```
                    5        10        15        20        25        30        35        40        45        50        55        60        65        70        75        80
conservation        ***** ******** ********************************** ************************* * **********
SCPP1˙CI            ATGAAACCTGTCATCCTGATTCTCTGTCTGCTGGGAGCAGCCTATGCTAATCCAATCTTGCATAAAATGGCTATGGAGAT    80
SCPP1˙AM            ................................................................................    0
SCPP1˙DR            ATGAAGTCTGCCCTATTGATTCTTTGTCTGTTGGGAGCAGCCTGTGCTAATCCAATCTTGCATAAAGTGGCTATGGAGAT    80
SCPP1˙IP            ATGAAGCTGGCTTTCGTGATTCTCTGCCTGCTCGGAGCTGCCGGTGCAAATCCAATTTTGCAC.........ACGGATAT    71
SCPP1˙LW            ATGAAACCTGCCATACTGATTCTATGTCTGCTGGGAACAGCCTATGCTAATCCAATCTTGCATAAAATGGCTATGGAGAT    80
SCPP1˙PN            ATGAAGTTTGCCATTGTAATTCTCTGCCTGCTGGGAACAACCAGTGCTAATCCCATCTTATACAAAGTGACCACTGAGAT    80
SCPP1˙PP            ATGAAACCTGCCATACTGATTCTATGTCTGCTGGGAGCAGCCTACGCTAATCCAATCTTGCATAAAATGGCTATGGAGAT    80
SCPP1-C1˙CC         ATGAAACCTGCCATGTTGATTCTCTGTCTGCTGGGAGCAGCCTATGCTAATCCAATCTTGCATAAAATGGCTATGGAGAT    80
SCPP1-C2˙CC         ATGAAACATGCCATATTGATTCTCTGTCTGCTGGGAGCAGCCTATGCTAATCCAATCTTGCATAAAACGGCTATGGAGAT    80
SCPP1-C1˙SR         ................................................................................    0
SCPP1-C2˙SR         ATGAAACCTGCCATATTGATTCTCTGTCTGCTGGGAGCAGCCTATGCTAATCCAATCTTGCATAAAAAGACTATGGAGAT    80
SCPP1-C1˙SA         ATGAACCCTGCCATATTGATTCTCTGTCTGCTGGGAGCAGCCTATGCTAATCCAATCTTGCATAAAATGGCTATGGAGGT    80
SCPP1-C2˙SA         ATGAAACCTGCCATACTGATTCTCTGTCTGCTGGGAGCAGCCTATGCTAATCCAATCTTGCATAAAAAGACTATGGAGAT    80
SCPP1-C1˙SG         ATGAAACCTGCCATATTGATTCTCTGTCTGCTGGGAGCAGCCTATGCTAATCCAATCTTGCATAAAATGGCTATGGAGGT    80
SCPP1-C2˙SG         ATGAAACCTGCCATATTGATTCTCTGTCTGCTGGGAGCAGCCTGTGCTAATCCAATCTTGCATAAAAAGACTATGGAGAT    80

                    85        90        95       100       105       110       115       120       125       130       135       140       145       150       155       160
conservation        ***********************************************  ***********  ********************  ********************  *****
SCPP1˙CI            GATTCAACATGCATCCAACTCTTCAGAAAGCTCTTCAATATCTGAATCCTCCGACCAGAGTAACACCTCAGAACCTTCAG   160
SCPP1˙AM            ................................................................................    0
SCPP1˙DR            GATTCAACATGCATCTAACTCTTCGGAAAGCTCCTCAATATCTGAATCCTCAGACCAGAGTAATACCTCAGAACCTTCGG   160
SCPP1˙IP            GATGGAGACCGCATCGAACTCTTCACAGACCTCCTCGATGTCTGCATCTACTGAAGAA......ACCGTTGCCATAGACC   145
SCPP1˙LW            GATTCAACATGCATCCAACTCTTCGGAAAGCTCTTCAATATCTGAATCCTCCGACCAGAGTAATACCTCAGAACCTTCAG   160
SCPP1˙PN            GATGGAACATGCATCCAACTCT......ACCTCATCAGTGTCTGAATCCTCTGAAGAAAGCGATACATCAGAA......C   148
SCPP1˙PP            GATTCAACATGCATCCAACTCTTCGGAAAGCTCTTCAATATCTGAATCCTCCGACCAGAGTAATACCTCAGAACCTTCAG   160
SCPP1-C1˙CC         GATTCAACATGCATCCAACTCTTCGGAGAGCTCTTCAATATCTGAATCCTCTGACCAGAGTAATACCTCAGAACCTTCAG   160
SCPP1-C2˙CC         TATTCAACATGCATCCAACTCTTCGGAAAGCTCTTCGATGTCTGAATCTTCCGACCAGAGTAATACCTCAGAACTTTCAG   160
SCPP1-C1˙SR         ...............................................CAGGATTCATCCCCACCAAGCG   22
SCPP1-C2˙SR         GATTCAACATGCATCCAACTCTTCGGAAAGCTCTTCAATGTCTGAATCCTCCGACCAGAGTAATACCTCAGAACTTTCAG   160
SCPP1-C1˙SA         GATTCAACATGCATCCAACTCTTCGGAAAGCTCTTCAATATCTGAATCCTCCGACCAGAGTAATATCTCAGAACCTTCAG   160
SCPP1-C2˙SA         GATTCAACATGCATCCAACTCTTCGGAAAGCTCTTCAATGTCTGAATCCTCCGACCAGAGTAATACCTCAGAACTTTCAG   160
SCPP1-C1˙SG         GATTCAACATGCATCCAACTCTTCGGAAAGCTCTTCAATATCTGAATCCTCCGACCAGAGTAATACCTCAGAACCTTCAG   160
SCPP1-C2˙SG         GATTCAACATGCATCCAGCTCTTCGGAAAGCTCTTCAATGTCTGAATCCTCCGACCAGAGTAATACCTCAGAACTTTCAG   160

                   165       170       175       180       185       190       195       200       205       210       215       220       225       230       235
conservation        * ***** ** ******* ****** ******* *** ********   ****** ***** ********** **
SCPP1˙CI            AAGAGAAGTCCAAGGAAAATGCCTCCGACAGCAGTAGCTCAGAATCGCTTGAATCTGACTCAGATGAACCGATCTCT...   237
SCPP1˙AM            ................................................................................    0
SCPP1˙DR            AAGAGAAGTCAGAGGAAAATGTTTCCGACAGCAATAGCTCTGAATCACTTGAATCTGAATCAGATGAACCGATATCTAAA   240
SCPP1˙IP            AAGATAGCTCTCAGGAAAATACTTCAGAGGACACCACCTCAGAATCCATGGAATCGAATTCATCTGAGAAGACCTTGGAA   225
SCPP1˙LW            AAGAGAAGTCCAAGGAAAATGTCTCAGACAGCAATAGCTCAGAATCGCTTGAATCTGACTCGGATGAAACGATCTCTAAA   240
SCPP1˙PN            ACAACAGCTCACCGGAAAATACTTCTGAGAACAATACGTCAGAATCGCTGGAGTCTGCATCCGATGATCAGACTTCAGAC   228
SCPP1˙PP            AAGAGGTAAGC.....................................................................   171
SCPP1-C1˙CC         ATGAGAAATCTAAGGAAACTGTCTCCGACAGCAGCAGCTCAGAATCAACTGAATCAGACTCTAATGAACCGATCTCA...   237
SCPP1-C2˙CC         AAGAGAAGTCTACGGAAACTGTCTCTGACAGCAATAGTTCAGAATCAATTGAATCGGCCTCAAATGAACCGATTTCT...   237
SCPP1-C1˙SR         AGACTAAGTCTAAGGAAACTGTCTCCGACAGCAATAGCTCAGAATCAACTGAATCAGACTCTAATGAACCGATCTCT...    99
SCPP1-C2˙SR         AAGAGAACTCTAAGGAAACTGTCTCCGACAGCAATAGTTCAGAATCAATTGAATCGGACTCAAATGAACCGATTTCA...   237
SCPP1-C1˙SA         ATGAGAAGTCTAAGGAAACTGTCTCCGACAGCAATAGCTCAGAATCAACTGAATCAGACTCTAATGAACCGATCTCT...   237
SCPP1-C2˙SA         AAGAGAAGTCTAAGGAAACTGTCTCCGACAGCAATAGTTCAGAATCAATTGAATATGACTCAAATGAACCGATTTCA...   237
SCPP1-C1˙SG         ATGAGAAGTCTAAGGAAACTGTCTCCGACAGCAATAGCTCAGAATCAACTGAATCAGACTCTAATGAACCGATCTCT...   237
SCPP1-C2˙SG         AAGAGAAGTCTAAGGAAACTGTCTCAGAGAGCAGTAGTTCAGAATCAATTGAATCAGACTCAAATGAACCGATTTCA...   237

                   240       245       250       255       260       265       270       275       280       285       290       295       300       305       310
conservation          ** ********** ********** ********** ********** ********** ********** ********* ********** ********** ********** ********
SCPP1˙CI               AGCGAAAGCCATTCTGTGGAAAGACTGATTGGAAAAAGCGACACTTCACTGACATCAGATAACACTCAAAGCAGTAA   314
SCPP1˙AM            .........TCAGATTCCATGTTGGTTCAGTTTGGTACAGGTGAGCCTGGAATGACCACAGACAACAGCCAAGGAAGCCA    71
SCPP1˙DR            GAGAGTGAAAGCCATTCTGTGGAAAGTCTGATTGGAAAAAGCGAGACCGCACTGACAGCCGACAACACTCAAAGCAGTAA   320
SCPP1˙IP            ACAAGCCAAAGCAACTCGCTGGAAGAACGGTTTGGCAATGGAGAGGCTGGAATGACCGTCGATAACAGCCAAGGAAGCAC   305
SCPP1˙LW            GAGAGCGAAAGCCATTCTGTGGAAAATTTGATTGGAAAAAGCGAGGCTGCACTGACATCGGATAACACTCAAAGCAGTAA   320
SCPP1˙PN            ATGAGCCGGAGTCACTCCCTGGAAGCTCGGTTTGGCACAGGTGAGACAGGAATGACCACAGATGACAGCCAAGGCAGCCT   308
SCPP1˙PP            .....................................................................................   171
SCPP1-C1˙CC            AGTGAAAGCCATTCTGTGGAAAGTCTGATTGGAAAAAGTGAGACTGCACTGACATCAGATAACACTCAAAGCAGTAA   314
SCPP1-C2˙CC            AGTGAAAGCCATTCAGTGGAAAGTCTGTCTGGAAAAAGCGAGACTGCACTGACATCAGACAACACTCAAAGCAGTAA   314
SCPP1-C1˙SR            AGTGAAAGCCATTCTGTGGAAAGTCTGATTGGAAAAAGTGAGACTGCACTGACATCAGATAACACTCAAAGCAGTAA   176
SCPP1-C2˙SR            AGTGAAAGCCATTCTGTGGAAAGTCTGACTGGAAAAAGCGAGACTGCACTGACATCAGATAACACTCAAAGCAGTAA   314
SCPP1-C1˙SA            AGTGAAAGCCATTCTGTGGAAAGTCTGATTGGAAAAAGTGAGACTGCACTGACATCAGATAACACTCAAAGCAGTAA   314
SCPP1-C2˙SA            AGTGAAAGCCATTCTGTCGAAAGTCTGACTGGAAAAAGCGAGACTGCACTGACATCAGATAACACTCAAAGCAGTAA   314
SCPP1-C1˙SG            AGTGAAAGCCATTCTGTGGAAAGTCTGATTGGAAAAAGTGAGACTGCACTTACATCAGATAACACTCAAAGCAGTAA   314
SCPP1-C2˙SG            AGTGAAAGCCATTCTGTGGAAAGTCTGACTGGAAAAAGCGAGACTGCACTGACATCAGATAACACTCAAAGCAGTAA   314

                   315       320       325       330       335       340       345       350       355       360       365       370       375       380
conservation        ****** ** ********** ********** ********** ********** ********* ********* * ** **** ****
SCPP1˙CI            AGAGAACGTTCGTAGGGGCTGGATCTACACCCTCAAA...........TGGGTCCATATGAACAATGGTGGCAAAGTGC   382
SCPP1˙AM            GGAGAACATGCGCAAGAACTGGGTTCATCTCATCAAC...........GTCAAAATGGCCAGTAAAGAGGACACGGAGG   139
SCPP1˙DR            AGAGAACATCCGTAGGGGCTGGATCTACACCCTCAAA...........TGGGTCCAACCAAACAACAAC...ATTGTGC   385
SCPP1˙IP            AGAGATCATGCGCAAGAGCCCAGAGAAAAACTCCAAG...........AGCATCAGCAGCAGTAGTGAGAGCAGTG   370
SCPP1˙LW            AGAGAATGTCCGTAGGGGCTGGATCTACACCCTCAAA...........TGGGTCCATACGAATAACAGCGGCAAAGTGC   388
SCPP1˙PN            GGAGAACATGCGCAAGCTCCACCCATTCATACTGAGGTTCTAATGAAGCTGGATTCATCTCATCAACGTCAAAGTGCTCA   388
SCPP1˙PP            .....................................................................................   171
SCPP1-C1˙CC         CGAGAACGTTCGTAAGGGCTGGATCTACACCCTCAAA...........TGGGTTCATAAGAACAACAATGCCATTGTGC   382
SCPP1-C2˙CC         AGAGAATGTTCGTAGGGGCTGGATCTACACCCTCAAA...........TGGGTCCATATTAACAACAACGGCAAAGTGC   382
SCPP1-C1˙SR         AGAGAACGTTCGTAGGGGCTGGATTTACACCCTCAAA...........TGGGTCCATAAGAACAACAATTCCATTGTGC   244
SCPP1-C2˙SR         AGAGAATGTTCGTAGGGGCTGGATCTACACCCTCAAA...........TGGGTCCATATTAACAACAACGGCAAAGTGC   382
SCPP1-C1˙SA         AGAGAACGTTCGTAGGGGCTGGATTTACACCCTCAAA...........TGGGTCCATAAGAACAACAATGCCATTGTGC   382
SCPP1-C2˙SA         AGAGAATGTTCGTAGGGGCTGGATCTACACCCTCAAA...........TGGGTCCATATTAACAACAACGGCAAAGTGC   382
SCPP1-C1˙SG         AGAGAACGTTCGTAGGGGCTGGATTTACACCCTCAAA...........TGGGTCCATAAGAACAACAATGCCATTGTGC   382
SCPP1-C2˙SG         AGAGAATGTTCGTAGGGGCTGGATCTACACCCTCAAA...........TGGGTCCATATTAACAACAACGGCAAAGTGC   382
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               385 390 395 400 405 410 415 420 425 430 435 440 445 450 455 460
conservation   ******************* ***********  * ***** ** **        *       ** *********** ** 
SCPP1_CI       AAGCGACCAGCCAGCCAGACGAGAATTACATCAAATCCACCAGTGATGCATCAAAAAAAAGTGAAAGCAGTGAATCGAGT   462
SCPP1_AM       AAGTGACCGACCAACCAGACGAGGAGGACAAAGACACGACAGAGGAC...............CAGACCTCCGAGAGCAGC   204
SCPP1_DR       AACCAACCGGCCAGCCACATGAAAATGACATCCAGTCCACTAGTGACGCATCCCAAATAAGTGAGAGCAGCGAATCAAGC   465
SCPP1_IP       AGTCTACTGAGGGTCAAGGCAACAATTCAACAAGCAGCAGCAGCGAGAGCAGTGAGAGCAGCGAAAGCAGTGAGAGCAGC   450
SCPP1_LW       AAGCGACCAGCCAGCCGGAC..............................ATCAAATCCAGTGAAAGCAGTGAATCAAGT   438
SCPP1_PN       GCGAGGAGGACAAAGACACGACAGAGGACCTCTCCTCTAAC........................AGCAGTGAATCTACA   444
SCPP1_PP       ................................................................................   171
SCPP1-C1_CC    AAGCGACCAGCCAGCCAGACGAGAATGACATCAAATCCACTAGTGACGCATCAAAAAAAAGTGAAAGCAGTGTTTCAAGT   462
SCPP1-C2_CC    AAGCGACCAGCCAGCCAGATGAGAATGACATCAAGTCCACTAGTGATGCGTCAAAAAAAAGTGAAAGCAGTGAATCAAGT   462
SCPP1-C1_SR    AAGCGACCAGCCAGCCAGAAGAGAATGACATAAAATCCACTAGTGACGAATCAAAAAAAAGTGAAAGCAGTGAATCAAGT   324
SCPP1-C2_SR    AAGCGACCAGCCAGCCAGAAGAGAATGACATCAAATCCACTAGTGACGCGTCAAAAAAAAGTGAAAGCAGTGAATCAAGT   462
SCPP1-C1_SA    AAGCGACCAGCCAGCCAGAAGAGAATGACATCAAATCCACTAGTGACGAATCAAAAAAAAGTGAAAGCAGTGAATCAAGT   462
SCPP1-C2_SA    AAGCGACCAGCCAGCCAGAAGAGAATGACATCAAATCCACTAGTGACGCATCAAAAAAA......AGCAGTGAATCAAGT   456
SCPP1-C1_SG    AAGCGACCAGCCAGCCAGAAGAGAATGACATCAAATCCACTAGT..................GAAAGCAGTGAATCAAGT   444
SCPP1-C2_SG    AAGTGACCAGCCAGCCAGAAGAGAATGAC...................................................   411

               465 470 475 480 485 490 495 500 505 510 515 520 525 530 535 540
conservation   ****** ****** ********* * *** *** * *************  ***** *********************  
SCPP1_CI       GAAAGCCAGGAAACGGTTGTTCTGAACACAAAGGAAGAGGATAACAGTGCCGACACGAGTGAAAGCACTGAACACAGT..   540
SCPP1_AM       GAGTCCACGGAGAAGCCCACCCCCAGCAGCAGCAGCAGCAGCACAGAGGACAGC...............AGGGCGGT   269
SCPP1_DR       GAAAGCCAGGAAAAGGTTGTCGTGAACAAGATGCAAGTGGATAACAGTGACGACACCAGTGAAAGCGCTGAACACAATGG   545
SCPP1_IP       GAGTCCACTGAAAACCCGGAGAAGAACTCCAACAGCAGCAGCAACAGCAGCAGTGAGAGCAAAAGCGTCGAGAACAGATC   530
SCPP1_LW       GAAAGCCAGGAAACAGTTGTTCTGAGCACAAAGGAAGAGAATAACGGTGCCGACACAAGTGAAAGC..............   504
SCPP1_PN       GAGAGTCAGGAGCAGCAAACCCACAGCAGCAGCAGCAGCAGCAGCACAGAGTCCAGC...AGCAAAGACAGCAGGGC   521
SCPP1_PP       ................................................................................   171
SCPP1-C1_CC    GAAAGCCAGGAAATGGCTGTTCTAAACACAAAGCAAGAGGATAACAGTGTGGACACAAGTGAAAGCACTGAAAACAGTTA   542
SCPP1-C2_CC    GAAAGCAAGGAAACAGTTGTTCTGAACACCAAGGAGGAGGATAACAGTGCCGATACAAGTGAAAGCACTGAAAACAGTTA   542
SCPP1-C1_SR    GAAAGCCAGGAAATGGTTGTTCTAAGCACAAAGGAAGAGGATAAGAGTGCGGACATAAGTGAAAGCACTGAAAACAGTTA   404
SCPP1-C2_SR    GAAAGCAAGGAAATGGTTGTTCTGAACACAAAAAAGGAGGATAACAGTGCCGACACAAGTGAAAGCACTGAAAACAGTTA   542
SCPP1-C1_SA    GAAAGCCAGGAAATGGTTGTTCTAAGCACAAAGGAAGAGGATAAGAGTGTGGACACAAGTGAAAGCACTGAAAACAGTTA   542
SCPP1-C2_SA    GAAAGCAAGGAAATGGTTGTTCTGAACACAAAGGAGGAGGATAACAGTGCCGACACAAGTGAAAGCACTGAAAACAGTTA   536
SCPP1-C1_SG    GAAAGCCAGGAAATGGTTGTTCTAAGCACAAAGGAAGAGGATAAGAGTGCGGACACAAGTGAAAGCACTGAAAACAGTTA   524
SCPP1-C2_SG    ...AGCAAGGAAATGGTTGTTCTGAACACAAAGGAGGAGGATAACAGTGCCGACACAAATGAAAGCACTGAAAACAGTTA   488

               545 550 555 560 565 570 575 580 585 590 595 600 605
conservation   *   *********************** *****          ***************** ************** ****
SCPP1_CI       ....GTTGAAGGTACTGAGTATAATACGAGCAAC.........AGCAGCAGCAGCAGCAGTAGCAGTGAAAGCACCGAGT   607
SCPP1_AM       GGTGGTGGACAGCTCTGAGAACAGCCACAGCAAC............AGCAGCAGCAGCAGCAGCAGCGAGAGCCACGAGT   337
SCPP1_DR       TGTTGTT...GCTACTGAATACAGCAACAGTAAC.........AGCAGCAGCAGCAGTAGCAGCAGTGAAAGCACCGAGT   613
SCPP1_IP       AACGATCGACAGCGATGAGAACAGCGCCCTGAAA...AGCAGCAACAGCTCCAGCAGCGAGAGTCACGAGTCCACTGAGA   607
SCPP1_LW       ....ATTGAAGGTACTGAGTATAGTACGAGCAAC...AGCAGCAGCAGCAGCAGCAGCAGCAGCAGCGAAAGCACTGAGT   577
SCPP1_PN       TGTGGTGGACAGCAACGAGACCAGTGAGAGCAGAAGCAGCAACAGCAGTAGCAGCAACGAGAGCCGAGAGTCCAGCCACA   601
SCPP1_PP       ................................................................................   171
SCPP1-C1_CC    TATTGTTGACGGAACTGAGTACAGTACCAGCAACAGCAGCAGCAGCAGCAGCAGCAGCAGTAGCAGTGAAAGCACAGAGT   622
SCPP1-C2_CC    TAGTGTTGATGGTACTGAGTACAGTACTAGCAACAGCAGCAGCAGCAGCAGTAGCAGCAGCAGTGAAAGCACCGAGT   622
SCPP1-C1_SR    TATTGTTGACGGTACTGAGTACAGTACTAGCAAT......AGCAGCAGCAGCAGCAGTAGCAGCAGTGAAAGCACCGAGT   478
SCPP1-C2_SR    TAGTGTTGACGGTACTGAGTACAGTACCAGCAAT.........AGCAGCAGCAGCAGCAGTAGCAGTGAAAGCACTGAGT   613
SCPP1-C1_SA    TATTGTTGACGGTACTGAGTACAGTACTAGCAAT.........AGCAGCAGCAGCAGCAGTAGCAGTGAAAGCACCGAGT   613
SCPP1-C2_SA    TAGTGTTGACGGTACTGAGTACAGTACCAGCAAT......AGCAGCAGCAGCAGCAGTAGCAGCAGTGAAAGCACTGAGT   610
SCPP1-C1_SG    TATTGTTGACGGTACTGAGTACAGTACTAGCAAT.........AGCAGCAGCAGCAGCAGTAGCAGTGAAAGCACCGAGT   595
SCPP1-C2_SG    TAGTGTTGACGGTACTGAGTACAGTACCAGCAAT.........AGCAGCAGCAGCAGTAGCAGCAGTGAAAGCACCGAGT   559

               610 615 620 625 630 635 640 645 650 655 660 665 670 675 680 685
conservation   ***** **** **** ****** ***** ******* ******** *** ******************************
SCPP1_CI       CCAAAAACAGCGCAGTGGATAGCGAAAGTCGCTCAACTGAGTGCCTTCCCGGGGCTGACAGCCAAGAATGTGACAGCGAG   687
SCPP1_AM       CCAGCCACAGCACAGAGAGCACAGAGAGCCAGTCCAAAGAGTGCCCGCCCGGAACAGACAGCAACGAGTGCGACAGCGAC   417
SCPP1_DR       CTAAAGACAGCGGCACGGATAGTGAAAGTCGCTCCACTGAGTGTGTGCCTGGCGATGACAGTCAAGATTGTGAGAGCGAG   693
SCPP1_IP       CCACCGAGAGCACCGAAAGCAAACAGAGTCGCTCTAACGAGTGCCAGCCTGGAGCCGACAGCCAGGACTGCGACAGCGAC   687
SCPP1_LW       CCAAAAGCAGTGCGATGGAGAGCGAAAGTCGCTCAACTGAGTGCCTTCCCGGGGCTGACAGCCAAGAATGTGACAGCGAG   657
SCPP1_PN       GCACAGAGAGCAGGGAAAGCACAGAGAGCCGCTCGAAAGAGTGCCCGTCTGAGGCCGACAGCGATGAGTGCGACAGCGAT   681
SCPP1_PP       ................................................................................   171
SCPP1-C1_CC    CCAAAAACAGTGCAGTGGACAGCGAAAGTCGCTTAAGTGAGTGCCAGCCAGGGGCTGACAGCCAAGAATGTGACAGCGAG   702
SCPP1-C2_CC    CCAAAAACAACGCAGTGGACAGTGAAAGTCGCTCAGCTGAGTGCCAGCCAGGGGCTAACAGCCAAGAATGTGACAGCGAG   702
SCPP1-C1_SR    CCAAAAACAGTGCAGTGGACAGCGAAAGTCGCTCAACTGAGTGCCAGCCAGGGGCTGACAGCCAAGAATGTGACAGCGAG   558
SCPP1-C2_SR    CCAAAAACAGTGCAGTGGACAGCGAAAGCCGCTCAGCTGAGTGCCAGTCAGGGGCTGACAGCCAAGAATGTGACAGCGAG   693
SCPP1-C1_SA    CCAAAGACAGCGCAGTGGACAGTGAAAGTCGCTCAATTGAGTGCCAGCCAGGGGCTGACAGCCAGGAATGTGACAGCGAG   693
SCPP1-C2_SA    CCAAAAACAGTGCAGCAGACAGCGAAAGCCGCTCAGCTGAGTGCCAGCCAGAGGCTGACAGCCAAGAATGTGACAGCGAG   690
SCPP1-C1_SG    CCAAACACAGCGCAGTGGACAGCGAAAGTCGCTCAACTGAGTGCCAGCCAGGGGCTGACAGCCAAGAATGTGACAGCGAG   675
SCPP1-C2_SG    CCAAAAACAGTGCACCGGACAGCGAAAGCCGCTCAGCTGAGTGCCAGCCAGAGGCTGACAGCCAAGAATGTGACAGCGAG   639

               690 695 700 705 710 715 720 725 730 735 740 745 750 755 760
conservation   ******      ************************************************ *******************
SCPP1_CI       GAAGAC......TTTCCTCAGGATATCGGTGATGATGGTGCAACCGACCCATTCAATGGCTTCCTAATGCCTGATGTCAC   761
SCPP1_AM       GAATAC......CAGTTCCACGACGTCGGAGACGATGGAGCCACTGACCCCTTCAACGGCTTCCACACGACCGACAACGC   491
SCPP1_DR       GAAAAC......CTTCCTCAGGATATCGGTGATGATGGAGCAACCGACCCTTTCAATGGCTTCTTAATGCCTGATGTCGC   767
SCPP1_IP       AGCGACGAGTACGTCCTCCAGAACGTCGGAGACGACGGAACCAACGACCCATTTGACGGCTTCCACGTGCCTGACAGCAC   767
SCPP1_LW       GAAGCC......TTTCCTCAGGATATCGGTGATGATGGTGCAACCGACCCTTTCAATGGCTTCCTAATGCCTGATGTCAC   731
SCPP1_PN       GAATAC......GACTTCCACGACATCGGAGACGATGGAGCCAGTGAACCGCTGAACGGCTTCCGTACGCCCGAT.....   750
SCPP1_PP       ................................................................................   171
SCPP1-C1_CC    GAAGAC......TTTCCTCAGGATATCGGTGATGATGGTGCAACCGACCCTTTCAATGGCATCCTAATGCCTGATGTCAC   776
SCPP1-C2_CC    GAAGAC......TTTCCTCAGGATATCGGTGATGATGGTGAAACCGACCCTTTCAATGGCATCCTAATGCCTGATGTCAC   776
SCPP1-C1_SR    GAAGAC......TTTCCTCAGGATATCGGTGATGATGGTGCAACCGACCCTTTCAATGGCATCCTAATGCCTGATGTCAC   632
SCPP1-C2_SR    GAAGAC......TTTCCTCAGGATATCGGTGATGATGGTGCAACCGACCCTTTCAATGGCATCCTAATGCCTGATGTCAC   767
SCPP1-C1_SA    GAAGAC......TTTCCTCAGGATATCGGTGATGATGGTGCAACCGACCCTTTCAATGGCATCCTAATGCCTGATGTCAC   767
SCPP1-C2_SA    GAAGAC......TTTCCTCAGGATATCGGTGATGATGGTGCAACCGACCCTTTCAATGGCATCCTAATGCCTGATGTCAC   764
SCPP1-C1_SG    GAAGAC......TTTCCTCAGGATATCGGTGATGATGGTGCAACGGACCCTTTCAATGGCATCCTAATGCCTGATGTCAC   749
SCPP1-C2_SG    GAAGAC......TTTCCTCAGGATATCGGTGATGATGGTGCAACCGACCCTTTCAATGGCATCCTAATGCCTGATGTCAC   713
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                    765       770
conservation        **********
SCPP1˙CI            TGAGCCCTAG   771
SCPP1˙AM            C.........   492
SCPP1˙DR            TGAGCCCTAG   777
SCPP1˙IP            CGAG......   771
SCPP1˙LW            TGAGCCCTAG   741
SCPP1˙PN            ..........   750
SCPP1˙PP            ..........   171
SCPP1-C1˙CC         TGAGCCCTAG   786
SCPP1-C2˙CC         TGAGCCCTAG   786
SCPP1-C1˙SR         TGAGCCCTAG   642
SCPP1-C2˙SR         TGAGCCCTAG   777
SCPP1-C1˙SA         TGAGCCCTAG   777
SCPP1-C2˙SA         TGAGCCCTAG   774
SCPP1-C1˙SG         TGAGCCCTAG   759
SCPP1-C2˙SG         TGAGCCCTAG   723

                    X  non conserved
                    X  ≥55% conserved
                    X  ≥85% conserved
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