

SOBA

Feature Count Table

Source	Type	Number	Attributes (Tag: Number)
.	contig	28	ID: 28 Name: 28
augustus_masked	match	9621	ID: 9621 Name: 9621
	match_part	25829	ID: 25829 Target: 25829 Parent: 25829
blastn	expressed_sequence_match	12983	ID: 12983 Name: 12983
	match_part	21869	ID: 21869 Target: 21869 Parent: 21869
blastx	match_part	38106	ID: 38106 Target: 38106 Parent: 38106
	protein_match	20579	ID: 20579 Name: 20579
est2genome	expressed_sequence_match	12942	ID: 12942 Name: 12942
	match_part	22286	ID: 22286 Target: 22286 Parent: 22286
genemark	match	10280	ID: 10280 Name: 10280
	match_part	27729	ID: 27729 Target: 27729 Parent: 27729
maker	CDS	11201	ID: 11201 Parent: 11201
	exon	11272	ID: 11272 Parent: 11272
	five_prime_UTR	1113	ID: 1113 Parent: 1113
	gene	3697	ID: 3697 Name: 3697
	mRNA	3697	_QI: 3697 ID: 3697 _eAED: 3697 _AED: 3697 Name: 3697 Parent: 3697
	three_prime_UTR	876	ID: 876 Parent: 876
protein2genome	match_part	35281	ID: 35281 Target: 35281 Parent: 35281
	protein_match	14961	ID: 14961 Name: 14961
repeatmasker	match	28383	ID: 28383 Target: 28383 Name: 28383
	match_part	28383	ID: 28383 Target: 28383 Parent: 28383
repeatrunner	match_part	26	ID: 26 Target: 26 Parent: 26
	protein_match	26	ID: 26 Target: 26 Name: 26
snap_masked	match	8829	ID: 8829 Name: 8829
	match_part	21500	ID: 21500 Target: 21500 Parent: 21500

Feature Lengths Table

Source	Type	min	max	median	mean	Footprint(%)
.	contig	1065	6064072	1252084.75	4548.5	100
augustus_masked	match	204	15347	1719.86	1449	47.198
	match_part	3	14757	575.91	309	42.43
blastn	expressed_sequence_match	44	10181	522.22	539	9.751
	match_part	31	1063	272.78	229	8.824
blastx	match_part	39	13128	704.05	447	11.281
	protein_match	66	13128	1371.81	1186	11.946
est2genome	expressed_sequence_match	95	10181	542.12	558	9.97
	match_part	7	1061	274.76	233	9.049
genemark	match	12	47771	1843.94	1426	54.069