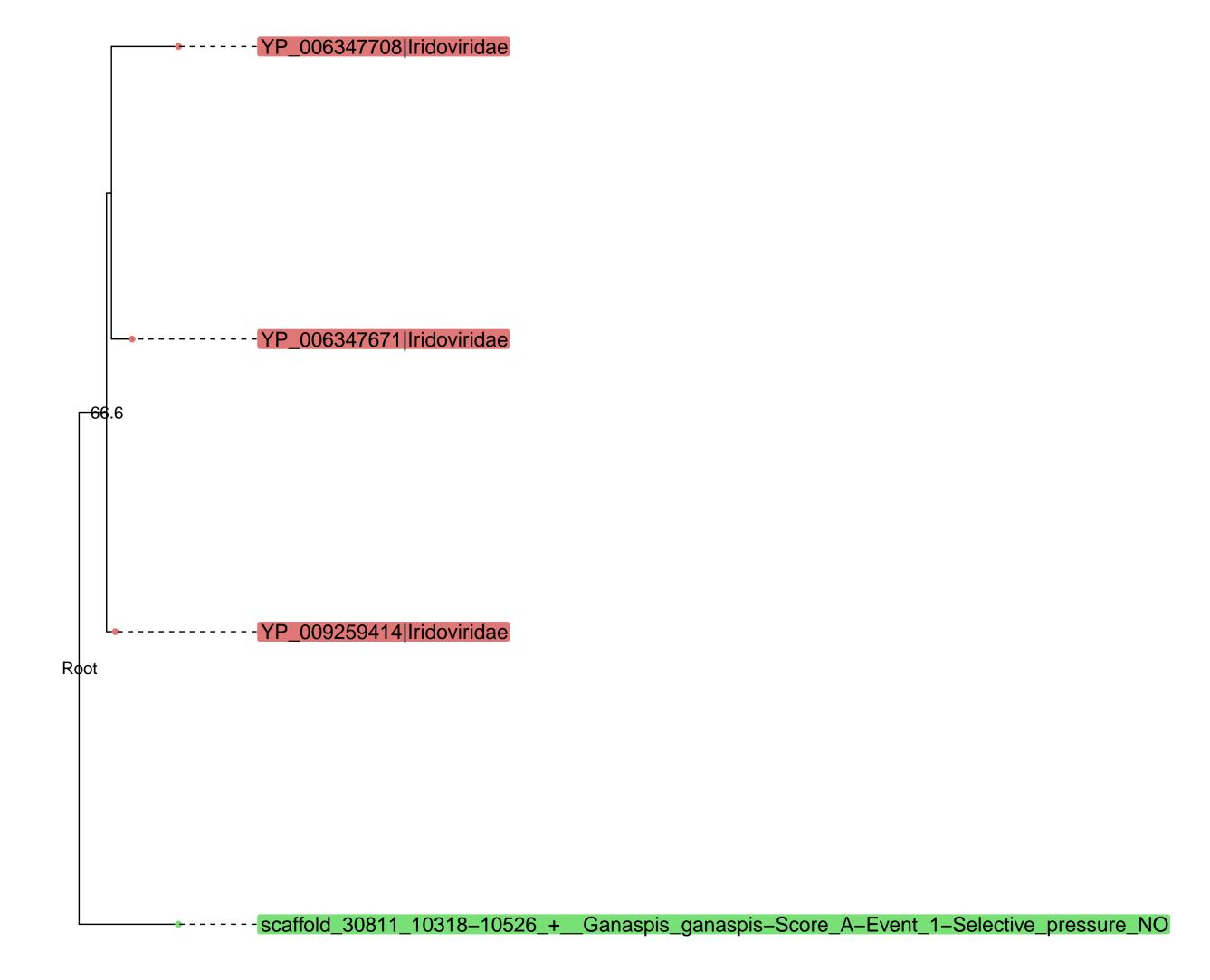


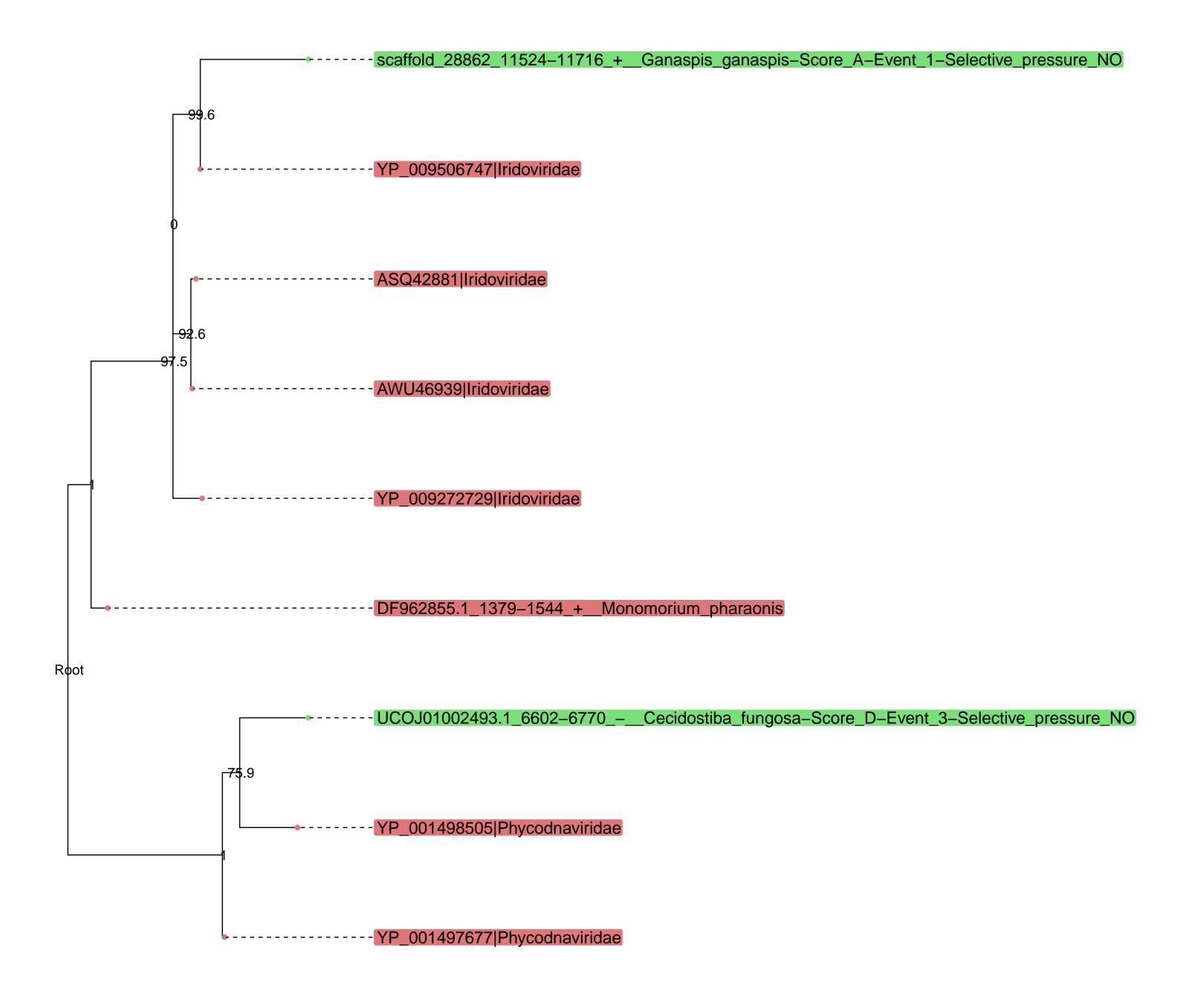


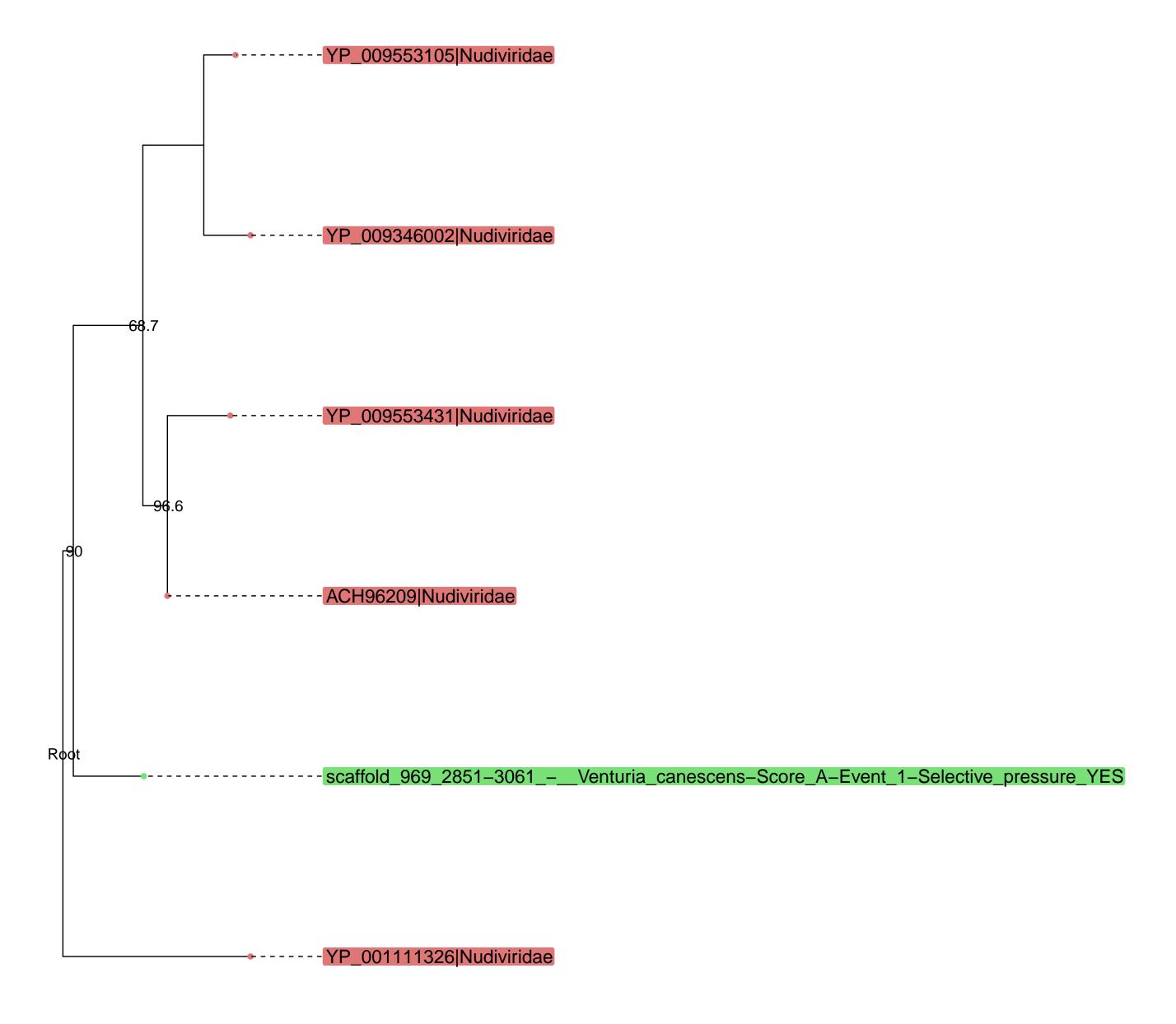


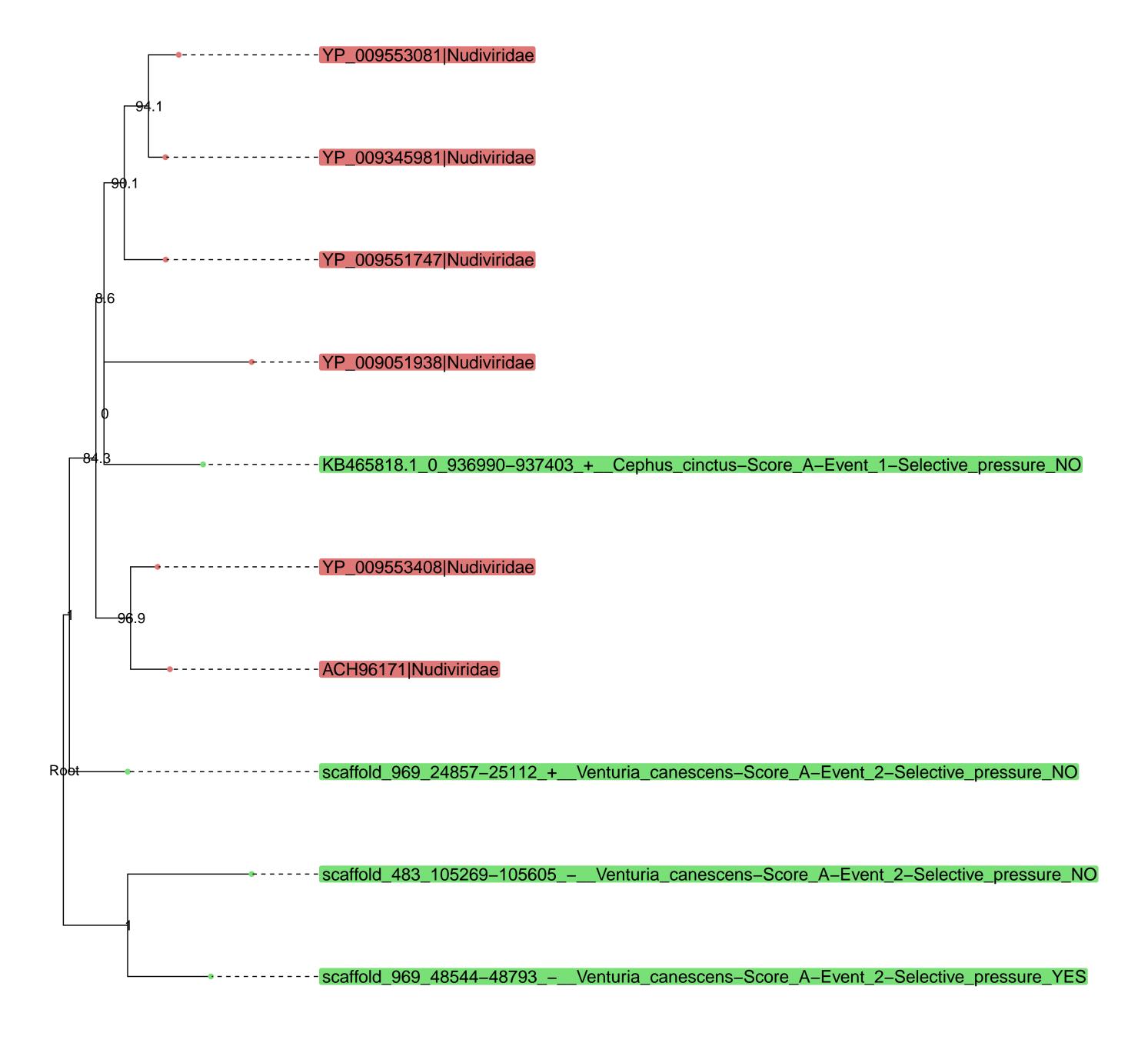


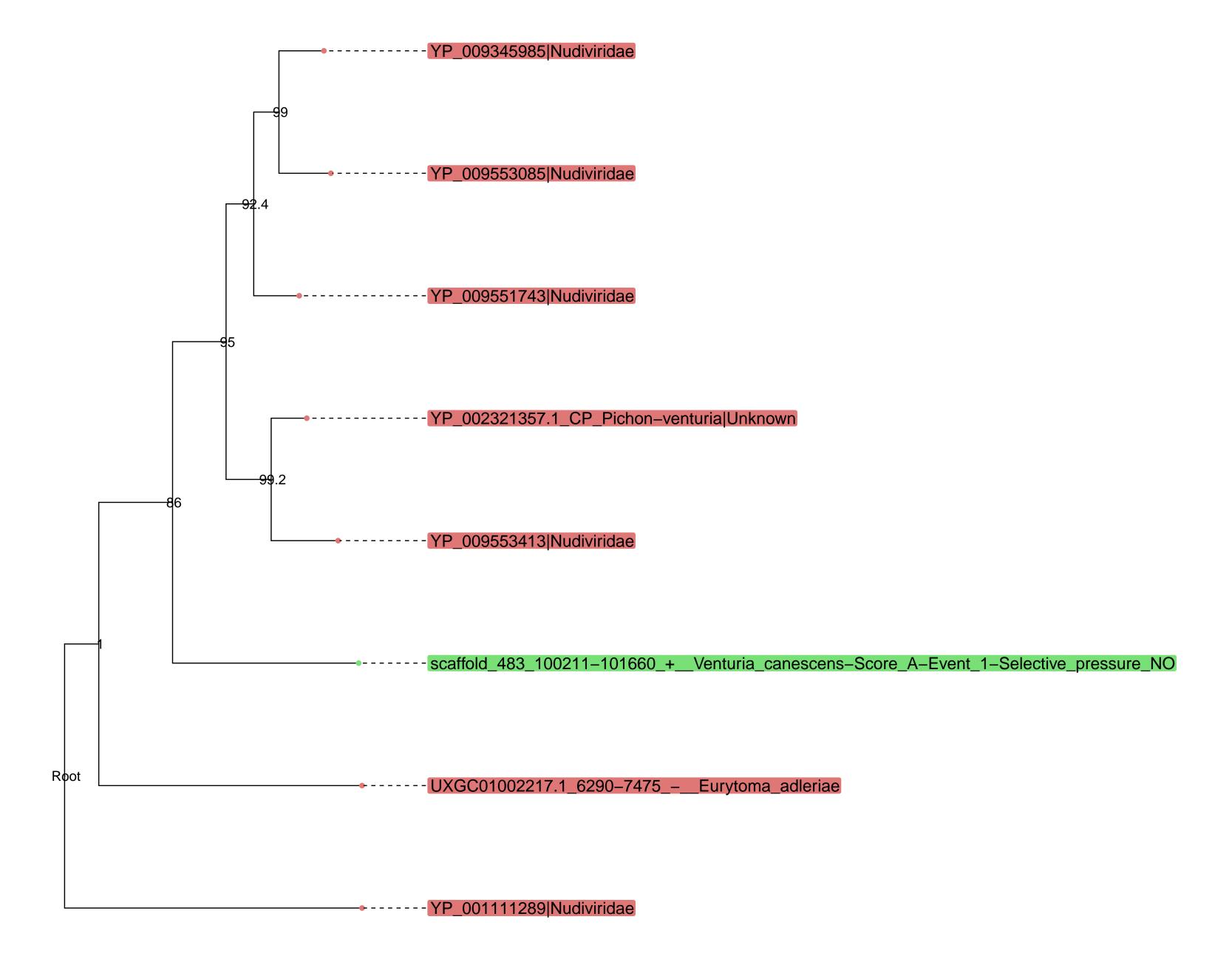


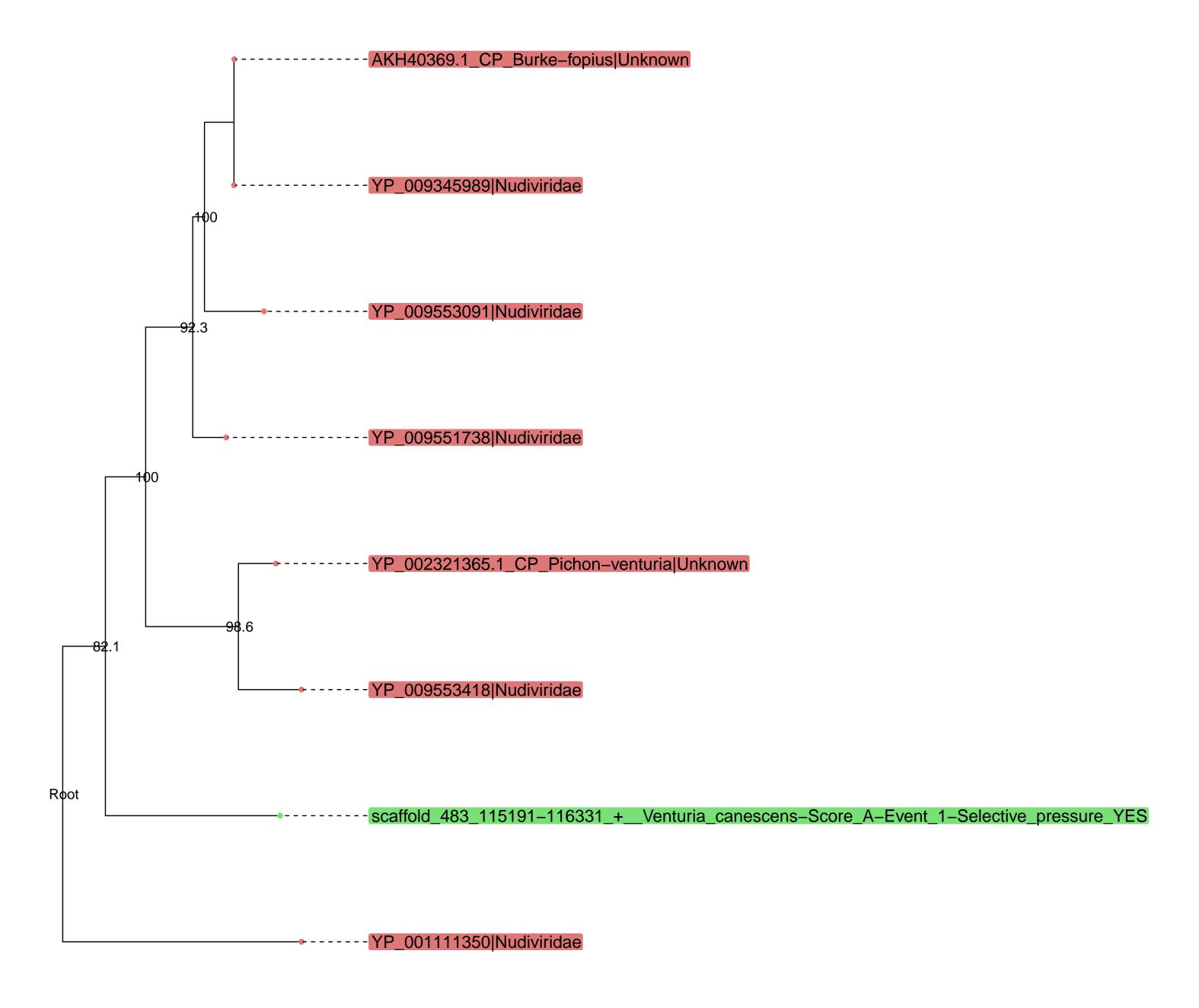


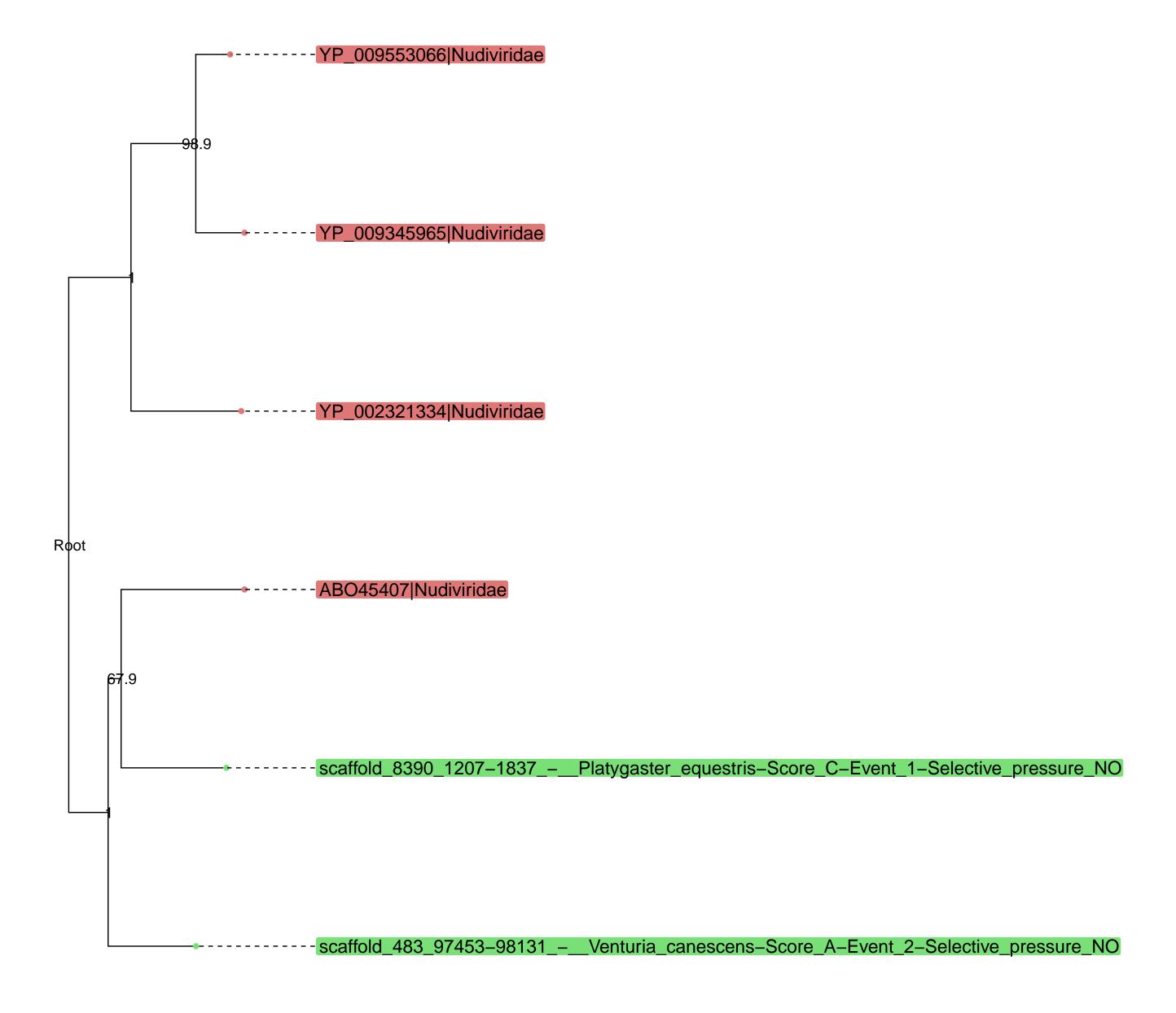


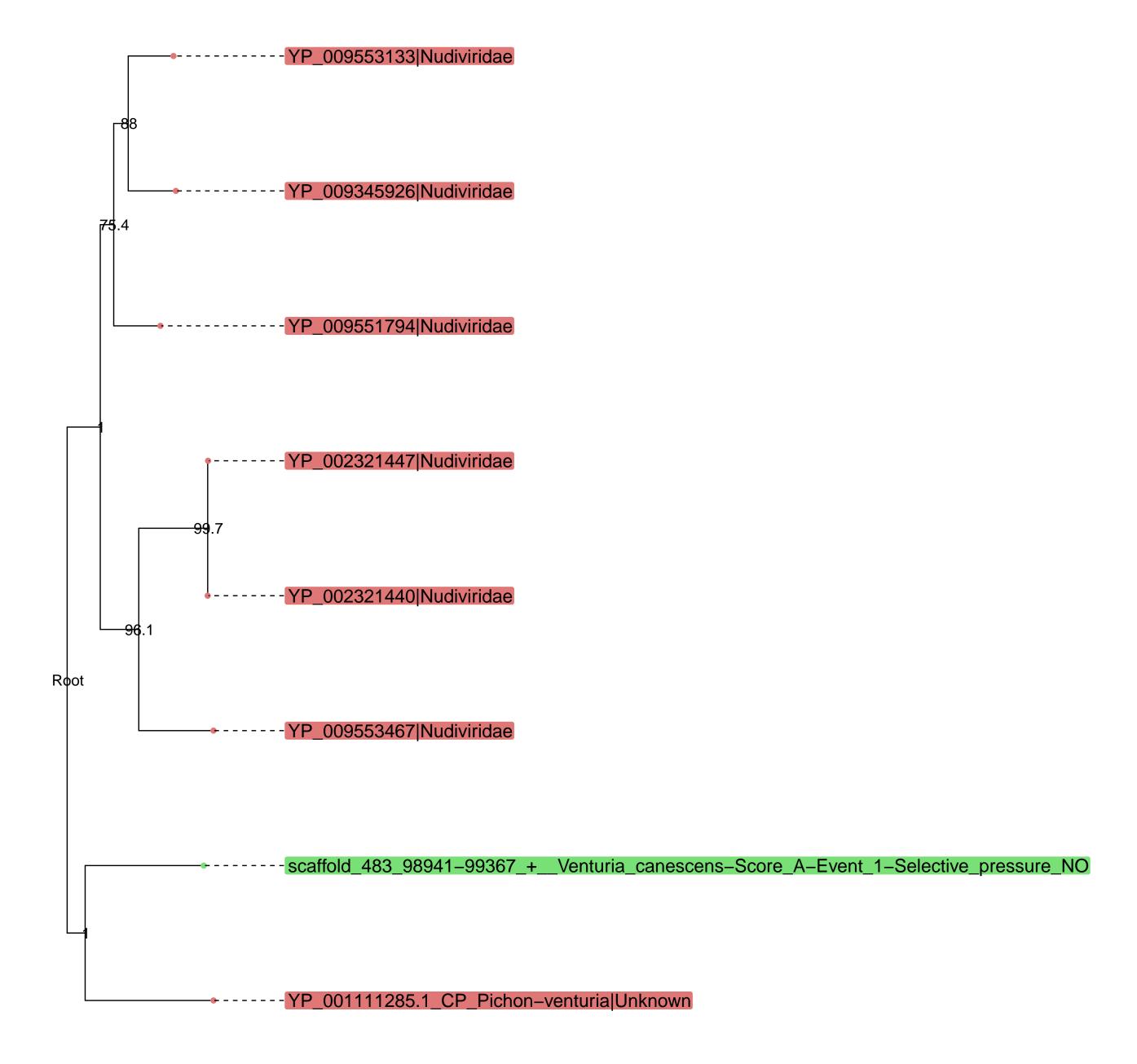


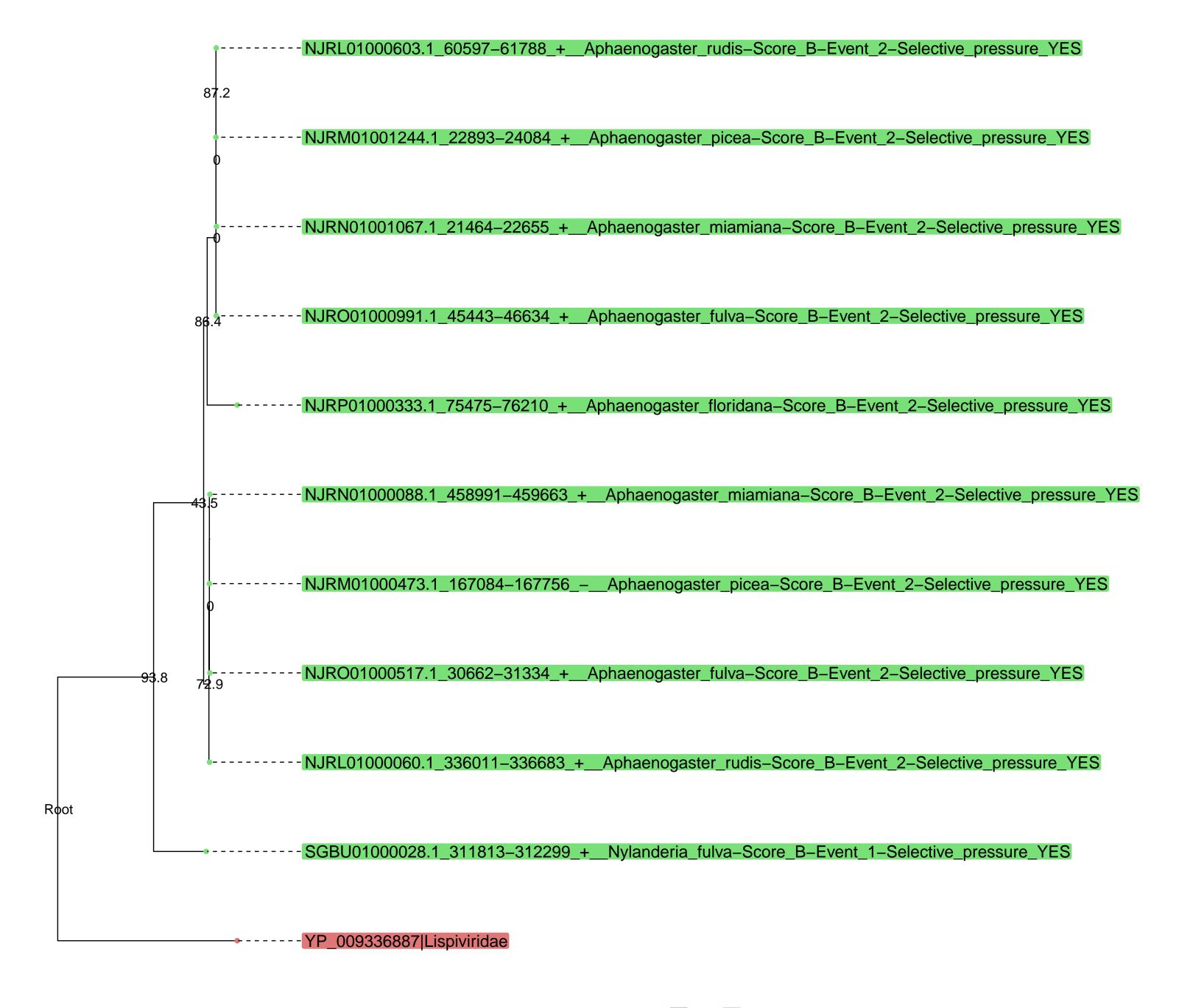


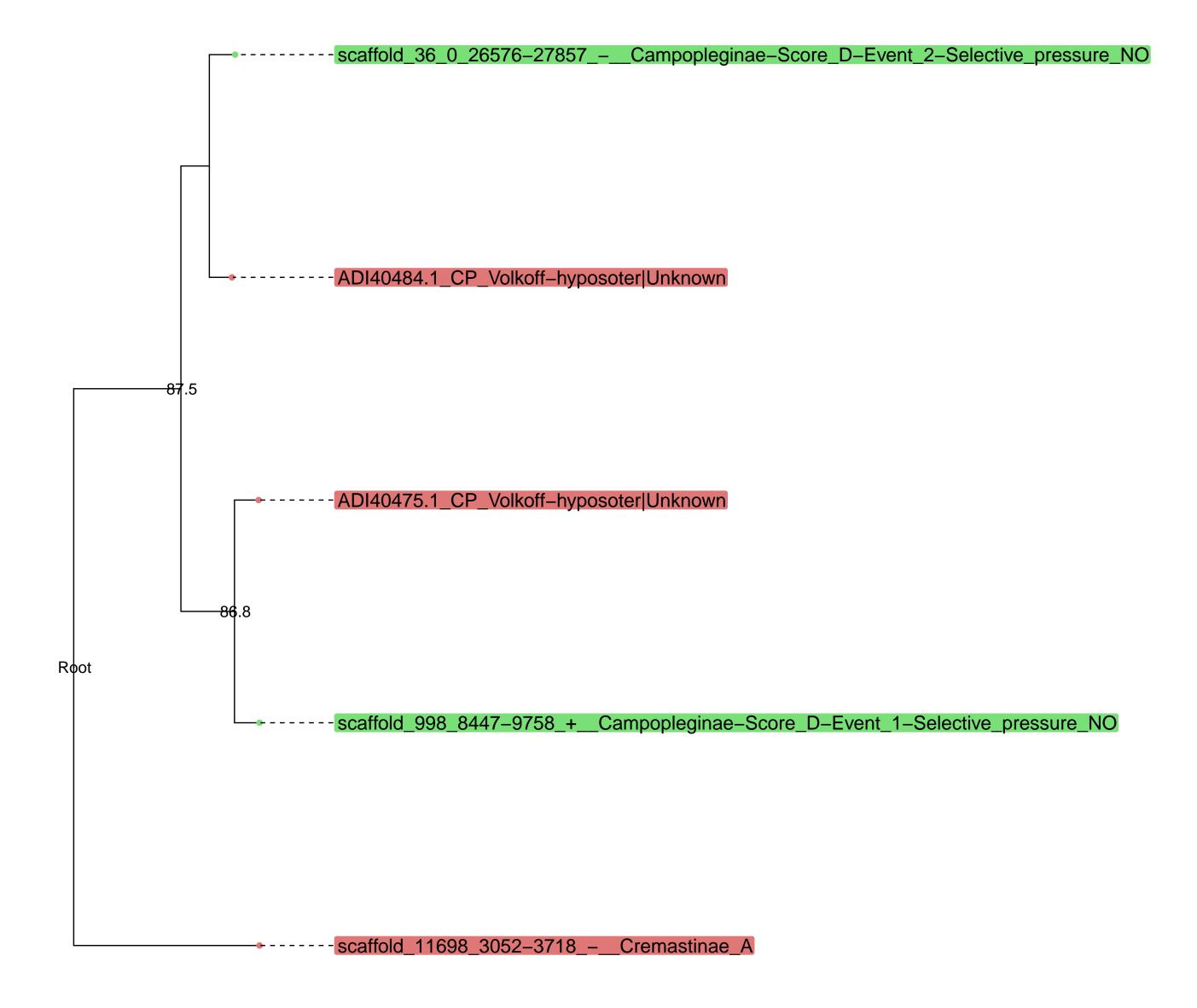






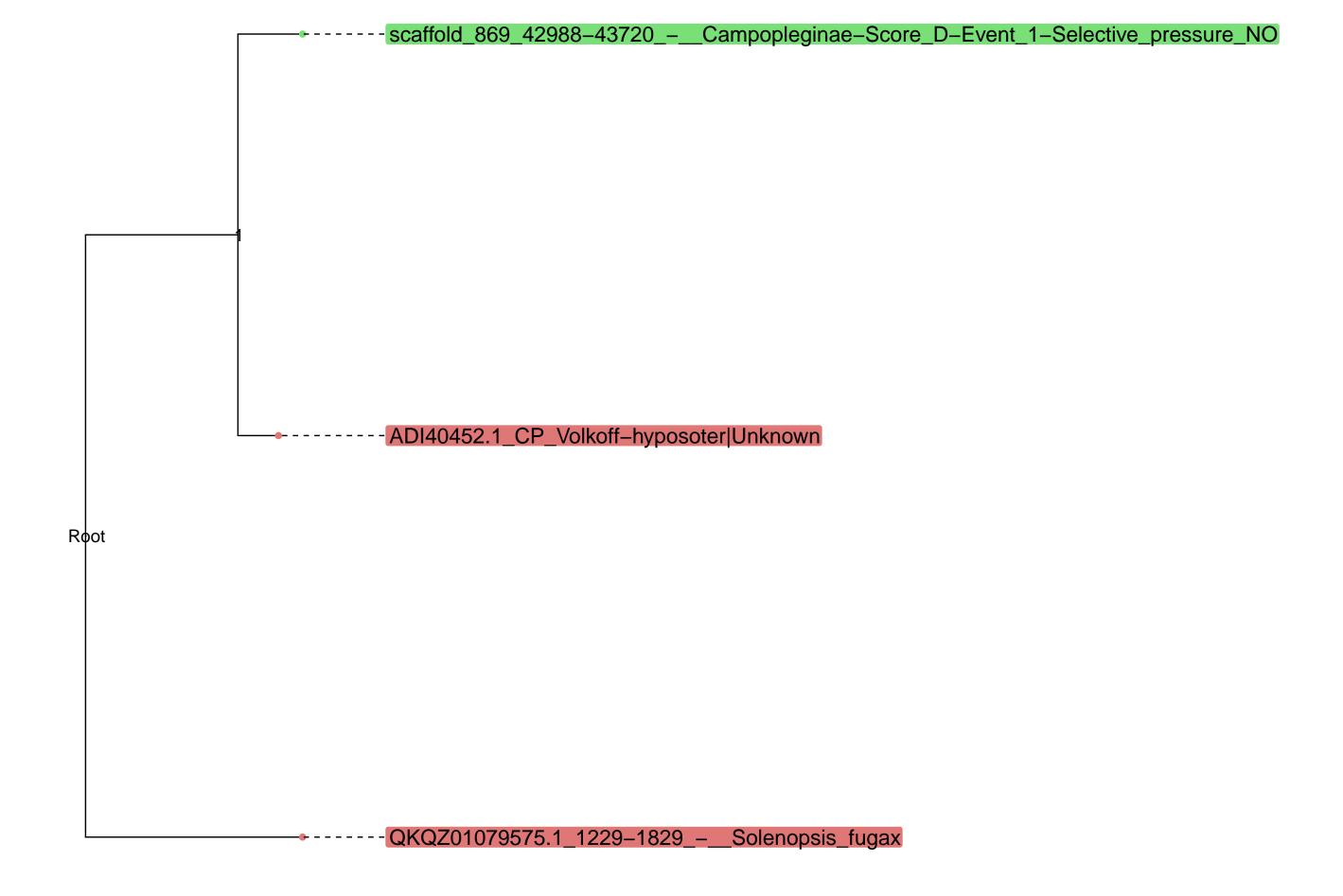




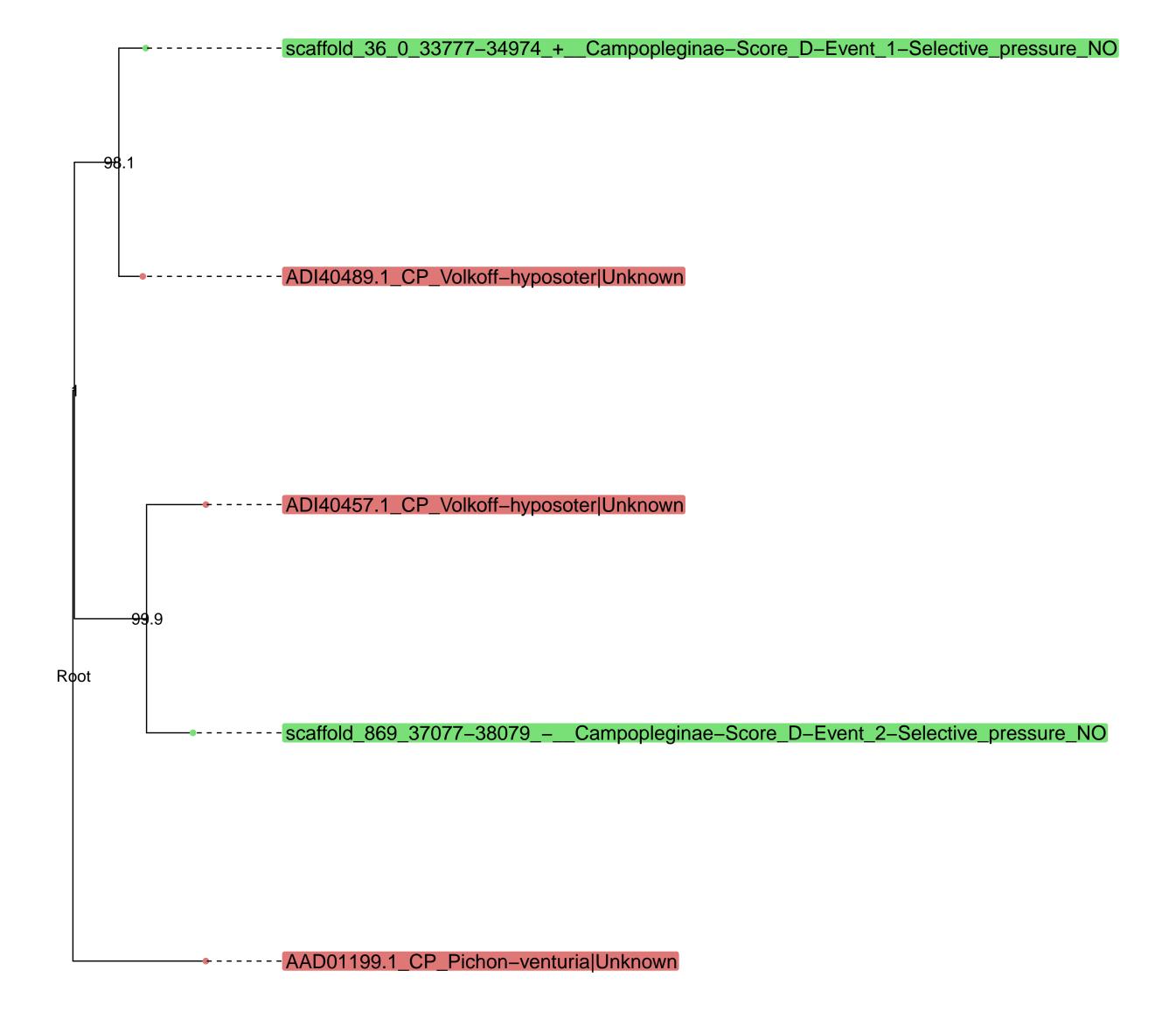


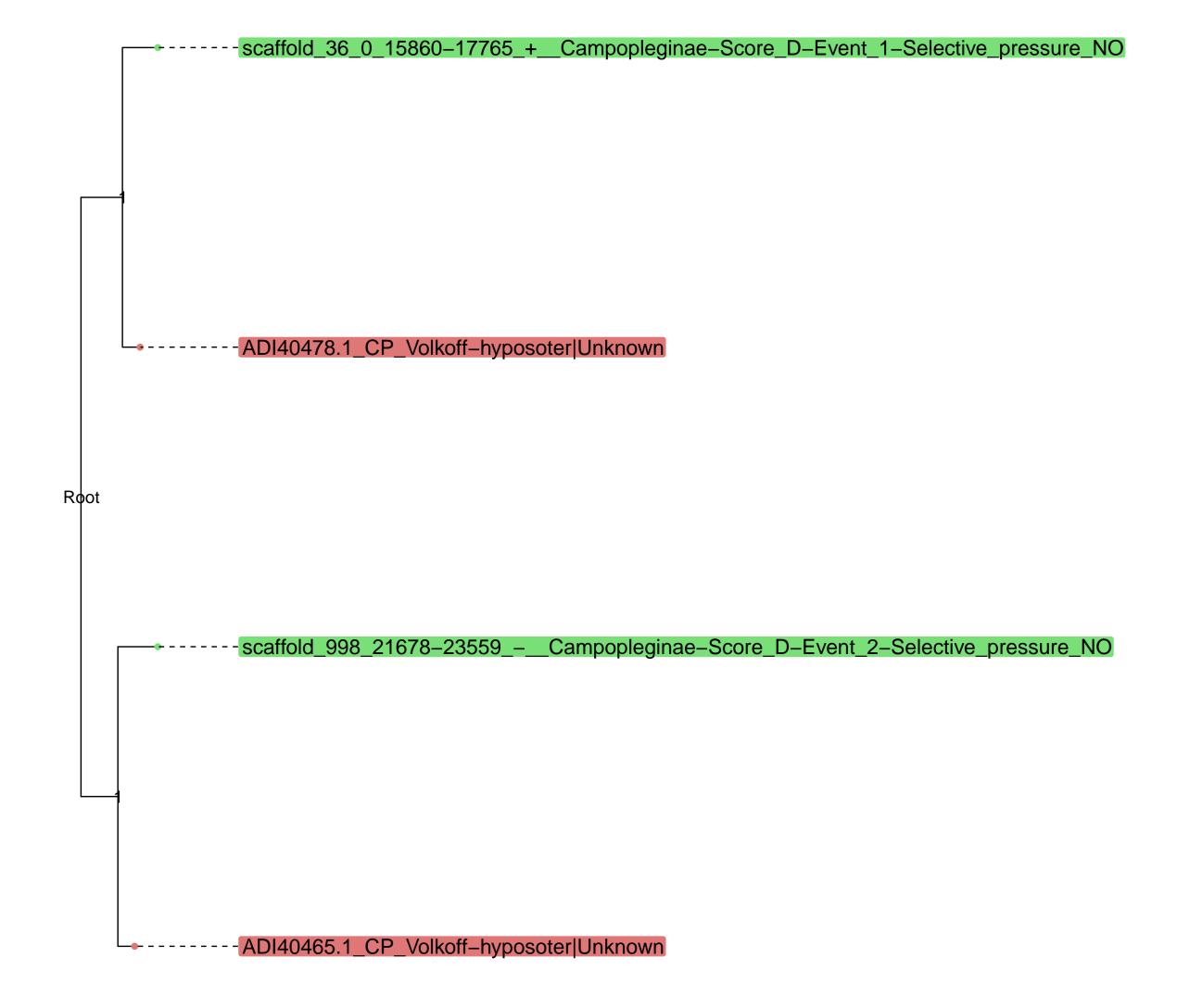


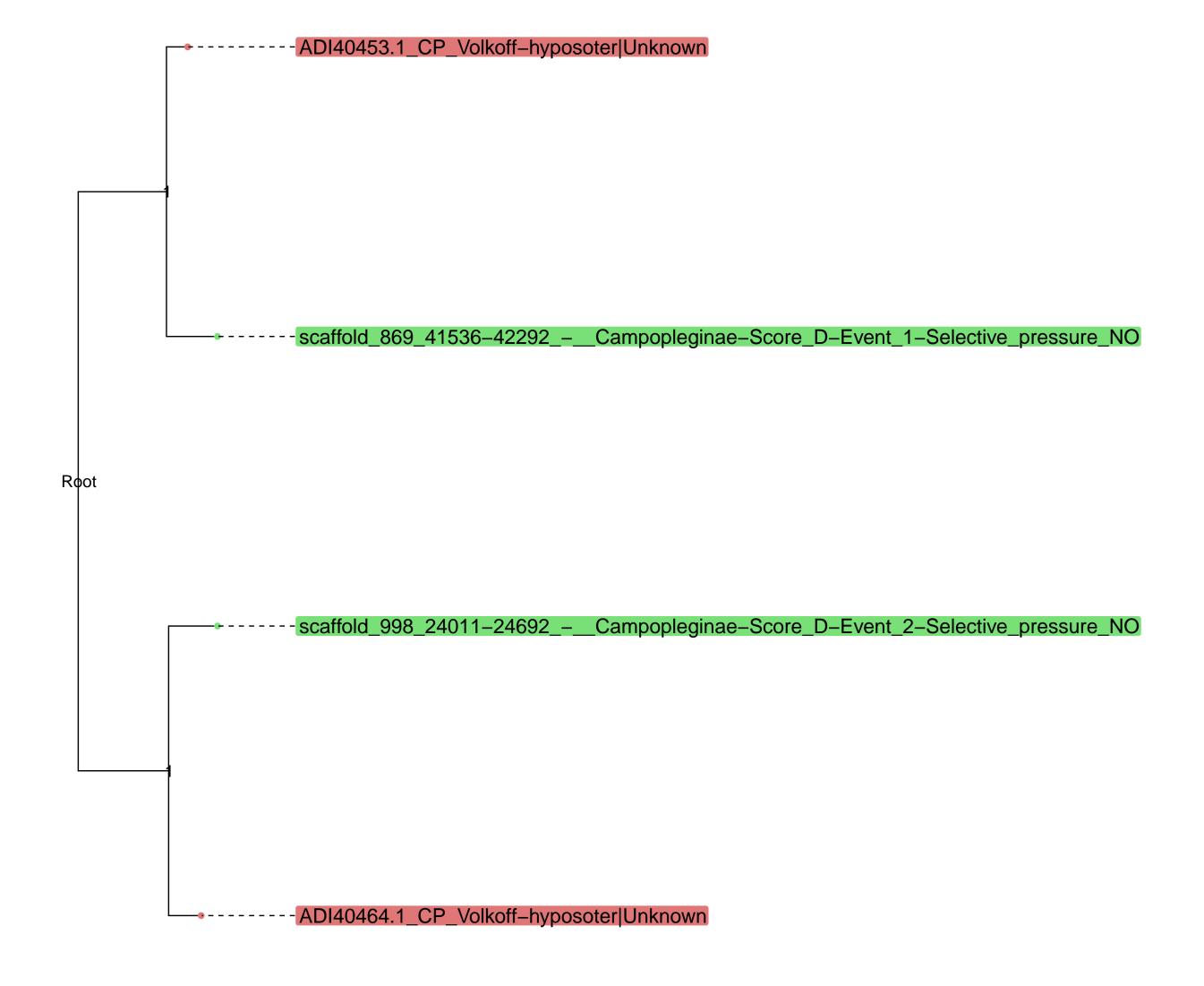




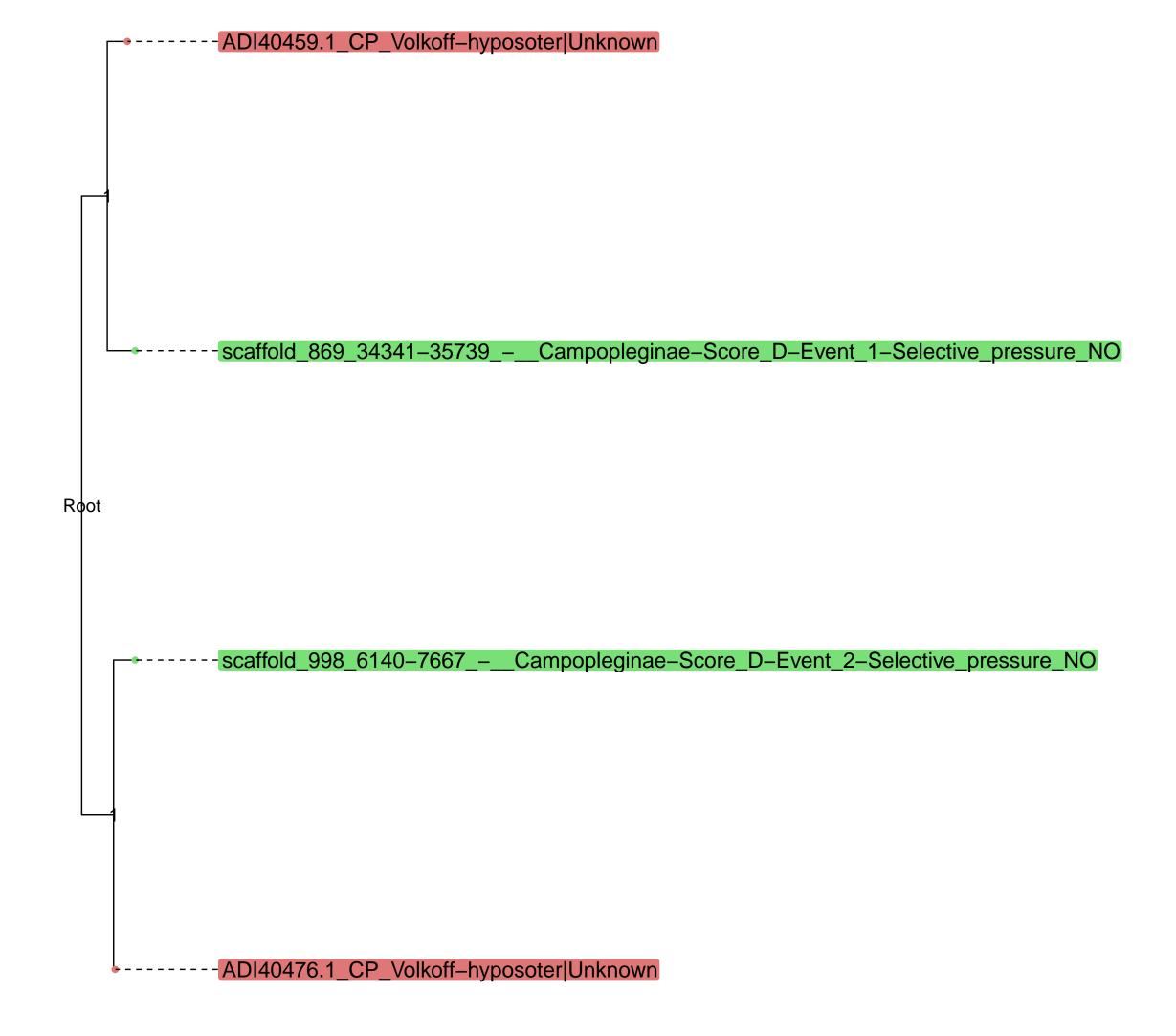
Cluster20806 – families involved : ['IVSPERs'] – putative protein : ['Cholesterol oxidase, substrate-binding']

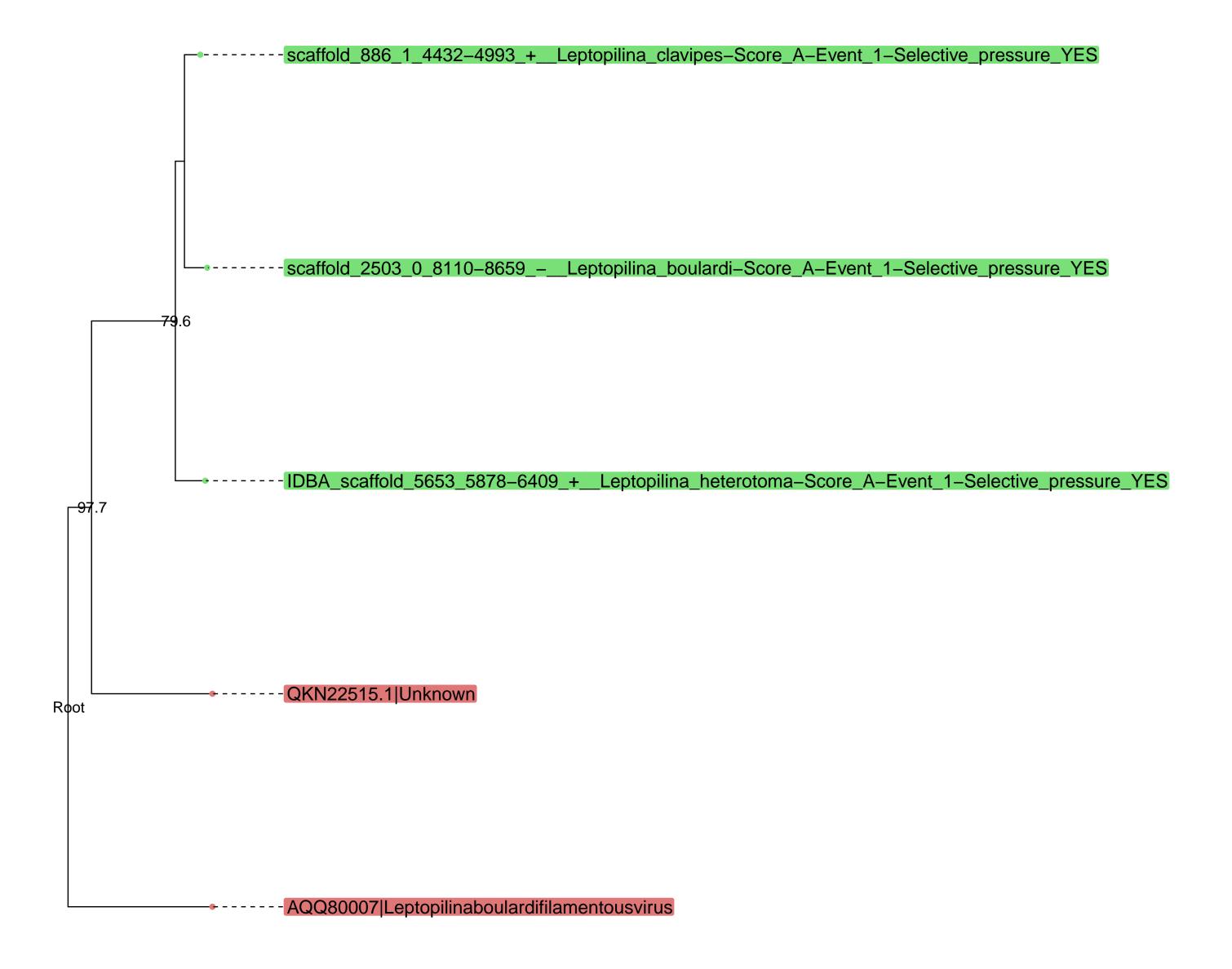




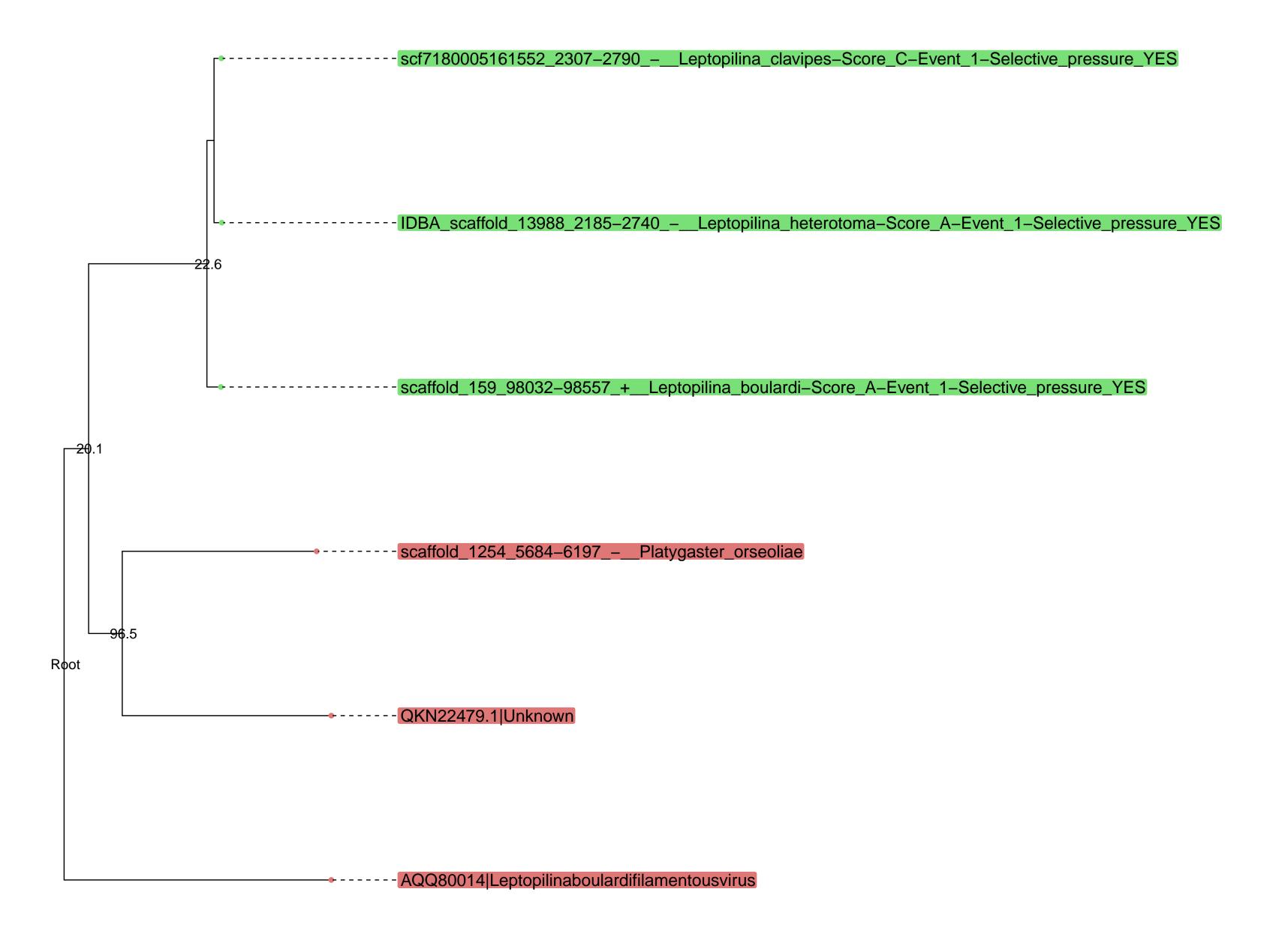


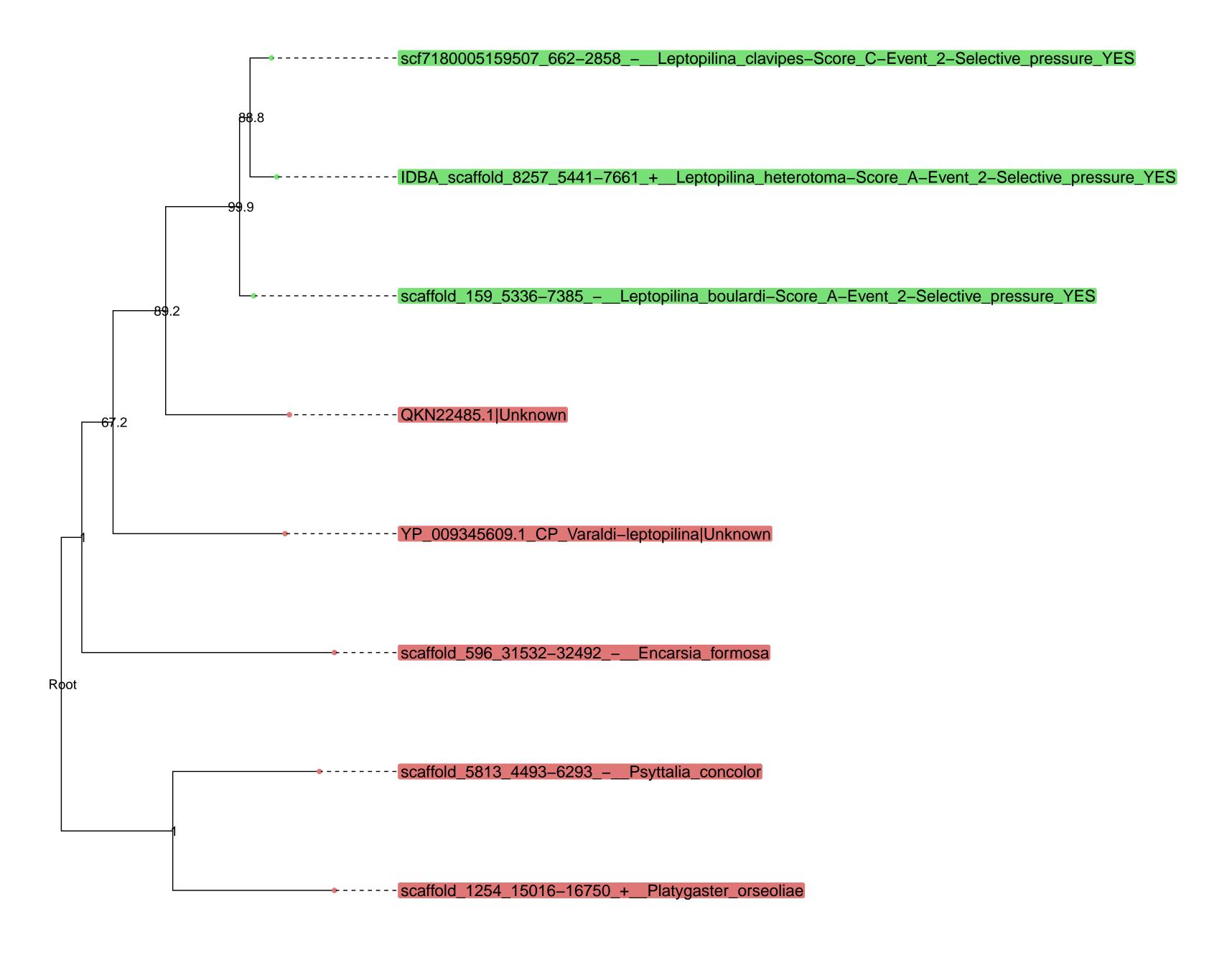
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                                 scaffold_143760_8-443_-_Platygaster_orseoliae
                                 scaffold_4517_8-698_-__Trissolcus_brochymenae
                                 scaffold_44109_738-1464_-__Copidosoma_sp
                                 -scaffold_32204_373-1099_+__Encarsia_formosa
                                 scaffold_9707_1981-2707_-__Cremastinae_A-Score_B-Event_8-Selective_pressure_NO
                                 -scaffold_8336_1-244_+_Campopleginae
                                 scaffold_9552_2-233_+__Campopleginae
                                 -scaffold_50530_287-974_+__Trichopria_sp_970989
                                 scaffold_8353_250-1033_-__Trissolcus_japonicus
                                 -scaffold_54608_3-255_+__Platygaster_orseoliae
                                -scaffold_42187_659-992_-_Platygaster_equestris
                                 scaffold_20133_1061-1631_-_Anagyrus_pseudococci
                                 scaffold_6624_2-263_-_Campopleginae
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                                 CM009916.1_13_878767-879412_-__Neodiprion_lecontei-Score_A-Event_14-Selective_pressure_NO
                                 KZ625021.1_2_614985-616080_-__Athalia_rosae-Score_A-Event_12-Selective_pressure_NO
                                 scaffold_36_0_37799-39269_+__Campopleginae-Score_D-Event_4-Selective_pressure_NO
                                 ADI40491.1 CP Pichon-venturia|Unknown
                                 scaffold_998_3230-4511_-__Campopleginae-Score_D-Event_5-Selective_pressure_NO
Root
                                 scaffold_869_32349-33729_-__Campopleginae-Score_D-Event_3-Selective_pressure_NO
     94.8
                                 ADI40460.1_CP_Volkoff-hyposoter|Unknown
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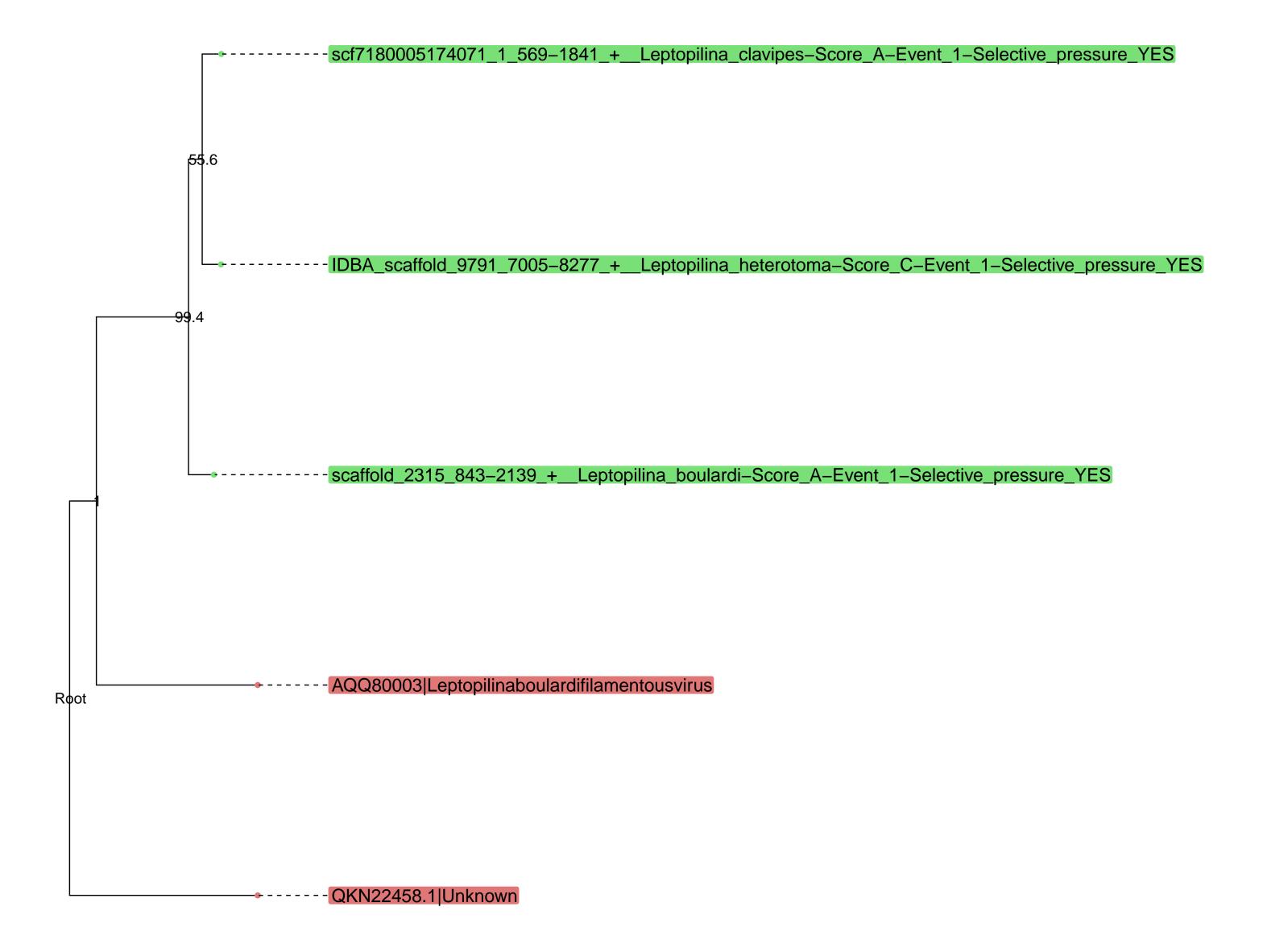


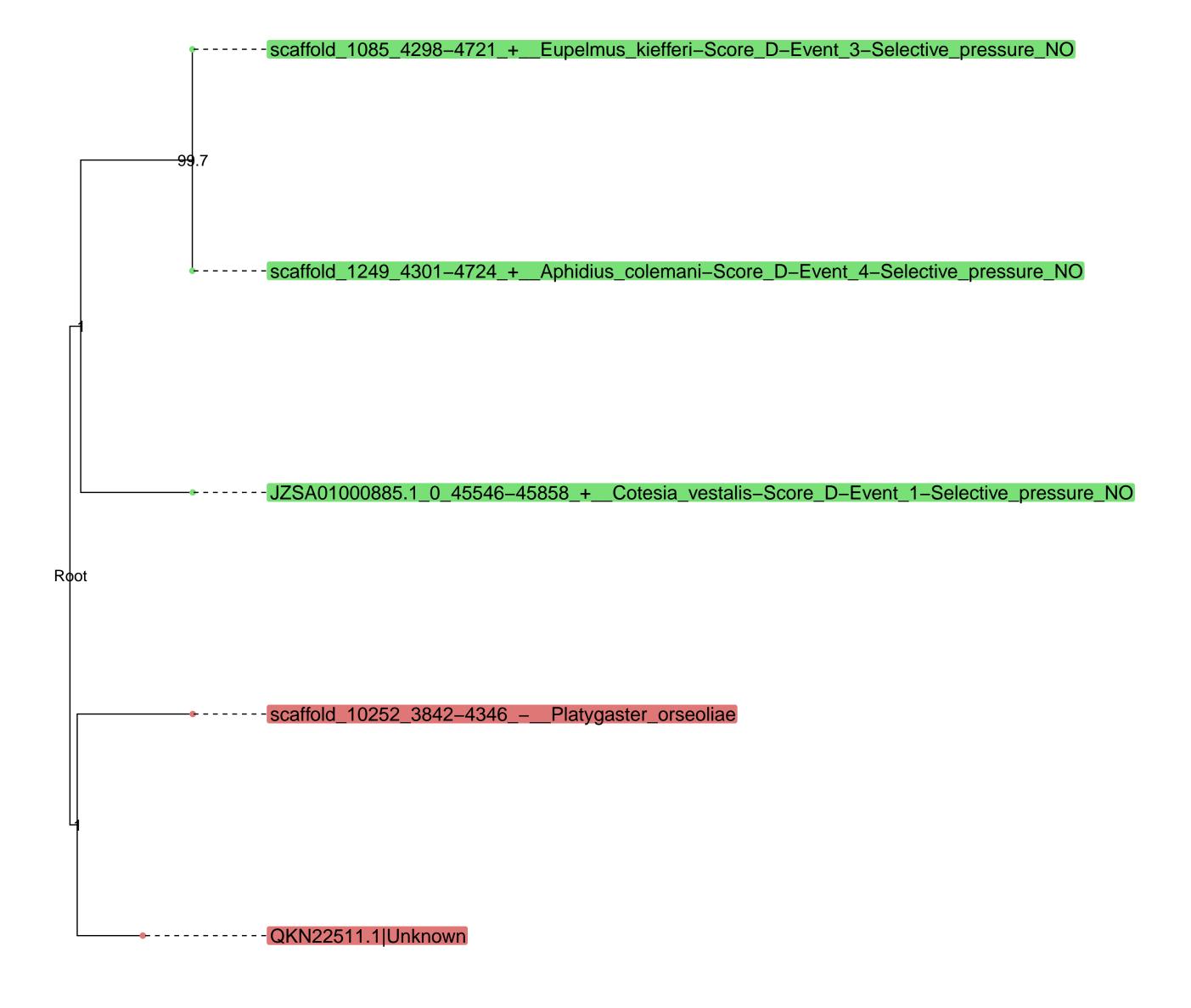


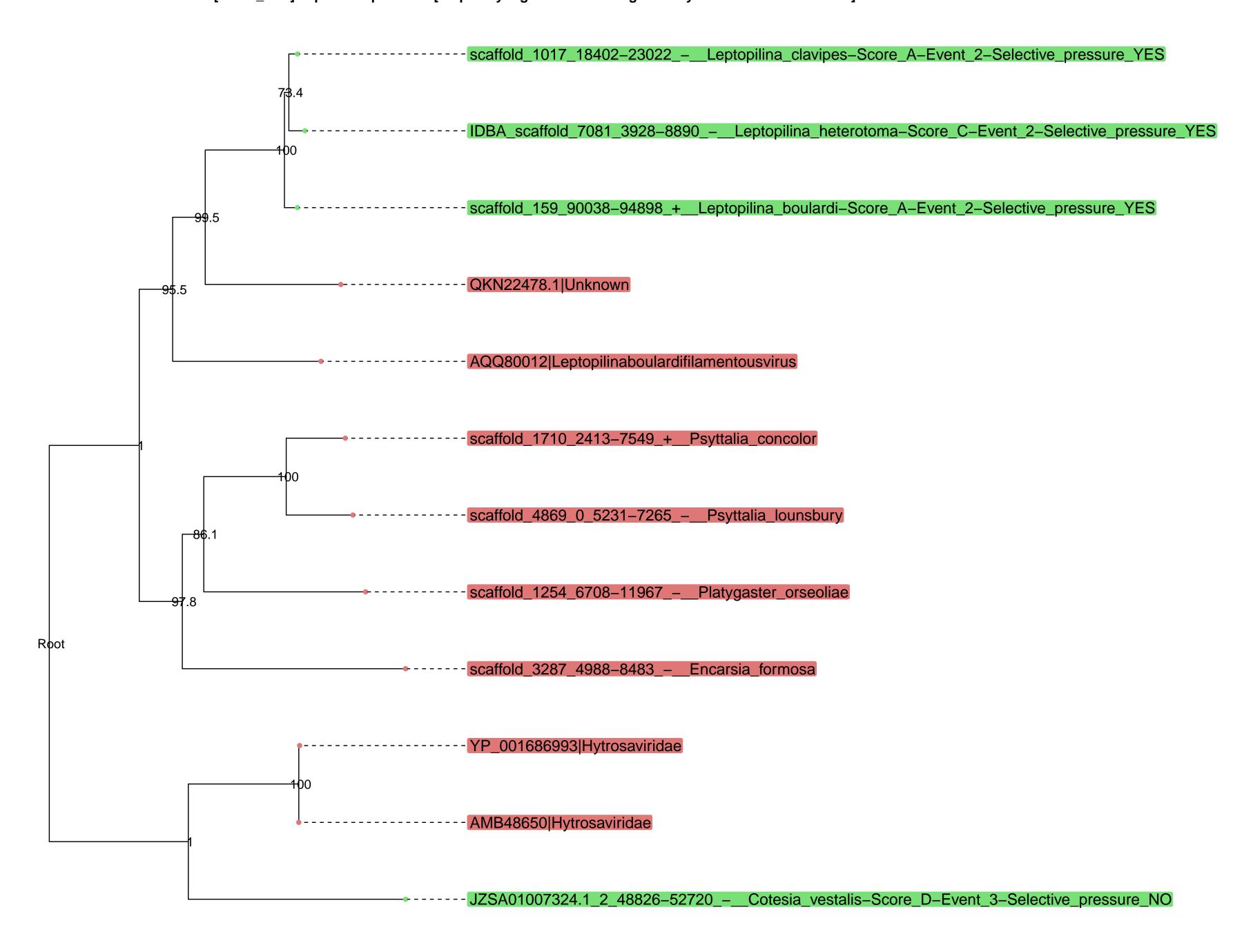


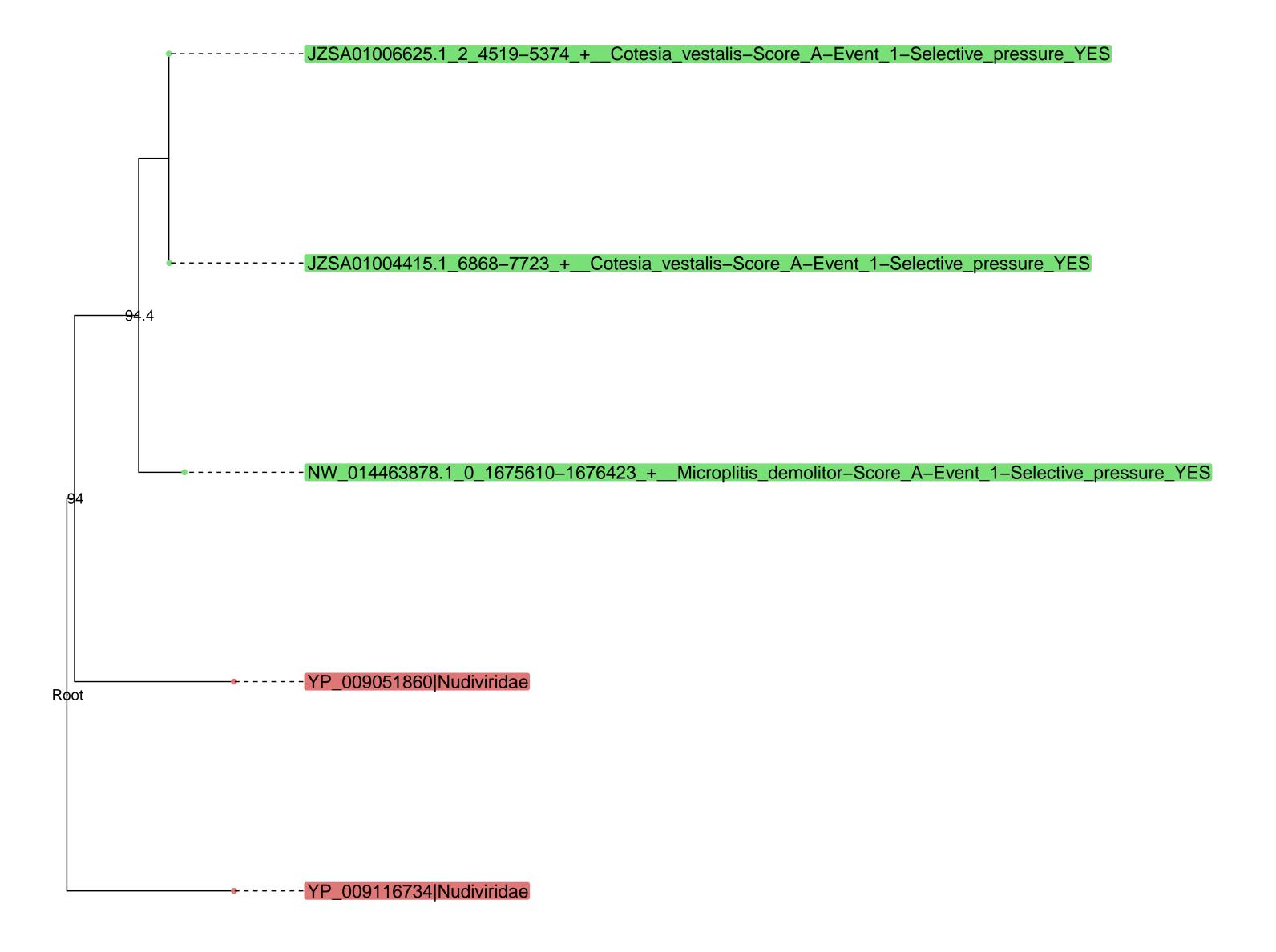


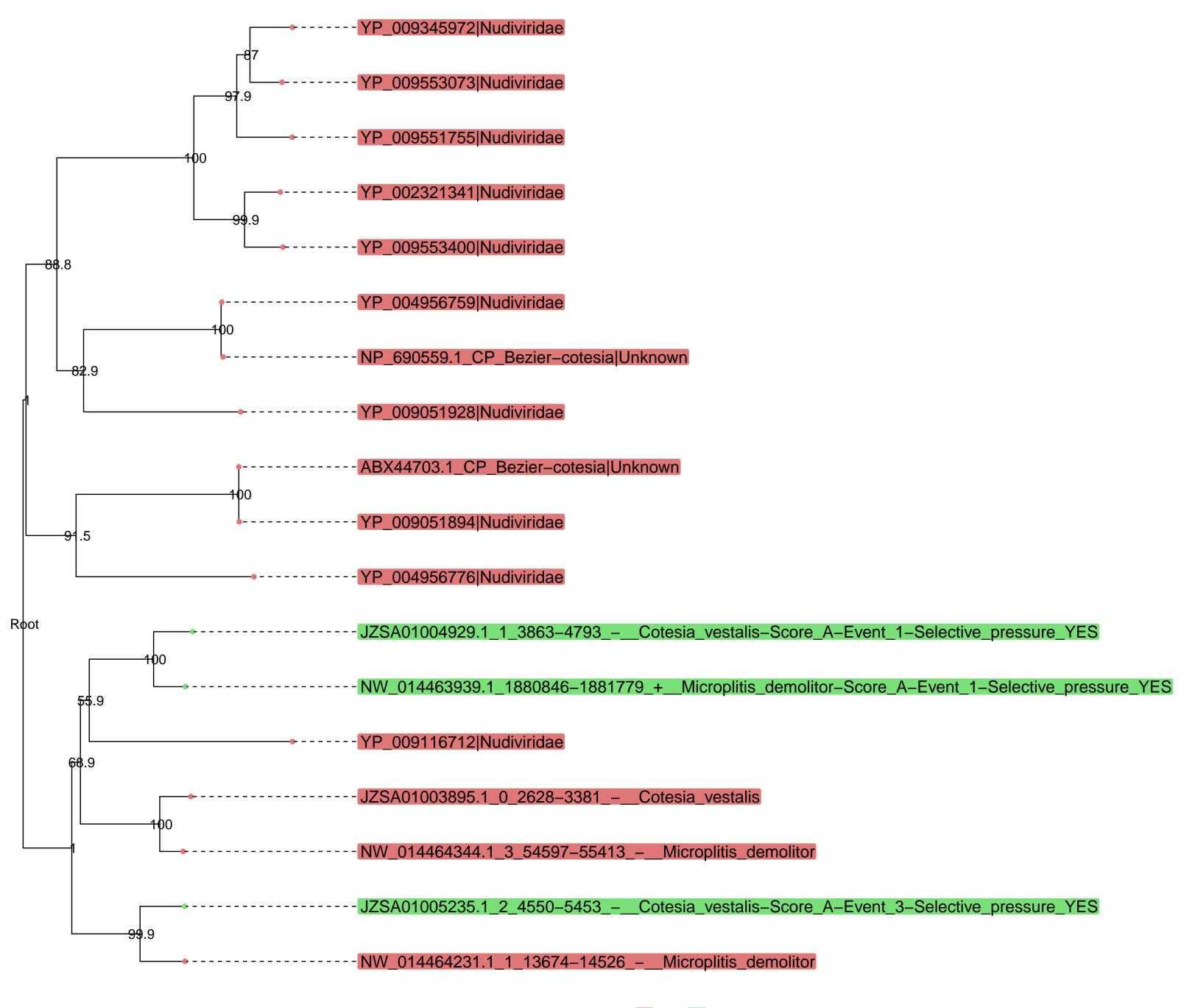




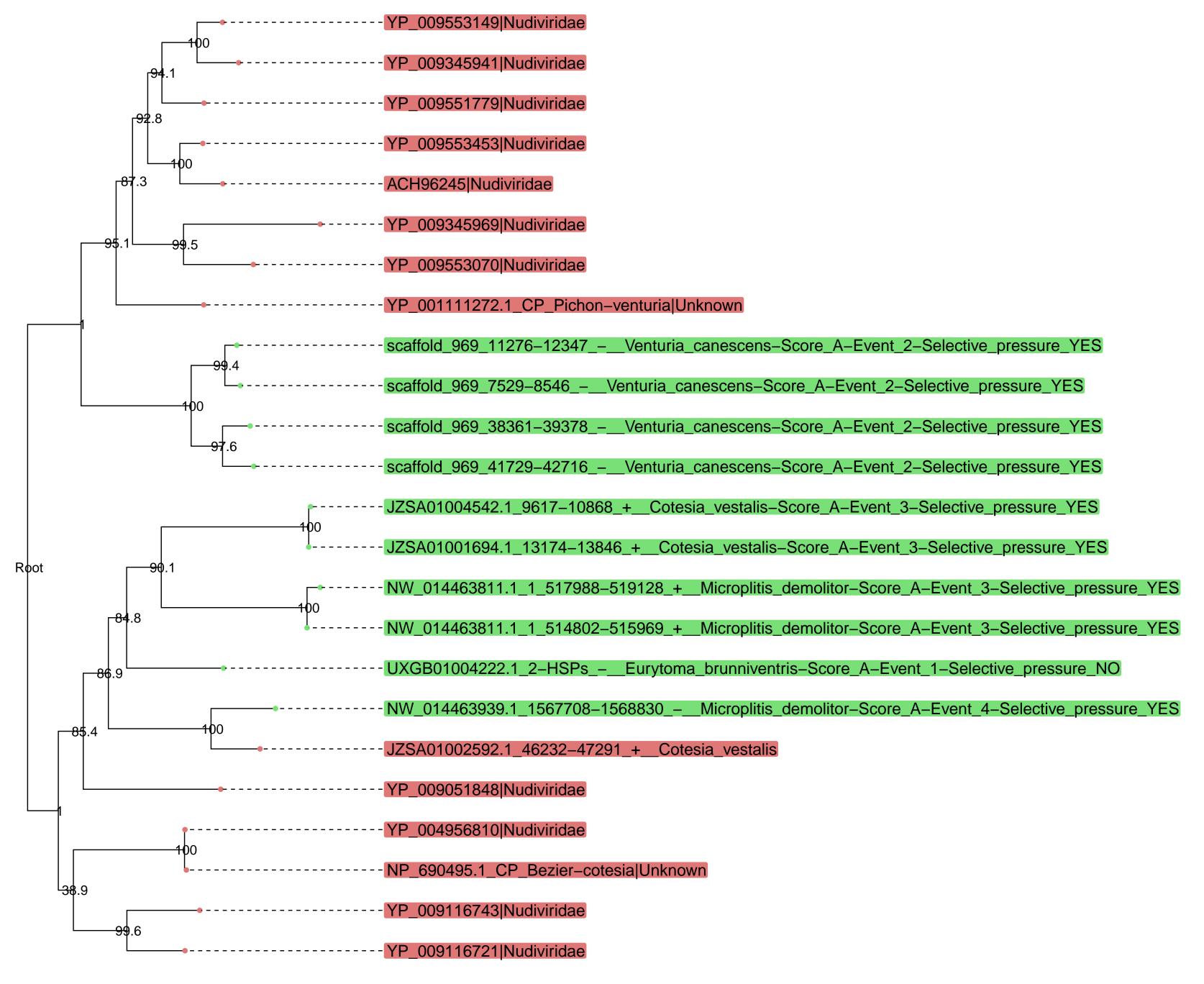


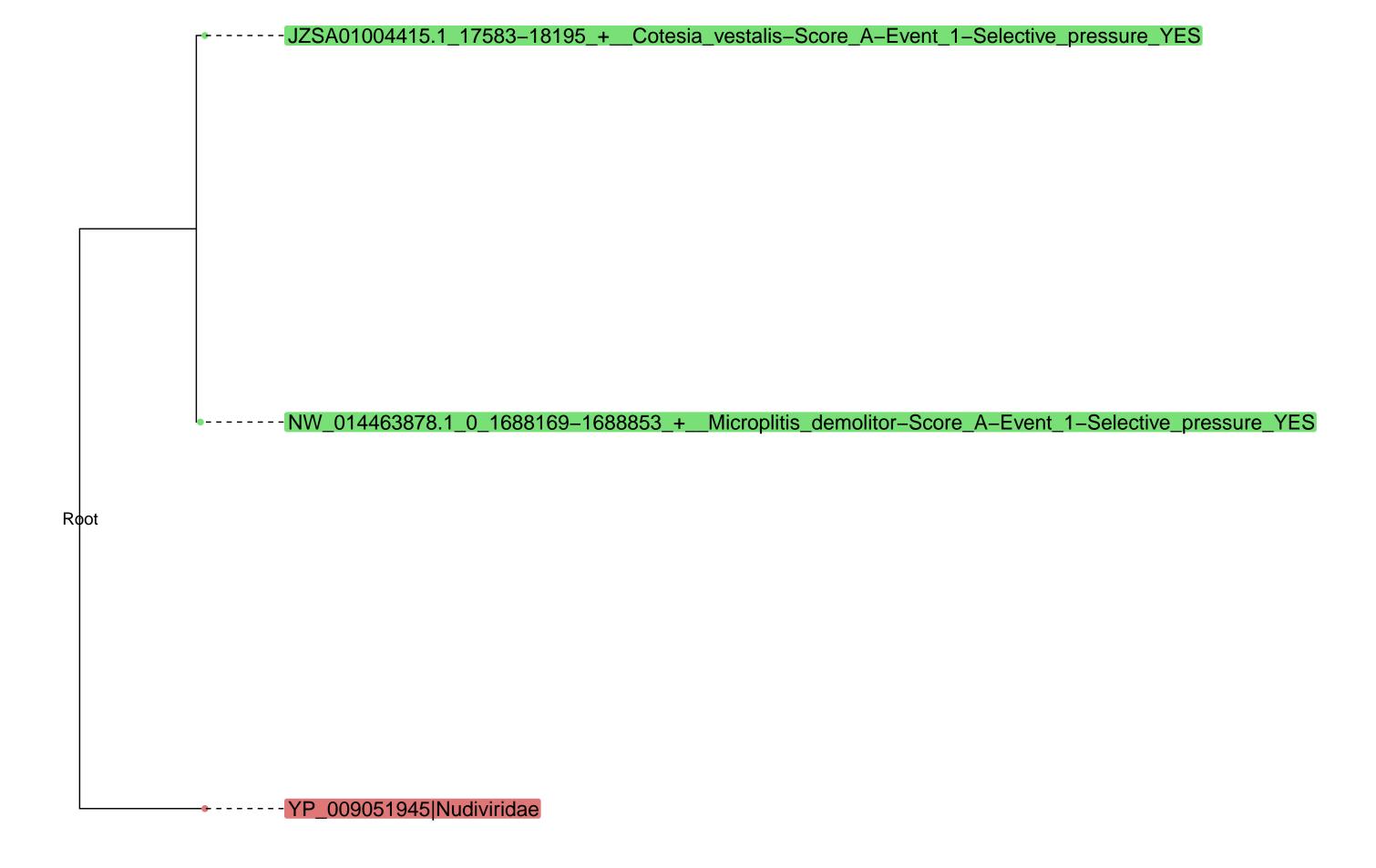


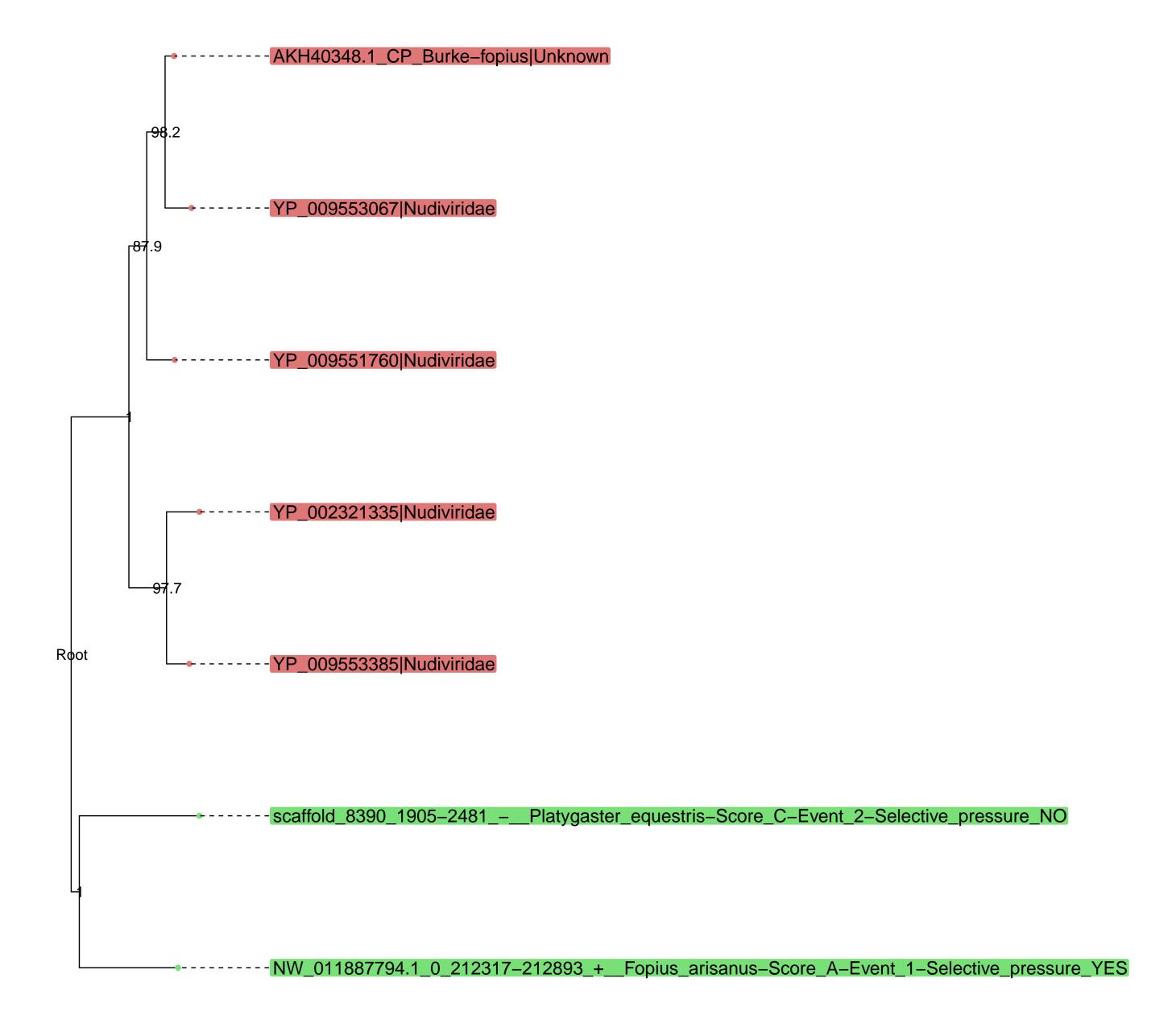


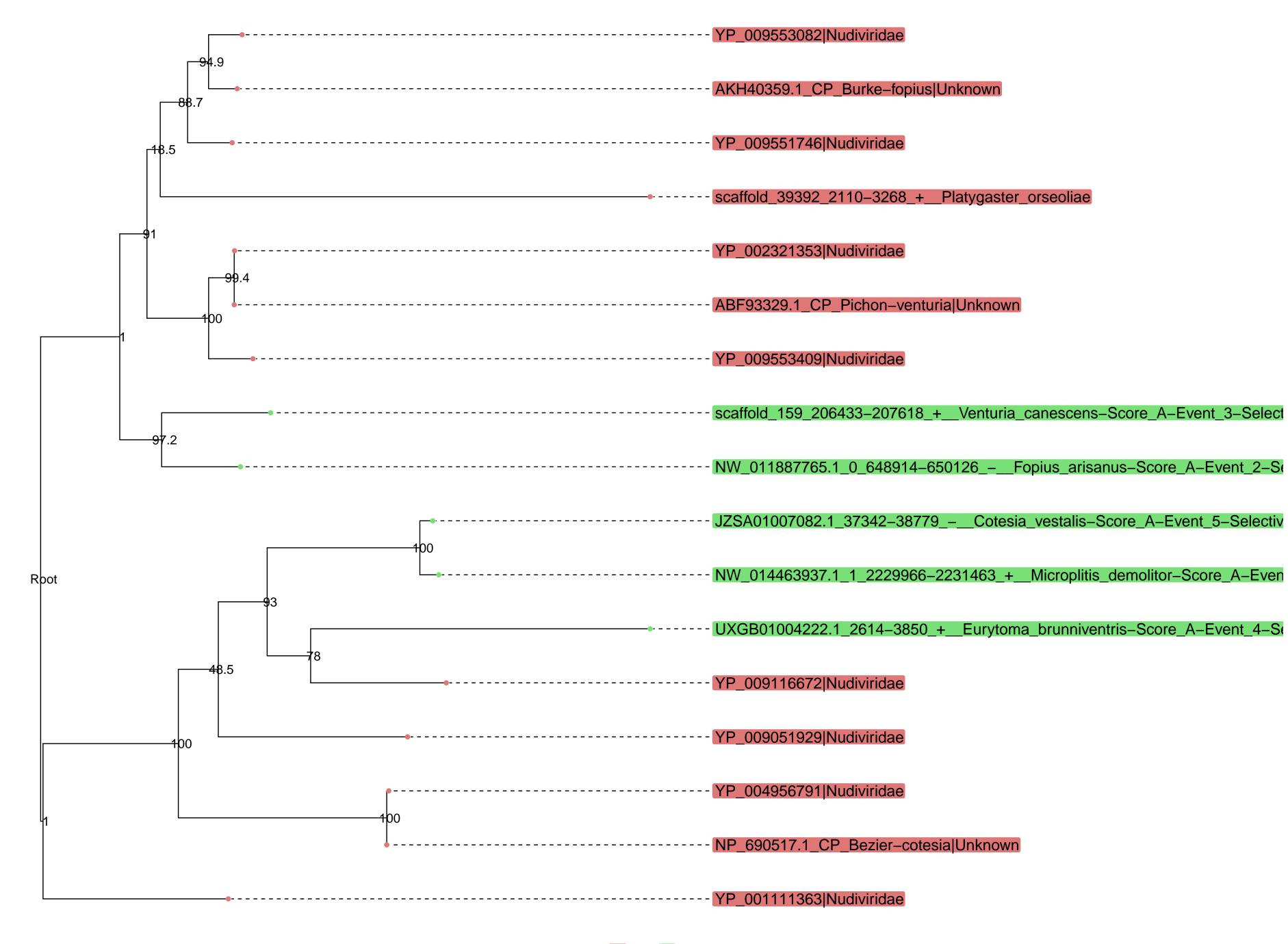


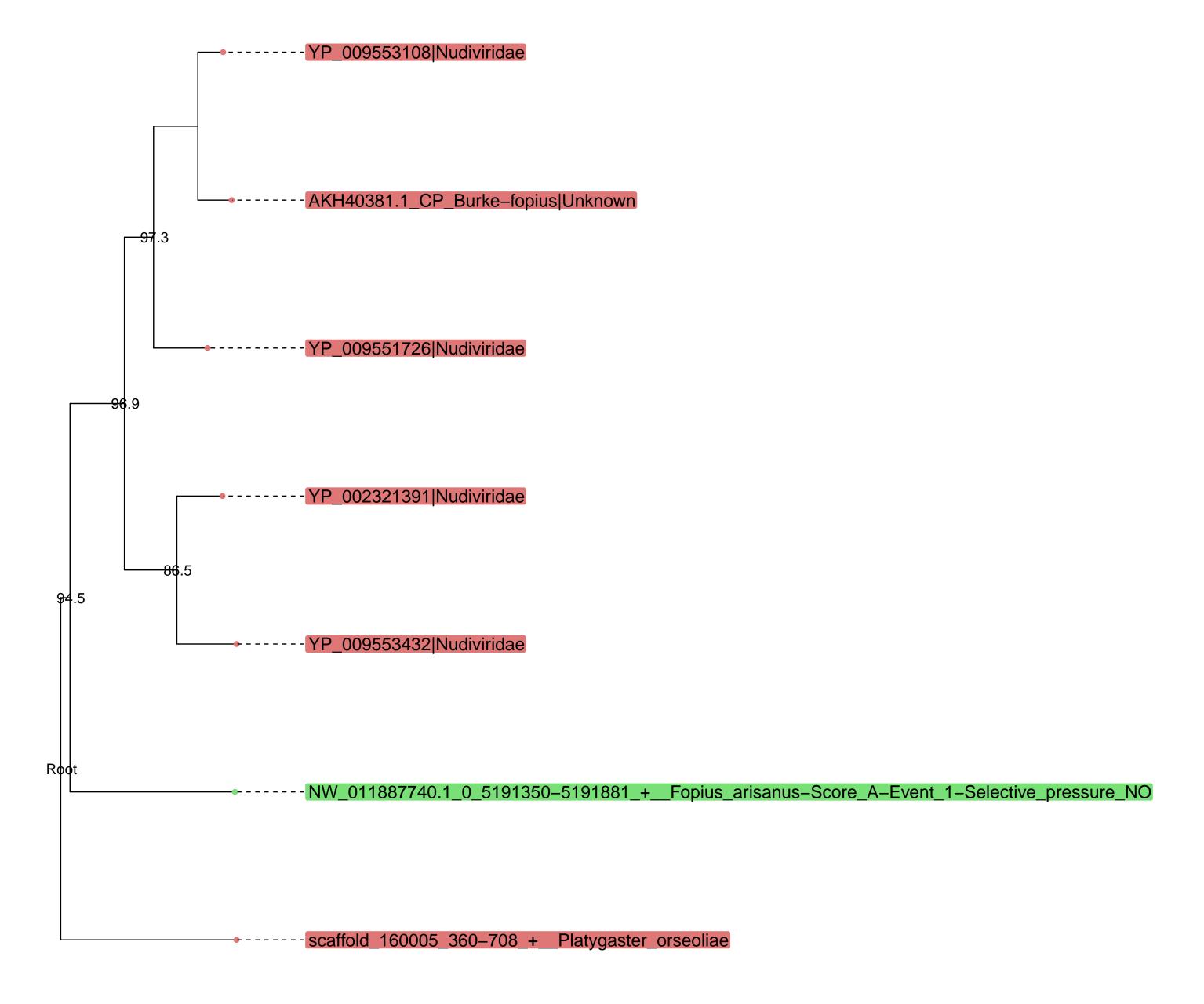
phylum a Virus a EVE

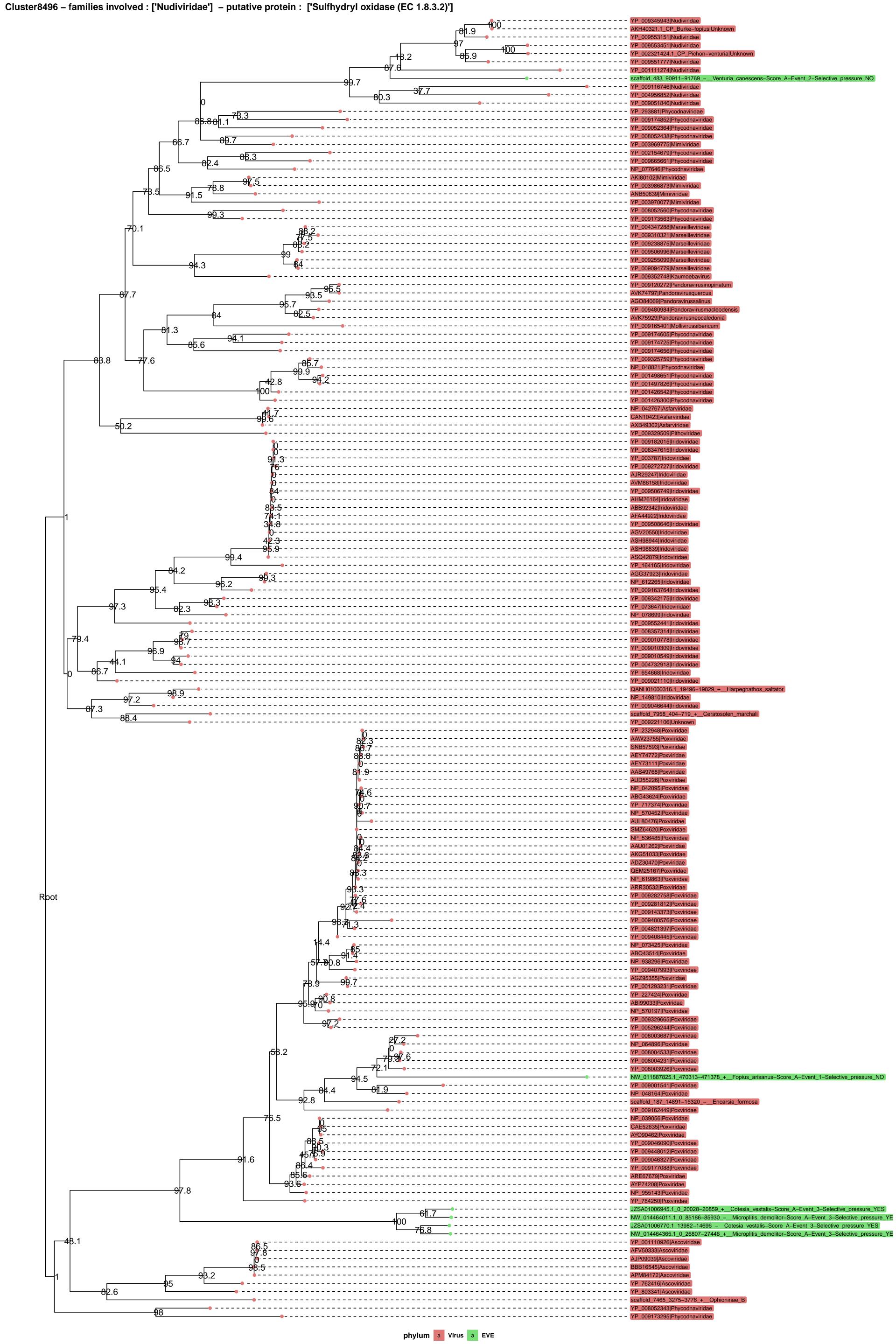


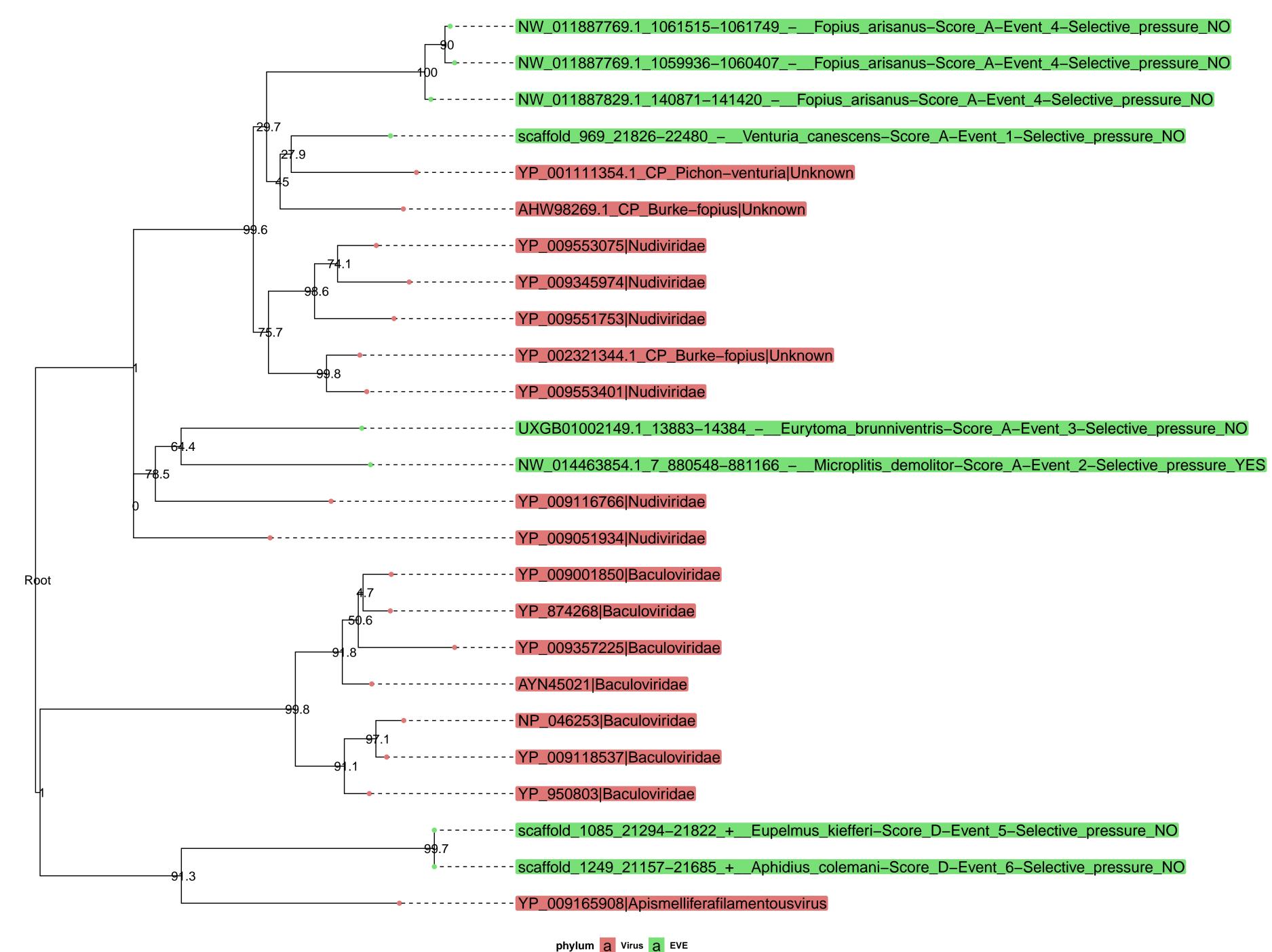


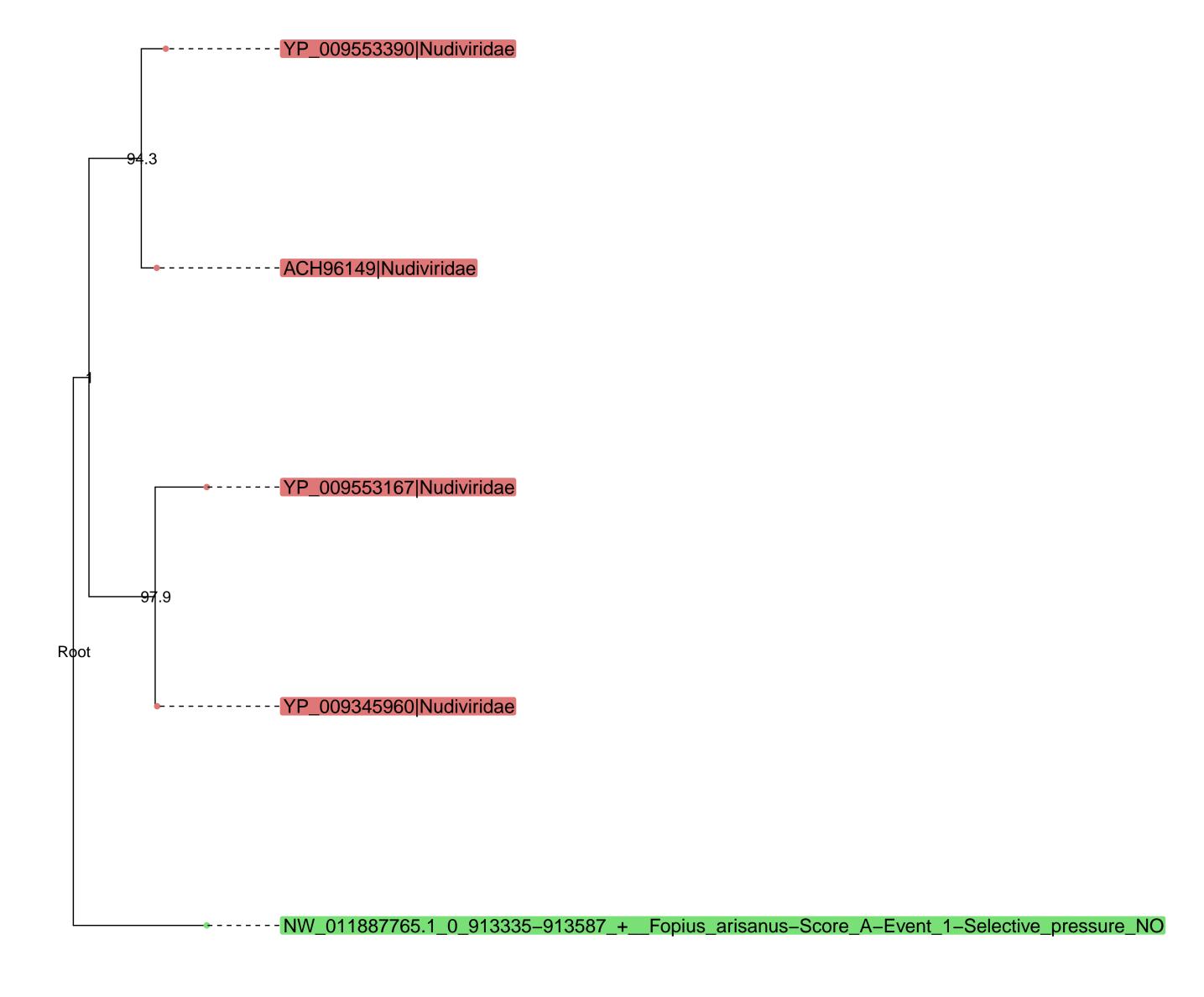


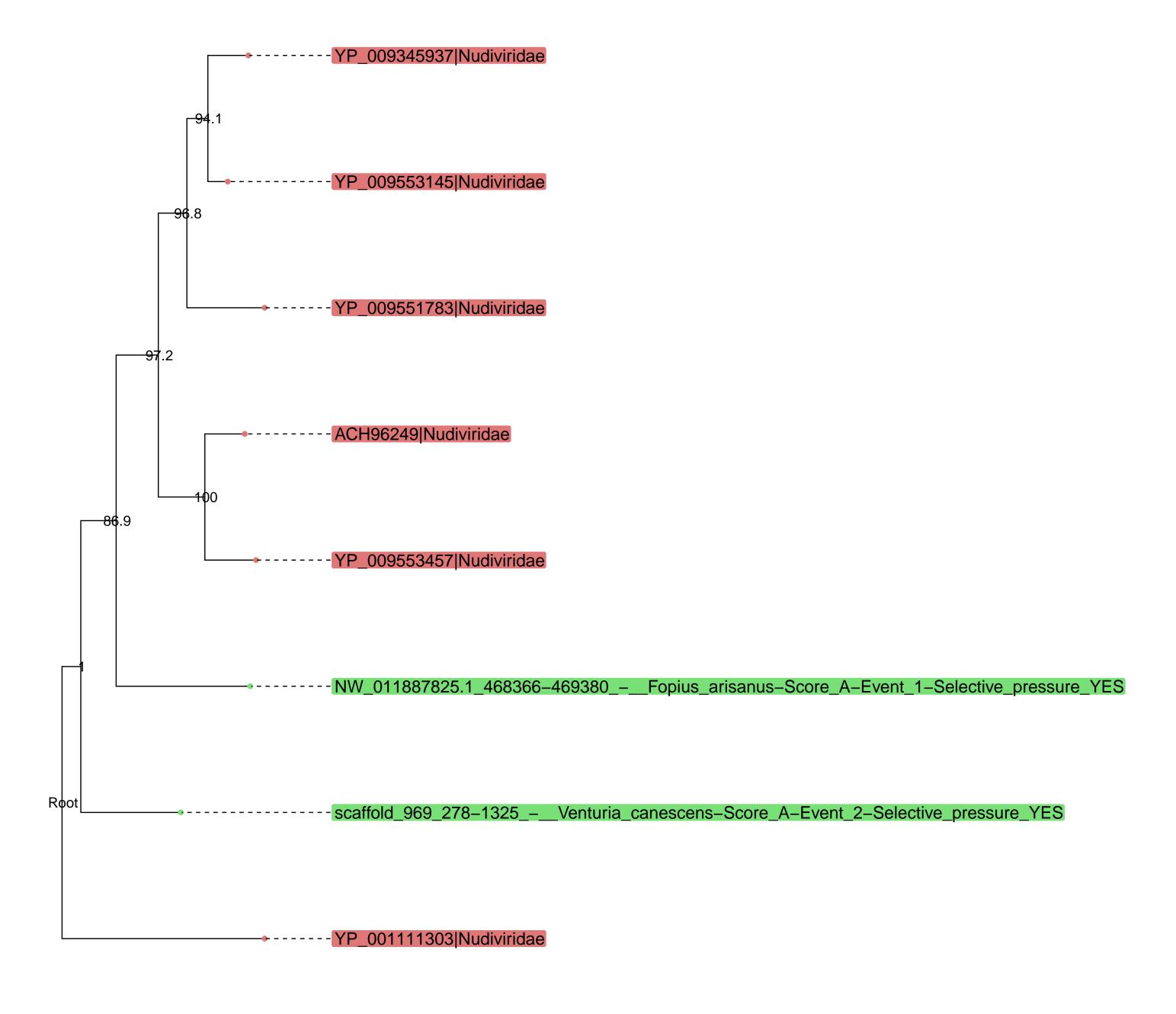


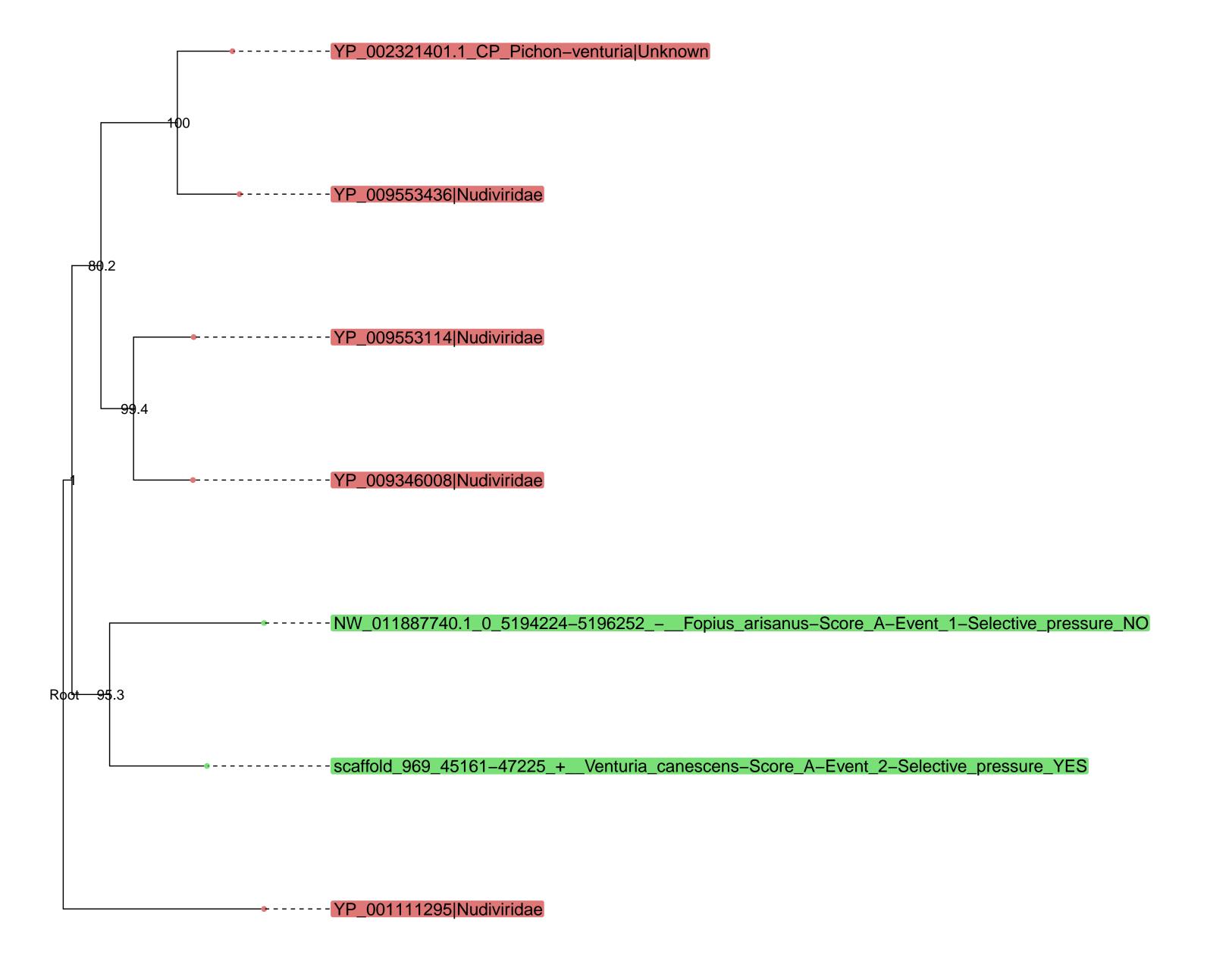


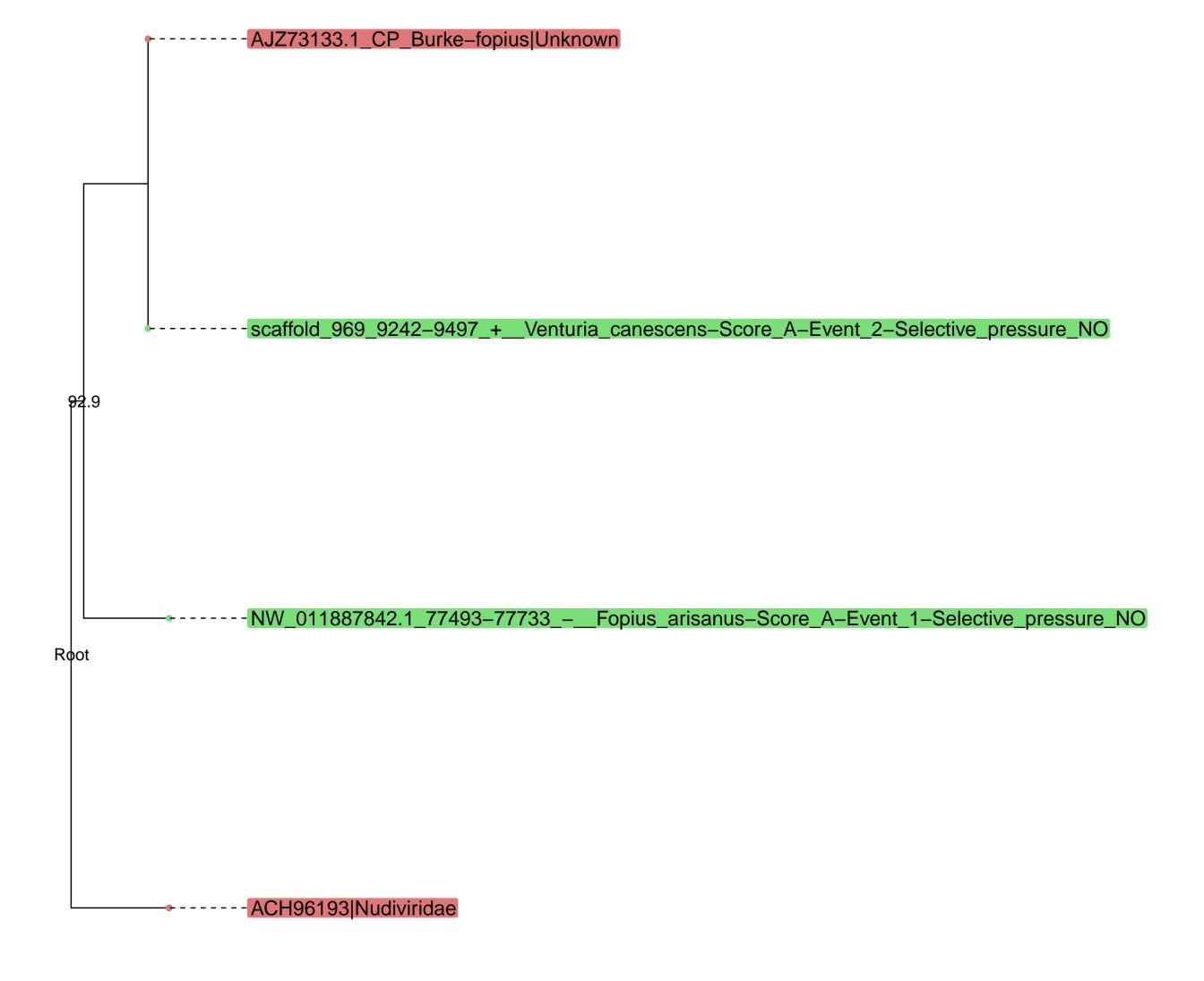


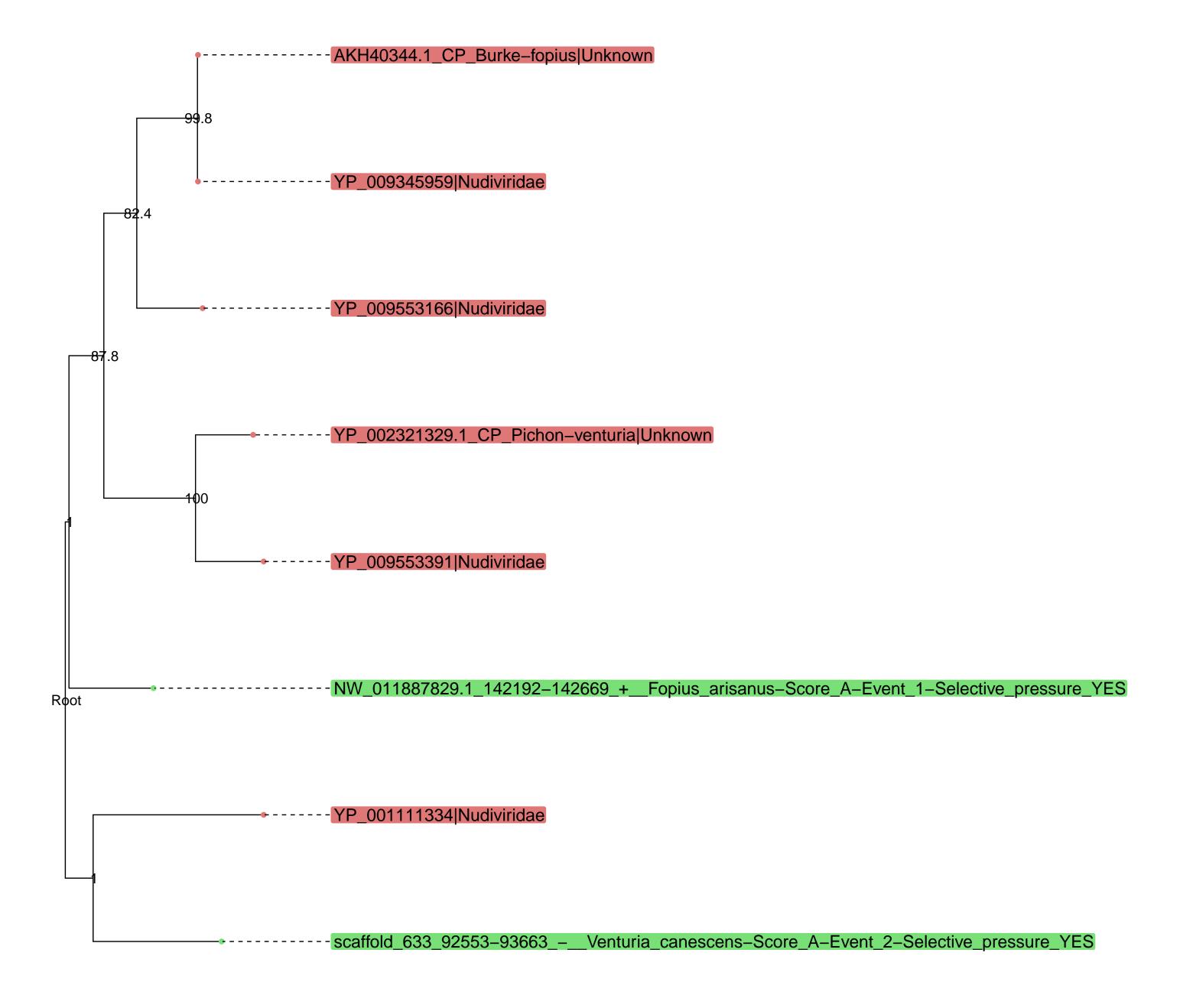


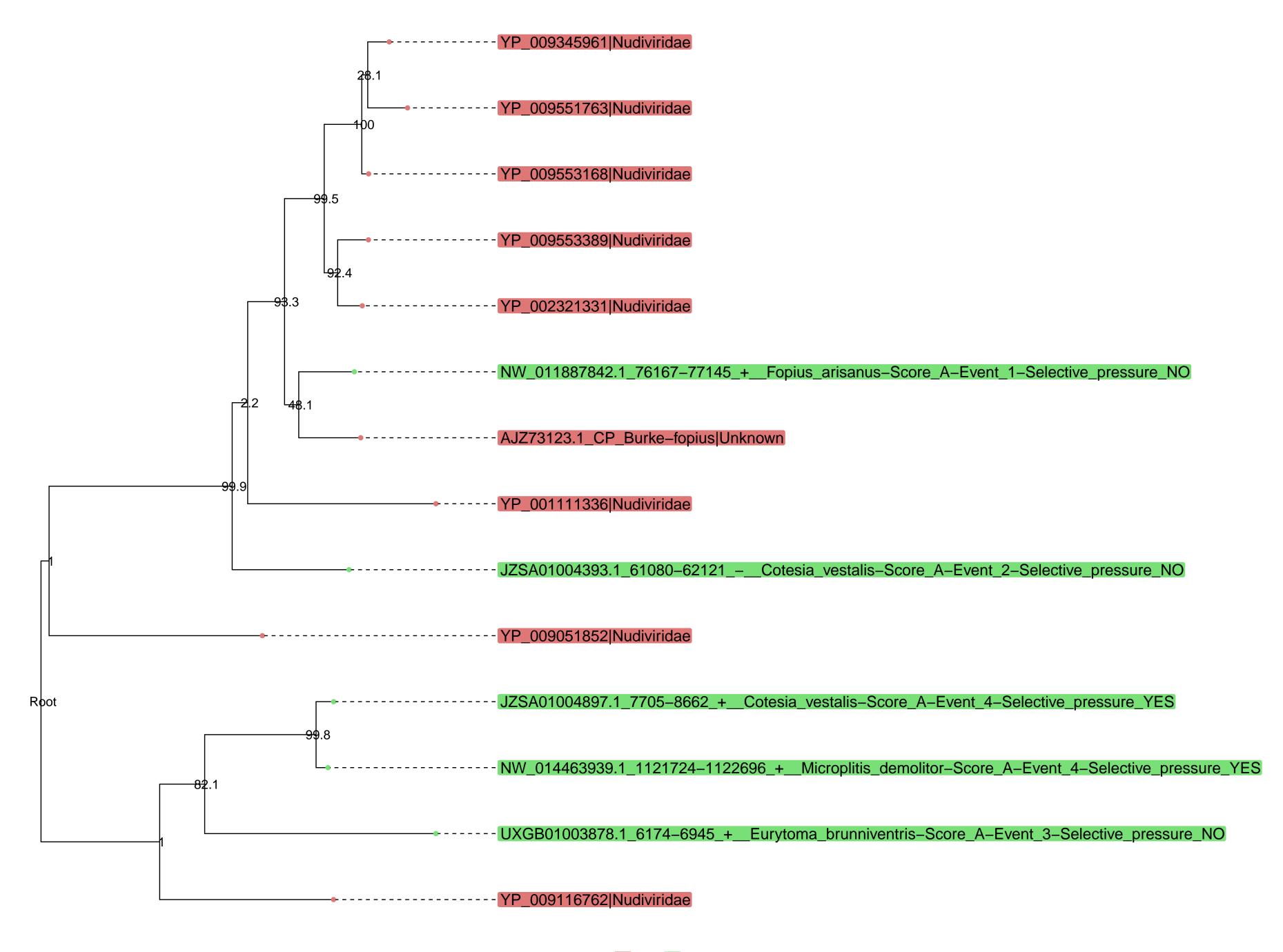


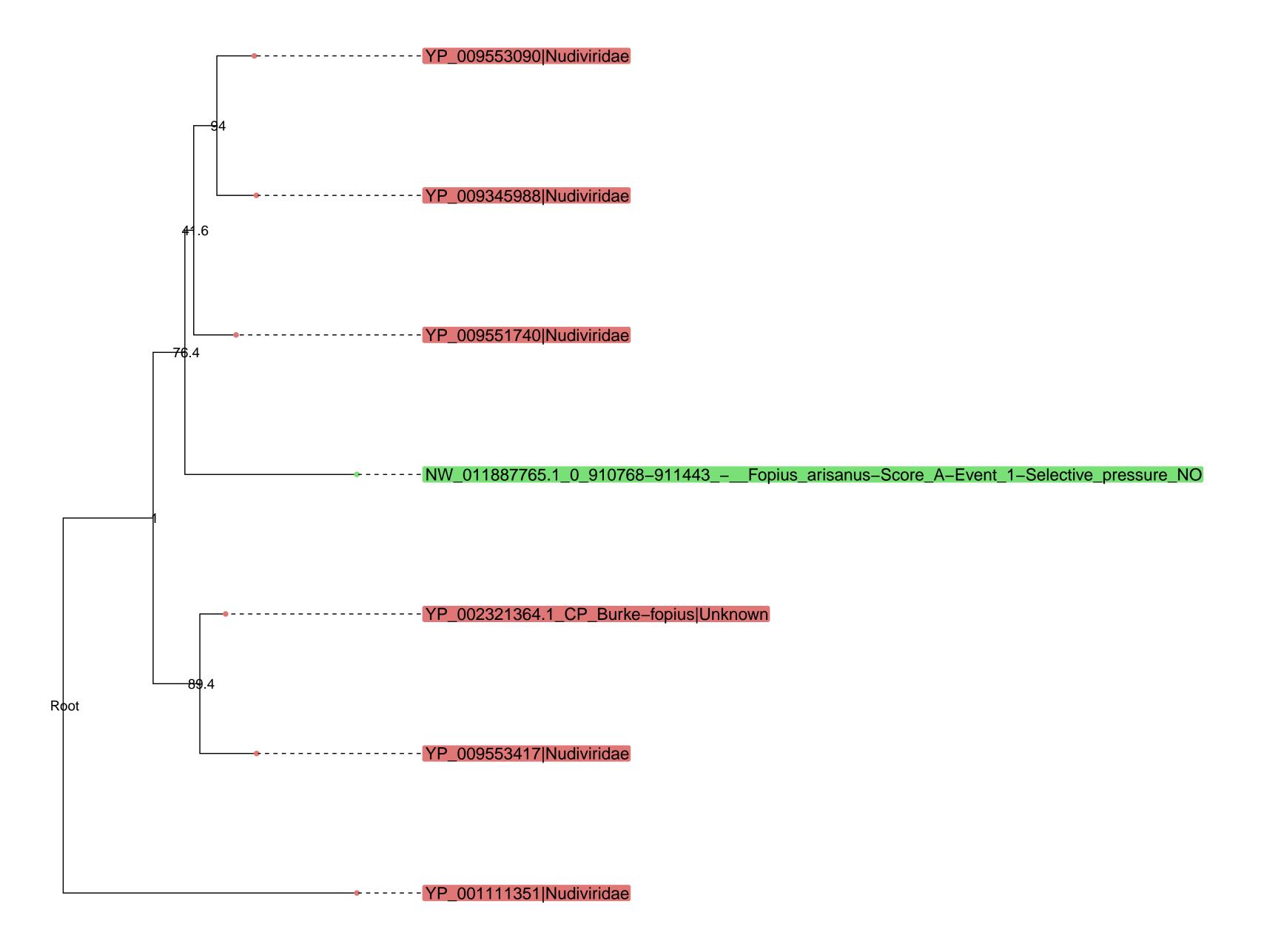




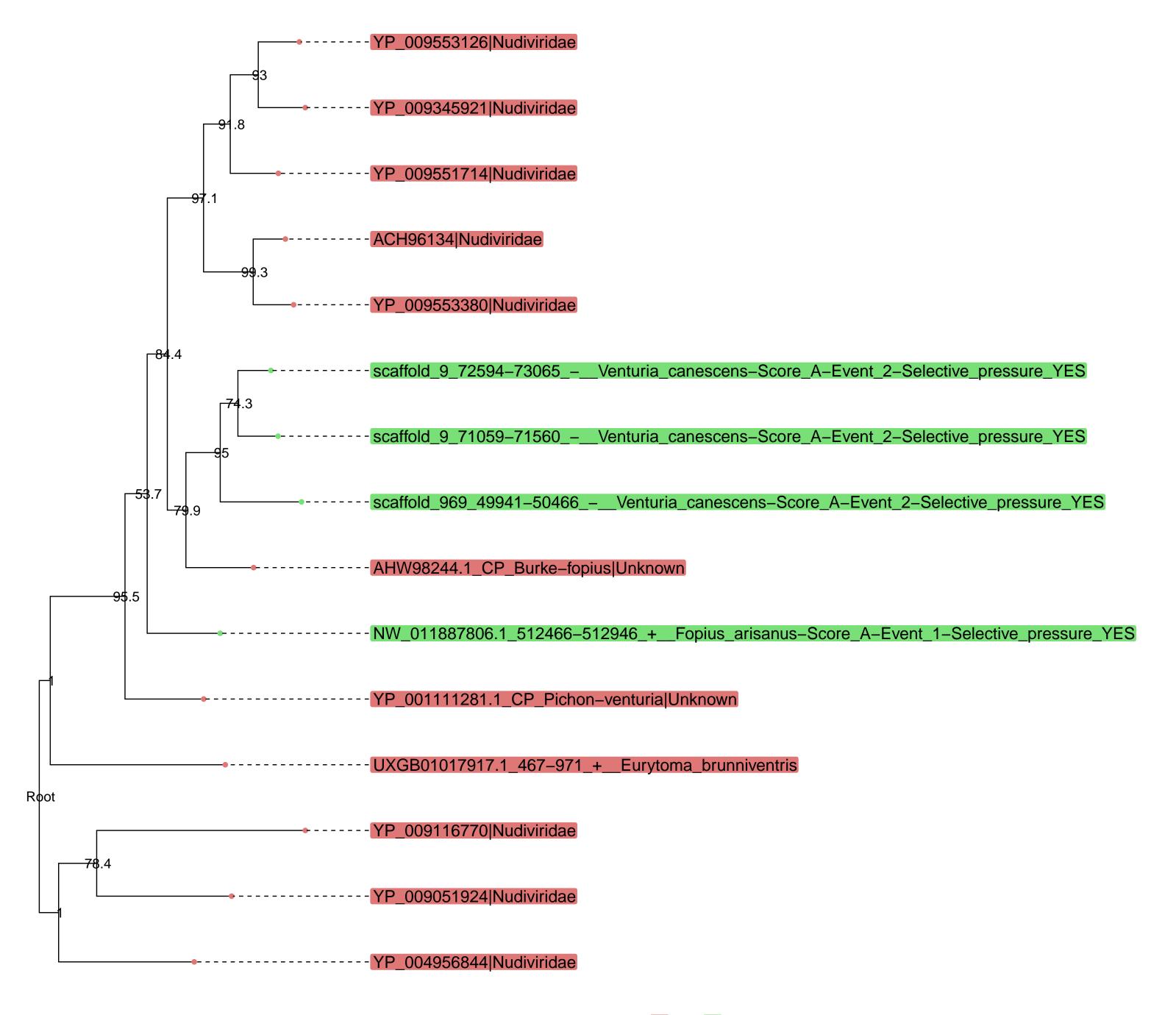


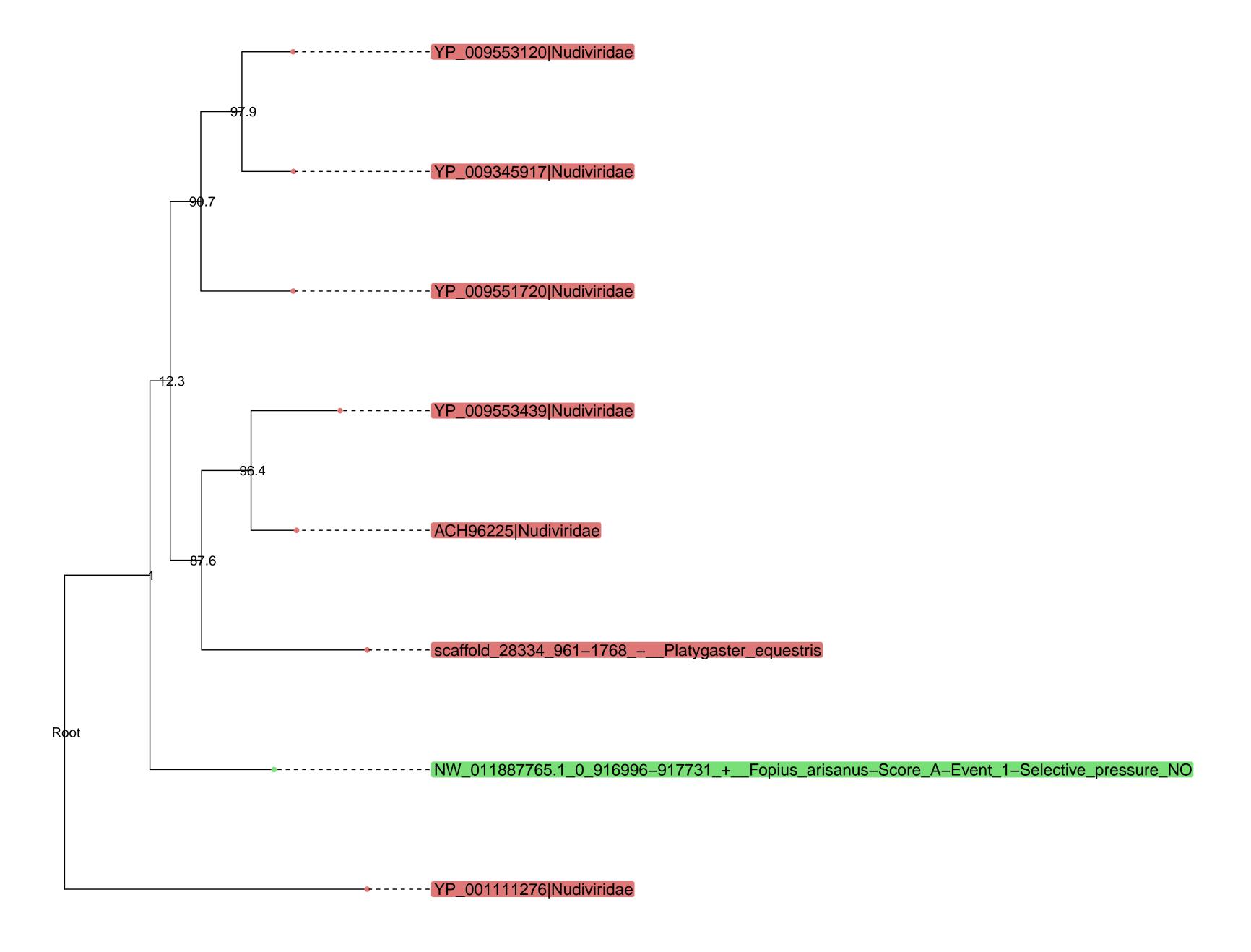


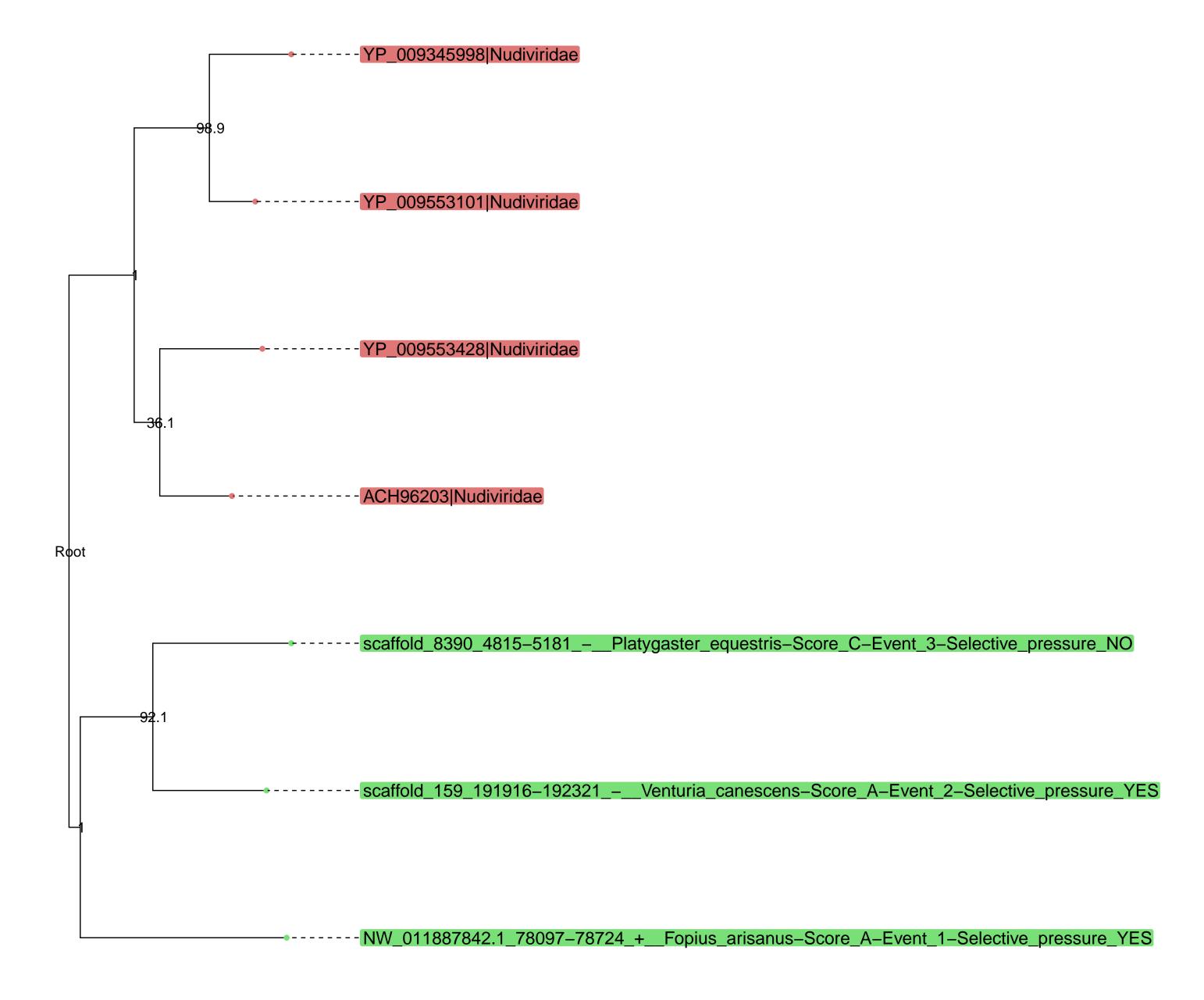


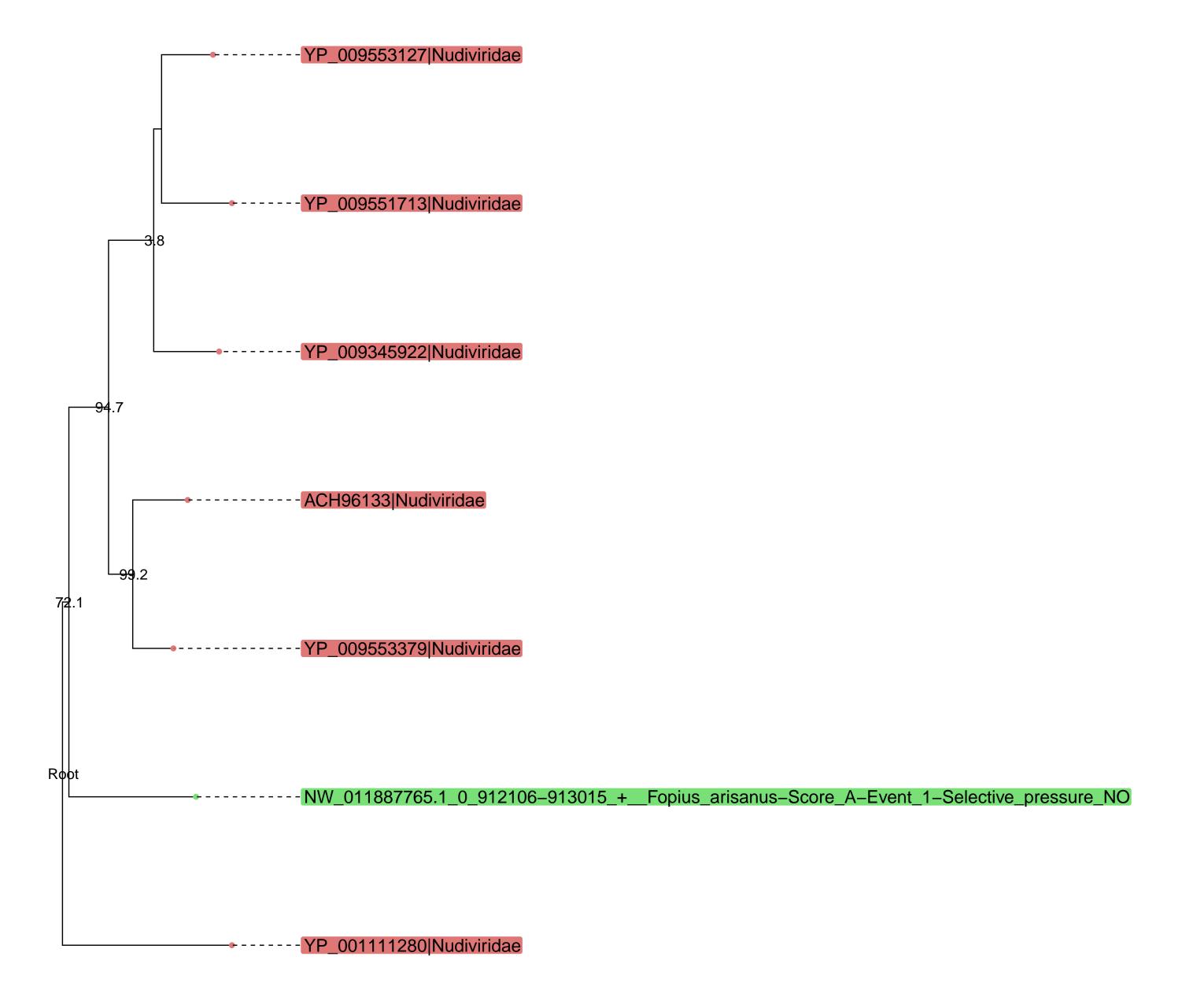


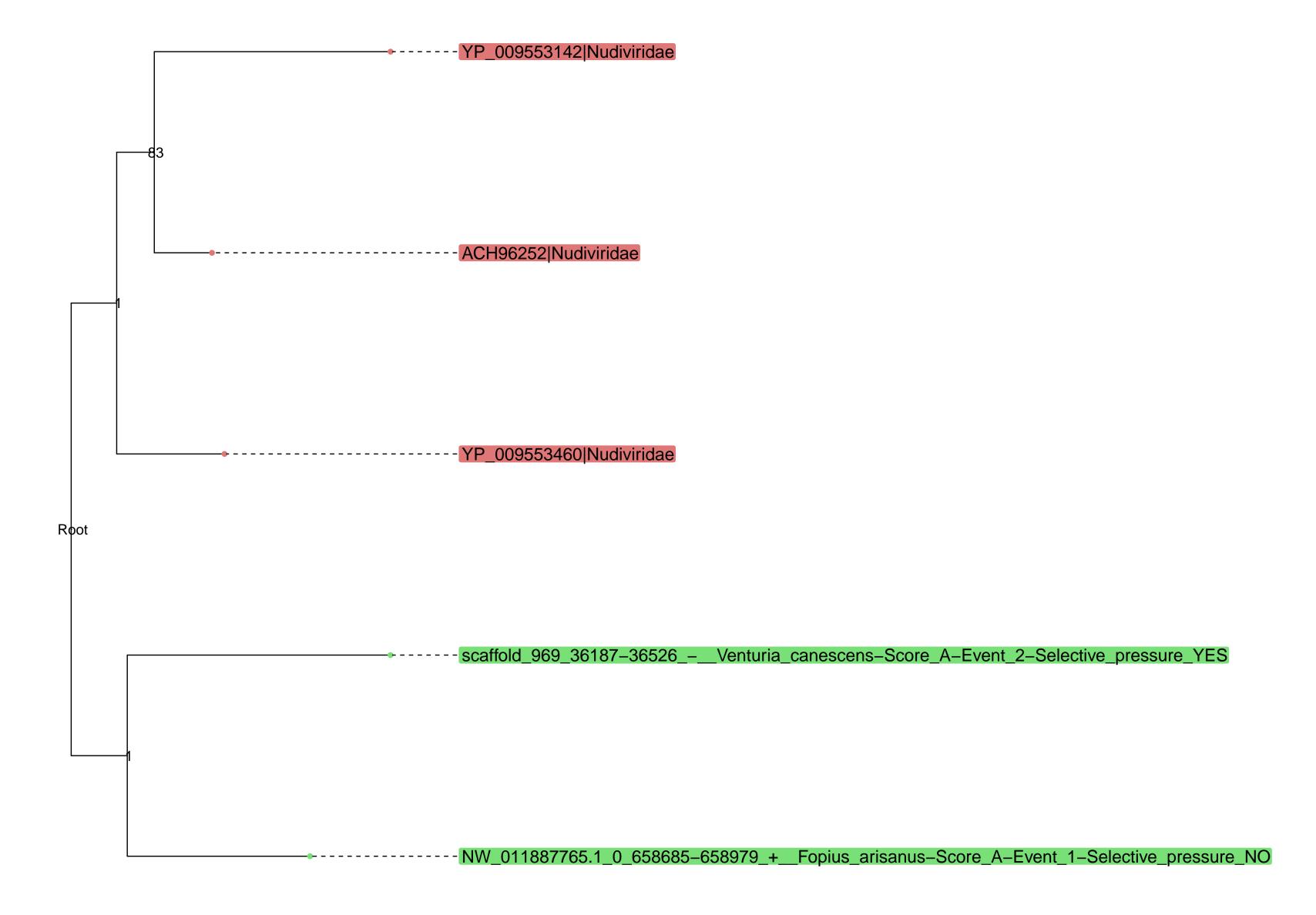


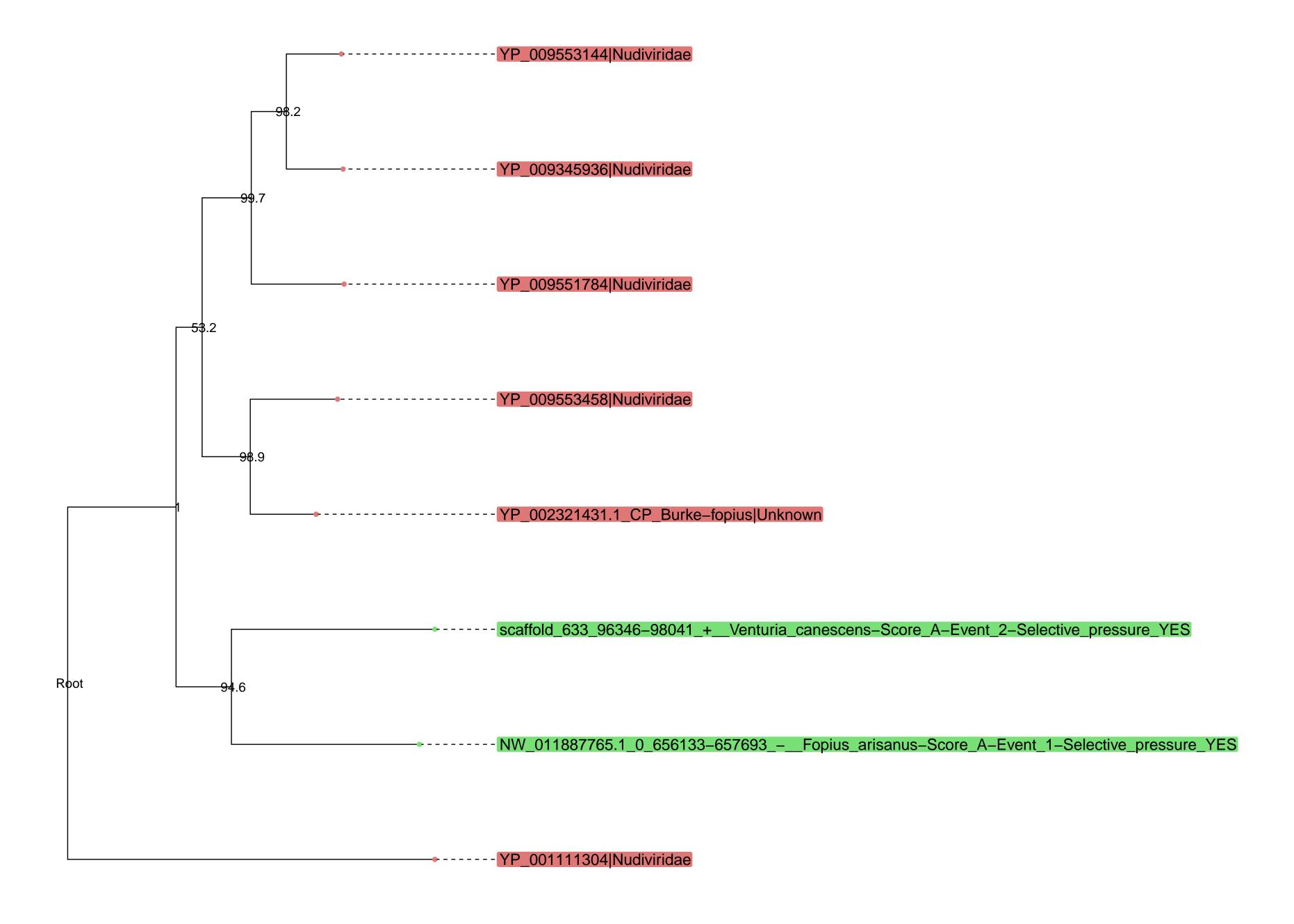


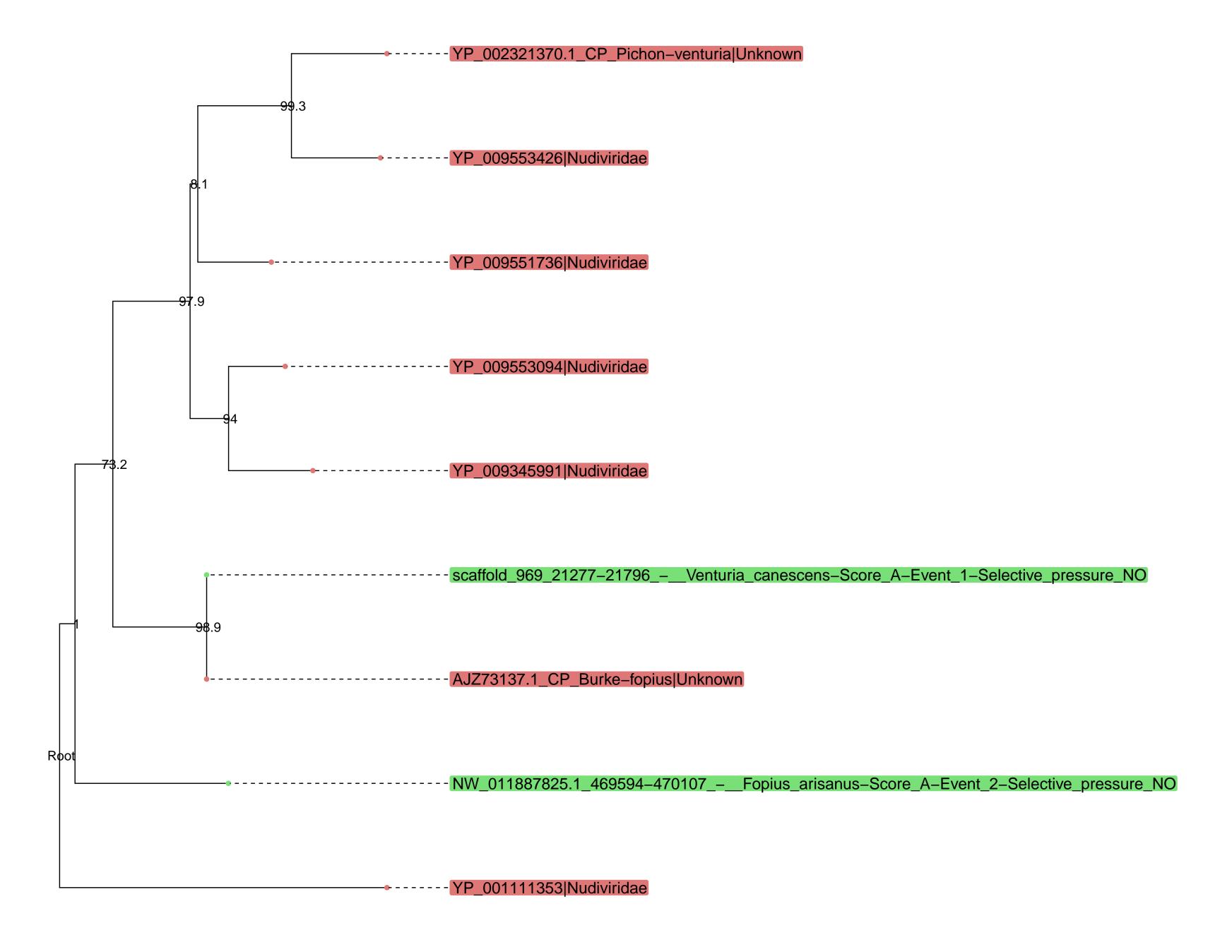


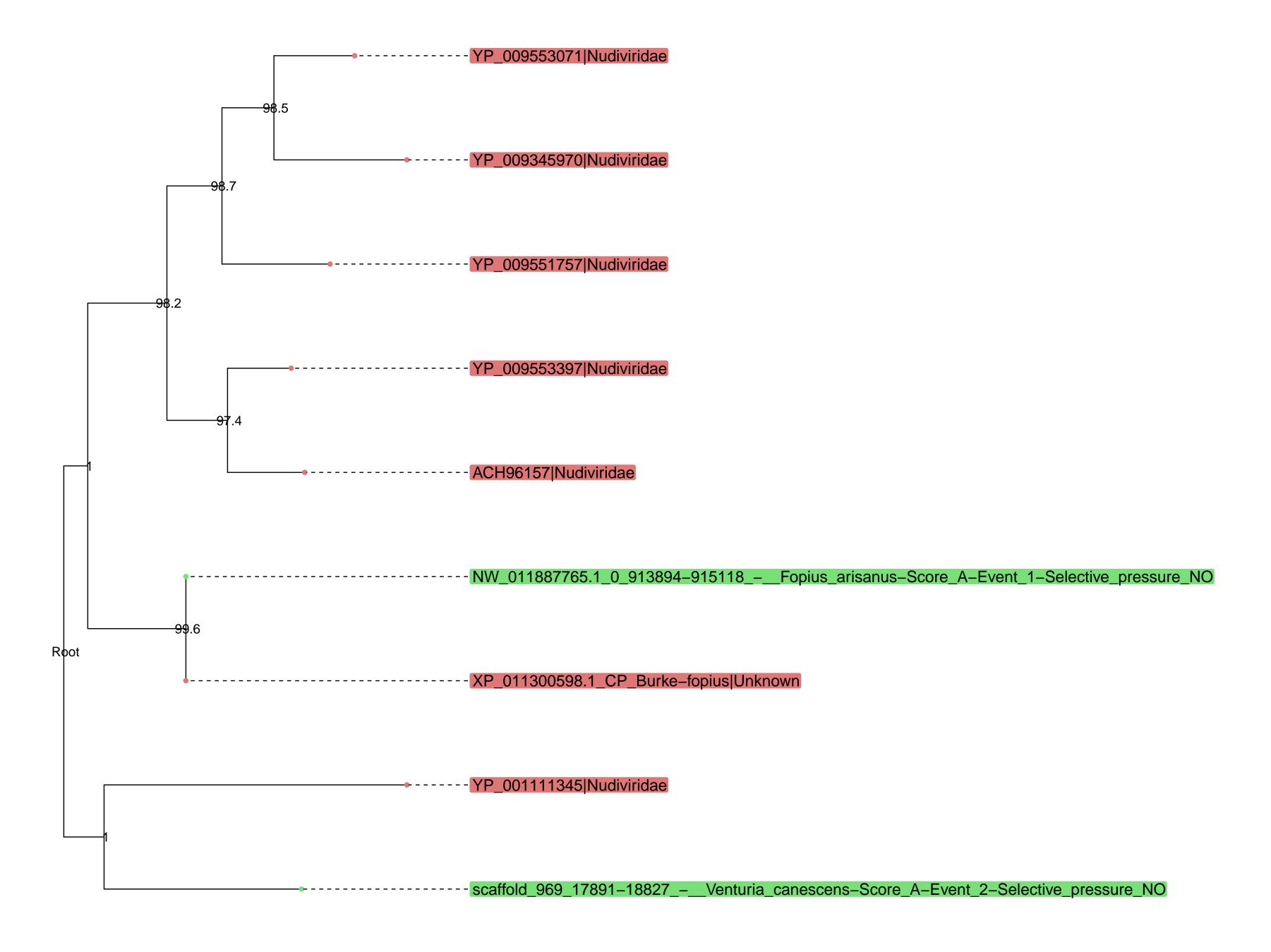


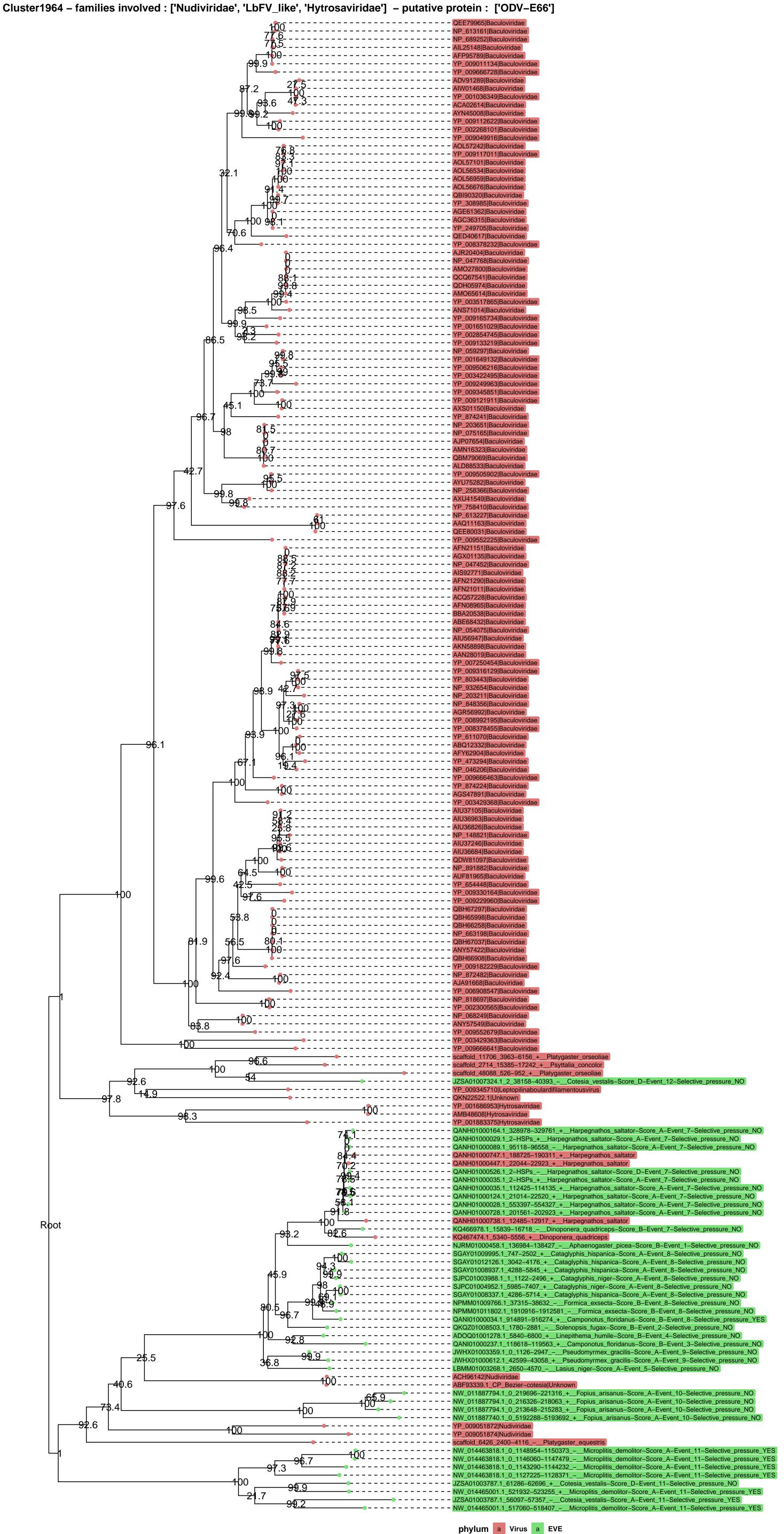


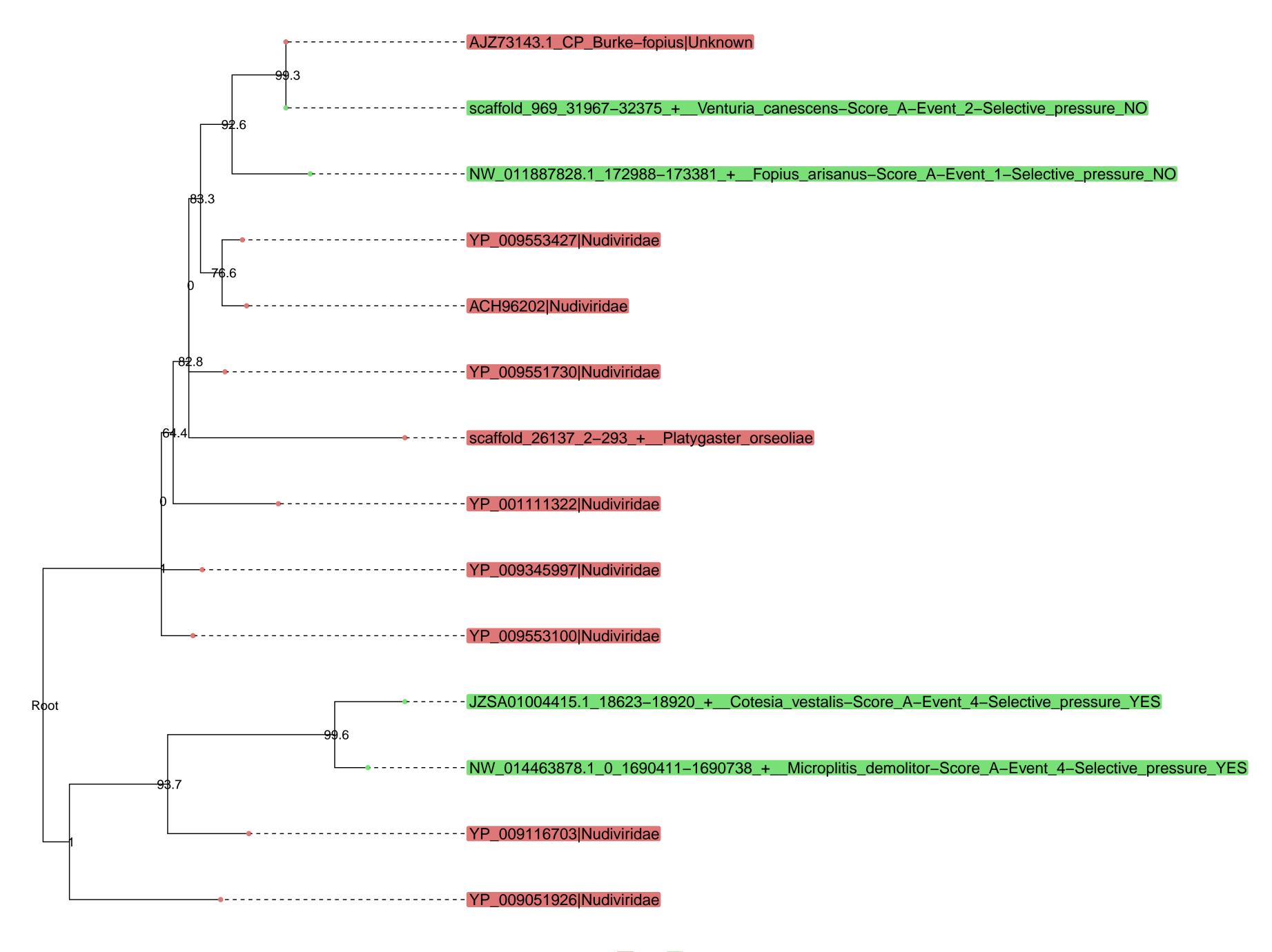


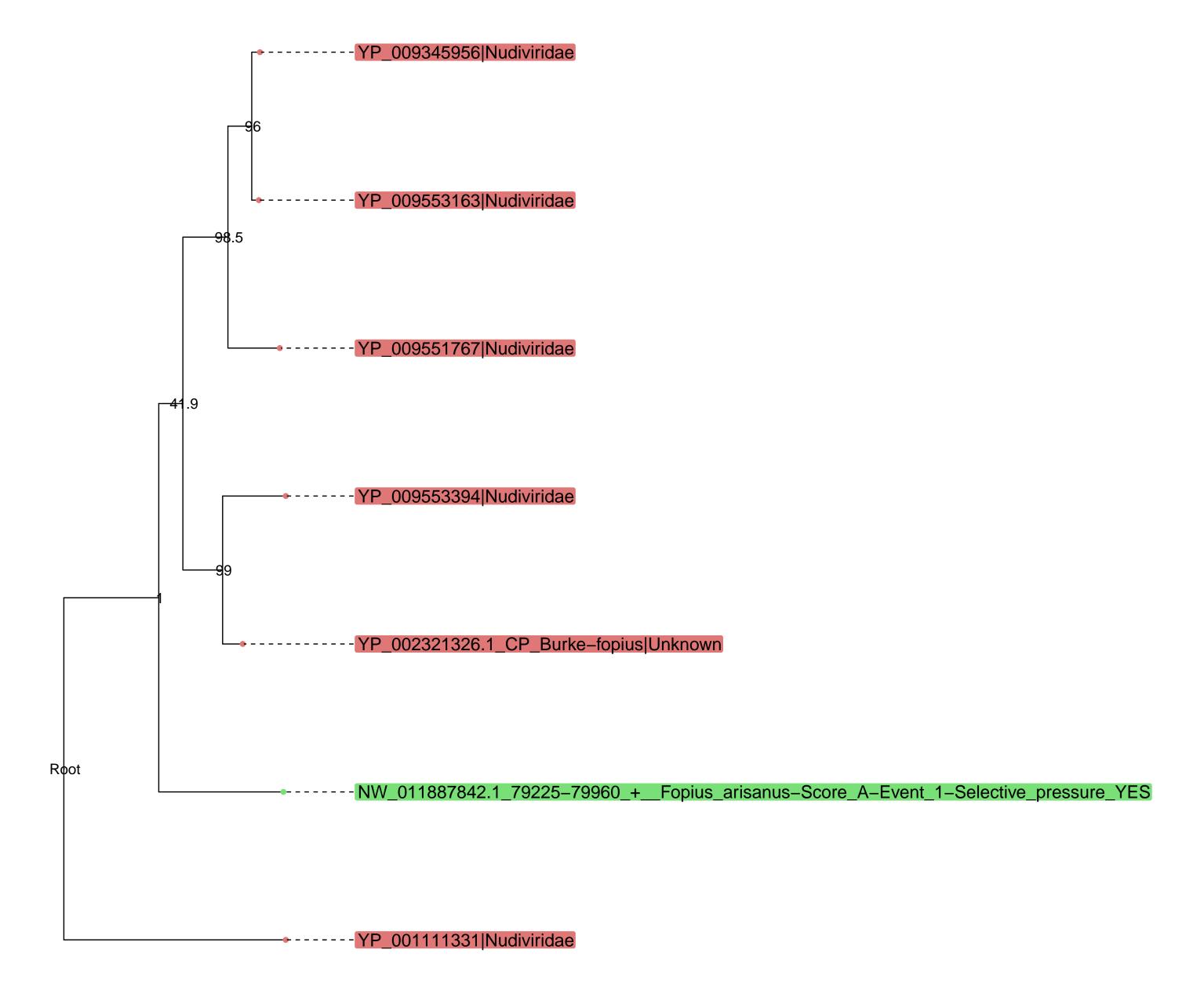


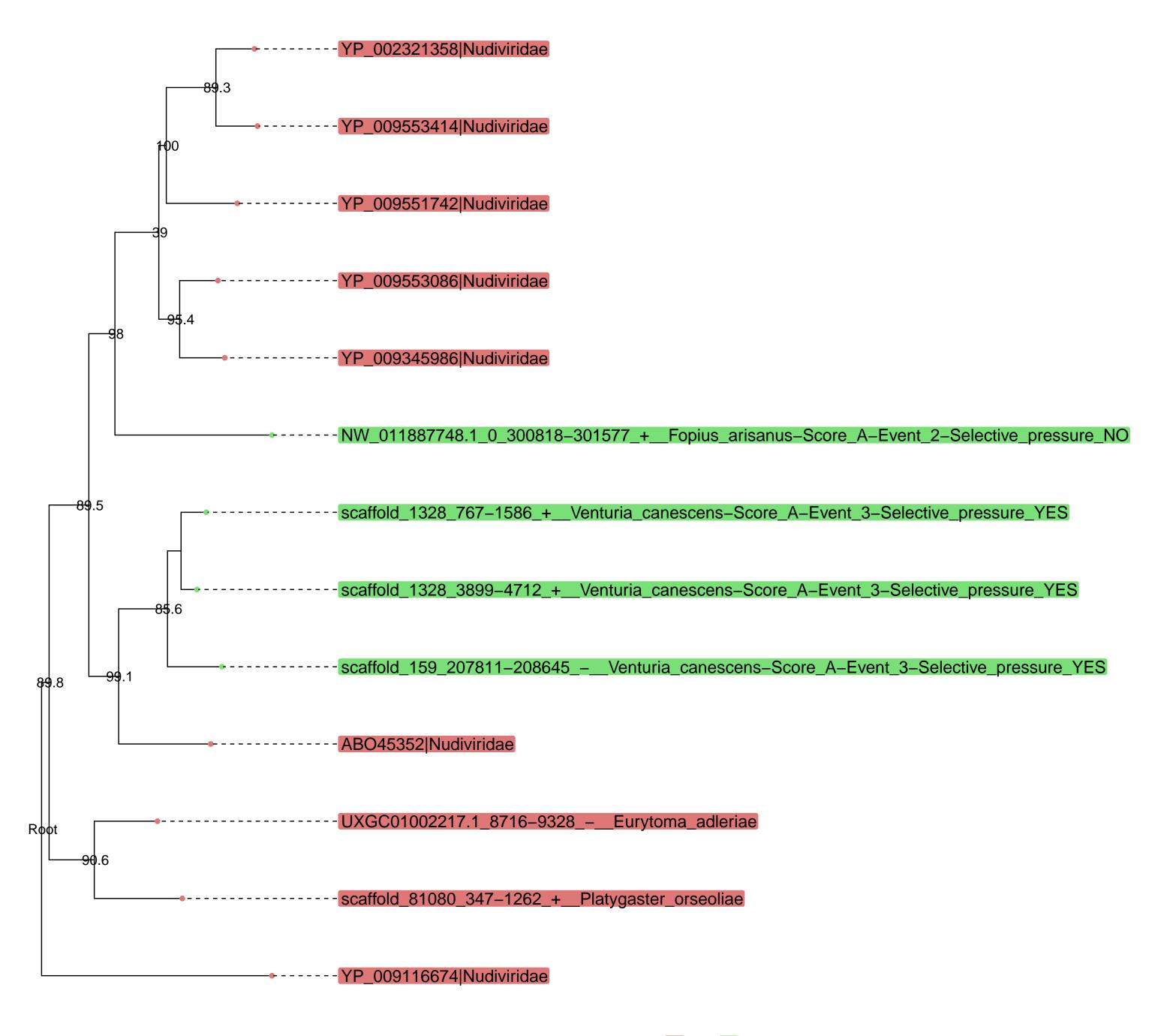


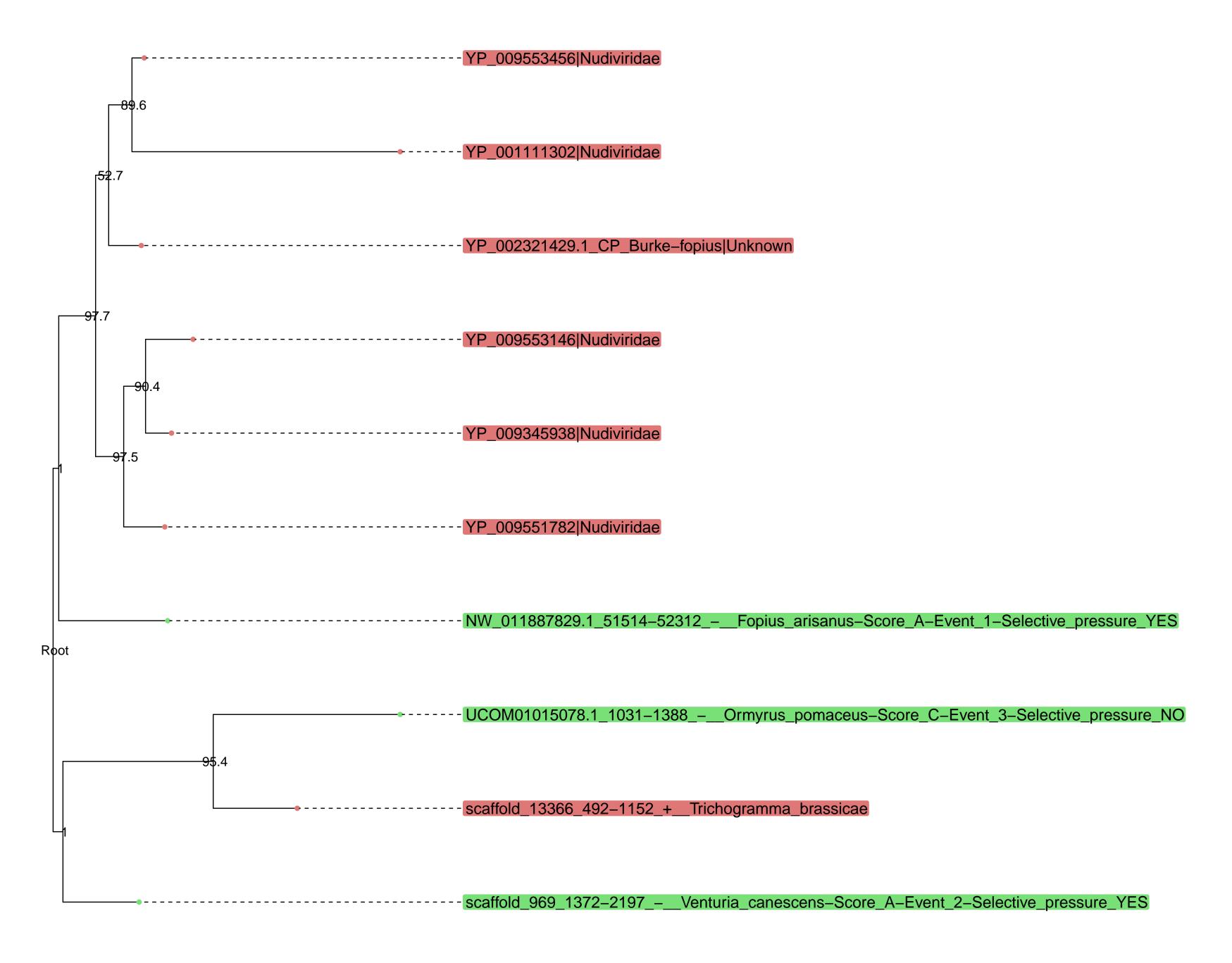


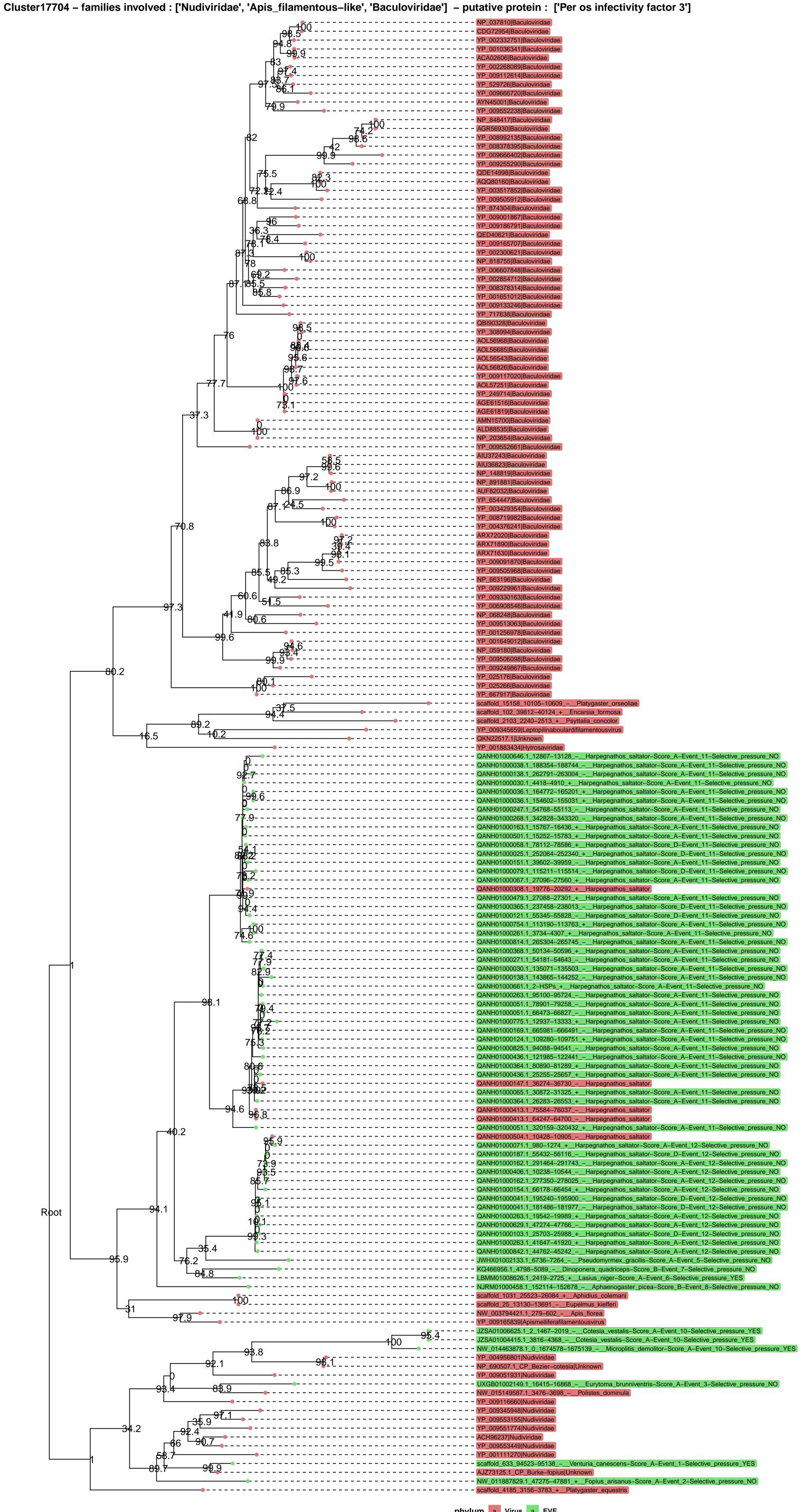


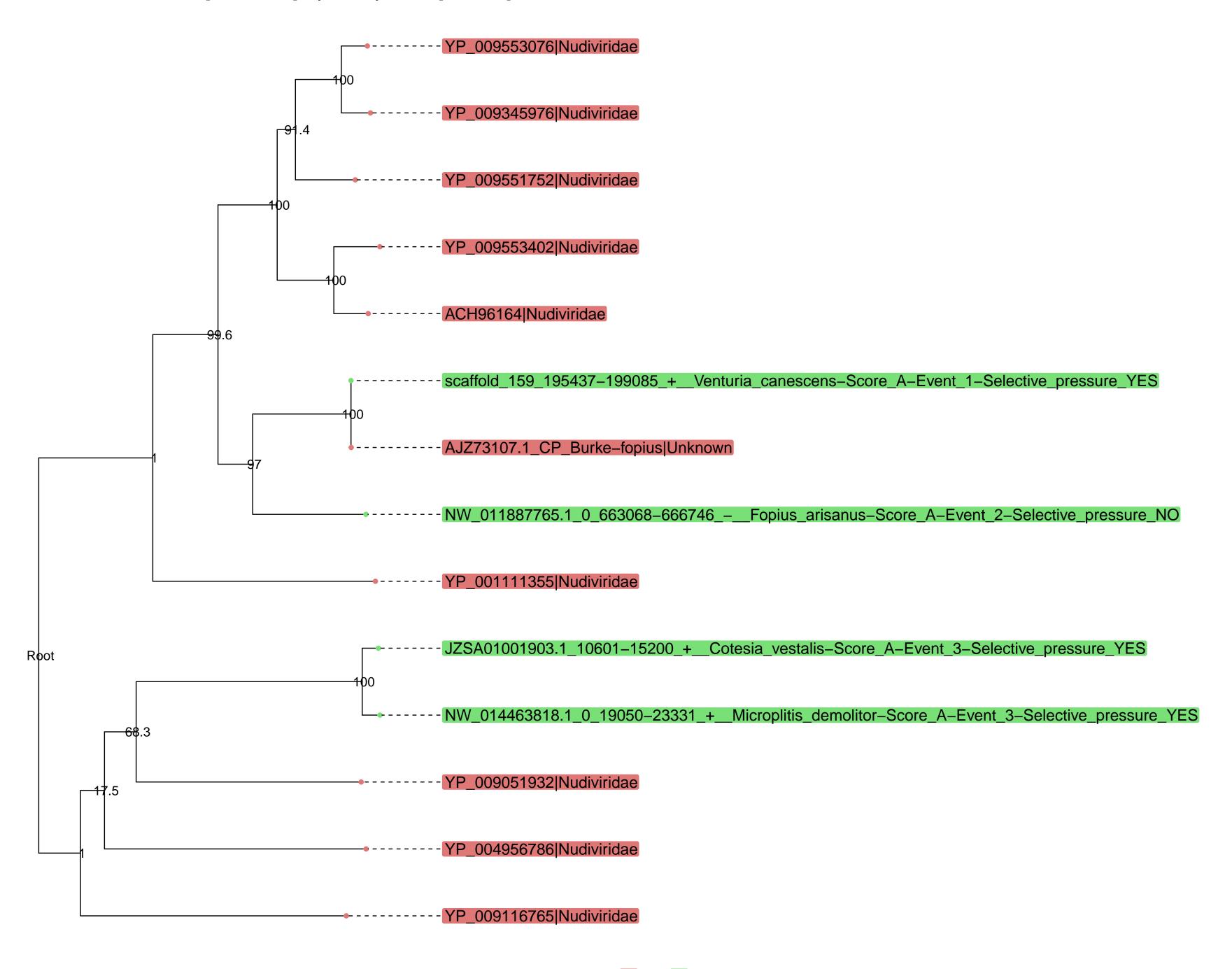


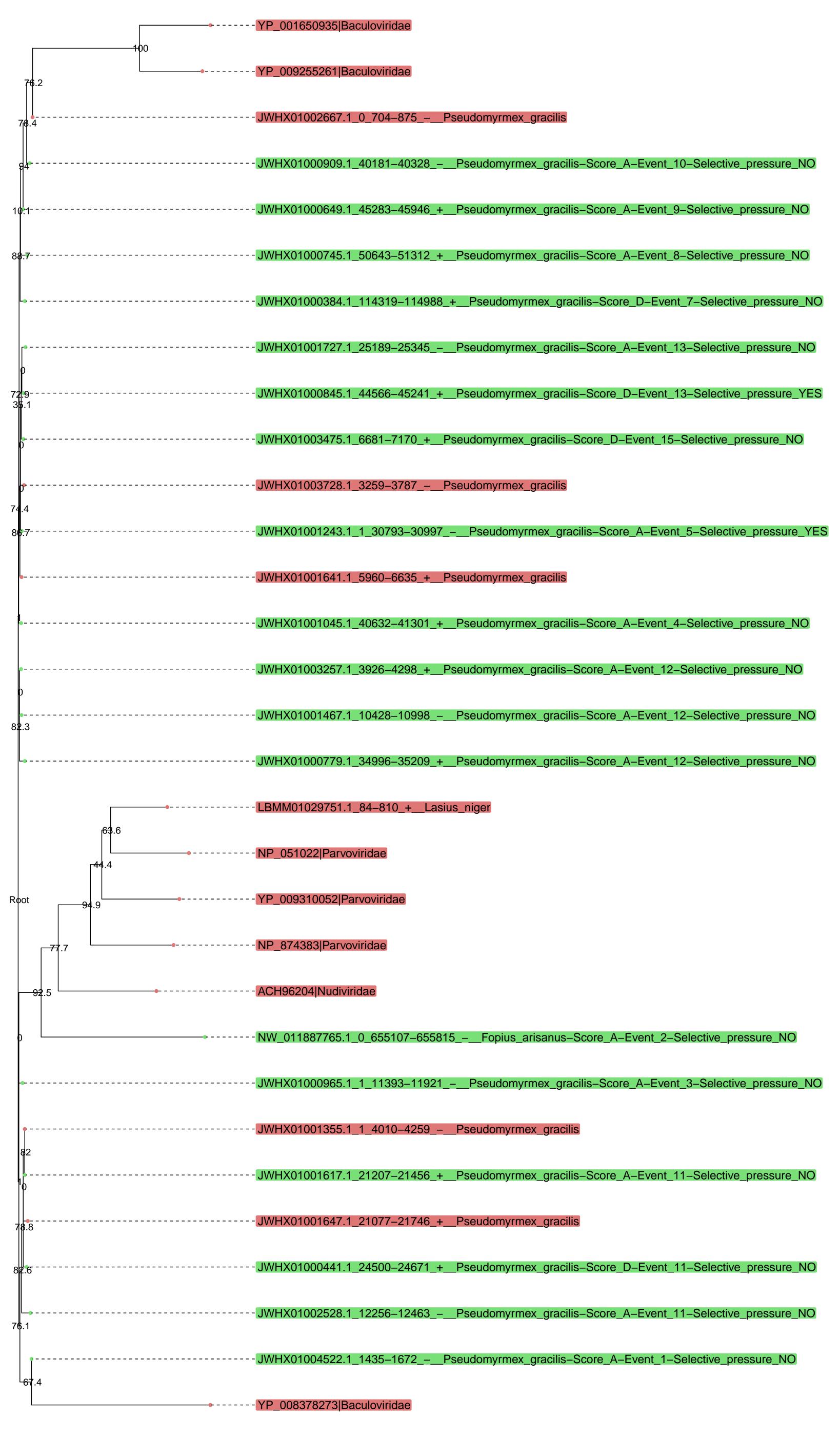


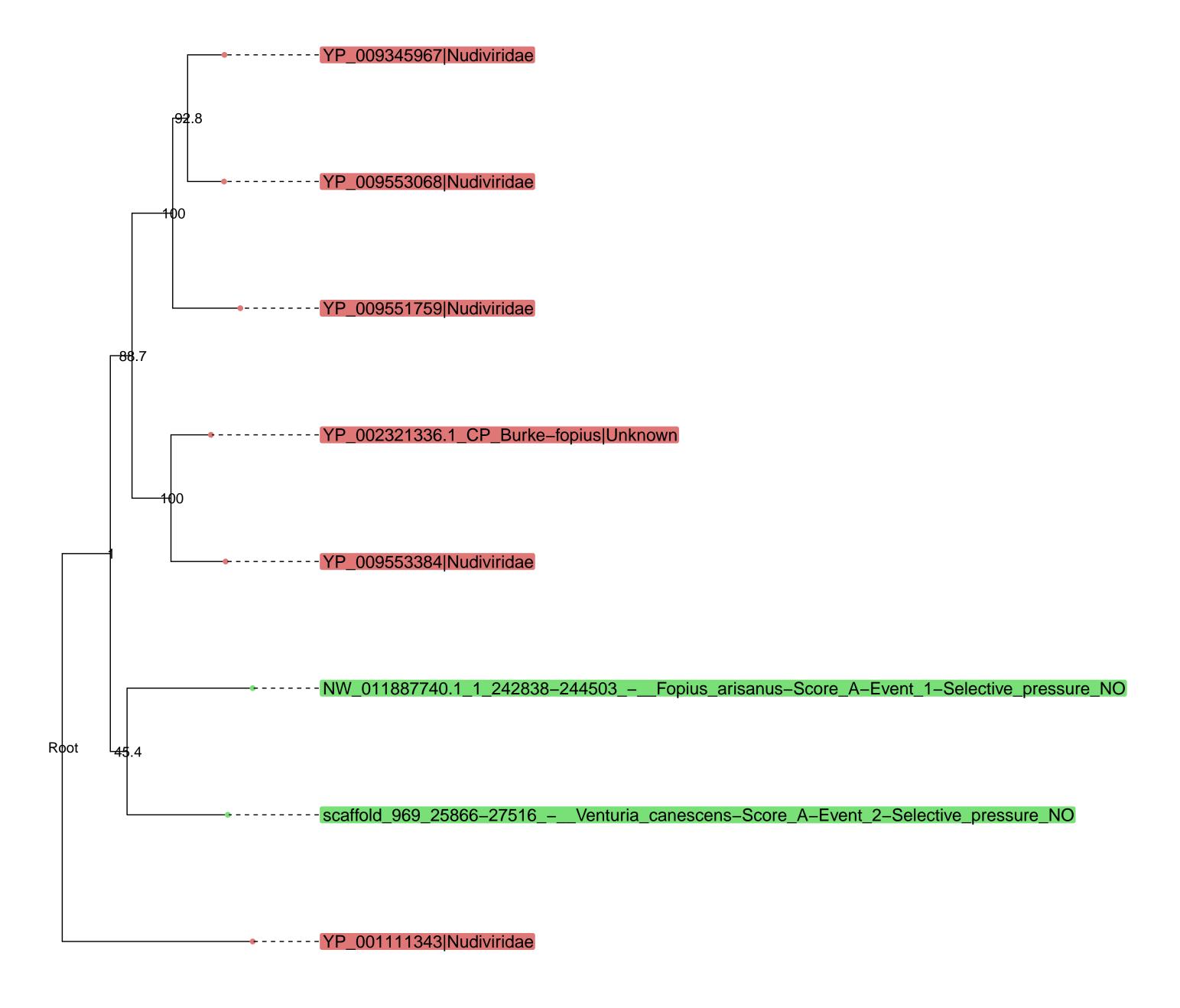


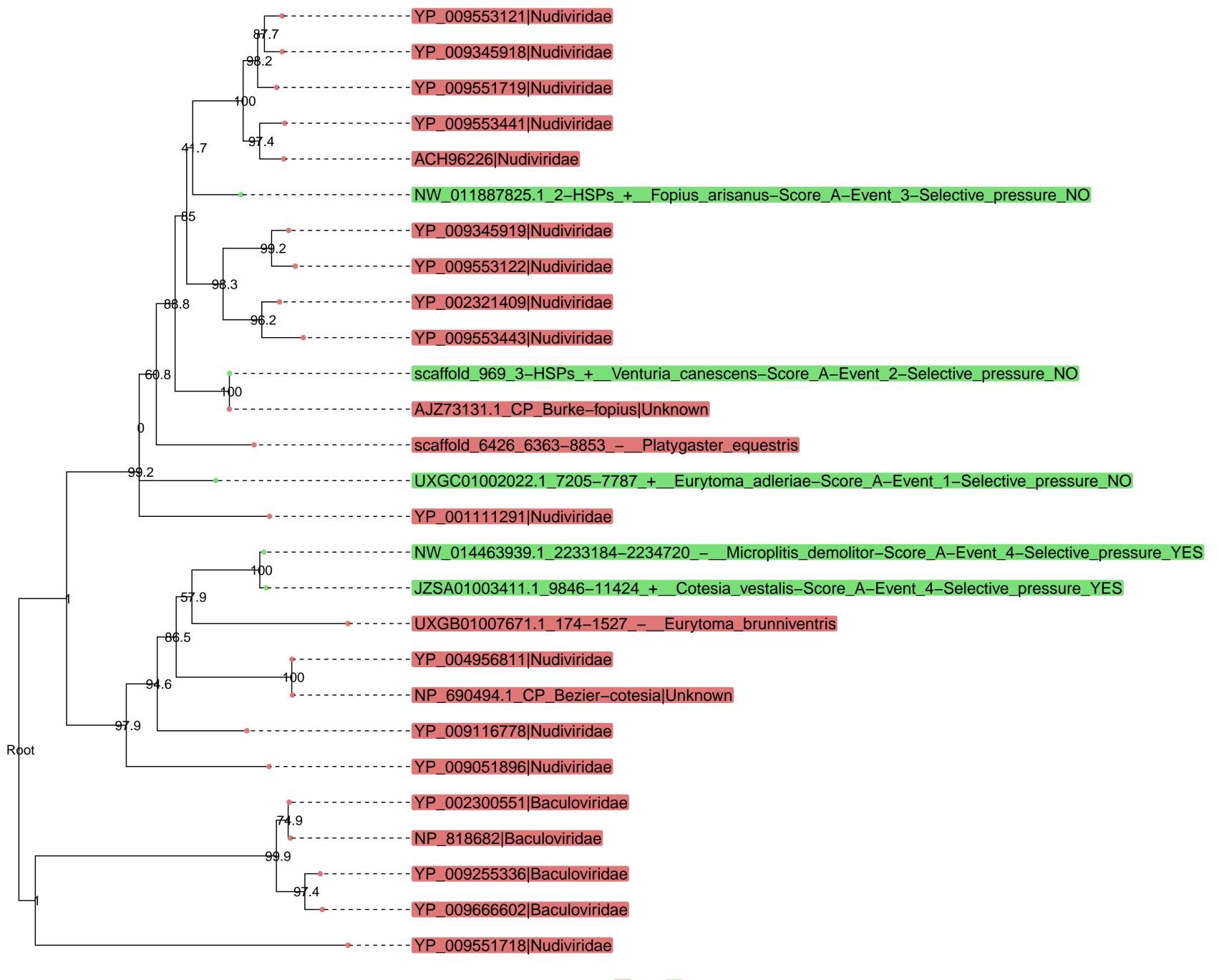


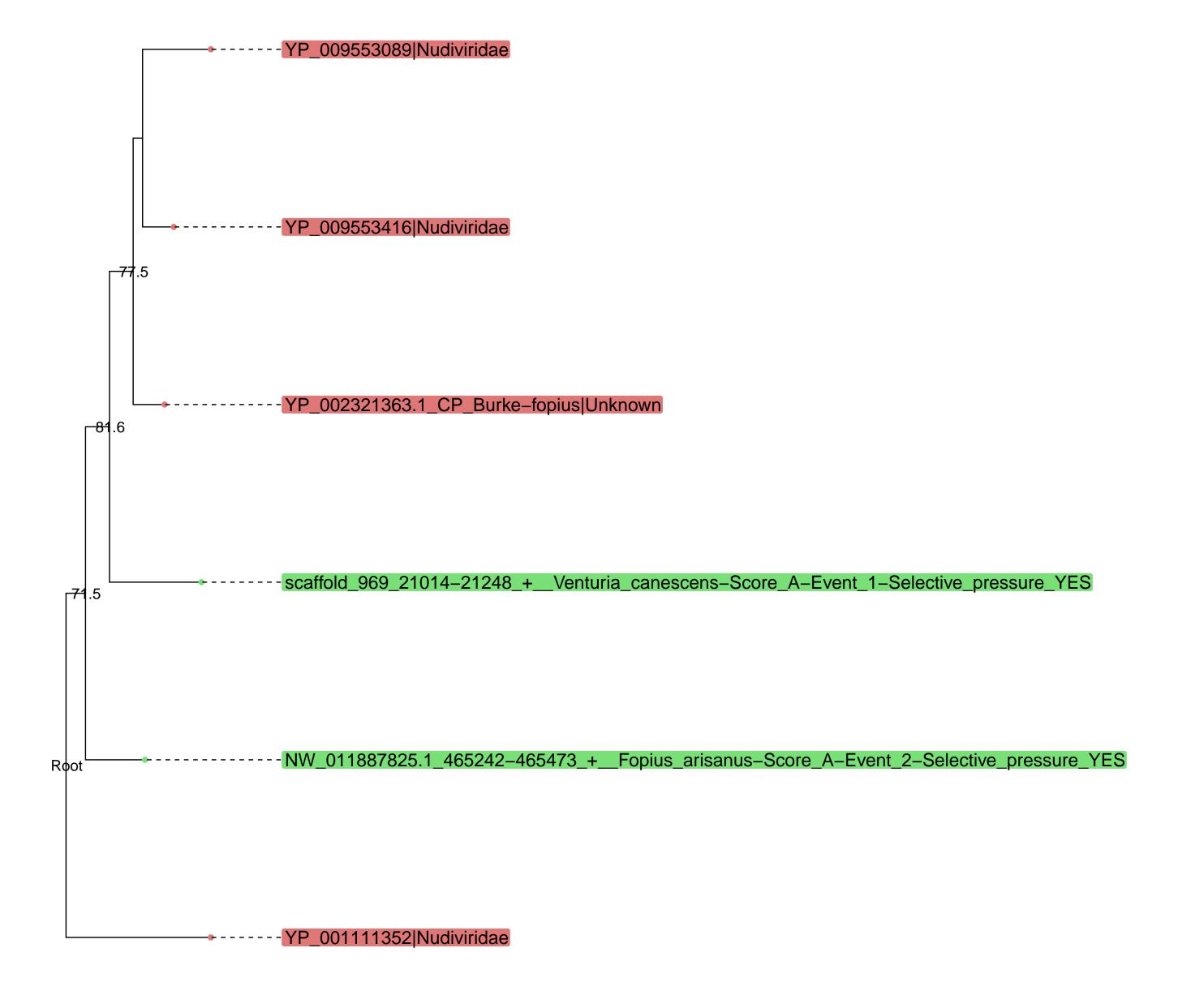


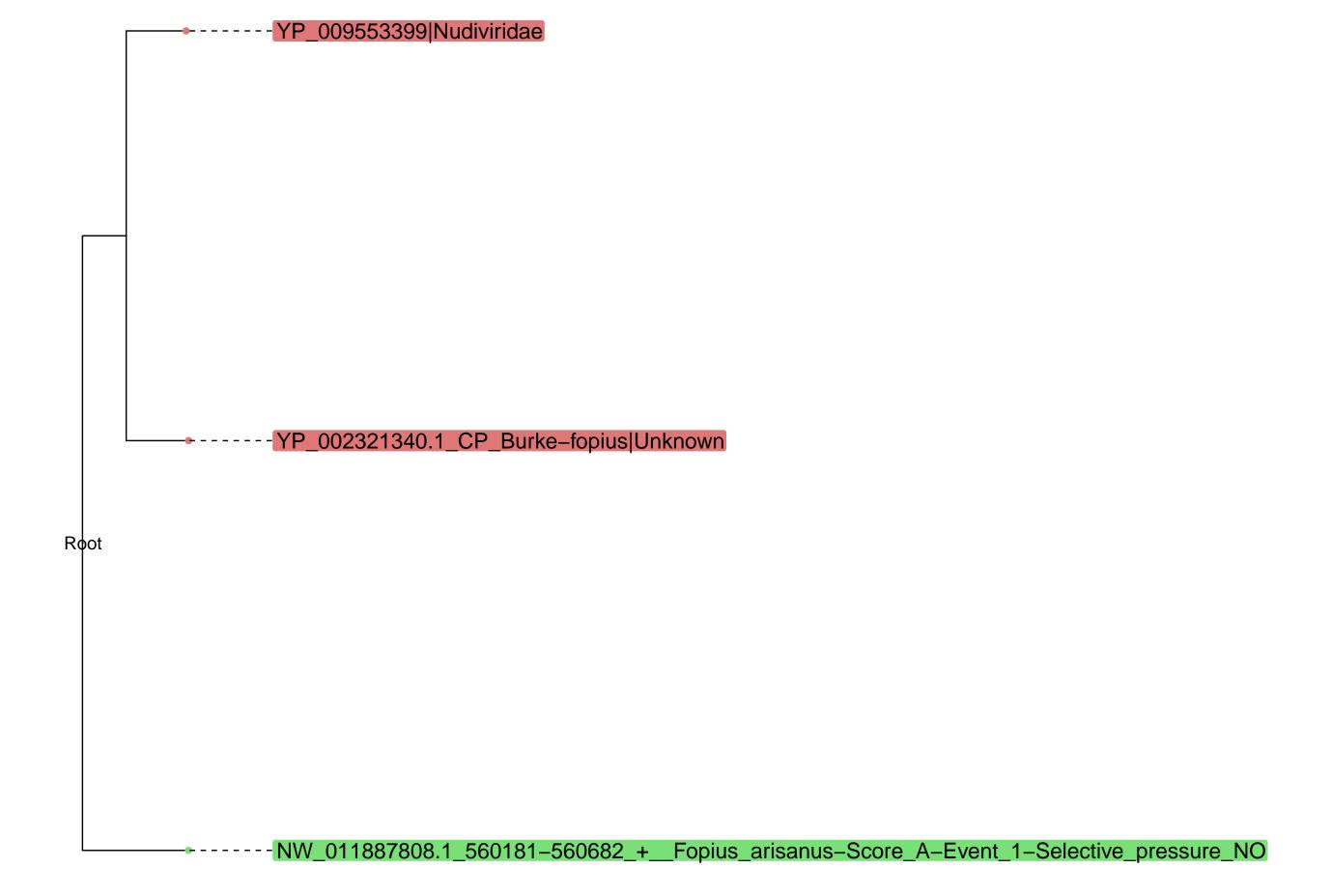


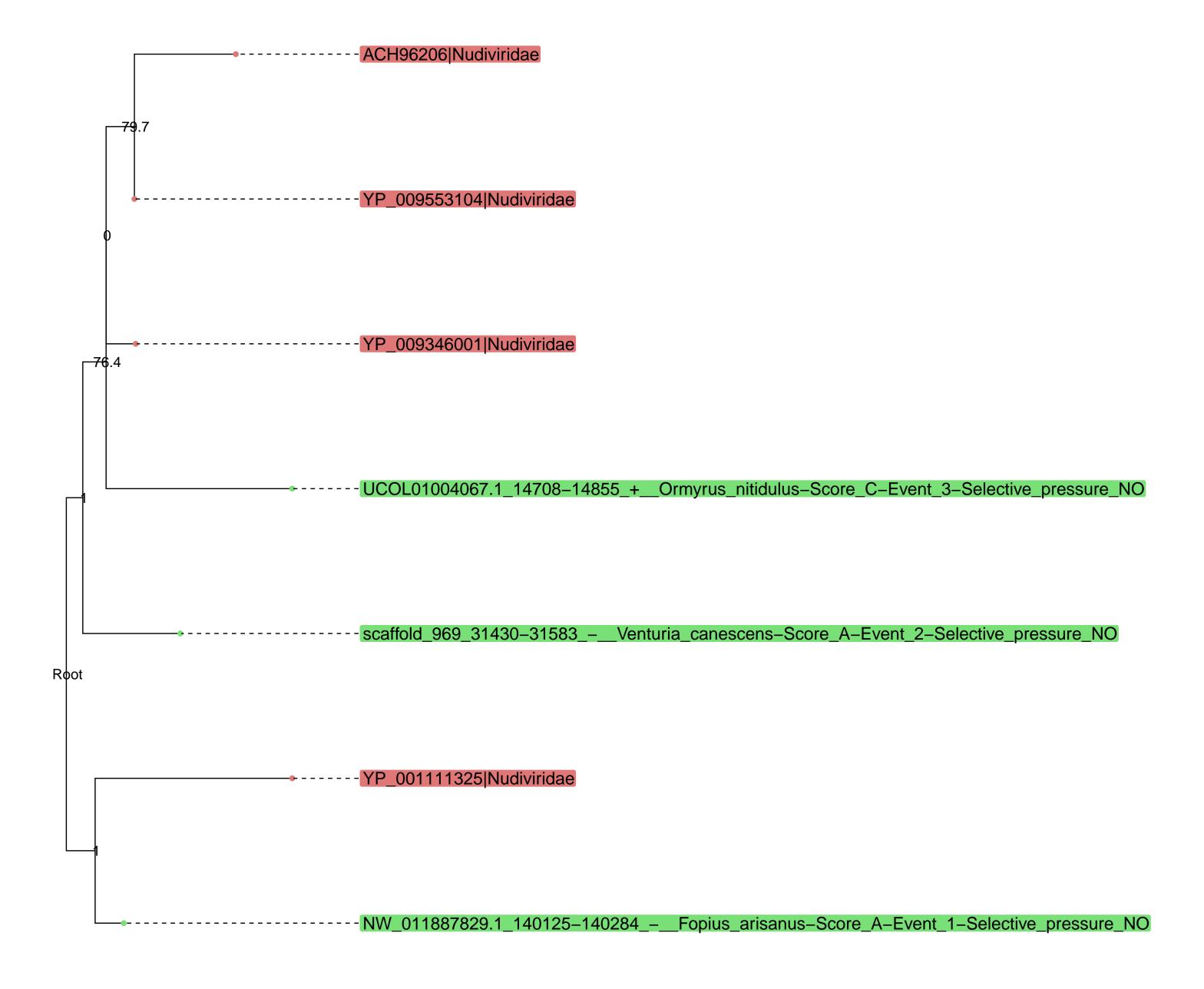


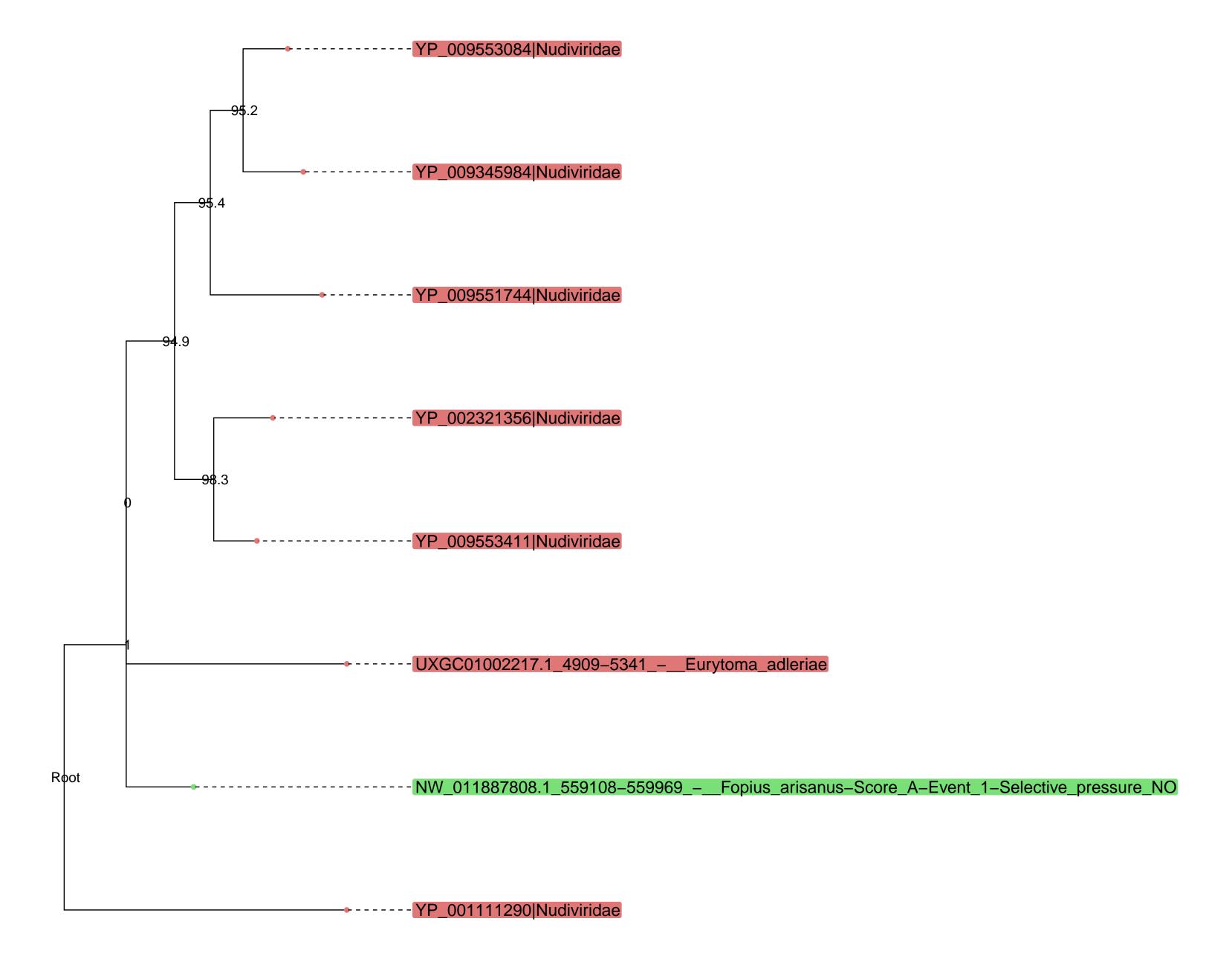


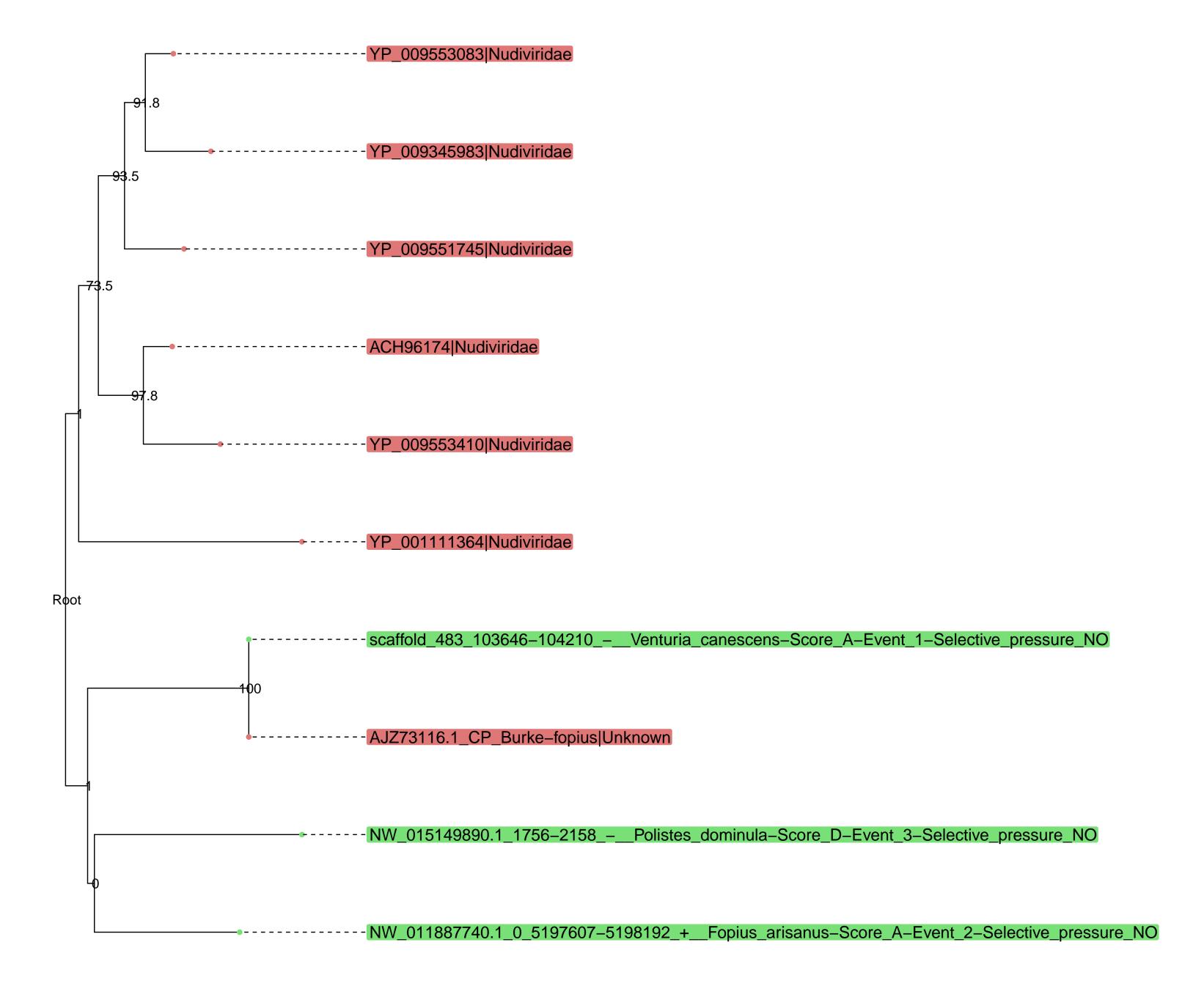


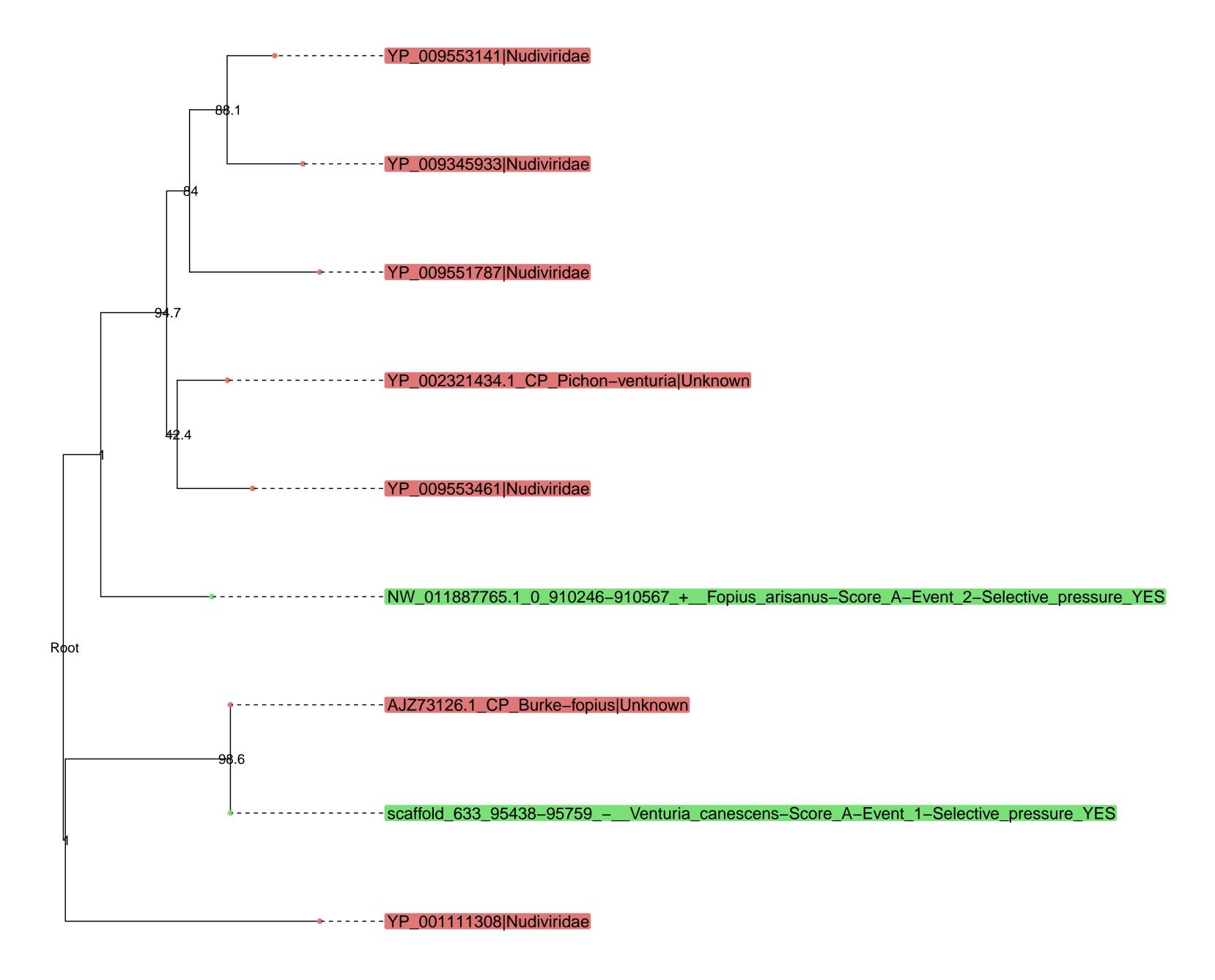


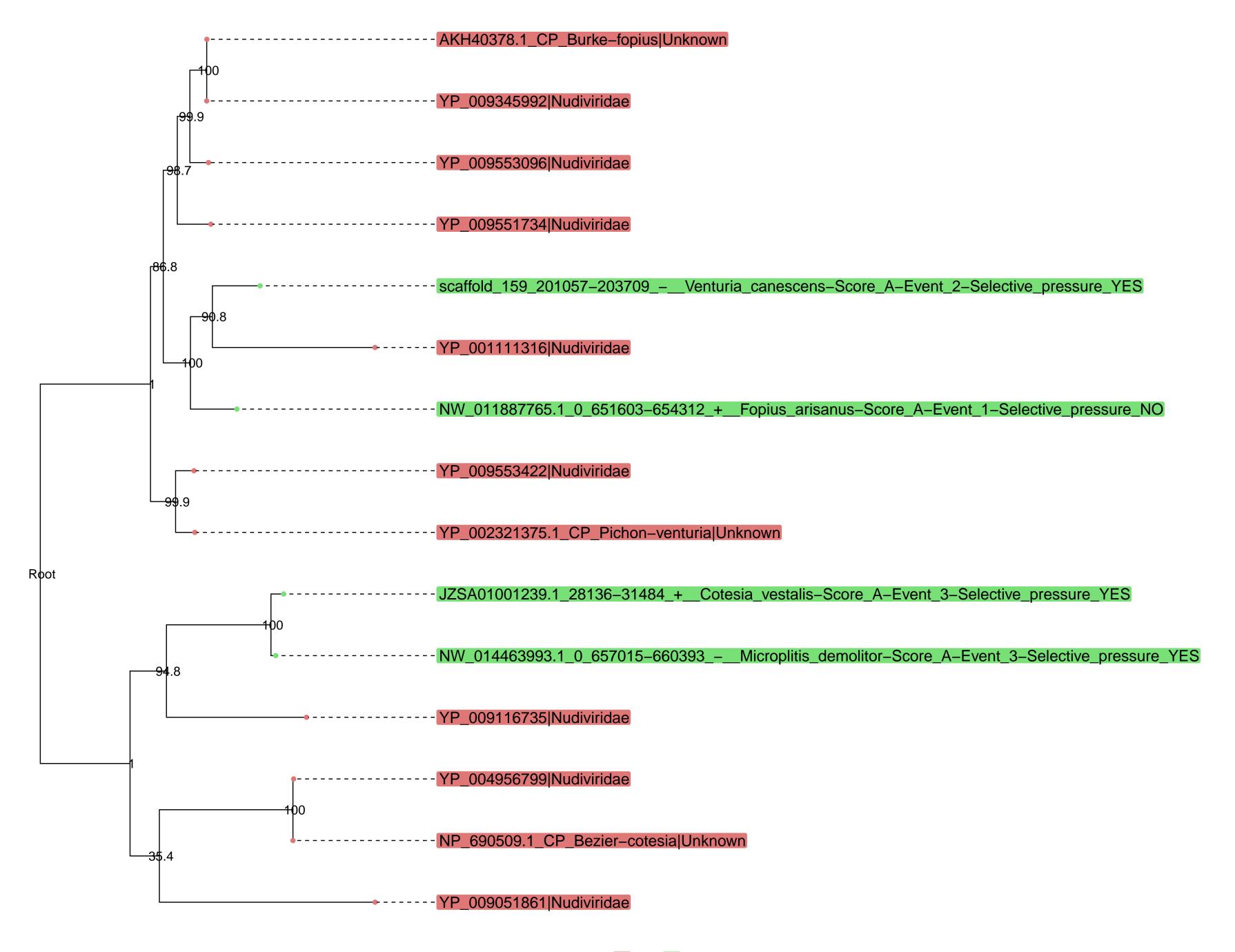


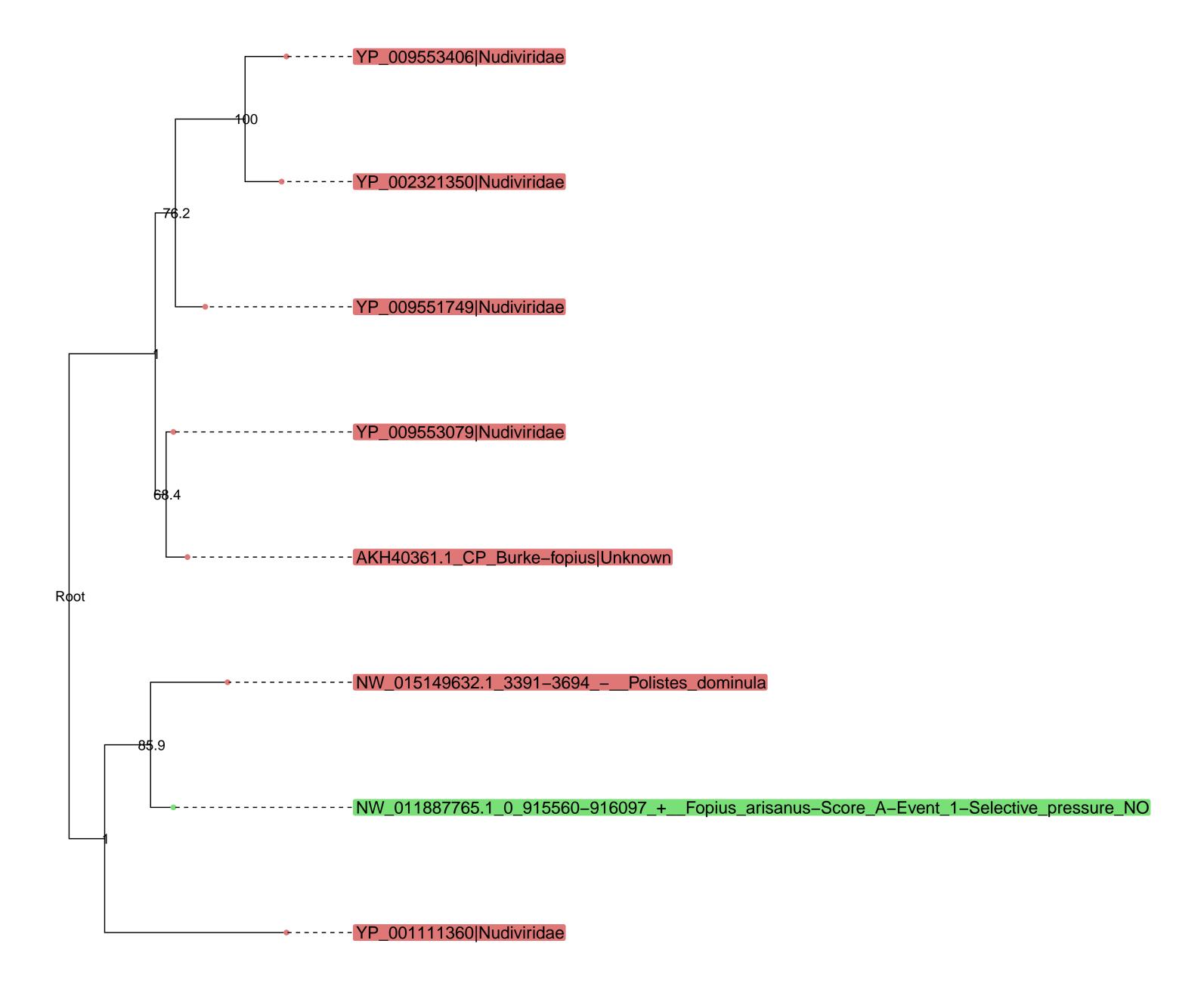


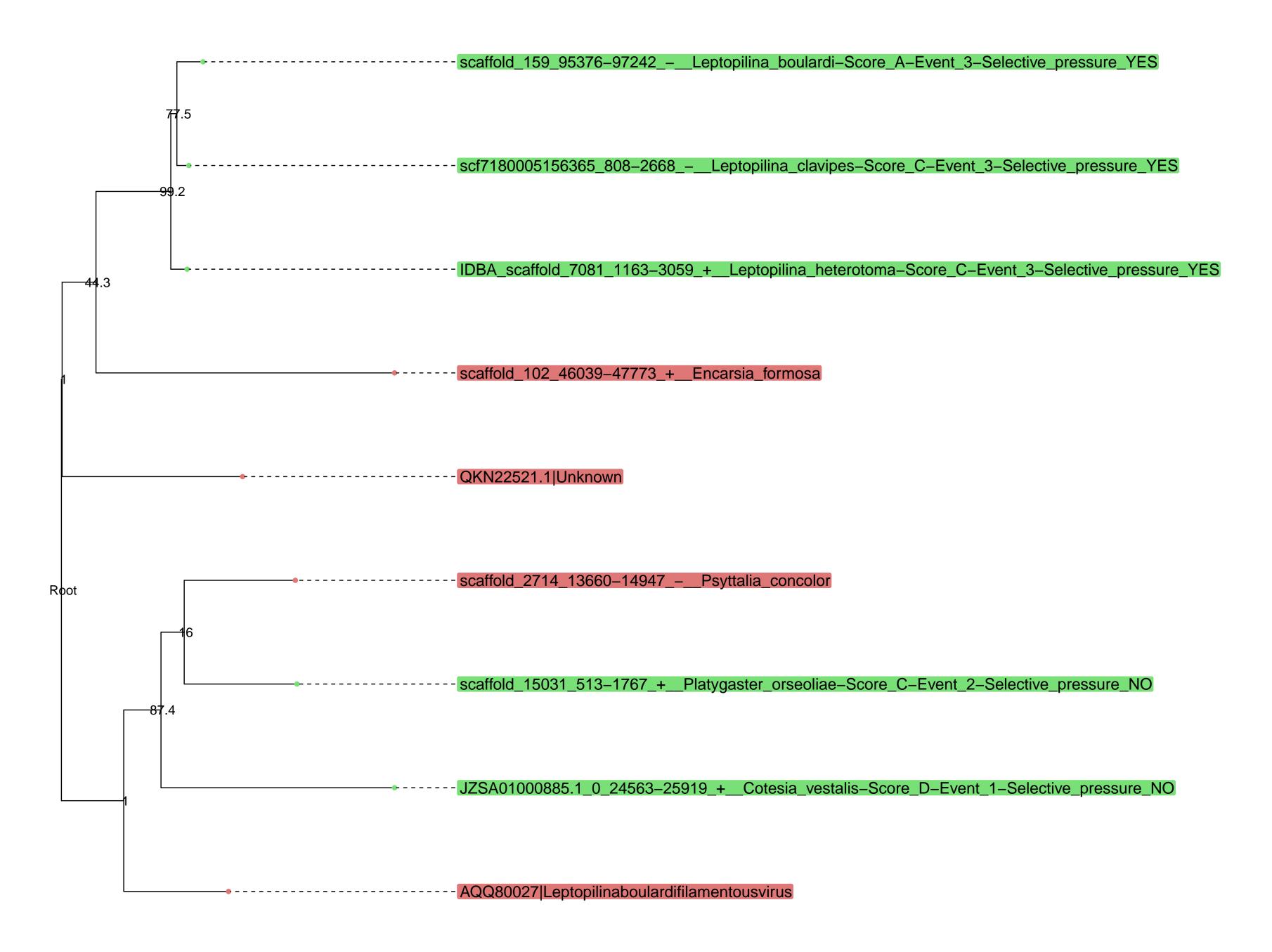




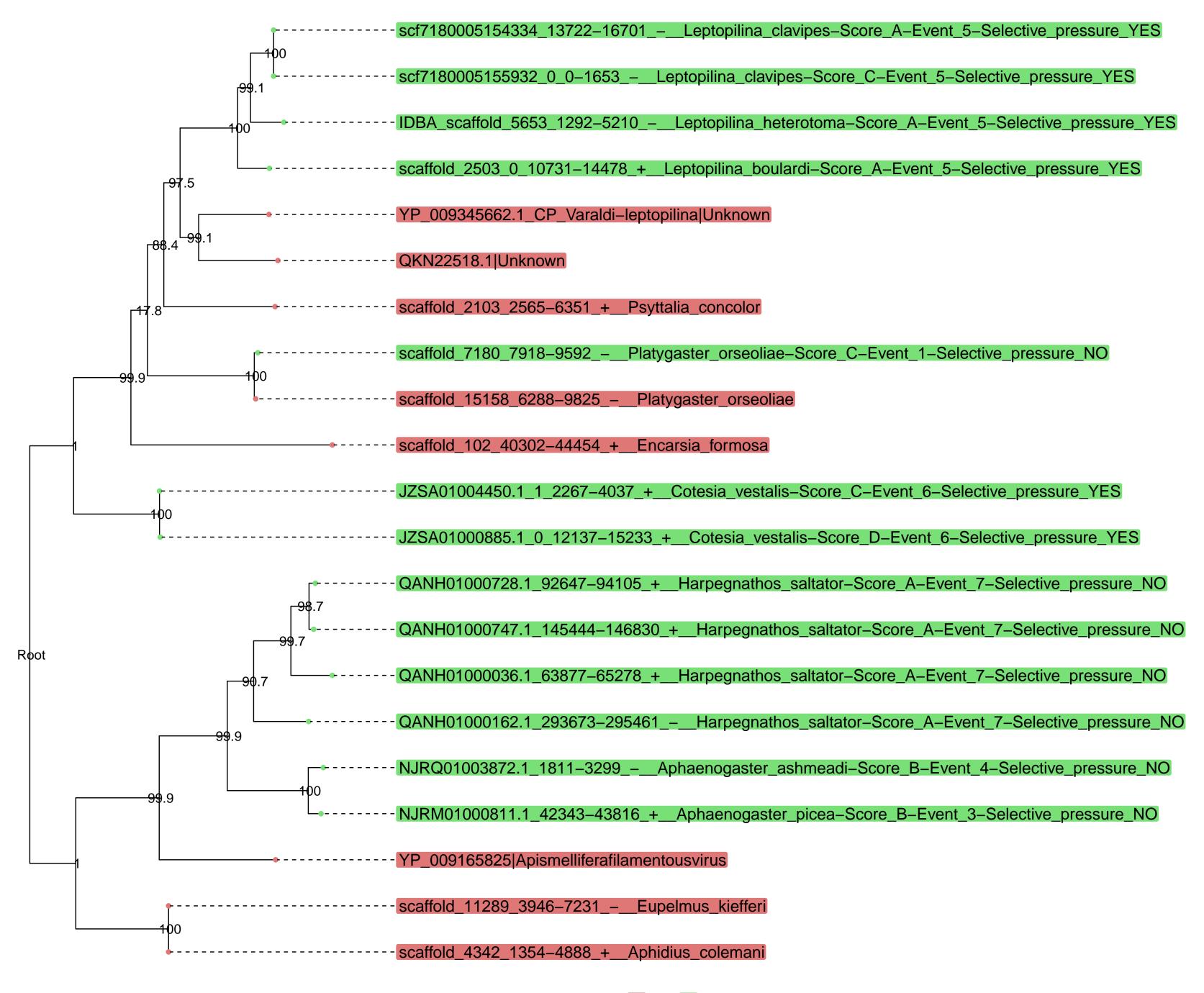


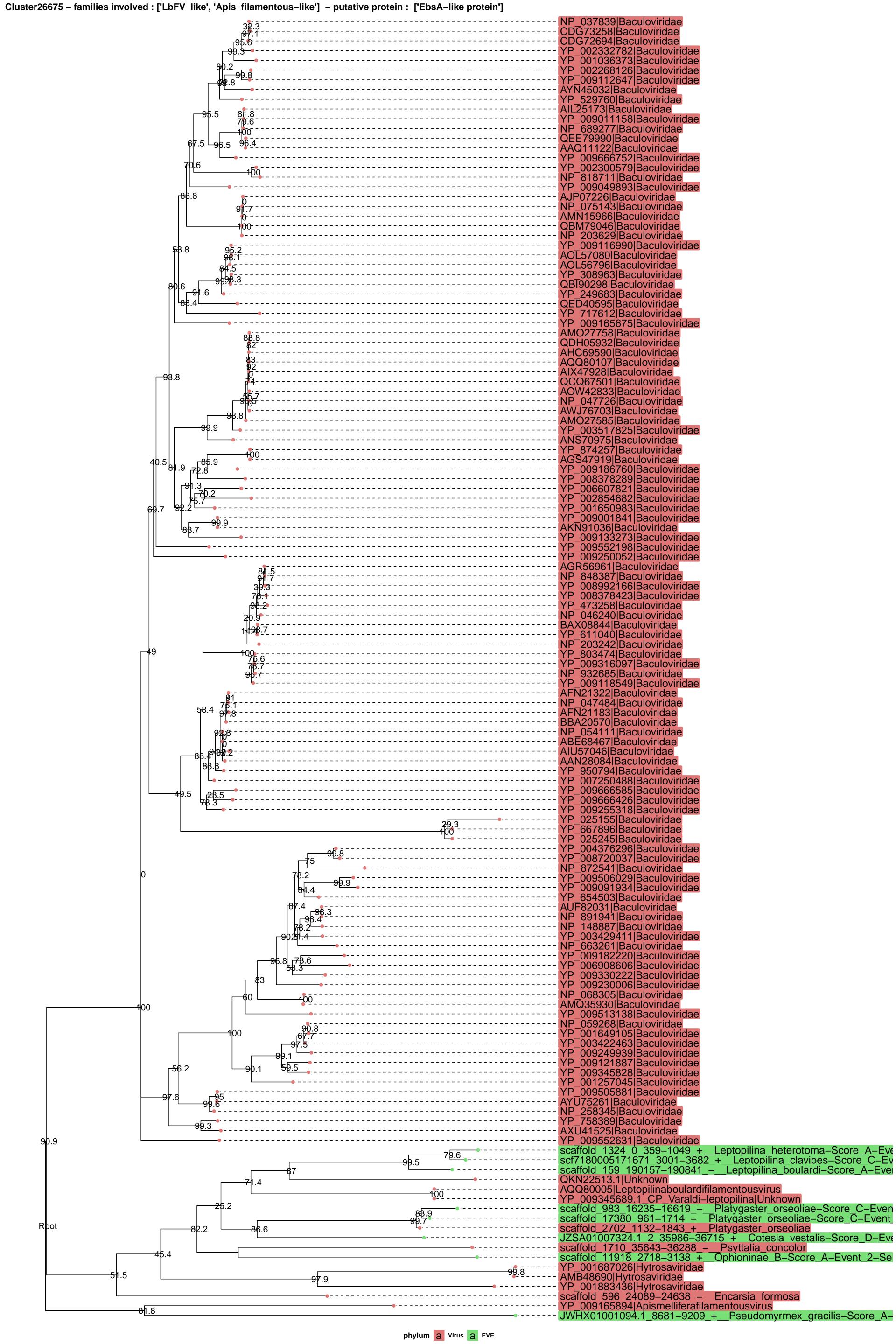


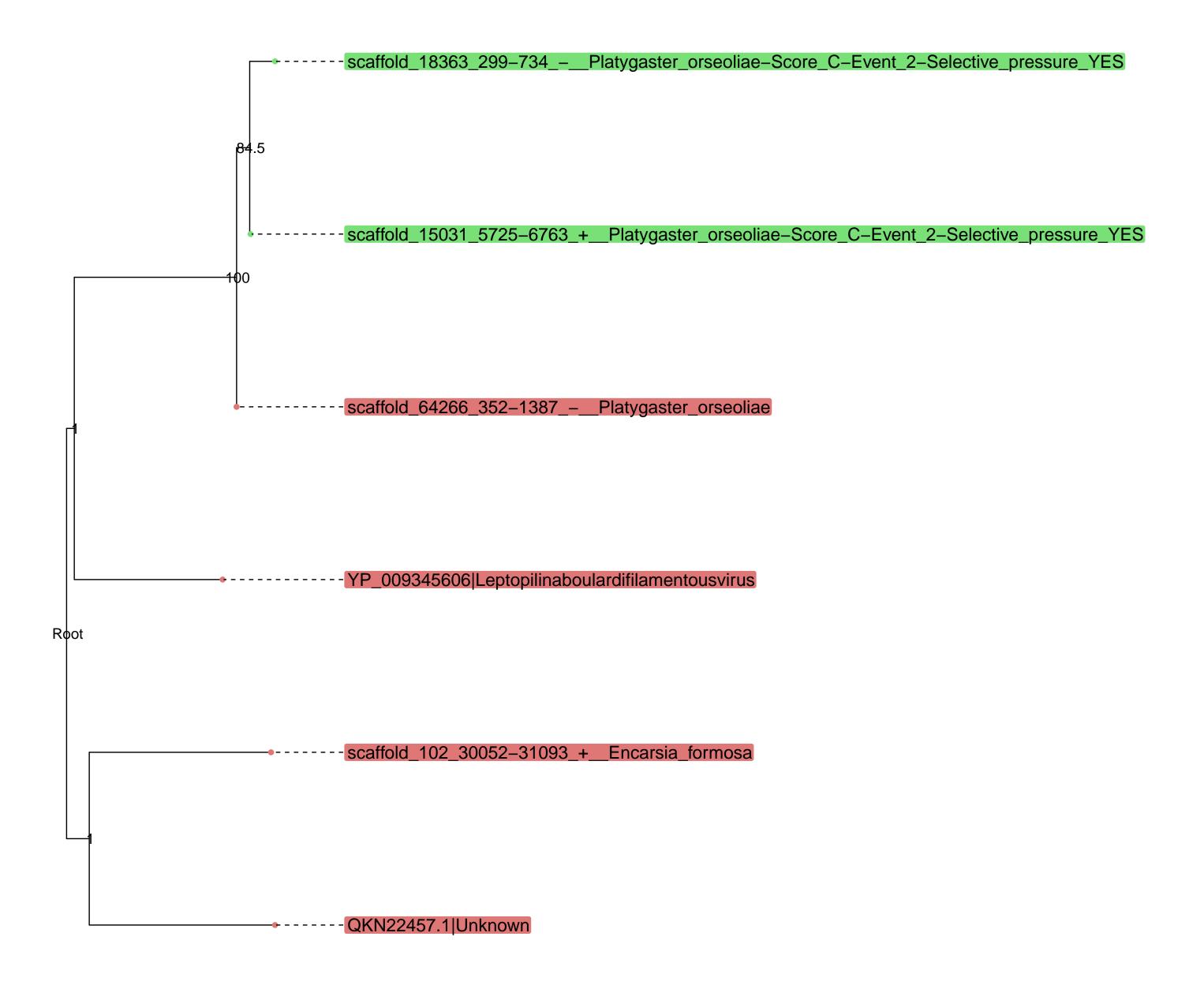


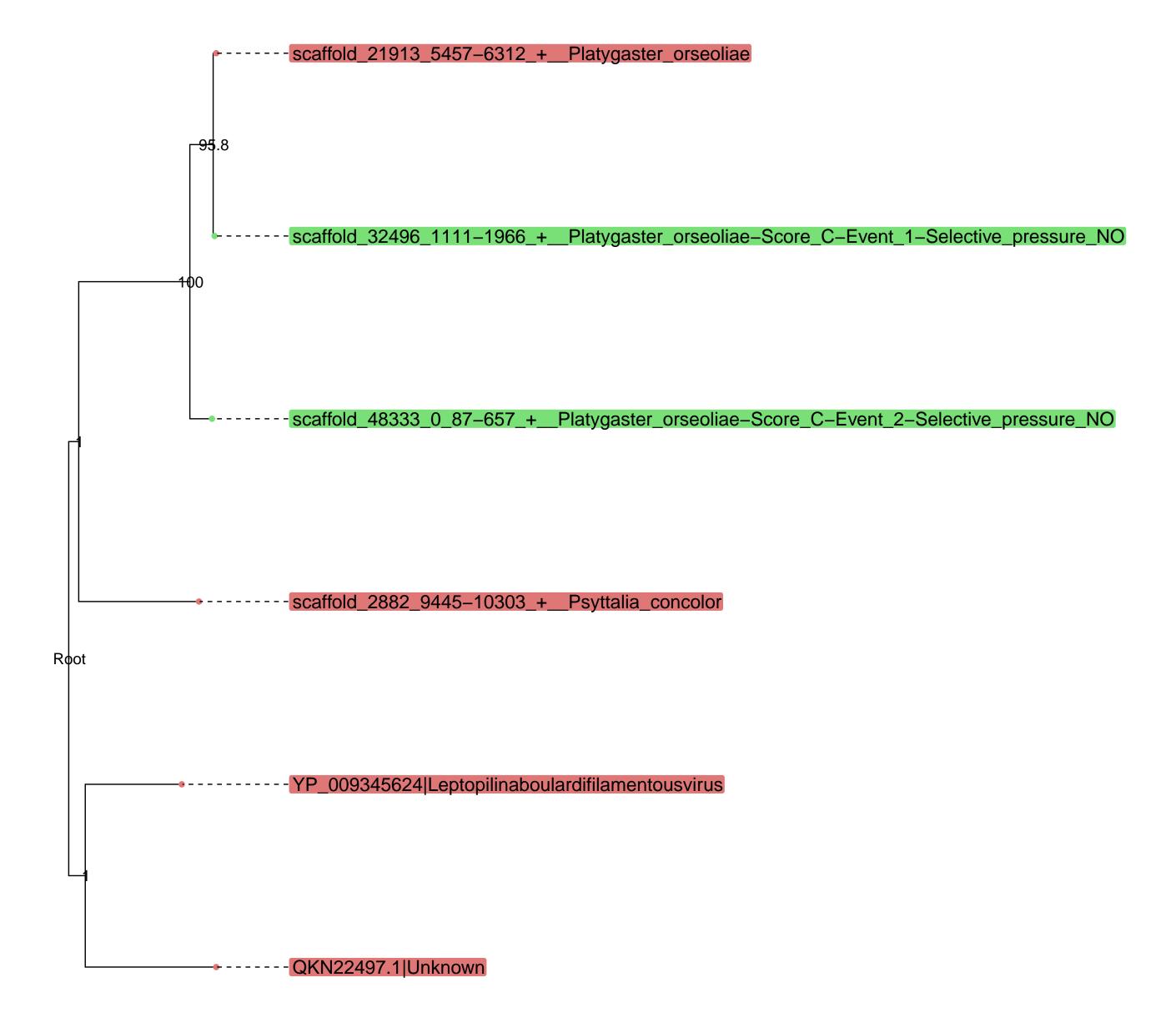


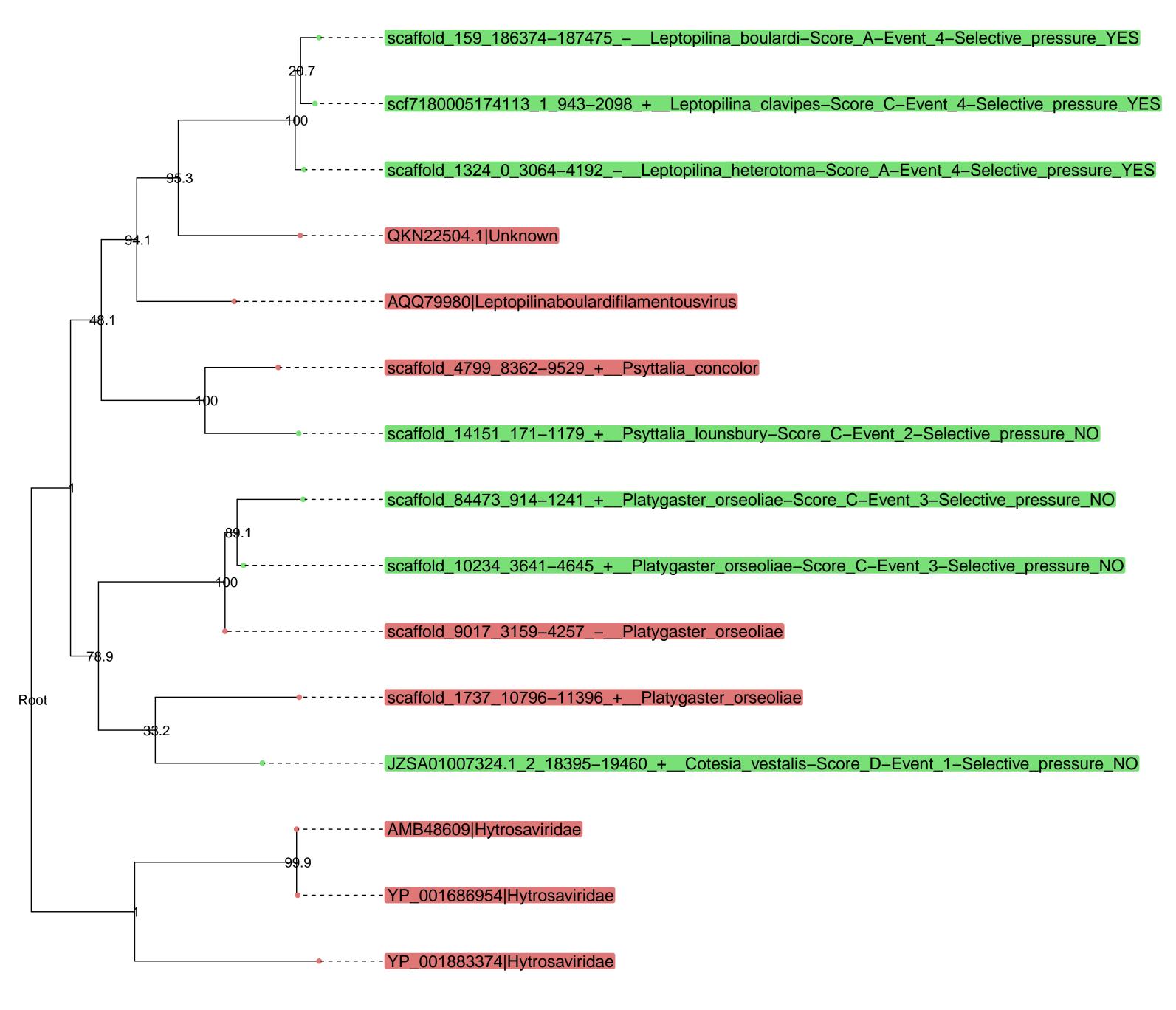
Cluster5269 – families involved : ['LbFV_like'] – putative protein : [nan]

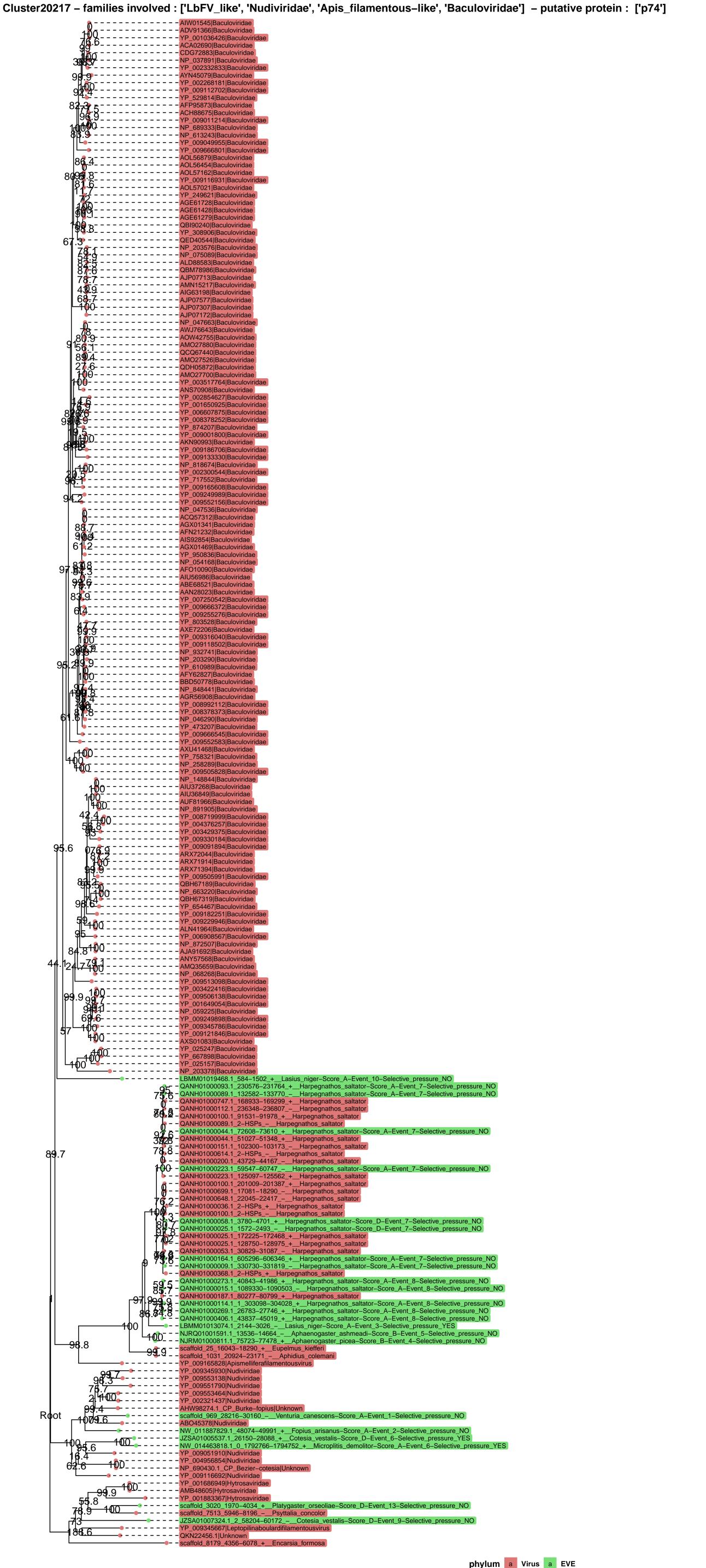


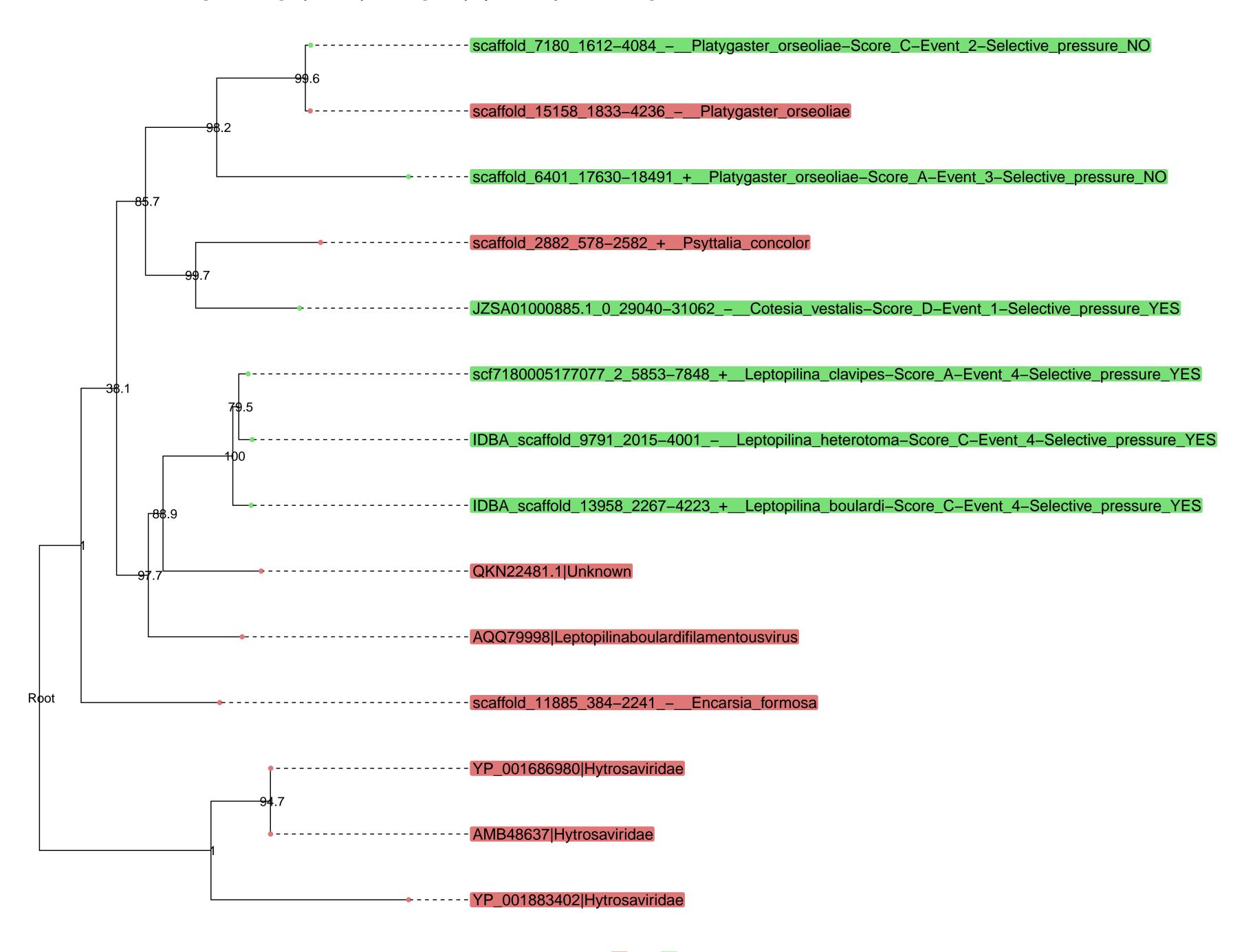


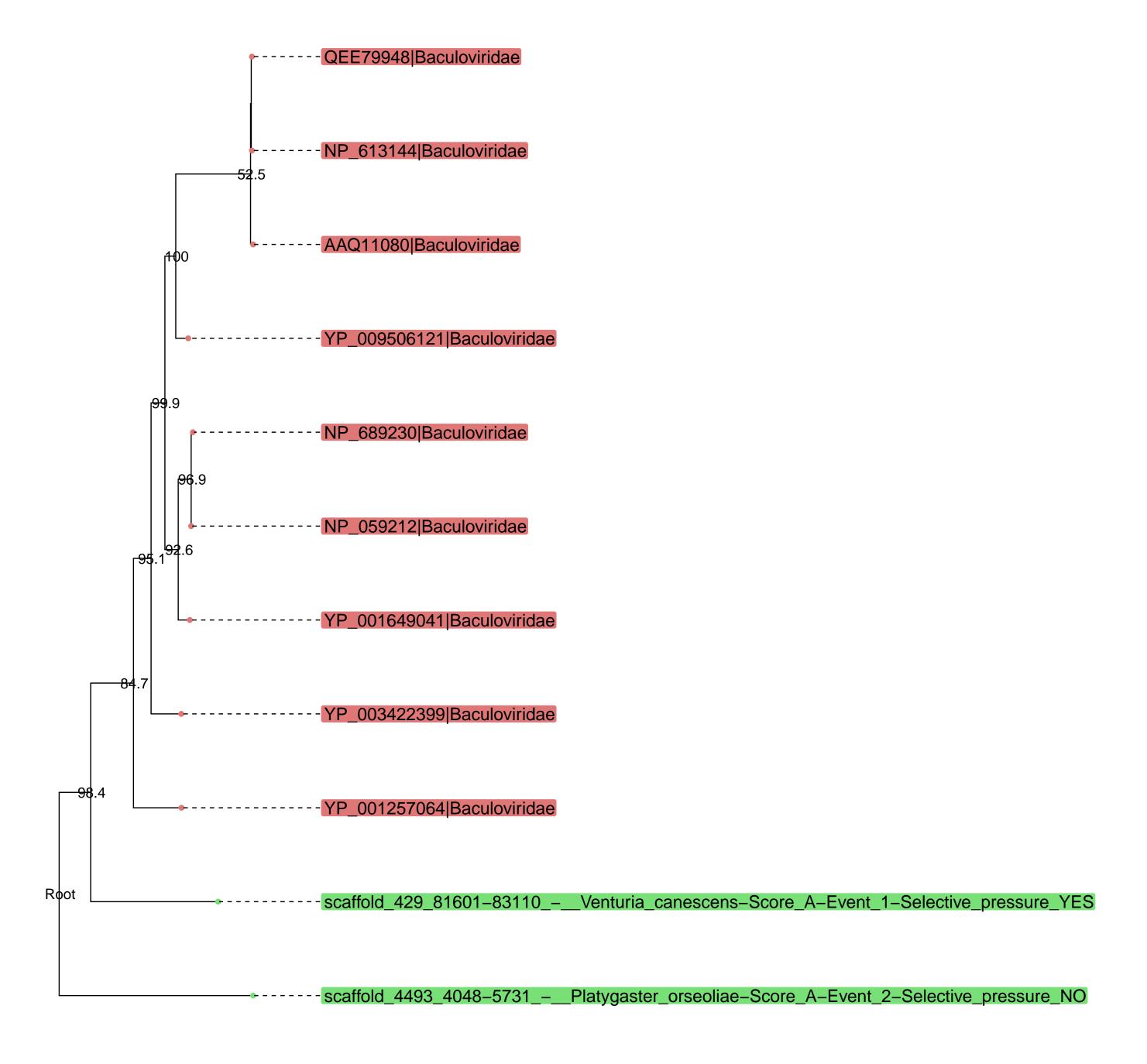




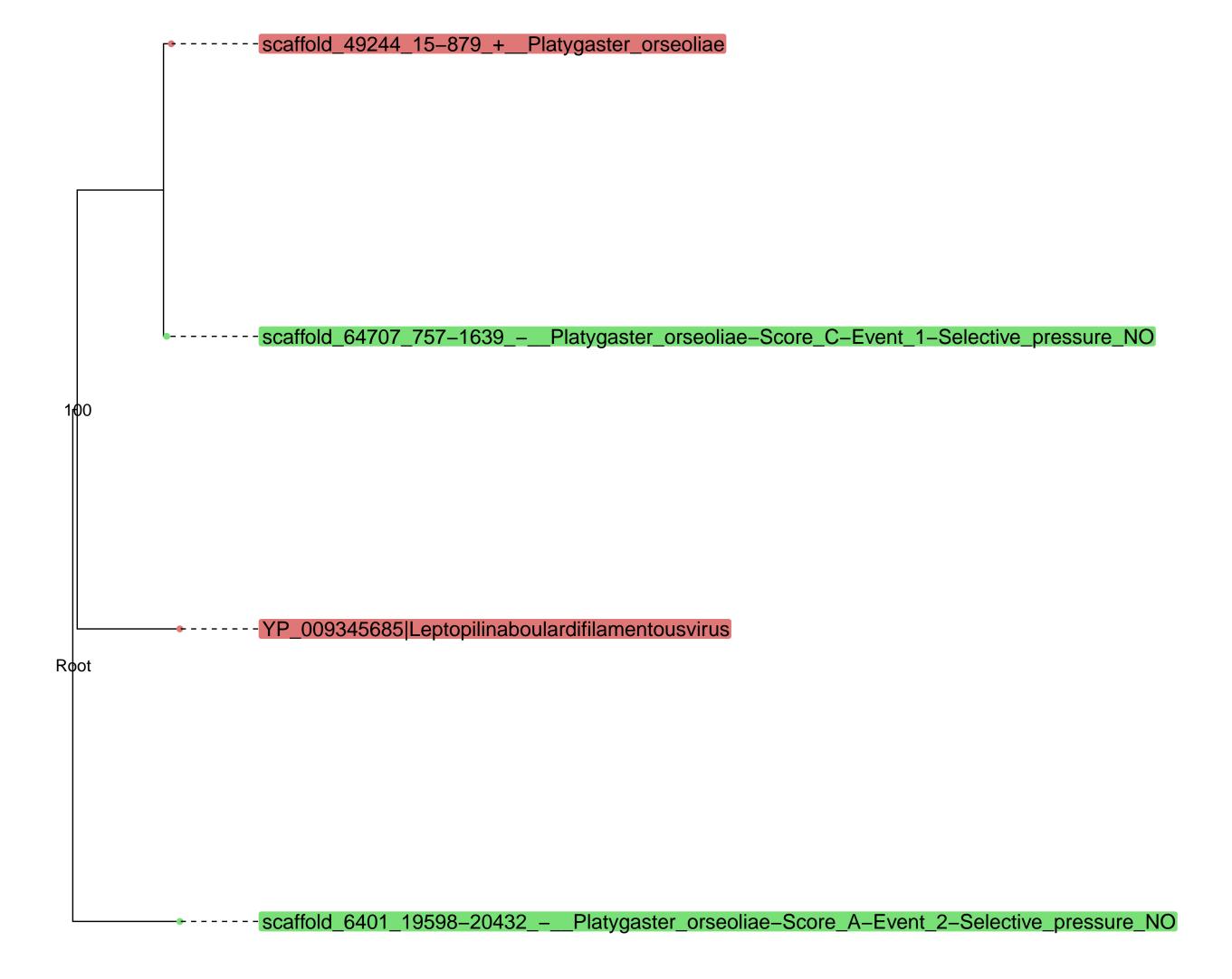


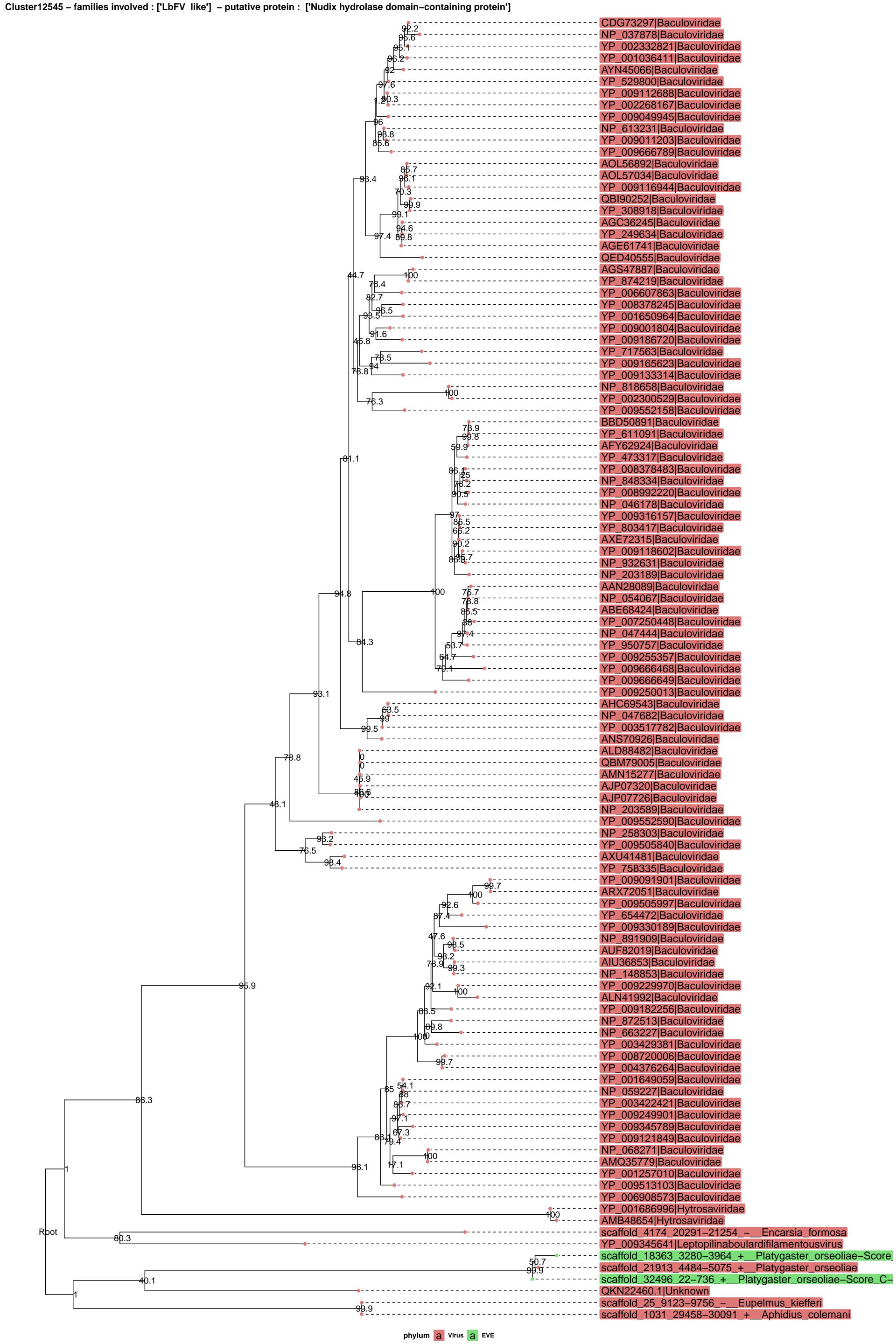


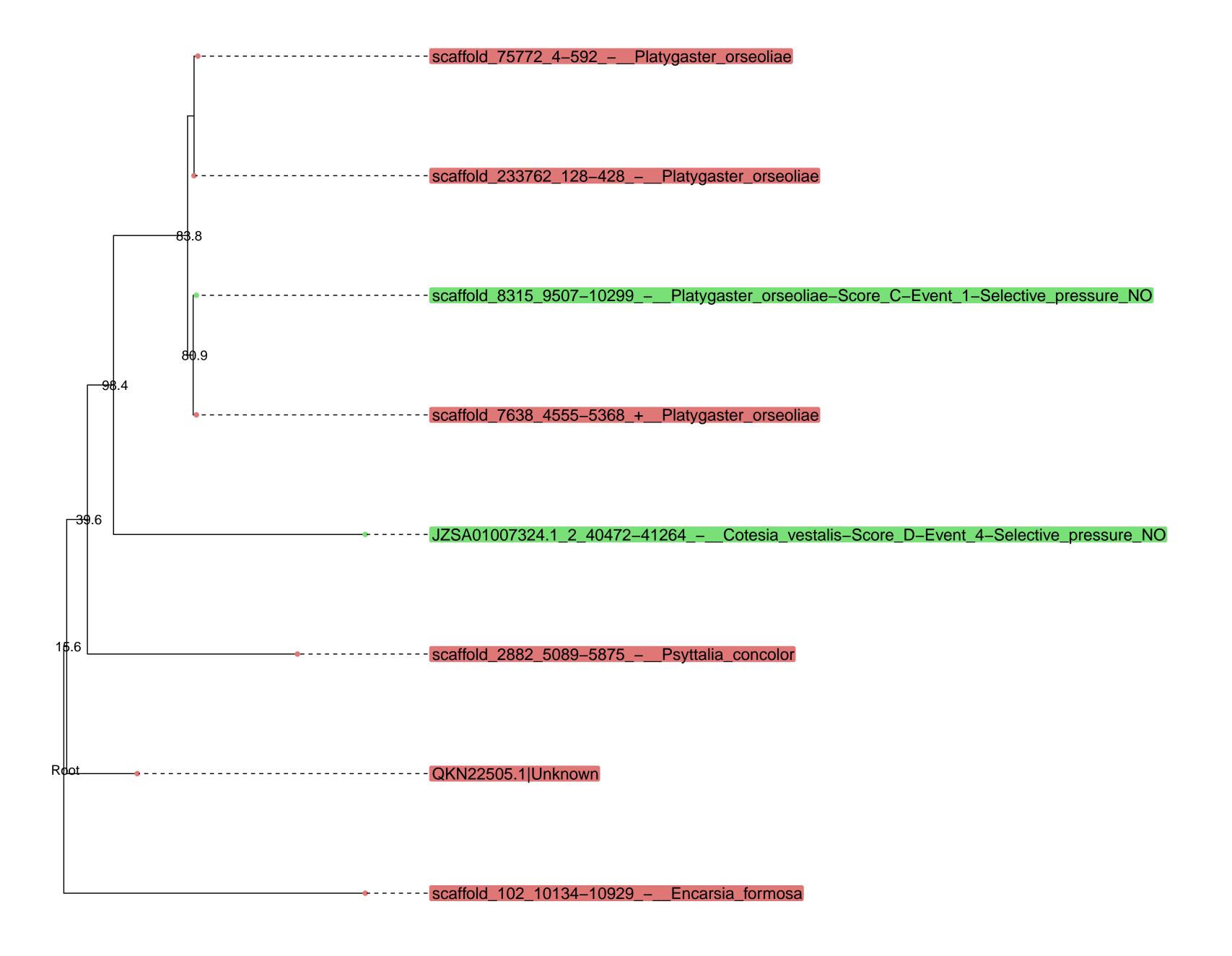


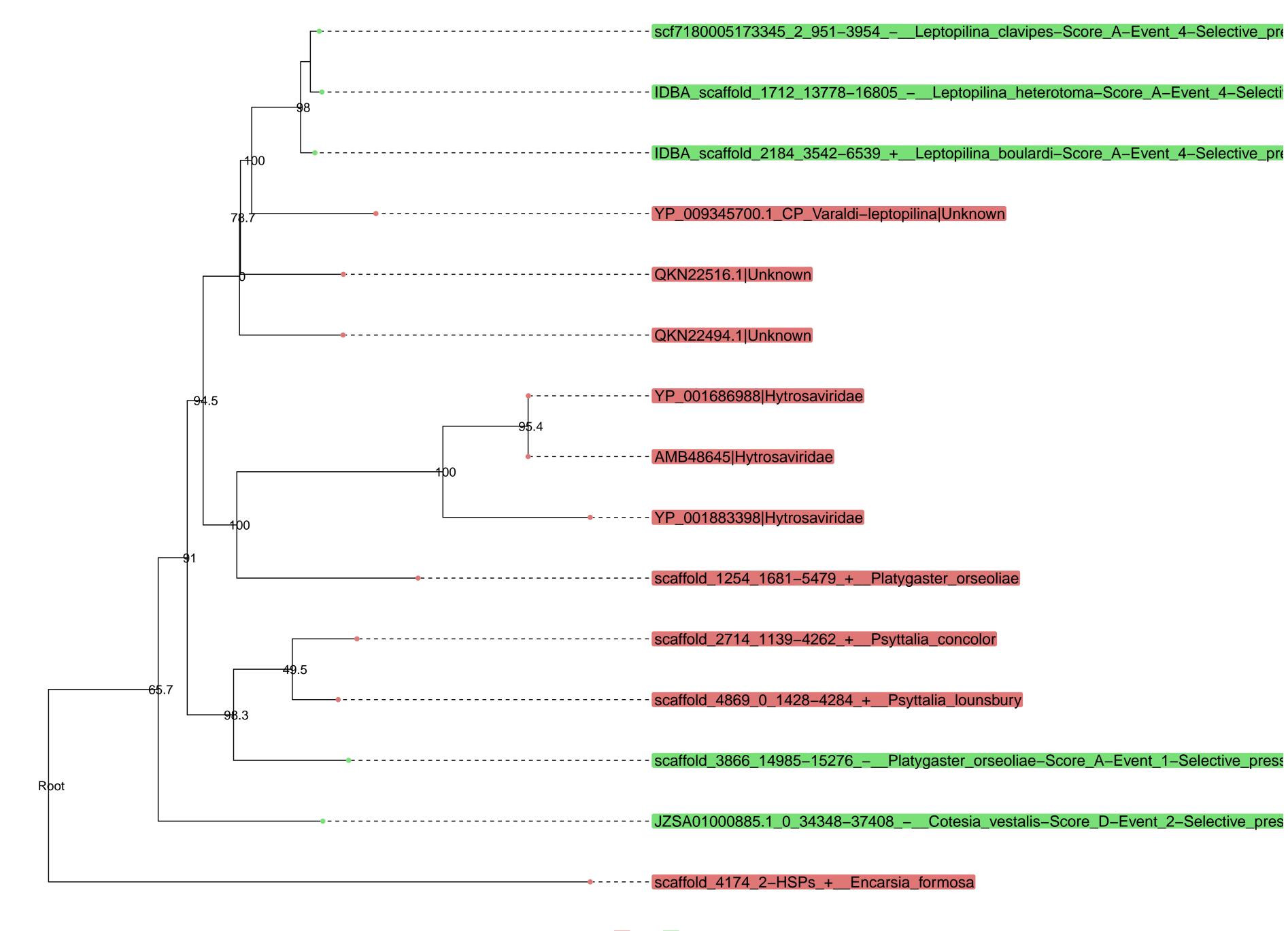


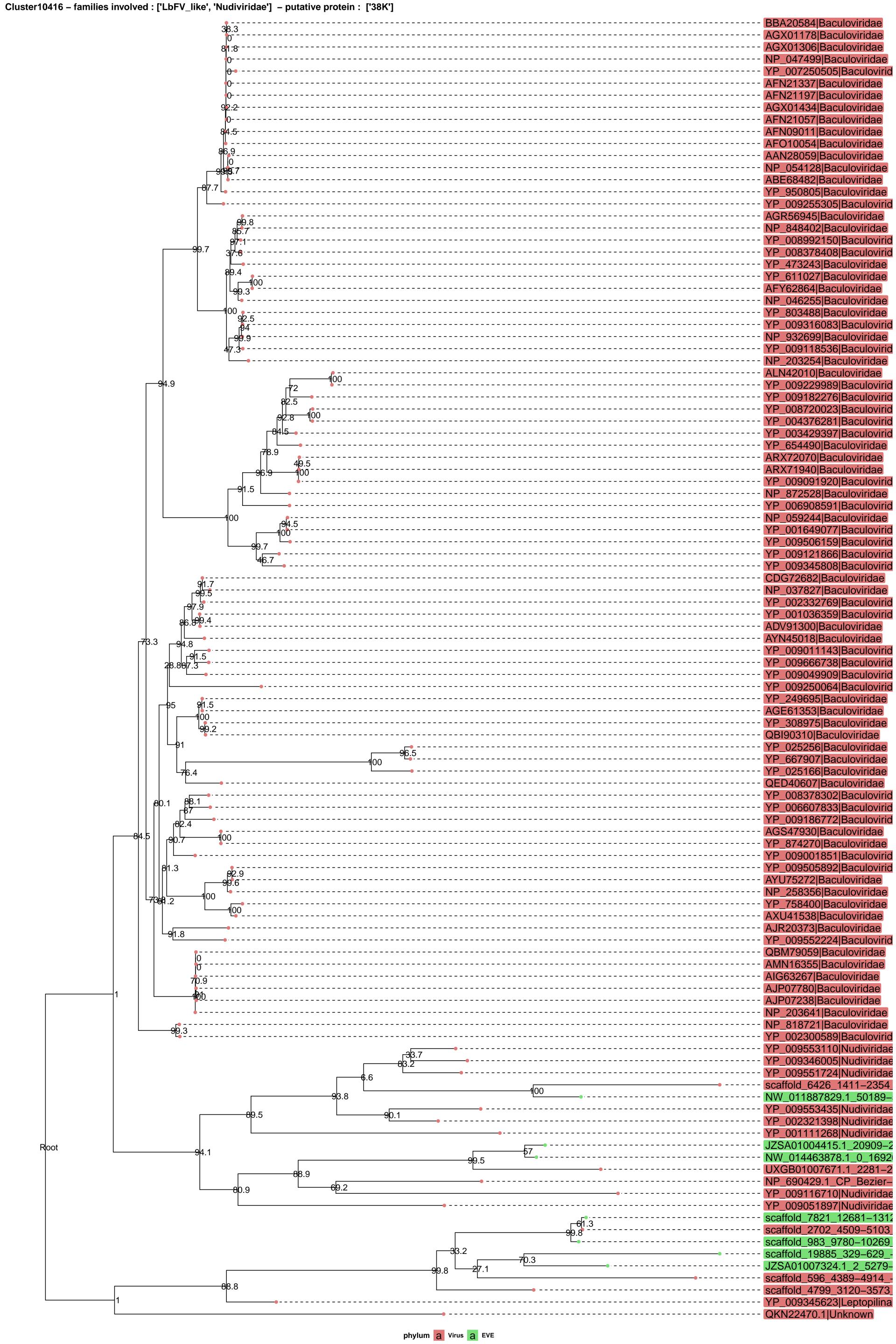


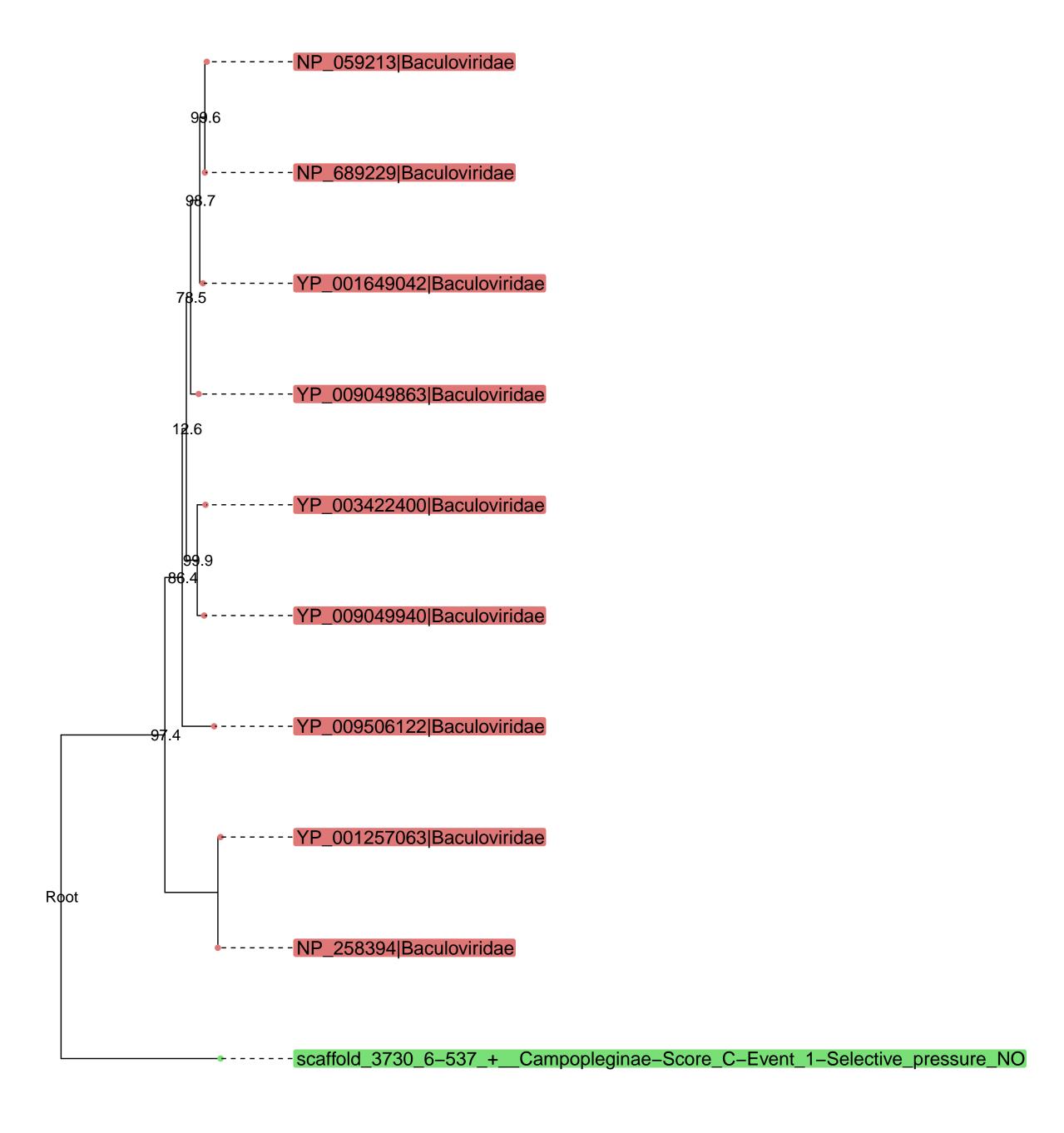


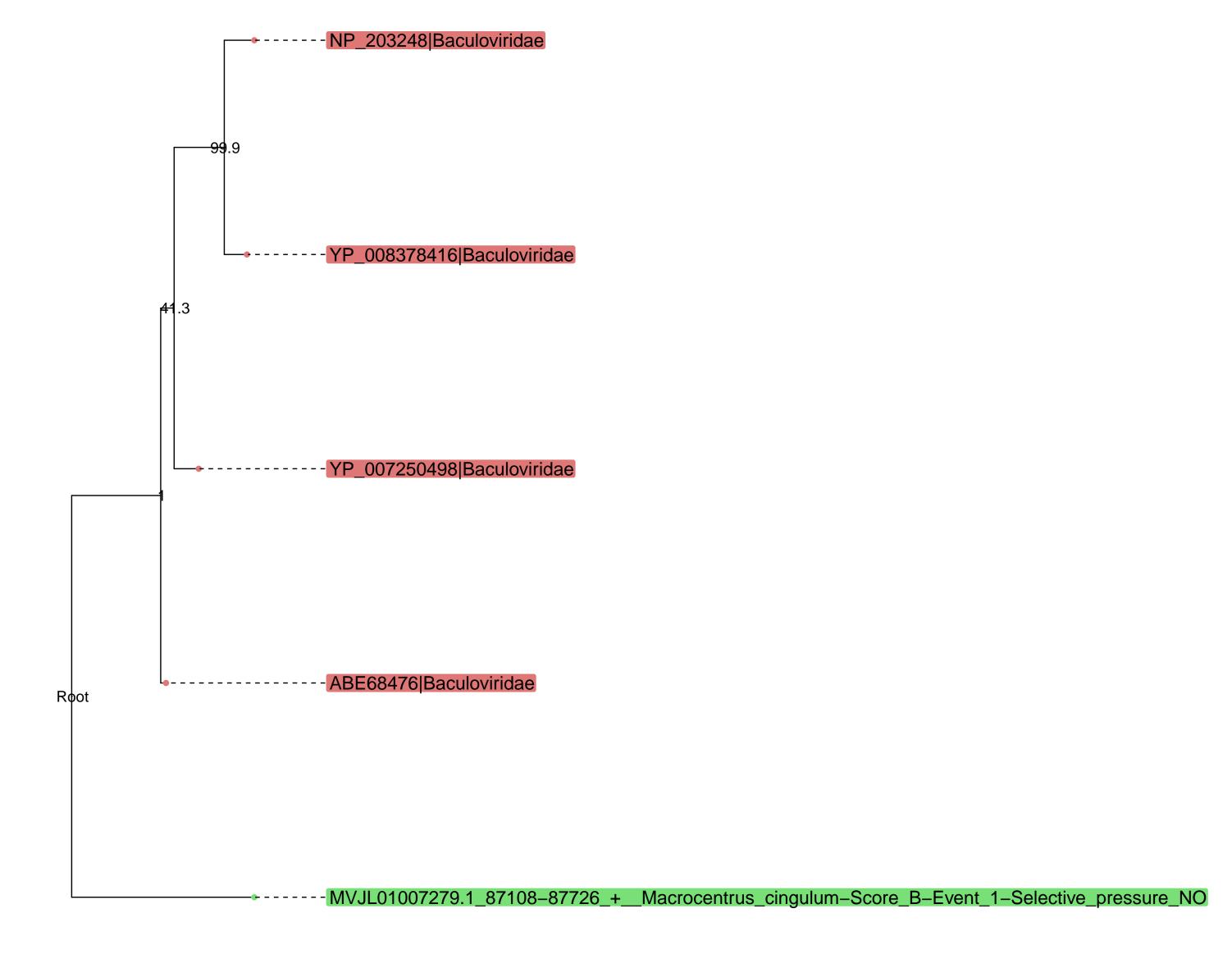












scaffold_11405_7622-8171_+__Copidosoma_sp-Score_A-Event_1-Selective_pressure_NO

-YP_009666553|Baculoviridae