

chromosome [1]	MCO448	PL448	BP57	CP367	CP61	SZMC21154	SZMC8029	SZMC8092	SZMC8094	SZMC8095	PL429 pe300	PL429 pe400ov	PL429 pe600	PL429 mp5000	PL429 fosmid	comments
scaffold1 size3459122	1.94	1.80	1.97	1.95	1.97	1.80	1.95	1.97	1.90	2.01	1.92	1.95	1.98	1.98	2.04	
scaffold2 size2959145	1.99	<b>2.16</b>	1.96	1.96	1.95	<b>2.39</b>	1.94	1.95	1.99	1.94	1.90	1.92	1.94	1.96	1.90	
scaffold7 size2161850	1.89	1.78	1.94	1.92	1.93	1.75	1.90	1.91	1.89	1.90	1.87	1.90	1.90	1.92	2.09	
scaffold3 size1388033	1.91	1.80	1.91	1.92	1.92	1.77	1.90	1.93	1.94	1.92	1.86	1.93	1.87	1.90	1.85	
scaffold5 size1063617 [2]	<b>2.39</b>	<b>3.08</b>	<b>2.32</b>	<b>2.34</b>	<b>2.34</b>	<b>2.25</b>	<b>2.32</b>	<b>2.35</b>	<b>2.38</b>	<b>2.36</b>	<b>2.21</b>	<b>2.44</b>	<b>2.24</b>	<b>2.44</b>	2.05	rDNA
scaffold4 size908642	1.91	1.79	1.94	1.96	1.94	1.77	1.92	1.93	1.95	1.91	1.87	1.94	1.91	1.94	1.74	
scaffold6 size820601	<b>2.23</b>	2.03	2.02	2.12	1.99	1.89	2.05	2.04	<b>2.20</b>	2.07	2.00	2.07	2.04	2.09	<b>2.38</b>	
scaffold8 size317314	<b>2.26</b>	2.01	2.00	1.99	1.98	1.88	2.01	2.02	2.16	2.13	1.97	2.03	2.01	2.11	2.15	
scaffold9 size263853	2.15	2.06	1.96	1.99	1.96	1.79	1.93	1.96	2.11	1.99	1.93	1.95	1.96	2.04	1.86	
scaffold10 size21864 [3]	1.80	1.99	<b>14.78</b>	<b>8.33</b>	<b>16.18</b>	<b>27.67</b>	<b>24.11</b>	<b>15.26</b>	<b>6.58</b>	<b>9.83</b>	<b>48.78</b>	<b>13.56</b>	<b>23.07</b>	<b>6.87</b>	0.00	mitochondrial chromosome
scaffold11 size20967	2.09	1.81	1.65	1.76	1.74	1.63	1.76	1.80	1.92	2.02	1.71	1.72	1.76	1.84	1.74	
scaffold12 size15030	<b>1.25</b>	<b>1.09</b>	<b>1.22</b>	<b>1.24</b>	<b>1.19</b>	<b>1.11</b>	<b>1.20</b>	<b>1.21</b>	<b>1.30</b>	<b>1.27</b>	<b>1.17</b>	<b>1.23</b>	<b>1.16</b>	<b>0.97</b>	<b>3.89</b>	
scaffold13 size9895	1.75	1.80	<b>1.34</b>	<b>1.98</b>	<b>1.22</b>	<b>2.12</b>	<b>0.38</b>	<b>1.31</b>	<b>1.46</b>	<b>1.27</b>	<b>2.02</b>	<b>1.69</b>	<b>1.76</b>	<b>1.74</b>	<b>0.00</b>	
scaffold14 size8524	1.65	<b>2.30</b>	1.76	1.71	1.73	1.74	1.86	1.83	1.77	1.76	1.84	1.67	1.69	1.79	0.69	
scaffold16 size6993	<b>0.87</b>	1.53	1.95	1.69	1.50	<b>2.89</b>	<b>3.63</b>	<b>2.22</b>	<b>1.24</b>	<b>1.49</b>	<b>3.56</b>	<b>2.61</b>	<b>3.47</b>	<b>2.66</b>	<b>8.39</b>	
scaffold15 size7016	2.02	2.16	1.70	1.91	1.72	1.76	1.62	1.65	1.83	1.86	<b>1.42</b>	1.80	<b>1.34</b>	<b>1.34</b>	0.97	
scaffold17 size5362	<b>2.97</b>	<b>2.88</b>	<b>2.46</b>	<b>3.04</b>	<b>1.07</b>	<b>3.37</b>	1.79	1.66	1.94	<b>1.46</b>	<b>2.79</b>	<b>2.50</b>	<b>2.54</b>	1.92	0.00	
scaffold18 size4846	1.83	2.80	<b>2.77</b>	<b>0.93</b>	1.75	<b>4.72</b>	<b>5.02</b>	<b>5.02</b>	<b>4.73</b>	<b>5.42</b>	0.00	8.87	8.67	1.73	0.00	
scaffold45 size1071	<b>22.49</b>	<b>23.70</b>	<b>9.07</b>	<b>52.99</b>	<b>8.56</b>	<b>7.76</b>	<b>8.98</b>	<b>8.80</b>	<b>48.46</b>	<b>6.70</b>	<b>14.62</b>	<b>9.47</b>	<b>18.46</b>	<b>4.66</b>	<b>12.93</b>	
scaffold48 size920	5.10	5.34	2.00	4.32	0.39	2.76	3.09	2.93	2.99	3.33	1.73	<b>2.42</b>	2.05	0.74	0.00	
scaffold54 size588	2.80	<b>3.35</b>	<b>1.34</b>	<b>5.47</b>	<b>1.08</b>	<b>2.39</b>	1.98	<b>2.85</b>	<b>4.29</b>	<b>1.43</b>	<b>5.50</b>	<b>7.95</b>	2.05	<b>2.98</b>	<b>29.87</b>	
scaffold19 size3739	0.48	1.04	1.55	2.05	1.68	0.26	0.29	0.28	1.97	0.33	1.86	<b>2.67</b>	1.50	2.14	3.13	
scaffold20 size3160	0.76	1.68	1.51	0.85	<b>2.36</b>	1.55	1.59	1.00	0.72	0.88	1.30	0.82	1.25	0.48	0.00	
scaffold21 size3072	1.37	1.31	0.97	1.25	1.00	0.94	0.96	0.96	1.07	1.14	1.06	1.55	0.89	0.91	<b>3.81</b>	
scaffold22 size2988	0.94	0.98	0.87	0.82	0.95	0.92	0.81	1.04	0.98	0.98	1.05	1.03	0.86	0.41	0.00	
scaffold23 size2932	0.33	0.26	1.27	1.40	0.68	1.06	0.71	0.60	<b>1.76</b>	0.70	0.55	<b>3.34</b>	0.15	0.56	<b>29.95</b>	
scaffold24 size2890	0.04	0.03	0.99	1.15	1.00	0.05	0.02	0.02	1.37	0.03	0.03	0.10	0.02	0.03	0.00	
scaffold25 size2889	0.74	0.77	0.95	0.88	0.95	0.69	0.85	0.83	0.90	0.74	0.72	0.68	0.79	0.56	0.00	
scaffold26 size2760	1.22	1.08	0.73	0.82	0.76	0.75	0.85	0.82	0.89	1.03	0.77	1.14	0.64	0.61	0.00	
scaffold27 size2613	0.01	0.01	0.97	1.02	1.00	0.00	0.00	0.00	0.99	0.00	0.01	0.01	0.01	0.01	0.00	
scaffold28 size2552	0.96	0.86	1.08	0.92	1.02	0.00	1.01	1.07	1.06	0.96	0.98	0.96	1.05	0.99	0.00	
scaffold29 size2468	1.53	1.48	1.62	1.49	1.62	<b>2.20</b>	1.59	1.58	1.41	1.51	0.87	0.82	0.79	0.42	<b>2.37</b>	
scaffold30 size2308	0.73	0.93	1.02	0.86	0.98	0.88	0.90	0.94	0.86	0.78	1.06	0.89	1.07	1.00	0.00	
scaffold31 size2209	0.52	0.46	1.64	1.31	1.64	0.73	0.82	0.84	<b>1.58</b>	0.82	0.66	0.78	0.52	0.34	<b>2.65</b>	
scaffold32 size2198	0.22	0.34	0.62	1.00	1.08	0.40	1.10	1.10	1.03	0.91	1.44	0.76	1.27	0.72	0.00	
scaffold33 size2195	1.03	0.98	1.57	0.86	0.39	1.67	1.83	1.00	0.89	0.97	1.10	1.09	1.03	0.82	<b>2.67</b>	
scaffold34 size2195	0.86	0.74	0.84	0.75	0.81	0.72	0.73	0.78	0.77	0.85	0.41	0.43	0.36	0.07	0.00	
scaffold35 size2060	0.99	0.82	0.81	0.88	0.82	0.70	0.77	0.77	0.86	0.82	0.94	0.87	0.82	0.30	0.00	
scaffold36 size1996	0.90	1.60	0.84	0.78	0.85	0.79	1.64	0.87	0.94	0.05	0.91	0.97	0.88	0.73	0.00	
scaffold37 size1869	1.39	1.53	1.08	1.79	1.80	1.88	2.00	2.13	<b>3.00</b>	1.97	1.86	1.80	1.90	1.38	<b>3.13</b>	

scaffold38 size1589	2.24	1.85	0.55	1.54	1.38	1.67	1.72	1.73	1.59	2.09	2.44	2.14	1.91	0.71	0.00
scaffold39 size1516	1.72	1.35	1.17	1.36	1.29	1.68	1.30	1.25	1.39	1.51	1.12	1.12	0.82	0.51	0.00
scaffold40 size1514	0.60	0.55	1.36	1.31	1.27	0.67	0.73	0.68	1.69	0.78	0.93	0.61	0.55	0.43	0.00
scaffold41 size1479	1.84	1.64	1.38	1.70	1.54	1.47	1.54	1.59	1.66	1.69	1.98	2.44	1.37	1.08	0.00
scaffold42 size1291	0.00	0.00	0.69	0.79	0.11	0.00	0.00	0.00	0.93	0.00	0.00	0.01	0.00	0.01	0.00
scaffold43 size1245	1.97	1.72	1.53	1.50	1.38	1.20	1.26	1.34	1.41	1.53	1.67	2.18	1.33	0.79	0.00
scaffold44 size1158	0.00	0.00	0.87	0.98	0.93	0.00	0.00	0.00	1.03	0.00	0.00	0.00	0.00	0.00	0.00
scaffold46 size1011	1.45	1.26	0.83	1.21	1.07	1.02	1.09	1.01	1.11	1.15	1.39	1.70	0.77	1.00	5.79
scaffold47 size994	0.53	0.82	1.27	1.05	1.31	1.27	1.21	1.19	0.66	0.60	1.74	1.46	1.30	1.22	11.78
scaffold49 size918	1.09	1.01	0.95	0.85	0.97	0.86	0.92	0.93	0.89	0.15	1.51	1.41	1.07	1.23	6.38
scaffold50 size882	0.53	0.87	1.41	1.43	0.73	0.71	0.79	0.85	0.83	0.47	0.81	0.74	0.92	1.05	0.00
scaffold51 size796	0.01	0.01	0.69	0.82	0.72	0.00	0.00	0.00	0.86	0.00	0.01	0.00	0.00	0.04	0.00
scaffold52 size792	0.00	0.00	1.08	0.91	1.10	0.00	0.00	0.00	1.02	0.00	0.00	0.00	0.00	0.00	0.00
scaffold53 size677	0.76	1.27	1.36	1.43	1.17	1.31	1.85	1.35	0.81	0.69	1.73	0.84	2.02	2.21	8.59
scaffold55 size569	0.71	1.03	0.96	1.14	1.10	1.09	1.17	1.14	0.83	0.76	1.23	1.06	1.07	1.13	20.58
scaffold56 size522	0.83	0.77	0.78	0.73	0.76	1.28	0.75	0.65	0.65	0.67	1.93	1.91	1.74	1.97	0.00
scaffold57 size518	0.01	0.18	0.05	0.02	0.08	0.03	0.01	0.04	0.00	0.02	0.05	0.02	0.10	1.69	0.00

[1] core chromosomes are in bold

[2] rDNA cluster

[3] mtDNA is highly dependant on DNA isolation technique, if nuclear DNA was enriched, little or no mtDNA is expected to be sequenced