

Second Generation Sequencing

Sequencing Machine	ABI 3730	Roche GSFLX	Illumina Solexa	AB SOLID	Helicos HeliScope	Pacific Bioscience
Launched	2000	2007	2006	2007	2008	2009
Read length (bp)	800-1100	250-400	35-50	25-35	28	?
Reads per run	96	400 K	60 M	85 M	85 M	?
Throughput per run	0.1 MB	100 MB	3 GB	3 GB	2 GB	?
Cost per GB	>\$2500k	\$84k	\$6k	\$5.8k	?	?









Size comparisons

	bp	distance	
Solexa read	35	1.9 m	me
Sanger read	900	50 m	swimming pool
BAC	120,000	6.7 Km	
rice genome	400,000,000	22,222 Km	distance to UK
canola genome	1,200,000,000	67, 000 Km	











Brassica genome sequencing

- A1 UK/China
- A2 Canada
- A3 Korea
- A4 UK/China
- A5 Japan
- A6 India? France?
- A7 Australia?
- A8 UK/China
- A9 Korea
- A10 Canada









http://www.brassica.info/resource/sequencing/status.php

Microsoft Excel - A8_bac_seq_prog_feb09a.xls [Read-Only]

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D	E	F	G	Н	J	К	L	м	N O	Р	Q	R	S	T U V W X
1	MBrG	SP BAC Re	aistry:	Chromosome A8 (R8)										
			J , J .											
2 Markers	Position(cM)	Markers	Position(cM) Markers	Markers	⊃osition (cN	1 Bin range		Year Status	Progress	Genbank No.	Project	date last undated	
.IVE30	JVE26	VESIMER	<u>усsзм.лн</u>	CKDH (Korea)	сконции	сконцкі	сконции						and tap Cabanca	
4 some walking		1000000		anas (rece)	ener (en)	Gries (Gri	enter (ent)		2007 Phase2	Complete	CII984540	IE-China	3-Dec-08	
5 some walking									2007 Phase2	Complete	CU984561	UK-China	3-Dec-08	
6 KBrB092P11	83.2								2007 Phase2	complete	not yet submitted	Korea	5-Dec-08	
7 m tiling		KBrB003007-6	48.6		_				2005 Phase3	complete	AC189194	Korea	5-Dec-08	
8 KS40490	77.1	-		nia_ssr084				Mapped to A08 by Chris Pires,	2005 Phase2	complete	AC189202	Korea	3-Dec-08	
9 KS10240	24.9	Velosso	50	msatfinder:KBrB006B05.19 (KBrB006B05:6197361994)	hri_akB.	1	68.3 -	75	2005 Phasel	complete	not yet submitted	Korea	S-Dec-08	
11 KC40700-2	72	KA40700-2	61.2	nsatfinder:KBrB006019.12 (KBrB006019:19739.19791)	hri mKB	30	26.9 -	44.1	2005 Phase2	complete	AC189205	Korea	5-Dec-08	
12 some walking									2008 Phase2	Complete	CU740081	UK-China	29-Apr-08	Brassica rapa Genome Sequencing Pro
13 KS10750	91.8	1		cnu ssr300 ; msatfinder:KBrB012024.10 (KBrB012024:3628236306)	hri mKB	13.7	8.1 - 1	6	2005 Phase2	complete	AC189228	Korea	5-Dec-08	Sequencing status
14 KC10680-1	60	KA10680-2	54.6	cnu_ssr537				Mapped to A08 by Chris Pires,	2005 Phase2	complete	AC189240	Korea	3-Dec-08	
15 KS10250	29.3	KS10250	32.3	nia_ssr050					2005 Phase3	complete	AC172858	Korea	5-Dec-08	
16 KS40210	90.6				hri_mKB	RB019A15	8.1 - 1	6	2005 Phase2	complete	AC189249	Korea	5-Dec-08	UK-China project
17 some waiking	;	KBrB023L08-4	51.3		-				2008 Phase2 2005 Phase3	complete	LU695257 AC189271	UK-Unina Korea	3-Sep-08	
19 some walking	;								2008 Phase2	Complete	CU914552	UK-China	3-Sep-08	Feb-09
20 some walking	;								2008 Phase2	Complete	CU695258	UK-China	3-Sep-08	
21 KS10710	59.1								2005 Phase3	complete	AC189296	Korea	5-Dec-08	data collated by Jun Wang and Graham King, Rothar
22 some walking									2007 Phase2	complete	CU915247	UK-China	3-Sep-08	
23 KS10160	17.4	K310160	12.6						2005 Phase3	complete	AC189331	Korea	5-Dec-08	
24 K510440	49.9	1 1210440	44.2						2005 Phases	complete	AC109346	Vorea	S=Dec=08	
26 some walking	40.5								2008 Phase2	Complete	CU695262	UK-China	29-Apr-08	
27 KBrB046G20	62	:		cnu_ssr459					2005 Phase3	complete	AC189366	Korea	5-Dec-08	
28 some walking									2008 Phase2	Complete	CU695263	UK-China	29-Apr-08	
29 KR40760-3	74.2	1			_			Mapped to A08 by Chris Pires,	2005 Phase3	complete	AC189375	Korea	3-Dec-08	
30 some walking					_				2008 Phase2	Complete	CU695264	UK-China	29-Apr-08	
31 KR10610-1a	35.9	KAIUGIU	49.4	ann ear190					2005 Phase3	complete	AC189385	Korea	S-Dec-US	
33 KS40300	81			cha_ssrood					2005 Phase2	complete	AC189398	Korea	5-Dec-08	
34 KR10430-1	44.7							Mapped to A08 by Chris Pires,	2005 Phase3	complete	AC189405	Korea	3-Dec-08	
35 KC40580-4	69.8	1							2005 Phase3	complete	AC189407	Korea	5-Dec-08	
36 m tiling		KBrB065D24-5	42						2007 Phase3	complete	not yet submitted	Korea	S-Dec-08	
37 m tiling; pre	viously 1	unassigned, up	dated Ja	n 2008	_				2005 Phase3	complete	not yet submitted	Rorea	24-Sep-08	
39 come walking	confirm	P072C19-4	E9 2		-				2008 Phase2	complete	CU695268	UK-Unina Voroo	29-Apr-08	
40 KBrB074G23-4	87.6	1	57.2						2005 Phase2	complete	AC189460	Korea	5-Dec-08	
41 KS40480	76.2								2005 Phase3	complete	AC189464	Korea	5-Dec-08	
42 some walking									2008 Phase2	Complete	CU695269	UK-China	29-Apr-08	
43 KS10320	32.5	KS10320	37	msatfinder:KBrB077F22.7 (KBrB077F22:3447034488)	hri_nRB	81.6	68.3 -	75	2005 Phase2	complete	AC189466	Korea	5-Dec-08	
44 KS10220	24.1								2007 Phase3	Complete	not vet submitted	Korea IIV-Chino	5-Dec-08	
46 KS40740	73.3	KS40740	63.1	nia ssr095					2005 Phase2	complete	AC189517	Korea	5-Dec-08	
47 KR10130-1	0	1							2005 Phase3	complete	AC189518	Korea	5-Dec-08	
48 m tiling		KS10170	15.2						2005 Phase3	complete	AC189519	Korea	5-Dec-08	
49 some walking									2008 Phase2	Complete	CU695274	UK-China	29-Apr-08	
50 some walking									2008 Phase2	Complete	CU695277	UK-China	29-Apr-08	
52 some walking									2008 Phase2	Complete	CU695280 CU928182	UK-China	29-Apr-08 3-Sen-09	
53 KC10200-3	21.3	KA10200-1	26.7				-		2007 Phase3	complete	not vet submitted	Korea	5-Dec-08	
54 KR10140-2	11.1								2005 Phase2	complete	AC189533	Korea	5-Dec-08	
55 m tiling ; al	so mappe	dKS40900	58.6	nsatfinder:KBrH004A18.21 (KBrH004A18:6898769002)	hri_nKB	38	26.9 -	44.1	2005 Phase3	complete	AC189541	Korea	5-Dec-08	
56 KS40981	65.1	KS40981	58	nia_ssr061 ; msatfinder:KBrH004J18.33 (KBrH004J18:9392494038)	hri_mRB	58.6	44.2 -	54	2005 Phase3	complete	AC189545	Korea	5-Dec-08	
57 KS40930	66.8			nia_ssr052 ; msatfinder:KBrH005C21.33 (KBrH005C21:5905059082)	hri_mKB	56.9	44.2 -	54	2005 Phase3	complete	AC189549	Korea	5-Dec-08	
59 m tiling: my	ziousla a	massimad	dated To:	2008					2008 PhaseZ	complete	LU075289	VA-UNINA Koree	29-Apr-08	
60 KS10150	21 9	anassigned, up	aaceu Ja	a 2000			-	Mapped to A08 by Chris Pires	2005 Phase?	complete	AC189590	Korea	3-Dec-08	
61 KS11010	93.3	8		nsat finder: KBrH013K13.10 (KBrH013K13:2085120881)	hri_nRB	0	<0		2005 Phase3	complete	AC189592	Korea	5-Dec-08	
62 some walking									2008 Phase2	Complete	CU695291	UK-China	29-Apr-08	
63 some walking									2008 Phase2	Complete	CU740085	UK-China	29-Apr-08	
54 some walking									2008 Phase2	Complete	CU914554	UK-China	30-Jun-08	
66 some walking									2008 PhaseZ	Complete	CU075345 CU928184	UK-Unina UK-Chine	29-Apr-08	
67 some walking		1					-		2008 Phase?	Complete	CU695295	UK-China	29-Apr-08	
68 some walking									2008 Phase2	Complete	CU695296	UK-China	29-Apr-08	
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New Chinese Initiative

Institute of Vegetables and Flowers (IVF)



Oil Crops Research Institute (OCRI) CAAS



Sanwen Huang, Xixiang Li

Shengyi Liu

生大基因

Peixiang Ni, **Hongkun Zheng**

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Ruiqiang Li

Korea: **Beom-Seok Park, Yong Pyo Lim** UK: Ian Bancroft Canada: **Isobel Parkin** Australia: **David Edwards** The Netherlands: Guusje Bonnema, Mark Aarts



IMB Institute for Molecular Bioscience







New Chinese Initiative

~50x coverage Illumina short reads

Contigs without N

- Longest = 45,025 bp
- N50 = 4,342 bp

Scaffolds containing N

- Longest = 293,912 bp
- N50 = 30,723 bp











Change of plan for completion

• Sequence BACs to phase 1, use paired read data to finish to phase 3

• Focus on assigned chromosomes but also sequence and map BACs from chromosomes 7 (and 6?)









New data, new capabilities

- Shotgun sequence *B. oleracea* and *B. napus*?
- Re-sequence different varieties to identify SNPs?
- Identify all gene variants in the genome and compare between genomes









GRDC related projects

ARC Linkage: Bayer CropScience

Developing technology for the cost effective *de novo* sequencing and analysis of complex genomes.

ARC Linkage: Bayer CropScience and DPI NWS Characterising genetic variation in *Brassica napus*.

ARC Linkage: Biomatters Ltd. Next generation metagenomics.

ARC Discovery: Jacqueline Batley and Regine Delourme Co-evolution of the host pathogen interaction between *Leptosphaeria maculans* and Brassica species.



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