

**Table S3 Detailed sequence pile-up (Li et al. 2009) for *scalloped* binding site regulating *cut* (Halder & Carrol 2001, Guss et al. 2001).**

Chrom.	Pos.	Ref.	# reads (SAM)	Read bases (SAM)	Base quality (SAM)	# reads (ORE)	Read bases (ORE)	Base quality (ORE)
X	7424817	T	15	,.....	HFIHIDIBI<GIII	15	.\$.....	DIIFIHIIHIII
X	7424818	G	15	,.....	HIHGIEIGIGGIIIG	15	,.....^],	GIIIIHGIIHIII
X	7424819	T	16	,\$.....^].	EHGGICHGIGDIIHGE	15	,\$.....	EIIIIIIIIII
X	7424820	C	15	.....	IIII4IBIBGIIIGI	16	.....^],^],	IGGIIIIIIIBIIEE
X	7424821	A	15	.....G.	IHIICIBIGGII*I	18	.....^],^],	IIIIIIIBIIIIIDBB
X	7424822	A	15	.....	FHHIEIGIDGIIGGI	18	.....	IIIIIIHIIIIHIII
X	7424823	T	15	.....	HIIDIGICGIIIGI	18	.....	IIIIIIHHI@IIIGII
X	7424824	G	15	.....	GIIIDI@IEGIGIGI	18	.....	IIIIIIIDIIIIHIIIG
X	7424825	T	15	,\$.....	EGGG=HDCGIIIGI	18	.....	IIIIIIIIIGIIEHII
X	7424826	A	14	.....	IHIBIGIDGGIIGI	18	.....	IIIIIIIIIGIIII
X	7424827	A	14	.....	HII:I4IDGIIIGI	19	.....^],	IIIIIIIIIIHIG>
X	7424828	T	14	.....	III=IEIEGIIII	22	.....^],^],^],	IHHIIIIIIIIHIEBBB
X	7424829	T	14	.....	III?IBICDIIII	23	.....^],	IIHIIIFIIHIIIFIHICEEED
X	7424830	C	14	.....	IIH?IDIDDIIII	23	.....	IIIIIIIIHIGIIHBFHF
X	7424831	G	14	,\$.....	EDIDIGICGIIII	24	.....^],	IIIIIIHIIIIIGBIHIE
X	7424832	A	14	.....^].	>>DFGI>EIIII7	25	.....^].	IIHIIIIIGIIIIHGHIGI>
X	7424833	A	14	,\$.....	;<IGIGHIIII:	25	.....	IIIIIIIIHIIIIIDI>II?
X	7424834	A	13	,\$.....	9?IBIGHIIII5	25	.....	IIIIIIIIIIIEIH>IIHIIA
X	7424835	A	12	.....	BIGIDHIIII5	25	.....	IIIIIIIIIIIIIGIHHIB
X	7424836	A	12	.....	DIGIDHIIII2	25	.....	IIIGIIIIIIHIIIEIIGIHHIE
X	7424837	T	12	.....	@IBGBEFBIII-	26	.....^].	IIIIIIIIII8IHHHI@ADHDEDIE
X	7424838	G	12	.....	GIGHDDIIII:	26	.....	HIIIIIIIIIIIIIGHHIIII
X	7424839	T	12	.....	GH@IGGHIFHI4	26	.....	HIIIIIIHIIIGIIII3IHHIHI
X	7424840	C	12	.....	GIGI<GIIII5	26	.....	IIIIIIIIIIHIG8IHDHII
X	7424841	G	13	.....^].	:IGIGGIIGII7E	26	.....	IIIIIIIIIIIIIGIHHII