

TABLE S2. Summary of information used to validate the enJSRV insertion scan procedure.

enJSRV	amorces		Position OAR v4.0	Detected by Whole Genome Survey
	Forward	Reverse		
enJSRV-6	ccagttccagaaggaaggag	caggggaataactggtgctacct	1:239929928-239926832	Y
enJSRV-7	tgtcacacgtggtgggagtc	actcgagaggaagcagagggtc	(multiple locus amplified)	N
enJSRV-8	tcagtggatcaatggtctgca	tgggtgagcatgcacacag	3:139105810-139106319	Y
enJSRV-15	gtgggaagaattccttggtacac	ccataaggggtgtatcgtctgca	(Sequence not detected)	N
enJSRV-16	tgctcagttcccaggtccca	gtgcaagagctagagctggaagg	UnplacedScaffold_004083582.1:13920-14138	Y
enJSRV-18	gggaagattcgtttcttaggcgctc	atgaaccggactccatggcag	11:32040949-32041591	Y
enJSF16	ggataagctacactataaaaccaag	ccatatgtaggattgggggtg	15:42218266-42211217	Y