Bootstrap 3.0 – Collapse – JSFiddle

(bp)	Quality	Impacted Gene	Product Name	Gene Length (aa)	Distance from Start Codon (aa)	Reference Base	AH067 Alternate Allele	ES021 Alternate Allele	AH068 Alternate Allele	ES022 Alternate Allele	Wildtype Base
101497	158.87	LdBPK_020190.1	Phosphoglycan Beta 1,2 Arabinosyltransferase	780	426	T	TGG	TGG	Ref	Ref	Ref
125070	101.9	LdBPK_030370.1	Hypothetical Protein, Conserved	359	163	GATAT	G	G	Ref	Ref	Ref
125228	133.75	LdBPK_030370.1	Hypothetical Protein, Conserved	359	216	TCC	Ref, T	Ref, T	Ref	Ref	Ref
268606	909.73	LdBPK_030670.1	Hypothetical Protein, Conserved	2209	2209	ATCT	A	A	Ref	Ref	Ref
156665	553.84	LdBPK_050490.1	Hypothetical Protein, Conserved	271	19	TCC	Ref	Ref	T	T	Ref
225080	2641.77	LdBPK_100521.1	Hypothetical Protein, Unknown Function	77	77	T	Ref	Ref	C	C	Ref
436547	109.9	LdBPK_101050.1	Histone H3	155	22	GAC	G	G	Ref	Ref	Ref
424027	298.74	LdBPK_181030.1	Hypothetical Repeat Protein	153	76	CTG	C	C	Ref	Ref	Ref
295122	43.73	LdBPK_220640.1	Hypothetical Protein, Conserved	977	165	A	Ref, AT	Ref, AT	Ref	Ref	Ref
295124	43.73	LdBPK_220640.1	Hypothetical Protein, Conserved	977	165	CA	C, Ref	C, Ref	Ref	Ref	Ref
378794	181.8	LdBPK_230920.1	Hypothetical Protein, Conserved	138	73	A	AGG	AGG	Ref	Ref	Ref
145586	39.7	LdBPK_240430.1	Hypothetical Protein, Unknown Function	89	61	A	ACC	ACC	Ref	Ref	Ref
102608	1514.77	LdBPK_250350.1	Hypothetical Protein, Conserved	141	0	A	C	C	Ref	Ref	Ref
579030	360.77	LdBPK_301640.1	Hypothetical Protein, Conserved	696	333	C	Ref	Ref	Ref, T	Ref, T	Ref
602800	3120.77	LdBPK_311390.1	Hypothetical Protein, Conserved	1486	0	C	Ref	Ref	A	A	Ref
1000447	509.73	LdBPK_312060.1	Succinyl-Diaminopimelate Desuccinylase-Like Protein	471	129	A	Ref, AG	Ref, AG	Ref	Ref	Ref
9103	754.75	LdBPK_320030.1	Hypothetical Protein, Unknown Function	367	266	C	CG	CG	Ref	Ref	Ref
174033	61.77	LdBPK_320480.1	Hypothetical Protein, Conserved	939	239	C	Ref, T	Ref, T	Ref	Ref	Ref
279828	171.76	LdBPK_320840.1	Map Kinase Pseudogene	130	100	AGAGG	Ref	Ref	A	A	Ref
729169	60.73	LdBPK_341700.1	Amastin-Like Surface Protein, Putative	272	71	CA	C, Ref	C, Ref	Ref	Ref	Ref
288971	190.8	LdBPK_350660.1	Hypothetical Protein, Unknown Function	123	110	TCC	T	T	Ref	Ref	Ref