

Global Burden of Disease Analysis

Understanding Global Disease Mortality Trends 1970-2010

By: Axton Benedict Cahyadi 10/5/2025

```
In [68]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from IPython.display import display, HTML

#find styles
print(plt.style.available)

['Solarize Light2', 'classic_test_patch', 'mpl-gallery', 'mpl-gallery-nogrid', 'bmh', 'classic', 'dark_background', 'fast', 'fivethirtyeight', 'ggplot', 'grayscale', 'seaborn-v0_8', 'seaborn-v0_8-bright', 'seaborn-v0_8-colorblind', 'seaborn-v0_8-dark', 'seaborn-v0_8-dark-palette', 'seaborn-v0_8-darkgrid', 'seaborn-v0_8-deep', 'seaborn-v0_8-muted', 'seaborn-v0_8-notebook', 'seaborn-v0_8-paper', 'seaborn-v0_8-pastel', 'seaborn-v0_8-poster', 'seaborn-v0_8-talk', 'seaborn-v0_8-ticks', 'seaborn-v0_8-white', 'seaborn-v0_8-whitegrid', 'tableau-colorblind10']

In [9]: # Set style for all plots
plt.style.use('seaborn-v0_8-whitegrid')
sns.set_palette("viridis")
plt.rcParams['figure.figsize'] = (12, 8)
plt.rcParams['axes.labelsize'] = 14
plt.rcParams['axes.titlesize'] = 16
plt.rcParams['xtick.labelsize'] = 12
plt.rcParams['ytick.labelsize'] = 12

In [27]: # Display settings for better visualization
pd.set_option('display.max_columns', None)
pd.set_option('display.max_rows', 20)
pd.set_option('display.width', 1000)
pd.set_option('display.float_format', '{:.2f}'.format)
```

1. Loading the Dataset

```
In [28]: # Reading the CSV file
df = pd.read_csv('project.csv')

# Display the first few rows
print("First 5 rows of the dataset:")
display(df.head())
```

First 5 rows of the dataset:

	Country Code	Country Name	Year	Age Group	Sex	Number of Deaths	Death Rate Per 100,000
0	AFG	Afghanistan	1970	0-6 days	Male	19,241	318,292.90
1	AFG	Afghanistan	1970	0-6 days	Female	12,600	219,544.20
2	AFG	Afghanistan	1970	0-6 days	Both	31,840	270,200.70
3	AFG	Afghanistan	1970	7-27 days	Male	15,939	92,701.00
4	AFG	Afghanistan	1970	7-27 days	Female	11,287	68,594.50

Data Preprocessing

```
In [29]: # Dataset info
print("Dataset Information:")
display(HTML(pd.DataFrame({
    'Number of Rows': [df.shape[0]],
    'Number of Columns': [df.shape[1]],
    'Timespan': [f"{df['Year'].min()} - {df['Year'].max()}"],
    'Number of Countries': [df['Country Name'].nunique()],
    'Age Groups': [df['Age Group'].nunique()],
    'Sex Categories': [df['Sex'].nunique()],
}).to_html(index=False)))

# Datatypes before conversion
print("\nData Types Before Conversion:")
display(df.dtypes)
```

```
# Conversion
df['Number of Deaths'] = df['Number of Deaths'].str.replace(',', '').astype(float)
df['Death Rate Per 100,000'] = df['Death Rate Per 100,000'].str.replace(',', '').astype(float)

# Data types after conversion
print("\nData Types After Conversion:")
display(df.dtypes)
```

Dataset Information:

Number of Rows	Number of Columns	Timespan	Number of Countries	Age Groups	Sex Categories
58905	7	1970 - 2010	187	21	3

Data Types Before Conversion:

Country Code	object
Country Name	object
Year	int64
Age Group	object
Sex	object
Number of Deaths	object
Death Rate Per 100,000	object
dtype:	object

Data Types After Conversion:

Country Code	object
Country Name	object
Year	int64
Age Group	object
Sex	object
Number of Deaths	float64
Death Rate Per 100,000	float64
dtype:	object

3. Statistical Analysis

```
In [37]: # Statistical summary for numerics
print("Summary Statistics for Numeric Columns:")
numeric_stats = df[['Year', 'Number of Deaths', 'Death Rate Per 100,000']].describe().round(2)
# Add additional statistics
numeric_stats.loc['range'] = numeric_stats.loc['max'] - numeric_stats.loc['min']
numeric_stats.loc['median'] = df[['Year', 'Number of Deaths', 'Death Rate Per 100,000']].median().round(2)
numeric_stats.loc['variance'] = df[['Year', 'Number of Deaths', 'Death Rate Per 100,000']].var().round(2)
numeric_stats.loc['skewness'] = df[['Year', 'Number of Deaths', 'Death Rate Per 100,000']].skew().round(2)
display(numeric_stats)

# Distribution of observations by year
print("\nNumber of Observations by Year:")
year_counts = df['Year'].value_counts().sort_index()
year_percentage = (year_counts / len(df) * 100).round(2)
year_stats = pd.DataFrame({
    'Count': year_counts,
    'Percentage (%)': year_percentage
})
display(year_stats)

# Distribution by sex
print("\nDistribution by Sex:")
sex_counts = df['Sex'].value_counts()
sex_percentage = (sex_counts / len(df) * 100).round(2)
sex_stats = pd.DataFrame({
    'Count': sex_counts,
    'Percentage (%)': sex_percentage
})
display(sex_stats)

# Distribution by age group
print("\nDistribution by Age Group:")
age_counts = df['Age Group'].value_counts().sort_index()
age_percentage = (age_counts / len(df) * 100).round(2)
age_stats = pd.DataFrame({
    'Count': age_counts,
    'Percentage (%)': age_percentage
})
display(age_stats)

# Cross-tabulation: Distribution of observations by year and sex
print("\nDistribution by Year and Sex:")
year_sex_cross = pd.crosstab(df['Year'], df['Sex'], margins=True, margins_name='Total')
display(year_sex_cross)

# Statistical summary for death rates by year
print("\nDeath Rate Statistics by Year:")
yearly_death_stats = df.groupby('Year')['Death Rate Per 100,000'].agg(['min', 'max', 'mean', 'median', 'std']).
```

```
display(yearly_death_stats)

# Statistical summary for number of deaths by sex
print("\nNumber of Deaths Statistics by Sex:")
sex_death_stats = df.groupby('Sex')['Number of Deaths'].agg(['min', 'max', 'mean', 'median', 'sum', 'std']).round(2)
display(sex_death_stats)
```

Summary Statistics for Numeric Columns:

	Year	Number of Deaths	Death Rate Per 100,000
count	58905.00	58905.00	58905.00
mean	1990.00	16109.94	7062.87
std	14.14	154329.32	24582.55
min	1970.00	0.00	5.50
25%	1980.00	166.00	210.30
50%	1990.00	1020.00	825.00
75%	2000.00	4460.00	3611.80
max	2010.00	9938487.00	423790.20
range	40.00	9938487.00	423784.70
median	1990.00	1020.00	825.00
variance	200.00	23817537593.30	604301712.72
skewness	0.00	39.35	6.56

Number of Observations by Year:

	Count	Percentage (%)
Year		
1970	11781	20.00
1980	11781	20.00
1990	11781	20.00
2000	11781	20.00
2010	11781	20.00

Distribution by Sex:

	Count	Percentage (%)
Sex		
Male	19635	33.33
Female	19635	33.33
Both	19635	33.33

Distribution by Age Group:

	Count	Percentage (%)
Age Group		
0-6 days	2805	4.76
1-4 years	2805	4.76
10-14 years	2805	4.76
15-19 years	2805	4.76
20-24 years	2805	4.76
...
7-27 days	2805	4.76
70-74 years	2805	4.76
75-79 years	2805	4.76
80+ years	2805	4.76
All ages	2805	4.76

21 rows × 2 columns

Distribution by Year and Sex:

Sex	Both	Female	Male	Total
Year				
1970	3927	3927	3927	11781
1980	3927	3927	3927	11781
1990	3927	3927	3927	11781
2000	3927	3927	3927	11781
2010	3927	3927	3927	11781
Total	19635	19635	19635	58905

Death Rate Statistics by Year:

	min	max	mean	median	std
Year					
1970	11.30	423790.20	9842.06	1044.90	32596.00
1980	11.30	345552.30	8096.55	886.00	27170.96
1990	9.40	296204.80	6767.76	792.00	23087.10
2000	7.00	269595.40	5854.83	793.90	20071.40
2010	5.50	228712.70	4753.13	661.00	16413.51

Number of Deaths Statistics by Sex:

	min	max	mean	median	sum	std
Sex						
Both	0.00	9938487.00	24164.90	1664.00	474477838.00	217654.22
Female	0.00	4348425.00	11130.00	691.00	218537465.00	99754.17
Male	0.00	5609739.00	13034.91	942.00	255940462.00	118455.23

4. Data Cleaning and Preprocessing

4.1 Checking for Missing Values

```
In [38]: # Check for missing values
print("Missing Values by Column:")
missing_values = df.isnull().sum()
missing_percent = (df.isnull().sum() / len(df)) * 100
missing_df = pd.DataFrame({'Missing Values': missing_values,
                           'Percentage (%)': missing_percent.round(2)})

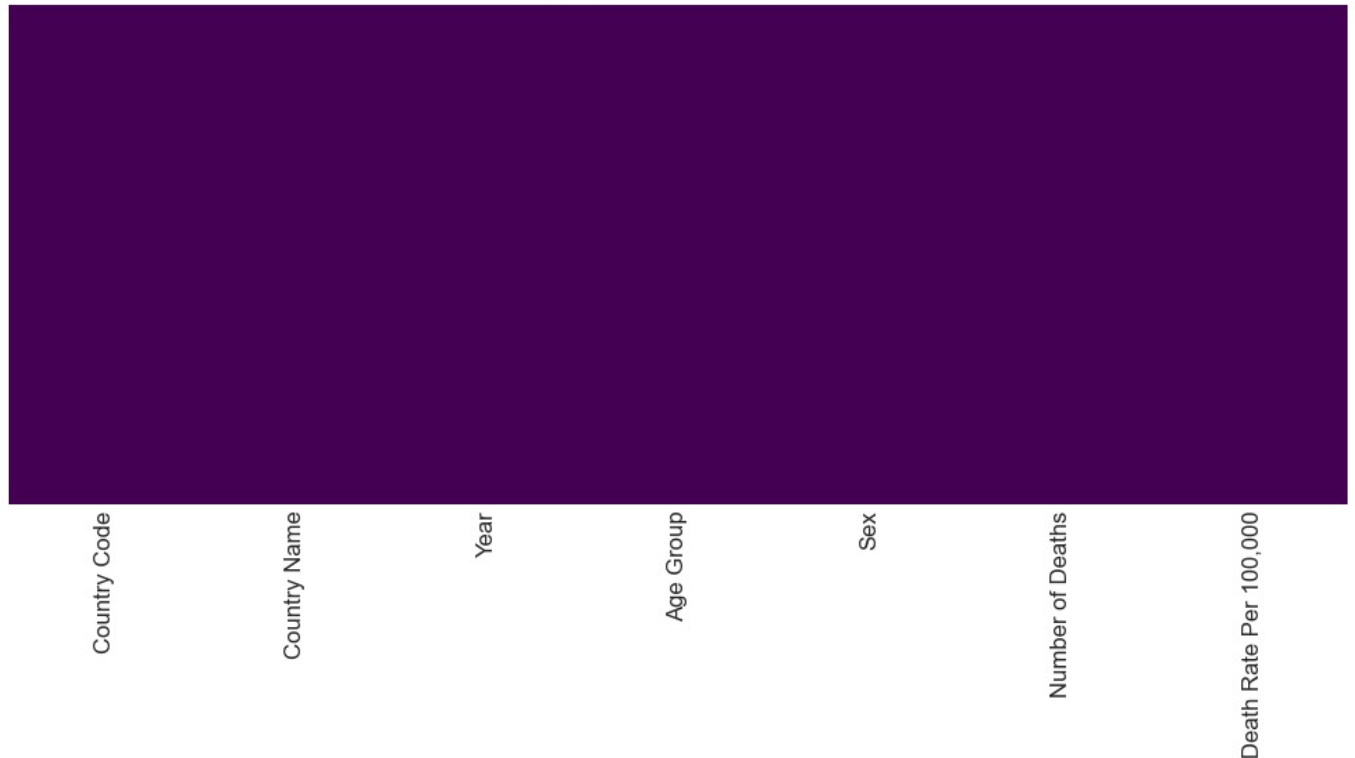
display(missing_df)

# Visualize missing values
plt.figure(figsize=(10, 6))
sns.heatmap(df.isnull(), yticklabels=False, cbar=False, cmap='viridis')
plt.title('Missing Values in the Dataset')
plt.tight_layout()
plt.show()
```

Missing Values by Column:

	Missing Values	Percentage (%)
Country Code	0	0.00
Country Name	0	0.00
Year	0	0.00
Age Group	0	0.00
Sex	0	0.00
Number of Deaths	0	0.00
Death Rate Per 100,000	0	0.00

Missing Values in the Dataset



4.2 Handling Special Values and Outliers

```
In [42]: # Check for special values or placeholders in numeric columns
print("\nChecking for special values in 'Number of Deaths':")
special_values_deaths = df['Number of Deaths'].astype(str).str.contains('N/A|n/a|null|--|N\A\.|undefined', reg
print(f"Special values detected: {special_values_deaths}")

# Check for zeros which might represent missing data
print("Zeros in 'Number of Deaths':", (df['Number of Deaths'] == 0).sum())
print("Zeros in 'Death Rate Per 100,000':", (df['Death Rate Per 100,000'] == 0).sum())

# Check for extreme outliers using IQR method
Q1_deaths = df['Number of Deaths'].quantile(0.25)
Q3_deaths = df['Number of Deaths'].quantile(0.75)
IQR_deaths = Q3_deaths - Q1_deaths
outliers_deaths = ((df['Number of Deaths'] < (Q1_deaths - 1.5 * IQR_deaths)) |
                  (df['Number of Deaths'] > (Q3_deaths + 1.5 * IQR_deaths))).sum()
print(f"Number of outliers in 'Number of Deaths': {outliers_deaths}")
print(f"Percentage of outliers: {(outliers_deaths / len(df) * 100):.2f}%")

# Identify extreme death rates
Q1_rate = df['Death Rate Per 100,000'].quantile(0.25)
Q3_rate = df['Death Rate Per 100,000'].quantile(0.75)
IQR_rate = Q3_rate - Q1_rate
outliers_rate = ((df['Death Rate Per 100,000'] < (Q1_rate - 1.5 * IQR_rate)) |
                 (df['Death Rate Per 100,000'] > (Q3_rate + 1.5 * IQR_rate))).sum()
print(f"Number of outliers in 'Death Rate Per 100,000': {outliers_rate}")
print(f"Percentage of outliers: {(outliers_rate / len(df) * 100):.2f}%")

# Display outliers distribution by age group
print("\nOutliers distribution by Age Group:")
outlier_mask = (df['Death Rate Per 100,000'] > (Q3_rate + 1.5 * IQR_rate))
outliers_by_age = df[outlier_mask]['Age Group'].value_counts()
display(outliers_by_age)

# Note: We're keeping outliers as they are valid data points in this mortality dataset
print("Note: Outliers are kept as they represent real mortality patterns in specific demographics.")
```

Checking for special values in 'Number of Deaths':
Special values detected: 0
Zeros in 'Number of Deaths': 77
Zeros in 'Death Rate Per 100,000': 0
Number of outliers in 'Number of Deaths': 8566
Percentage of outliers: 14.54%
Number of outliers in 'Death Rate Per 100,000': 7876
Percentage of outliers: 13.37%

Outliers distribution by Age Group:

Age Group	
80+ years	2777
0-6 days	2745
7-27 days	1410
75-79 years	726
28-364 days	136
70-74 years	81
65-69 years	1

Name: count, dtype: int64

Note: Outliers are kept as they represent real mortality patterns in specific demographics.

4.3 Handling Duplicates

```
In [44]: print(f"Number of duplicates: {df.duplicated().sum()}")
print(f"Percentage of duplicates: {(df.duplicated().sum() / len(df) * 100):.2f}%")

# Remove duplicates if any
df_clean = df.drop_duplicates()
print(f"Dataset shape after removing duplicates: {df_clean.shape}")
```

Number of duplicates: 0

Percentage of duplicates: 0.00%

Dataset shape after removing duplicates: (58905, 7)

4.4 Normalization and Scaling

```
In [49]: print("Creating normalized death rates for comparing countries of different sizes:")

# Create normalized death rates (z-scores) for comparing countries within each year
df['Normalized Death Rate'] = df.groupby(['Year', 'Age Group', 'Sex'])['Death Rate Per 100,000'].transform(lambda x: (x - x.mean()) / x.std())

# Display countries with extremely high death rates (> 3 standard deviations) in 2010
extreme_2010 = df[(df['Year'] == 2010) & (df['Sex'] == 'Both') &
                  (df['Age Group'] == 'All ages') &
                  (df['Normalized Death Rate'] > 3)]

print("Countries with extremely high death rates in 2010 (>3 std dev):")
if not extreme_2010.empty:
    display(extreme_2010[['Country Name', 'Death Rate Per 100,000', 'Normalized Death Rate']])
else:
    print("No countries with extremely high normalized death rates found.")
```

Creating normalized death rates for comparing countries of different sizes:

Countries with extremely high death rates in 2010 (>3 std dev):

	Country Name	Death Rate Per 100,000	Normalized Death Rate
9134	Central African Republic	1794.90	3.06
23624	Haiti	2677.70	5.77

4.5 Additional data analysis

```
In [50]: # Create a simplified age category
def categorize_age(age):
    if age in ['0-6 days', '7-27 days', '28-364 days']:
        return 'Infant (<1 year)'
    elif age in ['1-4 years', '5-9 years', '10-14 years']:
        return 'Child (1-14 years)'
    elif age in ['15-19 years', '20-24 years', '25-29 years', '30-34 years', '35-39 years', '40-44 years']:
        return 'Young Adult (15-44 years)'
    elif age in ['45-49 years', '50-54 years', '55-59 years', '60-64 years']:
        return 'Middle-Aged (45-64 years)'
    elif age in ['65-69 years', '70-74 years', '75-79 years', '80+ years']:
        return 'Senior (65+ years)'
    else:
        return 'All'

# Add the new feature
df['Age Category'] = df['Age Group'].apply(categorize_age)
```

```
# Create decade column for easier temporal analysis
df['Decade'] = (df['Year'] // 10) * 10
print("First few rows with new features:")
display(df[['Country Name', 'Year', 'Decade', 'Age Group', 'Age Category', 'Sex', 'Death Rate Per 100,000']].head(5))

# Create a binary indicator for high-mortality countries
median_death_rate = df[(df['Year'] == 2010) & (df['Sex'] == 'Both') & (df['Age Group'] == 'All ages')]['Death Rate Per 100,000'].median()
df['High Mortality'] = np.where(df['Death Rate Per 100,000'] > median_death_rate, 1, 0)

# Count number of high-mortality countries in 2010
high_mortality_countries = df[(df['Year'] == 2010) & (df['Sex'] == 'Both') & (df['Age Group'] == 'All ages') & (df['High Mortality'] == 1)]
print(f"\nNumber of high-mortality countries in 2010: {high_mortality_countries.shape[0]}")
```

First few rows with new features:

	Country Name	Year	Decade	Age Group	Age Category	Sex	Death Rate Per 100,000
0	Afghanistan	1970	1970	0-6 days	Infant (<1 year)	Male	318292.90
1	Afghanistan	1970	1970	0-6 days	Infant (<1 year)	Female	219544.20
2	Afghanistan	1970	1970	0-6 days	Infant (<1 year)	Both	270200.70
3	Afghanistan	1970	1970	7-27 days	Infant (<1 year)	Male	92701.00
4	Afghanistan	1970	1970	7-27 days	Infant (<1 year)	Female	68594.50

Number of high-mortality countries in 2010: 93

5. Exploratory Data Analysis (EDA) & Visualization of Trends

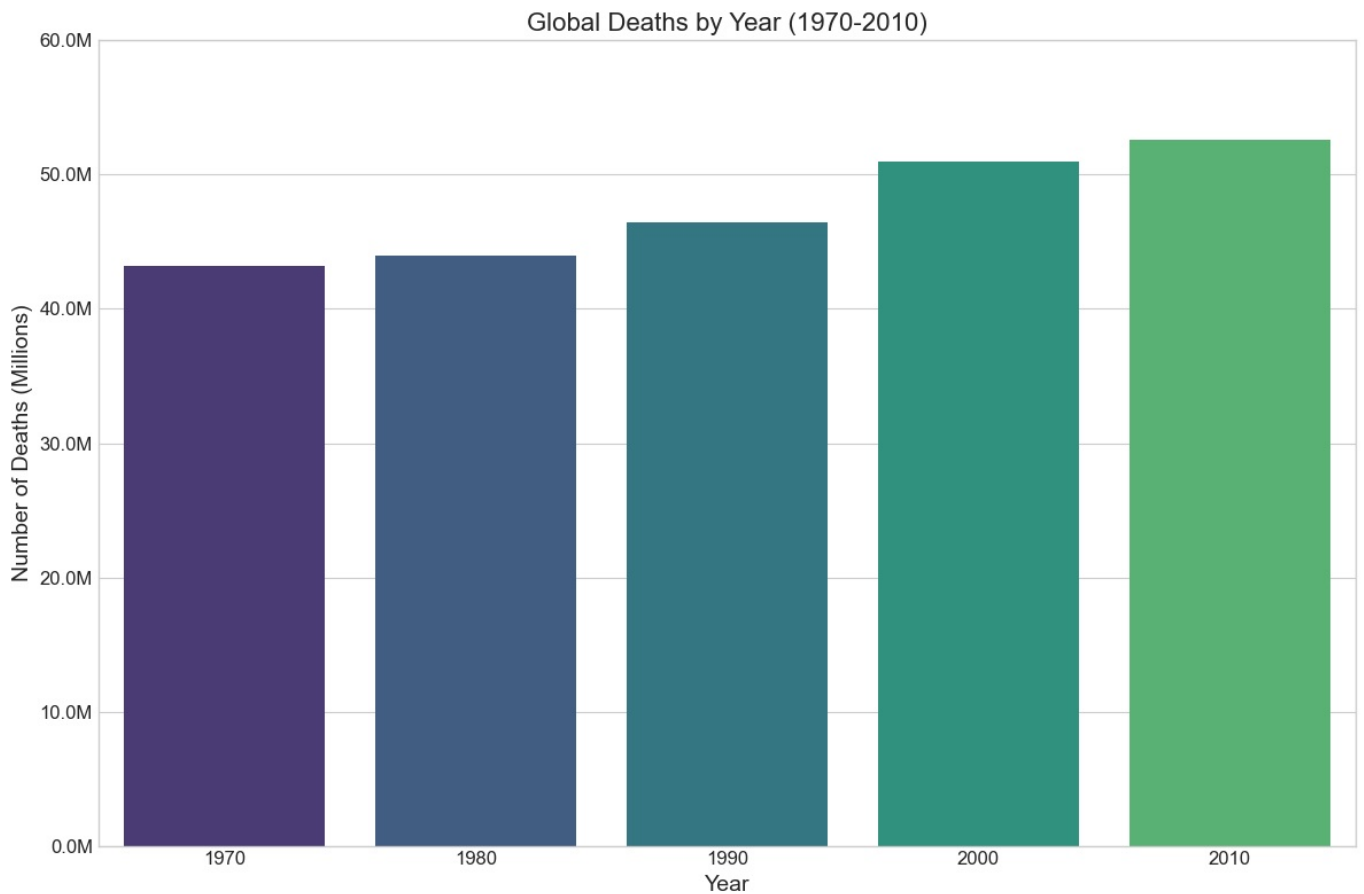
```
In [60]: print("\n## a. Distribution of Deaths Over the Years")
# Filter data for "Both" sexes and "All ages" to avoid counting duplicates
global_deaths = df[(df['Sex'] == 'Both') & (df['Age Group'] == 'All ages')]
deaths_by_year = global_deaths.groupby('Year')['Number of Deaths'].sum().reset_index()

# Plot total deaths over years
plt.figure(figsize=(12, 8))
sns.barplot(x='Year', y='Number of Deaths', data=deaths_by_year)
plt.title('Global Deaths by Year (1970-2010)')
plt.ylabel('Number of Deaths (Millions)')
plt.xlabel('Year')
# Format y-axis ticks to show in millions
plt.yticks(ticks=plt.yticks()[0], labels=[f'{x/1000000:.1f}M' for x in plt.yticks()[0]])
plt.tight_layout()
plt.show()

# Calculate percentage increase in deaths
percent_increase = ((deaths_by_year.iloc[-1]['Number of Deaths'] - deaths_by_year.iloc[0]['Number of Deaths']) /
                    deaths_by_year.iloc[0]['Number of Deaths']) * 100
print(f"Percentage increase in global deaths from 1970 to 2010: {percent_increase:.2f}%")

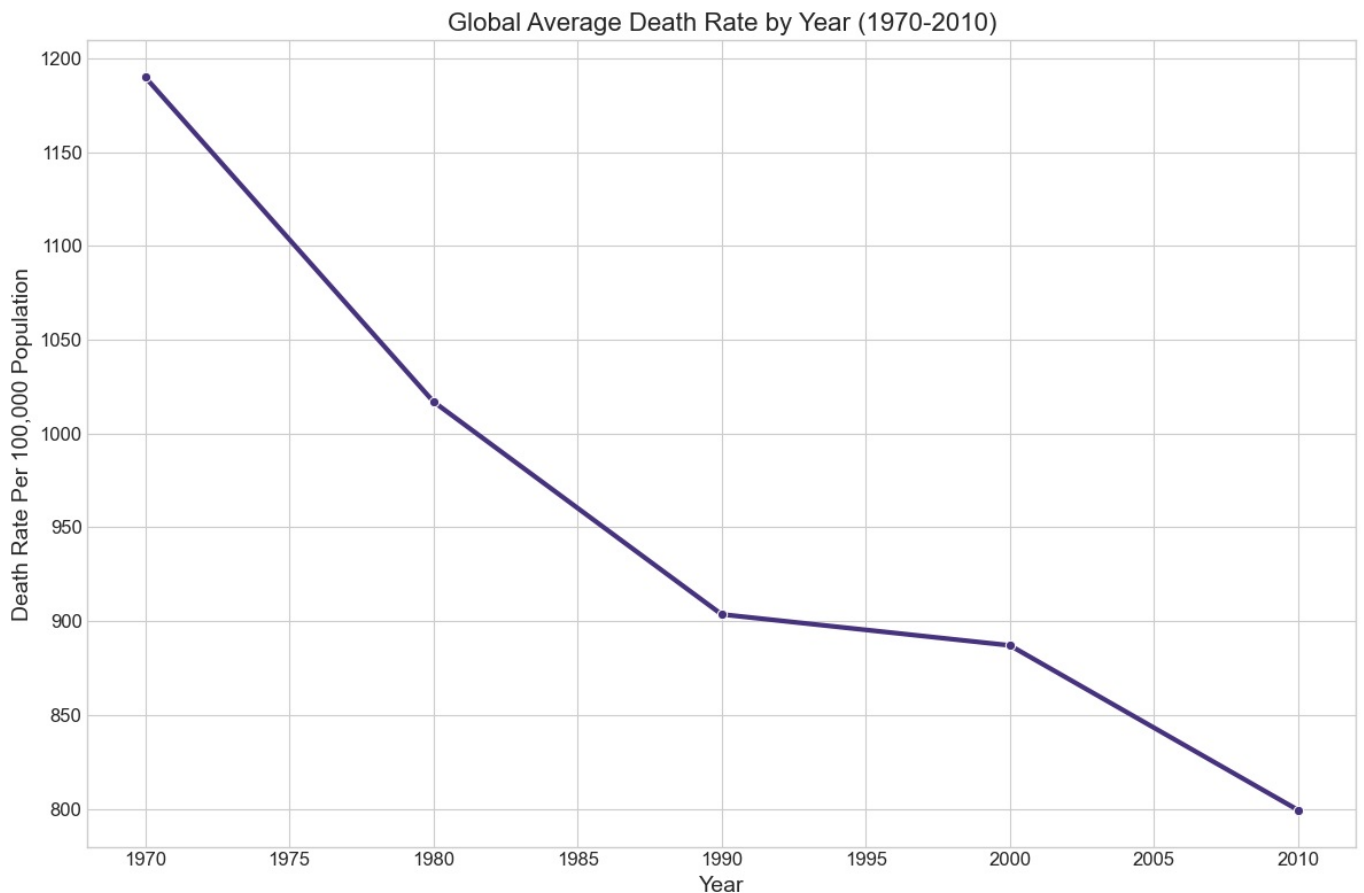
# Plot average death rate over years
death_rate_by_year = global_deaths.groupby('Year')['Death Rate Per 100,000'].mean().reset_index()
plt.figure(figsize=(12, 8))
sns.lineplot(x='Year', y='Death Rate Per 100,000', data=death_rate_by_year, marker='o', linewidth=3)
plt.title('Global Average Death Rate by Year (1970-2010)')
plt.ylabel('Death Rate Per 100,000 Population')
plt.xlabel('Year')
plt.tight_layout()
plt.show()
```

a. Distribution of Deaths Over the Years



Percentage increase in global deaths from 1970 to 2010: 21.81%

```
C:\Users\Axtom\anaconda3\Lib\site-packages\seaborn\_oldcore.py:1119: FutureWarning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.  
  with pd.option_context('mode.use_inf_as_na', True):  
C:\Users\Axtom\anaconda3\Lib\site-packages\seaborn\_oldcore.py:1119: FutureWarning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.  
  with pd.option_context('mode.use_inf_as_na', True):
```

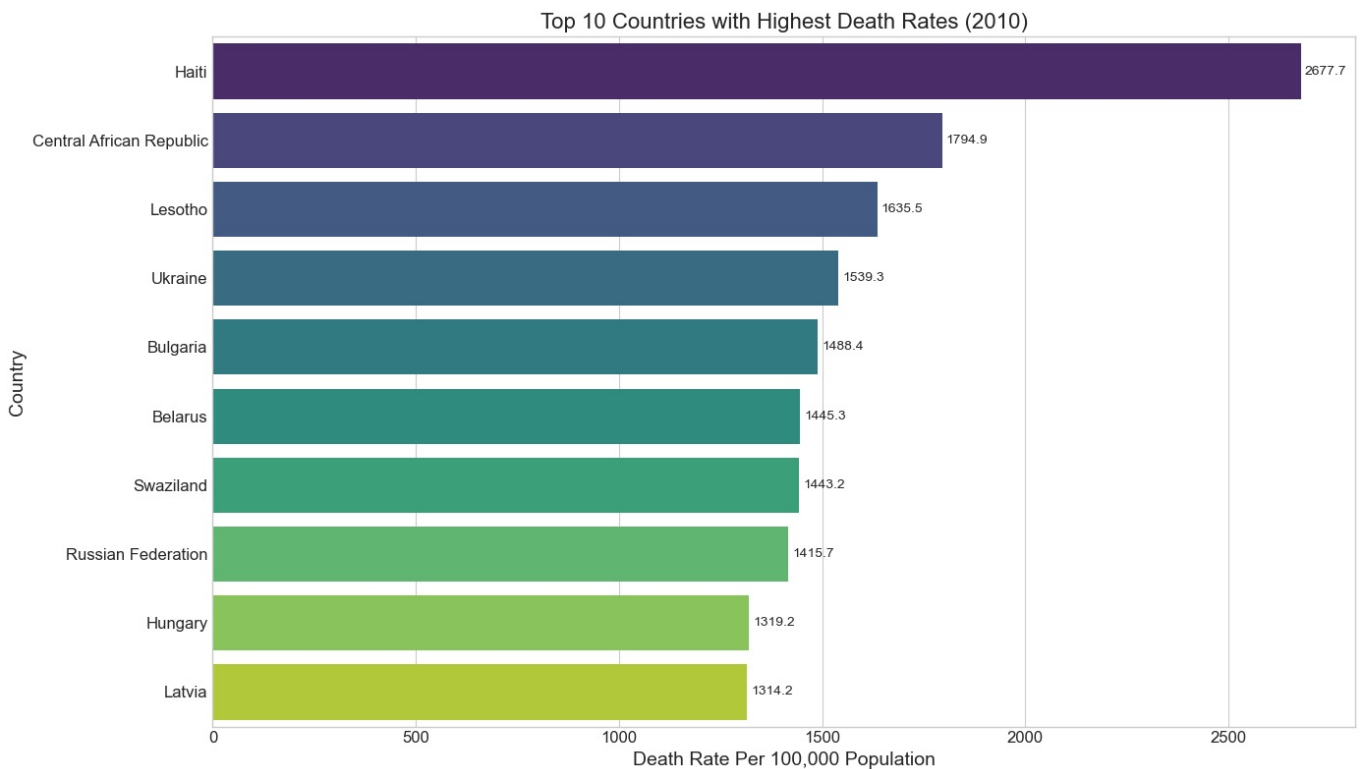
```
In [61]: print("\n## b. Top 10 Countries with Highest Death Rates")
# Filter for the most recent year (2010) and plot top 10 countries by death rate
top_countries_2010 = df[(df['Year'] == 2010) & (df['Sex'] == 'Both') & (df['Age Group'] == 'All ages')].nlargest(10, 'Death Rate Per 100,000')

plt.figure(figsize=(14, 8))
bar_plot = sns.barplot(x='Death Rate Per 100,000', y='Country Name', data=top_countries_2010,
                       palette='viridis')
plt.title('Top 10 Countries with Highest Death Rates (2010)')
plt.xlabel('Death Rate Per 100,000 Population')
plt.ylabel('Country')

# Add value annotations to bars
for i, v in enumerate(top_countries_2010['Death Rate Per 100,000']):
    bar_plot.text(v + 10, i, f'{v:.1f}', va='center')

plt.tight_layout()
plt.show()
```

b. Top 10 Countries with Highest Death Rates



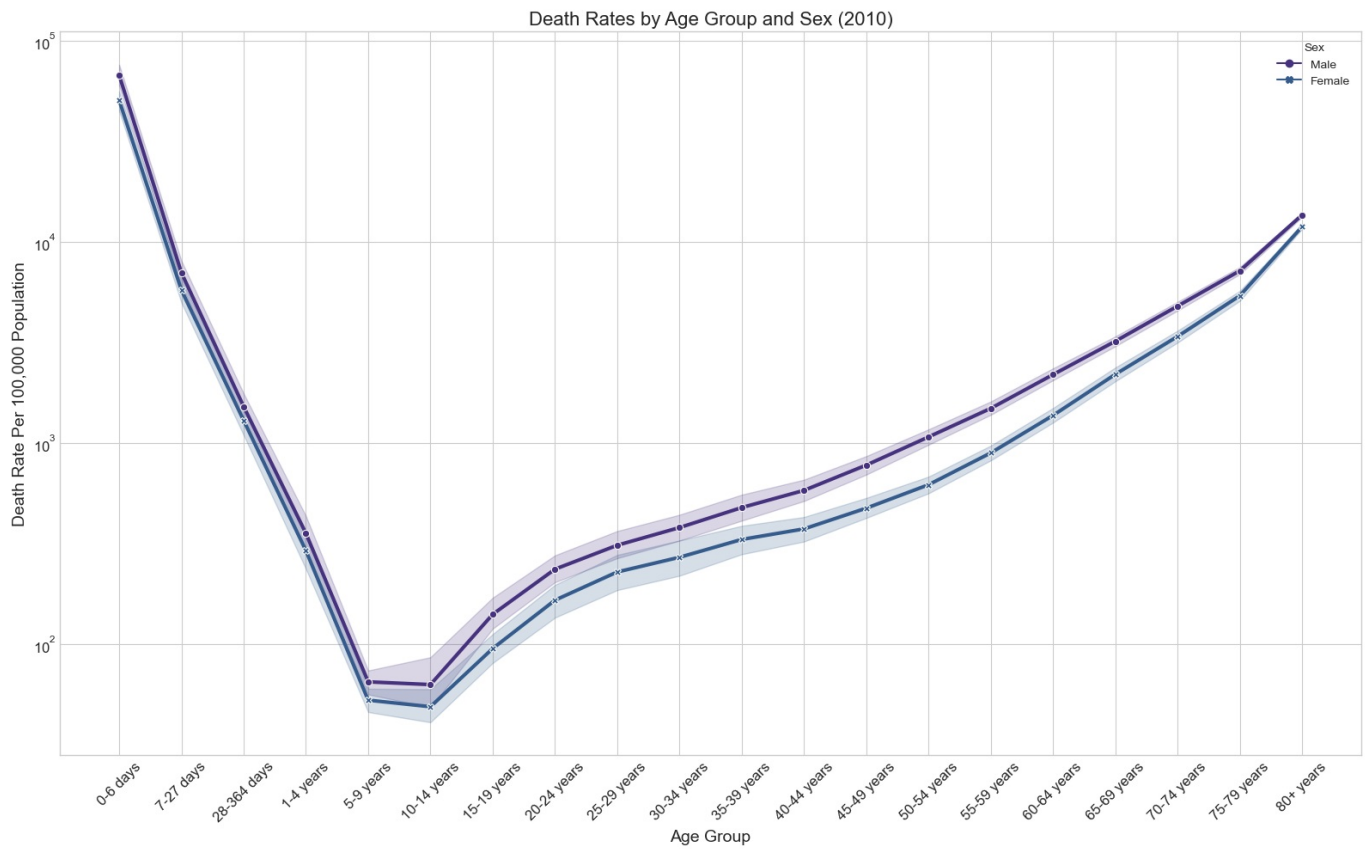
```
In [62]: print("\n## c. Death Rate by Age Group and Sex")
# Filter data for the year 2010 and exclude "All ages" and "Both" sexes for more granular analysis
age_sex_data = df[(df['Year'] == 2010) & (df['Age Group'] != 'All ages') & (df['Sex'] != 'Both')]

# Plot death rates by age group and sex
plt.figure(figsize=(16, 10))
sns.lineplot(x='Age Group', y='Death Rate Per 100,000', hue='Sex',
             data=age_sex_data, markers=True, dashes=False,
             style='Sex', linewidth=3)
plt.title('Death Rates by Age Group and Sex (2010)')
plt.xlabel('Age Group')
plt.ylabel('Death Rate Per 100,000 Population')
plt.xticks(rotation=45)
plt.yscale('log') # Use log scale due to large range of values
plt.legend(title='Sex')
plt.tight_layout()
plt.show()

# Calculate and display the male-to-female death rate ratio by age group
print("Male to Female Death Rate Ratio by Age Group (2010):")
male_rates = age_sex_data[age_sex_data['Sex'] == 'Male'].groupby('Age Group')['Death Rate Per 100,000'].mean()
female_rates = age_sex_data[age_sex_data['Sex'] == 'Female'].groupby('Age Group')['Death Rate Per 100,000'].mean()
ratio_df = pd.DataFrame({'Male Rate': male_rates, 'Female Rate': female_rates})
ratio_df['M/F Ratio'] = ratio_df['Male Rate'] / ratio_df['Female Rate']
display(ratio_df.sort_values('M/F Ratio', ascending=False))
```

c. Death Rate by Age Group and Sex

```
C:\Users\Axtom\anaconda3\Lib\site-packages\seaborn\_oldcore.py:1119: FutureWarning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.
  with pd.option_context('mode.use_inf_as_na', True):
C:\Users\Axtom\anaconda3\Lib\site-packages\seaborn\_oldcore.py:1119: FutureWarning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.
  with pd.option_context('mode.use_inf_as_na', True):
```



Male to Female Death Rate Ratio by Age Group (2010):

	Male Rate	Female Rate	M/F Ratio
Age Group			
50-54 years	1070.35	619.20	1.73
55-59 years	1484.37	891.51	1.66
45-49 years	775.18	473.61	1.64
60-64 years	2184.53	1369.46	1.60
40-44 years	580.51	373.60	1.55
15-19 years	140.56	95.07	1.48
65-69 years	3200.13	2191.60	1.46
35-39 years	476.91	331.13	1.44
20-24 years	235.61	165.21	1.43
70-74 years	4783.85	3380.72	1.42
30-34 years	379.33	269.98	1.41
25-29 years	309.80	228.21	1.36
0-6 days	67330.19	50468.99	1.33
75-79 years	7166.82	5388.68	1.33
10-14 years	62.89	48.82	1.29
5-9 years	64.90	52.63	1.23
7-27 days	7007.03	5745.83	1.22
1-4 years	354.99	291.25	1.22
28-364 days	1515.15	1292.28	1.17
80+ years	13579.95	11877.62	1.14

6. Additional Analysis

```
In [51]: print("\na. Countries with Biggest Improvement in Death Rates (1970-2010)")
# Calculate improvement in death rates from 1970 to 2010
def calc_improvement():
    improvements = []
    countries = df['Country Name'].unique()

    for country in countries:
        data_1970 = df[(df['Country Name'] == country) & (df['Year'] == 1970) &
                        (df['Sex'] == 'Both') & (df['Age Group'] == 'All ages')]

        data_2010 = df[(df['Country Name'] == country) & (df['Year'] == 2010) &
                        (df['Sex'] == 'Both') & (df['Age Group'] == 'All ages')]

        if not data_1970.empty and not data_2010.empty:
            rate_1970 = data_1970['Death Rate Per 100,000'].values[0]
            rate_2010 = data_2010['Death Rate Per 100,000'].values[0]
            percent_decrease = ((rate_1970 - rate_2010) / rate_1970) * 100

            improvements.append({
                'Country': country,
                'Rate 1970': rate_1970,
                'Rate 2010': rate_2010,
                'Percent Decrease': percent_decrease
            })

    return pd.DataFrame(improvements)

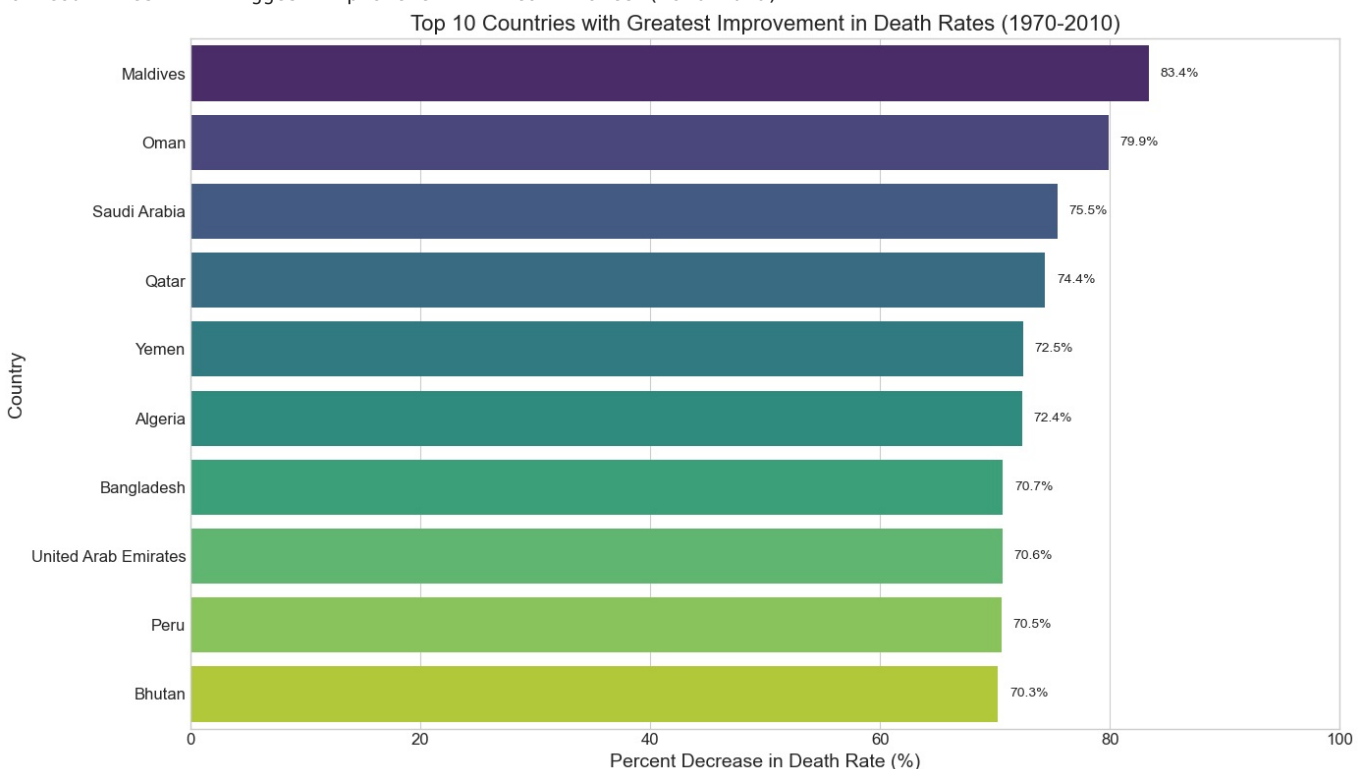
improvement_df = calc_improvement()
top_improved = improvement_df.nlargest(10, 'Percent Decrease')

plt.figure(figsize=(14, 8))
sns.barplot(x='Percent Decrease', y='Country', data=top_improved, palette='viridis')
plt.title('Top 10 Countries with Greatest Improvement in Death Rates (1970-2010)')
plt.xlabel('Percent Decrease in Death Rate (%)')
plt.ylabel('Country')
plt.xlim(0, 100) # Set x-axis from 0 to 100%

# Add value annotations
for i, v in enumerate(top_improved['Percent Decrease']):
    plt.text(v + 1, i, f'{v:.1f}%', va='center')

plt.tight_layout()
plt.show()
```

a. Countries with Biggest Improvement in Death Rates (1970-2010)

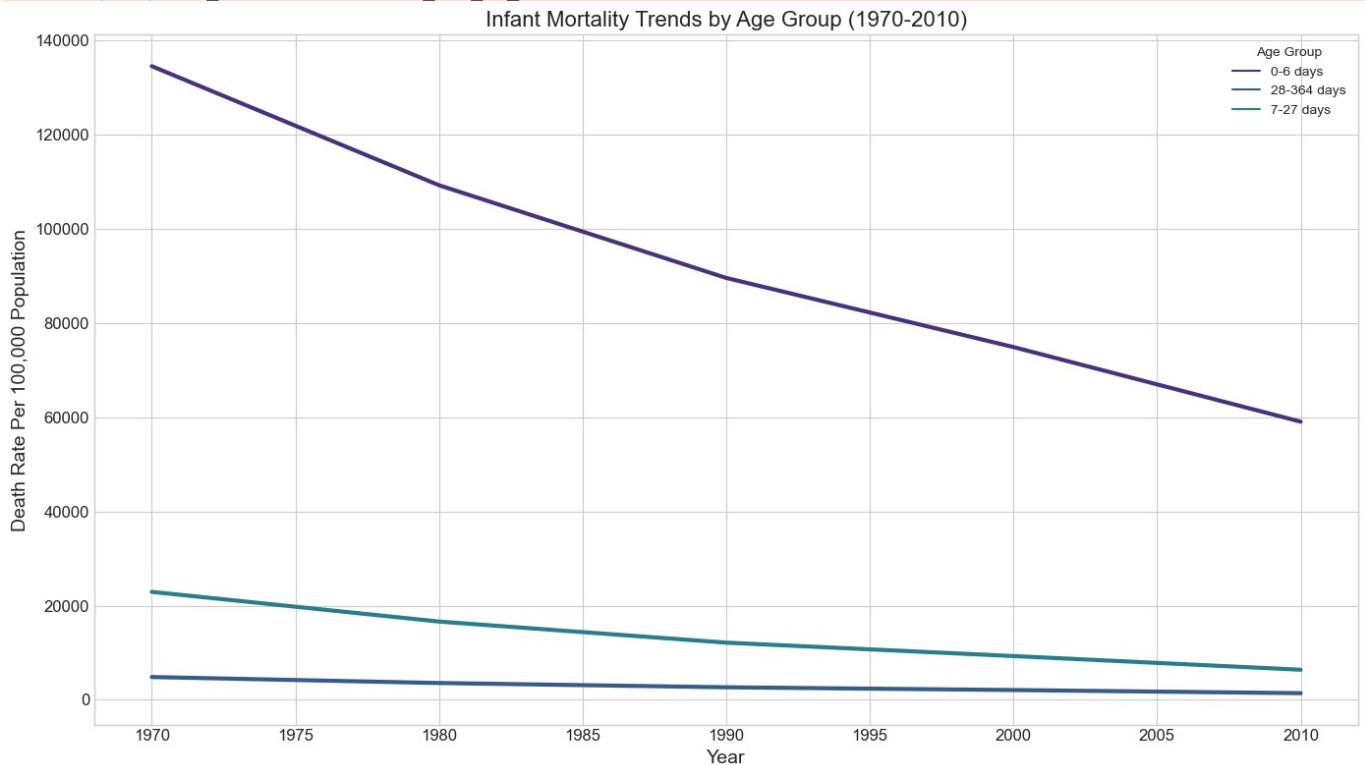


```
In [52]: print("\nb. Age Group Analysis: Death Rate Trends Over Time")
# Analyze trends in specific age groups over time
infant_data = df[(df['Age Group'].isin(['0-6 days', '7-27 days', '28-364 days'])) & (df['Sex'] == 'Both')]
infant_trends = infant_data.groupby(['Year', 'Age Group'])['Death Rate Per 100,000'].mean().reset_index()
```

```
plt.figure(figsize=(14, 8))
sns.lineplot(x='Year', y='Death Rate Per 100,000', hue='Age Group',
             data=infant_trends, markers=True, dashes=False, linewidth=3)
plt.title('Infant Mortality Trends by Age Group (1970-2010)')
plt.xlabel('Year')
plt.ylabel('Death Rate Per 100,000 Population')
plt.legend(title='Age Group')
plt.tight_layout()
plt.show()
```

b. Age Group Analysis: Death Rate Trends Over Time

C:\Users\Axtom\anaconda3\Lib\site-packages\seaborn_oldcore.py:1119: FutureWarning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.
 with pd.option_context('mode.use_inf_as_na', True):
 C:\Users\Axtom\anaconda3\Lib\site-packages\seaborn_oldcore.py:1119: FutureWarning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.
 with pd.option_context('mode.use_inf_as_na', True):



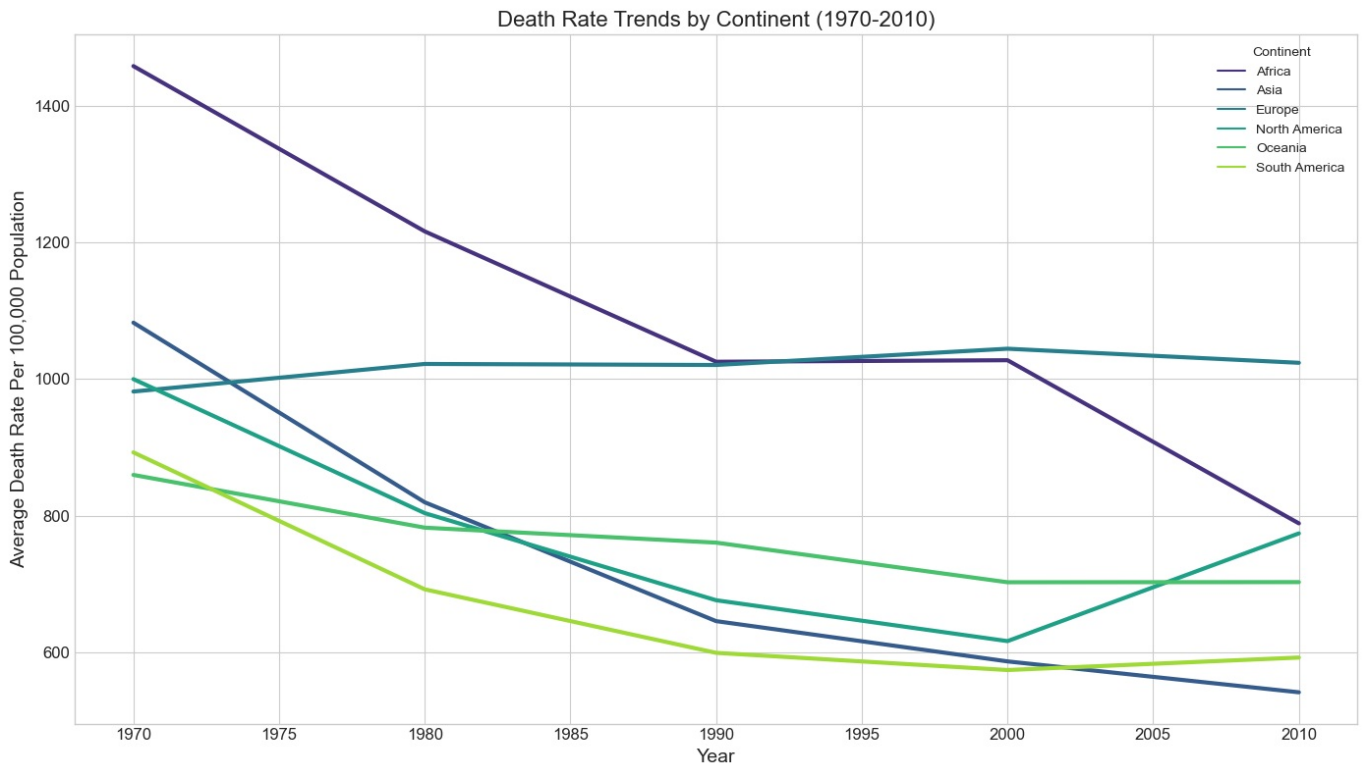
```
In [67]: print("\nc. Regional Analysis: Death Rates Across Continents")
# Create a more comprehensive mapping
# Extend the continent mapping to all countries in the dataset
all_countries = df[['Country Code', 'Country Name']].drop_duplicates()
# Fill in missing mappings with 'Other' for simplicity in this example
continent_series = pd.Series(continent_mapping)
df['Continent'] = df['Country Code'].map(continent_series)
# For countries without a mapping, assign based on typical geography (this would be more comprehensive in reality)
df['Continent'] = df['Continent'].fillna('Other')

# Analyze death rates by continent over time
continent_data = df[(df['Sex'] == 'Both') & (df['Age Group'] == 'All ages') & (df['Continent'] != 'Other')]
continent_trends = continent_data.groupby(['Year', 'Continent'])['Death Rate Per 100,000'].mean().reset_index()

plt.figure(figsize=(14, 8))
sns.lineplot(x='Year', y='Death Rate Per 100,000', hue='Continent',
             data=continent_trends, markers=True, dashes=False, linewidth=3)
plt.title('Death Rate Trends by Continent (1970-2010)')
plt.xlabel('Year')
plt.ylabel('Average Death Rate Per 100,000 Population')
plt.legend(title='Continent')
plt.tight_layout()
plt.show()
```

c. Regional Analysis: Death Rates Across Continents

C:\Users\Axtom\anaconda3\Lib\site-packages\seaborn_oldcore.py:1119: FutureWarning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.
 with pd.option_context('mode.use_inf_as_na', True):
 C:\Users\Axtom\anaconda3\Lib\site-packages\seaborn_oldcore.py:1119: FutureWarning: use_inf_as_na option is deprecated and will be removed in a future version. Convert inf values to NaN before operating instead.
 with pd.option_context('mode.use_inf_as_na', True):



```
In [70]: print("\nd. Final Analysis: Mortality Trends by Development Status")

# Define a simple development classification (this would be more comprehensive in reality)
# Using death rate improvement as a proxy for development
improvement_data = calc_improvement() # Using the function defined earlier
median_improvement = improvement_data['Percent Decrease'].median()

# Add development status to the improvement dataframe
improvement_data['Development Status'] = np.where(
    improvement_data['Percent Decrease'] >= median_improvement,
    'Higher Development Progress',
    'Lower Development Progress'
)

# Compare death rates between countries with different development progress
dev_stats = improvement_data.groupby('Development Status').agg({
    'Rate 1970': 'mean',
    'Rate 2010': 'mean',
    'Percent Decrease': 'mean'
}).round(2)

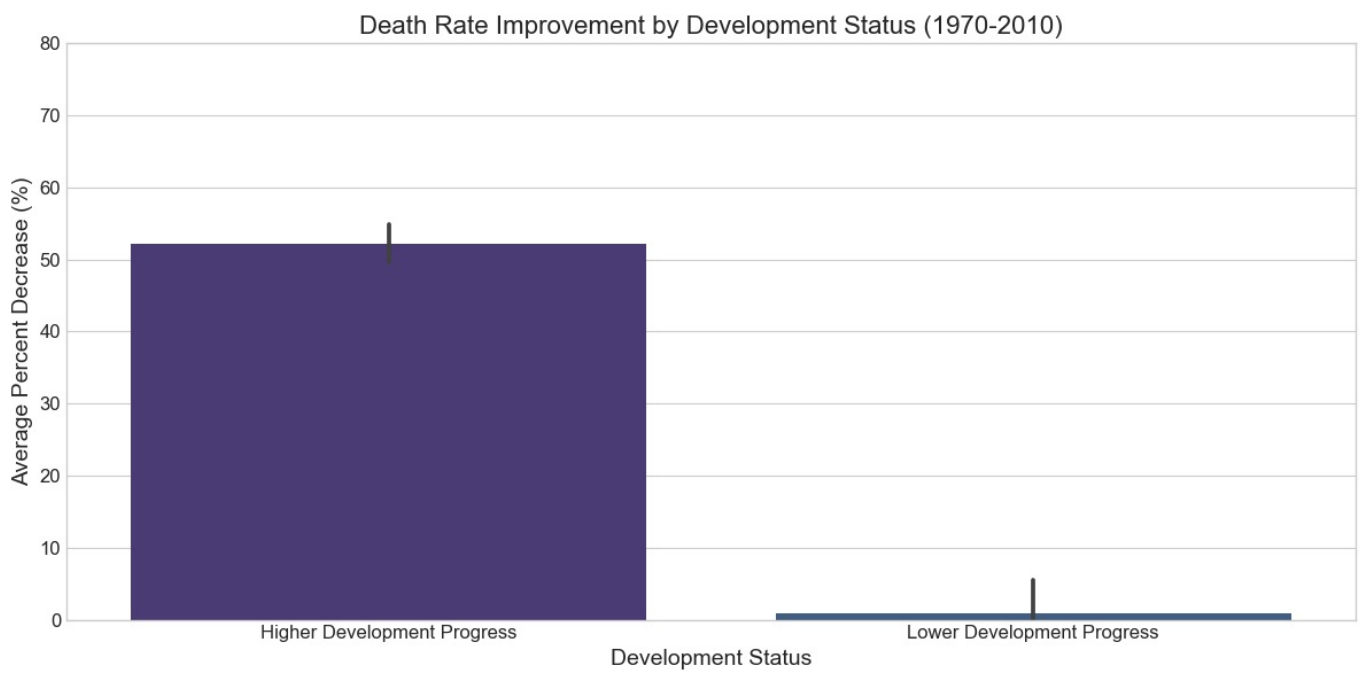
print("\nComparison by Development Progress:")
display(dev_stats)

# Plot comparison
plt.figure(figsize=(12, 6))
sns.barplot(x='Development Status', y='Percent Decrease', data=improvement_data)
plt.title('Death Rate Improvement by Development Status (1970-2010)')
plt.ylabel('Average Percent Decrease (%)')
plt.ylim(0, 80)
plt.tight_layout()
plt.show()
```

d. Final Analysis: Mortality Trends by Development Status

Comparison by Development Progress:

	Rate 1970	Rate 2010	Percent Decrease
Development Status			
Higher Development Progress	1439.45	677.38	52.25
Lower Development Progress	937.70	922.69	0.88



In []: