

Supplementary Table S2: Annotation file of mapped reads after initial filtering steps (see Supplementary Figure S1).

Chromosome: Chromosome to which the reads map.

Start: The location of the first 5' end nucleotide of the reads.

Length: Length of the read after removal of the adapter sequences.

Strand: The DNA strand to which the read was mapped. (+) and (-) denotes the top and bottom strand, respectively, as defined by dictyBase (www.dictybase.org).

Annotation: The mapped reads are annotated as: complex repeat regions i.e. DDT-A, DIRS 1, DDT-B, TRE3-A, Skipper, thug-S, TRE5-A1 or TRE5-A ((Glockner et al. 2001) and www.dictyBase.org); non-coding (nc)RNAs i.e. tRNAs, snoRNAs, rRNA, Class I RNAs (rNN), and predicted ncRNAs (drpNNN) with an e-value <0.001 (Larsson et al. 2008); protein coding genes (DDB_GNNN) as defined in the gff file (generated 2009-04-29) downloaded from dictyBase and; intergenic regions denoted intergenic_NNN. In the cases where one location have multiple/overlapping annotations the displayed annotation was chosen in the following hierarchical order: complex repeat regions, annotated ncRNAs, annotated protein coding genes, predicted ncRNAs and intergenic regions.

Chromosome	Start	Length	Strand	Annotation
1	74662	21	+	DDB_G0267258
1	74667	21	+	DDB_G0267258
1	74755	21	-	DDB_G0267258
1	75324	21	-	DDB_G0267258
1	75337	21	-	DDB_G0267258
1	114454	21	+	DDB_G0267304
1	116152	21	+	DDB_G0267304
1	116154	21	+	DDB_G0267304
1	216234	21	+	DDB_G0267492
1	1319053	21	+	DDB_G0268022
1	597508	21	-	DDB_G0268502
1	2143202	21	-	DDB_G0268808
1	2388251	21	-	DDB_G0269032
1	4858512	21	+	DDB_G0270432
1	4775888	21	+	DDB_G0270804
1	4775889	21	+	DDB_G0270804
1	4776109	21	+	DDB_G0270804
1	4776110	21	+	DDB_G0270804
1	4776112	21	+	DDB_G0270804
1	4776113	21	+	DDB_G0270804
1	4776114	21	+	DDB_G0270804
1	4776118	21	+	DDB_G0270804
1	4776119	21	+	DDB_G0270804
1	4776121	21	+	DDB_G0270804
1	4776122	21	+	DDB_G0270804
1	91708	21	+	DDT-A 1207 3051

1	12633	21	-	DDT-A 1780 2784
1	4914154	21	+	DDT-A 1936 2749
1	71791	21	+	DDT-A 588 3967
1	40231	21	+	DDT-B 1616 5424
1	175649	21	-	DIRS-1 1 1329
1	132976	21	-	DIRS-1 1 3821
1	132978	21	-	DIRS-1 1 3821
1	136485	21	-	DIRS-1 1 3821
1	131140	21	-	DIRS-1 1 4826
1	131706	21	+	DIRS-1 1 4826
1	131708	21	+	DIRS-1 1 4826
1	131709	21	+	DIRS-1 1 4826
1	28085	21	+	DIRS-1 1 4826
1	31872	21	+	DIRS-1 1 4826
1	4424	21	+	DIRS-1 1 4826
1	46886	21	-	DIRS-1 1 4826
1	46887	21	-	DIRS-1 1 4826
1	4956	21	+	DIRS-1 1 4826
1	49853	21	-	DIRS-1 1 4826
1	6418	21	-	DIRS-1 1 4826
1	7401	21	+	DIRS-1 1 4826
1	7402	21	+	DIRS-1 1 4826
1	7403	21	+	DIRS-1 1 4826
1	88671	21	-	DIRS-1 1 687
1	139684	21	+	DIRS-1 1 913
1	178848	21	+	DIRS-1 1 913
1	24026	21	+	DIRS-1 1417 1761
1	185021	21	-	DIRS-1 1816 3353
1	145857	21	-	DIRS-1 1816 4826
1	147971	21	+	DIRS-1 1816 4826
1	148567	21	+	DIRS-1 1816 4826
1	90662	21	-	DIRS-1 1968 2016
1	11364	21	-	DIRS-1 2419 4826
1	11618	21	+	DIRS-1 2419 4826
1	11619	21	+	DIRS-1 2419 4826
1	11620	21	+	DIRS-1 2419 4826
1	8923	21	-	DIRS-1 2496 3300
1	8924	21	-	DIRS-1 2496 3300
1	112756	21	+	DIRS-1 2533 3627
1	25314	21	+	DIRS-1 2635 3821
1	2120	21	+	DIRS-1 2910 4450
1	2740	21	-	DIRS-1 2910 4450
1	2996	21	+	DIRS-1 2910 4450
1	56201	21	-	DIRS-1 34 1084
1	56472	21	+	DIRS-1 34 1084
1	89833	21	+	DIRS-1 3432 4781

1	61785	21	+	DIRS-1 3671 4445
1	109218	21	+	DIRS-1 3701 4451
1	109219	21	+	DIRS-1 3701 4451
1	127412	21	+	DIRS-1 3821 4826
1	35147	21	+	DIRS-1 511 970
1	52791	21	-	DIRS-1 610 4826
1	52792	21	-	DIRS-1 610 4826
1	52793	21	-	DIRS-1 610 4826
1	56001	21	+	DIRS-1 610 4826
1	3642470	21	+	TRE3-A 1 2339
1	3642471	21	+	TRE3-A 1 2339
1	3643103	21	-	TRE3-A 1 2339
1	3643107	21	-	TRE3-A 1 2339
1	3643177	21	-	TRE3-A 1 2339
1	4706600	21	-	TRE3-A 1 5248
1	4707253	21	-	TRE3-A 1 5248
1	4707981	21	-	TRE3-A 1 5248
1	4708012	21	-	TRE3-A 1 5248
1	4708016	21	-	TRE3-A 1 5248
1	4708086	21	-	TRE3-A 1 5248
1	742243	21	+	TRE3-A 1 5248
1	2849768	21	-	TRE5-A1 4631 5652
1	3220178	21	-	TRE5-A1 4672 5653
1	4552023	21	+	TRE5-A1 4677 5118
1	2012269	21	+	Tdd-4 1 3387
1	2164925	21	-	drp130
1	2166447	21	+	drp132
1	2414279	21	+	drp154
1	1758380	21	-	drp91
1	3181051	21	-	forE
1	268196	21	+	intergenic_126
1	3515179	21	-	intergenic_1375
1	1756429	21	+	intergenic_694
1	1756430	21	+	intergenic_694
1	4044570	21	-	mgp2
1	4044571	21	-	mgp2
1	3001192	21	+	r35
1	3001196	21	+	r35
1	122288	21	+	skipper 1 1869
1	818921	21	+	tRNA-Cys-GCA-1
1	2850098	21	+	tRNA-Gln-UUG-3
1	2850099	21	+	tRNA-Gln-UUG-3
1	2558702	21	+	tRNA-Ser-AGA-1
1	2342379	21	+	wdr75
2	291162	21	-	DDB_G0271208
2	1632037	21	+	DDB_G0272292

2	1610720	21	+	DDB_G0272448
2	1611441	21	+	DDB_G0272448
2	1970378	21	+	DDB_G0272690
2	2071452	21	-	DDB_G0272861
2	2592367	21	-	DDB_G0273093
2	2497645	21	-	DDB_G0273323
2	2783014	21	+	DDB_G0273375
2	2783015	21	+	DDB_G0273375
2	2784164	21	+	DDB_G0273375
2	3247422	21	-	DDB_G0273663
2	3248571	21	-	DDB_G0273663
2	3248572	21	-	DDB_G0273663
2	3248573	21	-	DDB_G0273663
2	3439221	21	+	DDB_G0273813
2	3533941	21	+	DDB_G0273897
2	6472786	21	-	DDB_G0276055
2	6605362	21	-	DDB_G0276307
2	6769333	21	-	DDB_G0276525
2	6769337	21	-	DDB_G0276525
2	7127115	21	-	DDB_G0276687
2	7127116	21	-	DDB_G0276687
2	7432987	21	+	DDB_G0276983
2	8227642	21	+	DDB_G0277621
2	8481118	21	+	DDB_G0277785
2	923363	21	+	DDB_G0294587
2	367497	21	+	TRE3-A 1 5248
2	14140	21	+	TRE3-A 4216 4478
2	7424699	21	+	TRE3-C 3988 4751
2	333877	21	+	TRE5-A_ModA 236 523
2	1183699	21	+	drp446
2	2555210	21	-	fip-1
2	3476378	21	+	fip-2
2	1128	21	-	intergenic_0
2	2874	21	-	intergenic_0
2	407	21	-	intergenic_0
2	2889992	21	-	intergenic_1129
2	2889993	21	-	intergenic_1129
2	2890098	21	-	intergenic_1129
2	3141488	21	+	intergenic_1219
2	3141594	21	+	intergenic_1219
2	292293	21	-	intergenic_128
2	312906	21	+	intergenic_140
2	3693136	21	-	intergenic_1441
2	4787322	21	+	intergenic_1864
2	4787323	21	+	intergenic_1864
2	4787809	21	-	intergenic_1864

2	4982370	21	-	intergenic_1943
2	4982374	21	-	intergenic_1943
2	4982425	21	-	intergenic_1943
2	5492360	21	-	intergenic_2134
2	5985687	21	-	intergenic_2315
2	7174782	21	+	intergenic_2787
2	7174840	21	+	intergenic_2787
2	7876894	21	+	intergenic_3075
2	7876901	21	+	intergenic_3075
2	7876921	21	+	intergenic_3075
2	7876923	21	+	intergenic_3075
2	7876924	21	+	intergenic_3075
2	7876925	21	+	intergenic_3075
2	7876926	21	+	intergenic_3075
2	7876934	21	+	intergenic_3075
2	7876935	21	+	intergenic_3075
2	7876936	21	+	intergenic_3075
2	8477165	21	+	intergenic_3311
2	886496	21	+	intergenic_352
2	1630880	21	+	intergenic_629
2	1630881	21	+	intergenic_629
2	2338450	21	+	intergenic_906
2	5917517	21	+	med26
2	8069109	21	-	nubp1
2	8304234	21	-	r29
2	8304270	21	-	r29
2	5946893	21	+	r41
2	4084553	21	-	r42
2	4084554	21	-	r42
2	4315375	21	+	rnf160
2	471469	21	-	rpc5
2	1825553	21	+	tRNA-Gln-UUG-6
2	1825554	21	+	tRNA-Gln-UUG-6
2	6364438	21	-	tRNA-Gln-UUG-8
2	8290778	21	+	tRNA-His-GUG-2
2	8084355	21	-	tRNA-His-GUG-3
2	5947115	21	-	tRNA-His-GUG-4
2	5289937	21	-	tRNA-His-GUG-5
2	3515072	21	-	tRNA-Met-CAU-10
2	2513178	21	-	tRNA-Met-CAU-11
2	3518408	21	+	tRNA-Met-CAU-6
2	4399816	21	+	tRNA-Ser-AGA-2
2	6379240	21	+	tRNA-Ser-AGA-3
2	6571286	21	+	tRNA-Ser-AGA-4
2	7878647	21	-	tRNA-Ser-AGA-5
2	7878648	21	-	tRNA-Ser-AGA-5

2	7878649	21	-	tRNA-Ser-AGA-5
2	7237150	21	-	tRNA-Ser-AGA-6
2	7237151	21	-	tRNA-Ser-AGA-6
2	7237152	21	-	tRNA-Ser-AGA-6
2	2516514	21	+	thug-S 579 726
2	4109189	21	-	thug-S 580 730
2F	158010	21	-	DDB_G0294310
2F	136332	21	+	DDT-A 3576 4116
2F	69578	21	+	DIRS-1 1 1099
2F	79691	21	+	DIRS-1 1 2183
2F	79692	21	+	DIRS-1 1 2183
2F	104017	21	-	DIRS-1 1 496
2F	104020	21	-	DIRS-1 1 496
2F	95259	21	-	DIRS-1 1 816
2F	95352	21	-	DIRS-1 1 816
2F	113817	21	-	DIRS-1 1387 1773
2F	113823	21	-	DIRS-1 1387 1773
2F	113973	21	+	DIRS-1 1387 1773
2F	142831	21	+	DIRS-1 2156 4322
2F	144942	21	-	DIRS-1 2156 4322
2F	73364	21	-	DIRS-1 3267 3328
2F	73255	21	-	DIRS-1 3519 3797
2F	152168	21	-	DIRS-1 3933 4826
2F	61964	21	+	DIRS-1 4140 4799
2F	61968	21	+	DIRS-1 4140 4799
2F	145891	21	+	DIRS-1 512 970
2F	96241	21	+	DIRS-1 512 970
2F	67806	21	-	DIRS-1 83 396
2F	103152	21	-	DIRS-1 936 1667
2F	124322	21	+	Tdd-4 2422 2589
2F	152718	21	+	intergenic_73
2F	154067	21	-	intergenic_74
2F	157488	21	-	intergenic_79
2F	111586	21	+	skipper 790 1795
3	2528768	21	+	CYP513D1
3	819604	21	-	DDB_G0278395
3	885814	21	-	DDB_G0278443
3	692580	21	-	DDB_G0278659
3	692582	21	-	DDB_G0278659
3	1861766	21	+	DDB_G0279267
3	2901166	21	-	DDB_G0280079
3	3010158	21	-	DDB_G0280373
3	3748873	21	-	DDB_G0280809
3	3748876	21	-	DDB_G0280809
3	3986006	21	-	DDB_G0281057
3	4885572	21	+	DDB_G0281727

3	5346066	21	+	DDB_G0282057
3	1667483	21	-	TRE3-A 1 5248
3	1670245	21	-	TRE3-A 1 5248
3	2633580	21	+	TRE3-A 1 5248
3	2633700	21	+	TRE3-A 1 5248
3	4607142	21	-	TRE3-A 1 5248
3	4607490	21	+	TRE3-A 1 5248
3	4607493	21	+	TRE3-A 1 5248
3	4607494	21	+	TRE3-A 1 5248
3	4607525	21	+	TRE3-A 1 5248
3	4607758	21	+	TRE3-A 1 5248
3	4524347	21	+	TRE3-A 1265 1718
3	4524467	21	+	TRE3-A 1265 1820
3	2875040	21	-	TRE3-A 13 5248
3	2155591	21	-	TRE3-A 7 2339
3	2155595	21	-	TRE3-A 7 2339
3	2155665	21	-	TRE3-A 7 2339
3	1570069	21	+	TRE3-A 85 5248
3	890777	21	-	TRE3-D 719 822
3	5869696	21	-	TRE5-A1 4468 5652
3	1776470	21	+	TRE5-A1 4760 5652
3	2626480	21	-	TRE5-A1 4855 4958
3	2626236	21	-	TRE5-A1 5234 5653
3	5674586	21	-	Tdd-4 1 2600
3	2684844	21	+	Tdd-4 633 3843
3	3378221	21	+	ascc2
3	1574153	21	-	drp1003
3	2103520	21	-	drp1024
3	2330677	21	+	drp1034
3	2450297	21	+	drp1039
3	2623293	21	-	drp1045
3	2632482	21	+	drp1046
3	3739296	21	-	drp1151
3	4409131	21	+	drp1196
3	4409414	21	+	drp1196
3	5325958	21	-	drp1272
3	5706992	21	-	drp1300
3	549118	21	+	drp949
3	1098365	21	+	drp981
3	1098366	21	+	drp981
3	1101921	21	+	drp984
3	1101922	21	+	drp984
3	2602338	21	+	intergenic_1003
3	2910215	21	+	intergenic_1113
3	4883105	21	+	intergenic_1930
3	5268560	21	+	intergenic_2082

3	568298	21	-	intergenic_228
3	1650073	21	-	metS
3	1101113	21	-	r25
3	4352741	21	-	r33
3	4352742	21	-	r33
3	2300535	21	-	r36
3	2300573	21	-	r36
3	2300577	21	-	r36
3	2300578	21	-	r36
3	2300579	21	-	r36
3	1100445	21	-	r46
3	1100446	21	-	r46
3	1671409	21	-	tRNA-Cys-GCA-5
3	3366200	21	+	tRNA-Gln-UUG-11
3	3366201	21	+	tRNA-Gln-UUG-11
3	4240817	21	+	tRNA-Gly-UCC-1
3	4240820	21	+	tRNA-Gly-UCC-1
3	4240821	21	+	tRNA-Gly-UCC-1
3	5706555	21	-	tRNA-Gly-UCC-3
3	4247198	21	-	tRNA-Gly-UCC-5
3	4523857	21	+	tRNA-Leu-AAG-2
3	4660226	21	-	tRNA-Met-CAU-12
3	3143646	21	+	tRNA-Ser-AGA-7
3	2539836	21	-	tRNA-Ser-AGA-8
3	2539837	21	-	tRNA-Ser-AGA-8
3	2539838	21	-	tRNA-Ser-AGA-8
3	5902934	21	+	tRNA-Ser-GCU-4
3	5781598	21	-	tRNA-Ser-GCU-6
3	5339384	21	-	tRNA-Ser-GCU-7
3	5335140	21	-	tRNA-Ser-GCU-8
3	4919674	21	+	thug-S 1 877
3	4529840	21	+	thug-S 1 905
3	4529841	21	+	thug-S 1 905
3	4530987	21	+	thug-S 1396 2192
3F	8208	21	-	DIRS-1 3654 4826
4	232738	21	+	DDB_G0283133
4	1213248	21	+	DDB_G0283893
4	1213249	21	+	DDB_G0283893
4	2028768	21	+	DDB_G0284481
4	3149962	21	-	DDB_G0285397
4	3228184	21	+	DDB_G0285441
4	3392612	21	-	DDB_G0285567
4	3392613	21	-	DDB_G0285567
4	3228749	21	-	DDB_G0285737
4	4106791	21	-	DDB_G0286165
4	4657170	21	+	DDB_G0286585

4	4659586	21	+	DDB_G0286587
4	4666827	21	+	DDB_G0286789
4	693004	21	+	DIRS-1 2168 2298
4	693005	21	+	DIRS-1 2168 2298
4	4137433	21	+	DIRS-1 3322 3359
4	1622766	21	+	E2F
4	3944988	21	-	TRE3-A 1 1197
4	3716463	21	-	TRE3-A 1 1579
4	3073776	21	+	TRE3-A 1305 3837
4	3089661	21	+	TRE3-A 3084 5248
4	3089663	21	+	TRE3-A 3084 5248
4	3089664	21	+	TRE3-A 3084 5248
4	3711014	21	+	TRE3-A 3176 3866
4	1665692	21	+	TRE3-A 3535 5248
4	2845926	21	+	TRE5-A1 5358 5653
4	2734258	21	-	TRE5-B 5560 5667
4	735747	21	-	drp1411
4	735761	21	-	drp1411
4	735768	21	-	drp1411
4	736031	21	-	drp1412
4	866594	21	-	drp1420
4	897144	21	+	drp1423
4	897145	21	+	drp1423
4	957510	21	+	drp1430
4	957511	21	+	drp1430
4	960597	21	+	drp1431
4	960598	21	+	drp1431
4	1663000	21	+	drp1458
4	1663001	21	+	drp1458
4	1667276	21	+	drp1459
4	1667277	21	+	drp1459
4	2792500	21	+	drp1541
4	2792501	21	+	drp1541
4	2927366	21	+	drp1556
4	2931837	21	+	drp1557
4	5064054	21	+	drp1702
4	2673735	21	+	fsjC
4	2673757	21	-	fsjC
4	3952261	21	+	gmsA
4	3385812	21	+	intergenic_1315
4	4486771	21	+	intergenic_1749
4	4487122	21	+	intergenic_1749
4	1476931	21	-	intergenic_553
4	2377641	21	-	polB
4	2377642	21	-	polB
4	956841	21	-	r21

4	956876	21	-	r21
4	956877	21	-	r21
4	956878	21	-	r21
4	956879	21	-	r21
4	956880	21	-	r21
4	960338	21	-	r22
4	1668863	21	-	r23A
4	1667001	21	-	r23B
4	1662746	21	-	r23C
4	1663884	21	+	r24A
4	1675683	21	-	r24B
4	1675709	21	-	r24B
4	1668202	21	+	r26
4	1668241	21	+	r26
4	2793917	21	+	r32
4	3575039	21	+	r34
4	3575070	21	+	r34
4	959146	21	-	r50
4	959605	21	-	r51
4	961592	21	+	r52
4	2792238	21	-	r56
4	2767691	21	+	srpA
4	2767940	21	+	srpA
4	2767952	21	+	srpA
4	5435372	21	+	tRNA-Leu-AAG-4
4	2220215	21	-	tRNA-Ser-GCU-11
4	317650	21	+	thug-S 1 123
5	172203	21	+	DDB_G0287355
5	507091	21	-	DDB_G0287667
5	734682	21	+	DDB_G0287863
5	878289	21	+	DDB_G0287951
5	1174684	21	+	DDB_G0288167
5	1432679	21	+	DDB_G0288355
5	1590937	21	-	DDB_G0288505
5	1590968	21	-	DDB_G0288505
5	1590972	21	-	DDB_G0288505
5	1591042	21	-	DDB_G0288505
5	1677020	21	-	DDB_G0288547
5	1934525	21	+	DDB_G0288763
5	2053828	21	+	DDB_G0288841
5	2744649	21	-	DDB_G0289399
5	2321841	21	+	TRE3-A 1 891
5	3759178	21	-	TRE3-A 3058 5248
5	3759179	21	-	TRE3-A 3058 5248
5	3058765	21	-	TRE5-A1 5239 5651
5	849496	21	-	drp1787

5	849811	21	-	drp1787
5	984430	21	+	drp1797
5	984802	21	+	drp1798
5	1458870	21	-	drp1816
5	2684954	21	-	intergenic_1064
5	593956	21	+	intergenic_242
5	594068	21	+	intergenic_242
5	850067	21	+	intergenic_349
5	850068	21	+	intergenic_349
5	850349	21	+	intergenic_349
5	850350	21	+	intergenic_349
5	1592356	21	+	intergenic_629
5	1592371	21	+	intergenic_629
5	172141	21	+	intergenic_76
5	4859054	21	-	pus1
5	1473461	21	+	r31
5	1225000	21	+	roco9
5	2600331	21	-	sno13
5	1594020	21	-	thug-S 1 167
5	1594101	21	-	thug-S 1 167
5	425235	21	-	thug-S 1 905
5	425236	21	-	thug-S 1 905
5	425237	21	-	thug-S 1 905
5	4796458	21	+	thug-S 1 905
5	4796459	21	+	thug-S 1 905
5	424089	21	-	thug-S 1396 2037
5	4797617	21	+	thug-S 1396 2192
5	1515666	21	+	thug-S 16 333
5	2966818	21	+	thug-S 2128 2176
5	3226493	21	+	thug-S 585 730
6	778784	21	-	DDB_G0291806
6	1077869	21	-	DDB_G0292046
6	1009495	21	-	DDB_G0292154
6	1349663	21	-	DDB_G0292236
6	1661339	21	+	DDB_G0292450
6	3599369	21	-	DDT-A 1909 2782
6	8965	21	+	DDT-B 1944 2634
6	2460	21	-	DIRS-1 1 2285
6	6739	21	+	DIRS-1 1 2456
6	12962	21	+	DIRS-1 3331 3449
6	11647	21	+	DIRS-1 3443 4445
6	11648	21	+	DIRS-1 3443 4445
6	147	21	-	DIRS-1 4484 4826
6	1527988	21	-	TRE3-A 1 5248
6	1528709	21	-	TRE3-A 1 5248
6	878021	21	+	TRE3-A 1 5248

6	870708	21	-	TRE3-A 2 5248
6	871091	21	+	TRE3-A 2 5248
6	2768548	21	-	TRE3-A 2273 2920
6	1126040	21	+	TRE3-A 2727 5248
6	1777093	21	-	TRE3-A 2776 3756
6	1777094	21	-	TRE3-A 2776 3756
6	1777103	21	-	TRE3-A 2776 3756
6	1777223	21	+	TRE3-A 2776 3756
6	859970	21	-	TRE3-A 3 886
6	2954254	21	+	TRE3-A 3694 3895
6	875717	21	+	TRE3-A 4347 5248
6	610434	21	-	TRE3-A 937 5248
6	610465	21	-	TRE3-A 937 5248
6	610469	21	-	TRE3-A 937 5248
6	610494	21	-	TRE3-A 937 5248
6	1379361	21	+	Tdd-4 1 3504
6	2258615	21	+	drp2192
6	2268749	21	+	drp2194
6	2270952	21	+	drp2195
6	2340690	21	+	drp2202
6	2728505	21	+	drp2223
6	632323	21	-	intergenic_246
6	746280	21	+	intergenic_290
6	2295180	21	+	r59
6	2298172	21	+	r60
6	3354232	21	+	r62
6	850626	21	+	rfc2
6	1664160	21	-	tRNA-Cys-GCA-8
6	1927259	21	-	tRNA-Gly-GCC-17
6	1838015	21	-	tRNA-His-GUG-10
6	2840444	21	+	tRNA-His-GUG-8
6	3173679	21	+	tRNA-His-GUG-9
6	1948288	21	-	tRNA-Met-CAU-17
6	2347735	21	-	tRNA-Ser-GCU-12
6	3233660	21	-	thug-S 2026 2126
6	2905301	21	-	thug-S 23 375
BF	2779	21	+	DDT-B 1942 2433
BF	8570	21	-	DIRS-1 1 2372
BF	8573	21	-	DIRS-1 1 2372
BF	8574	21	-	DIRS-1 1 2372
BF	68485	21	+	intergenic_28
BF	70759	21	+	intergenic_28
BF	71078	21	+	intergenic_28
BF	71329	21	+	intergenic_28
M	5355	21	+	DidioMp05
M	54145	21	+	intergenic_28

M	22134	21	+	intergenic_9
M	25402	21	+	tRNA-Asn-GUU-1
M	55535	21	+	tRNA-Glu-UUC-1
M	30112	21	+	tRNA-Met-CAU-2
R	8668	21	-	26S_rRNA-1
R	75913	21	+	26S_rRNA-2
R	76232	21	+	26S_rRNA-2
R	76483	21	+	26S_rRNA-2
R	11512	21	-	5.8S_rRNA-1
R	73639	21	+	5.8S_rRNA-2
R	5306	21	-	5S_rRNA-1
R	5307	21	-	5S_rRNA-1
R	5310	21	-	5S_rRNA-1
R	5315	21	-	5S_rRNA-1
R	5316	21	-	5S_rRNA-1
R	5317	21	-	5S_rRNA-1
R	5318	21	-	5S_rRNA-1
R	5319	21	-	5S_rRNA-1
R	5340	21	-	5S_rRNA-1
R	5347	21	-	5S_rRNA-1
R	5383	21	-	5S_rRNA-1
R	5396	21	-	5S_rRNA-1
R	5398	21	-	5S_rRNA-1
R	79753	21	+	5S_rRNA-2
R	79768	21	+	5S_rRNA-2
R	79804	21	+	5S_rRNA-2
R	79811	21	+	5S_rRNA-2
R	79831	21	+	5S_rRNA-2
R	79833	21	+	5S_rRNA-2
R	79834	21	+	5S_rRNA-2
R	79835	21	+	5S_rRNA-2
R	79836	21	+	5S_rRNA-2
R	79844	21	+	5S_rRNA-2
R	79845	21	+	5S_rRNA-2
R	79846	21	+	5S_rRNA-2

Supplementary references

- Glockner G, Szafranski K, Winckler T, Dingermann T, Quail MA, Cox E, Eichinger L, Noegel AA, Rosenthal A. 2001. The complex repeats of *Dictyostelium discoideum*. *Genome Res* 11: 585-594.
- Larsson P, Hinas A, Ardell DH, Kirsebom LA, Virtanen A, Soderbom F. 2008. De novo search for non-coding RNA genes in the AT-rich genome of *Dictyostelium discoideum*: performance of Markov-dependent genome feature scoring. *Genome Res* 18: 888-899