

Results

ANOVA - freq

ANOVA - freq

Cases	Sum of Squares	df	Mean Square	F	p
freq_cat	983.199	1	983.199	2135.619	< .001
mismatch	1.109	2	0.555	1.205	0.302
freq_cat * mismatch	0.049	2	0.025	0.053	0.948
Residuals	107.729	234	0.460		

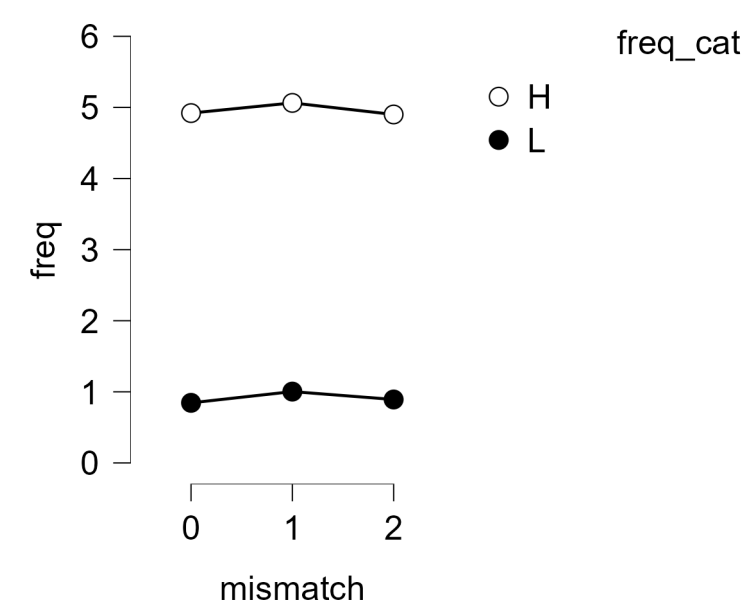
Note. Type III Sum of Squares

Descriptives

Descriptives - freq

freq_cat	mismatch	Mean	SD	N
H	0	4.920	0.764	40
	1	5.064	1.062	40
	2	4.901	0.510	40
L	0	0.845	0.495	40
	1	1.003	0.461	40
	2	0.892	0.576	40

Descriptives plots



Post Hoc Tests

Standard

Post Hoc Comparisons - mismatch

		Mean Difference	SE	t	P _{tukey}
0	1	−0.151	0.107	−1.404	0.340
	2	−0.014	0.107	−0.130	0.991
1	2	0.137	0.107	1.275	0.411

Note. P-value adjusted for comparing a family of 3
Note. Results are averaged over the levels of: freq_cat

Post Hoc Comparisons - freq_cat

		Mean Difference	SE	t	P _{tukey}
H	L	4.048	0.088	46.213	< .001

Note. Results are averaged over the levels of: mismatch

ANOVA - nletters

ANOVA - nletters

Cases	Sum of Squares	df	Mean Square	F	p
freq_cat	0.104	1	0.104	0.179	0.673
mismatch	161.008	2	80.504	138.387	< .001
freq_cat * mismatch	0.758	2	0.379	0.652	0.522
Residuals	136.125	234	0.582		

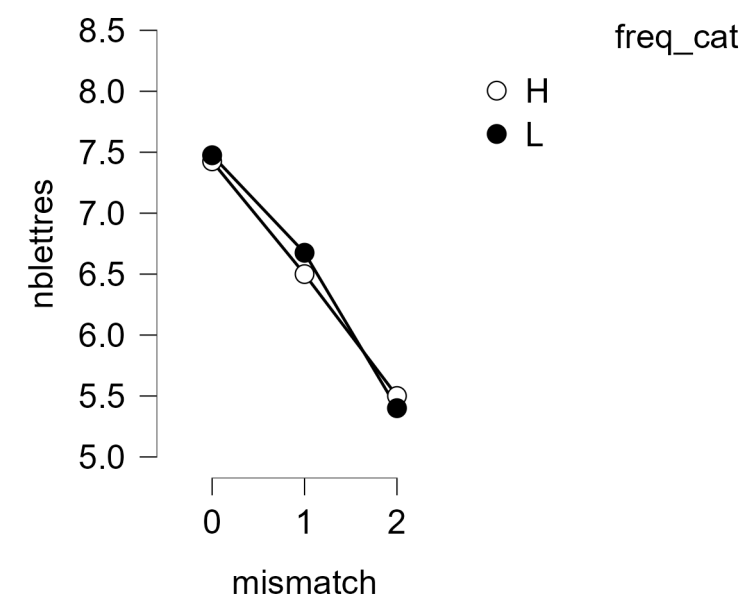
Note. Type III Sum of Squares

Descriptives

Descriptives - nletters

freq_cat	mismatch	Mean	SD	N
H	0	7.425	0.747	40
	1	6.500	0.816	40
	2	5.500	0.716	40
L	0	7.475	0.816	40
	1	6.675	0.829	40
	2	5.400	0.632	40

Descriptives plots



Post Hoc Tests

Standard

Post Hoc Comparisons - freq_cat

		Mean Difference	SE	t	P _{tukey}
H	L	-0.042	0.098	-0.423	0.673

Note. Results are averaged over the levels of: mismatch

Post Hoc Comparisons - mismatch

		Mean Difference	SE	t	P _{tukey}
0	1	0.863	0.121	7.152	< .001
	2	2.000	0.121	16.584	< .001
1	2	1.137	0.121	9.432	< .001

Note. P-value adjusted for comparing a family of 3

Note. Results are averaged over the levels of: freq_cat

Post Hoc Comparisons - freq_cat * mismatch

		Mean Difference	SE	t	P _{tukey}
H 0	L 0	-0.050	0.171	-0.293	1.000
	H 1	0.925	0.171	5.424	< .001
	L 1	0.750	0.171	4.398	< .001
	H 2	1.925	0.171	11.287	< .001
	L 2	2.025	0.171	11.874	< .001
L 0	H 1	0.975	0.171	5.717	< .001
	L 1	0.800	0.171	4.691	< .001
	H 2	1.975	0.171	11.580	< .001
	L 2	2.075	0.171	12.167	< .001
H 1	L 1	-0.175	0.171	-1.026	0.909
	H 2	1.000	0.171	5.863	< .001
	L 2	1.100	0.171	6.450	< .001
L 1	H 2	1.175	0.171	6.890	< .001
	L 2	1.275	0.171	7.476	< .001
H 2	L 2	0.100	0.171	0.586	0.992

Note. P-value adjusted for comparing a family of 6

