Summarized results of (A) INDUS (B) TACOA (C) SORT-ITEMS (D) MEGAN and (E) SPHINX, obtained for simLC, simMC, and simHC data sets. All numbers given in this table are with respect to the total number of sequences in the respective data set.

(A) INDUS

	Percentage of Sequences Assigned					
Assignment Category/Taxon	SimLC		SimMC		SimHC	
	Complete RefDB®	Modified RefDB ^s	Complete RefDB [®]	Modified RefDB ^s	Complete RefDB [@]	Modified RefDB ^s
Non-Specific* Levels	8.16	14.75	8.5	14.03	13.06	15.25
Specific [#] Levels	81.38	66.75	84.26	72.12	76.58	63.65
Phylum	15.61	21.51	14.85	19.82	13.06	13.53
Class	4.73	12.77	7.2	14.55	6.21	9.1
Order	3.24	7.08	3.81	7.56	2.9	3.39
Family	8.87	6.27	10.42	7.97	10.69	8.03
Genus & Lower Levels	48.93	19.12	47.98	22.21	43.72	29.6
Total Correct	89.54	81.49	92.76	86.15	89.64	78.91
Wrong	2.61	7.28	1	4.51	1.89	5.8
Unassigned	7.85	11.23	6.24	9.34	8.47	15.29

(B) TACOA

	Percentage of Sequences Assigned					
Assignment Category/Taxon	SimLC		SimMC		SimHC	
	Complete RefDB®	Modified RefDB ^s	Complete RefDB [@]	Modified RefDB ^s	Complete RefDB [@]	Modified RefDB ^s
Non-Specific* Levels	54.9	53.5	58.8	56.0	53.4	48.8
Specific [#] Levels	24.4	21.0	23.7	23.0	18.7	18.2
Phylum	19.7	15.3	19.0	16.2	11.3	8.9
Class	0.9	1.6	1.3	2.6	1.6	2.6
Order	0.8	0.9	0.8	1.1	0.9	1.2
Family	0.04	1.04	0.02	1.0	0.1	1.1
Genus & Lower Levels	2.89	2.13	2.6	2.1	4.8	4.5
Total Correct	79.3	74.4	82.5	79.0	72.1	67.0
Wrong	9.0	13.9	7.3	10.8	11.4	16.0
Unassigned	11.6	11.6	10.2	10.2	16.5	17.0

*Non-specific: Assignments made at super kingdom level

[#]Specific : Assignments made at phylum, class, order, family and genus levels

[®]Complete RefDB: Complete reference database constructed from sequences belonging to 952 genomes

^sModified RefDB: Modified reference database created by removing sequences of 300 genomes from the complete reference database

Note that the subtotals may vary by a value of 0.1, since the individual values are rounded off to single decimals.

(C) SOrt-ITEMS

	Percentage of Sequences Assigned					
Assignment Category/Taxon	SimLC		SimMC		SimHC	
	Complete RefDB®	Modified RefDB ^s	Complete RefDB [@]	Modified RefDB ^s	Complete RefDB [@]	Modified RefDB ^s
Non-Specific* Levels	0.0	0.0	0.0	0.0	0.0	0.0
Specific [#] Levels	94.9	78.1	93.7	79.1	92.0	78.1
Phylum	2.4	10.5	2.3	8.8	3.7	9.1
Class	2.6	12.4	3.0	10.9	4.0	8.4
Order	0.2	0.1	0.2	0.1	0.3	0.1
Family	5.0	17.1	5.3	16.5	6.4	7.8
Genus & Lower Levels	84.6	37.9	83.0	42.9	77.6	52.7
Total Correct	94.9	78.1	93.7	79.1	92.0	78.1
Wrong	2.0	9.2	3.1	9.1	3.6	9.2
Unassigned	3.1	12.6	3.2	11.8	4.3	12.6

(D) MEGAN

	Percentage of Sequences Assigned						
Assignment Category/Taxon	SimLC		SimMC		SimHC		
	Complete RefDB®	Modified RefDB ^s	Complete RefDB [@]	Modified RefDB ^s	Complete RefDB [@]	Modified RefDB ^s	
Non-Specific* Levels	1.1	7.6	1.0	6.7	1.5	8.4	
Specific [#] Levels	93.9	65.0	93.1	63.2	90.3	69.5	
Phylum	0.8	5.3	0.7	4.6	3.7	3.9	
Class	0.6	2.7	0.6	2.7	4.0	2.2	
Order	1.4	4.1	1.2	3.3	0.3	1.6	
Family	5.4	13.4	5.4	11.1	4.4	2.9	
Genus & Lower Levels	85.8	39.6	85.1	41.5	77.9	58.9	
Total Correct	95.0	72.6	94.0	69.9	91.9	75.9	
Wrong	2.7	23.6	3.5	26.6	4.9	17.6	
Unassigned	2.3	3.7	2.4	3.5	3.2	4.5	

*Non-specific: Assignments made at super kingdom level #Specific : Assignments made at phylum, class, order, family and genus levels @Complete RefDB: Complete reference database constructed from sequences belonging to 952 genomes

^{\$}Modified RefDB: Modified reference database created by removing sequences of 300 genomes from the complete reference database

Note that the subtotals may vary by a value of 0.1, since the individual values are rounded off to single decimals.

	Percentage of Sequences Assigned					
Assignment Category/Taxon	SimLC		SimMC		SimHC	
	Complete RefDB®	Modified RefDB ^s	Complete RefDB [®]	Modified RefDB ^s	Complete RefDB [®]	Modified RefDB ^s
Non-Specific* Levels	10.3	12.7	11.5	11.6	6.4	6.4
Specific [#] Levels	82.8	71.0	81.1	70.1	79.6	71.2
Phylum	4.1	13.7	5.9	17.4	5.1	12.0
Class	5.4	14.6	4.4	16.2	6.1	13.9
Order	0.8	0.2	0.5	0.1	0	0.5
Family	14.6	17.4	16.1	12.5	18.2	17.2
Genus & Lower Levels	57.9	25.1	54.2	23.9	50.2	27.6
Total Correct	93.1	83.7	92.6	81.8	86.0	77.6
Wrong	2.8	11.5	3.6	13.5	7.1	12.1
Unassigned	4.1	4.8	3.8	4.7	6.9	10.3

*Non-specific: Assignments made at super kingdom level #Specific : Assignments made at phylum, class, order, family and genus levels @Complete RefDB: Complete reference database constructed from sequences belonging to 952 genomes

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Note that the subtotals may vary by a value of 0.1, since the individual values are rounded off to single decimals.