

Statistics of watermelon Charleston Gray genome assembly

(version 1, 04/12/2014)

category	scaffold (>=200bp)			contig (>=200bp)		
	length (with N)	length (AGCT)	number	length (with N)	length (AGCT)	number
N01	23422029	22487609	1	235198	235198	1
N05	23422029	22487609	1	118089	118089	130
N10	21106771	20111028	2	93909	93909	311
N15	14209552	13504365	4	79590	79590	530
N25	12112137	11450730	7	61706	61706	1072
N50	7471260	7216157	17	36551	36551	3093
N60	5237419	5062134	23	29066	29066	4255
N70	4153776	3930082	32	22142	22142	5746
N80	3079070	2899440	43	15980	15980	7741
N90	1606462	1485283	61	9378	9378	10767
N95	519471	500949	81	5591	5591	13323
N96	328671	319776	90	4747	4747	14055
N97	83599	66392	113	3867	3867	14932
N98	27765	27765	202	2854	2854	16059
N99	8376	8046	477	1443	1443	17840
N100	200	200	6893	84	84	26406
Total bases	397803803	377267184		377267184	377267184	