Name:

1 WI23 Midterm Problem 6

The DataFrame tv_excl (right) contains information about a group of TV shows, and DataFrame counts (left) contains the number of TV shows for *every* combination of "Age" and "Service" in tv_excl.

Service	Disney+	Hulu	Netflix	Prime Video		Title	Year	Age	IMDb	Rotten Tomatoes	Service
Age					0	Jersey Shore	2009	16+	3.6	54	Hulu
13+	NaN	4.0	2.0	1.0	4	Honry Hugglomonstor	2012	011	5.2	10	Dispovu
16+	13.0	405.0	320.0	147.0		Henry Hugglemonster	2013	an	5.5	42	Disney+
18+	NaN	223.0	445.0	134.0	2	Fast & Furious Spy Racers	2019	7+	5.5	62	Netflix
7+	91.0	246.0	245.0	149.0	3	Atlanta	2016	18+	8.6	84	Hulu
all	116.0	97.0	151.0	144.0	4	Played	2013	NaN	6.4	45	Prime Video

Given the above information, what does the following expression evaluate to? tv_excl.groupby(["Age", "Service"]).sum().shape[0]

Tiffany would like to compare the distribution of Age for Hulu and Netflix. Specifically, she'd like to test the following hypotheses:

- Null Hypothesis: The distributions of Age for Hulu and Netflix are drawn from the same population distribution, and any observed differences are due to random chance.
- Alternative Hypothesis: The distributions of Age for Hulu and Netflix are drawn from different population distributions.

Is this a hypothesis test, or a permutation test? Why?

Consider the DataFrame distr, defined below.

hn = counts[["Hulu", "Netflix"]]
distr = (hn / hn.sum()).T # Note that distr has 2 rows and 5 columns.

To test the hypotheses above, Tiffany decides to use the total variation distance as her test statistic. Which of the following expressions **DO NOT** correctly compute the observed statistic for her test?

[] distr.diff().iloc[-1].abs().sum() / 2 [] distr.diff().sum().abs().sum() / 2 [] distr.diff().sum().sum().abs() / 2 [] (distr.sum() - 2 * distr.iloc[0]).abs().sum() / 2 [] distr.diff().abs().sum(axis=1).iloc[-1] / 2

2 WI23 Final Problem 2.5

Suppose $\vec{a} = \begin{bmatrix} a_1 & a_2 & \dots & a_n \end{bmatrix}^T$ and $\vec{b} = \begin{bmatrix} b_1 & b_2 & \dots & b_n \end{bmatrix}^T$ are both vectors containing proportions that add to 1. As we've seen before, the TVD is defined as follows:

$$\text{TVD}(\vec{a}, \vec{b}) = \frac{1}{2} \sum_{i=1}^{n} |a_i - b_i|$$

The TVD is not the only metric that can quantify the distance between two categorical distributions. Here are three other possible distance metrics:

• dis1 $(\vec{a}, \vec{b}) = \vec{a} \cdot \vec{b} = a_1 b_1 + a_2 b_2 + \dots + a_n b_n$

• dis2
$$(\vec{a}, \vec{b}) = \frac{\vec{a} \cdot \vec{b}}{|\vec{a}||\vec{b}|} = \frac{a_1b_1 + a_2b_2 + \dots + a_nb_n}{\sqrt{a_1^2 + a_2^2 + \dots + a_n^2}\sqrt{b_1^2 + b_2^2 + \dots + b_n^2}}$$

• dis3 $(\vec{a}, \vec{b}) = 1 - \frac{\vec{a} \cdot \vec{b}}{|\vec{a}||\vec{b}|}$

Of the above three possible distance metrics, only one of them has the same range as the TVD (i.e. the same minimum possible value and the same maximum possible value) and has the property that smaller values correspond to more similar vectors. Which distance metric is it?

3 FA23 Midterm Problem 4

In this question, we will work with the DataFrame donkeys, about the health of various donkeys. (Don't worry about the WeightAlt column for now.)

	id	BCS	Age	Weight	WeightAlt	id	A unique identifier for each donkey (d01, d02,
0	d01	3.0	<2	77	NaN		etc.).
1	d02	2.5	<2	100	NaN	BCS	Body condition score: from 1 (emaciated) to 3 (healthan) to 5 (above) in increments of 0.5
2	d03	1.5	<2	74	NaN	Age	(nearthy) to 5 (obese) in increments of 0.5. Age in vears: $<2, 2-5, 5-10, 10-15, 15-20$, and
						U	over 20 years.
						Weight	Weight in kilograms.
						WeightAlt	Second weight measurement taken for 30 don-
							keys. NaN if the donkey was not reweighed.

Alan wants to see whether donkeys with $BCS \ge 3$ have larger Weight values on average compared to donkeys that have BCS < 3. Select all the possible test statistics that Alan could use to conduct this hypothesis test. Let μ_1 be the mean weight of donkeys with $BCS \ge 3$ and μ_2 be the mean weight of donkeys with BCS < 3.

[] μ_1 [] $\mu_1 - \mu_2$ [] $2\mu_2 - \mu_1$ [] $|\mu_1 - \mu_2|$ [] Total variation distance