

Comparing NHL Salaries: A Random Permutation Test

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Are salaries "fair" in the National Hockey League (NHL)? If so, we would expect that salaries are comparable for the right-wing and left-wing positions. To address this question, I downloaded data from the *USA Today* salary database for the 2007-2008 NHL season and used JMP to run a random permutation test.

jmp 2007NHLSalary.jmp

PermutationTest.jsl

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Permutation Tests

* Also called randomization tests, re-randomization tests, or exact tests

GENERAL PROCEDURE FOR PERMUTATION TESTS

To carry out a permutation test based on a statistic that measures the size of an effect of interest:

1. Compute the statistic for the original data.

2. Choose permutation resamples from the data without replacement in a way that is consistent with the null hypothesis of the test and with the study design. Construct the permutation distribution of the statistic from its values in a large number of resamples.

3. Find the *P*-value by locating the original statistic on the permutation distribution.

From Moore, McCabe, & Craig (2009)

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Advantages of Random Permutation Tests

Sexier" than traditional nonparametric tests

- ☑ Can be used for <u>any</u> test statistic, regardless of whether or not its distribution is known
- Solution More general hypotheses than those used for *t*-tests *For example:*
 - H₀: Right-wing and left-wing hockey players have the same salary distribution.
 H₁: The salaries of right-wing and left-wing hockey players are systematically different.

☑ Very accurate *P*-values if enough permutations are used

☑ Useful for checking the normality assumption of the twosample *t*-test

Primary Disadvantage

Custom computer code is required.

 My JMP script can only be used for hypothesis tests involving two independent samples.

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Procedure

- First, find the difference $\overline{x}_{RW} \overline{x}_{LW}$ for the original sample data. We find that $\overline{x}_{RW} \overline{x}_{LW} = \$400,102$. This is our test statistic.
- We don't compute t or any other standardized statistic.
- To perform a permutation test, we randomly regroup the salary data into two groups that are the same sizes as the two original samples (114 LW and 114 RW). This is consistent with the null hypothesis that left-wing and right-wing player salaries have the same distribution. Each salary appears once in each resample, but some salaries move from the LW group to the RW group and vice versa.
- Open up the script named *PermutationTest.jsl* and the data table named 2007NHLSalary.jmp.
- Bring the script window to the front, and then click on Edit > Run Script.

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- After a short delay, the script will generate a summary table containing 999 resample means for the two groups.
- •Add a new column named "Difference (RW LW)" to the summary table and define it as *Total Salary_RW Total Salary_LW*.
- •The last row in the table includes the test statistic. Select this row, then click on Rows > Exclude/Unexclude.

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For each permutation resample, the script calculates the means for the "new" RW group and the "new" LW group. Then, we find the difference between the two means, $\overline{x}_{RW} - \overline{x}_{LW}$. We repeat the process 999 times and construct the <u>permutation</u> <u>distribution</u> of $\overline{x}_{RW} - \overline{x}_{LW}$. The permutation distribution estimates the sampling distribution under the condition that H_0 is true.





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Select Columns Total Salary_LW Total Salary_RW Difference (RW - LW) Replace table Output table name:	By A Difference (RW Remove A Difference (RW <i>a</i> V Copy formula Copy formula Suppress formula evaluation	Action OK Cancel Help

	00	😝 🔿 🔿 📑 Sorted Resamples 🖉				
	Sorted Resamples					ŕ
	Source		Total Salary_LW	Total Salary_RW	Difference (RW - LW)	- 1
		949	1635139.65	2020440.44	385300.789	- 1
		950	1632628.86	2022951.23	390322.368	- 1
		951	1631504.3	2024075.79	392571.491	- 1
		952	1629187.02	2026393.07	397206.053	- 1
	Gelenne (2.0)	953	1628821.05	2026759.04	397937.982	- 1
The <i>P</i> -value is the proportion of the 999 resamples that	Total Salary_LW	S 954	1627694.04	2027886.05	400192.018	- 1
		955	1627562.46	2028017.63	400455.175	- 1
vield $ \overline{x}_{pw} - \overline{x}_{rw} $ values that are at least as extreme as the	Difference (PW = 1W)	956	1626044.39	2029535.7	403491.316	- 1
Jiera n _{Rw} - n _{Lw} fundes that are at reast as endedness are		957	1625252.81	2030327.28	405074.474	-
observed difference of 400192 018		958	1624845.09	2030735	405889.912	- 11
		959	1624793.25	2030786.84	405993.596	
		960	1623284.21	2032295.88	409011.667	
	Rows	961	1618309.65	2037270.44	418960.789	-
For a two sided test, we need to count how many	All rows 1000	962	1614193.25	2041386.84	427193.596	
For a two-sided test, we need to count now many	Selected 83	963	1614187.11	2041392.98	427205.877	
r_{a} results have differences that are either < 400102019	Hidden	964	1612769.21	2042810.88	430041.667	
resamples have unreferences that are either \geq -400192.018	Labelled (965	1612076.93	2043503.16	431426.228	-
$2\pi > 400102.018$		966	1611781.67	2043798.42	432016.754	Ă
$0f \ge 400192.018.$		967	1611449.39	2044130.7	432681.316	¥ ► //.
To improve the accuracy of the <i>P</i> -value, add 1 to both the numerator and the denominator.	• When I ran the script, there were 83 resamples that resulted in $ \bar{x}_{RW} - \bar{x}_{LW} $ that was at least as extreme as 400192.018. (Your results may vary.)					
	• The two-sided P-value for my random permutation test is					
	$\frac{83+1}{999+1} = .084$					
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Moore, McCabe, & Craig (2009) on the robustness of the two-sample *t*-test procedure: "When the sizes of the two samples are equal and the distributions of the two populations being compared have similar shapes [not necessarily normal], probability values from the *t* table are quite accurate for a broad range of distributions when the sample sizes are as small as $n_1 = n_2 = 5$."

6th Edition, p. 456



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