Quality control data for the 7 species demonstrated

LSTrAP-Kingdom automatically generates:

- 1. A summary of recommended quality control (QC) thresholds for each species (<u>Table 1A</u>)
- 2. A matrix of the number of experiments passing QC for combinations of thresholds used (<u>Table 1B</u> shows the example matrix for taxid3218)
- 3. Jointplots visualizing the quality of experiments in each species (Table 1C)

The user can edit the QC thresholds in Table 1A (example in Table 2A) and pass it back to LSTrAP-Kingdom, which will then update the following:

1. Updated jointplots visualizing the quality of experiments in each species, with labels updated for the new thresholds (<u>Table 2B</u>)

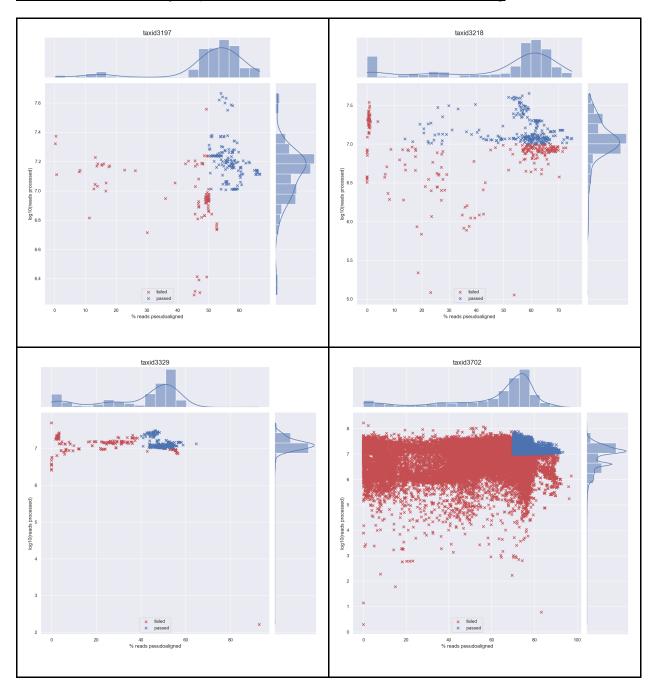
<u>Table 1A: An automatically generated summary of recommended quality control (QC) thresholds for each species</u>

	log10(processed) cutoff	% pseudoaligned cutoff	Number of Run IDs passed
3197	7	50	144
3218	7	10	252
3329	7	40	254
3702	7	70	11795
4081	7	20	4086
4530	7	50	5681
4577	7	70	1578

Table 1B: Example of matrix of different combinations of cutoffs for log10(processed) and % pseudoaligned for taxid3218

	% pseudoaligned								
log10(number processed)	10	20	30	40	50	60	70	80	90
6	227	212	210	207	152	26	0	0	0
6.2	227	212	210	207	152	26	0	0	0
6.4	223	208	206	203	152	26	0	0	0
6.6	221	206	204	201	152	26	0	0	0
6.8	216	201	199	197	148	26	0	0	0
7	175	162	160	159	144	26	0	0	0
7.2	74	73	73	73	67	1	0	0	0
7.4	9	9	9	9	8	0	0	0	0
7.6	4	4	4	4	4	0	0	0	0
7.8	0	0	0	0	0	0	0	0	0
8	0	0	0	0	0	0	0	0	0

Table 1C: Generated jointplots for QC for visualization before editting



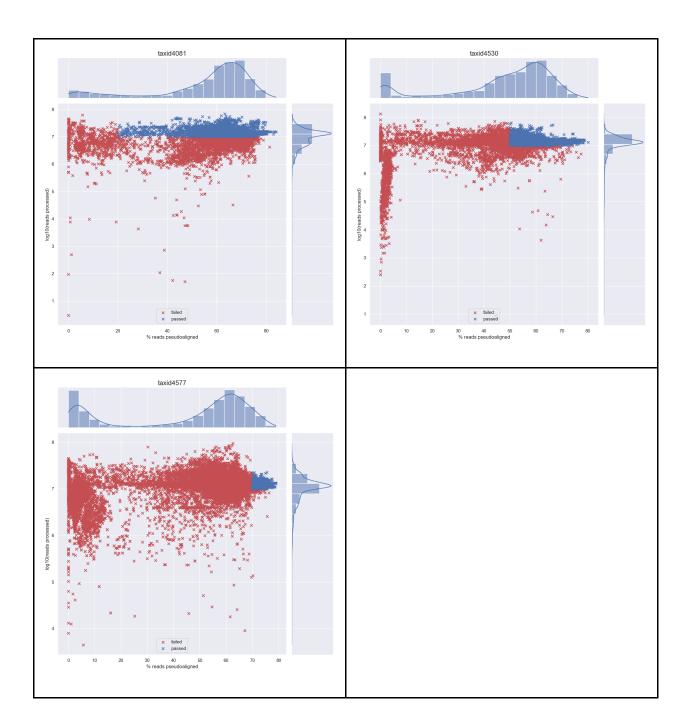


Table 2A: User edited summary of recommended quality control (QC) thresholds for each species

	log10(processed) cutoff	% pseudoaligned cutoff	Number of Run IDs passed
3197	7	50	144
3218	6.5	55	318
3329	6.8	42	304
3702	7	80	2117
4081	6.4	70	1416
4530	6	65	1449
4577	7	70	1578

Table 2B: Updated jointplots for QC for visualization after user had edited the thresholds

