

Table S5 Gene action of QTLs identified by CIM in HSBCF₁ population across four environments

Trait ^a	QTL ^b	Env. ^c	Position ^d	D/A or					
				A ^e	D ^e	A+D ^e	2 D / (A+D) ^f	GA ^g	Population
FL	qFL-C02-2	2015Yc	78.11		0.18			OD	HSBCF ₁ MPH
	qFL-Chr05-3	2014Bg	11.21	-1.16					RIL
	qFL-Chr05-4	2014Bg	40.01	-0.32					RIL
	qFL-C05-2	2015Bg	43.21			0.04		A	HSBCF ₁
	qFL-Chr05-1	2014Yc	45.31	-0.37					RIL
	qFL-C05-3	2015Bg	50.81			0.24	6.21	OD	HSBCF ₁
		2014Yc	51.61			1.25	1.21		HSBCF ₁
		2014Yc	50.81		0.76				HSBCF ₁ MPH
	qFL-Chr05-2	2014Yc	52.11	-0.33					RIL
	qFL-Chr09-1	2014Bg	3.81	-0.30					RIL
	qFL-C09-1	2014Bg	12.91			-0.17		A	HSBCF ₁
	qFL-C09-3	2015Bg	46.51			0.86		A	HSBCF ₁
	qFL-Chr10-1	2014Yc	44.51	0.29					RIL
	qFL-C11-1	2015Yc	4.81			-1.73		A	HSBCF ₁
		2014Bg	44.51	0.26					RIL
	qFL-Chr12-1	2014Yc	16.21	-0.33					RIL
	qFL-C13-2	2015Yc	30.61			0.56		A	HSBCF ₁
	qFL-Chr14-1	2014Yc	14.21	-0.50					RIL
	qFL-Chr14-2	2014Yc	20.91	-0.41					RIL
	qFL-C14-4	2015Bg	23.21			-0.83		A	HSBCF ₁
FU	qFU-Chr01-1	2014Yc	23.31	-0.46					RIL
		2014Bg	25.71	-0.27					RIL
	qFL-Chr15-1	2014Yc	13.11	-0.36					RIL
	qFL-C16-1	2014Yc	6.21			0.84		A	HSBCF ₁
		2014Bg	12.31	-0.38					RIL
	qFL-C17-1	2014Bg	31.31			-0.08		A	HSBCF ₁
	qFL-C18-1	2014Bg	59.01		-2.68			OD	HSBCF ₁ MPH
	qFL-C19-1	2014Yc	16.01			0.51	3.13	OD	HSBCF ₁
		2014Yc	16.01		0.80				HSBCF ₁ MPH
	qFL-Chr19-1	2014Yc	22.51	0.46					RIL
FU	qFL-C19-3	2014Yc	26.51			-0.73		A	HSBCF ₁
	qFL-C20-4	2014Yc	42.11			-0.31		A	HSBCF ₁
		2015Bg	44.11			-0.45			HSBCF ₁
	qFL-C21-1	2014Bg	44.61		-0.76			OD	HSBCF ₁ MPH
		2015Yc	30.31			0.89		A	HSBCF ₁
	qFU-C01-2	2014Bg	41.71			0.50	29.41	OD	HSBCF ₁
		2015Bg	45.71			-4.99	2.97		HSBCF ₁
		2015Bg	45.71		-7.40				HSBCF ₁ MPH
FU	qFU-C01-3	2015Bg	50.41			-1.09		A	HSBCF ₁
	qFU-C02-1	2015Bg	44.81			-4.44	2.72	OD	HSBCF ₁

		2015Bg	44.81	-6.04			HSBCF _i MPH	
qFU-Chr05-1	2014Bg	45.31	-0.22				RIL	
qFU-C06-2	2015Bg	35.01		0.29	6.15	OD	HSBCF _i	
	2015Yc	36.01	-0.90				HSBCF _i MPH	
qFU-C06-3	2015Bg	47.21		0.19		A	HSBCF _i	
qFU-Chr09-1	2014Yc	3.81	-0.24				RIL	
	2014Bg	3.81	-0.29				RIL	
			27.7					
qFU-C09-1	2014Bg	17.81	0	-113.49		OD	HSBCF _i MPH	
qFU-Chr09-5	2014Bg	18.11	-0.24				RIL	
qFU-C09-2	2015Bg	25.01		9.04		A	HSBCF _i	
qFU-Chr09-2	2014Yc	47.11	-0.24				RIL	
	2014Bg	47.11	-0.24				RIL	
qFU-C09-4	2014Yc	49.61		1.35		A	HSBCF _i	
	2015Bg	53.01		9.05			HSBCF _i	
qFU-Chr09-3	2014Yc	52.61	0.30				RIL	
	2014Bg	52.61	0.29				RIL	
qFU-Chr09-4	2014Yc	61.41	-0.20				RIL	
qFU-C13-1	2014Bg	32.41		0.31		A	HSBCF _i	
qFU-C14-2	2015Yc	37.41		0.16		A	HSBCF _i	
qFU-C16-1	2015Bg	72.91		-0.04	233.55	OD	HSBCF _i	
	2015Bg	72.91	4.93				HSBCF _i MPH	
qFU-C17-1	2014Bg	27.91		-0.19		A	HSBCF _i	
qFU-C18-1	2015Bg	57.11		-9.44		A	HSBCF _i	
qFU-Chr19-1	2014Yc	20.81	-0.22				RIL	
qFU-C19-1	2014Bg	37.41		1.04		A	HSBCF _i	
qFU-C20-2	2015Bg	38.01		-0.09	107.50	OD	HSBCF _i	
	2015Bg	38.01	4.69				HSBCF _i MPH	
qFU-C20-3	2015Yc	49.11	-0.70			OD	HSBCF _i MPH	
qFU-C20-4	2015Bg	57.81		-3.54	3.04	OD	HSBCF _i	
	2015Bg	57.81	-5.39				HSBCF _i MPH	
qFU-C22-1	2015Yc	20.81		-0.72		A	HSBCF _i	
qFU-C26-1	2015Yc	2.51	0.39			OD	HSBCF _i MPH	
qFU-C26-2	2014Yc	38.31		0.32		A	HSBCF _i	
MIC	qMIC-Chr01-1	2014Yc	14.51	0.24			RIL	
	qMIC-C02-1	2015Yc	32.21		0.97	A	HSBCF _i	
	qMIC-Chr05-1	2014Yc	12.61	-0.25			RIL	
	qMIC-C05-1	2014Bg	48.31		0.53	A	HSBCF _i	
	qMIC-Chr07-1	2014Bg	59.31	0.10			RIL	
	qMIC-C08-1	2015Bg	35.91	0.68		OD	HSBCF _i MPH	
	qMIC-C09-1	2015Yc	18.11		0.24	A	HSBCF _i	
	qMIC-C11-1	2015Yc	17.11		0.41	0.03	PD	HSBCF _i
		2015Yc	18.11	0.01			HSBCF _i MPH	
	qMIC-Chr10-1	2014Yc	62.61	0.16			RIL	

	qMIC-C13-3	2014Yc	44.11	0.08	A	HSBCF _i
	qMIC-C14-1	2015Yc	4.01	0.74	A	HSBCF _i
	qMIC-Chr14-1	2014Yc	23.31	0.08		RIL
		2014Bg	20.91	0.11		RIL
	qMIC-C14-2	2014Yc	27.21	0.21	A	HSBCF _i
		2015Bg	30.91	0.01		HSBCF _i
		2015Yc	32.71	0.30		HSBCF _i
	qMIC-C14-3	2015Yc	43.81	-0.01	A	HSBCF _i
	qMIC-C15-1	2015Bg	20.51	0.27	PD	HSBCF _i
		2015Yc	20.51	0.07	0.50	HSBCF _i MPH
		2015Bg	20.51	0.07	0.50	HSBCF _i MPH
	qMIC-Chr16-1	2014Yc	51.01	0.08		RIL
		2014Bg	49.31	0.09		RIL
	qMIC-Chr16-2	2014Yc	57.01	0.09		RIL
		2014Bg	57.41	0.15		RIL
	qMIC-Chr17-1	2014Bg	44.81	0.12		RIL
	qMIC-C19-1	2014Yc	26.21	-0.06	A	HSBCF _i
	qMIC-Chr24-1	2014Bg	16.81	-0.11		RIL
	qMIC-C24-4	2015Yc	73.31	0.51	A	HSBCF _i
	qMIC-Chr24-2	2014Bg	73.31	0.44		RIL
FE	qFE-C01-2	2015Bg	15.91	0.19	OD	HSBCF _i MPH
	qFE-C01-3	2014Bg	25.71	-0.38	OD	HSBCF _i MPH
	qFE-C09-3	2015Bg	50.61	0.09	OD	HSBCF _i MPH
	qFE-Chr11-1	2014Bg	5.31	1.16		RIL
	qFE-Chr14-2	2014Bg	1.11	1.09		RIL
	qFE-C14-1	2014Yc	5.31	1.75	A	HSBCF _i
	qFE-Chr14-3	2014Bg	6.31	0.25		RIL
		2014Bg	15.71	0.26		RIL
	qFE-Chr14-1	2014Yc	16.81	0.30		RIL
	qFE-C14-4	2015Yc	20.51	0.01	A	HSBCF _i
	qFE-C15-1	2015Yc	27.81	-0.05	A	HSBCF _i
	qFE-Chr16-1	2014Bg	1.11	1.17		RIL
	qFE-C17-1	2014Yc	33.41	-0.17	A	HSBCF _i
	qFE-Chr17-1	2014Bg	42.31	0.18		RIL
	qFE-C19-1	2015Yc	52.21	0.08	A	HSBCF _i
	qFE-Chr18-1	2014Bg	57.51	0.76		RIL
	qFE-C20-1	2014Yc	29.71	0.18	A	HSBCF _i
		2014Bg	29.71	0.01		HSBCF _i
	qFE-Chr20-1	2014Yc	41.51	0.20		RIL
		2014Bg	47.11	0.44		RIL
	qFE-C21-2	2014Yc	20.11	-1.83	A	HSBCF _i
	qFE-C24-2	2015Bg	40.11	0.15	OD	HSBCF _i MPH
	qFE-Chr24-1	2014Yc	73.31	0.75		RIL
		2014Bg	73.31	0.62		RIL

FS	qFS-Chr05-1	2014Bg	54.81	0.52		RIL
	qFS-C08-1	2014Bg	45.01		-0.05	A HSBCF ₁
	qFS-C09-1	2015Bg	38.71		-0.78	A HSBCF ₁
	qFS-C09-2	2015Yc	50.61	-2.05		OD HSBCF ₁ MPH
	qFS-C13-1	2014Yc	19.41		-0.27	A HSBCF ₁
	qFS-C13-3	2015Yc	39.21	-2.12		OD HSBCF ₁ MPH
	qFS-Chr14-1	2014Yc	6.71	-0.21		RIL
	qFS-Chr14-2	2014Yc	14.21	-0.28		RIL
	qFS-Chr14-3	2014Yc	21.61	-0.26		RIL
	qFS-Chr14-4	2014Yc	40.11	-0.20		RIL
	qFS-C14-1	2014Yc	50.51		-3.12	A HSBCF ₁
	qFS-C15-1	2014Yc	15.61		-0.01	A HSBCF ₁
	qFS-Chr19-1	2014Bg	22.51	0.51		RIL
	qFS-Chr19-2	2014Bg	27.81	0.61		RIL
	qFS-C19-3	2015Yc	57.11		-2.76	A HSBCF ₁
	qFS-C20-1	2014Yc	39.01		-0.82	A HSBCF ₁
		2015Bg	39.01		-1.10	HSBCF ₁
	qFS-Chr20-1	2014Yc	42.11	-0.50		RIL
	qFS-Chr20-2	2014Yc	59.61	-0.48		RIL
	qFS-C25-2	2015Yc	39.51		-3.91	A HSBCF ₁
		2015Bg	39.81		-3.67	HSBCF ₁

^a FL: fiber length; FU: fiber uniformity; MIC: micronaire; FE: fiber elongation; FS: fiber strength

^b QTLs in bold are those identified by CIM in RILs in our previous study (Li et al. 2016), which was just used to estimate the gene action of HSBCF₁ population

^c 2014Yc: Yacheng of Hainan Province in 2014; 2014Bg: Baogang of Hainan Province in 2014; 2015Yc: Yacheng of Hainan Province in 2015; 2015Bg: Baogang of Hainan Province in 2015

^d Position of QTL located on chromosome: as cM distance from the top of each chromosome

^e The genetic expectation of a QTL effect obtained is the additive effect (A) from the RILs, the additive and dominance effects (A+D) from the BCF₁s, and the dominance effect (D) from the MPH values

^f |D/A|: |dominance/additive|; 2|D|/(A+D)|: 2| dominance |/(additive + dominance)|

^g GA: gene action; PD/D partial dominance (|d/a| <= 1 or 2|d|/(a+d) | <= 1); OD overdominance(|d/a| >1 or 2|d|/(a+d) | > 1), here, 2|d|/(a+d) | > 1 same to 2|d| > |a + d|; A: when QTL detected only in BCF₁ or both BCF₁ and RIL was referred to as additive (A).