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table 3: evaluation of concordance - non-metric dental traits  
frequencies

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the table presents data concerning the evaluation of discrepancies in the observation of non-metric dental traits

these data are organized as follows:

three summary tables by variable indicate the degree of disagreement in recording between two sessions: O1a-O1b (intra-observer discrepancy), O1a-O2 and O1b-O2 (inter-observer discrepancy)

assessment differences are termed error and the associated number indicates the amplitude of disagreement (number of grades), thus:

error 0 indicates that the observations are identical between the two sessions

error 1 indicates that one grade of difference was observed

error 2 indicates two grades of difference, etc.

OSO (one session only) error indicates that a trait was recorded during only one of the two sessions

- the first table presents the raw counts of discrepancy between sessions  
all types of discrepancy are noted here, these can be a difference in attribution of grade or recording of a trait (OSO)

- the second table indicates the error percentages calculated from the first table

- the last table indicates the percentage of errors in attribution of grades during observation  
calculations are thus based on all errors, excluding the OSO error

to keep the uniformity of the tables, these latter all have the same number of types of error (including, for example, dichotomous traits): this is why the real gradation of each variable is shown by yellow fill enabling us to make the difference between the absence of discrepancies and the absence of grades for each trait

table 3

## evaluation concordance - non-metric dental traits

frequencies

T1 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	119	116	118
error 1	1	1	0
error 2	1	1	1
error 3	1	1	0
error 4			
error 5			
error 6			
error oso	30	33	33
%0	78.29	76.32	77.63
%1	0.66	0.66	0.00
%2	0.66	0.66	0.66
%3	0.66	0.66	0.00
%4			
%5			
%6			
%oso	19.74	21.71	21.71
%0	97.54	97.48	99.16
%1	0.82	0.84	0.00
%2	0.82	0.84	0.84
%3	0.82	0.84	0.00
%4			
%5			
%6			

T10 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	80	42	53
error 1	8	29	31
error 2	0	8	12
error 3	0	0	0
error 4	0	0	0
error 5			
error 6			
error oso	19	28	11
%0	74.77	39.25	49.53
%1	7.48	27.10	28.97
%2	0.00	7.48	11.21
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5			
%6			
%oso	17.76	26.17	10.28
%0	90.91	53.16	55.21
%1	9.09	36.71	32.29
%2	0.00	10.13	12.50
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5			
%6			

T12 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	51	48	63
error 1	1	1	0
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	36	39	25
%0	57.95	54.55	71.59
%1	1.14	1.14	0.00
%2			
%3			
%4			
%5			
%6			
%oso	40.91	44.32	28.41
%0	98.08	97.96	100.00
%1	1.92	2.04	0.00
%2			
%3			
%4			
%5			
%6			

T9 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	28	17	23
error 1	0	0	0
error 2	0	7	6
error 3	0	0	0
error 4	0	0	0
error 5			
error 6			
error oso	11	15	10
%0	71.79	43.59	58.97
%1	0.00	0.00	0.00
%2	0.00	17.95	15.38
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5			
%6			
%oso	28.21	38.46	25.64
%0	100.00	70.83	79.31
%1	0.00	0.00	0.00
%2	0.00	29.17	20.69
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5			
%6			

T11 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	101	89	89
error 1	0	2	2
error 2	0	1	1
error 3	0	0	0
error 4	0	0	0
error 5			
error 6			
error oso	0	9	9
%0	100.00	88.12	88.12
%1	0.00	1.98	1.98
%2	0.00	0.99	0.99
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5			
%6			
%oso	0.00	8.91	8.91
%0	100.00	96.74	96.74
%1	0.00	2.17	2.17
%2	0.00	1.09	1.09
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5			
%6			

T13 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	72	60	74
error 1	9	17	19
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	20	24	8
%0	71.29	59.41	73.27
%1	8.91	16.83	18.81
%2			
%3			
%4			
%5			
%6			
%oso	19.80	23.76	7.92
%0	88.89	77.92	79.57
%1	11.11	22.08	20.43
%2			
%3			
%4			
%5			
%6			

table 3

## evaluation concordance - non-metric dental traits

frequencies

T20 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	28	16	18
error 1	0	0	0
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	4	16	14
%0	87.50	50.00	56.25
%1	0.00	0.00	0.00
%2			
%3			
%4			
%5			
%6			
%oso	12.50	50.00	43.75
%0	100.00	100.00	100.00
%1	0.00	0.00	0.00
%2			
%3			
%4			
%5			
%6			

T21 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	59	34	45
error 1	2	2	0
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	9	34	25
%0	84.29	48.57	64.29
%1	2.86	2.86	0.00
%2			
%3			
%4			
%5			
%6			
%oso	12.86	48.57	35.71
%0	96.72	94.44	100.00
%1	3.28	5.56	0.00
%2			
%3			
%4			
%5			
%6			

T23 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	136	119	131
error 1	0	0	0
error 2	0	0	0
error 3	0	0	0
error 4	0	1	1
error 5	0	0	0
error 6	0	0	0
error oso	12	28	16
%0	91.89	80.41	88.51
%1	0.00	0.00	0.00
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4	0.00	0.68	0.68
%5	0.00	0.00	0.00
%6	0.00	0.00	0.00
%oso	8.11	18.92	10.81
%0	100.00	99.17	99.24
%1	0.00	0.00	0.00
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4	0.00	0.83	0.76
%5	0.00	0.00	0.00
%6	0.00	0.00	0.00

T24 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	108	90	120
error 1	4	9	10
error 2	1	3	3
error 3	0	1	0
error 4	0	1	1
error 5	0	0	0
error 6	0	0	0
error oso	42	51	21
%0	69.68	58.06	77.42
%1	2.58	5.81	6.45
%2	0.65	1.94	1.94
%3	0.00	0.65	0.00
%4	0.00	0.65	0.65
%5	0.00	0.00	0.00
%6	0.00	0.00	0.00
%oso	27.10	32.90	13.55
%0	95.58	86.54	89.55
%1	3.54	8.65	7.46
%2	0.88	2.88	2.24
%3	0.00	0.96	0.00
%4	0.00	0.96	0.75
%5	0.00	0.00	0.00
%6	0.00	0.00	0.00

T25 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	112	76	99
error 1	11	41	42
error 2	5	8	3
error 3	0	1	1
error 4	1	1	0
error 5	0	0	0
error 6	1	1	0
error oso	27	29	12
%0	71.34	48.41	63.06
%1	7.01	26.11	26.75
%2	3.18	5.10	1.91
%3	0.00	0.64	0.64
%4	0.44	0.64	0.00
%5	0.00	0.00	0.00
%6	0.64	0.64	0.00
%oso	17.20	18.47	7.64
%0	86.15	59.38	68.28
%1	8.46	32.03	28.97
%2	3.85	6.25	2.07
%3	0.00	0.78	0.69
%4	0.77	0.78	0.00
%5	0.00	0.00	0.00
%6	0.77	0.78	0.00

T26 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	96	62	100
error 1	13	30	22
error 2	9	12	6
error 3	2	5	2
error 4	4	4	2
error 5	1	1	0
error 6	1	1	1
error oso	26	37	19
%0	63.16	40.79	65.79
%1	8.55	19.74	14.47
%2	5.92	7.89	3.95
%3	1.32	3.29	1.32
%4	1.85	2.63	1.32
%5	0.66	0.66	0.00
%6	0.66	0.66	0.66
%oso	17.11	24.34	12.50
%0	76.19	53.91	75.19
%1	10.32	26.09	16.54
%2	7.14	10.43	4.51
%3	1.59	4.35	1.50
%4	3.17	3.48	1.50
%5	0.79	0.87	0.00
%6	0.79	0.87	0.75

table 3

## evaluation concordance - non-metric dental traits

frequencies

T33 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	149	142	137
error 1	0	1	0
error 2	0	1	2
error 3	1	0	0
error 4			
error 5			
error 6			
error oso	13	19	24
%0	91.41	87.12	84.05
%1	0.00	0.61	0.00
%2	0.00	0.61	1.23
%3	0.61	0.00	0.00
%4			
%5			
%6			
%oso	7.98	11.66	14.72
%0	99.33	98.61	98.56
%1	0.00	0.69	0.00
%2	0.00	0.69	1.44
%3	0.67	0.00	0.00
%4			
%5			
%6			

T44 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	100	102	99
error 1	2	0	2
error 2	0	0	0
error 3	0	0	0
error 4	0	0	0
error 5	0	0	0
error 6	0	0	0
error oso	1	1	2
%0	97.09	99.03	96.12
%1	1.94	0.00	1.94
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6	0.00	0.00	0.00
%oso	0.97	0.97	1.94
%0	98.04	100.00	98.02
%1	1.96	0.00	1.98
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6	0.00	0.00	0.00

T46 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	80	55	59
error 1	0	0	0
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	6.00	31.00	27.00
%0	93.02	63.95	68.60
%1	0.00	0.00	0.00
%2			
%3			
%4			
%5			
%6			
%oso	6.98	36.05	31.40
%0	100.00	100.00	100.00
%1	0.00	0.00	0.00
%2			
%3			
%4			
%5			
%6			

T43 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	102	98	99
error 1	1	5	4
error 2	0	0	0
error 3	0	0	0
error 4			
error 5			
error 6			
error oso	0	0	0
%0	99.03	95.15	96.12
%1	0.97	4.85	3.88
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4			
%5			
%6			
%oso	0.00	0.00	0.00
%0	99.03	95.15	96.12
%1	0.97	4.85	3.88
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4			
%5			
%6			

T45 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	31	24	30
error 1	0	0	0
error 2	0	0	0
error 3	0	2	2
error 4	0	0	0
error 5	0	0	0
error 6			
error oso	8	13	7
%0	79.49	61.54	76.92
%1	0.00	0.00	0.00
%2	0.00	0.00	0.00
%3	0.00	5.13	5.13
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6			
%oso	20.51	33.33	17.95
%0	100.00	92.31	93.75
%1	0.00	0.00	0.00
%2	0.00	0.00	0.00
%3	0.00	7.69	6.25
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6			

T47 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	75	30	37
error 1	6	28	24
error 2	2	4	4
error 3	0	0	0
error 4	0	3	3
error 5	1	1	0
error 6	0	0	0
error oso	2	19	17
%0	86.21	34.48	42.53
%1	6.90	32.18	27.59
%2	2.30	4.60	4.60
%3	0.00	0.00	0.00
%4	0.00	3.45	3.45
%5	1.15	1.15	0.00
%6	0.00	0.00	0.00
%oso	2.30	21.84	19.54
%0	88.24	44.12	52.86
%1	7.06	41.18	34.29
%2	2.35	5.88	5.71
%3	0.00	0.00	0.00
%4	0.00	4.41	4.29
%5	1.18	1.47	0.00
%6	0.00	0.00	0.00

table 3

## evaluation concordance - non-metric dental traits

frequencies

T54 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	129	106	100
error 1	8	15	22
error 2	2	2	1
error 3	0	1	1
error 4	0	0	0
error 5	0	0	0
error 6			
error oso	0	15	15
%0	92.81	76.26	71.94
%1	5.76	10.79	15.83
%2	1.44	1.44	0.72
%3	0.00	0.72	0.72
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6			
%oso	0.00	10.79	10.79
%0	92.81	85.48	80.65
%1	5.76	12.10	17.74
%2	1.44	1.61	0.81
%3	0.00	0.81	0.81
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6			

T56 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	142	123	123
error 1	0	0	0
error 2	0	0	0
error 3	0	0	0
error 4	0	0	0
error 5	0	0	0
error 6			
error oso	0	19	19
%0	100.00	86.62	86.62
%1	0.00	0.00	0.00
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6			
%oso	0.00	13.38	13.38
%0	100.00	100.00	100.00
%1	0.00	0.00	0.00
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6			

T58 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	110	86	86
error 1	0	1	1
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	0	23	23
%0	100.00	78.18	78.18
%1	0.00	0.91	0.91
%2			
%3			
%4			
%5			
%6			
%oso	0.00	20.91	20.91
%0	100.00	98.85	98.85
%1	0.00	1.15	1.15
%2			
%3			
%4			
%5			
%6			

T55 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	134	90	97
error 1	3	12	12
error 2	0	4	4
error 3	0	3	3
error 4	1	0	0
error 5	1	3	1
error 6			
error oso	5	32	27
%0	93.06	62.50	67.36
%1	2.08	8.33	8.33
%2	0.00	2.78	2.78
%3	0.00	2.08	2.08
%4	0.69	0.00	0.00
%5	0.69	2.08	0.69
%6			
%oso	3.47	22.22	18.75
%0	96.40	80.36	82.91
%1	2.16	10.71	10.26
%2	0.00	3.57	3.42
%3	0.00	2.68	2.56
%4	0.72	0.00	0.00
%5	0.72	2.68	0.85
%6			

T57 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	146	131	131
error 1	0	0	0
error 2	0	0	0
error 3	0	0	0
error 4	0	0	0
error 5	0	0	0
error 6			
error oso	0	15	15
%0	100.00	89.73	89.73
%1	0.00	0.00	0.00
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6			
%oso	0.00	10.27	10.27
%0	100.00	100.00	100.00
%1	0.00	0.00	0.00
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6			

T59 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	139	122	125
error 1	0	0	0
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	3	20	17
%0	97.89	85.92	88.03
%1	0.00	0.00	0.00
%2			
%3			
%4			
%5			
%6			
%oso	2.11	14.08	11.97
%0	100.00	100.00	100.00
%1	0.00	0.00	0.00
%2			
%3			
%4			
%5			
%6			

table 3

## evaluation concordance - non-metric dental traits

frequencies

T14 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	91	89	105
error 1	1	2	0
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	60	61	47
%0	59.87	58.55	69.08
%1	0.66	1.32	0.00
%2			
%3			
%4			
%5			
%6			
%oso	39.47	40.13	30.92
%0	98.91	97.80	100.00
%1	1.09	2.20	0.00
%2			
%3			
%4			
%5			
%6			

T15 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	88	46	75
error 1	6	38	34
error 2	3	4	4
error 3	0	0	0
error 4	0	0	0
error 5	0	0	0
error 6	0	0	0
error oso	31	40	15
%0	68.75	35.94	58.59
%1	4.69	29.69	26.56
%2	2.34	3.13	3.13
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6	0.00	0.00	0.00
%oso	24.22	31.25	11.72
%0	90.72	52.27	66.37
%1	6.19	43.18	30.09
%2	3.09	4.55	3.54
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6	0.00	0.00	0.00

T16 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	127	71	71
error 1	1	10	10
error 2	0	18	18
error 3	0	0	0
error 4	0	0	0
error 5	0	2	2
error 6	0	0	0
error oso	0	27	27
%0	99.22	55.47	55.47
%1	0.78	7.81	7.81
%2	0.00	14.06	14.06
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5	0.00	1.56	1.56
%6	0.00	0.00	0.00
%oso	0.00	21.09	21.09
%0	99.22	70.30	70.30
%1	0.78	9.90	9.90
%2	0.00	17.82	17.82
%3	0.00	0.00	0.00
%4	0.00	0.00	0.00
%5	0.00	1.98	1.98
%6	0.00	0.00	0.00

T17 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	172	153	174
error 1	6	8	9
error 2	1	5	4
error 3	2	1	2
error 4	0	0	0
error 5	0	0	0
error 6	0	0	0
error oso	34	48	26
%0	80.00	71.16	80.93
%1	2.79	3.72	4.19
%2	0.47	2.33	1.86
%3	0.93	0.47	0.93
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6	0.00	0.00	0.00
%oso	15.81	22.33	12.09
%0	95.03	91.62	92.06
%1	3.31	4.79	4.76
%2	0.55	2.99	2.12
%3	1.10	0.60	1.06
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6	0.00	0.00	0.00

T18 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	67	44	50
error 1	3	6	4
error 2	1	0	0
error 3	1	2	0
error 4			
error 5			
error 6			
error oso	10	30	28
%0	81.71	53.66	60.98
%1	3.66	7.32	4.88
%2	1.22	0.00	0.00
%3	1.22	2.44	0.00
%4			
%5			
%6			
%oso	12.20	36.59	34.15
%0	93.06	84.62	92.59
%1	4.17	11.54	7.41
%2	1.39	0.00	0.00
%3	1.39	3.85	0.00
%4			
%5			
%6			

T19 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	23	11	12
error 1	1	0	0
error 2	1	1	1
error 3	0	4	6
error 4	0	0	0
error 5	0	0	0
error 6			
error oso	5	14	11
%0	76.67	36.67	40.00
%1	3.33	0.00	0.00
%2	3.33	3.33	3.33
%3	0.00	13.33	20.00
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6			
%oso	16.67	46.67	36.67
%0	92.00	68.75	63.16
%1	4.00	0.00	0.00
%2	4.00	6.25	5.26
%3	0.00	25.00	31.58
%4	0.00	0.00	0.00
%5	0.00	0.00	0.00
%6			

table 3

## evaluation concordance - non-metric dental traits

frequencies

T27 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	118	93	93
error 1	0	4	4
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	0	21	21
%0	100.00	78.81	78.81
%1	0.00	3.39	3.39
%2			
%3			
%4			
%5			
%6			
%oso	0.00	17.80	17.80
%0	100.00	95.88	95.88
%1	0.00	4.12	4.12
%2			
%3			
%4			
%5			
%6			

T29 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	101	65	65
error 1	0	1	1
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	0	35	35
%0	100.00	64.36	64.36
%1	0.00	0.99	0.99
%2			
%3			
%4			
%5			
%6			
%oso	0.00	34.65	34.65
%0	100.00	98.48	98.48
%1	0.00	1.52	1.52
%2			
%3			
%4			
%5			
%6			

T31 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	99	65	65
error 1	0	1	1
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	0	33	33
%0	100.00	65.66	65.66
%1	0.00	1.01	1.01
%2			
%3			
%4			
%5			
%6			
%oso	0.00	33.33	33.33
%0	100.00	98.48	98.48
%1	0.00	1.52	1.52
%2			
%3			
%4			
%5			
%6			

T28 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	101	71	71
error 1	5	10	10
error 2	4	4	4
error 3	1	3	3
error 4	1	1	1
error 5	0	1	1
error 6			
error oso	35	57	57
%0	68.71	48.30	48.30
%1	3.40	6.80	6.80
%2	2.72	2.72	2.72
%3	0.68	2.04	2.04
%4	0.44	0.68	0.68
%5	0.00	0.68	0.68
%6			
%oso	23.81	38.78	38.78
%0	90.18	78.89	78.89
%1	4.46	11.11	11.11
%2	3.57	4.44	4.44
%3	0.89	3.33	3.33
%4	0.89	1.11	1.11
%5	0.00	1.11	1.11
%6			

T30 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	95	47	47
error 1	0	0	0
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	0	48	48
%0	100.00	49.47	49.47
%1	0.00	0.00	0.00
%2			
%3			
%4			
%5			
%6			
%oso	0.00	50.53	50.53
%0	100.00	100.00	100.00
%1	0.00	0.00	0.00
%2			
%3			
%4			
%5			
%6			

T32 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	98	61	61
error 1	0	3	3
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	0	34	34
%0	100.00	62.24	62.24
%1	0.00	3.06	3.06
%2			
%3			
%4			
%5			
%6			
%oso	0.00	34.69	34.69
%0	100.00	95.31	95.31
%1	0.00	4.69	4.69
%2			
%3			
%4			
%5			
%6			



table 3

## evaluation concordance - non-metric dental traits

frequencies

T48 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	18	14	15
error 1	0	0	0
error 2	0	0	0
error 3	0	0	0
error 4			
error 5			
error 6			
error oso	1.00	5.00	4.00
<hr/>			
%0	94.74	73.68	78.95
%1	0.00	0.00	0.00
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4			
%5			
%6			
%oso	5.26	26.32	21.05
<hr/>			
%0	100.00	100.00	100.00
%1	0.00	0.00	0.00
%2	0.00	0.00	0.00
%3	0.00	0.00	0.00
%4			
%5			
%6			

T50 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	108	90	93
error 1	0	1	1
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	9	26	23
<hr/>			
%0	92.31	76.92	79.49
%1	0.00	0.85	0.85
%2			
%3			
%4			
%5			
%6			
%oso	7.69	22.22	19.66
<hr/>			
%0	100.00	98.90	98.94
%1	0.00	1.10	1.06
%2			
%3			
%4			
%5			
%6			

T52 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	140	119	122
error 1	2	10	8
error 2	0	0	0
error 3			
error 4			
error 5			
error 6			
error oso	1	14	13
<hr/>			
%0	97.90	83.22	85.31
%1	1.40	6.99	5.59
%2	0.00	0.00	0.00
%3			
%4			
%5			
%6			
%oso	0.70	9.79	9.09
<hr/>			
%0	98.59	92.25	93.85
%1	1.41	7.75	6.15
%2	0.00	0.00	0.00
%3			
%4			
%5			
%6			

T49 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	19	6	11
error 1	0	3	3
error 2	0	0	0
error 3	0	1	1
error 4	0	0	0
error 5			
error 6			
error oso	7.00	16.00	11.00
<hr/>			
%0	73.08	23.08	42.31
%1	0.00	11.54	11.54
%2	0.00	0.00	0.00
%3	0.00	3.85	3.85
%4	0.00	0.00	0.00
%5			
%6			
%oso	26.92	61.54	42.31
<hr/>			
%0	100.00	60.00	73.33
%1	0.00	30.00	20.00
%2	0.00	0.00	0.00
%3	0.00	10.00	6.67
%4	0.00	0.00	0.00
%5			
%6			

T51 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	110	90	90
error 1	0	1	1
error 2			
error 3			
error 4			
error 5			
error 6			
error oso	4	23	23
<hr/>			
%0	96.49	78.95	78.95
%1	0.00	0.88	0.88
%2			
%3			
%4			
%5			
%6			
%oso	3.51	20.18	20.18
<hr/>			
%0	100.00	98.90	98.90
%1	0.00	1.10	1.10
%2			
%3			
%4			
%5			
%6			

T53 concordance

	O1a-O1b	O1a-O2	O1b-O2
error 0	101	66	66
error 1	1	11	12
error 2	1	7	8
error 3			
error 4			
error 5			
error 6			
error oso	6	25	23
<hr/>			
%0	92.66	60.55	60.55
%1	0.70	7.69	8.39
%2	0.70	4.90	5.59
%3			
%4			
%5			
%6			
%oso	4.20	17.48	16.08
<hr/>			
%0	98.06	78.57	76.74
%1	0.97	13.10	13.95
%2	0.97	8.33	9.30
%3			
%4			
%5			
%6			

