

**Table S4 Detailed sequence pile-up (Li et al. 2009) for *scalloped* binding site regulating *sal* (Guss et al. 2001).**

Chrom.	Pos.	Ref.	# reads (SAM)	Read bases (SAM)	Base quality (SAM)	# reads (ORE)	Read bases (ORE)	Base quality (ORE)
2L	11454659	G	17	,,.....,	II+IIIIIGIIIIHII	17	,,.....,	77977<97777787777
2L	11454660	G	18	,,.....,^],	II3IIIIIGIIIIIEIIE	17	,,.....,	34644964444454444
2L	11454661	A	18	,,.....,	IH6@HIHIBIIIIIIHH	17	ggGGggGgGggGGgggg	)644964444454444
2L	11454662	C	18	,\$,.....,	EI8IIIIIBHIIIIIII	18	,\$,.....,^>],	)HFF?IIIIIIIIIC
2L	11454663	A	17	,,.....,	I38IIIIIGGHIIIIHII	17	,\$,.....,	)HFFDIHIIIIIIH8
2L	11454664	T	17	,,.....,	IBGIIIIIGIIIIHHIII	16	,\$,\$,\$,.....,	>>>DIGIIIIHIGIIC
2L	11454665	T	18	,,.....,^],	IAEIIIIIGIIIIIIHIE	13	,\$,.....,	BIIIIIIIIII?
2L	11454666	C	18	,\$,.....,	EAEIIIIIGIHHIIGIIIIH	12	,,.....,	5GIIIIIIHI@
2L	11454667	G	17	,,.....,	A@IIIIIGIIIIIGIIII	13	,,.....,^],	-8888988888?!
2L	11454668	T	17	,\$,\$,.....,	?BIIIIIDIIII@IFI	13	CcCccCCcccccc	!44445444442!
2L	11454669	G	15	,,.....,	IIIGIIIIIGIHI	13	,,.....,	I3333433333<!
2L	11454670	G	15	,,.....,	IIIBIIIIHGIIII	13	A\$aAaaAAaaaaaa	!1111211111;!
2L	11454671	G	15	,,.....,	IIIEIIIIIGIIII	13	,,.....,^>.	7777877777@0E
2L	11454672	A	15	,,.....,	IIIGIIIIIDIIIIH	14	,,.....,^],	IGIIIIII46HE
2L	11454673	T	16	,\$,\$,.....,^],	>>IIGHIIIIIGIIIFB	15	,,.....,^],	GIIIIIIHII;>HIB
2L	11454674	T	14	,,.....,	IIGIIIIHGIIHIE	15	,,.....,	HIIIIIEII,<HIE
2L	11454675	C	14	,\$,.....,	>E@IIIIIGIIIFG	15	,,.....,a,...	EIIIIIIII,IGII
2L	11454676	C	13	,\$,.....,	BGIIIIIIIIII	15	,\$,.....,	BIIIIIGII;IHII