

Supplementary Table 10. 42 scaffolds (derived from unmapped reads) were matched to the EST database but unmatched to the NR-nucleotide database

rank	contig_id	pscore	query_length	target_length	target_accession
1	C53546 6.0	2.86E-24	301	609	84588596
1	C53766 6.0	8.91E-77	304	653	18817627
1	C53830 5.0	2.30745E-06	304	651	112589012
1	C53974 6.0	1.42E-149	306	596	50529253
1	C54826 8.0	5.08E-29	316	551	50449893
1	C55664 4.0	8.02E-99	326	443	50440720
1	C55674 7.0	3.39E-67	326	551	50449893
1	C56050 7.0	9.95762E-06	330	557	50528320
1	C56702 7.0	2.32E-179	340	561	50526540
1	C58026 7.0	1.81E-118	359	641	60541493
1	C59870 8.0	6.64E-69	389	609	84588596
1	C59878 5.0	8.51E-16	389	863	84586154
1	C60164 6.0	1.54E-134	394	569	60585400
1	C60562 10.0	1.86E-38	401	582	50449823
1	C60650 4.0	7.89021E-07	403	658	18823537
1	C61634 7.0	1.41E-48	422	768	84584849
1	C63244 7.0	1.66E-54	459	263	84589393
1	C63536 6.0	2.30E-115	467	923	84589511
1	C64908 9.0	0	507	664	60563913
1	C65602 4.0	1.01E-40	528	501	15911712
1	C65648 7.0	2.59E-78	530	609	84588596
1	C65718 4.0	1.47E-178	533	329	60582512
1	scaffold48 7.5	1.58E-110	540	483	18824422
1	C66232 10.0	9.88E-72	550	840	84588081
1	C66268 9.0	7.21E-156	551	618	50532189
1	C66792 7.0	1.02E-34	573	653	18817627
1	C67146 7.0	3.85E-28	588	609	84588596
1	C68202 9.0	1.75E-104	639	551	50449893
1	C68702 4.0	0	672	860	84588879
1	C68814 10.0	9.38134E-14	680	406	60527815
1	C69100 6.0	7.46E-30	698	609	84588596
1	C69258 5.0	0	708	673	60539614
1	C69624 9.0	5.17E-142	737	277	60635979
1	C69972 10.0	6.33E-46	764	443	50440720
1	C70632 7.0	2.89E-85	841	620	84589119
1	C70970 7.0	6.72E-111	890	412	62146151
1	C72338 9.0	3.93939E-11	1155	609	84588596
1	C72362 7.0	2.46E-46	1162	412	62146151
1	C72408 67.0	1.43E-75	1174	900	84586140
1	C72662 9.0	2.05E-25	1257	181	60636521
1	C73018 55.0	4.66E-42	1456	620	84589119
1	C73228 7.0	2.36E-16	1613	609	84588596