



Research article

Identification of auxin response factor in the pan-genome and their expression pattern analysis in cucumber

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Abstract: Auxin response factors (ARFs) are important transcription factors that regulate the expression of early genes in response to auxin. Auxin is a crucial phytohormone that controls plant growth and development. Even though *ARF* genes have been identified in the cucumber (*Cucumis sativus*) genome version 3.0, the features of this gene family are unclear in the cucumber pan-genome and in the newly assembled genome v4.0. We identified a total of 18 CsARF genes, including two CsARF13s in Gy14. All ARF proteins, except ARF5 and 7, showed differences in protein length; all of them showed amino acid variation in the cucumber pan-genome, including 13 accessions. We also analyzed their organ-specific expression and their expression patterns in fruit development and in response to biotic stresses. Importantly, we found that the overexpression of CsARF12 significantly enhanced the resistance of cucumber cotyledons to gray mold infection. Therefore, this study lays a foundation for understanding the biological functions of CsARF genes and provides *CsARF12* as a candidate gene to improve the cucumber resistance to gray mold.

Keywords: cucumber; pan-genome; *ARF* gene; biotic stress response; gray mold

Supplementary

Table S1. The variation of CsARF amino acids in 13 cucumber accessions (See file named “molsci-12-03-019-s002”).

Table S2. Primers used in instantaneous infection experiment (See file named “molsci-12-03-019-s002”).

Figure S1. The gene developmental tree, conserved protein motifs, and gene structure of *CsARF* genes.

Figure S2. The cis-acting elements of *CsARF* genes in the pan-genome.

Figure S3. Single amino acid differences with amino acid properties in the pan-genome.

Figure S4. Single amino acid differences without changes in the amino acid properties in the pan-genome.

Figure S5. The absence of amino acid fragments in the pan-genome.

Figure S6. Insertions of amino acid fragments in the pan-genome.

Figure S7. The phylogenetic tree of ARF proteins from the cucumber variety 9930 and *Arabidopsis* genomes.

Figure S8. The conserved protein motifs and gene structure of CsaV4_2G000004 and CsaV3_2G001050.

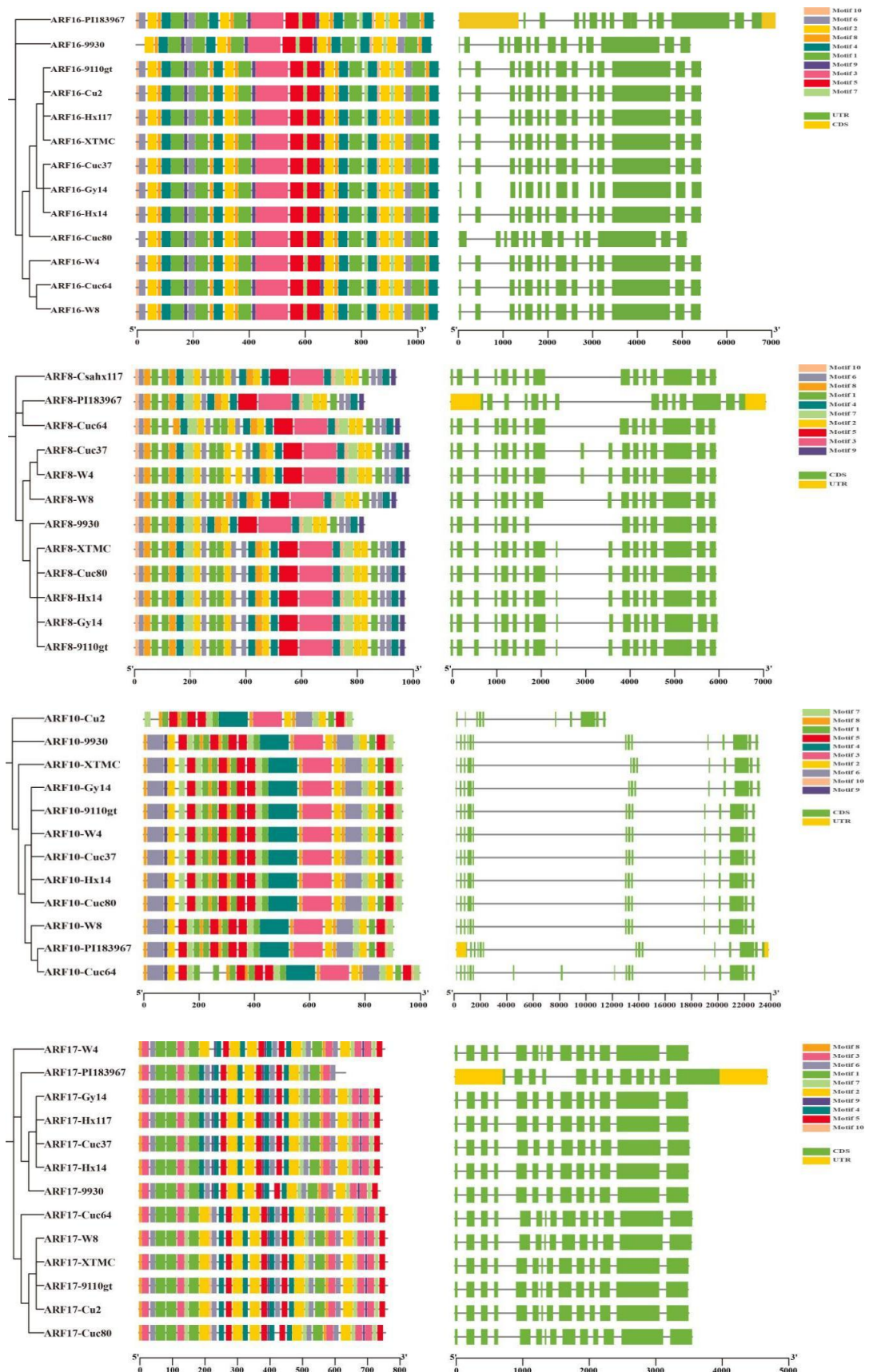
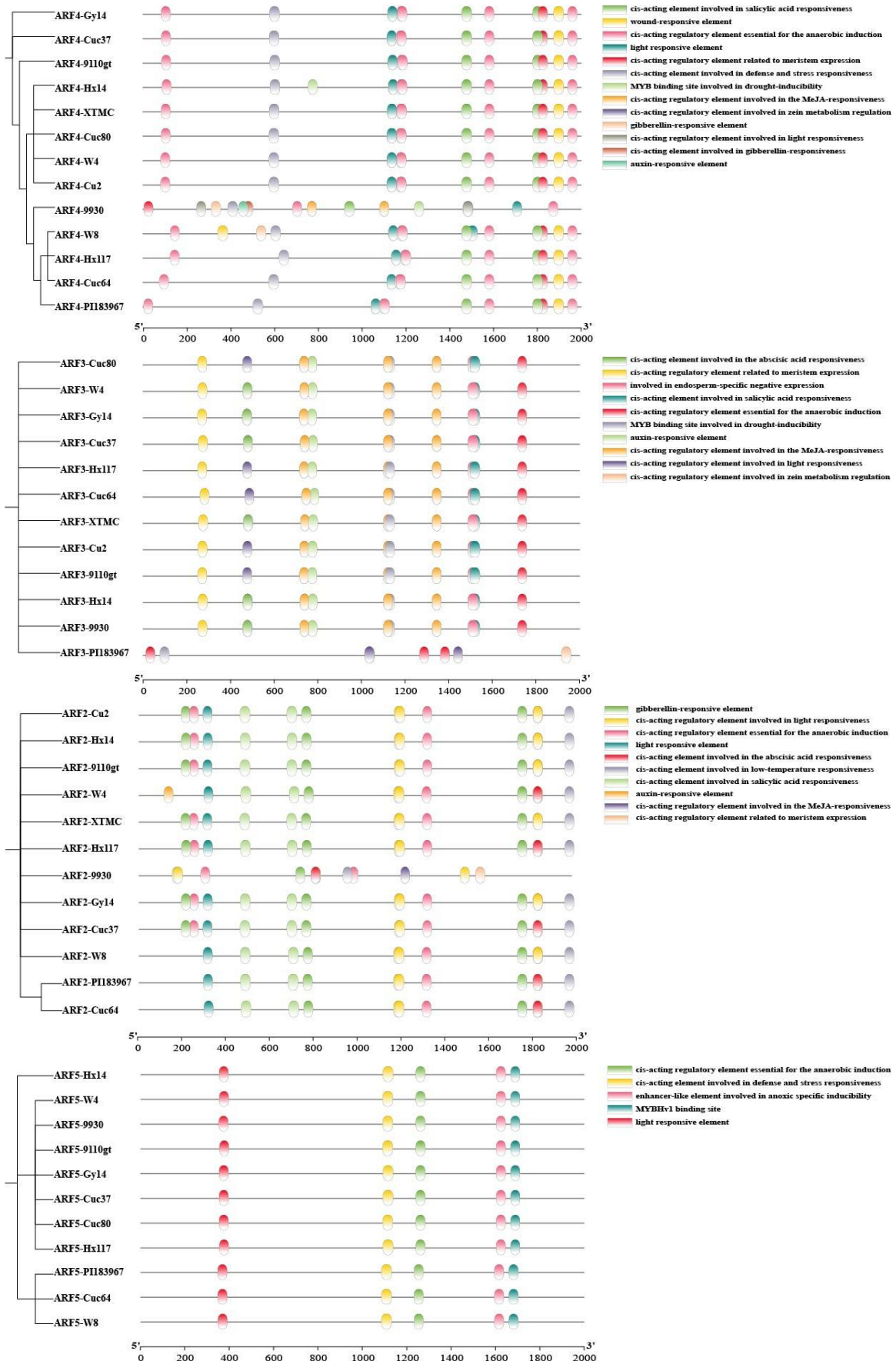
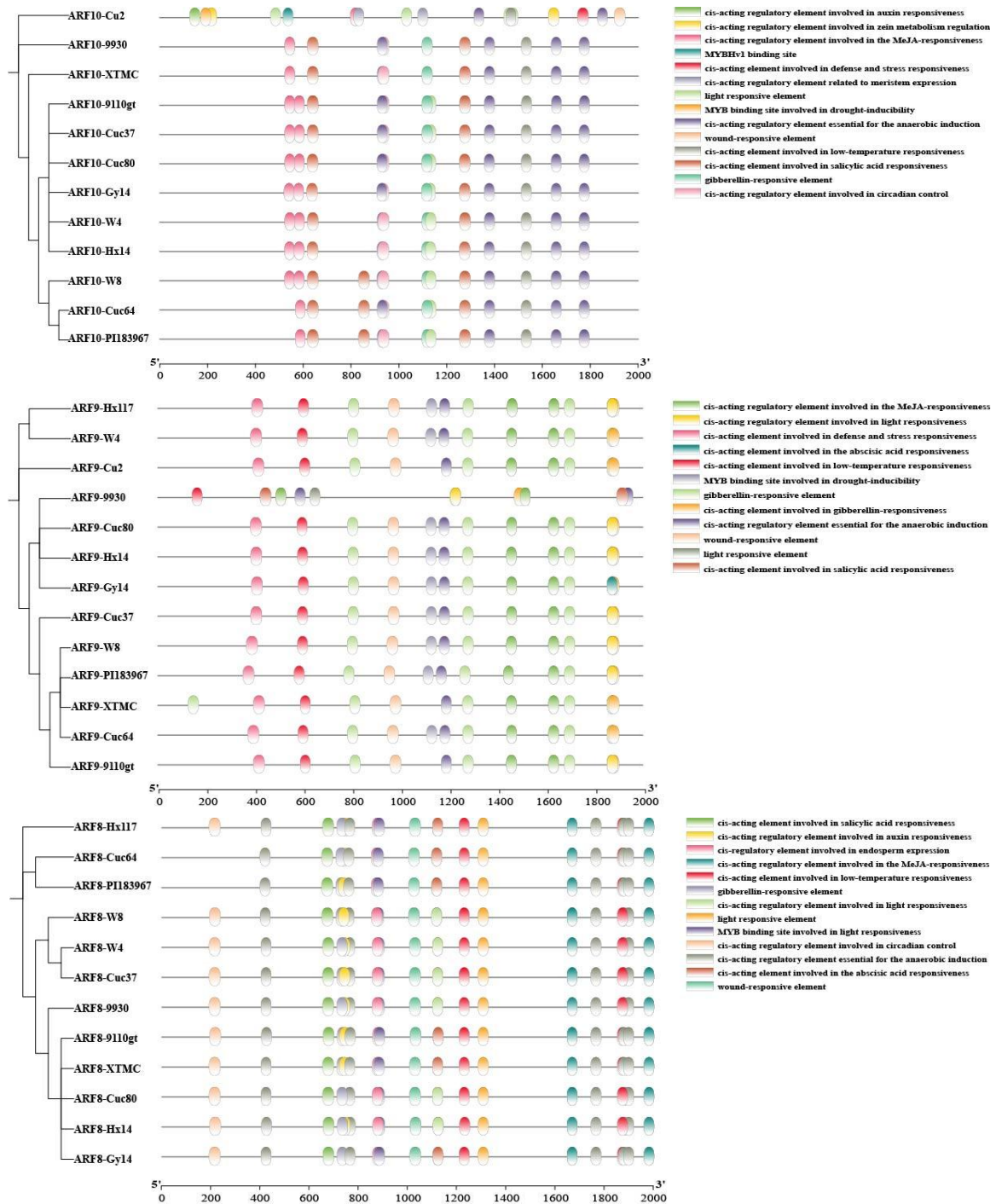
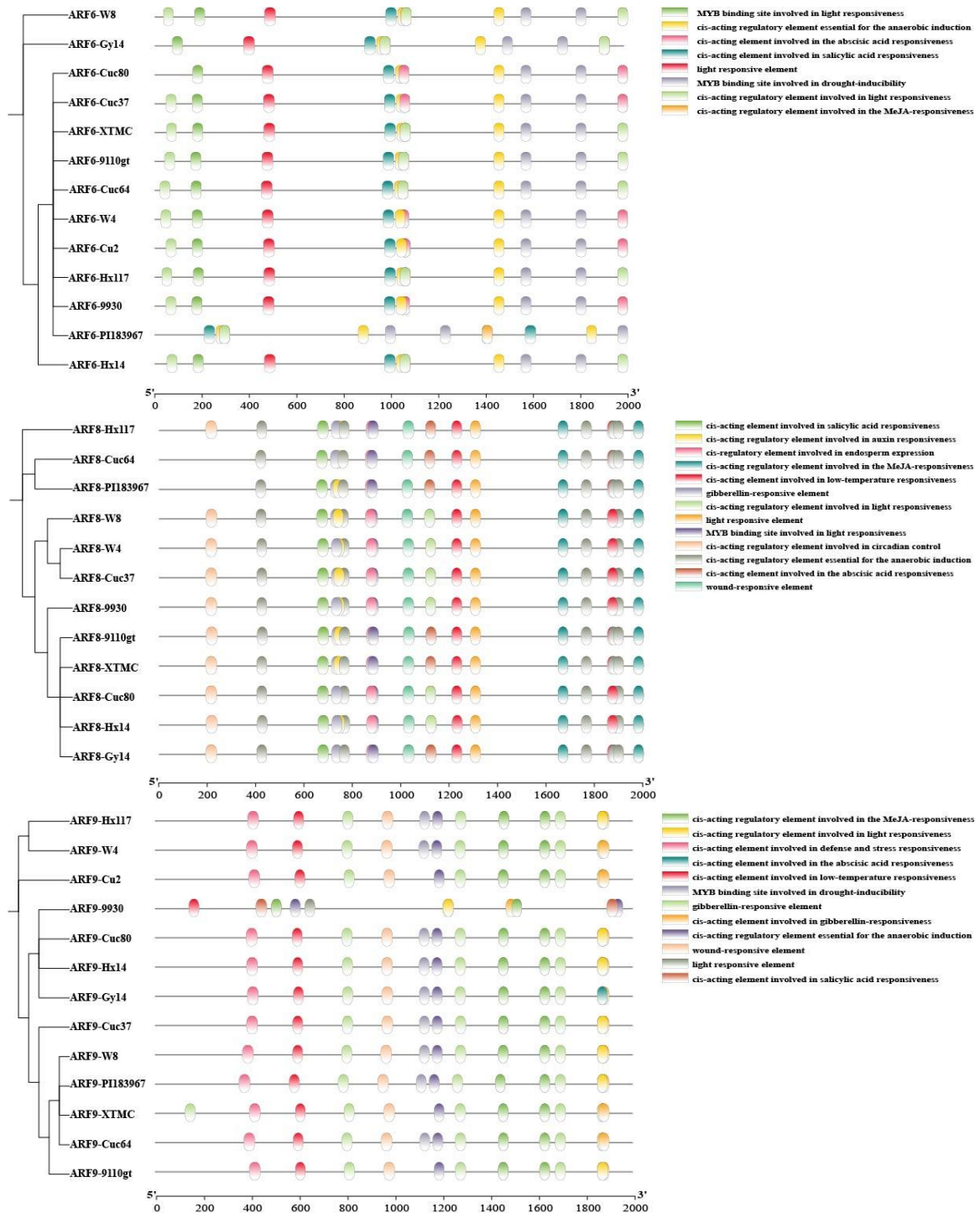


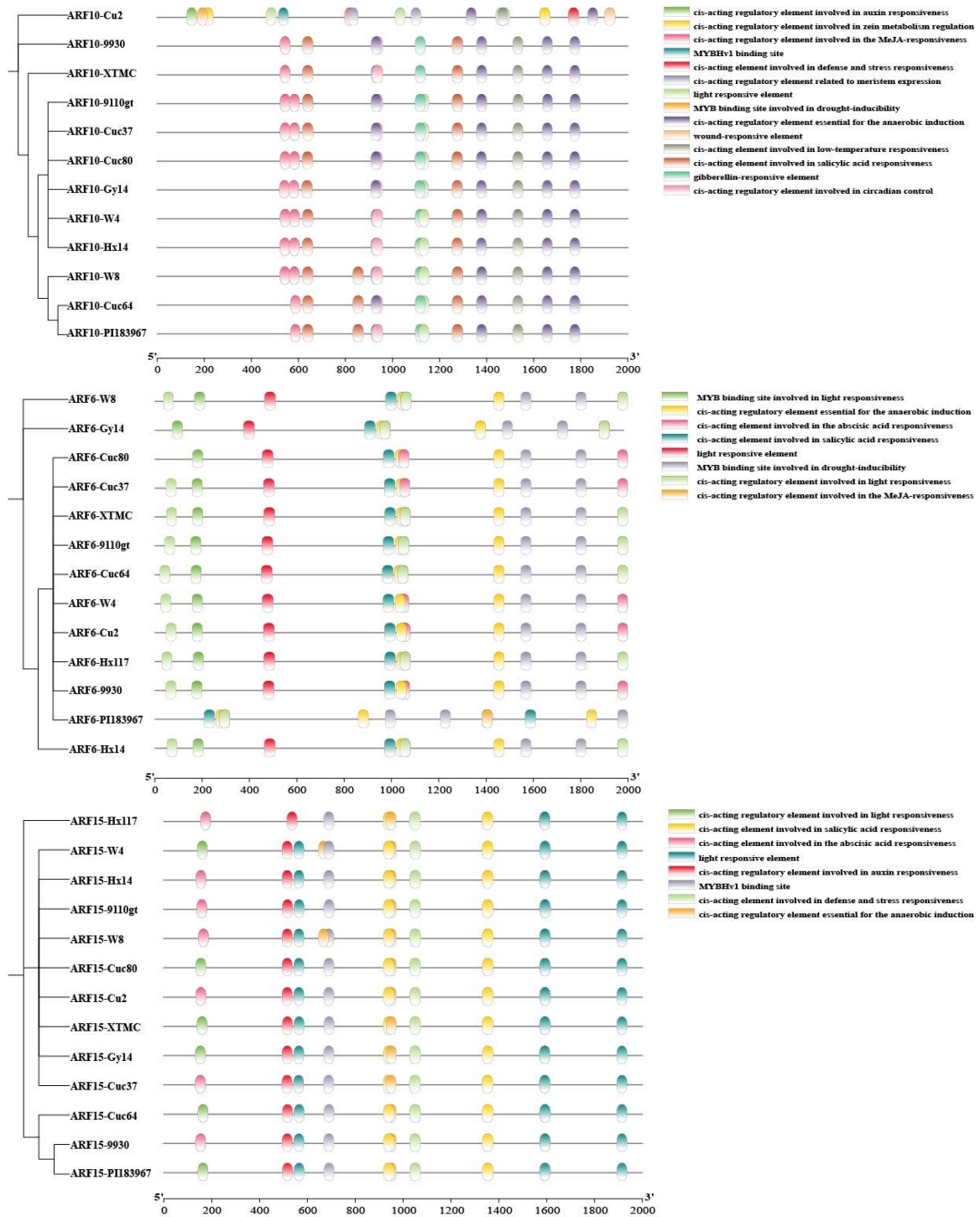
Figure S1. The gene developmental tree, conserved protein motifs, and gene structure of *CsARF*

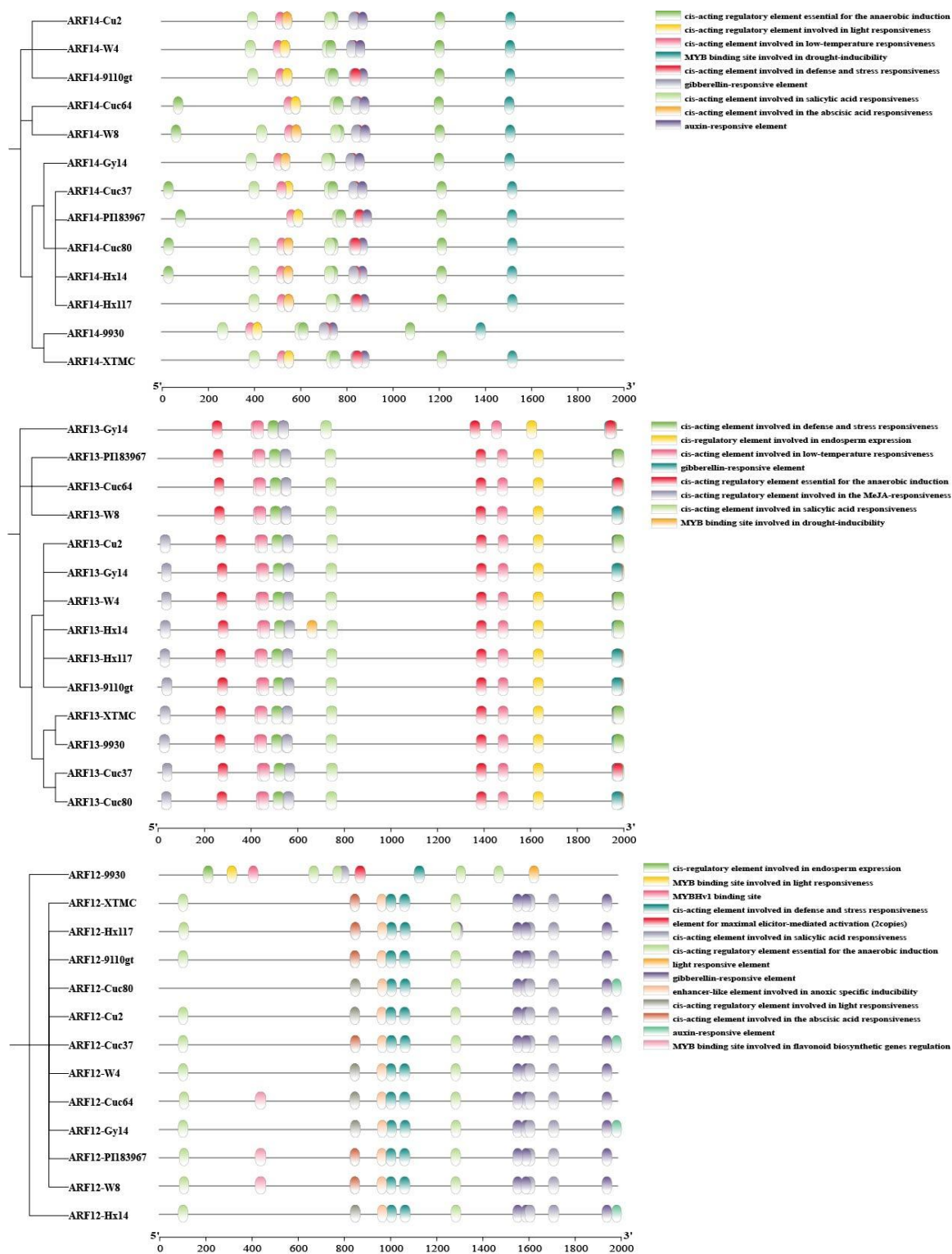
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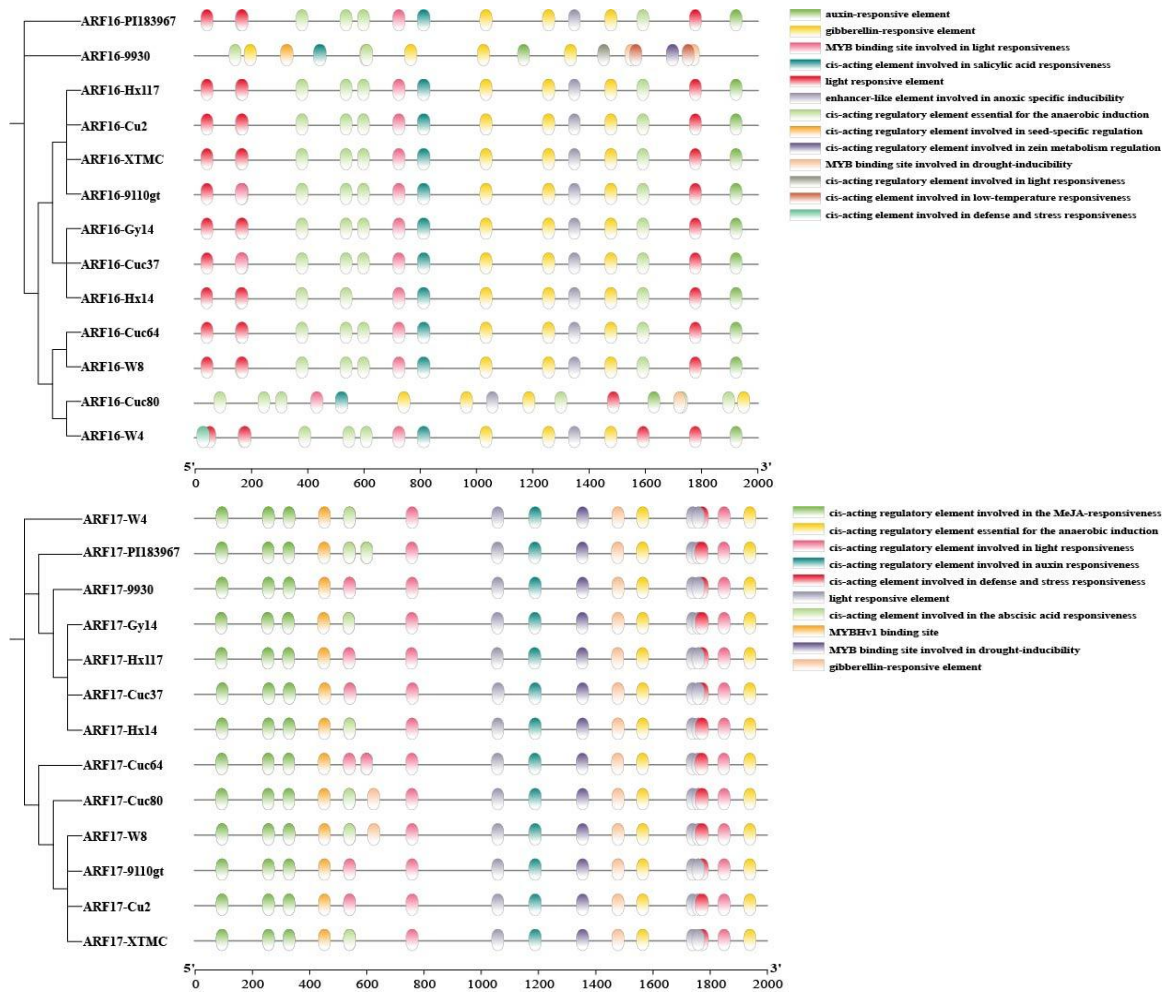
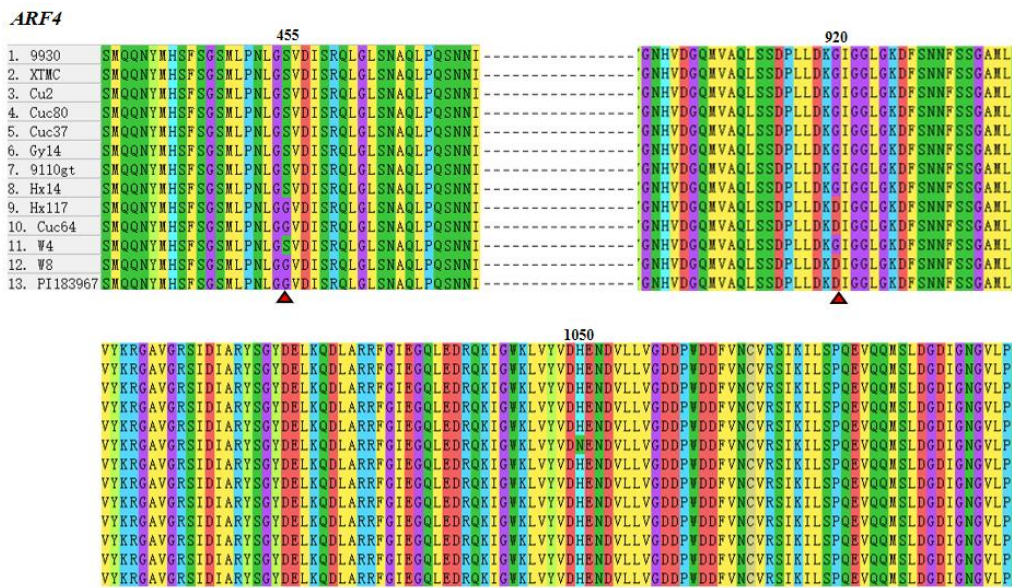


Figure S2. The cis-acting elements of *CsARF* genes in the pan-genome.



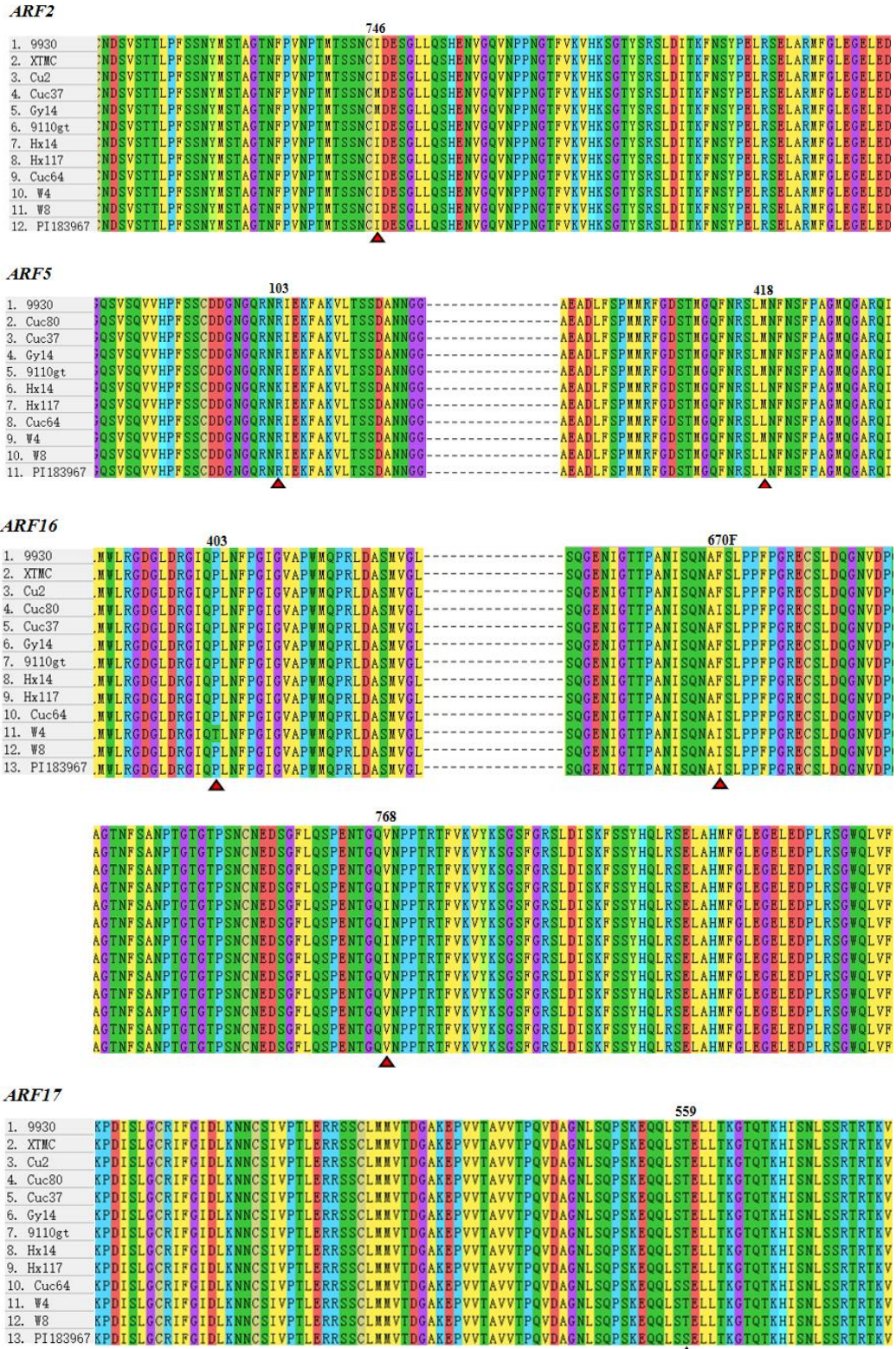


Figure S4. Single amino acid differences without changes in the amino acid properties in the pan-genome.

ARF1

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2. XTMC	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
3. Cuc2	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
4. Cuc80	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
5. Cuc37	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
6. Gy14	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
7. 9110gt	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
8. Hx14	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
9. Hx117	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
10. Cuc64	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
11. W8	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
12. W4	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		
13. PI183967	AELDQLFEPGGELQAPKKNLVIVYDDEGDHMLVGDPPREFCGMVRKIFIIYTRREEVQKMPGSLNLKGDENPSVEGEEVKETKSGAVPSMSAPESS		

ARF2

	1	84
1. 9930	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
2. XTMC	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
3. Cuc2	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
4. Cuc37	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
5. Gy14	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
6. 9110gt	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
7. Hx14	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
8. Hx117	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
9. Cuc64	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
10. W4	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
11. W8	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	
12. PI183967	MRLSAGGFSPPAQEGERRVNLSELWHACAPLVSLPAVGSRVVYFPQGHSEQVAASTNREVDQIPNYPNLPPLQLICQLHNLTHADAETDEVYAGM	

ARF3

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1. 9930	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
2. XTMC	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
3. Cuc2	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
4. Cuc80	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
5. Cuc37	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
6. Gy14	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
7. 9110gt	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
8. Hx14	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
9. Hx117	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
10. Cuc64	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
11. W4	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	
12. PI183967	MKAPSNGLFLANSGEGERKINSELWHACAPLVSLPPVGLVVYFPQGHSEQVAASMKETDFIPNYPNLPKSLICMLHNVTLHADPETDEVYAGM	

ARF6

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2. XTMC	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
3. Cuc2	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
4. Cuc80	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
5. Cuc37	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
6. Gy14	-----NKKLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
7. 9110gt	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
8. Hx14	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
9. Hx117	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
10. Cuc64	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
11. W4	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
12. W8	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	
13. PI183967	MGSVEEKLKTSGGLINNAPQTNLLDENKLLKENQDQSGARKAINSELWHACAPLVSLPHVGLVYVYFPQGHSEQVAVSTKRTATSIQIPNYPNLPSC	

ARF16

4 5
1. 9930 MIHS-----LERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
2. XTMC MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
3. Cu2 MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
4. Cuc80 MKCWDSELVSKFSGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
5. Cuc37 MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
6. Gy14 MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
7. 9110gt MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
8. Hx14 MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
9. Hx117 MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
10. Cuc64 MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
11. W4 MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
12. W8 MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA
13. PI183967 MRLSTAGFSPQAPBEGERRVLSNELWHACAGPLVSLPAVGSRRVYFPOGHSEQVAISTNREVDHIPSYPSPLPQLICQLHNVVTHADIETDEVYA

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AALQEMRTVDFAKAQAASLLQFQQTQNLNRPANFMPQMLQQPQPQP--PQTFLOGDENQHLSSHQAQSQPTAVLQQEIKHOTFNNHPQQQQ
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AALQEMRTVDFAKAQAASLLQFQQTQNLNRPANFMPQMLQQPQPQP--PQTFLOGDENQHLSSHQAQSQPTAVLQQEIKHOTFNNHPQQQQ
AALQEMRTVDFAKAQAASLLQFQQTQNLNRPANFMPQMLQQPQPQP--PQTFLOGDENQHLSSHQAQSQPTAVLQQEIKHOTFNNHPQQQQ
AALQEMRTVDFAKAQAASLLQFQQTQNLNRPANFMPQMLQQPQPQP--PQTFLOGDENQHLSSHQAQSQPTAVLQQEIKHOTFNNHPQQQQ
AALQEMRTVDFAKAQAASLLQFQQTQNLNRPANFMPQMLQQPQPQP--PQTFLOGDENQHLSSHQAQSQPTAVLQQEIKHOTFNNHPQQQQ
AALQEMRTVDFAKAQAASLLQFQQTQNLNRPANFMPQMLQQPQPQP--PQTFLOGDENQHLSSHQAQSQPTAVLQQEIKHOTFNNHPQQQQ

ARF12

453 454
1. 9930 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
2. XTMC EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
3. Cu2 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
4. Cuc80 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
5. Cuc37 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
6. Gy14 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
7. 9110gt EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
8. Hx14 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
9. Hx117 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
10. Cuc64 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
11. W4 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
12. W8 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI
13. PI183967 EDSVRSKVLQGGQENVVIVSFFYCDTVKRSLEFDVRSAAQGNQVSGGVEKLNIAIYVYKVHANSSFTGFMESDRFLKVLQGGQICSLRPPTSKPEYI

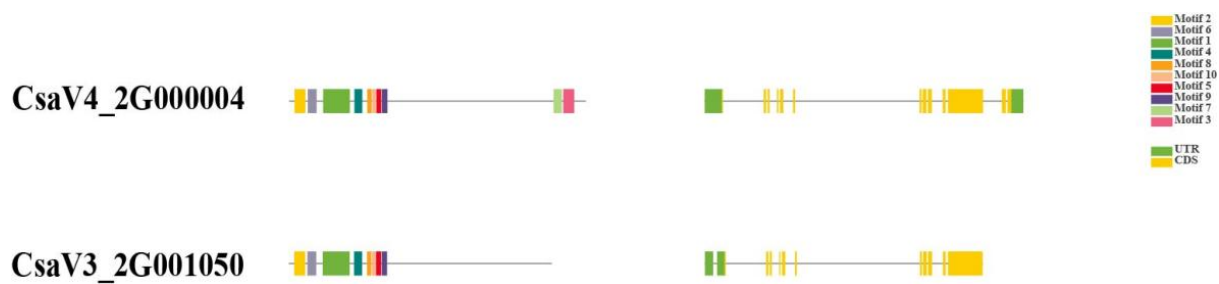


Figure S8. The conserved protein motifs, and gene structure of CsaV4_2G000004 and CsaV3_2G001050.



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