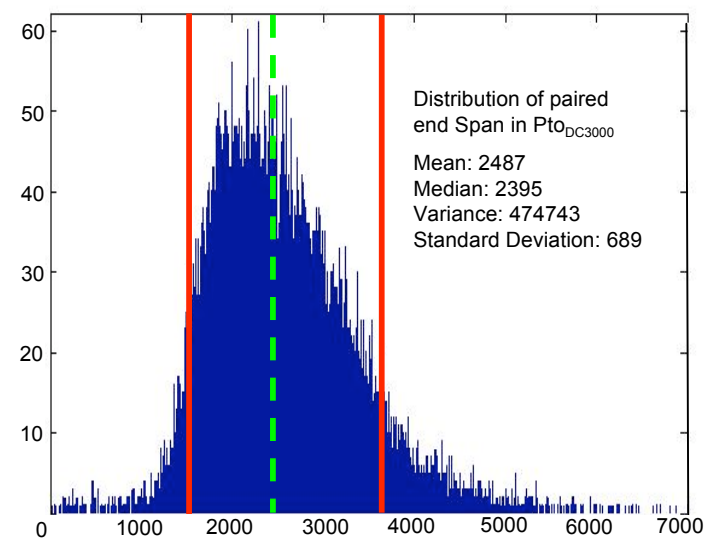
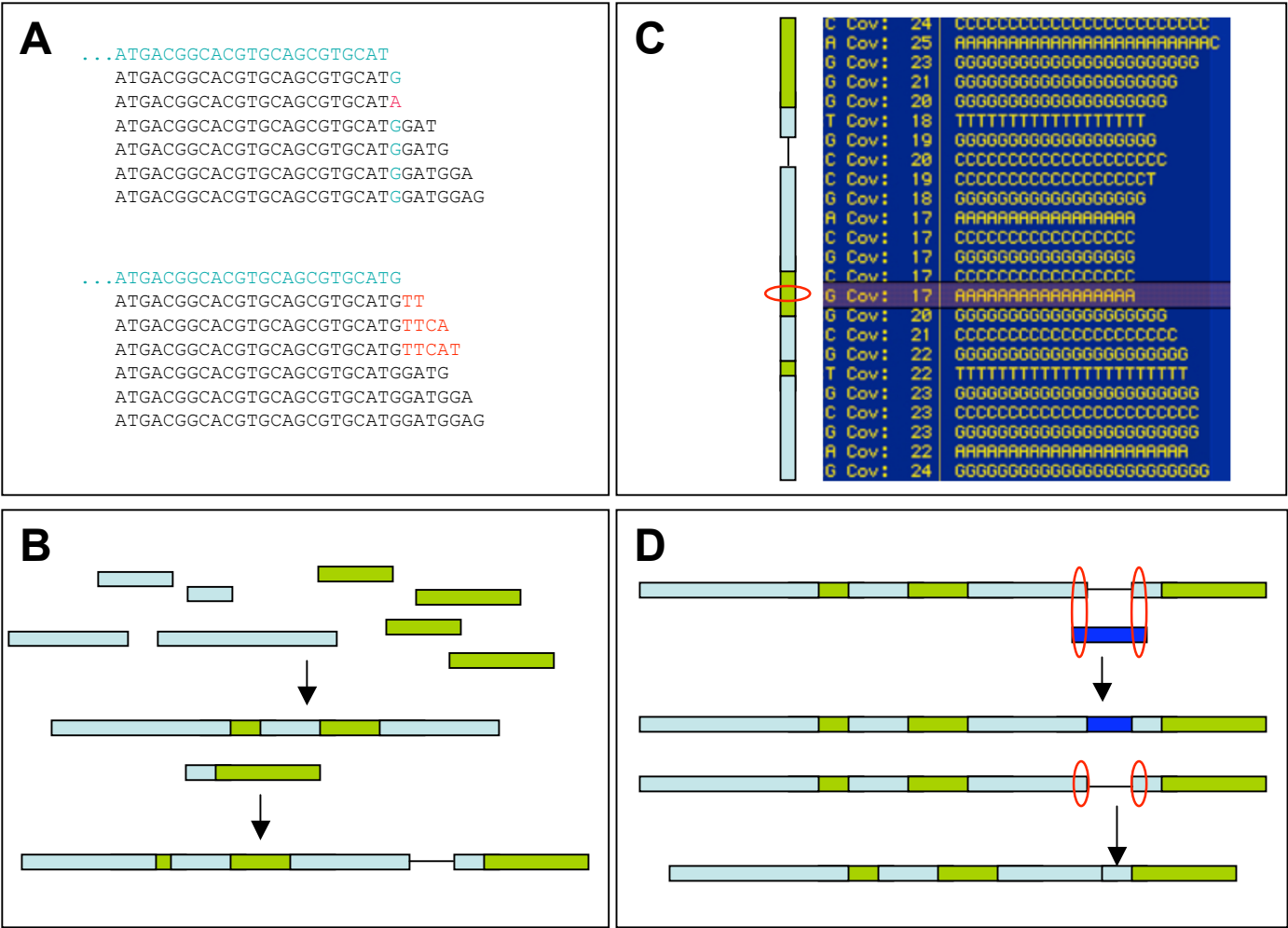
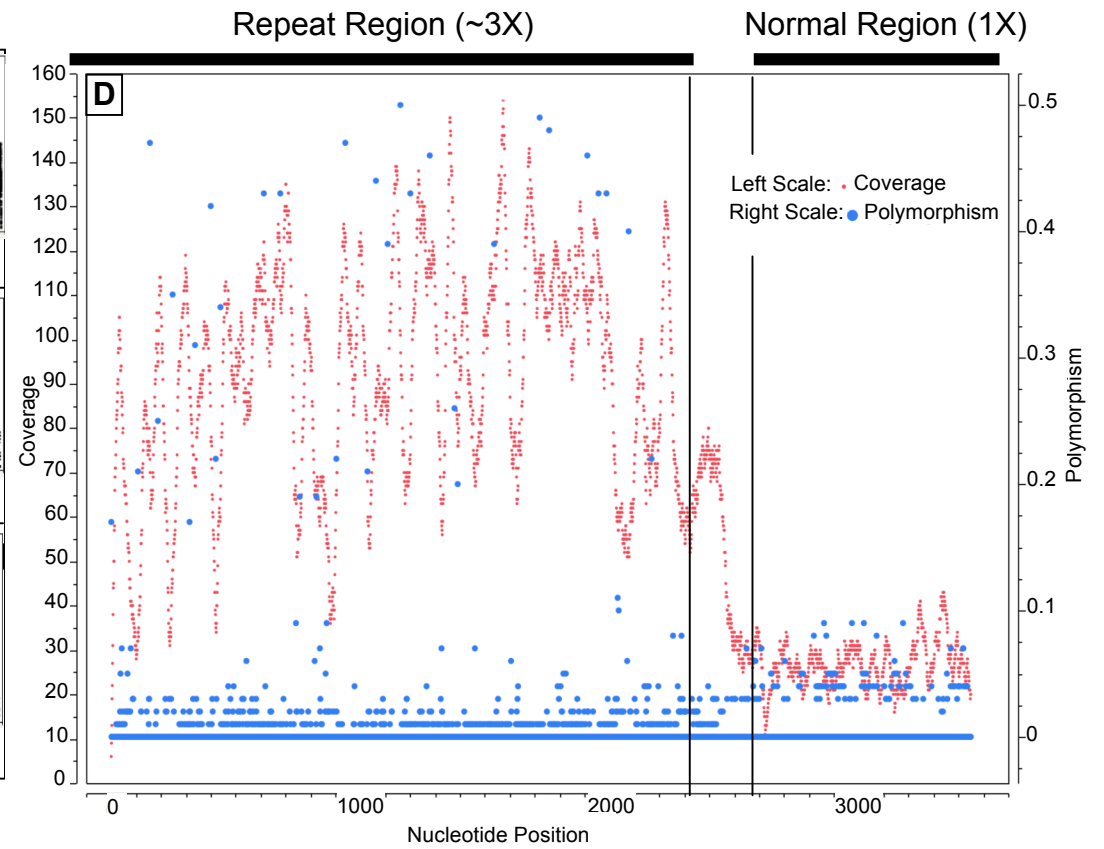
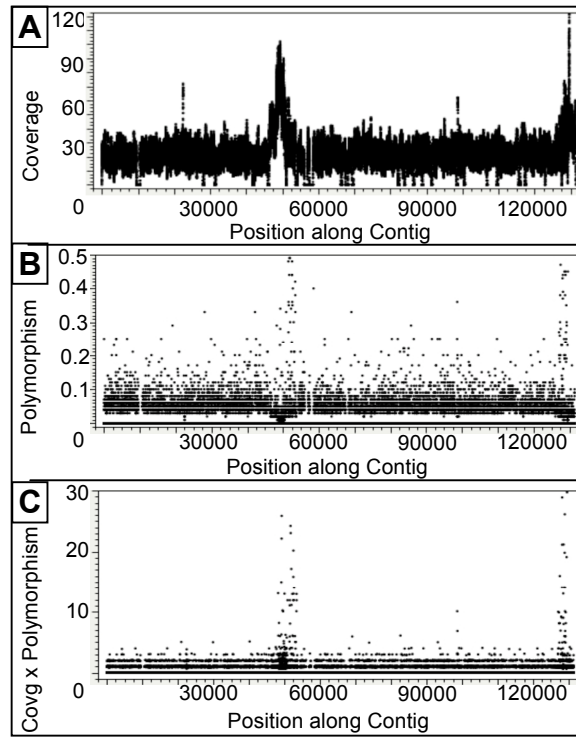
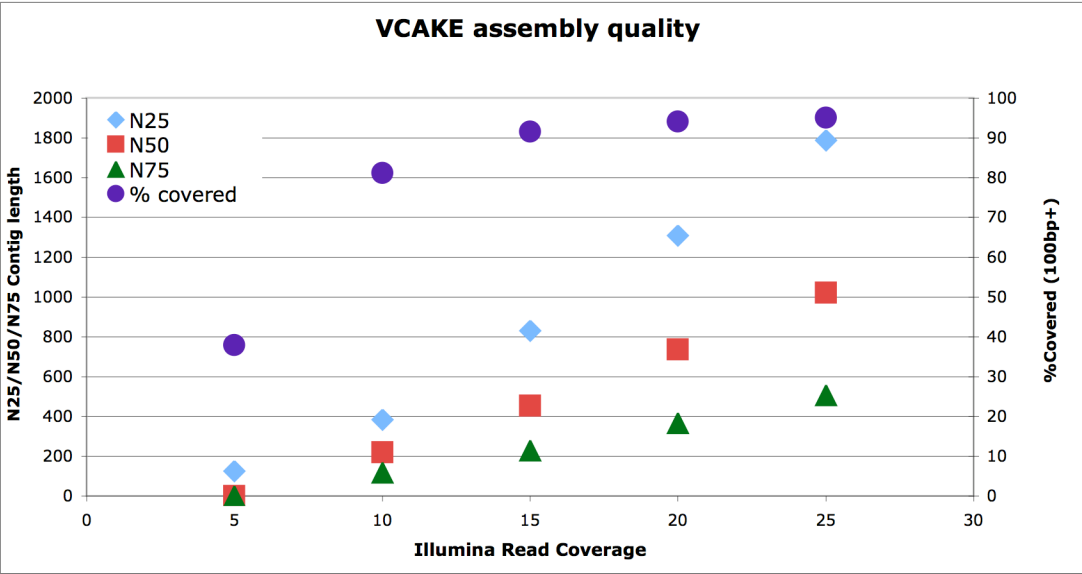




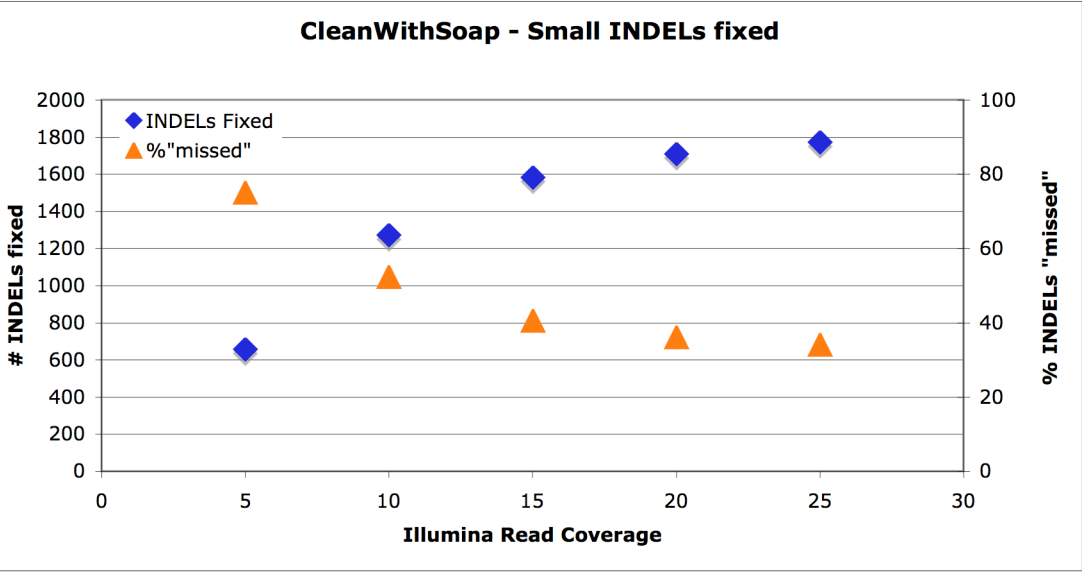


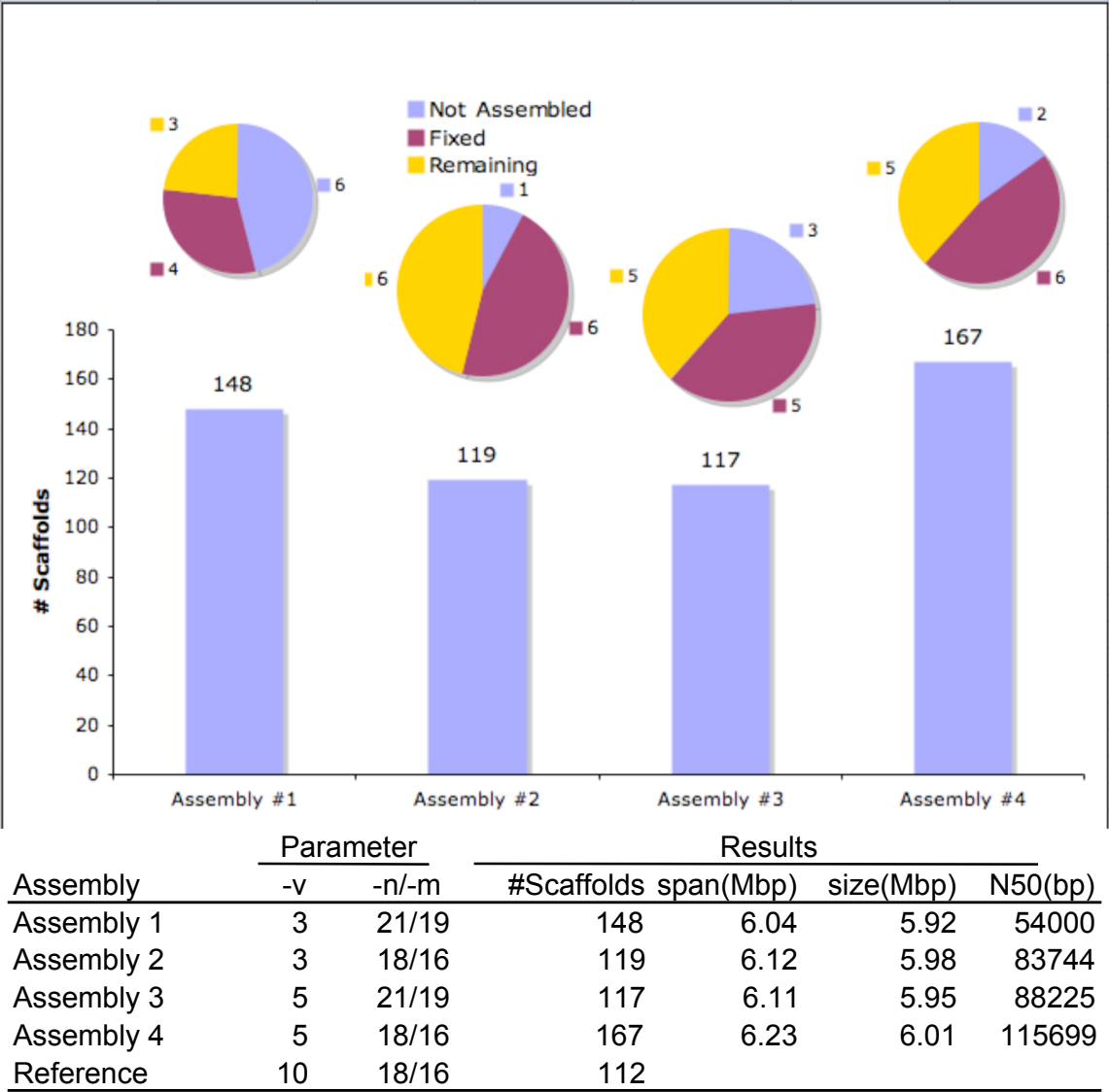


A



B





Supplemental Table 1: Verification of Core-genome deletions in *Por*_{1,6}

contig	observed amplicon	expected gap	observed gap	bases checked	SNPs	InDels	comments
Absent_PSPTO_0226	740	69	-17	690	0	0	17bp unbridged overlap
Absent_PSPTO_0444	800	0	0	750	0	0	
Absent_PSPTO_0824	500	0	0	447	0	0	
Absent_PSPTO_1945	966	378	509	404	3	1bp x1	
Absent_PSPTO_2015	798	22	341	748	0	0	not PCR amplified
Absent_PSPTO_2409	500	0	0	445	0	0	
Absent_PSPTO_2409_2	500	0	0	450	0	0	
Absent_PSPTO_3355	no product	760	nd	nd	nd	nd	
Absent_PSPTO_5010	500	0	0	449	0	0	
Absent_PSPTO_5010_2	500	0	0	450	0	0	
Absent_PSPTO_5113	500	0	0	446	0	0	
Absent_PSPTO_5458	500	0	0	450	0	0	
Absent_PSPTO_5458_2	1437	612	980	348	1	0	
Filled_16_1	391	0	0	343	0	0	
Filled_16_5	629	0	0	581	0	0	6bp unbridged overlap
Filled_6_1	490	0	0	442	0	0	
Filled_6_2	581	0	0	533	0	0	
Filled_9_2	314	0	0	266	0	0	
Gap_10_1	548	59	32	468	0	0	
Gap_10_2	540	203	209	255	0	0	
Gap_16_2	1027	22	510	454	0	0	
Gap_16_3	527	49	33	446	0	0	
Gap_16_6	484	134	27	408	0	0	
Gap_4_1	392	119	-6	345	0	0	
Gap_4_3	361	179	2	312	1	0	3bp unbridged overlap
Gap_4_5	576	47	-3	528	0	0	
Gap_6_3_3'	~1.5kb (gel)	259	282	477	0	0	
Gap_6_3_5'	~1.5kb (gel)	190	41	249	0	0	
Gap_9_1	845	322	313	484	0	0	25bp unbridged overlap
Gap_93_2	419	109	-25	371	0	0	
Gap_93_3	476	70	-9	428	0	0	
Gap_93_5	458	92	70	339	0	0	
				total bases	total SNP	total InDel	
				13806	5	1	
				error rate	0.0004346		
				Phred score	33.619166		

closest homolog	% identity	Scaffold	start	end	comments
avrB1_Pgy_race4	84	scaffold00041_length=53134_Fill	46798	47760	
avrB3_Psy_B728A	67.45	scaffold00005_length=172712_Fill	625	1608	
avrPto1_Pto_JL1065	41.1	scaffold00118_length=520441_Fill	93541	94017	ORF complete, but highly diverged
avrPto1Psy_B728a	54.94	scaffold00041_length=53134_Fill	45422	44946	ORF complete, but highly diverged
hopD1_Psv_ITM317	97.55	scaffold00015_length=796594_Fill	757328	755196	
hopG1Pto_DC3000	99.8	scaffold00015_length=796594_Fill	753478	754956	
hopH1_Psy_B728A	50	scaffold00035_length=822470_Fill	598670	599293	ORF complete, but highly diverged
hopJ1_Psy_B728A	70.22	scaffold00035_length=822470_Fill	149206	148766	
hopM1Psy_B728a	68.58	scaffold00035_length=822470_Fill	333311	335515	
hopQ1Pph_1448A	89	scaffold00035_length=822470_Fill	592849	594120	
hopR1Pph_1448A	82.65	scaffold00015_length=796594_Fill	45685	51549	
hopV1Pto_DC3000	99.7	scaffold00015_length=796594_Fill	761178	762347	
hopX1Pto_DC3000	83.55	scaffold00118_length=520441_Fill	369022	370167	
hopY1Pto_DC3000	63.38	scaffold00015_length=796594_Fill	43577	42720	
hopAA1_Pma_ES4326	79	scaffold00035_length=822470_Fill	330746	331921	C terminus in the middle of scaffold break
HopAB2_Pto_DC3000	55.36	scaffold00035_length=822470_Fill	414325	415806	
hopAC1_Psy_B728A	78.16	scaffold00015_length=796594_Fill	498487	492371	
hopAD1_Pto_DC3000	99.68	scaffold00100_length=113995_Fill	105671	107566	
hopAE1Pph_1448A	80.22	scaffold00100_length=113995_Fill	35578	32840	
hopAG1_Psy_B728A	65.89	scaffold00116_length=544280_Fill	451249	453438	Scaffold break in middle
hopAH1_Pto_DC3000	78.7	scaffold00116_length=544280_Fill	453546	454841	
hopAH2_Pto_DC3000	74	scaffold00097_length=271775_Fill	59921	58680	
hopAI1_Pto_DC3000	74.52	scaffold00116_length=544280_Fill	455028	455810	
hopAJ2_Pto_DC3000	97.49	scaffold00015_length=796594_Fill	701229	699901	
hopAK1_Pph_race6-1448A	70.22	scaffold00019_length=704711_Fill	226545	224953	
hopAN1_Pph_race6-1448A	95.1	scaffold00015_length=796594_Fill	412718	411507	
hopAO1Pto_DC3000	100	scaffold00015_length=796594_Fill	758907	760310	
hopAQ1_Pto_DC3000	100	scaffold00015_length=796594_Fill	774959	774804	C terminus in the middle of scaffold break
hopAS1_Pph_race6-1448A	86.09	scaffold00118_length=520441_Fill	318599	314517	
Potential Pseudogenes					
hopI1Pto_DC3000	86.32	scaffold00015_length=796594_Fill	732839	733390	frameshift caused by InDel
hopAF1Pto_DC3000	91.76	scaffold00019_length=704711_Fill	259211	258666	frameshift caused by InDel
hopZ1_Psy_A2	81	scaffold00035_length=822470_Fill	386393	385287	in-frame stop codon
avrE1Pto_DC3000	77	scaffold00035_length=822470_Fill	346607	341193	in-frame stop codon
hopH1_PtoDC3000	82.86	scaffold00015_length=796594_Fill	752666	752037	in-frame stop codon