



# Introduction

In recent years, in low-mortality countries, the causes of death have become more diverse. This entails:

- Increased uncertainties;
- Lower predictability;
- More complex and fragmented efforts to reduce mortality.

# Introduction

Studies of causes diversity are relatively recent, but are growing in numbers.

- Most focus on the underlying causes of death (UCD), but ....
- Various measures have been used and proposed:
  - ▶ Diversity indices
  - ▶ Distance measures
- Focus on low mortality countries, but...

# Learning from biodiversity studies

Biodiversity research offers valuable tools to study diversity. There are two key aspects:

- Richness: Numbers/types
- Evenness: Diversity/complexity

# Objective

To study diversity in causes of death using multiple causes of death (MCoD) data using the two aspects of diversity.

# Framework

We need a framework that:

- allow to quantify both metrics,
- is age-standardized.

Solution: A life table that accounts for MCoD.

# MCoD life tables

Multiple decrement life tables calculate the death probability at age  $x$  and UCD  $u$ , as:

$$q(x, u) = q(x) \frac{D(x, u)}{D(x)}$$

The death probability of underlying causes  $u$  with contributing cause  $c$  can be estimated as:

$$q(x, u, c) = q(x) \frac{D(x, u, c)}{D(x)}$$

# Other indicators in an MCoD life tables

Life table deaths (distribution):  $d(x, u, c) = q(x, u, c) l(x)$

Mortality rates:  $m(x, u, c) = \frac{d(x, u, c)}{L(x)}$

Prevalence at death:  $PD_{c/u} = \sum_x d(x, u, c) / d(x, u)$

Cause of death association indicator:  $CDAI_{u,c} = \sum_x \frac{d(x, u, c)}{d(x, u)} / \sum_x \frac{d(x, c)}{d(x)}$



# MCoD life table - Example

Age	$d_{u1}$	$d_{u1,c2}$	$d_{u1,c3}$	$d_{u2}$	$d_{u2,c1}$	$d_{u2,c3}$	$d_{u3}$	$d_{u3,c1}$	$d_{u3,c2}$
0	12	0	2	2	0	0	539	41	0
1	2	0	1	2	0	0	34	5	0
2	1	0	0	2	0	0	20	2	0
...									
65	299	9	113	336	79	113	429	187	14
66	314	10	132	359	88	140	454	203	18
67	341	11	162	375	89	152	471	213	18
...									
109	4	0	2	0	0	0	6	1	0
110+	5	0	2	0	0	0	5	1	0

# Data

## Application to US data:

- 1 MCoD data from the Multiple Causes of Death data on CDC Wonder
- 2 Causes grouping by main ICD-10 chapter
- 3 Life tables from the Human Mortality Database
- 4 Period: 2006 to 2021
- 5 Both sexes combined
- 6 If a death had multiple entry of the same cause, only one was kept.

Upcoming results for Denmark, Spain and France and by sex.

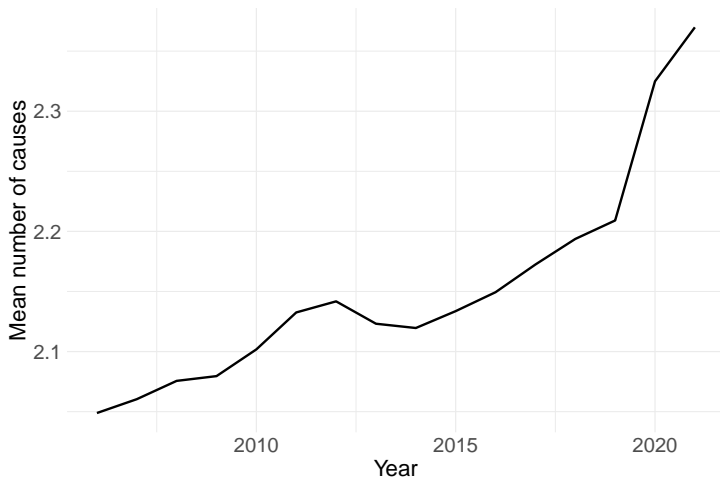
# Richness and number of causes

The average number of causes reported on the death certificate is estimated as:

$$N = \sum_x \sum_u \sum_c d(x, u, c)$$

with  $N(x) = \sum_u \sum_c d(x, u, c)$  and  $N(u) = \sum_x \sum_c d(x, u, c)$

# Richness: Time trend



# Richness: Decomposition

$N$  can be decomposed by changes in  $d(x)$ ,  $d(u)$  and  $N(x, u)$ , using the Das Gupta decomposition.

	$\Delta d(x)$	$\Delta d(u)$	$\Delta N(x, u)$	Total
2006-2019	-0.004	0.019	0.114	0.129
2019-2021	0.006	-0.024	0.175	0.157

# Evenness and diversity: Two approaches

What is diversity in a MCoD context? How to consider the contributing causes?

- 1 All causes are part of the mortality process, no matter their position.
- 2 The UCD all come with their own set of contributing causes with a specific diversity.

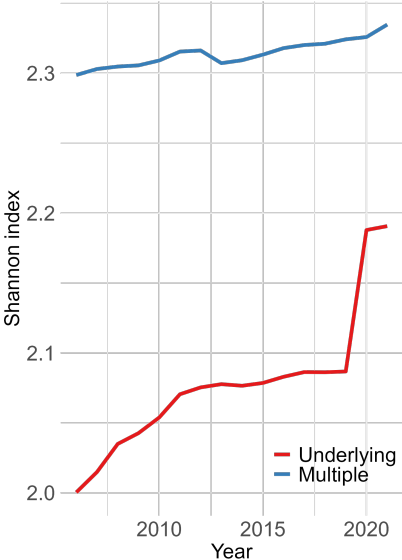
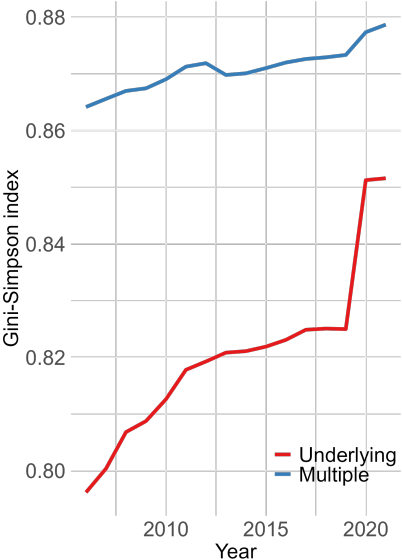
# Diversity, First approach

We are interested in studying the distribution  $\hat{d}_i$ , where  $i$  is a given cause no matter its position on the death certificate, such that

$$\hat{d}_i = \frac{d_i}{N}$$

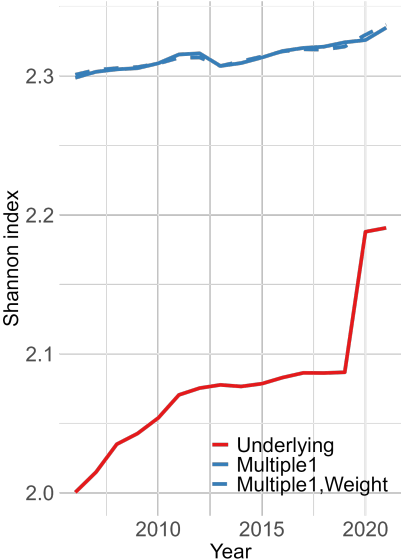
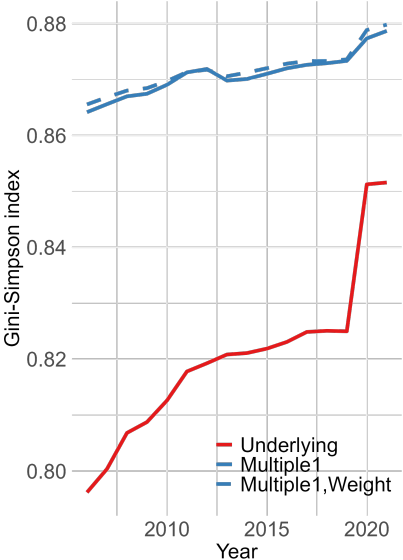
Diversity can be calculated with standard diversity index, such as the Shannon entropy and the Simpson index applied to  $\hat{d}_i$ .

# Diversity, First approach, Results



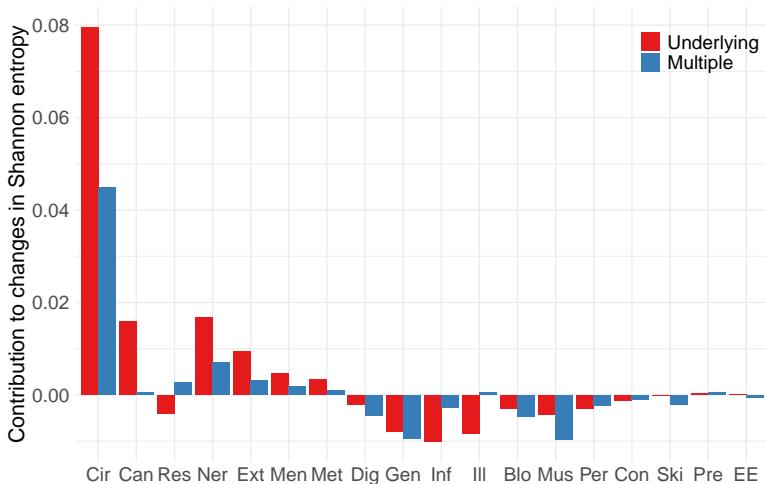


# Diversity, First approach, Results with weights



# Diversity, First approach, Decomposition

Both indices can be decomposed into the contribution of changes in mortality by age and causes, using the Horiuchi method.



## Diversity, Second approach

Let's assume that each UDC has its own diversity profile of contributing causes, measured with the Simpson index:

$$S(u) = \sum_c (\hat{d}^u(c))^2$$

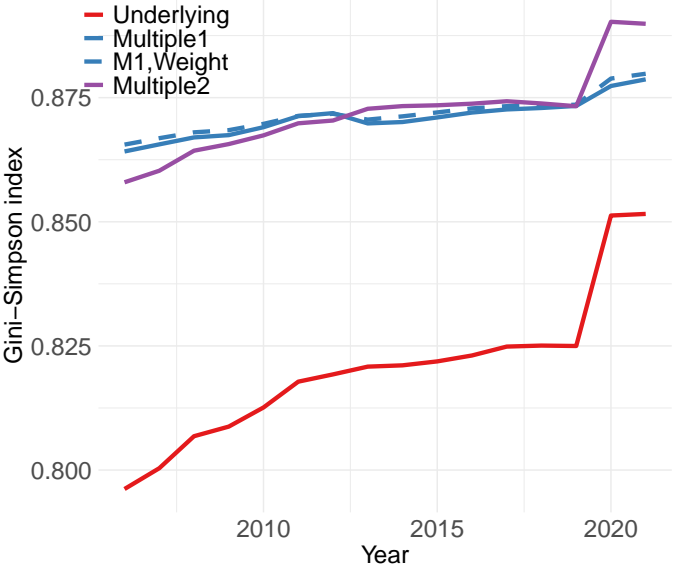
An modified Simpson index can be obtained for all UCD with:

$$S = \sum_u (d(u))^2 S(u)$$

The Gini-Simpson index will then be:

$$GS = 1 - S$$

# Diversity, Second approach, Results



# Diversity, Second approach, Decomposition

The index can be decomposed into changes in  $d(u)$  and changes in  $S(u)$ , using the Kitagawa method.

	$\Delta S(u)$	$\Delta d(u)$	Total
2006-2019	-0.005	0.020	0.015
2019-2021	-0.006	0.022	0.016

# Discussion

- Increased diversity due to both richness and evenness;
- Higher level of diversity when MCoD are considered, with both concepts;
- The increase is, in part, due to a shift towards UCD with more complex and numerous contributing causes;
- Increase in diversity indices explained mainly by mortality decrease from diseases of the circulatory system;
- Increase in  $N(x,u)$  might be due to changes in CoD reporting or changes in the prevalence of multi-morbidity at death;
- All tested variant of S provide similar results.

# Limitations

- Data quality?
- Sensitivity to how we group the causes.
- Which variant of S should be recommended?
- No long-term trends (yet).

# Next steps

- Application to other countries;
- Application by sex;
- Testing distance measures;
- Mapping changes by underlying and contributing causes.



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