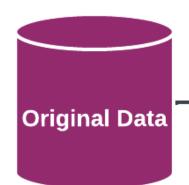
# Hyperdimensional Change Detection for Novel Data Set **Exploration and Continuous Unsupervised Monitoring** Sean Coffinger

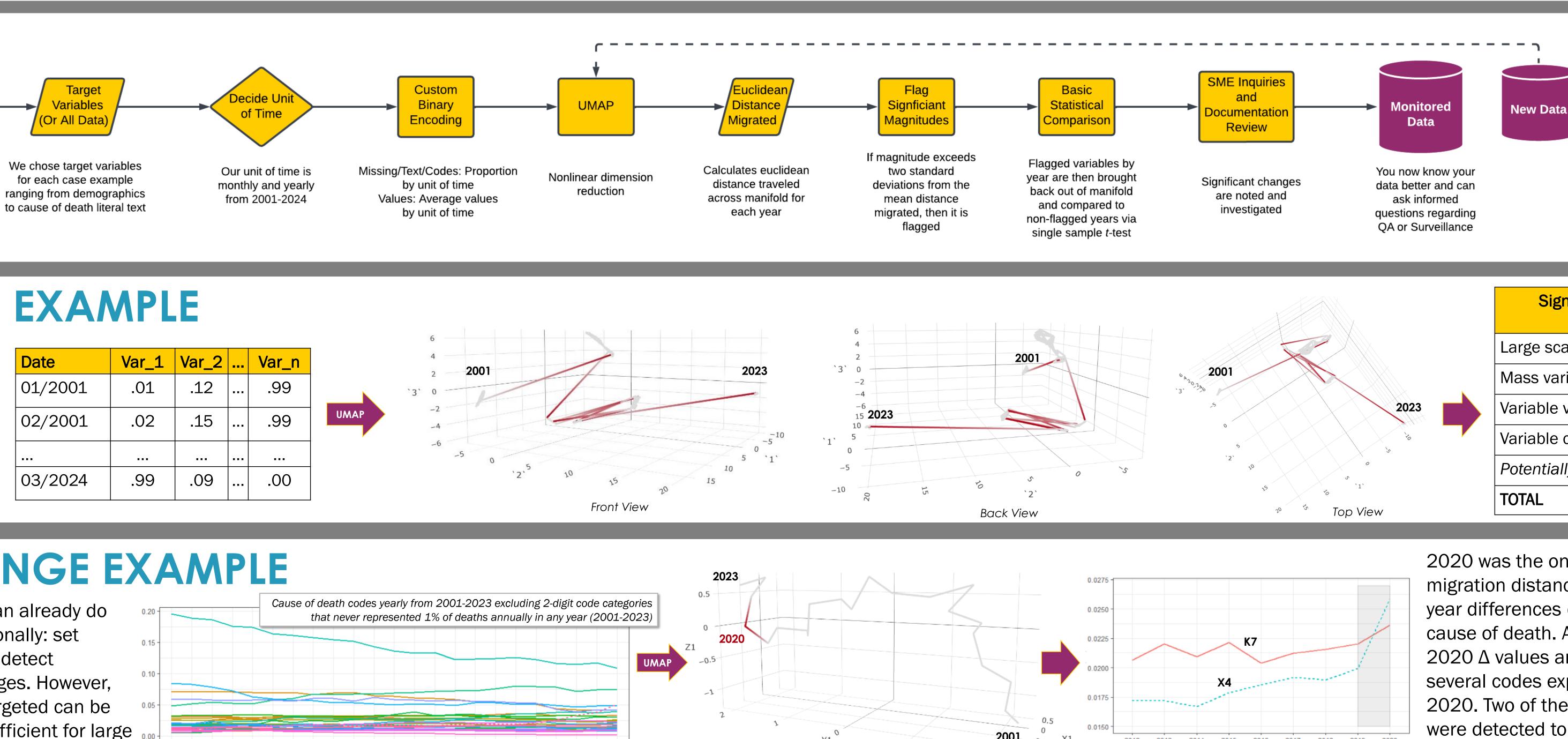
#### THE PROBLEM

Practitioners who interact with data often utilize unfamiliar longitudinal data sets and continuously monitor for new and unexpected changes. These data can vary across time in quality, values, coding, etc. The lack of comprehensive documentation of historical variable changes independently to pursue clarification of key changes from responsible subject matter experts. In addition to retroactive analyses of data, the practice of continuous monitoring for quality assurance, and/or informing robust and targeted surveillance, is warranted. Here we suggest a global, unsupervised, and untargeted approach to identify significant changes in individual and sets of variables. The goal is to inform practitioners during the 'exploratory' phase of a project of significant changes. Furthermore, the aim is not to quantify all abnormalities, but rather monitor when and where interesting changes have occurred with the benefit of generalizability, flexibility, control, speed and automation.

#### **OUR SOLUTION**

Change detection via manifold migration can be employed to identify points of interest (Time x Variable). This can be applied to any time scale and number/type of variables.





Here we use WA DOH Death Records

### CASE #1 – MISSINGNESS EXAMPLE

To investigate global system changes or large shifts in variable usage, we used this strategy to look at global missingness. Here we ingested all descriptive demographic variables used in a typical linkage project and evaluation from the death table and calculated proportion missing by month.



#### CASE #2.1 – VALUE CHANGE EXAMPLE

To demonstrate how we can track value changes in variable(s) we used this strategy to monitor significant changes in coded trends. Here we chose the non-quantitative value of Cause of Death (COD). Note this is even easier with quantitative values.

Of course we can already do this 2- dimensionally: set thresholds and detect abnormal changes. However, doing this untargeted can be tedious and inefficient for large variable change detection.

## CASE #2.2 – COD LITERALS EXAMPLE

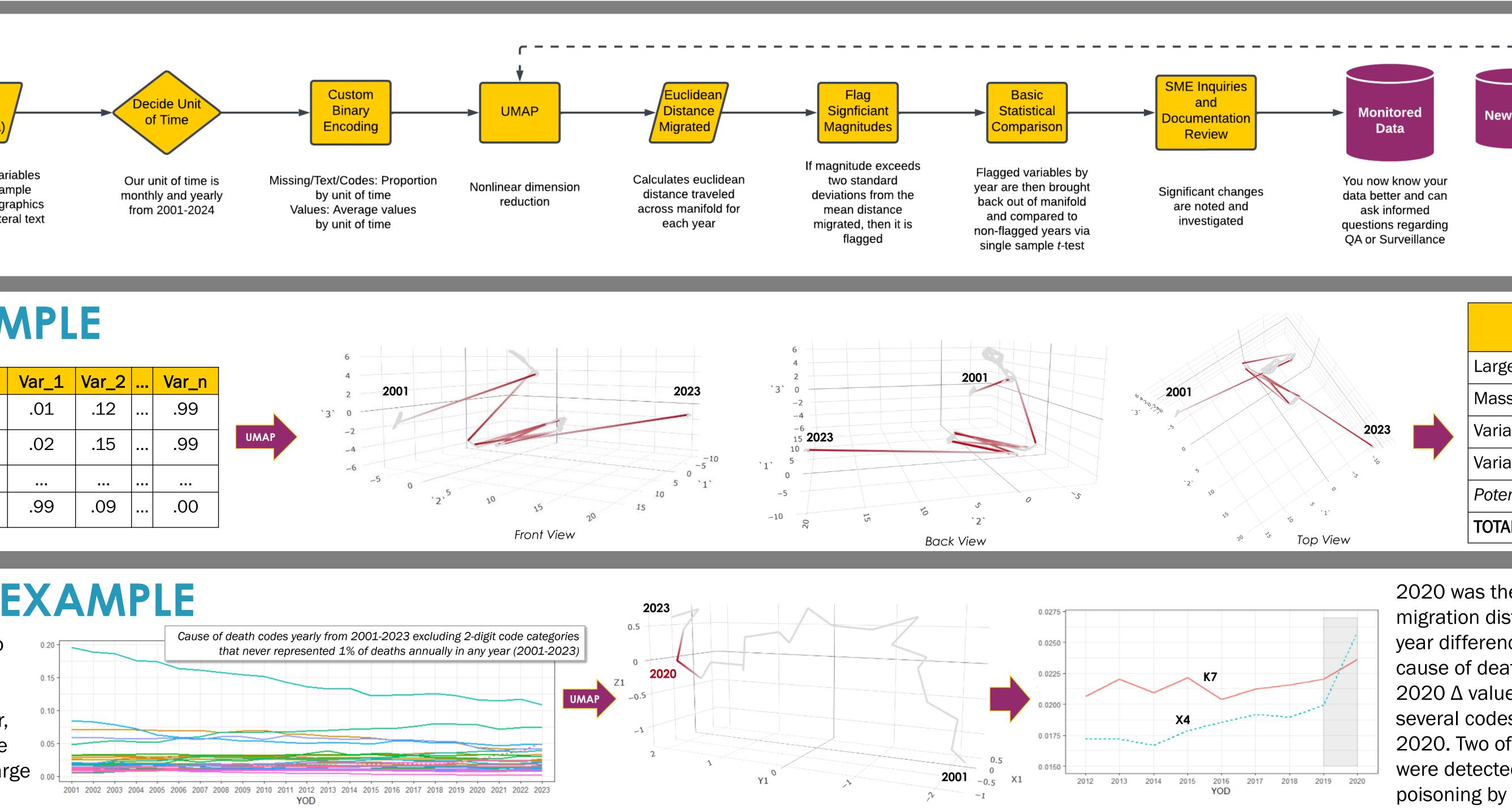
In Case 2.1 we discovered that a year flagged via migration magnitude identified two CODs that significantly increased their trajectory in coded proportion. To demonstrate the ability to utilize these methods and strategy on literals, qualitative and text fields, we employ similar methods to analyze individual words in COD literal text fields that are changing in terms of  $\Delta$  frequency.

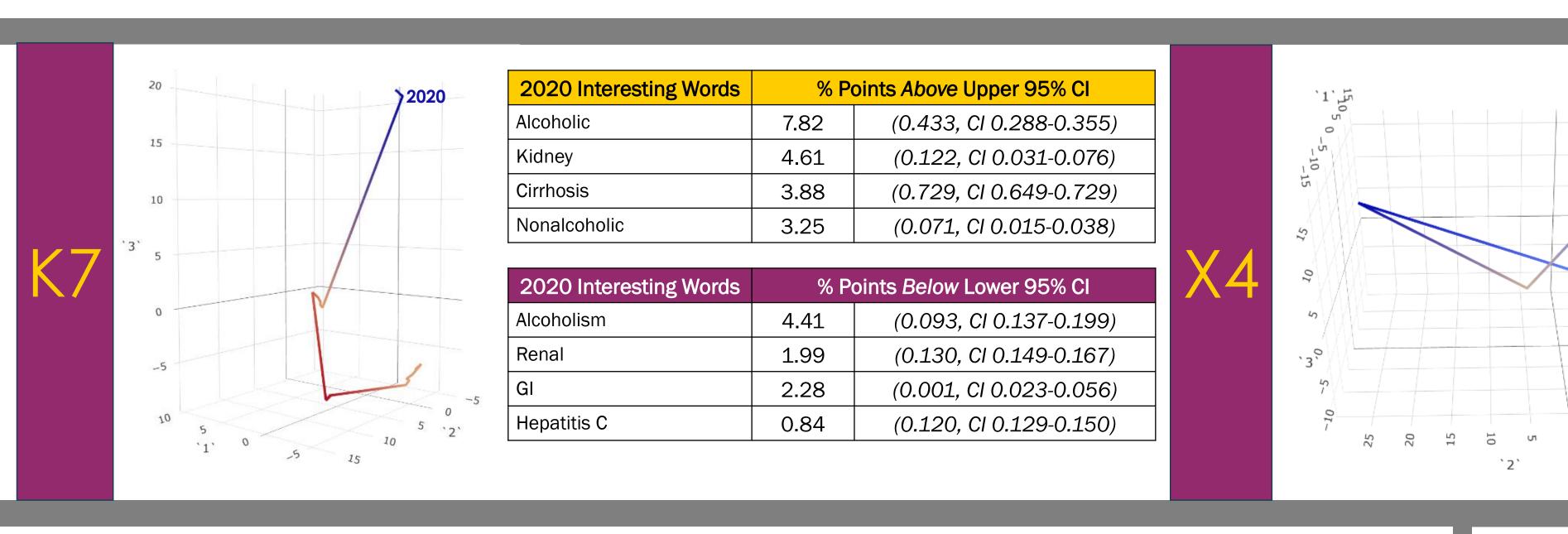
For both K7 and X4 COD records, literals were queried then tokenized into individual words. A medical coding "common garbage word" list was used to filter out common words (i.e., and, or, additional, etc.). Additionally, infrequent words and numeric values were excluded (<1% max annual proportion per code). Annual proportion of incidence for each word was calculated then fed into 4-dimensional UMAP manifold generation.

#### SUMMARY

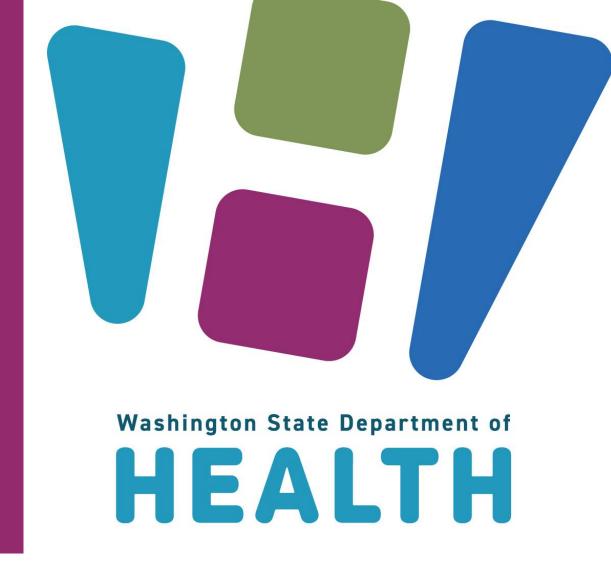
Case #1 successfully demonstrated how we can use manifold migration monitoring to successfully investigate and QA a novel historical dataset. Eight significant changes were detected in demographic variables, six were confirmed by SMEs and documentation. Case #2 demonstrated the generalizability of hyperdimensional change detection by looking at yearly death trends to discover two notable increases in 2020. Then, a separate manifold was constructed to delve further into our initial finding to gain insight on possible contributing factors. In all, manifold migration has limitations compared to targeted analyses, but can provide fast, generalized, untargeted information to any data user.

#### Washington State Department of Health, The Center for Health Statistics









UMAP can be easily used for future data, enabling continuous monitoring on an ongoing basis

Once defined, the constructed manifold output from

Note: Manifold Migration involves calculated Euclidean distance of units of time in a nonlinear manifold. This is not typically recommended and can produce some inaccurate quantifications; however, for the purpose of detection based on magnitude changes, it has limits but work sufficiently.

2023	
ં	

Significant Changes Identified Post-SME Inquiry	Count
Large scale system changes/migrations	2
Mass variable discontinuations	2
Variable value changes	1
Variable category addition	1
Potentially unidentified change	2
TOTAL	8

2020 was the only 'flagged' year based on migration distance. We then compared the year-toyear differences ( $\Delta$ ) of proportions coded for each cause of death. A *t*-Test ( $\alpha$ =.001) between 2019 to 2020  $\Delta$  values and all previous  $\Delta$ s confirmed that several codes experienced a significant change in 2020. Two of these 2-digit ICD-10 mortality codes were detected to have increases: X4 (Accidental poisoning by an exposure to noxious substances) and K7 (Diseases of Liver).

	2020 Interesting Words	% Points Above Upper 95% Cl		
	Methamphetamine	12.00	(0.441, CI 0.171-0.321)	
	Fentanyl	11.39	(0.383, CI 0.039-0.269)	
	Heroin	3.13	(0.241, CI 0.110-0.210)	
	Obesity	0.15	(0.036, CI 0.024-0.343)	
	2020 Interesting Words	% Points Below Lower 95% Cl		
	Oxycodone	3.44	(0.052, CI 0.087-0.130)	
	Diazepam	2.20	(0.010, CI 0.032-0.055)	
۱ <sub>2020</sub>	Morphine	1.93	(0.041, CI 0.060-0.088)	

CONTACT Sean Coffinger, MA **Health Statistics Manager** Center for Health Statistics Sean.coffinger@doh.wa.gov

Publication # DOH 422-284 To request this document in another format, call 1-800-525-0127. Deaf or hard of hearing customers, please call 711 (Washington Relay) or email doh.information@doh.wa.gov