

Table S4. Sanger sequencing results of amplified single-cell cDNA library^a.

Unenriched (0.1-1.5 kb)		Unenriched (1.5-4.0 kb)		Enriched (0.1-1.5 kb)		Enriched (1.5-4.0 kb)	
Gene ID ^b	Matched Length ^c	Gene ID ^b	Matched Length ^c	Gene ID ^b	Matched Length ^c	Gene ID ^b	Matched Length ^c
BTH_I1096	374	BTH_I0083	685	BTH_II0015	206	BTH_II1723	523
BTH_I1099	332	BTH_I0132	692	BTH_II1865	188	BTH_I0927	638
BTH_I2684	274	BTH_II2048	774	BTH_I0024	305	BTH_I1284	802
BTH_I1064	336	BTH_I2199	691	BTH_II0003	489	BTH_II0572	628
BTH_II0025	285	BTH_I1385	672	BTH_I0014	220	BTH_I0232	721
BTH_I1096	638	BTH_I1099	694	BTH_II0575	306	BTH_II0935	628
BTH_I1046	572	No match	-	BTH_II1253	212	BTH_I0143	627
BTH_I2298	497	BTH_I1096	687	No match	-	BTH_II1212	838
BTH_I0648	396	BTH_I1096	882	BTH_I2948	436	BTH_I0091	739
BTH_I0482	712	BTH_I2828	791	BTH_I0450	649	BTH_I1927	652
BTH_I1096	385	BTH_I0088	685	BTH_I0046	283	BTH_I2124	573
BTH_II1483	212	BTH_II0444	643	BTH_I0273	389	BTH_I0285	675
BTH_I0287	626	BTH_II1448	792	BTH_I2846	492	BTH_I1006	594
No match	-	BTH_I1096	626	BTH_II2195	673	BTH_I1096	709
BTH_I1096	254	BTH_II1653	730	BTH_I3075	608	BTH_II1313	783
BTH_I0286	191	BTH_I1005	682	BTH_II2237	439	BTH_II0440	684
BTH_I1778	323	BTH_I1099	682	BTH_I0476	497	BTH_II0438	585
BTH_I1741	323	BTH_I1656	609	BTH_II0374	282	BTH_II0441	602
BTH_II1334	587	BTH_II1567	582	BTH_II0949	221	BTH_I0026	571
BTH_II1574	396	BTH_II0066	684	BTH_II1654	486	BTH_II1572	686
BTH_I2975	103	BTH_I1573	529	BTH_I2287	336	BTH_I3292	573
BTH_I1096	288	BTH_I1099	622	BTH_I2837	478	BTH_II0028	590
BTH_I2572	492	BTH_I0483	640	BTH_II1715	291	BTH_II2233	582
BTH_II0864	642	BTH_I2409	628	No match	-	BTH_I2394	703
No match	-	BTH_I3072	615	BTH_II0577	721	BTH_I1643	737
BTH_I2253	668	BTH_I3055	581	BTH_II2154	623	BTH_I2795	683
BTH_II0708	395	BTH_I0181	607	BTH_I2023	723	No match	-
BTH_II0941	367	BTH_II1163	669	BTH_I0418	701	BTH_I1574	692
BTH_I1096	491	BTH_I2037	623	BTH_II2207	174	BTH_II1973	663
BTH_I1560	502	BTH_I1096	648	BTH_II0100	472	BTH_I0378	608
BTH_I2684	104	BTH_I1265	637	BTH_I2803	582	BTH_I1318	602
BTH_I0539	288	BTH_I1099	626	BTH_II2371	626	BTH_I2951	663
BTH_I0553	672	BTH_I0882	595	BTH_II1976	561	BTH_I1117	672
BTH_I1096	640	BTH_II0528	633	BTH_II1531	589	BTH_II1570	652
BTH_I1112	696	BTH_I1073	594	BTH_II0536	563	BTH_I3016	649
BTH_I1096	384	BTH_II1226	667	BTH_I0810	375	BTH_II1631	532
BTH_I1175	357	BTH_I0949	692	BTH_II1756	374	BTH_I3151	706
BTH_I1099	552	BTH_II1135	626	BTH_II2071	501	BTH_I1487	561
BTH_II0665	598	BTH_I3003	647	BTH_II1105	466	BTH_II0380	592
BTH_I2925	462	BTH_I1227	658	BTH_I2804	462	BTH_II2039	639
BTH_I0366	476	BTH_I1096	655	BTH_I0001	682	BTH_I3025	673
BTH_II1834	592	BTH_I1099	641	BTH_II2040	703	BTH_I1033	662
BTH_I2230	364	No match	-	BTH_I1942	220	BTH_II2108	677

BTH_I12149	330	BTH_I1663	703	BTH_I0819	265	BTH_I10020	593
BTH_I12183	369	BTH_I1938	736	BTH_I2092	195	BTH_I1195	502
BTH_I2618	472	BTH_I10562	593	No match	-	BTH_I3079	690
BTH_I11468	581	BTH_I11225	802	BTH_I11586	301	BTH_I0415	717
BTH_I1099	390	BTH_I2038	745	BTH_I10047	699	BTH_I2367	599

^aApproximately, 4% of the cDNA clones did not match to any sequences in Genbank.

^bSequences that match the rRNAs (BTH_I1096, 16S rRNA; BTH_I1099, 23S rRNA) or tRNA (BTH_I1741, tRNA-Gly) are highlighted in red.

^cLength of sequences that match continuously to sequences of the *B. thailandensis* genome.