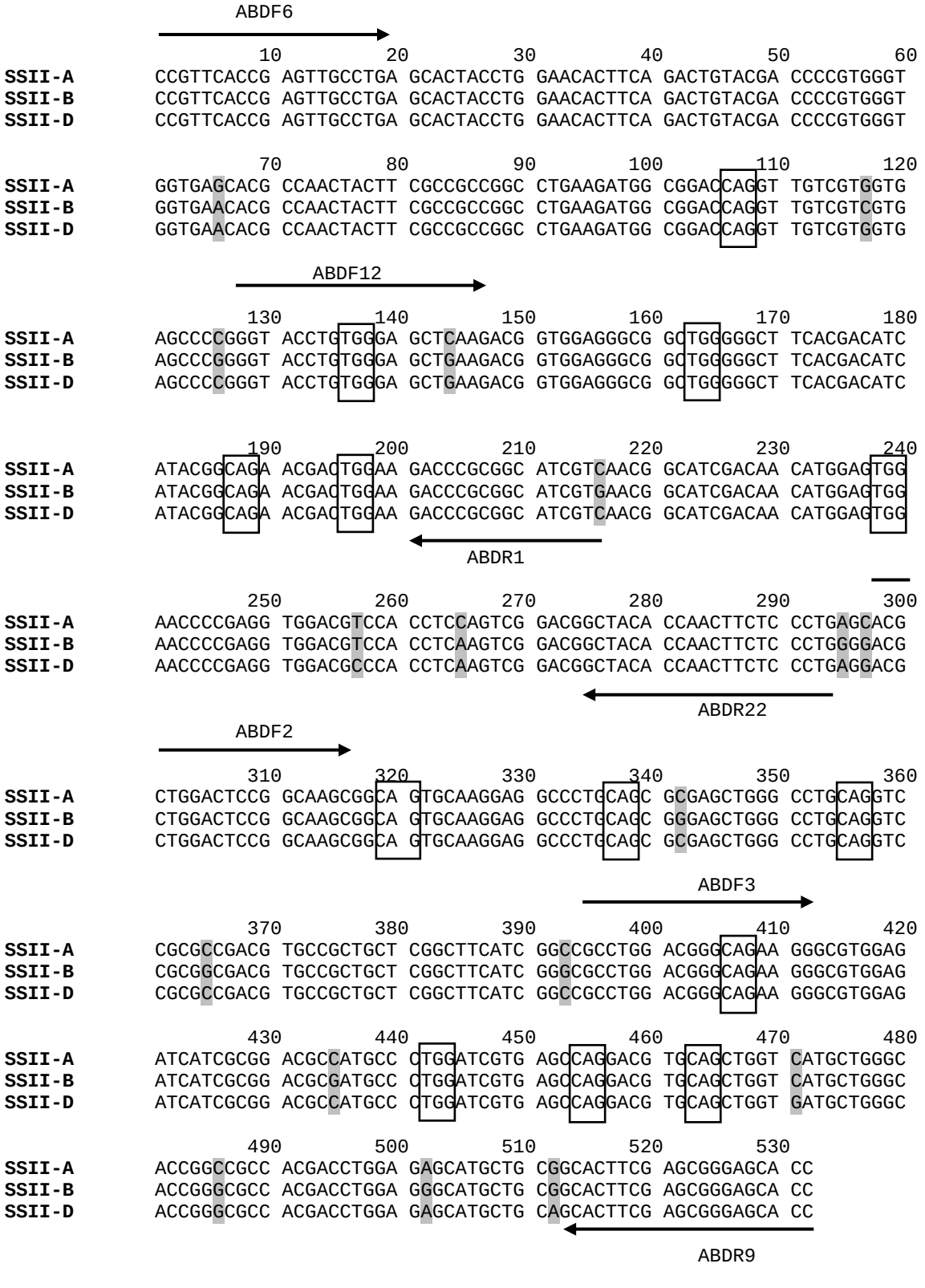
ABDF6
10 20 30 40 50 60
SSII-A CCGTTCACCG AGTTGCCTGA GCACTACCTG GAACACTTCA GACTGTACGA CCCCGTGGGT
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SSII-D CCGTTCACCG AGTTGCCTGA GCACTACCTG GAACACTTCA GACTGTACGA CCCCGTGGGT
70 80 90 100 110 120
SSII-A GGTGAGCACG CCAACTACTT CGCCGCCGGC CTGAAGATGG CGGACCAGGT TGTCGTGGTG
SSII-B GGTGAACACG CCAACTACTT CGCCGCCGGC CTGAAGATGG CGGACCAGGT TGTCGTCGTG
SSII-D GGTGAACACG CCAACTACTT CGCCGCCGGC CTGAAGATGG CGGACCAGGT TGTCGTGGTG
ABDF12
130 140 150 160 170 180
SSII-A AGCCCCGGGT ACCTGTGGGA GCTCAAGACG GTGGAGGGCG GCTGGGGGCT TCACGACATC
SSII-B AGCCCGGGGT ACCTGTGGGA GCTGAAGACG GTGGAGGGCG GCTGGGGGCT TCACGACATC
SSII-D AGCCCCGGGT ACCTGTGGGA GCTGAAGACG GTGGAGGGCG GCTGGGGGCT TCACGACATC
190 200 210 220 230 240
SSII-A ATACGGCAGA ACGACTGGAA GACCCGCGGC ATCGTCAACG GCATCGACAA CATGGAGTGG
SSII-B ATACGGCAGA ACGACTGGAA GACCCGCGGC ATCGTGAACG GCATCGACAA CATGGAGTGG
SSII-D ATACGGCAGA ACGACTGGAA GACCCGCGGC ATCGTCAACG GCATCGACAA CATGGAGTGG
ABDR1
250 260 270 280 290 300
SSII-A AACCCCGAGG TGGACGTCCA CCTCCAGTCG GACGGCTACA CCAACTTCTC CCTGAGCACG
SSII-B AACCCCGAGG TGGACGTCCA CCTCAAGTCG GACGGCTACA CCAACTTCTC CCTGGGGACG
SSII-D AACCCCGAGG TGGACGCCCA CCTCAAGTCG GACGGCTACA CCAACTTCTC CCTGAGGACG
ABDR22
ABDF2
310 320 330 340 350 360
SSII-A CTGGACTCCG GCAAGCGGCA GTGCAAGGAG GCCCTGCAGC GCGAGCTGGG CCTGCAGGTC
SSII-B CTGGACTCCG GCAAGCGGCA GTGCAAGGAG GCCCTGCAGC GGGAGCTGGG CCTGCAGGTC
SSII-D CTGGACTCCG GCAAGCGGCA GTGCAAGGAG GCCCTGCAGC GCGAGCTGGG CCTGCAGGTC
ABDF3
370 380 390 400 410 420
SSII-A CGCGCCGACG TGCCGCTGCT CGGCTTCATC GGCCGCCTGG ACGGGCAGAA GGGCGTGGAG
SSII-B CGCGGCGACG TGCCGCTGCT CGGCTTCATC GGGCGCCTGG ACGGGCAGAA GGGCGTGGAG
SSII-D CGCGCCGACG TGCCGCTGCT CGGCTTCATC GGCCGCCTGG ACGGGCAGAA GGGCGTGGAG
430 440 450 460 470 480
SSII-A ATCATCGCGG ACGCCATGCC CTGGATCGTG AGCCAGGACG TGCAGCTGGT CATGCTGGGC
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490 500 510 520 530
SSII-A ACCGGCCGCC ACGACCTGGA GAGCATGCTG CGGCACTTCG AGCGGGAGCA CC
SSII-B ACCGGGCGCC ACGACCTGGA GGGCATGCTG CGGCACTTCG AGCGGGAGCA CC
SSII-D ACCGGGCGCC ACGACCTGGA GAGCATGCTG CAGCACTTCG AGCGGGAGCA CC
ABDR9