Using the Years-of-Healthy-Life Measure to Calculate QALYs

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**Background:** The quality-adjusted life year (QALY) is an attractive outcome measure because it captures both health-related quality of life (HRQL) and life expectancy in a single metric. We present a method for calculating QALYs that is simple, utilizes data that are free of charge, and may improve consistency in burden-of-disease investigations.

**Methods:** For purposes of illustration, we calculated the burden of disease due to stroke using two abridged life tables, each adjusted for HRQL. The first life table was generated using all-cause mortality and morbidity data (a reference cohort) and the second was generated using all diseases except stroke (a stroke-free cohort). The difference in total QALYs and in quality-adjusted life expectancy (QALE) was determined by subtraction.

**Results:** Approximately 61,328 (95% CI = 60,272, 62,383) QALYs were lost to stroke in the life-table cohort. Stroke is responsible for a decrement of 0.03 years of life expectancy and 0.61 years of QALE in the United States.

**Conclusions:** The “years of health life” measure affords a rapid, inexpensive, and sensitive means for estimating the burden of disease for local health priorities and may assist research efforts in including QALYs as an outcome measure.


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**Introduction**

Because the quality-adjusted life year (QALY) can capture both health-related quality of life (HRQL) and life expectancy in a single metric, it is an attractive outcome measure for cost-effectiveness analyses and burden-of-disease calculations.1 However, since calculating QALYs may be complicated, expensive, and time consuming,2 some investigators may be reluctant to include them in their research. This may be especially true when limited funding is available with which to conduct a study or when rapid burden-of-disease assessments are needed. Moreover, burden-of-disease estimates may not be comparable when different data sources are used and may not fully capture comorbid illness or disability when data are limited. Techniques that permit ready calculation of QALYs using comprehensive data are therefore helpful to investigations intended to support public health agency resource allocation decisions under conditions of uncertainty.

The National Health Interview Survey (NHIS) is an annual survey of the U.S. population containing self-reported illness, role function, and perceived health status. A QALY-compatible research tool, the “years of healthy life” (YHL) measure has been used to generate a simple “off-the-shelf” list of HRQL scores for the calculation of quality-adjusted life expectancy (QALE) in the United States using mortality data and data from the NHIS.3–5 The National Center for Health Statistics (NCHS) has created a matrix of 30 possible HRQL values by combining five levels of self-reported health and six levels of role function and linking these states to a multiattribute utility model, described in detail by Erickson et al.4 The HRQL values for the different cells have been linked to self-reported conditions.3 Though these data may be limited by the use of only two dimensions of HRQL and self-report bias, it is possible to adjust these parameters for demographic factors and to thereby capture the average HRQL of specific populations.5,6

Given that the NHIS data may be combined with mortality data that are simple to access and use, developing a means of using the YHL measure to calculate both QALYs and QALE may shorten the time to com-
Table 1. QALY and QALE in the reference life table cohort

<table>
<thead>
<tr>
<th>Age interval</th>
<th>Probability of death in interval</th>
<th>Number of person-years beginning interval</th>
<th>Number of person-years dying in interval</th>
<th>Person-years in interval</th>
<th>Cumulative person-years</th>
<th>Life expectancy</th>
<th>HRQL *</th>
<th>QALYs</th>
<th>Cumulative QALYs</th>
<th>QALE</th>
</tr>
</thead>
<tbody>
<tr>
<td>&lt;45</td>
<td>0.050</td>
<td>100,000</td>
<td>5,004</td>
<td>4,405,191</td>
<td>7,601,026</td>
<td>76.0</td>
<td>0.91</td>
<td>4,019,648</td>
<td>6,449,217</td>
<td>64.5</td>
</tr>
<tr>
<td>45–64</td>
<td>0.142</td>
<td>94,996</td>
<td>13,486</td>
<td>1,765,060</td>
<td>3,195,834</td>
<td>33.6</td>
<td>0.82</td>
<td>1,445,507</td>
<td>2,429,570</td>
<td>25.6</td>
</tr>
<tr>
<td>65–74</td>
<td>0.225</td>
<td>81,510</td>
<td>18,348</td>
<td>723,360</td>
<td>1,430,774</td>
<td>17.6</td>
<td>0.75</td>
<td>543,005</td>
<td>984,063</td>
<td>12.1</td>
</tr>
<tr>
<td>75+</td>
<td>1.000</td>
<td>63,162</td>
<td>63,162</td>
<td>707,414</td>
<td>707,414</td>
<td>11.2</td>
<td>0.62</td>
<td>441,058</td>
<td>441,058</td>
<td>7.0</td>
</tr>
</tbody>
</table>

*Average HRQL for persons in age interval.

HRQL, health-related quality of life; QALE, quality-adjusted life expectancy; QALYs, quality-adjusted life years

Methods

Overview

To illustrate the use of the YHL measure for the calculation of QALYs and QALE, we calculated the burden of disease due to stroke. We first calculated quality-adjusted person-years remaining at birth in an abridged life table cohort using the methods forwarded by Erickson et al.4,7,8 (Table 1.) This cohort serves as a reference against which marginal QALYs due to stroke may be calculated. We then generated a second abridged life table in which the cohort is subjected to the risk of dying from all diseases but stroke (a “stroke-free” cohort). Finally, we subtracted the quality-adjusted person-years remaining at birth in the reference life table cohort from the quality-adjusted person-years remaining at birth in the stroke-free cohort. All calculations were conducted using abridged life tables on Excel 98 for the Macintosh (Microsoft Inc., Redmond, WA).

Reference Cohort

An abridged life table is constructed using a hypothetical population in which 100,000 people are born each year. It is assumed that all people in this cohort are at risk of dying or becoming ill, but that there is no migration into or out of the population. Because 100,000 people are born into the cohort each year, the number of persons in the cohort is equal to the number of person-years in the cohort.7,8 When person-years are adjusted for HRQL, they become QALYs. Table 1 illustrates how a life table is constructed using the YHL measure to determine the number of QALYs in a cohort.

Use of Published Data

With reference to Table 1, Column 1 represents the probability of death in a given interval; Column 2, the number of people alive at the beginning of the age interval; and Column 3, the number of persons dying during the interval. Using data from an abridged life table, Column-3 values for each interval are entered first.7 Column-2 values are calculated as s_x=s_x−1−d_x−1, where s_x is the number of survivors at the beginning of age interval x, d_x is the number of persons dying in the interval, and x−1 denotes the previous interval. Because all life tables begin with 100,000 persons alive at the beginning of the first age interval, the first value of Column 2 is 100,000. Dividing the total number of deaths in the interval (Column 3) by the number of persons alive at the beginning of the interval (Column 2), yields the probability of death during the interval (Column 1).

Column 4 row values represent the total number of person-years in the interval and are calculated using the formula n(s_x−0.5d_x), where n is the length of the interval. This formula assumes that all deaths occurred at the midpoint of the interval—an assumption that does not hold for the first year of life (because of infant mortality) or the last age interval (because the interval is of indeterminate length).

The number of person-years in the first year of life is calculated as 100,000f+(1−f)s_1, where f is the separation factor7 and s_1 is equal to the number of persons surviving to age 1. The separation factor is equal to the proportion of infant deaths in the base year occurring to infants born the previous year and is available from the NCHS8 and s_1 may be obtained directly from an existing life table.7,8 The number of person-years in the first age interval in Table 1 is thus (n−1)(s_x−0.5d_x)+s_xf+(1−f)s_1, where d_x is the number of deaths occurring in persons aged 1 to 44. The probability of death for the ≥75 age interval is equal to the reciprocal of the life expectancy at age 75.

To obtain cumulative person-years (Column 5) the values for Column 4 are summed backwards such that the value for the first age interval is equal to the sum of all values in Column 4. Each row in Column 6 is calculated by dividing the row value of Column 5 by the corresponding row value for Column 2. The value of the first row in Column 6 is equal to the life expectancy at birth of the cohort and serves to validate the spreadsheet (this should be similar to the life expectancy at birth in abridged life tables). The values in Column 7 represent the HRQL score for the average person in the United States in each corresponding age interval and are available from the NCHS.4 Column 8 is equal to the product of Column 7 and Column 4. Columns 9 and 10 are derived in a manner analogous to Columns 4 and 5.

Since the age intervals must be the same across prevalence, mortality, and HRQL data, it is best to generate age intervals...
Table 2. QALY and QALE in an abridged life table cohort for persons without stroke

<table>
<thead>
<tr>
<th>Age interval</th>
<th>Probability of death in interval</th>
<th>Number dying in interval</th>
<th>Person-years in interval</th>
<th>Life expectancy</th>
<th>Prevalence</th>
<th>YHL score</th>
<th>HRQL</th>
<th>QALYs</th>
<th>Cumulative QALYs</th>
<th>QALE</th>
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<tbody>
<tr>
<td>&lt;45</td>
<td>0.050</td>
<td>5,002</td>
<td>4,405,283</td>
<td>76.0</td>
<td>0.0017</td>
<td>0.62</td>
<td>0.913</td>
<td>4,024,338</td>
<td>6,510,545</td>
<td>65.11</td>
</tr>
<tr>
<td>45–64</td>
<td>0.142</td>
<td>13,460</td>
<td>1,765,369</td>
<td>33.7</td>
<td>0.0149</td>
<td>0.47</td>
<td>0.825</td>
<td>1,458,123</td>
<td>2,486,208</td>
<td>26.17</td>
</tr>
<tr>
<td>65–74</td>
<td>0.224</td>
<td>18,238</td>
<td>724,195</td>
<td>17.6</td>
<td>0.0519</td>
<td>0.45</td>
<td>0.772</td>
<td>559,793</td>
<td>1,028,085</td>
<td>12.61</td>
</tr>
<tr>
<td>75+</td>
<td>1.000</td>
<td>63,300</td>
<td>708,964</td>
<td>11.2</td>
<td>0.988</td>
<td>0.38</td>
<td>0.660</td>
<td>468,291</td>
<td>468,291</td>
<td>7.40</td>
</tr>
</tbody>
</table>

*HRQL score associated with the YHL measure.

HRQL, health-related quality of life; QALE, quality-adjusted life expectancy; QALYs, quality-adjusted life years; YHL, years of healthy life

Use of Electronic Data

Although the use of electronic data substantially increases the cost and time requirements of cost-effectiveness or burden-of-disease studies, it is possible to use much smaller age intervals, and thus improve the accuracy of model outputs. To generate reliable prevalence and HRQL estimates using electronic data, it is necessary to aggregate 3 to 4 years of NHIS data.

The first row value for Column 1 (q0) is approximated using the infant mortality rate, which is equal to the number of deaths in the first year of life divided by the number of live births. Because some deaths occur in infants born the previous year and the number of live births varies slightly from year to year, a more accurate estimate may be obtained using the separation factor, f. The more accurate estimator of q0 is D0/(1-f)/B0+D0/Bx-1, where D0 is the total number of infant deaths and B0 is the total number of births in year x.

The remaining values for Column 1 are generated using the formula qx=Dx/(Px+0.5Dx) where qx is the probability of death during the interval, Dx is the total number of deaths observed at age x, and Px is the midyear population for persons aged x. The survivors at age x (sx) may then be calculated as sx=(1-qx). Column 3 values are simply the product of Columns 1 and 2 values for the corresponding rows.

All other values are calculated in an identical manner to those derived from a published life table, with the exception of total person-years in the final age interval. This is equal to the number living at the beginning of the interval divided by the probability of death during the interval (sx/qx).

Stroke-Free Cohort

Table 2 represents an abridged life table for all persons except those with cerebrovascular disease listed as an underlying cause of death (International Classification for Disease, 9th Revision codes 430 to 438). Table 2 is calculated in the same manner as Table 1 with a few exceptions. In Column 1, the probability of death in the interval due to the disease under study (in this case, stroke) is subtracted from the all-cause mortality rate.

Column 5 contains age-specific prevalence rates for the disease under study. These are multiplied by (1–HRQL) for the disease in question (in this case, the age-specific YHL score for stroke) in Column 6. The product of Columns 5 and 6 for each age interval is then added to the average HRQL (Column 7). This is mathematically equivalent to averaging out the HRQL lost to stroke in the cohort. The remainder of Table 2 is calculated in a manner analogous to Table 1.

Finally, we calculated the burden of disease due to stroke by subtracting QALYs remaining in the cohort with all diseases (see Table 1, Column 9, Row 1) from QALYs remaining in the cohort with all diseases except stroke (see Table 2, Column 9, Row 1). To obtain the QALE lost to stroke, the QALE in the reference cohort (see Table 1, Column 10, Row 1) is subtracted from the QALE in the stroke-free cohort (see Table 2, Column 10, Row 1). We validated the spreadsheet model by comparing our QALE value with published reports.

Sensitivity Analysis

Of the two data sets used to calculate QALYs, only the NHIS data are subject to random error. Since the NHIS is used to generate both HRQL scores and disease prevalence rates, it is possible to calculate the standard error for these parameters. A discussion of random and nonrandom errors may obtained from the NCDS.9 Although it is possible to estimate the standard error of HRQL scores obtained from the YHL measure, the standard error provides no information about the degree to which the scores represent the preferences of particular subgroups for the conditions specified. We have noted previously that the scores are developed from a two-domain model, which necessarily represents a relatively coarse description of a health state. In addition, scores are calculated rather than measured. Finally, the structure of the NHIS does not allow comorbidities to be broken out separately. For all of these reasons, investigators may wish to conduct sensitivity analyses for the HRQL values by using 25th and 75th percentile values for the conditions studied. These values have been previously published and are available in tabular form. The extent to which death certificate data are subject to misclassification bias or other types of nonrandom error may be estimated from the...
medical literature\textsuperscript{10–12} and sensitivity analyses conducted, as appropriate.

**Results**

Life expectancy, QALE, and QALYs remaining at birth for the general U.S. population cohort and the stroke-free cohort are presented in Table 3. The burden of disease due to stroke was approximately 61,328 QALYs. Assuming that a negligible number of deaths due to stroke would be attributed to other causes and that the years of healthy life measure produces HRQL scores that are representative of the condition-specific values of the general U.S. population, the 95\% confidence interval for the burden of disease due to stroke in our life table cohort would be approximately 60,272 QALYs to 62,383 age-adjusted QALYs. If stroke were eliminated, persons in the United States would gain approximately 0.61 years of QALE but only approximately 0.03 additional years of life.

The reference cohort produced a QALE at birth of 64.5 years—0.5 years longer than the 1990 NCHS estimate of QALE, 64 years.\textsuperscript{4} Our estimate of life expectancy in 1997, 76 years, differed from the NCHS estimate, 76.5 years, by approximately seven tenths a percent—a variation that is likely attributable to the large age intervals used in this study.

**Conclusions**

In this paper, we illustrated a simple technique for calculating QALYs using data accessible by the Internet. To illustrate this method, we calculated QALYs lost to stroke in our cohort, an estimate affected by random error. The model predicted the loss of approximately 61,328 QALYs to stroke in the hypothetical life-table cohort. On average, persons residing in the United States could expect to live approximately 0.61 additional years of quality-adjusted life if strokes were eliminated, but only approximately 0.03 additional years of life when quality adjustments are not included. The difference between absolute and quality-adjusted years gained underscores the importance of including a valuation of quality years lost to morbidity. This method not only allows for rapid tabulation of the burden of disease due to most illnesses, but it also provides a convenient way of calculating the number of QALYs gained due to an intervention in cost-effectiveness analyses.

For researchers wishing to conduct a cost-effectiveness analysis, the differential in QALE for a group with and without a condition will usually provide sufficient data for the construction of a decision analysis model.\textsuperscript{1} Moreover, most interventions will not require de novo generation of HRQL scores using data from the NHIS since these data are readily available in the medical literature.\textsuperscript{4,5}

Calculations for the YHL measure are based on the technique of correspondence analysis\textsuperscript{13} which is linked to values drawn from the Health Utilities Index (HUI),\textsuperscript{14} a relatively complex, multi-attribute, health status instrument. Multi-attribute models typically utilize multiple health domains. Given its two-domain structure (role function and self-reported health), HRQL scores derived from the YHL measure may be less precise than scores generated from measures that have a more complex formulation of health such as the HUI, the Quality of Well-Being Index,\textsuperscript{15} and the EuroQol.\textsuperscript{16} On the other hand, the measure’s ability to include the average HRQL of a population and to generate HRQL scores specific to a demographically defined group may provide advantages over other HRQL indexes in particular settings. Ideally, future population surveys would contain more complex preference-based instruments that could be coupled to specific disease prevalence data.

The NHIS is an annual survey conducted by the NCHS.\textsuperscript{17} Data are collected in face-to-face interviews in a nationally representative sample of households. When conducting our sensitivity analysis, we only included sources of random error in estimating the 95\% confidence interval for the burden of disease due to stroke. There is debate surrounding the sensitivity and specificity of self-report data which are subject to recall bias.\textsuperscript{18,19} A subject’s inability to accurately remember the illnesses from which he or she suffers may affect both the tabulation of prevalence rates and the HRQL score. Moreover, the use of death certificates to ascertain mortality rates is subject to misclassification bias.\textsuperscript{10–12} Death certificates may fail to correctly classify

<table>
<thead>
<tr>
<th></th>
<th>U.S. population cohort</th>
<th>Stroke-free cohort</th>
<th>Difference</th>
<th>95% CI**</th>
</tr>
</thead>
<tbody>
<tr>
<td>Life expectancy</td>
<td>76.01</td>
<td>76.04</td>
<td>0.03</td>
<td>–</td>
</tr>
<tr>
<td>QALE</td>
<td>64.49</td>
<td>65.11</td>
<td>0.61*</td>
<td>0.60–0.61</td>
</tr>
<tr>
<td>QALY</td>
<td>6,449,217</td>
<td>6,510,545</td>
<td>61,328</td>
<td>60,272–62,383</td>
</tr>
</tbody>
</table>

*Difference is not exact due to rounding.

**Based on sampling error in prevalence estimate alone.

QALE, quality-adjusted life expectancy; QALY, quality-adjusted life years.

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Table 3. Life expectancy, QALE, and QALY remaining at birth for the general U.S. population cohort and stroke-free cohort.
the primary or underlying cause of death or the demographic characteristics of the decedents.

Because the sample size of the NHIS is small (approximately 103,000 people in 1997), relative to the total number of possible diseases subjects might be afflicted with, the researcher often faces a trade-off between unreasonably large age intervals or too few subjects to generate reliable estimates of prevalence or HRQL. Although African Americans and Hispanics are over-sampled in the NHIS, researchers interested in obtaining prevalence rates for other races/ethnicities or for specific geographic regions will not likely be able to generate reliable estimates using a single year of NHIS data. Sample size issues may be overcome by aggregating NHIS data over a number of years. However, year-to-year changes in disease prevalence may limit this method. Finally, the age intervals we used were broad. When infant mortality is accounted for separately, however, the use of the broad age intervals we present here only produces a slight underestimate of life expectancy (data not shown).

The Panel on Cost-Effectiveness in Health and Medicine recommends that cost-effectiveness analyses designed to inform resource allocation decisions should present outcomes in the form of QALYs. The method we present for calculating QALYs is simple enough to permit their use in resource-limited research efforts. Using the techniques discussed in this paper, it is possible for public health researchers to generate burden-of-disease estimates specific to demographically defined communities within the span of a few hours. This method may, therefore, also have great utility in assisting public health agencies in making rapid decisions surrounding the allocation of resources to local health priorities.

We are indebted to Marianne Fahs, Clyde Schecter, Peter Franks, and Karla Hansen for their assistance with verifying the statistical accuracy of the calculations in this paper.

References