Main: Iterating through fileSet...

Adding Calls from file /home/zhou4/PhATE/Code/PhATE_pipeline/PipelineOutput/Ec_P2/genemark.cgc

Adding Calls from file /home/zhou4/PhATE/Code/PhATE_pipeline/PipelineOutput/Ec_P2/glimmer.cgc

Adding Calls from file /home/zhou4/PhATE/Code/PhATE_pipeline/PipelineOutput/Ec_P2/glimmer.cgc

Adding Calls from file /home/zhou4/PhATE/Code/PhATE_pipeline/PipelineOutput/Ec_P2/prodigal.cgc

Adding Calls from file /home/zhou4/PhATE/Code/PhATE_pipeline/PipelineOutput/Ec_P2/phanotate.cgc

Main: CallerList is genemarks , glimmer , prodigal , phanotate ,

Main: Conting gene calls for each caller...

Main: Comparing accross the call sets...

The following gene callers were considered: , genemarks , glimmer , prodigal , phanotate

The number of distinct gene calls over all gene callers is 62

The number of gene calls in common among all callers is 29

The number of genemarks produced 41 gene calls.

Caller genemarks produced 41 gene calls.

Caller genemarks pene-call length stats: min: 141 , max: 2448 , ave: 747

Caller glimmer produced 42 gene calls.

Caller prodigal gene-call length stats: min: 114 , max: 2448 , ave: 729

Caller prodigal gene-call length stats: min: 114 , max: 2448 , ave: 735

Caller phanotate produced 49 gene calls.

Caller phanotate gene-call length stats: min: 141 , max: 2448 , ave: 653

count caller strand leftEnd rightEnd length conting caller strand leftEnd rightEnd 1

1 genemarks + 1 141 141 E_P2_NC_001895

2 genemarks - 187 1221 1035 E_P2_NC_001895

3 genemarks - 1221 2993 173 E_P2_NC_001895

4 glimmer + 3101 4021 921 to 100 glimmer - 1221 3023 1803 to 100 gli

Caller	phanotate produc				112 ,		, ,.	,,,																	
	phanotate gene-c																								
count	caller strand	leftEnd					caller		leftEnd	rightEn	nd	length	contig	caller strand	leftEnd			length	contig cal	ler str	and lef	tEnd righ	tEnd	len	gth contig
1	genemarks	+	1	141	141		NC_001895							prodigal	+	1		141	unknown						
2	genemarks	-	187	1221	1035		NC_001895			187	1221	1035	unknown	prodigal	-	187		1035	unknown phar		-	187	1221	1035	Ec_P2_NC_001895
3	genemarks	-	1221	2993	1773		NC_001895		2022	1002				prodigal	-	1221	2993	1773	unknown phar	notate	-	1221	2993	1773	Ec_P2_NC_001895
5						glimme		1221 3101	3023 4021	1803 921	unknown unknown						phanotat	e +	3101	4021	921	E . D . N	C 00100E		
6	genemarks	+	3167	4021	855	glimme	r + NC 001895		4021	921	unknown			prodigal	+	3167		.e + 855	unknown	4021	921	EC_PZ_IV	IC_001895)	
7	genemarks	+	4080	5153	1074		NC_001895 NC_001895			4080	5153	1074	unknown	prodigat	+	4080		1074	unknown phar	201210	+	4080	5153	1074	Ec P2 NC 001895
8	genellarks	т	4000	3133	1074	alimme		5157	5900	744		prodiga		+ 5157	5900	744	unknown			5157	5900	744		IC 00189	
9	genemarks	+	5181	5900	720		NC 001895		3300	,	unknown	proutgu		. 5157	3300	,	unknown	phanoca		3137	3300	,	20_12_1	·c_00105.	,
10	genemarks		3101	3300	, 20	qlimme		5940	6509	570	unknown														
11	genemarks	+	6000	6509	510		NC 001895							prodigal	+	6000	6509	510	unknown phar	notate	+	6000	6509	510	Ec P2 NC 001895
12	genemarks	+	6509	6712	204		NC 001895		+	6509	6712	204	unknown	prodigal	+	6509		204	unknown phar		+	6509	6712	204	Ec P2 NC 001895
13	genemarks	+	6716	6997	282		NC_001895			6716	6997	282		prodigal	+	6716	6997	282	unknown phar		+	6716	6997	282	Ec_P2_NC_001895
14	genemarks	+	6997	7494	498	Ec_P2_f	NC_001895	glimmer	+	6997	7494	498	unknown	prodigal	+	6997	7494	498	unknown phar	notate	+	6997	7494	498	Ec_P2_NC_001895
15	genemarks	+	7509	7934	426	Ec_P2_I	NC_001895		+	7509	7934	426		prodigal	+	7509		426	unknown phar	notate	+	7509	7934	426	Ec_P2_NC_001895
16						glimme			8347	426	unknown	prodiga	l	+ 7922	8347	426	unknown	phanota	ite +	7922	8347	426	Ec_P2_N	IC_00189	5
17	genemarks	+	7934	8347	414		NC_001895																		
18						glimme		8334	8492	159		prodiga		+ 8334	8492	159	unknown			8334	8492	159		IC_00189	
19	genemarks	+	8455	8922	468		NC_001895			8455	8922	468		prodigal	+	8455		468	unknown phar		+	8455	8922	468	Ec_P2_NC_001895
20	genemarks	+	8915	9367	453		NC_001895			8915	9367	453	unknown	prodigal	+	8915	9367	453	unknown phar	notate	+	8915	9367	453	Ec_P2_NC_001895
21			0.420	10224	706	glimme			10125	687	unknown					0.420	10224	706	unlena en mhaur			0.420	10224	706	En DO NC 00100E
22 23	genemarks	+	9439 10308	10224 10943	786 636		NC_001895 NC 001895			10308	10943	636		prodigal prodigal	+	9439 10308		786 636	unknown phar		+	9439 10308	10224 10943	786 636	Ec_P2_NC_001895 Ec P2 NC 001895
23	genemarks genemarks	+	10940	11287	348		NC_001895 NC_001895			10940	11287	348		prodigat	+	10940		348	unknown phar unknown phar		+	10940	11287	348	Ec_P2_NC_001895
25	genemarks	I	11292	12200	909		NC_001895 NC_001895			11292	12200	909		prodigat	+	11292		909	unknown phar		+	11292	12200	909	Ec P2 NC 001895
26	genemarks	+	12193	12723	531		NC 001895			12193	12723	531		prodigat	·	12193		531	unknown phar		+	12193	12723	531	Ec_P2_NC_001895
27	genemarks	+	12734	14743	2010		NC_001895			12734	14743	2010		prodigal	+	12734		2010	unknown phar		+	12734	14743	2010	Ec P2 NC 001895
28	genemarks	+	14747	15274	528		NC 001895			14747	15274	528		prodigal	+	14747		528	unknown phar		+	14747	15274	528	Ec P2 NC 001895
29	3							9						p			phanotat		15276		90		IC_001895		
30																	phanotat			15559	105		IC_001895		
31						glimme	r –	15474	15587	114	unknown												_		
32	genemarks	-	15675	17261	1587	Ec_P2_I	NC_001895	glimmer		15675	17261	1587	unknown	prodigal	-	15675	17261	1587	unknown phar	notate	-	15675	17261	1587	Ec_P2_NC_001895
33																	phanotat	:e -	17269	17430	162		IC_001895		
34																	phanotat	:e -	17402	17632	231	Ec_P2_N	IC_001895	5	
35	genemarks	+	17652	18842	1191		NC_001895			17652	18842	1191		prodigal	+	17652		1191	unknown phar		+	17652		1191	Ec_P2_NC_001895
36	genemarks	+	18855	19373	519		NC_001895			18855	19373	519		prodigal	+	18855		519	unknown phar		+	18855	19373	519	Ec_P2_NC_001895
37	genemarks	+	19430	19705	276		NC_001895			19430	19705	276	unknown	prodigal	+	19430	19705	276	unknown phar	notate	+	19430	19705	276	Ec_P2_NC_001895
38 39						glimme	r +	19702	19857	156	unknown								10720	10057	120	F - D2 N			
39 40	aonomarko	+	19850	22297	2448	Ec D2 I	NC 001895	alimmor		19850	22297	2448	unknown	prodigal	+	19850	phanotat 22297	2448	19738 unknown phar		120 +		IC_001895 22297	2448	Ec P2 NC 001895
41	genemarks genemarks	+	22312	22791	480		NC_001895 NC_001895			22312	22791	480		prodigat	+	22312		480	unknown phar		+		22791	480	Ec P2 NC 001895
42	genemarks	+	22791	23954	1164		NC 001895			22791	23954	1164		prodigat	+	22791		1164	unknown phar		+		23954	1164	Ec_P2_NC_001895
43	genemarks		22/31	23334	1104	20_12_1	vc_001033	germiner		22/31	23334	1104	unknown	proutgut		22/31	phanotat			24254	219		IC_001895		20_12_10_001033
44																	phanotat		24335		111		IC 001895		
45	genemarks	_	24462	25475	1014	Ec P2 I	NC 001895	glimmer		24462	25475	1014	unknown	prodigal	-	24462		1014	unknown phar		-	24462	25475	1014	Ec P2 NC 001895
46	genemarks	-	25591	25890	300	Ec_P2_f	NC_001895	5						prodigal	-	25591	25890	300	unknown phar	notate	-	25591	25890	300	Ec_P2_NC_001895
47	genemarks	+	26005	26280	276	Ec_P2_I	NC_001895	5						prodigal	+	26005	26280	276	unknown phar	notate	+	26005	26280	276	Ec_P2_NC_001895
48						glimme			26280	186	unknown														
49	genemarks	+	26291	26461	171		NC_001895			26291	26461	171		prodigal	+	26291	26461	171	unknown phar	notate	+	26291	26461	171	Ec_P2_NC_001895
50	genemarks	+	26452	26958	507	Ec_P2_I	NC_001895	glimmer	+	26452	26958	507	unknown						phanotate	+	26452	26958	507	Ec_P2_I	NC_001895
51												prodiga		+ 26458	26958	501	unknown								
52	genemarks	+	27022	27246	225		NC_001895			27022	27246	225		prodigal	+	27022		225	unknown phar		+	27022	27246	225	Ec_P2_NC_001895
53	genemarks	+	27246	27548	303		NC_001895			27246	27548	303		prodigal	+	27246		303	unknown phar		+	27246	27548	303	Ec_P2_NC_001895
54 55	genemarks	+	27548	27772 28044	225		NC_001895			27548	27772 28044	225 276		prodigal	+	27548		225 276	unknown phar		+	27548	27772 28044	225	Ec_P2_NC_001895
55 56	genemarks genemarks	+	27769 28034	30319	276 2286		NC_001895 NC_001895			27769 28034	28044 30319	276		prodigal prodigal	+	27769 28034		2286	unknown phar unknown	iotate	+	27769	20044	276	Ec_P2_NC_001895
50 57	geriellar KS	т	20034	20219	2200	LC_FZ_I	AC_001033	, g crimiler	*	20034	20219	2200	ulikilowii	pi ouigat	*	20034	phanotat			30319	2223	Fc P2 M	IC 001895	;	
58	genemarks	+	30316	30645	330	Fc P2 I	NC_001895	alimmer	. +	30316	30645	330	unknown	prodigal	+	30316		330	unknown phar		+	30316	30645	330	Ec_P2_NC_001895
59	genemarks	_	30684	31445	762		NC 001895			30684	31445	762		prodigat	_	30684		762	unknown phar		_	30684	31445	762	Ec_P2_NC_001895
60	genemarks	_	31620	33314	1695		NC_001895			31620	33314	1695	unknown	r		5000.			pilai			30007	515		
61	•							,				prodiga		- 31620	33380	1761	unknown	phanota	ite –	31620	33380	1761	Ec_P2_N	IC_00189	5
62																	phanotat	e -	33380	33593	214		IC_001895		
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