

Supplementary Table 1 Samples included in the analysis according to BLIND ordering.

Sample name	Annotated time/stage	Counted reads (genes)	Counted reads (ERCC)	Counted genes
<i>P. dumerilii</i>				
PT0005	300	681568	41980	6640
PT0001	60	577220	70082	6385
PT0003	180	804010	124567	7157
PT0004	240	709031	118360	7076
PT0006	360	810083	107398	7091
PT0007	420	597310	112207	7435
PT0008	480	1377866	106535	8527
PT0009	540	900560	91019	7576
PT0010	600	998981	110855	7853
PT0012	720	700919	66042	7342
PT0013	780	465169	113705	7396
PT0014	840	572886	70201	7192
PT0016	960	922296	104135	8030
PT0017	1020	1059582	127037	8224
PT0015	900	765433	109974	8217
PT0020	1200	467940	100472	7000
PT0019	1140	620799	113260	7915
PT0018	1080	670991	93554	8242
PT0025	1560	538879	93809	6826
PT0026	1620	747435	94828	8275
PT0024	1500	614968	76925	8099
PT0030	1860	985442	87280	8665
PT0027	1680	1268911	136535	8899
PT0028	1740	1090567	95544	8871
PT0029	1800	1273122	122872	8874
PT0031	1920	1084301	81723	8769
PT0032	1980	662822	63513	8323
PT0033	2040	464284	46182	7416
PT0034	2100	628178	59307	8301
PT0035	2160	832949	75593	8763
PT0037	2280	966112	64257	8111
PT0036	2220	743666	58717	8605
PT0038	2340	717946	50838	8357
PT0039	2400	687201	59238	8354
PT0040	2460	739542	59378	8530
PT0041	2520	1403309	141475	9638
PT0042	2580	1653093	117712	9578
PT0043	2640	1318067	111798	9502
PT0044	2700	1373586	102004	9495
PT0045	2760	1390536	112141	9552

PT0046	2820	1195204	111805	9423
PT0048	2940	1205721	93256	9585
PT0049	3000	1394118	130792	9261
PT0050	3060	725309	71418	8584
PT0051	3120	743197	62149	4700
PT0052	3180	860190	64738	5336
PT0057	3480	2577933	115021	7420
PT0054	3300	978923	74567	5510
PT0053	3240	1249381	102656	5903
PT0059	3600	1153911	103399	5693
PT0056	3420	858784	77454	5132
PT0058	3540	1147598	86748	5764
PT0060	3660	824226	81604	5145
PT0062	3780	547222	41772	4032
PT0064	3900	804163	78776	4988
PT0065	3960	866680	82850	5158
PT0063	3840	928299	106576	5008
PT0066	4020	868486	66620	4953
PT0067	4080	1248041	124545	5506
PT0068	4140	680387	68975	5033
PT0070	4260	875054	83558	5420
PT0072	4380	514979	72971	4656
PT0069	4200	750849	78969	4619
PT0073	4440	766357	101721	4518
PT0075	4560	712341	91694	4394
PT0080	4860	795394	93077	4919
PT0074	4500	443912	54317	3759
PT0076	4620	676428	76424	5263
PT0077	4680	790667	105479	4755
PT0078	4740	679305	85814	4625
PT0079	4800	752087	98014	4529
PT0082	4980	454756	66183	3840
PT0091	5520	687313	118701	4729
PT0081	4920	556274	93160	4139
PT0084	5100	438558	52641	3853
PT0093	5640	752933	114832	4778
PT0092	5580	498673	77603	4146
PT0088	5340	407727	34788	3508
PT0087	5280	440506	60557	2927
PT0090	5460	437535	53939	3532

S. polychroa

FW0108	1	471534	11501	8582
FW0130	3	433389	31223	8735
FW0122	3	638065	22752	10082
FW0112	2	553706	21409	9829
FW0114	2	417217	29273	9039
FW0132	3	806561	22969	10091
FW0121	3	468622	22247	9528
FW0115	2	519937	49135	9500
FW0125	3	1032887	32095	11339
FW0049	5	433472	74393	10237
FW0042	4	501400	91382	10732
FW0041	4	776945	90151	11521
FW0038	4	715199	43534	12299
FW0057	5	548386	46422	11484
FW0043	4	844644	89355	11685
FW0036	4	1032949	57962	12446
FW0047	5	727616	68217	11949
FW0050	5	512407	41458	11877
FW0073	6	621893	60913	11663
FW0065	6	412765	40131	11318
FW0054	5	1112761	60194	12824
FW0066	6	593949	74948	11557
FW0061	6	513889	53023	11692
FW0070	6	533586	61713	12387
FW0035	4	788421	29769	13168
FW0055	5	841916	58440	13127
FW0067	6	562982	43189	11664
FW0058	6	601292	38260	11684
FW0051	5	1055676	47043	13393
FW0064	6	622145	25725	12274
FW0052	5	1251141	63796	13432
FW0059	6	685246	41617	12496
FW0069	6	1068561	66513	13311
FW0084	7	832728	22879	13855
FW0085	7	671228	19659	12584
FW0088	7	1301374	38769	13849
FW0089	7	956826	30829	13438
FW0075	7	1091667	32793	13321
FW0081	7	700371	33044	12329
FW0080	7	1512296	34827	13854
FW0078	7	1075723	49817	13109
FW0079	7	1531600	37809	13726
FW0077	7	672560	23161	12571
FW0087	7	1623149	49002	14282
FW0083	7	610429	33740	12963

FW0092	8	1093508	13301	13451
FW0090	8	563653	9625	12667
FW0082	7	717279	28335	13338
FW0100	8	801213	13495	12346
FW0095	8	1035899	12617	12930
FW0091	8	682548	13362	12022
FW0103	8	863110	9278	12397
FW0093	8	916690	12236	12689
FW0098	8	847292	10568	13465
FW0074	7	533076	16792	12713
FW0096	8	741443	11368	12046
FW0102	8	654591	11661	12013
FW0101	8	609778	14848	11859

C. elegans

CE0046	30	648258	3421	7024
CE0053	30	1168406	5598	6743
CE0054	30	1762351	7725	6902
CE0096	-30	1697207	1882	8996
CE0051	0	1349669	5381	6152
CE0052	0	978778	3271	5781
CE0095	-50	2206884	3715	8861
CE0101	40	2317349	4399	10086
CE0098	10	2360636	3662	9408
CE0103	60	1585271	4126	10453
CE0104	70	896597	3306	9987
CE0072	20	4557314	17625	10429
CE0102	50	1499248	3392	10298
CE0055	70	554674	3525	5508
CE0056	70	699720	3837	5886
CE0108	110	711957	2118	9696
CE0057	110	1205412	7297	6785
CE0107	100	1927968	6911	10775
CE0109	120	1996361	6800	10502
CE0060	140	692649	4325	5844
CE0110	130	1264572	3738	10287
CE0111	140	513094	1832	8462
CE0112	150	1187968	3537	9822
CE0113	160	1525934	4731	10120
CE0114	170	1822764	3151	10460
CE0074	200	1282092	10897	9251
CE0115	180	1516342	3890	9986
CE0086	200	959327	3331	9435
CE0088	220	1550505	7086	9723
CE0076	220	2867909	8550	10338
CE0116	190	1128714	2388	9732
CE0087	210	1396974	4220	10564

CE0120	230	1598295	2667	10576
CE0092	290	514765	1324	9594
CE0091	280	1309976	4121	10082
CE0080	290	3169191	9077	11049
CE0123	260	2225197	3765	11634
CE0127	300	2664028	17494	11689
CE0124	270	1824186	2771	11576
CE0063	310	1566714	8092	7538
CE0064	310	1143535	4420	7103
CE0128	310	814212	1901	9957
CE0081	380	760663	1935	11186
CE0129	320	2758725	10587	12143
CE0066	340	1104506	4214	6786
CE0133	360	1936400	9040	11321
CE0130	330	2254120	8435	11866
CE0132	350	1933195	9289	11760
CE0134	370	1426465	2610	10637
CE0067	380	1908726	8823	7890
CE0082	400	1258477	3768	11111
CE0131	340	842531	2119	10921
CE0068	380	1913256	6810	8250
CE0136	390	2600646	11881	12378
CE0069	400	1312991	6630	7458
CE0138	410	1763959	6422	11426
CE0139	420	2247757	9363	12184
CE0020	410	855654	2582	9511
CE0005	440	531473	2006	8369
CE0015	450	807643	2412	9059
CE0013	430	1661072	3784	10748
CE0017	490	910328	2516	9931
CE0016	510	718413	1657	9359
CE0034	540	662718	2642	8281
CE0021	550	891965	3063	9507
CE0024	580	686643	2006	9226
CE0014	600	559134	1220	9612
CE0033	620	821268	1734	9091
CE0023	630	785772	2826	9125
CE0035	690	506608	1672	8367
CE0025	660	623795	2060	7983
CE0032	700	1006677	3472	8207
CE0028	740	634443	2089	7536
CE0042	640	1216706	4155	9550
CE0045	750	999983	2326	9421
CE0048	840	662565	2662	8122
CE0050	830	1306826	3789	9474
CE0049	800	750425	1611	8886

CE0040	760	878648	2612	8705
CE0037	780	511198	1077	8503
CE0041	810	877461	1578	9249
		<i>H. dujardini</i>		
HD0008	270	2175441	312559	8789
HD0010	360	1604242	350364	8379
HD0005	120	1637670	176151	8392
HD0007	240	4196340	530616	9002
HD0006	180	2613662	307598	8642
HD0112	960	774196	3367	9294
HD0110	900	1482692	6462	9901
HD0111	930	832592	3213	9315
HD0113	990	1056906	3903	9203
HD0004	90	4331054	388647	9059
HD0009	300	4972036	414583	8885
HD0002	30	3039305	245146	8725
HD0012	420	1587171	187253	8447
HD0001	0	3650555	346205	8875
HD0003	60	3426322	517121	9369
HD0013	450	3699814	696436	8830
HD0014	480	1442738	180134	7639
HD0015	510	4104690	436657	8680
HD0016	540	1699612	213395	7884
HD0119	1170	848682	5905	9563
HD0124	1320	425334	3084	9233
HD0123	1290	1186475	7350	10034
HD0125	1350	1204582	7023	10103
HD0126	1380	523004	3072	9631
HD0055	2280	752928	154266	6233
HD0026	1410	609737	133449	5222
HD0027	1440	1087654	183117	6382
HD0029	1500	1756168	332678	7264
HD0028	1470	1010468	151303	6459
HD0030	1530	610275	127237	5692
HD0031	1560	1067481	356034	6729
HD0032	1590	1208796	222402	6899
HD0033	1620	811742	135209	5516
HD0034	1650	657814	137793	5657
HD0035	1680	1156732	232701	6585
HD0038	1770	748268	145285	6007
HD0041	1860	1292816	292470	6391
HD0036	1710	796306	158187	5723
HD0042	1890	835553	180827	5946
HD0043	1920	790928	178789	5645
HD0048	2070	537381	82532	4702
HD0051	2160	500395	115970	4651

HD0054	2250	1632110	233475	7475
HD0053	2220	1100872	205535	6381
HD0056	2310	2400687	346938	7874
HD0057	2340	1016999	134358	6974
HD0060	2430	2177144	349696	8043
HD0063	2520	705171	118899	6158
HD0067	2640	1220231	205754	7479
HD0066	2610	1263915	184793	7607
HD0071	2760	799844	119118	6393
HD0069	2700	921083	149880	7054
HD0070	2730	1124678	216625	7274
HD0075	2880	791611	146607	6319
HD0072	2790	694302	123316	5630
HD0078	2970	521121	121616	5659
HD0073	2820	733073	118264	6760
HD0068	2670	2077092	383030	8198
HD0074	2850	546807	94611	5952
HD0022	2760	2026191	183845	9327
HD0023	2850	915069	97116	7401
HD0025	2940	1511717	194827	7381

D. melanogaster

DM0002	0	2687595	85237	7223
DM0001	-15	2286826	79587	7173
DM0004	30	1670023	85718	7176
DM0003	15	2331605	80147	7221
DM0005	45	1213557	54308	6661
DM0012	150	609408	51642	4144
DM0007	75	2182589	104198	7320
DM0008	90	2388874	102946	7467
DM0009	105	1638267	66695	7056
DM0011	135	561108	32150	5581
DM0014	180	1497773	73309	7025
DM0018	240	2027556	89589	7374
DM0013	165	1438210	83594	6936
DM0006	60	1979477	89999	7542
DM0016	210	439350	86085	5173
DM0015	195	2197202	104762	7785
DM0019	255	2078411	112206	7656
DM0020	270	2819385	129801	8113
DM0021	285	2006424	114188	7636
DM0024	330	1300032	81786	6450
DM0023	315	1849586	139236	7416
DM0022	300	1416117	89966	7222
DM0042	600	584875	79754	4014
DM0041	585	815158	77702	5361
DM0037	525	1217773	94130	5907

DM0031	435	1587831	106806	6706
DM0034	480	631243	47551	5268
DM0025	345	1740242	74823	7182
DM0028	390	839049	45894	6112
DM0026	360	2186242	104990	7458
DM0033	465	960366	79692	6299
DM0039	555	1506619	98682	6339
DM0036	510	1523878	114449	6549
DM0035	495	898278	52110	6256
DM0027	375	2407313	92392	7717
DM0030	420	2501751	141167	7519
DM0043	615	1503175	96367	6366
DM0040	570	2747282	171154	7732
DM0032	450	2081128	99747	7479
DM0044	630	1110109	118742	5921
DM0029	405	1947457	95440	7853
DM0047	675	1065624	64189	8312
DM0052	750	1706293	98886	8721
DM0038	540	2486403	89722	8229
DM0045	645	2094989	107703	7908
DM0054	780	1615614	92521	9200
DM0060	870	1611063	85511	9082
DM0048	690	1647145	84474	9452
DM0046	660	1502524	66495	9194
DM0053	765	1611588	72886	9169
DM0059	855	1624813	79244	9313
DM0064	930	1245187	71134	8975
DM0050	720	823617	35184	8247
DM0055	795	1644069	71798	9312
DM0049	705	2441235	98570	9848
DM0061	885	1665387	68338	9186
DM0062	900	1872826	105635	9304
DM0066	960	1932002	107735	9488
DM0058	840	1741904	91362	9325
DM0051	735	1964421	79688	9581
DM0071	1035	2518821	125287	9698
DM0056	810	902574	33528	8542
DM0065	945	2571015	99631	9825
DM0068	990	2030130	103458	9382
DM0067	975	1609392	65257	9165
DM0070	1020	3308015	109493	10184
DM0063	915	1376068	56603	8991
DM0084	1230	2843277	118768	9469
DM0078	1140	2774847	120424	9759
DM0081	1185	2577832	136355	9408
DM0085	1245	1193766	55219	8394

DM0077	1125	2896821	117952	9702
DM0090	1320	2173903	103114	9494
DM0087	1275	3097439	132949	9936
DM0088	1290	2483315	120684	9739
DM0083	1215	1868625	79879	9370
DM0089	1305	1468253	110700	8008
		<i>S. purpuratus</i>		
SU0005	200	694425	222946	17078
SU0011	760	491960	152687	16672
SU0141	360	401022	37653	16203
SU0008	320	1415058	232786	18948
SU0052	360	672588	239782	17363
SU0054	440	1585131	292221	19412
SU0055	480	725898	249692	18998
SU0060	1000	990532	327474	18381
SU0146	1200	488216	31363	17603
SU0070	1720	2280500	183101	20075
SU0149	1680	489874	33934	16575
SU0023	1520	486542	133325	13861
SU0020	1400	895132	217758	18620
SU0064	1160	874126	354280	17203
SU0017	1280	725490	259257	16619
SU0018	1320	413657	108452	13752
SU0021	1440	551938	150014	15703
SU0025	1600	675210	169115	15443
SU0067	1280	547609	153792	17020
SU0066	1240	648934	161810	15412
SU0024	1560	789626	181915	16121
SU0068	1640	985780	258540	17770
SU0026	1960	459678	117522	15944
SU0071	1760	458624	118326	15225
SU0027	2000	556810	126239	14435
SU0074	1880	890317	257775	17343
SU0072	1800	689737	216902	16242
SU0031	2480	2451801	446254	19424
SU0078	2120	834394	164665	16268
SU0028	2360	533340	129351	13431
SU0080	2200	666546	214543	16339
SU0079	2160	405203	90149	14569
SU0082	2280	472338	121927	14376
SU0077	2080	565569	138318	16565
SU0083	2320	542577	124452	15681
SU0033	2560	657655	184995	14696
SU0076	2040	429260	103793	15710
SU0034	2600	572913	153177	13825
SU0035	2640	748366	177795	17021

SU0081	2240	528366	142691	15330
SU0088	2840	769838	250473	17189
SU0085	2720	408219	107583	15825
SU0038	3080	1472470	300715	18581
SU0089	2880	455466	98749	15430
SU0086	2760	405528	117327	14902
SU0036	3000	457402	100956	13472
SU0040	3160	467079	112262	13194
SU0092	3320	1021627	264444	18264
SU0094	3400	1154311	253894	18282
SU0099	3600	596623	129034	17130
SU0100	3640	650766	170427	16934
SU0095	3440	467521	91969	16997
SU0097	3520	466908	111014	17239
SU0098	3560	640479	162232	17593
SU0043	3280	615618	141404	16690
SU0093	3360	601745	120555	17570
SU0128	4200	1010615	204083	18595
SU0110	4120	498397	128284	16128
SU0103	3840	430828	100009	15970
SU0111	4160	548351	124248	17044
SU0124	4520	621433	173921	16767
SU0122	4440	650927	173151	16979
SU0116	4360	1096769	292663	17937
SU0125	4560	535307	185498	16043
SU0108	4040	697217	172571	16293
SU0109	4080	403679	97297	16219
SU0114	4280	821493	275302	17374
SU0096	3480	785953	198521	17230
<i>D. rerio</i>				
ZF0007	280	727866	53207	6673
ZF0003	120	1620208	73343	9398
ZF0004	160	1197803	44692	8054
ZF0001	40	689209	41857	5761
ZF0006	240	1015423	57356	8345
ZF0013	520	678107	38723	6397
ZF0016	640	511848	23998	5942
ZF0015	600	908481	35065	7194
ZF0017	680	2138819	81564	8277
ZF0020	800	1552826	47213	8075
ZF0018	720	958813	29556	7121
ZF0019	760	1454934	54578	7281
ZF0022	880	950626	34192	6701
ZF0025	1000	905647	27730	7040
ZF0026	1040	1255829	26818	8140
ZF0027	1080	1342410	29771	8621

ZF0023	920	2279740	69876	8649
ZF0029	1160	785984	23900	6641
ZF0028	1120	858627	18800	7686
ZF0030	1200	1277654	24414	8313
ZF0034	1360	584710	12263	5004
ZF0033	1320	1002283	24633	6203
ZF0031	1240	1723148	39577	9376
ZF0024	960	1391813	34465	8022
ZF0032	1280	851981	15390	7600
ZF0036	1440	1128533	18408	7225
ZF0037	1480	1164889	36831	6915
ZF0038	1520	1610955	19712	8006
ZF0039	1560	1443913	23388	7890
ZF0040	1600	1047890	15738	6991
ZF0042	1680	2859897	35617	11253
ZF0041	1640	1528963	27948	8850
ZF0043	1720	1312704	19097	8660
ZF0045	1800	2061063	28180	9779
ZF0044	1760	2656648	31808	10440
ZF0046	1840	1221576	16993	8746
ZF0047	1880	1088575	13819	7686
ZF0048	1920	1075421	15507	8452
ZF0054	2160	980215	13041	7158
ZF0052	2080	3394933	46611	10403
ZF0049	1960	4577296	71264	10991
ZF0050	2000	2005815	36280	9461
ZF0051	2040	4570469	68806	10776
ZF0057	2280	3934051	34680	11558
ZF0055	2200	2276658	21254	10690
ZF0056	2240	1509127	18364	8925
ZF0059	2360	1896378	17889	10027
ZF0053	2120	3062606	48685	9795
ZF0058	2320	1360520	16528	8586
ZF0062	2480	2917992	39453	9476
ZF0060	2400	720648	8306	7052
ZF0061	2440	1744615	17657	8908
ZF0074	2960	1520112	31596	8422
ZF0082	3280	1683627	49090	7213
ZF0084	3360	2653951	59333	8624
ZF0064	2560	3945928	39056	10578
ZF0073	2920	2368533	27347	10779
ZF0076	3040	1837187	24342	9642
ZF0083	3320	2661693	29003	8945
ZF0063	2520	2256553	25498	9457
ZF0065	2600	810657	10645	5918
ZF0069	2760	3023994	38138	10383

ZF0068	2720	4709802	53410	10819
ZF0071	2840	1741004	18074	9835
ZF0072	2880	2647030	27937	10665
ZF0078	3120	3485547	41786	10397
ZF0070	2800	1668034	18621	8784
ZF0081	3240	2585745	41132	8938
ZF0077	3080	1466102	17802	9400
ZF0087	3480	1343935	17005	7988
ZF0092	3680	1055016	13398	8181
ZF0086	3440	591653	13417	5377
ZF0104	4160	1397155	24460	8014
ZF0094	3760	10217895	113742	13333
ZF0090	3600	1771105	22862	9193
ZF0079	3160	2679793	29229	9708
ZF0085	3400	2325011	21449	9462
ZF0089	3560	1477199	12459	9392
ZF0095	3800	4164173	35311	11898
ZF0101	4040	5788896	52233	12595
ZF0099	3960	4234659	40472	12279
ZF0098	3920	3583393	37101	10984
ZF0102	4080	2568150	23451	11003
ZF0097	3880	3213241	28746	10891
ZF0088	3520	2347639	23857	10841
ZF0105	4200	554087	7714	5481
ZF0093	3720	1204765	9868	8750
ZF0103	4120	1480502	16683	9634
		<i>N. vectensis</i>		
NV0008	220	896172	361538	19834
NV0004	80	594349	214018	18115
NV0050	-20	1018123	370482	18177
NV0002	20	787233	219590	18629
NV0003	40	888605	351895	19419
NV0007	200	912053	369267	19144
NV0012	400	1012741	422783	20577
NV0016	560	509348	218071	17169
NV0019	680	1333573	468437	19805
NV0024	880	772685	192510	17997
NV0022	800	542925	163051	16619
NV0023	840	862248	242205	18427
NV0026	960	1341689	301184	22008
NV0028	1060	766545	164216	19390
NV0030	1160	1706574	432428	23076
NV0029	1120	1676973	353943	22457
NV0036	1400	1402882	266953	22497
NV0035	1360	535853	151843	19044
NV0039	1520	811057	235267	20263

NV0033	1280	946697	207461	21491
NV0031	1200	1541825	314689	22294
NV0045	1760	612056	189041	19720
NV0049	1920	591373	183623	19787
NV0034	1320	1602804	237707	22615
NV0032	1240	1368668	258069	22118
NV0037	1440	1572334	186245	22907
NV0038	1480	1173944	182756	21659
NV0052	2000	2051815	406131	25843
NV0053	2040	518980	169426	20840
NV0041	1600	1076231	126558	22012
NV0073	2840	741450	99210	22102
NV0065	2520	1041722	241318	23846
NV0060	2320	921595	222789	23314
NV0058	2240	745031	283383	21826
NV0055	2120	1342897	297289	24030
NV0062	2400	986182	197858	22893
NV0063	2440	658387	142336	21246
NV0051	1960	1755470	254241	23785
NV0059	2280	738715	180503	21696
NV0044	1720	1296502	210154	22270
NV0047	1840	736539	118472	20362
NV0042	1640	1124264	139030	21742
NV0048	1880	812631	112153	21410
NV0072	2800	792349	160601	21599
NV0069	2680	1206247	218295	23451
NV0040	1560	2646585	330586	23553
NV0054	2080	1554742	201762	23919
NV0067	2600	1678711	185449	25046
NV0070	2720	1460701	222754	24902
NV0107	3000	701998	71247	17336
NV0071	2760	1247919	149037	24098
NV0117	2600	1412429	98334	20407
NV0113	2760	871727	41458	18818
NV0066	2560	1970889	175561	25283
NV0118	2560	1634079	93557	20516
NV0064	2480	1405665	181293	24451
NV0110	2880	649289	87303	16897
NV0079	4120	758031	70374	18521
NV0075	4280	953494	60170	19921
NV0121	2440	1063925	53738	19173
NV0105	3080	1492204	65827	21791
NV0116	2640	1335235	89529	20292
NV0114	2720	1162650	75428	19165
NV0100	3280	1470447	88916	20338
NV0120	2480	729577	64933	17271

NV0084	3920	604729	53095	17105
NV0104	3120	1629528	109223	20469
NV0092	3600	520066	68675	15544
NV0102	3200	2029785	127671	22175
NV0098	3360	1251692	54897	20500
NV0108	2960	1672773	83075	20575
NV0109	2920	827571	36112	18143
NV0097	3400	1016876	57309	19147
NV0091	3640	796298	61806	18948
NV0078	4160	677007	42633	16850
NV0093	3560	1827588	102199	21854
NV0087	3800	843433	55310	18312
NV0090	3680	2113426	91948	21906
NV0095	3480	1212798	90500	19807
NV0088	3760	1910978	121762	21386
NV0099	3320	1234653	56370	19921
NV0089	3720	1175599	69637	19340
NV0094	3520	1379810	93300	19699

A. queenslandica

AQ0100	1	944757	54991	14168
AQ0065	1	1700085	83980	15923
AQ0099	1	1587647	71348	14468
AQ0003	1	2591430	79498	21795
AQ0028	2	2116187	32683	19571
AQ0004	2	849645	8919	18516
AQ0059	2	1806685	15390	14626
AQ0063	2	3563847	84922	17165
AQ0005	2	915998	23753	15713
AQ0060	2	2361730	49107	17099
AQ0009	2	1975705	60079	16690
AQ0051	3	1678767	29131	14632
AQ0010	3	2449683	40721	17602
AQ0068	3	2026721	40438	17228
AQ0008	3	2716028	60261	17959
AQ0017	3	4997392	68359	24835
AQ0064	3	2862907	37234	17450
AQ0006	3	925328	21712	14393
AQ0074	4	1625998	48350	14157
AQ0078	4	2596022	94898	14898
AQ0030	4	693474	11504	15143
AQ0097	5	1045072	49556	12359
AQ0066	4	2764340	50498	15566
AQ0048	4	1051721	21361	15242
AQ0079	5	1286013	13662	13647
AQ0007	5	1738658	39833	16727
AQ0040	4	1812023	28369	16594

AQ0070	6	1894790	11660	14669
AQ0002	6	1033268	16287	14097
AQ0095	5	4359554	85937	17112
AQ0026	6	1552142	21443	16548
AQ0085	6	2860800	63374	15779
AQ0042	6	1482777	20315	15134
AQ0088	5	2753331	84296	15700
AQ0013	6	2002920	34096	17599
AQ0019	7	2576299	66771	16002
AQ0032	7	923915	14340	14370
AQ0038	7	965698	18612	13973
AQ0001	7	1254080	29743	14651
AQ0027	7	1433618	23955	14710
AQ0057	7	1683613	71677	15969
AQ0029	8	1017487	25489	16946
AQ0084	8	1505063	68667	14334
AQ0034	8	849712	22642	15224
AQ0025	8	1387416	73188	16099
AQ0022	8	1469511	63132	16109
AQ0081	8	1118943	52746	13913
AQ0094	8	1307495	72324	14293
AQ0096	8	1057642	56701	12784
AQ0053	8	801275	51360	11896
AQ0098	7	1000900	48920	12023
		<i>M. leidy</i>		
ML0043	120	401652	5866	6246
ML0022	60	1861129	5023	8201
ML0042	60	1151164	10020	7760
ML0024	180	2036972	4598	8530
ML0023	120	1622510	4936	7956
ML0021	0	2423255	6028	8753
ML0001	0	1663802	3814	8406
ML0044	180	864377	8207	7486
ML0025	240	1592641	4490	8268
ML0002	60	2539697	4510	9048
ML0003	120	3205977	4701	9412
ML0004	180	2762924	4495	9267
ML0045	240	554706	6577	7319
ML0046	270	953736	10091	8116
ML0026	270	1765162	4089	8688
ML0047	300	678408	5677	7739
ML0006	270	1854929	3620	9418
ML0007	300	999350	3301	8587
ML0008	330	1434231	4476	9348
ML0029	360	2377873	7259	9655
ML0049	360	872390	8753	8668

ML0011	420	1382830	4254	9287
ML0027	300	2299919	8262	9794
ML0009	360	1994545	4825	10246
ML0030	390	1424209	6057	8847
ML0051	420	507616	6414	7806
ML0050	390	925981	6964	8838
ML0010	390	2447153	3824	10621
ML0031	420	4508618	8670	11248
ML0053	480	519768	8205	8123
ML0032	450	1865779	3920	9642
ML0052	450	923820	9521	9175
ML0012	450	1368424	2798	10227
ML0054	510	797187	11526	9283
ML0033	480	3052642	7544	10984
ML0014	510	2474678	6415	10999
ML0013	480	1592021	3929	10546
ML0055	540	842919	8660	9599
ML0035	540	974408	5763	9231
ML0015	540	1401639	3899	10224
ML0056	600	484496	7847	8533
ML0057	660	743177	10651	9249
ML0036	600	1926181	5260	10746
ML0058	720	1265333	9891	10758
ML0059	780	414804	5282	8565
ML0038	720	1908109	6017	10577
ML0037	660	2217519	5895	10812
ML0017	660	3148306	4579	12359
ML0019	780	1536966	3495	11093
ML0039	780	2087047	7136	10963
ML0020	1200	1339223	4206	10869
ML0060	1200	606527	7106	9609
ML0040	1200	1342583	4567	11017

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