

# Comparing Age Assignments

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## Preliminaries

```
> library(FSA) # for ageBias(), agePrecision()
```

## Loading Data

```
> SB <- read.csv("StripedBass4.csv") # appropriately set the working directory before this
> str(SB)
'data.frame': 1202 obs. of 2 variables:
 $ reader1: int 2 2 2 2 2 2 2 2 2 2 ...
 $ reader2: int 2 2 2 2 2 2 2 2 2 2 ...
```

## Examine Age Bias

```
> ab <- ageBias(reader2~reader1,data=SB)
```

```
> summary(ab,what="table",flip.table=TRUE)
```

	reader1																			
reader2	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	
20	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	1	-	
19	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	
18	-	-	-	-	-	-	-	-	-	-	-	1	-	-	-	-	1	-	-	
17	-	-	-	-	-	-	-	-	-	-	-	-	-	1	-	1	-	-	-	
16	-	-	-	-	-	-	-	-	-	-	-	-	-	-	2	1	-	-	-	
15	-	-	-	-	-	-	-	-	-	-	1	2	2	3	-	-	-	-	-	
14	-	-	-	-	-	-	-	-	-	2	6	8	5	4	-	-	-	-	-	
13	-	-	-	-	-	-	1	-	-	3	5	8	1	-	-	-	-	-	-	
12	-	-	-	-	-	-	-	1	17	13	23	9	1	-	-	-	-	-	-	
11	-	-	-	-	-	1	1	4	22	25	4	1	-	-	-	-	-	-	-	
10	-	-	-	-	-	2	15	51	144	24	2	1	-	-	-	-	-	-	-	
9	-	-	-	-	1	1	29	89	32	4	-	-	-	-	-	-	-	-	-	
8	-	-	-	-	3	21	97	25	9	-	-	-	-	-	-	-	-	-	-	
7	-	-	-	3	23	149	38	5	-	-	-	-	-	-	-	-	-	-	-	
6	-	-	-	6	51	15	2	-	-	-	-	-	-	-	-	-	-	-	-	
5	-	-	5	45	10	1	-	1	-	-	-	-	-	-	-	-	-	-	-	
4	-	6	25	5	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	
3	4	25	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	
2	50	-	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	

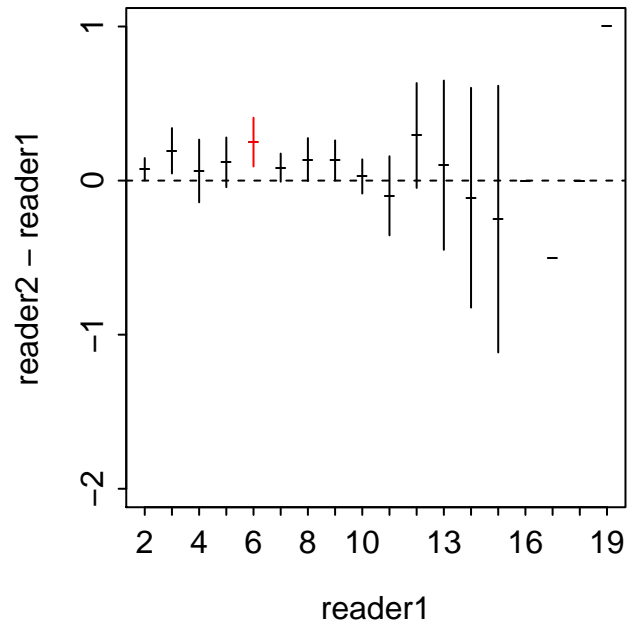
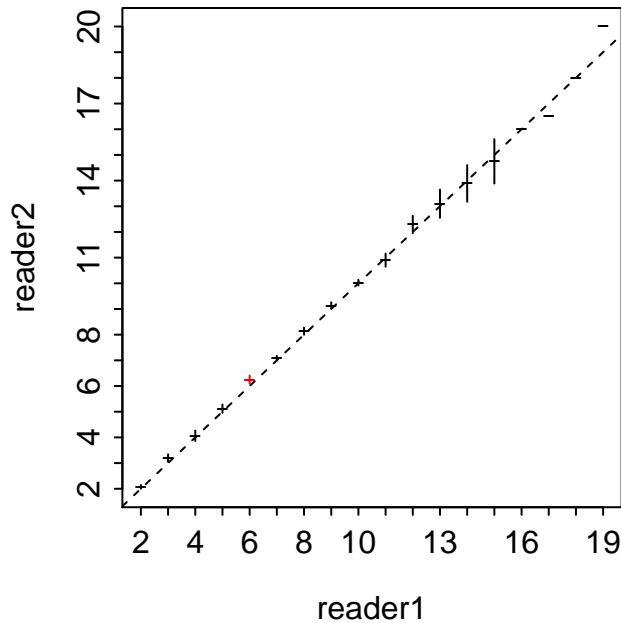
```
> summary(ab,what="symmetry")
```

	symTest	df	chi.sq	p
1	McNemars	1	9.204793	0.0024138229
2	EvansHoenig	5	19.824421	0.0013481675
3	Bowkers	37	72.685469	0.0004126986

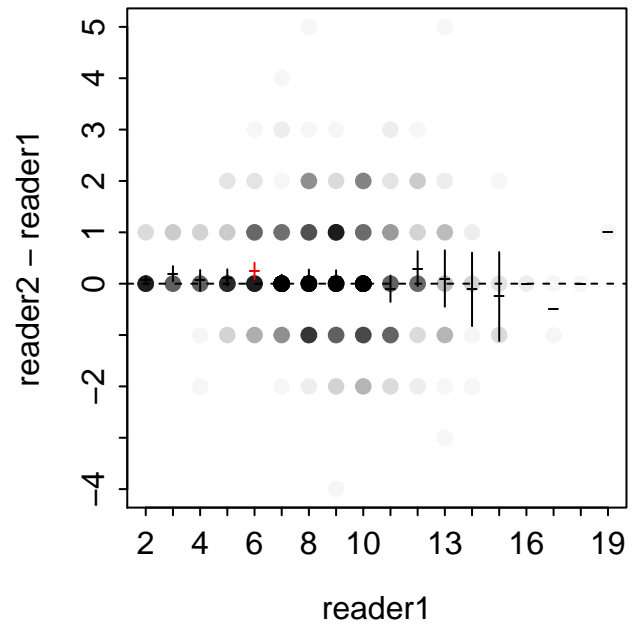
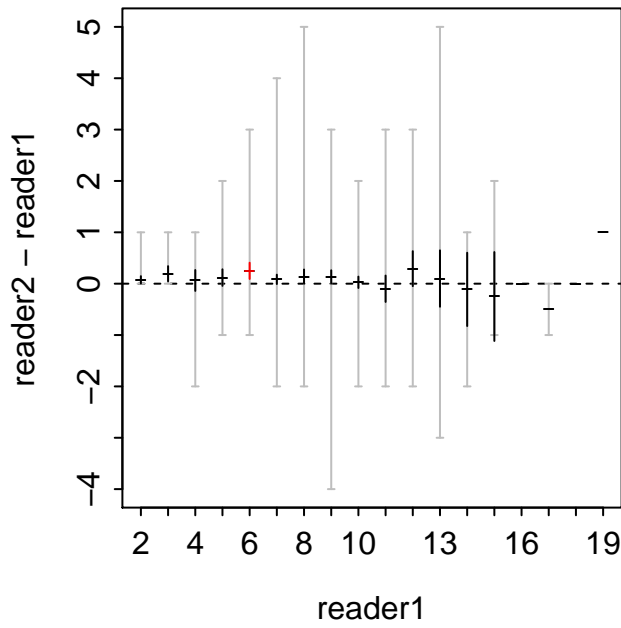
```
> summary(ab,what="bias")
```

reader1	n	min	max	mean	SE	t	adj.p	sig	LCI	UCI
2	54	2	3	2.07	0.0360	2.059	0.5329	FALSE	2.00	2.15
3	31	3	4	3.19	0.0721	2.683	0.1527	FALSE	3.05	3.34
4	32	2	5	4.06	0.0998	0.626	1.0000	FALSE	3.86	4.27
5	59	4	7	5.12	0.0805	1.474	1.0000	FALSE	4.96	5.28
6	88	5	9	6.25	0.0796	3.141	0.0322	TRUE	6.09	6.41
7	190	5	11	7.08	0.0462	1.823	0.6294	FALSE	6.99	7.18
8	183	6	13	8.14	0.0705	1.937	0.5423	FALSE	8.00	8.28
9	176	5	12	9.13	0.0660	1.981	0.5404	FALSE	9.00	9.26
10	224	8	12	10.03	0.0562	0.477	1.0000	FALSE	9.92	10.14
11	71	9	14	10.90	0.1287	-0.766	1.0000	FALSE	10.64	11.16
12	41	10	15	12.29	0.1684	1.738	0.7187	FALSE	11.95	12.63
13	30	10	18	13.10	0.2685	0.372	1.0000	FALSE	12.55	13.65
14	9	12	15	13.89	0.3093	-0.359	1.0000	FALSE	13.18	14.60
15	8	14	17	14.75	0.3660	-0.683	1.0000	FALSE	13.88	15.62
16	2	16	16	16.00	NA	NA	NA	FALSE	NA	NA
17	2	16	17	16.50	NA	NA	NA	FALSE	NA	NA
18	1	18	18	18.00	NA	NA	NA	FALSE	NA	NA
19	1	20	20	20.00	NA	NA	NA	FALSE	NA	NA

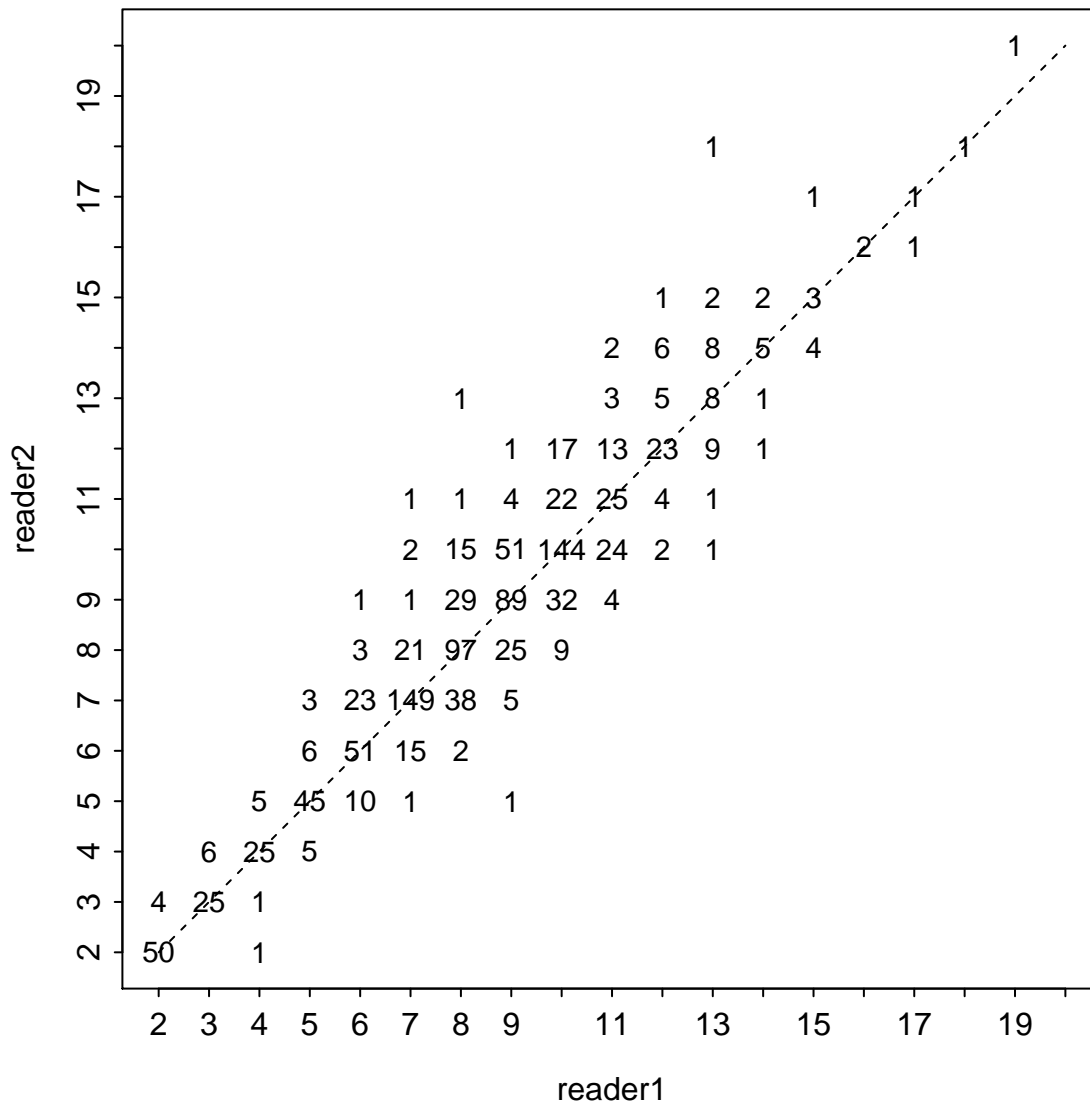
```
> plot(ab) # Left
> plot(ab,diff=TRUE) # Right
```



```
> plot(ab,diff=TRUE,show.range=TRUE) # Left
> plot(ab,diff=TRUE,show.pts=TRUE,transparency=1/25) # Right
```



```
> plot(ab,what="numbers",xlim=c(2,20),ylim=c(2,20))
```



## Examine Age Precision

```
> ap <- agePrecision(reader2~reader1,data=SB)
> summary(ap,what="difference",digits=1)
  -4   -3   -2   -1    0    1    2    3    4    5
0.08 0.08 2.16 14.06 61.81 16.31 4.58 0.67 0.08 0.17

> summary(ap,what="absolute difference",digits=2)
    0    1    2    3    4    5
61.81 30.37 6.74 0.75 0.17 0.17

> summary(ap,what="precision")
      n R  ACV   APE PercAgree
1202 2 3.98 2.815    61.81

> summary(ap,what="detail") # only some rows shown
```

	reader2	reader1	avg	sd	APE	ACV
1	2	2	2.0	0.0000000	0.000000	0.000000
2	2	2	2.0	0.0000000	0.000000	0.000000
3	2	2	2.0	0.0000000	0.000000	0.000000
1200	18	13	15.5	3.5355339	16.129032	22.809896
1201	18	18	18.0	0.0000000	0.000000	0.000000
1202	20	19	19.5	0.7071068	2.564103	3.626189

## Application Assignment

Herbst and Marsden (2012) [reprint is here](#) compared the precision, bias, and reader uncertainty of scales, dorsal fin rays, and otolith age estimates from 151 lake whitefish (*Coregonus clupeaformis*) from Lake Champlain in 2009. The data for their comparisons were recorded in `WhitefishLC.csv`. This file contains initial age assessments for two readers on three structures (variable names are the structure name with a “1” or “2” appended to denote the reader). In addition, the two readers developed a consensus age (variable name is the structure name with a “C” appended).

Create a script that performs the following tasks:

1. Use a variety of methods (tabular, graphical, and statistical) to describe any apparent bias in **consensus** ages between scales and otoliths.
2. Describe any apparent bias in age assessment for otoliths between the two readers.
3. Describe precision of age assessment for otoliths between the two readers.
4. (*Time Permitting*) Describe any apparent bias in age assessment for scales between the two readers.
5. (*Time Permitting*) Describe precision of age assessment for scales between the two readers.
6. (*Time Permitting*) Use a variety of methods (tabular, graphical, and statistical) to describe any apparent bias in **consensus** ages between fin rays and otoliths.
7. (*Time Permitting*) Describe any apparent bias in age assessment for fin rays between the two readers.
8. (*Time Permitting*) Describe precision of age assessment for fin rays between the two readers.

**Save your script!**