REL606A: ge	neration 0															
LOUNT et al	. 2 sequenced at	M\$ single end,	35 or 36 bp					Tenaillon et a	I. 2 Population: Ara	a-3						
TRAIN	Generation	Clade	Population	Mtator	Run	Sequencing dep	CIT	strain	generation	clade	mutator	facility	run	read_type	read_length	sequencing_dept cit
DB464	200	00 (C1,C2)	ara-3	None	SRR098285	29.7	unknown	REL2181A	500		none	MSU RTSF	SRR2589044	paired	150	0 60.2 unknow
REL10979	400	00 C3+H	ARA-3	"+"	SRR098029	30.1	"+"	REL966A	100	00	none	IntraGen	SRR2589001	paired	10	1 67.3 unknow
REL10988	400	00 C2	Ara-3	"+"	SRR098030	30.2	minus	REL764B	50	00	None	IntraGen	SRR2584853	paired	10	1 82.4 unknow
DB429	100	00 UC	Ara-3	None	SRR098282	87.3	unknown	REL1166A	200	00	None	IntraGen	SRR2584859	paired	10	1 85.9 unknow
DB357	300	00 C2	Ara-3	None	SRR098280	111.2	unknown	REL7179B	1500	00	None	MSU RTSF	SRR2584863	paired	150	0 88 unknow
DB16	300	00 C1	Ara-3	None	SRR098031	113.9	unknown	REL1070A	150	00	None	IntraGen	SRR2584857	paired	10	1 92 unknow
DB458	200	00 (C1,C2)	Ara-3	None	SRR098284	126.8	unknown	REL4538A	1000	0 UC	None	MSU RTSF	SRR2589045	paired	150	99.5 unknow
DB446	150	00 UC	Ara-3	None	SRR098283	141.1	unknown	REL966B	100	00	None	IntraGen	SRR2584856	paired	10	1 101.2 unknow
DB409	50	00 unknown	Ara-3		SRR098281	144.2	unknown	REL1070B	150	00	None	IntraGen	SRR2584858	paired	10	1 102.4 unknow
DB467	200	00 (C1,C2)	Ara-3		SRR098286		unknown	REL1166B	200	00	None	MSU RTSF	SRR2591041	paired	150	0 108.5 unknow
DB477	250	00 C1	Ara-3		SRR098287		unknown	REL764A	50	00	None	IntraGen	SRR2584852	paired	10	1 113.4 unknow
DB483	250	00 C3	Ara-3		SRR098288		unknown	REL11365	5000	00 C3+H	plus	MSU RTSF	SRR2584866	paired	150	138.3 plus
DB199	315	00 C1	Ara-3	None	SRR098044		minus	REL11364	5000	0 C3+H	plus	MSU RTSF	SRR2584864	single	5	1 156.6 plus
DB200	315	00 C2	Ara-3	None	SRR098279		minus									
DB564	315	00 C3+	Ara-3	None	SRR098289		"+"	Leon et al. 20	18 read length: 10	1; population: AF	RA- clade: C3	sequenced at	JTA GSAF			
DB172	320	00 C3+	Ara-3	None	SRR098042		plus	strain	generation	run	cit					
DB30	320	00 C3	Ara-3	None	SRR098032		minus	ZDB1	1000	0 SRR6178299	unk					
DB143	325	00 C2	Ara-3	None	SRR098041		minus	ZDB425	1000	0 SRR6178304	unk					
DB158	325	00 C2	Ara-3	None	SRR098040		minus	ZDB445	1500	0 SRR6178301	unk					
ZB152	330	00 C3+	Ara-3	None	SRR098027		lus	ZDB478	2500	0 SRR6178302	unk					
ZB154	330	00 C3+	Ara-3	None	SRR097977		plus	ZDB486	2500	0 SRR6178309	unk					
ZB199	330	00 C1	Ara-3	None	SRR098026		minus	ZDB488	2500	0 SRR6178310	unk					
DB83	340	00 C3+	Ara-3	None	SRR098034		plus	ZDB309	2700	0 SRR6178307	unk					
DB87	340	00 C2	Ara-3	None	SRR098035		minus	ZDB310	2700	0 SRR6178308	unk					
DB96	360	00 C3+H	Ara-3	plus	SRR098036		plus	ZDB317	2700	0 SRR6178305	unk					
DB99	360	00 C1	Ara-3	None	SRR098037		minus	ZDB334	2800	0 SRR6178306	unk					
DB107	380	00 C3+H	Ara-3	plus	SRR098038		plus	ZDB339	2800	0 SRR6178303	unk					
ZDB111	380	00 C2	Ara-3	None	SRR098039		minus	ZDB13	2900	00 SRR6178300	unk					
								ZDB14	2900	0 SRR6178297	unk					
								ZDB17	3000	0 SRR6178298	unk					
								ZDB18	3000	0 SRR6178295	unk					
								ZDB19	3050	0 SRR6178296	unk					
								ZDB20	3050	0 SRR6178293	unk					
								ZDB23	3100	0 SRR6178294	unk					
								ZDB25	3150	0 SRR6178291	minus					
								ZDB27	3150	0 SRR6178292	minus					