

-	wf_da	2 items
-	uclust_picked_otus	4 items
-	rep_set	4 items
-	pynast_aligned_seqs	5 items
-	fasttree_phylogeny	2 items
	seqs_rep_set.tre	54.7 KB
	seqs_rep_set_phylogeny.log	173 bytes
	seqs_rep_set_aligned.fasta	15.1 MB
	seqs_rep_set_aligned_pfiltered.fasta	1.5 MB
	seqs_rep_set_failures.fasta	10.3 KB
	seqs_rep_set_log.txt	81.5 KB
-	rdp_assigned_taxonomy	3 items
-	otu_table	1 item
	seqs_otu_table.txt	301.5 KB
	seqs_rep_set_tax_assignments.log	401 bytes
	seqs_rep_set_tax_assignments.txt	187.0 KB
	seqs_rep_set.fasta	545.2 KB
	seqs_rep_set.log	257 bytes
	seqs_clusters.uc	55.0 MB
	seqs_otus.log	399 bytes
	seqs_otus.txt	4.1 MB
	log_20110823081355.txt	4.0 KB