

Table S5 Sequence alignment of variable region in the promoter of *CsCBF1* in 10 cucumber lines

Lines	Sequence
CC	GATAGTAAAAGAGAAAAGTAGAATTTTCGATGGTGAAGAACAAGTCAAAGTGACACATGTCAGTTCCTGA
R1461	GATAGTAAAAGAGAAAAGTAGAATCCTTCGATGGTGAAGAACAAGTCAAAGTGACACATGTCAGTTCCTGA
9930	GATAGTAAAAGAGAAAAGTAGAATCCTTCGATGGTGAAGAACAAGTCAAAGTGACACATGTCAGTTCCTGA
HN	GATAGTAAAAGAGAAAAGTAGAATCCTTCGATGGTGAAGAACAAGTCAAAGTGACACATGTCAGTTCCTGA
XSBN	GATAGTAAAAGAGAAAAGTAGAATCCTTCGATGGTGAAGAACAAGTCAAAGTGACACATGTCAGTTCCTGA
HNHY	GATAGTAAAAGAGAAAAGTAGAATCCTTCGATGGTGAAGAACAAGTCAAAGTGACACATGTCAGTTCCTGA
HNHP	GATAGTAAAAGAGAAAAGTAGAATCCTTCGATGGTGAAGAACAAGTCAAAGTGACACATGTCAGTTCCTGA
ZN	GATAGTAAAAGAGAAAAGTAGAATCCTTCGATGGTGAAGAACAAGTCAAAGTGACACATGTCAGTTCCTGA
NT	GATAGTAAAAGAGAAAAGTAGAATCCTTCGATGGTGAAGAACAAGTCAAAGTGACACATGTCAGTTCCTGA
HNND	GATAGTAAAAGAGAAAAGTAGAATCCTTCGATGGTGAAGAACAAGTCAAAGTGACACATGTCAGTTCCTGA
CC	AGTTGACTCATTGTTTATCAATTTTCATTTTCAAATATCTTTTAGTATTTTATGAATTTAAAAA
R1461	AGTTGACTCATTGTTTATCAATTTTCATTTTCAAATATCTTTTAGTATTTTATGAATTTAAAAA
9930	AGTTGACTCATTGTTTATCAATTTTCATTTTCAAATATCTTTTAGTATTTTATGAATTTAAAAA
HN	AGTTGACTCATTGTTTATCAATTTTCATTTTCAAATATCTTTTAGTATTTTATGAATTTAAAAA
XSBN	AGTTGACTCATTGTTTATCAATTTTCATTTTCAAATATCTTTTAGTATTTTATGAATTTAAAAA
HNHY	AGTTGACTCATTGTTTATCAATTTTCATTTTCAAATATCTTTTAGTATTTTATGAATTTAAAAA
HNHP	AGTTGACTCATTGTTTATCAATTTTCATTTTCAAATATCTTTTAGTATTTTATGAATTTAAAAA
ZN	AGTTGACTCATTGTTTATCAATTTTCATTTTCAAATATCTTTTAGTATTTTATGAATTTAAAAA
NT	AGTTGACTCATTGTTTATCAATTTTCATTTTCAAATATCTTTTAGTATTTTATGAATTTAAAAA
HNND	AGTTGACTCATTGTTTATCAATTTTCATTTTCAAATATCTTTTAGTATTTTATGAATTTAAAAA
CC	AATAGAAAAGGGCCAAAGAGAAAAAATAAAAAAATAAAAAAAGGGAAGGTGTCGTTGATGGCGTGGG
R1461	AATAGAAAAGGGCCAAAGAGAAAAAATAAAAAAATAAAAAAAGGGAAGGTGTCGTTGATGGCGTGGG
9930	AATAGAAAAGGGCCAAAGAGAAAAAATAAAAAAATAAAAAAAGGGAAGGTGTCGTTGATGGCGTGGG
HN	AATAGAAAAGGGCCAAAGAGAAAAAATAAAAAAATAAAAAAAGGGAAGGTGTCGTTGATGGCGTGGG
XSBN	AATAGAAAAGGGCCAAAGAGAAAAAATAAAAAAATAAAAAAAGGGAAGGTGTCGTTGATGGCGTGGG
HNHY	AATAGAAAAGGGCCAAAGAGAAAAAATAAAAAAATAAAAAAAGGGAAGGTGTCGTTGATGGCGTGGG
HNHP	AATAGAAAAGGGCCAAAGAGAAAAAATAAAAAAATAAAAAAAGGGAAGGTGTCGTTGATGGCGTGGG
ZN	AATAGAAAAGGGCCAAAGAGAAAAAATAAAAAAATAAAAAAAGGGAAGGTGTCGTTGATGGCGTGGG
NT	AATAGAAAAGGGCCAAAGAGAAAAAATAAAAAAATAAAAAAAGGGAAGGTGTCGTTGATGGCGTGGG
HNND	AATAGAAAAGGGCCAAAGAGAAAAAATAAAAAAATAAAAAAAGGGAAGGTGTCGTTGATGGCGTGGG
CC	CAGCCAACACAGGCTGGCTCCTTATCATTTAACTCCGTGTTTCAGTGTTTCACTCTGTGCTTCAATT
R1461	CAGCCAACACAGGCTGGCTCCTTATCATTTAACTCCGTGTTTCAGTGTTTCACTCTGTGCTTCAATT
9930	CAGCCAACACAGGCTGGCTCCTTATCATTTAACTCCGTGTTTCAGTGTTTCACTCTGTGCTTCAATT
HN	CAGCCAACACAGGCTGGCTCCTTATCATTTAACTCCGTGTTTCAGTGTTTCACTCTGTGCTTCAATT
XSBN	CAGCCAACACAGGCTGGCTCCTTATCATTTAACTCCGTGTTTCAGTGTTTCACTCTGTGCTTCAATT
HNHY	CAGCCAACACAGGCTGGCTCCTTATCATTTAACTCCGTGTTTCAGTGTTTCACTCTGTGCTTCAATT
HNHP	CAGCCAACACAGGCTGGCTCCTTATCATTTAACTCCGTGTTTCAGTGTTTCACTCTGTGCTTCAATT
ZN	CAGCCAACACAGGCTGGCTCCTTATCATTTAACTCCGTGTTTCAGTGTTTCACTCTGTGCTTCAATT
NT	CAGCCAACACAGGCTGGCTCCTTATCATTTAACTCCGTGTTTCAGTGTTTCACTCTGTGCTTCAATT
HNND	CAGCCAACACAGGCTGGCTCCTTATCATTTAACTCCGTGTTTCAGTGTTTCACTCTGTGCTTCAATT