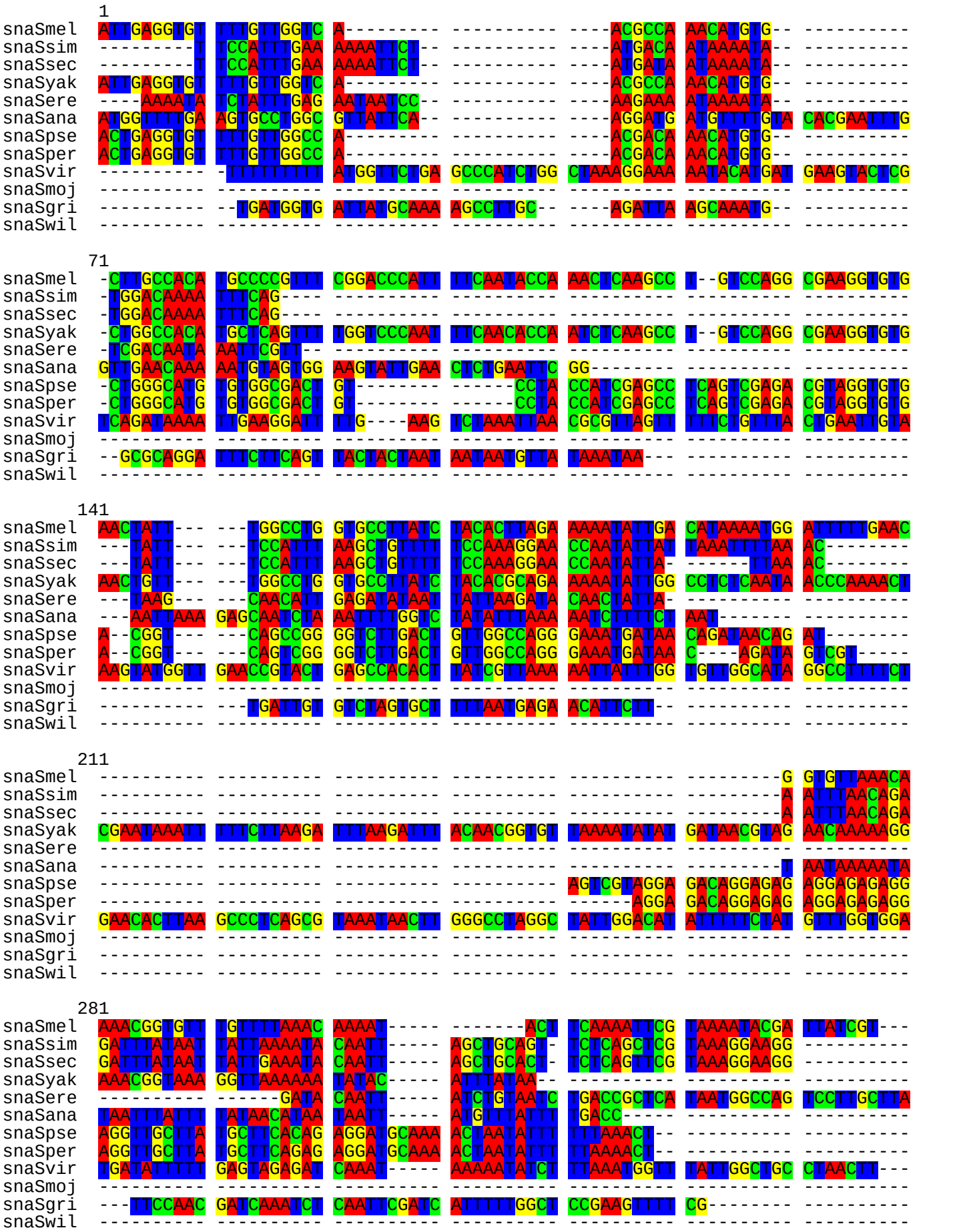
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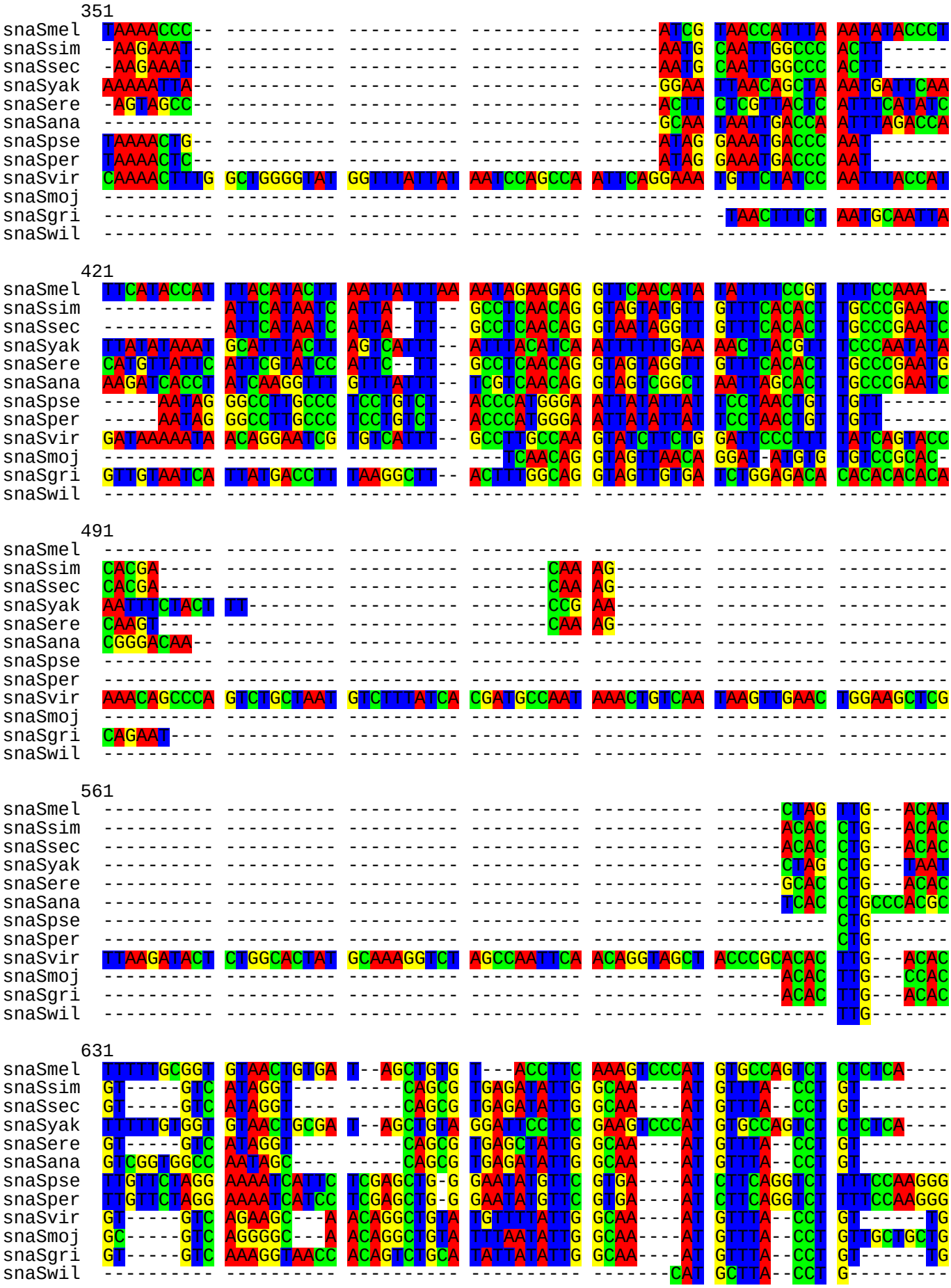
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snaSsim -AAGAAAT-- ---------- ---------- ---------- ------AATG CAATTGGCCC ACTT------
snaSsec -AAGAAAT-- ---------- ---------- ---------- ------AATG CAATTGGCCC ACTT------
snaSyak AAAAATTA-- ---------- ---------- ---------- ------GGAA TTAACAGCTA AATGATTCAA
snaSere -AGTAGCC-- ---------- ---------- ---------- ------ACTT CTCGTTACTC ATTTCATATC
snaSana ---------- ---------- ---------- ---------- ------GCAA TAATTGACCA ATTTAGACCA
snaSpse TAAAACTG-- ---------- ---------- ---------- ------ATAG GAAATGACCC AAT-------
snaSper TAAAACTC-- ---------- ---------- ---------- ------ATAG GAAATGACCC AAT-------
snaSvir CAAAACTTTG GCTGGGGTAT GGTTTATTAT AATCCAGCCA ATTCAGGAAA TGTTCTATCC AATTTACCAT
snaSmoj ---------- ---------- ---------- ---------- ---------- ---------- ----------
snaSgri ---------- ---------- ---------- ---------- ---------- -TAACTTTCT AATGCAATTA
snaSwil ---------- ---------- ---------- ---------- ---------- ---------- ----------
421
snaSmel TTCATACCAT TTACATACTT AATTATTTAA AATAGAAGAG GTTCAACATA TATTTTCCGT TTTCCAAA--
snaSsim ---------- ATTCATAATC ATTA--TT-- GCCTCAACAG GTAGTATGTT GTTTCACACT TGCCCGAATC
snaSsec ---------- ATTCATAATC ATTA--TT-- GCCTCAACAG GTAATAGGTT GTTTCACACT TGCCCGAATC
snaSyak TTATATAAAT GCATTTACTT AGTCATTT-- ATTTACATCA ATTTTTTGAA AACTTACGTT TCCCAATATA
snaSere CATGTTATTC ATTCGTATCC ATTC--TT-- GCCTCAACAG GTAGTAGGTT GTTTCACACT TGCCCGAATG
snaSana AAGATCACCT ATCAAGGTTT GTTTATTT-- TCGTCAACAG GTAGTCGGCT AATTAGCACT TGCCCGAATC
snaSpse -----AATAG GGCCTTGCCC TCCTGTCT-- ACCCATGGGA ATTATATTAT TCCTAACTGT TGTT------
snaSper -----AATAG GGCCTTGCCC TCCTGTCT-- ACCCATGGGA ATTATATTAT TCCTAACTGT TGTT------
snaSvir GATAAAAATA ACAGGAATCG TGTCATTT-- GCCTTGCCAA GTATCTTCTG GATTCCCTTT TATCAGTACC
snaSmoj ---------- ---------- ---------- ---TCAACAG GTAGTTAACA GGAT-ATGTG TGTCCGCAC-
snaSgri GTTGTAATCA TTATGACCTT TAAGGCTT-- ACTTTGGCAG GTAGTTGTGA TCTGGAGACA CACACACACA
snaSwil ---------- ---------- ---------- ---------- ---------- ---------- ----------
491
snaSmel ---------- ---------- ---------- ---------- ---------- ---------- ----------
snaSsim CACGA----- ---------- ---------- -------CAA AG-------- ---------- ----------
snaSsec CACGA----- ---------- ---------- -------CAA AG-------- ---------- ----------
snaSyak AATTTCTACT TT-------- ---------- -------CCG AA-------- ---------- ----------
snaSere CAAGT----- ---------- ---------- -------CAA AG-------- ---------- ----------
snaSana CGGGACAA-- ---------- ---------- ---------- ---------- ---------- ----------
snaSpse ---------- ---------- ---------- ---------- ---------- ---------- ----------
snaSper ---------- ---------- ---------- ---------- ---------- ---------- ----------
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snaSmoj ---------- ---------- ---------- ---------- ---------- ---------- ----------
snaSgri CAGAAT---- ---------- ---------- ---------- ---------- ---------- ----------
snaSwil ---------- ---------- ---------- ---------- ---------- ---------- ----------
561
snaSmel ---------- ---------- ---------- ---------- ---------- ------CTAG TTG---ACAT
snaSsim ---------- ---------- ---------- ---------- ---------- ------ACAC CTG---ACAC
snaSsec ---------- ---------- ---------- ---------- ---------- ------ACAC CTG---ACAC
snaSyak ---------- ---------- ---------- ---------- ---------- ------CTAG CTG---TAAT
snaSere ---------- ---------- ---------- ---------- ---------- ------GCAC CTG---ACAC
snaSana ---------- ---------- ---------- ---------- ---------- ------TCAC CTGCCCACGC
snaSpse ---------- ---------- ---------- ---------- ---------- ---------- CTG-------
snaSper ---------- ---------- ---------- ---------- ---------- ---------- CTG-------
snaSvir TTAAGATACT CTGGCACTAT GCAAAGGTCT AGCCAATTCA ACAGGTAGCT ACCCGCACAC TTG---ACAC
snaSmoj ---------- ---------- ---------- ---------- ---------- ------ACAC TTG---CCAC
snaSgri ---------- ---------- ---------- ---------- ---------- ------ACAC TTG---ACAC
snaSwil ---------- ---------- ---------- ---------- ---------- ---------- TTG-------
631
snaSmel TTTTTGCGGT GTAACTGTGA T--AGCTGTG T---ACCTTC AAAGTCCCAT GTGCCAGTCT CTCTCA----
snaSsim GT-----GTC ATAGGT---- -----CAGCG TGAGATATTG GCAA----AT GTTTA--CCT GT--------
snaSsec GT-----GTC ATAGGT---- -----CAGCG TGAGATATTG GCAA----AT GTTTA--CCT GT--------
snaSyak TTTTTGTGGT GTAACTGCGA T--AGCTGTA GGATTCCTTC GAAGTCCCAT GTGCCAGTCT CTCTCA----
snaSere GT-----GTC ATAGGT---- -----CAGCG TGAGCTATTG GCAA----AT GTTTA--CCT GT--------
snaSana GTCGGTGGCC AATAGC---- -----CAGCG TGAGATATTG GCAA----AT GTTTA--CCT GT--------
snaSpse TTGTTCTAGG AAAATCATTC TCGAGCTG-G GAATATGTTC GTGA----AT CTTCAGGTCT TTTCCAAGGG
snaSper TTGTTCTAGG AAAATCATCC TCGAGCTG-G GAATATGTTC GTGA----AT CTTCAGGTCT TTTCCAAGGG
snaSvir GT-----GTC AGAAGC---A ACAGGCTGTA TGTTTTATTG GCAA----AT GTTTA--CCT GT------TG
snaSmoj GC-----GTC AGGGGC---A ACAGGCTGTA TTTAATATTG GCAA----AT GTTTA--CCT GTTGCTGCTG
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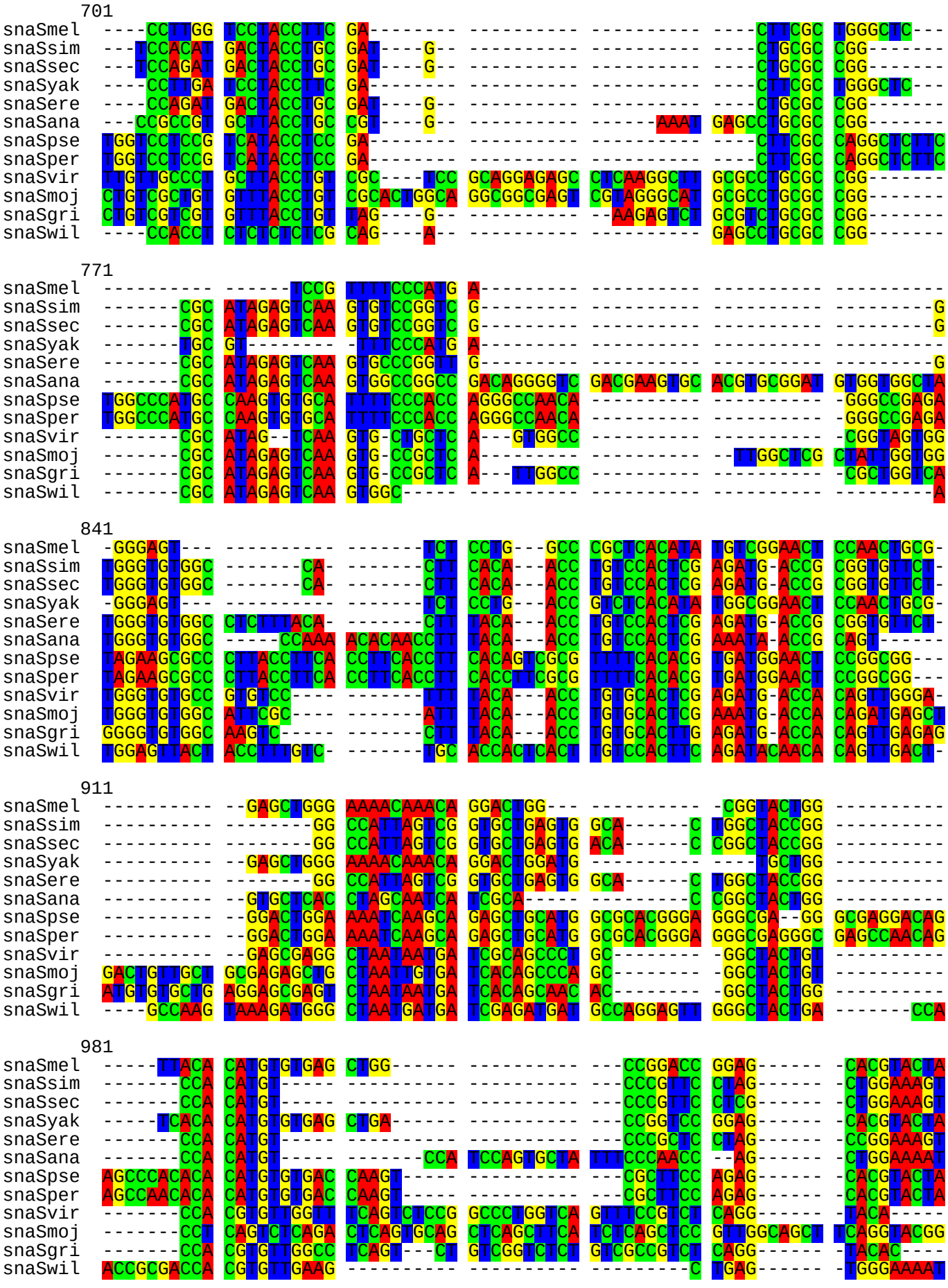
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snaSsec ---TCCAGAT GACTACCTGC GAT----G-- ---------- ---------- ----CTGCGC CGG-------
snaSyak ----CCTTGA TCCTACCTTC GA--------- ---------- ---------- ----CTTCGC TGGGCTC---
snaSere ----CCAGAT GACTACCTGC GAT----G-- ---------- ---------- ----CTGCGC CGG-------
snaSana ---CCGCCGT GCTTACCTGC CGT----G-- ---------- ------AAAT GAGCCTGCGC CGG-------
snaSpse TGGTCCTCCG TCATACCTCC GA--------- ---------- ---------- ----CTTCGC CAGGCTCTTC
snaSper TGGTCCTCCG TCATACCTCC GA--------- ---------- ---------- ----CTTCGC CAGGCTCTTC
snaSvir TTGTTGCCCT GCTTACCTGT CGC----TCC GCAGGAGAGC CTCAAGGCTT GCGCCTGCGC CGG-------
snaSmoj CTGTCGCTGT GTTTACCTGT CGCACTGGCA GGCGGCGAGT CGTAGGGCAT GCGCCTGCGC CGG-------
snaSgri CTGTCGTCGT GTTTACCTGT TAG----G-- ---------- --AAGAGTCT GCGTCTGCGC CGG-------
snaSwil ----CCACCT CTCTCTCTCG CAG----A-- ---------- ---------- GAGCCTGCGC CGG-------
771
snaSmel ---------- ------TCCG TTTTCCCATG A--------- ---------- ---------- ----------
snaSsim -------CGC ATAGAGTCAA GTGTCCGGTC G--------- ---------- ---------- ---------G
snaSsec -------CGC ATAGAGTCAA GTGTCCGGTC G--------- ---------- ---------- ---------G
snaSyak -------TGC GT--------- -TTTCCCATG A--------- ---------- ---------- ----------
snaSere -------CGC ATAGAGTCAA GTGCCCGGTT G--------- ---------- ---------- ---------G
snaSana -------CGC ATAGAGTCAA GTGGCCGGCC GACAGGGGTC GACGAAGTGC ACGTGCGGAT GTGGTGGCTA
snaSpse TGGCCCATGC CAAGTGTGCA TTTTCCCACC AGGGCCAACA ---------- ---------- -GGGCCGAGA
snaSper TGGCCCATGC CAAGTGTGCA TTTTCCCACC AGGGCCAACA ---------- ---------- -GGGCCGAGA
snaSvir -------CGC ATAG--TCAA GTG-CTGCTC A---GTGGCC ---------- ---------- -CGGTAGTGG
snaSmoj -------CGC ATAGAGTCAA GTG-CCGCTC A--------- ---------- --TTGGCTCG CTATTGGTGG
snaSgri -------CGC ATAGAGTCAA GTG-CCGCTC A---TTGGCC ---------- ---------- -CGCTGGTCA
snaSwil -------CGC ATAGAGTCAA GTGGC----- ---------- ---------- ---------- ---------A
841
snaSmel -GGGAGT--- ---------- ------TCT CCTG---GCC CGCTCACATA TGTCGGAACT CCAACTGCG-
snaSsim TGGGTGTGGC -------CA- ------CTT CACA---ACC TGTCCACTCG AGATG-ACCG CGGTGTTCT-
snaSsec TGGGTGTGGC -------CA- ------CTT CACA---ACC TGTCCACTCG AGATG-ACCG CGGTGTTCT-
snaSyak -GGGAGT--- ---------- ------TCT CCTG---ACC GTCTCACATA TGGCGGAACT CCAACTGCG-
snaSere TGGGTGTGGC CTCTTTACA- ------CTT TACA---ACC TGTCCACTCG AGATG-ACCG CGGTGTTCT-
snaSana TGGGTGTGGC -----CCAAA ACACAACCTT TACA---ACC TGTCCACTCG AAATA-ACCG CAGT------
snaSpse TAGAAGCGCC CTTACCTTCA CCTTCACCTT CACAGTCGCG TTTTCACACG TGATGGAACT CCGGCGG---
snaSper TAGAAGCGCC CTTACCTTCA CCTTCACCTT CACCTTCGCG TTTTCACACG TGATGGAACT CCGGCGG---
snaSvir TGGGTGTGCC GTGTCC----- -------TTT TACA---ACC TGTGCACTCG AGATG-ACCA CAGTTGGGA-
snaSmoj TGGGTGTGGC ATTCGC----- -------ATT TACA---ACC TGTGCACTCG AAATG-ACCA CAGATGAGCT
snaSgri GGGGTGTGGC AAGTC----- -------CTT TACA---ACC TGTGCACTTG AGATG-ACCA CAGTTGAGAG
snaSwil TGGAGTTACT ACCTTTGTC- ------TGC ACCACTCACT TGTCCACTTC AGATACAACA CAGTTGACT-
911
snaSmel ---------- --GAGCTGGG AAAACAAACA GGACTGG--- ---------- -CGGTACTGG ----------
snaSsim ---------- --------GG CCATTAGTCG GTGCTGAGTG GCA------C TGGCTACCGG ----------
snaSsec ---------- --------GG CCATTAGTCG GTGCTGAGTG ACA------C CGGCTACCGG ----------
snaSyak ---------- --GAGCTGGG AAAACAAACA GGACTGGATG ---------- ----TGCTGG ----------
snaSere ---------- ---------- CCATTAGTCG GTGCTGAGTG GCA------C TGGCTACCGG ----------
snaSana ---------- --GTGCTCAC CTAGCAATCA TCGCA----- ---------- CGGCTACTGG ----------
snaSpse ---------- --GGACTGGA AAATCAAGCA GAGCTGCATG GCGCACGGGA GGGCGA--GG GCGAGGACAG
snaSper ---------- --GGACTGGA AAATCAAGCA GAGCTGCATG GCGCACGGGA GGGCGAGGGC GAGCCAACAG
snaSvir ---------- --GAGCGAGG CTAATAATGA TCGCAGCCCT GC-------- -GGCTACTGT ----------
snaSmoj GACTGTTGCT GCGAGAGCTG CTAATTGTGA TCACAGCCCA GC-------- -GGCTACTGT ----------
snaSgri ATGTGTGCTG AGGAGCGAGT CTAATAATGA TCACAGCAAC AC-------- -GGCTACTGG ----------
snaSwil ----GCCAAG TAAAGATGGG CTAATGATGA TCGAGATGAT GCCAGGAGTT GGGCTACTGA -------CCA
981
snaSmel -----TTACA CATGTGTGAG CTGG------ ---------- ---CCGGACC GGAG------ -CACGTACTA
snaSsim -------CCA CATGT----- ---------- ---------- ---CCCGTTC CTAG------ -CTGGAAAGT
snaSsec -------CCA CATGT----- ---------- ---------- ---CCCGTTC CTCG------ -CTGGAAAGT
snaSyak -----TCACA CATGTGTGAG CTGA------ ---------- ---CCGGTCC GGAG------ -CACGTACTA
snaSere -------CCA CATGT----- ---------- ---------- ---CCCGCTC CTAG------ -CCGGAAAGT
snaSana -------CCA CATGT----- -------CCA TCCAGTGCTA TTTCCCAACC --AG------ -CTGGAAAAT
snaSpse AGCCCACACA CATGTGTGAC CAAGT----- ---------- ---CGCTTCC AGAG------ -CACGTACTA
snaSper AGCCAACACA CATGTGTGAC CAAGT----- ---------- ---CGCTTCC AGAG------ -CACGTACTA
snaSvir -------CCA CGTGTTGGTT TCAGTCTCCG GCCCTGGTCA GTTTCCGTCT CAGG------ -TACA-----
snaSmoj -------CCT CAGTCTCAGA CTCAGTGCAG CTCAGCTTCA TCTCAGCTCC GTTGGCAGCT TCAGGTACGG
snaSgri -------CCA CGTGTTGGCC TCAGT---CT GTCGGTCTCT GTCGCCGTCT CAGG------ -TACAC-----
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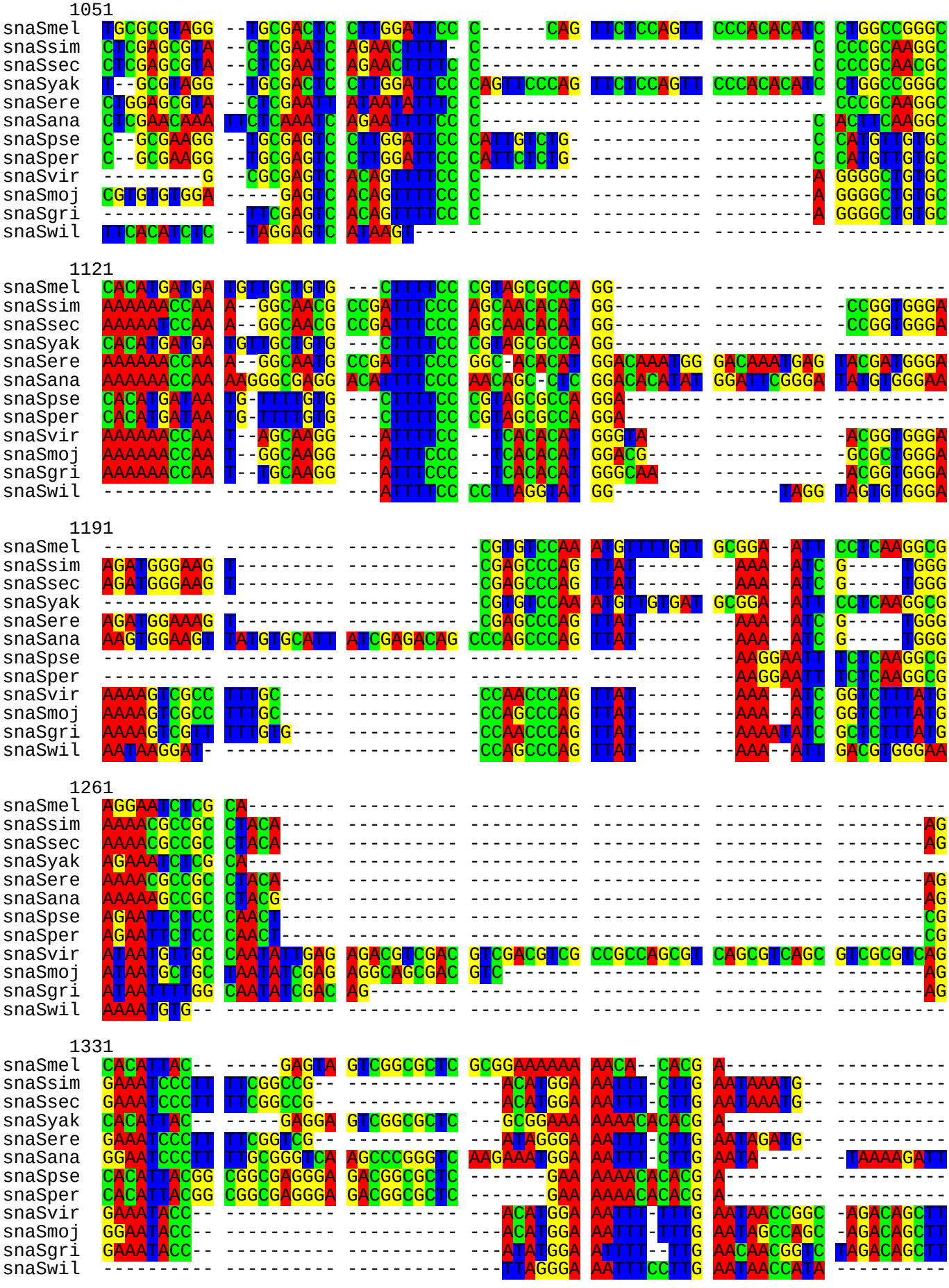
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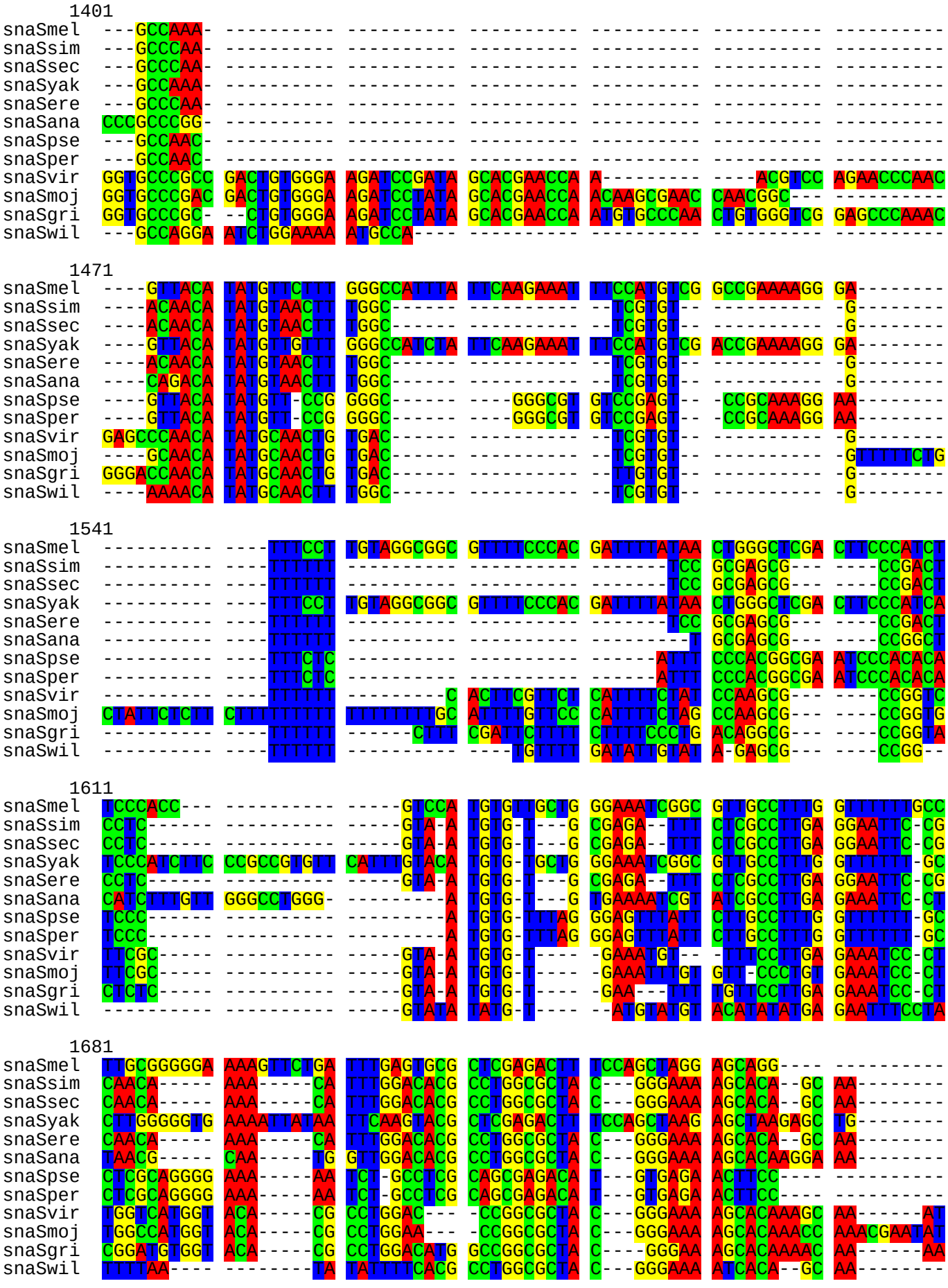
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snaSsec ---GCCCAA- ---------- ---------- ---------- ---------- ---------- ----------
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snaSere ---GCCCAA- ---------- ---------- ---------- ---------- ---------- ----------
snaSana CCCGCCCGG- ---------- ---------- ---------- ---------- ---------- ----------
snaSpse ---GCCAAC- ---------- ---------- ---------- ---------- ---------- ----------
snaSper ---GCCAAC- ---------- ---------- ---------- ---------- ---------- ----------
snaSvir GGTGCCCGCC GACTGTGGGA AGATCCGATA GCACGAACCA A---------- ----ACGTCC AGAACCCAAC
snaSmoj GGTGCCCGAC GACTGTGGGA AGATCCTATA GCACGAACCA ACAAGCGAAC CAACGGC--- ----------
snaSgri GGTGCCCGC- --CTGTGGGA AGATCCTATA GCACGAACCA ATGTGCCCAA CTGTGGGTCG GAGCCCAAAC
snaSwil ---GCCAGGA ATCTGGAAAA ATGCCA---- ---------- ---------- ---------- ----------
1471
snaSmel ----GTTACA TATGTTCTTT GGGCCATTTA TTCAAGAAAT TTCCATGTCG GCCGAAAAGG GA--------
snaSsim ----ACAACA TATGTAACTT TGGC------ ---------- --TCGTGT-- ---------- -G--------
snaSsec ----ACAACA TATGTAACTT TGGC------ ---------- --TCGTGT-- ---------- -G--------
snaSyak ----GTTACA TATGTTGTTT GGGCCATCTA TTCAAGAAAT TTCCATGTCG ACCGAAAAGG GA--------
snaSere ----ACAACA TATGTAACTT TGGC------ ---------- --TCGTGT-- ---------- -G--------
snaSana ----CAGACA TATGTAACTT TGGC------ ---------- --TCGTGT-- ---------- -G--------
snaSpse ----GTTACA TATGTT-CCG GGGC------ ----GGGCGT GTCCGAGT-- -CCGCAAAGG AA--------
snaSper ----GTTACA TATGTT-CCG GGGC------ ----GGGCGT GTCCGAGT-- -CCGCAAAGG AA--------
snaSvir GAGCCCAACA TATGCAACTG TGAC------ ---------- --TCGTGT-- ---------- -G--------
snaSmoj ----GCAACA TATGCAACTG TGAC------ ---------- --TCGTGT-- ---------- -GTTTTTCTG
snaSgri GGGACCAACA TATGCAACTG TGAC------ ---------- --TTGTGT-- ---------- -G--------
snaSwil ----AAAACA TATGCAACTT TGGC------ ---------- --TCGTGT-- ---------- -G--------
1541
snaSmel ---------- ----TTTCCT TGTAGGCGGC GTTTTCCCAC GATTTTATAA CTGGGCTCGA CTTCCCATCT
snaSsim ---------- ----TTTTTT ---------- ---------- -------TCC GCGAGCG--- ----CCGACT
snaSsec ---------- ----TTTTTT ---------- ---------- -------TCC GCGAGCG--- ----CCGACT
snaSyak ---------- ----TTTCCT TGTAGGCGGC GTTTTCCCAC GATTTTATAA CTGGGCTCGA CTTCCCATCA
snaSere ---------- ----TTTTTT ---------- ---------- -------TCC GCGAGCG--- ----CCGACT
snaSana ---------- ----TTTTTT ---------- ---------- ---------T GCGAGCG--- ----CCGGCT
snaSpse ---------- ----TTTCTC ---------- ---------- ------ATTT CCCACGGCGA ATCCCACACA
snaSper ---------- ----TTTCTC ---------- ---------- ------ATTT CCCACGGCGA ATCCCACACA
snaSvir ---------- ----TTTTTT ---------C ACTTCGTTCT CATTTTCTAT CCAAGCG--- ----CCGGTC
snaSmoj CTATTCTCTT CTTTTTTTTT TTTTTTTGC ATTTTGTTCC CATTTTCTAG CCAAGCG--- ----CCGGTG
snaSgri ---------- ----TTTTTT ------CTTT CGATTCTTTT CTTTTCCCTG ACAGGCG--- ----CCGGTA
snaSwil ---------- ----TTTTTT ---------- ----TGTTTT GATATTGTAT A-GAGCG--- ----CCGG--
1611
snaSmel TCCCACC--- ---------- -----GTCCA TGTGTTGCTG GGAAATCGGC GTTGCCTTTG GTTTTTTGCC
snaSsim CCTC------ ---------- -----GTA-A TGTG-T---G CGAGA--TTT CTCGCCTTGA GGAATTC-CG
snaSsec CCTC------ ---------- -----GTA-A TGTG-T---G CGAGA--TTT CTCGCCTTGA GGAATTC-CG
snaSyak TCCCATCTTC CCGCCGTGTT CATTTGTACA TGTG-TGCTG GGAAATCGGC GTTGCCTTTG GTTTTTT-GC
snaSere CCTC------ ---------- -----GTA-A TGTG-T---G CGAGA--TTT CTCGCCTTGA GGAATTC-CG
snaSana CATCTTTGTT GGGCCCTGGG- ---------A TGTG-T---G TGAAAATCGT ATCGCCTTGA GAAATTC-CT
snaSpse TCCC------ ---------- ---------A TGTG-TTTAG GGAGTTTATT CTTGCCTTTG GTTTTTT-GC
snaSper TCCC------ ---------- ---------A TGTG-TTTAG GGAGTTTATT CTTGCCTTTG GTTTTTT-GC
snaSvir TTCGC----- ---------- -----GTA-A TGTG-T---- -GAAATGT-- -TTTCCTTGA GAAATCC-CT
snaSmoj TTCGC----- ---------- -----GTA-A TGTG-T---- -GAAATTTGT GTT-CCCTGT GAAATCC-CT
snaSgri CTCTC----- ---------- -----GTA-A TGTG-T---- -GAA---TTT TGTTCCTTGA GAAATCC-CT
snaSwil ---------- ---------- -----GTATA TATG-T---- --ATGTATGT ACATATATGA GAATTTCCTA
1681
snaSmel TTGCGGGGGA AAAGTTCTGA TTTGAGTGCG CTCGAGACTT TCCAGCTAGG AGCAGG---- ----------
snaSsim CAACA----- AAA-----CA TTTGGACACG CCTGGCGCTA C---GGGAAA AGCACA--GC AA--------
snaSsec CAACA----- AAA-----CA TTTGGACACG CCTGGCGCTA C---GGGAAA AGCACA--GC AA--------
snaSyak CTTGGGGGTG AAAATTATAA TTCAAGTACG CTCGAGACTT TCCAGCTAAG AGCTAAGAGC TG--------
snaSere CAACA----- AAA-----CA TTTGGACACG CCTGGCGCTA C---GGGAAA AGCACA--GC AA--------
snaSana TAACG----- CAA-----TG GTTGGACACG CCTGGCGCTA C---GGGAAA AGCACAAGGA AA--------
snaSpse CTCGCAGGGG AAA-----AA TCT-GCCTCG CAGCGAGACA T---GTGAGA ACTTCC---- ----------
snaSper CTCGCAGGGG AAA-----AA TCT-GCCTCG CAGCGAGACA T---GTGAGA ACTTCC---- ----------
snaSvir TGGTCATGGT ACA-----CG CCTGGAC--- -CCGGCGCTA C---GGGAAA AGCACAAAGC AA------AT
snaSmoj TGGCCATGGT ACA-----CG CCTGGAA--- -CCGGCGCTA C---GGGAAA AGCACAAACC AAACGAATAT
snaSgri CGGATGTGGT ACA-----CG CCTGGACATG GCCGGCGCTA C----GGGAA AGCACAAAAC AA------AA
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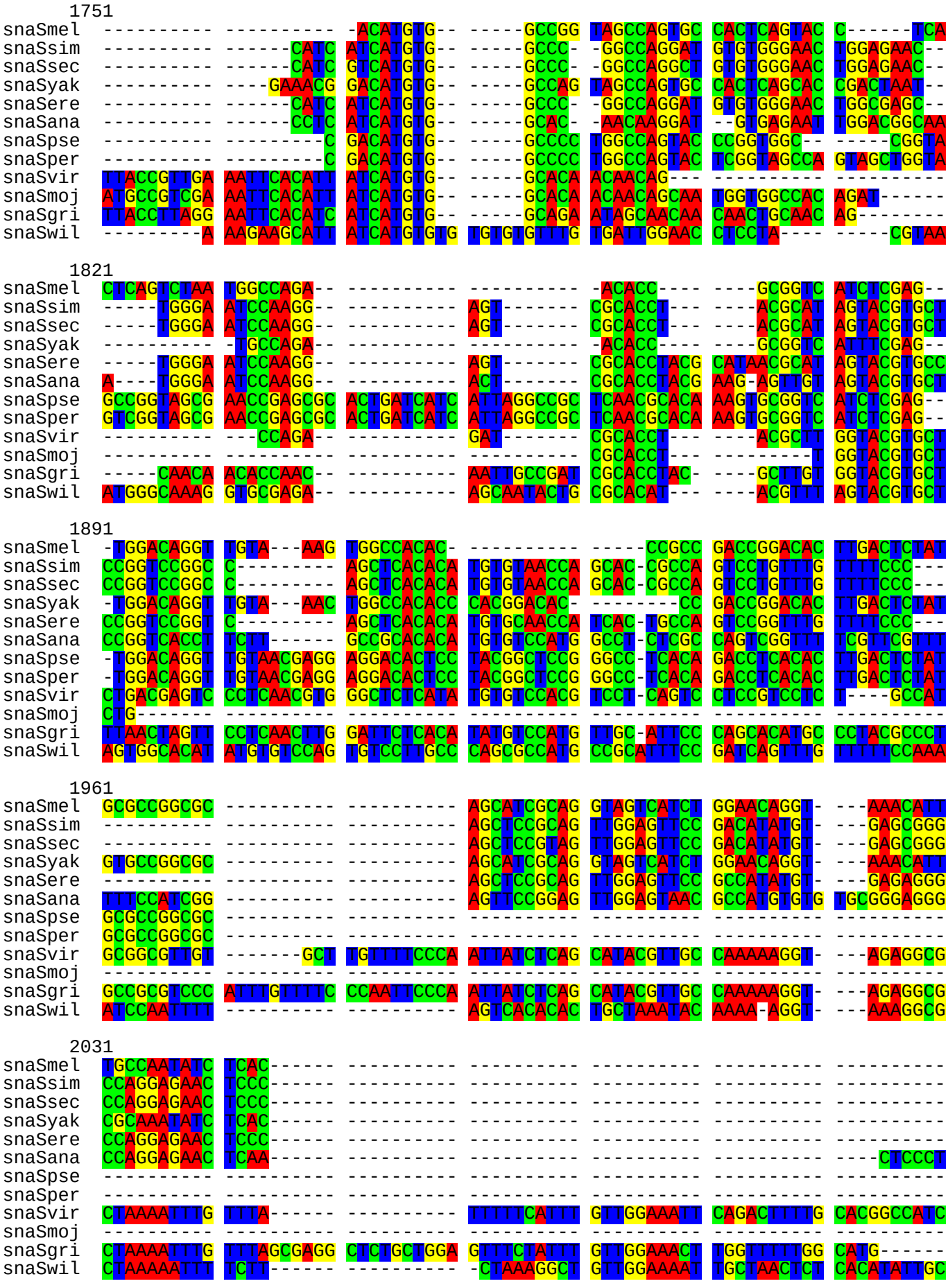
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snaSsec ---------- ------CATC GTCATGTG-- -----GCCC- -GGCCAGGCT GTGTGGGAAC TGGAGAAC--
snaSyak ---------- ----GAAACG GACATGTG-- -----GCCAG TAGCCAGTGC CACTCAGCAC CGACTAAT--
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snaSpse ---------- ---------C GACATGTG-- -----GCCCC TGGCCAGTAC CCGGTGGC-- -----CGGTA
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snaSwil ---------A AAGAAGCATT ATCATGTGTG TGTGTGTTTG TGATTGGAAC CTCCTA---- -----CGTAA
1821
snaSmel CTCAGTCTAA TGGCCAGA-- ---------- ---------- -ACACC---- ----GCGGTC ATCTCGAG--
snaSsim -----TGGGA ATCCAAGG-- ---------- AGT------- CGCACCT--- ----ACGCAT AGTACGTGCT
snaSsec -----TGGGA ATCCAAGG-- ---------- AGT------- CGCACCT--- ----ACGCAT AGTACGTGCT
snaSyak ---------- -TGCCAGA-- ---------- ---------- -ACACC---- ----GCGGTC ATTTCGAG--
snaSere -----TGGGA ATCCAAGG-- ---------- AGT------- CGCACCTACG CATAACGCAT AGTACGTGCC
snaSana A----TGGGA ATCCAAGG-- ---------- ACT------- CGCACCTACG AAG-AGTTGT AGTACGTGCT
snaSpse GCCGGTAGCG AACCGAGCGC ACTGATCATC ATTAGGCCGC TCAACGCACA AAGTGCGGTC ATCTCGAG--
snaSper GTCGGTAGCG AACCGAGCGC ACTGATCATC ATTAGGCCGC TCAACGCACA AAGTGCGGTC ATCTCGAG--
snaSvir ---------- ---CCAGA-- ---------- GAT------- CGCACCT--- ----ACGCTT GGTACGTGCT
snaSmoj ---------- ---------- ---------- ---------- CGCACCT--- ---------T GGTACGTGCT
snaSgri -----CAACA ACACCAAC-- ---------- AATTGCCGAT CGCACCTAC- ----GCTTGT GGTACGTGCT
snaSwil ATGGGCAAAG GTGCGAGA-- ---------- AGCAATACTG CGCACAT--- ----ACGTTT AGTACGTGCT
1891
snaSmel -TGGACAGGT TGTA---AAG TGGCCACAC- ---------- -----CCGCC GACCGGACAC TTGACTCTAT
snaSsim CCGGTCCGGC C---------- AGCTCACACA TGTGTAACCA GCAC-CGCCA GTCCTGTTTG TTTTCCC---
snaSsec CCGGTCCGGC C---------- AGCTCACACA TGTGTAACCA GCAC-CGCCA GTCCTGTTTG TTTTCCC---
snaSyak -TGGACAGGT TGTA---AAC TGGCCACACC CACGGACAC- --------CC GACCGGACAC TTGACTCTAT
snaSere CCGGTCCGGT C---------- AGCTCACACA TGTGCAACCA TCAC-TGCCA GTCCGGTTTG TTTTCCC---
snaSana CCGGTCACCT TCTT------ GCCGCACACA TGTGTCCATG GCCT-CTCGC CAGTCGGTTT TCGTTCGTTT
snaSpse -TGGACAGGT TGTAACGAGG AGGACACTCC TACGGCTCCG GGCC-TCACA GACCTCACAC TTGACTCTAT
snaSper -TGGACAGGT TGTAACGAGG AGGACACTCC TACGGCTCCG GGCC-TCACA GACCTCACAC TTGACTCTAT
snaSvir CTGACGAGTC CCTCAACGTG GGCTCTCATA TGTGTCCACG TCCT-CAGTC CTCCGTCCTC T----GCCAT
snaSmoj CTG------- ---------- ---------- ---------- ---------- ---------- ----------
snaSgri TTAACTAGTT CCTCAACTTG GATTCTCACA TATGTCCATG TTGC-ATTCC CAGCACATGC CCTACGCCCT
snaSwil AGTGGCACAT ATGTGTCCAG TGTCCTTGCC CAGCGCCATG CCGCATTTCC GATCAGTTTG TTTTTCCAAA
1961
snaSmel GCGCCGGCGC ---------- ---------- AGCATCGCAG GTAGTCATCT GGAACAGGT- ---AAACATT
snaSsim ---------- ---------- ---------- AGCTCCGCAG TTGGAGTTCC GACATATGT- ---GAGCGGG
snaSsec ---------- ---------- ---------- AGCTCCGTAG TTGGAGTTCC GACATATGT- ---GAGCGGG
snaSyak GTGCCGGCGC ---------- ---------- AGCATCGCAG GTAGTCATCT GGAACAGGT- ---AAACATT
snaSere ---------- ---------- ---------- AGCTCCGCAG TTGGAGTTCC GCCATATGT- ---GAGAGGG
snaSana TTTCCATCGG ---------- ---------- AGTTCCGGAG TTGGAGTAAC GCCATGTGTG TGCGGGAGGG
snaSpse GCGCCGGCGC ---------- ---------- ---------- ---------- ---------- ----------
snaSper GCGCCGGCGC ---------- ---------- ---------- ---------- ---------- ----------
snaSvir GCGGCGTTGT -------GCT TGTTTTCCCA ATTATCTCAG CATACGTTGC CAAAAAGGT- ---AGAGGCG
snaSmoj ---------- ---------- ---------- ---------- ---------- ---------- ----------
snaSgri GCCGCGTCCC ATTTGTTTTC CCAATTCCCA ATTATCTCAG CATACGTTGC CAAAAAGGT- ---AGAGGCG
snaSwil ATCCAATTTT ---------- ---------- AGTCACACAC TGCTAAATAC AAAA-AGGT- ---AAAGGCG
2031
snaSmel TGCCAATATC TCAC------ ---------- ---------- ---------- ---------- ----------
snaSsim CCAGGAGAAC TCCC------ ---------- ---------- ---------- ---------- ----------
snaSsec CCAGGAGAAC TCCC------ ---------- ---------- ---------- ---------- ----------
snaSyak CGCAAATATC TCAC------ ---------- ---------- ---------- ---------- ----------
snaSere CCAGGAGAAC TCCC------ ---------- ---------- ---------- ---------- ----------
snaSana CCAGGAGAAC TCAA------ ---------- ---------- ---------- ---------- ----CTCCCT
snaSpse ---------- ---------- ---------- ---------- ---------- ---------- ----------
snaSper ---------- ---------- ---------- ---------- ---------- ---------- ----------
snaSvir CTAAAATTTG TTTA------ ---------- TTTTTCATTT GTTGGAAATT CAGACTTTTG CACGGCCATC
snaSmoj ---------- ---------- ---------- ---------- ---------- ---------- ----------
snaSgri CTAAAATTTG TTTAGCGAGG CTCTGCTGGA GTTTCTATTT GTTGGAAACT TGGTTTTTGG CATG------
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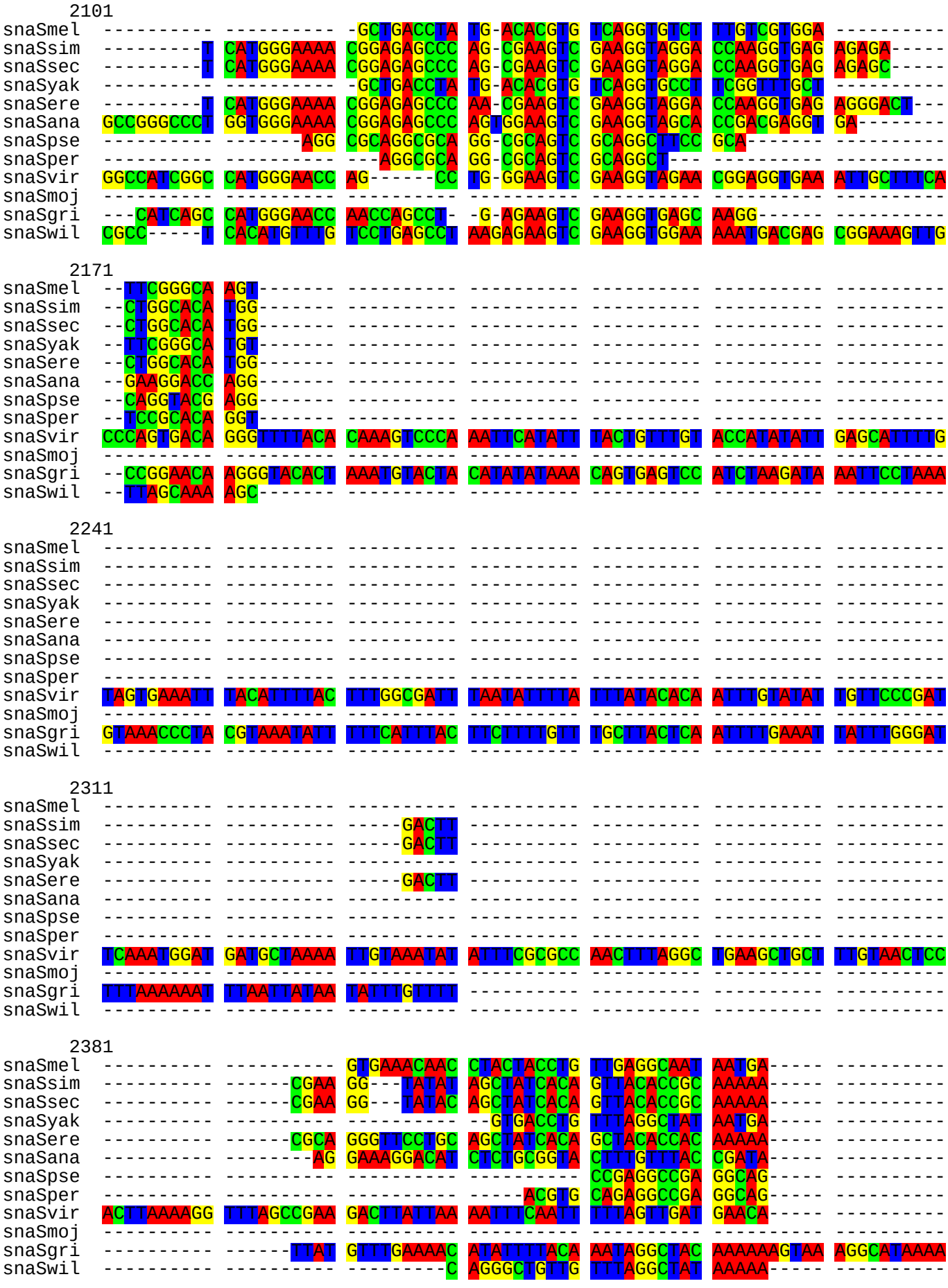
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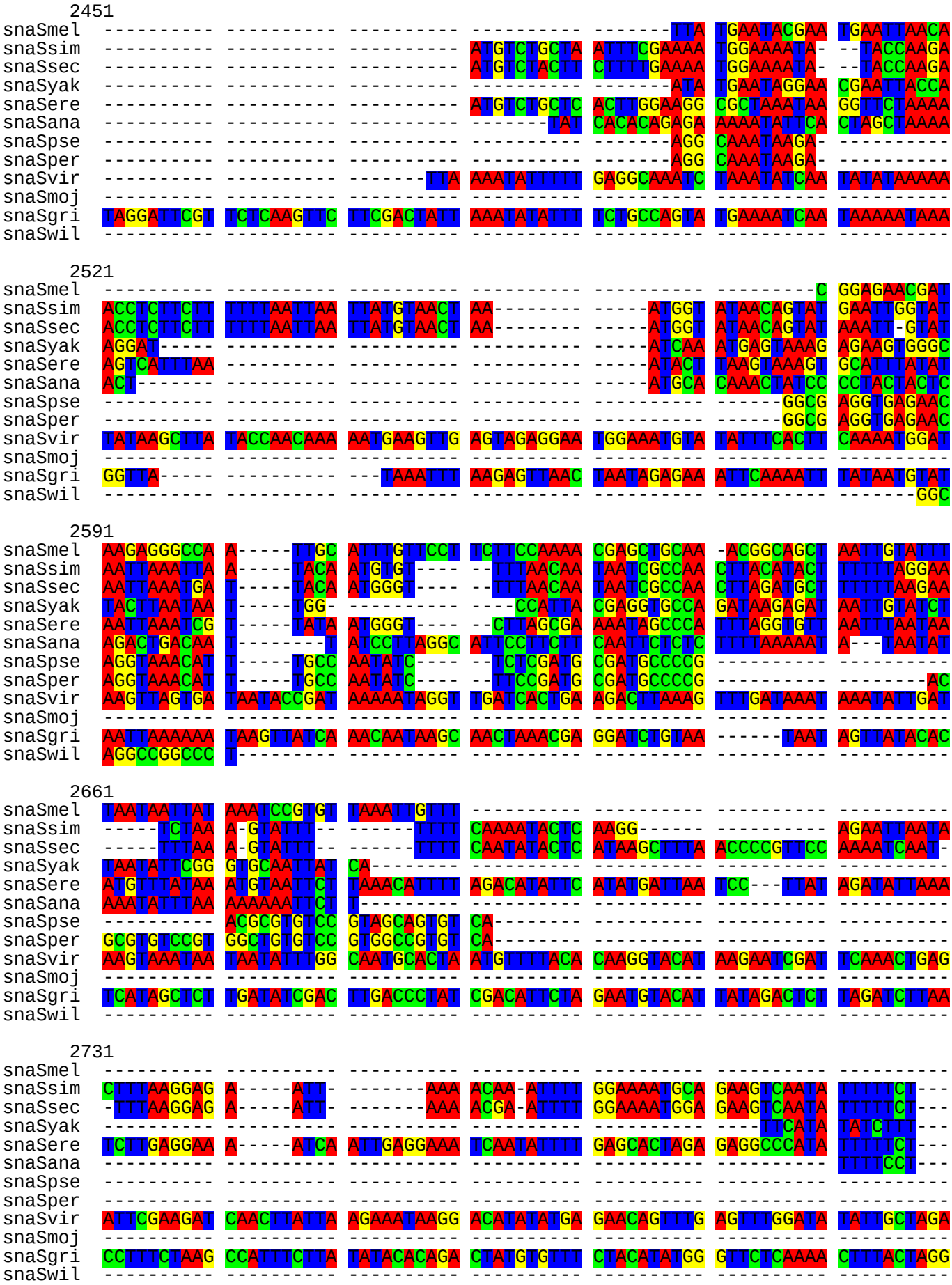
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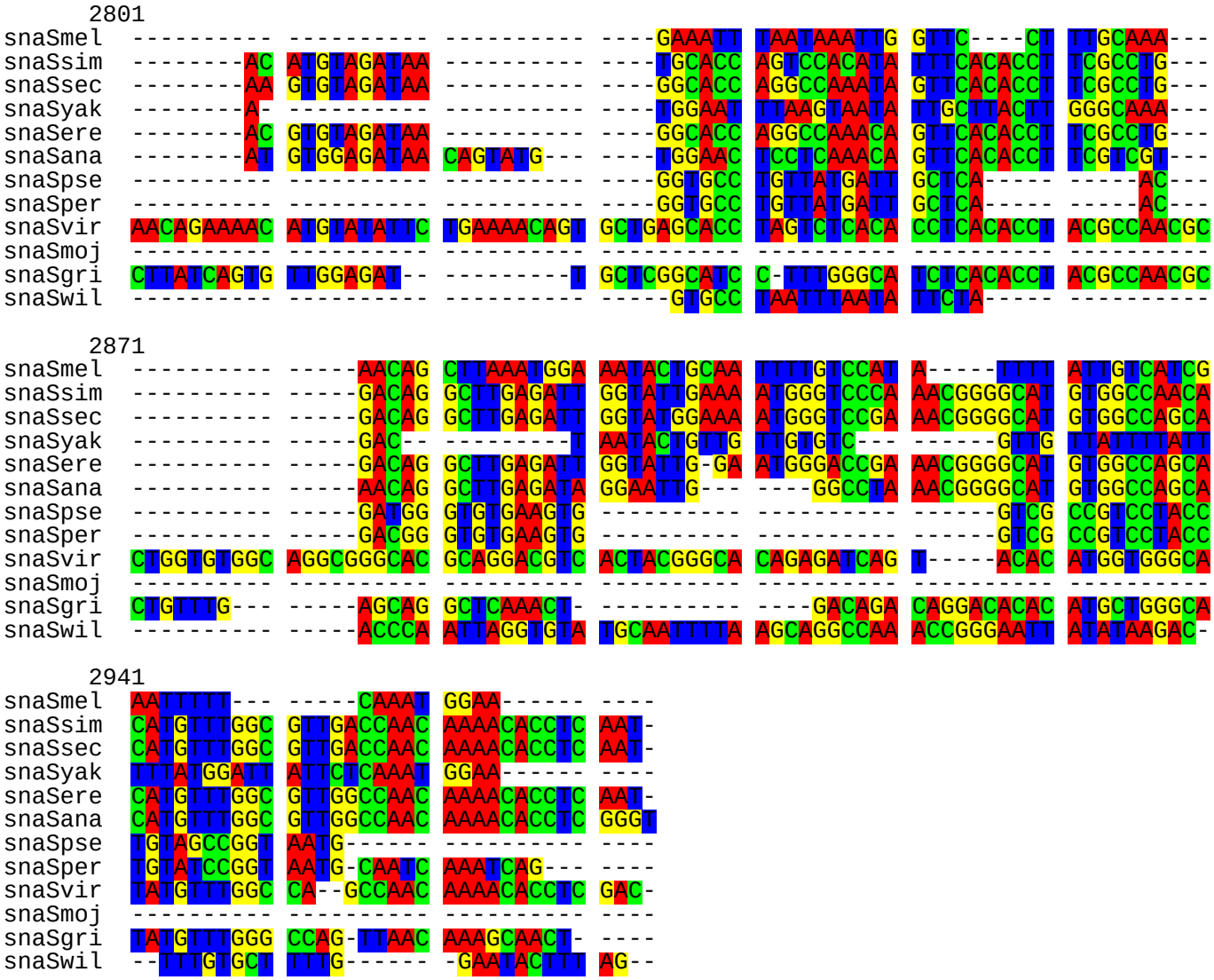
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