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# Age at Diagnosis and the Utility of Prognostic Testing in Patients with Chronic Lymphocytic Leukemia (CLL)

Tait D. Shanafelt, M.D., Kari G. Rabe, MS, Neil E. Kay, M.D., Clive S. Zent, M.D., Diane F. Jelinek, Ph.D., Megan S. Reinalda, Susan M. Schwager, Debbie A. Bowen, F.NP, Susan L. Slager, Ph.D., Curtis A. Hanson, M.D., and Timothy G. Call, M.D. Mayo Clinic, Rochester, MN

## Abstract

**PURPOSE**—To analyze the survival of CLL patients relative to age-matched individuals in the general population and determined the age-stratified utility of prognostic testing.

**METHODS**—All 2487 patients diagnosed with CLL between January 1995 and June 2008 and cared for in the Mayo Division of Hematology were categorized by age at diagnosis and evaluated for differences in clinical characteristics, time to first treatment(TFT), and overall survival(OS).

**RESULTS**—Among Rai stage 0 patients, survival was shorter than the age-matched general population for patients age<55 years(p<0.001), 55-64 years(p<0.001), and 65-74 years(p<0.001) but not those age 75 at diagnosis(p=NS). CD38, IGHV mutation, and ZAP-70 each predicted TFT independent of stage for all age groups(all p <0.04) but had less value for predicting OS, particularly as age increased. IGHV and FISH predicted OS independent of stage for patients <a href="#reliable"></a> (p <0.001), 55-64(p 0.004), and 65-74(p 0.001) but not those 75. CD38 and ZAP-70 each predicted OS independent of stage for only 2 of 4 age categories. Among Rai 0 patients age<75, survival was shorter than the age-matched population only for IGHV unmutated(p<0.001) patients or those with unfavorable FISH(p<0.001).

**CONCLUSIONS**—Survival of CLL patients age<75 is shorter than the age-matched general population regardless of disease stage. Among patients age<75, the simple combinations of stage and IGHV or stage and FISH identifies those with excess risk of death relative to the age-matched population. Although useful for predicting TFT independent of stage for patients of all ages, prognostic testing had little utility for predicting OS independent of stage among patients age 75.

## Background

Chronic lymphocytic leukemia/small lymphocytic lymphoma (CLL) is one of the most common lymphoid malignancies accounting for approximately 11% of hematologic cancers

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Corresponding Author: Tait D. Shanafelt, M.D., Mayo Clinic, 200 First Street, Rochester, MN 55902, Phone: (507) 266-1154, FAX: (507) 266-4972, shanafelt.tait@mayo.edu.

Author Contributions

Concept: Tait Shanafelt, Neil Kay, Tim Call, Clive Zent

Designed research: Tait Shanafelt, Kari Rabe, Neil Kay, Tim Call, Clive Zent, Diane Jelinek

Performed research: Tait Shanafelt, Diane Jelinek, Susan Schwager, Curt Hanson

Analyzed and interpreted data: Kari Rabe, Megan Reinalda, Susan Slager, Tait Shanafelt, Neil Kay, Clive Zent, Diane Jelinek, Debbie Bowen, Curt Hanson, Tim Call

Critical review and revision of paper: Neil Kay, Kari Rabe, Tim Call, Clive Zent, Diane Jelinek, Debbie Bowen, Susan Slager, Curt Hanson

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in the Western World.<sup>1</sup> The prevalence of CLL increases with age and the median age at the time of diagnosis is between 65 and 70 years.<sup>2-6</sup> Recent studies suggest that the 5 year survival of CLL patients of all ages has increased over the last two decades,<sup>6-8</sup> likely due in part to early stage at diagnosis.<sup>3,4,6,9</sup> The absolute 10 year survival of patients with CLL has increased by ~10% for patients of all ages except those over age 80 years<sup>7,8</sup>.

While the observed improvement in the survival of CLL patients at the population level is encouraging, the clinical course of individual patients is heterogeneous. Even among individuals with early stage disease, there remains significant heterogeneity in clinical behavior and stage alone does not adequately predict the risk of progression for a given patient.<sup>10</sup> Although numerous clinical and biologic parameters are able to predict survival and time to first treatment (TFT),<sup>10-15</sup> the utility of these prognostic parameters may vary based on age given the higher mortality from competing health problems in older individuals.<sup>7,16,17</sup> Indeed there remains a strong age gradient in the survival among CLL patients<sup>2,7,8,18</sup> where the expected 10 year survival for those less than age 60 is 59% compared to 6% for those over age 80.<sup>7</sup> Accordingly, while risk stratification using leukemia cell biomarkers (e.g. ZAP-70, FISH, IGHV testing) may provide useful information for counseling a newly diagnosed 50 year old patient with Rai stage 0 disease, its usefulness to a 75 year old patient in the same clinical circumstance is less clear<sup>13,16</sup>.

These facts have important implications for use of prognostic testing and counseling regarding life expectancy for older individuals with CLL who represent the majority of CLL patients world-wide. Most of the data on the ability of prognostic parameters to predict outcome is derived from cohorts of CLL patients with a median age <65 years<sup>10-15</sup> and the median age of patients in many series is <60 years<sup>12,14,15</sup>. In the present study, we evaluated the clinical outcome of 2487 patients diagnosed with CLL between January 1995 and June 2008 to: i) evaluate differences in natural history based on age at diagnosis, ii) compare survival to age-matched individuals in the general population, and iii) determine the age-stratified utility of prognostic testing.

## Methods

#### Patients

The Mayo Clinic CLL Database includes all patients with a diagnosis of CLL<sup>1932</sup> seen in the Division of Hematology at Mayo Clinic Rochester (MCR) who permit their records to be used for research purposes.<sup>20-26</sup> Clinical information regarding date of diagnosis, physical examination, clinical stage (Rai), prognostic parameters, treatment history, and disease-related complications are abstracted from clinical records on all patients at the time of inclusion and maintained on an ongoing, prospective basis. For staging purposes, patients with SLL who have cytopenias at diagnosis are grouped with Rai stage III/IV patients while those with palpable lymphadenopathy without cytopenias are grouped with Rai stage I patients. Results of prognostic testing performed as part of clinical or research studies are also included in the database. This includes evaluation of absolute lymphocyte count (ALC), IGHV gene mutation analysis, ZAP-70 status, CD38 status, and cytogenetics abnormalities by interphase FISH testing using methods previously described by our group<sup>20,27-29</sup>.

With the approval of the Mayo Clinic Institutional Review Board and in accord with federal regulations and the Declaration of Helsinki, we used this database to identify all patients diagnosed with CLL between January 1995 and June 2008. All these patients had an ALC  $5.0 \times 10^{9}$ /L and fulfilled the 1996 criteria for CLL which were in effect throughout the study period<sup>32</sup> and/or fulfilled the WHO criteria for CLL/SLL<sup>19</sup>. Patients were categorized by age at the time of CLL diagnosis (<55 years, 55-64 years, 65-74 years, 75 years) with categories based in part on the previous designations of "young" CLL as individuals age 55

at diagnosis<sup>16,33</sup> and stratification of patients >age 55 in 10 year intervals up to age 75. Differences in clinical characteristics, time to first treatment (TFT), and overall survival (OS) based on age at diagnosis were assessed. Since FISH can change during the course of the disease or after treatment,<sup>29</sup> only FISH analysis obtained prior to first treatment was included in the present analysis. Based on evidence that VH 3-21 family usage is associated with poor outcome independent of mutation status <sup>30,31</sup>, patients with VH 3-21 family usage were considered to have high risk IGHV status regardless of percent mutation.

#### **Statistical methods**

OS was defined as the time between the date of diagnosis to the date of death or last followup. TFT was defined as the time between date of diagnosis and the date of initiation of first treatment or date of last follow-up at which the patient was known to be untreated. The accepted indications to initiate treatment were based on the NCI-WG 1996 criteria.<sup>32</sup> Patients receiving early treatment as part of experimental protocols prior to meeting NCI-WG 1996 criteria to initiate therapy were censored as untreated on the date experimental therapy was administered. Estimates of survival were calculated using the Kaplan-Meier method. Cox proportional hazard models were used to model the relationship of multiple variables simultaneously including age at diagnosis with OS and TFT. Expected survival was calculated using the Cohort (Hakulinen) method<sup>34</sup>; estimates are based on the "Minnesota White" population<sup>35</sup>. Likelihood ratio tests were used to test effects of individual factors either individually or jointly. Hazard ratios (HR) and 95% confidence intervals (CI) were calculated from the Cox models. In addition to individual cytogenetic categories, FISH results were classified as 'unfavorable' (17p-, 11q-) or 'favorable' (normal, trisomy 12, 13q-, other) for some analyses. P-values <0.05 were considered significant. All statistical analyses were performed using the SAS 9.1 software package (SAS Institute; Cary, North Carolina).

### Results

There were 2487 patients who qualified for inclusion in this study. The median age at diagnosis was 64 years old. When grouped by age, 593 patients were age <55, 713 age 55-64, 748 age 65-74, and 433 age 75 at diagnosis. The demographic and prognostic characteristics of patients by age at diagnosis are shown in Table 1. Because CD38, ZAP-70, IGHV gene mutation status, and FISH analysis were not routinely performed during the entirety of the study period, results were not available for all patients. Patients under the age of 55 were more likely to have intermediate stage (Rai I or II) at diagnosis while those >age 65 were less likely to have an ALC>30 ×10<sup>9</sup>/L; however no statistically significant differences were observed in CD38, IGHV mutation, ZAP-70 or the frequency of common cytogenetic abnormalities as identified by FISH based on age. TFT among Rai stage 0 patients was shorter for patients <55, however no difference in TFT by age was observed among Rai stage I or II patients.

Median follow-up was 9.7 years. As of last follow-up, 954 patients have been treated and 727 patients have died. Median TFT was 4.8 years. Although survival decreased as the age at diagnosis increased (Figure 1A), the survival of CLL patients was significantly shorter than that of the age-matched general population for patients age <55 years (p<0.001), 55-64 years (p<0.001), and 65-74 years (p<0.001) at diagnosis but not those 75 years (p=0.14; Figure 1B-E). Among Rai stage 0 patients, survival was also shorter than that of the age-matched general population for CLL patients age <55 years (p<0.001), 55-64 years (p<0.001), and 65-74 years (p<0.001) but not those age 75 (p=0.07) at diagnosis (Figure 2A-D).

We next evaluated the relationship between prognostic parameters and TFT and OS. Consistent with prior reports, stage, ALC ( $or >30 \times 10^9/L$ ), CD38, IGHV mutation, ZAP-70 and cytogenetic analysis by FISH were all powerful predictors of both TFT and OS on univariate analysis (all p 0.003 for both TFT and OS; Table 2). In a multi-variate analysis in the 585 patients who had results for all prognostic variables, stage (HR high Rai risk vs. low Rai risk=16.1; HR intermediate Rai risk vs. low Rai risk=2.3), IGHV unmutated (HR=2.8; p<0.001), ALC >30 × 10<sup>9</sup>/L (HR=2.0), and CD38 positive (HR=1.9) remained independent predictors of TFT (all p<0.001) while ZAP-70 (HR=1.3; p=0.16) and FISH (HR=1.3; p=0.264) were no longer statistically significant. With respect to OS, only high risk FISH (HR=2.9; p=0.008) and CD38 positive (HR=2.2; p=0.041) remained independent predictors of OS where IGHV (HR 2.8; p=0.072), stage (HR high Rai risk=2.8, p=0.20; HR intermediate Rai risk=1.4, p=0.37), ALC (HR 1.2; p=0.62), and ZAP-70 (HR 0.9; p=0.77) were no longer statistically significant.

Given the variation in TFT and OS based on age at diagnosis as well as differences in the magnitude of effect of a CLL diagnosis on survival relative to age-matched controls, we next evaluated whether ALC, CD38, IGHV mutation, ZAP-70 and FISH remained useful predictors of TFT and OS for CLL patients in all age categories. Stage, ALC, CD38, ZAP-70, and IGHV remained powerful predictors of TFT for CLL patients of all ages including those age 75 (all p 0.005). FISH was also a powerful predictor of TFT for CLL patients of all ages except those over age 75 where it failed to reach statistical significance (p=0.09). The hazard ratio of the individual prognostic parameter for predicting TFT was generally similar across age groups.

In contrast to the near uniform value of ALC, CD38, IGHV mutation, ZAP-70 and FISH for predicting TFT in all age categories, their ability to predict OS was less consistent. Stage, CD38, and IGHV were statistically significant predictors of OS in CLL patients of all age categories including those – age 75 (all p<0.05). FISH predicted OS in all age categories <a href="mailto:</a> <a href="mailto:seage">seage 75</a> (all p<0.05). FISH predicted OS in all age categories <a href="mailto:</a> <a href="mailto:seage">seage 75</a> (all p<0.03) but not patients age 75 (p=0.34). ZAP-70 predicted OS for patients who were <55 (p=0.007) and 55-64 (p=0.004) but not patients\_age 65-74 (p=0.28) or >75 (p=0.97). ALC>30 ×10<sup>9</sup>/L was only a significant predictor of OS for patients in the 65-74 year old group.

Next, we evaluated the ability of ALC, CD38, IGHV mutation, ZAP-70 and FISH to predict TFS and OS in each age category after adjusting for stage. With respect to TFT, CD38, IGHV mutation, and ZAP-70, each predicted TFT independent of stage for all age categories (Top Table 3). FISH predicted TFT independent of stage for patients age<55, 55-64 and 65-74 but not for those age 75 (p=0.08). The utility of these parameters for predicting OS independent of stage for patients <a href="https://www.age">age</a> 55, 55-64 and 65-74 but not for those age 75 (p=0.08). The utility of these parameters for predicting OS independent of stage for patients <a href="https://www.age">age</a> 55, 55-64 and 65-74 but not for those age 75 (p=0.08). The utility of these parameters for predicting OS independent of stage for patients <a href="https://www.age">age</a> 55, 55-64, and 65-74 but not for those age 75 (p=0.08). The utility of these parameters for predicting OS independent of stage for patients <a href="https://www.age">age</a> 55, 55-64, and 65-74 but not for those age 75 (p=0.08). The utility of these parameters for predicted OS independent of stage for patients <a href="https://www.age">age</a> 55, 55-64, and 65-74 but not those 75 (Bottom Table 3). CD38 and ZAP-70 each predicted survival independent of stage for only 2 of the 4 age categories (ZAP-70 for age <55 and 55-64; CD38 for 55-64 and 75).

Finally, since OS was only shorter than the age matched population for CLL patients <age 75, we evaluated whether prognostic testing could identify which Rai 0 CLL patients <age 75 had a survival shorter than the age matched population. IGHV and FISH testing were used for this analysis based on the ability of these tests to identify CLL patients with shorter OS independent of stage in all age categories <75. The survival of Rai 0 IGHV unmutated CLL patients <age 75 was shorter than that of the age-matched general population (p<0.001) while Rai 0 IGHV M CLL patients had a survival similar to the population (Figure 3A and B). Similarly, the survival of Rai 0 CLL patients <age 75 with unfavorable FISH was shorter than that of the age-matched general population (p<0.001) while Rai 0 CLL patients with

favorable FISH had a survival similar to the age-matched general population (Figure 3C and D).

## Discussion

Age has repeatedly been shown to be an independent predictor of survival in CLL patients<sup>16,17,36-38</sup> and has the potential to alter the utility of prognostic testing given the higher mortality from competing health problems in older individuals. While the majority of patients with CLL are over age 65 at the time of diagnosis and have early stage disease,<sup>3,4,6,9</sup> most of the published data on prognostic parameters is derived from younger patient cohorts. This incongruity has led to uncertainty regarding if and when to use prognostic testing in routine clinical practice for patients with CLL. In the present study, the survival of CLL patients < age 75 at diagnosis was shorter than that of the age-matched general population among CLL patients age 75 at diagnosis. Prognostic testing USA, IGHV mutation, and ZAP-70 was useful for predicting TFT independent of stage for CLL patients of all ages (including those age 75) but had less value for predicting OS, particularly as the age at diagnosis increased. Among Rai 0 patients <a href="https://was.shorter.survival-shor

These findings have a number of important implications for the use of prognostic tests in patients with CLL as well as the use of test results to select patients for clinical trials testing the value of early treatment. First, although the life expectancy of CLL patients <age 75 is substantially shorter than that of the age matched general population, the simple combination of stage and IGHV mutation status or stage and FISH can identify those with excess risk of death. Second, with regard to TFT, CD38, ZAP-70, IGHV, and FISH each provide useful information for CLL patients of all ages *independent* of stage. Third, prognostic markers appear to be a sound basis upon which to select "high risk" early stage patients <age 75 for clinical trials of early intervention. In the present cohort, both IGHV mutation status and FISH predicted OS independent of stage with relatively large hazard ratios (range 2.8-6.2) among patients in all age categories < age 75. Fourth, while useful for predicting OS independent of stage among patients in this age category. Accordingly, it does not appear appropriate to enroll patients >age 75 in clinical trials of early intervention based on prognostic testing if the aim is to improve OS.

Several other aspects of this analysis are noteworthy. The study included comprehensive multi-variate analysis of both traditional (age, stage, ALC) and biologic (ZAP-70, CD38, FISH, IGHV) parameters in a large cohort of CLL patients. Consistent with prior reports demonstrating that the biologic parameters contain complementary prognostic information,<sup>14,39,40</sup> multiple biologic prognostic parameters were independent predictors of TFT and/or OS in multivariate analyses including traditional parameters (stage, ALC). It is also notable that the distribution of biologic prognostic parameter results did not differ by age at diagnosis arguing against the notion that CLL in younger patients is more biologically aggressive.

How do these results relate to previous studies? Although age has repeatedly been shown to be an independent predictor of survival in CLL patients,<sup>7,8,16,17</sup> few prior studies have evaluated interactions between age and the utility of prognostic testing. Mauro and colleagues previously demonstrated that lymphocyte doubling time is a predictor of OS among CLL patients both age 55 and over age 55 but provided no further age stratification of those over 55.<sup>33</sup> Dohner and colleagues observed that the presence of del(11q22) was a

profound stratifier of survival in patients <age 55 (median survival del(11q22)=64 months vs. no del(11q22)=209 months; p<0.001) but not patients > age 55 (median survival del(11q22)=94 months vs. no del(11q22)=111 months; p=0.82).<sup>16</sup> Due to the limited data demonstrating utility in older patients, many hematologist/oncologists do not routinely use prognostic testing for older individuals with early stage CLL which limits the accuracy of counseling on natural history and life expectancy for these patients. The present study provides more comprehensive data regarding the utility of prognostic testing for classifying risk among CLL patients over age 65. To our knowledge, it also is the first study to evaluate how prognostic test results can be used to stratify the risk of death among patients with CLL relative to the age-matched general population.

Our study has several important limitations. Given that the natural history of CLL is changing,<sup>7,8</sup> the fact that all patients in the current cohort were diagnosed in the last 15 years is a strength of the study. However, not all patients had all the molecular/biologic prognostic parameters measured since they were only discovered/used routinely in the last 5-10 years.<sup>10-14</sup> Second, the diagnosis of CLL was based on the 1996 criteria for CLL<sup>32</sup> which were in effect throughout the study interval but recently underwent revision<sup>41</sup>. Third, as in other analyses of clinical outcome,<sup>36,42-44</sup> we evaluated overall survival rather than disease-specific survival. Because overall survival is the outcome of greatest interest to patients, we believe that this is the most appropriate outcome for survival analysis with use of TFT as a secondary measure of disease specific outcomes. Fourth, our study focused on the ability of prognostic tests to predict TFT and OS. The utility of these assays for predicting other outcomes, such as response to treatment, was not the focus of the current study. Finally, the study represents a single center experience that requires validation in independent series of patients monitored prospectively.

In aggregate, these findings suggest that survival of CLL patients <age 75 at diagnosis is shorter than that of the age-matched general population regardless of disease stage. Prognostic testing had little utility for predicting OS independent of stage among patients age 75, although it remained useful for predicting TFT. In settings where the goal is to identify patients with excess risk of death relative to the age matched population, it appears that clinical staging is the only test necessary for patients age 75 at diagnosis while limited testing with a combination of stage and IGHV or stage and FISH are appropriate strategies for most patients <a href="#age75"></a>.

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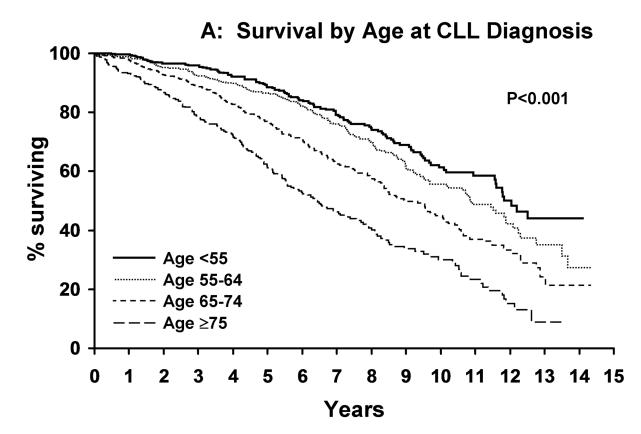
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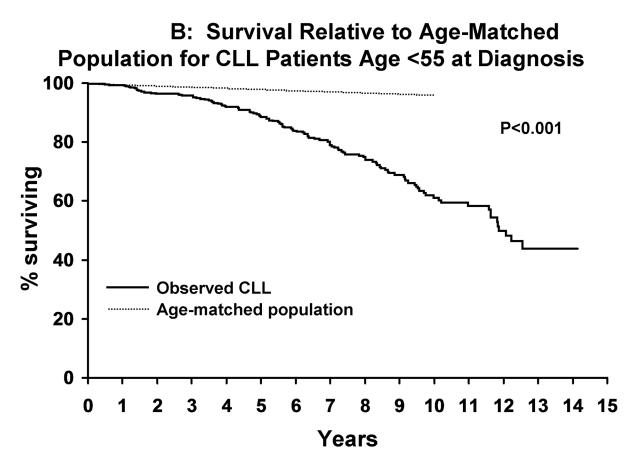
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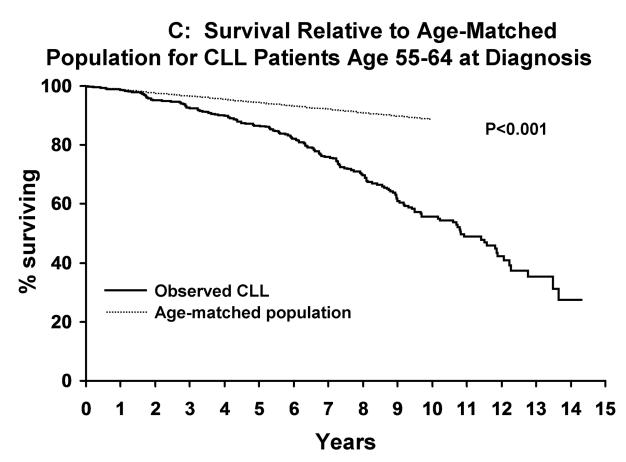
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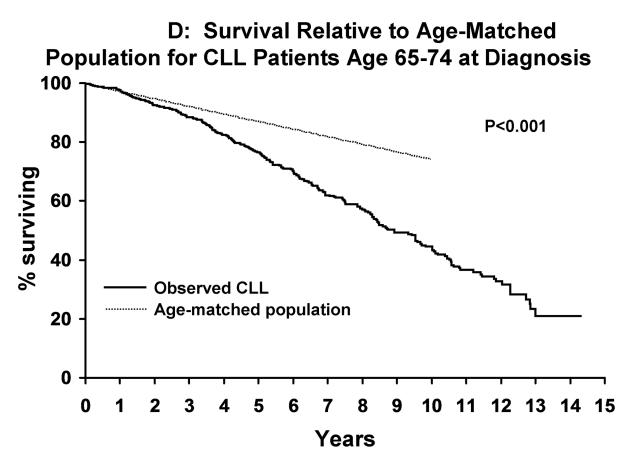
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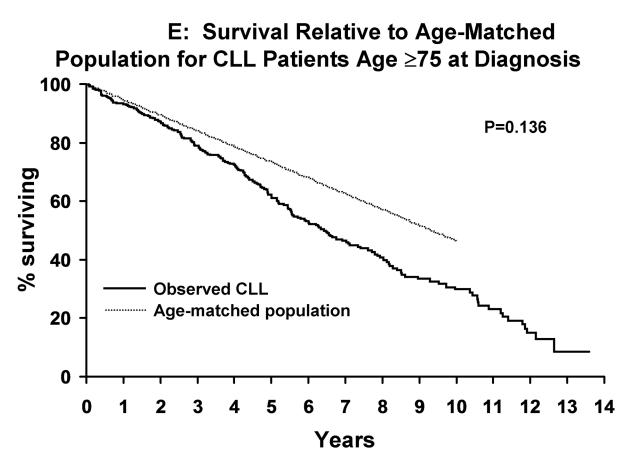


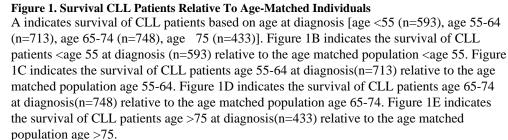
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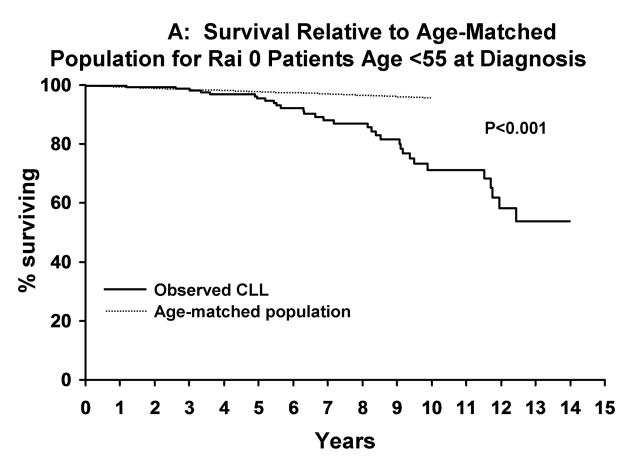


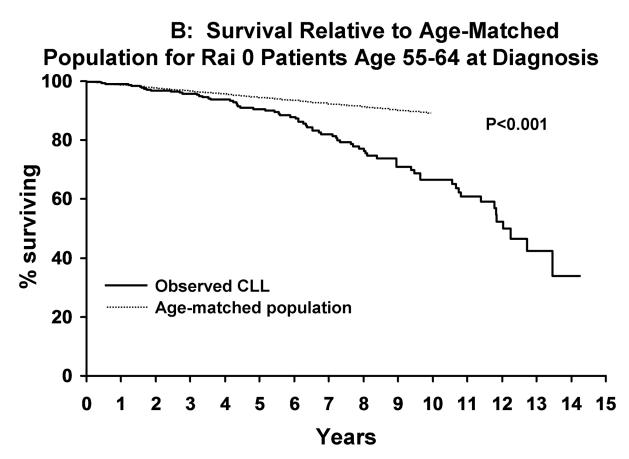


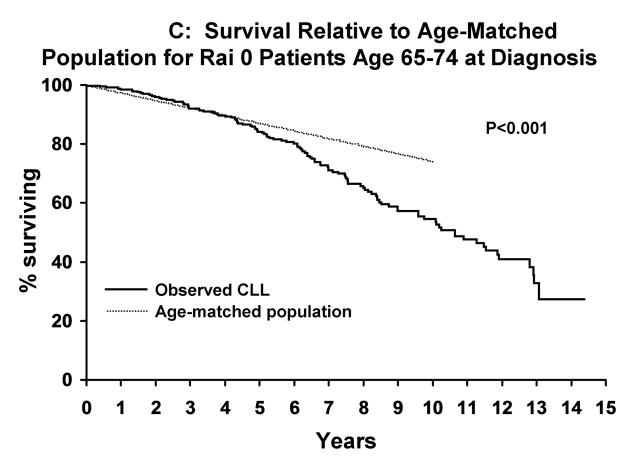


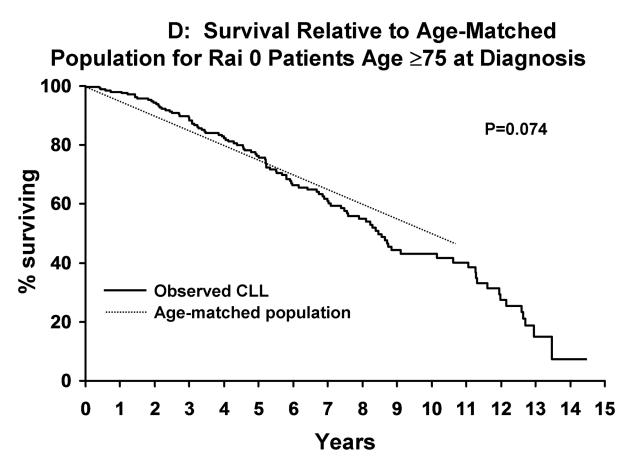


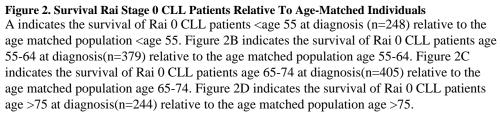
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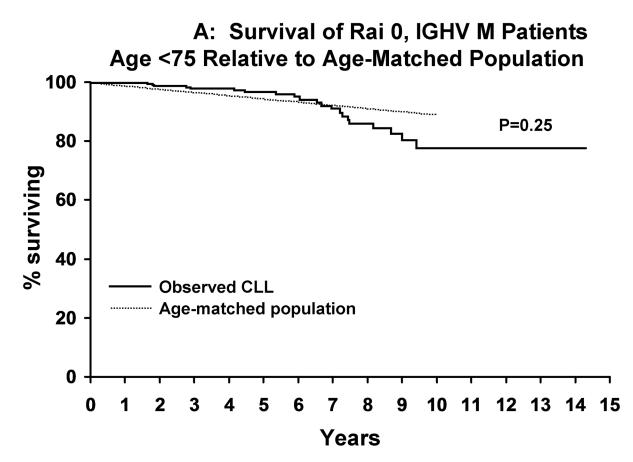


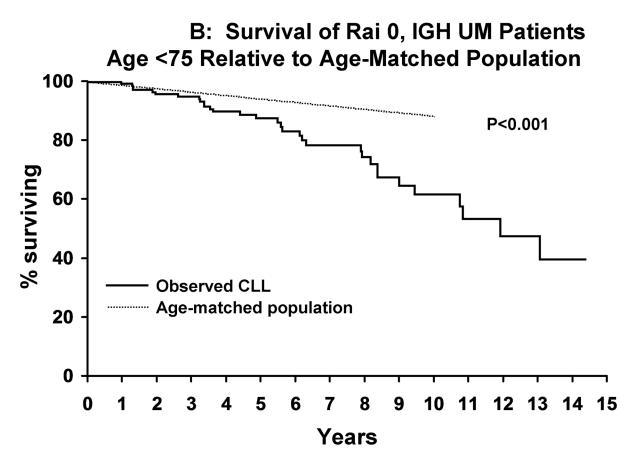




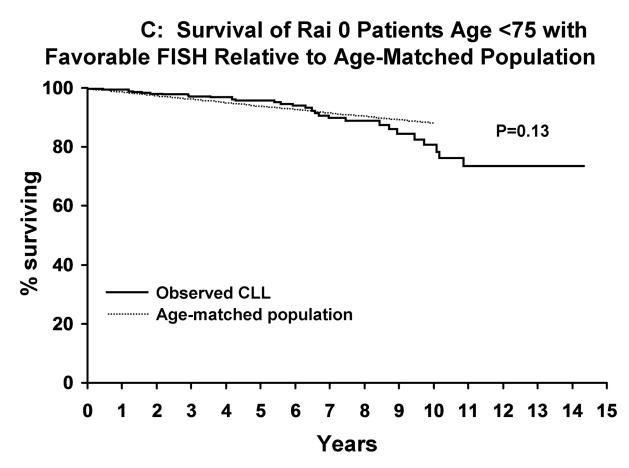


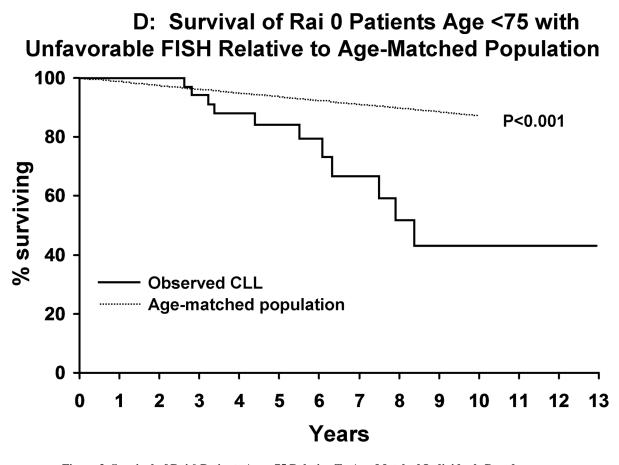


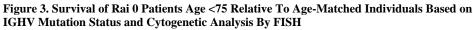




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A: Survival of Rai 0, IGHV mutated CLL patients Age <75 (n=344) relative to the agematched population. Figure 3B: Survival of Rai 0, IGHV unmutated CLL patients Age <75 (n=160) relative to the age-matched population. Figure 3C: Survival of Rai 0, CLL patients Age <75 (n=317) with favorable FISH (no 17p- or 11q-) relative to the age-matched population. Figure 3D: Survival of Rai 0, CLL patients Age <75 (n=27) with unfavorable FISH (e.g. 17p- or 11q-) relative to the age-matched population.

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|                               | Age <55 N=593 | Age 55-64 N=713 | Age 65-74 N=748 | Age 75 N=433 | P Value |
|-------------------------------|---------------|-----------------|-----------------|--------------|---------|
| Median Age at Diagnosis       | 67            | 09              | 69              | 08           |         |
| Male                          | 404 (68%)     | 493 (69%)       | (%99) /67       | 295 (68%)    | 0.739   |
| Rai Risk at diagnosis         |               |                 |                 |              |         |
| Low (Rai stage 0)             | 251 (43%)     | 383 (56%)       | 408 (57%)       | 245 (58%)    | <0.001  |
| Intermediate (Rai stage I-II) | 308 (53%)     | 257 (38%)       | (%88) 697       | 129 (31%)    |         |
| High (Rai stage III-IV)       | 21 (4%)       | 40 (6%)         | 40 (6%)         | 46 (11%)     |         |
| Missing                       | 13            | 33              | 31              | 13           |         |
| ALC ( $\times 10^{9}$ /L)     |               |                 |                 |              |         |
| 30                            | 449 (77%)     | (%62) (220)     | 631 (85%)       | 370 (86%)    | <0.001  |
| >30                           | 133 (23%)     | 148 (21%)       | 109 (15%)       | 60 (14%)     |         |
| Missing                       | 11            | 15              | 8               | 3            |         |
| CD38                          |               |                 |                 |              |         |
| Negative                      | 294 (66%)     | 376 (69%)       | 365 (68%)       | 195 (65%)    | 0.588   |
| Positive                      | 152 (34%)     | 169 (31%)       | 172 (32%)       | 105 (35%)    |         |
| Missing                       | 147           | 168             | 211             | 133          |         |
| ZAP-70                        |               |                 |                 |              |         |
| Negative                      | 203 (61%)     | 220 (60%)       | 219 (66%)       | 95 (66%)     | 0.340   |
| Positive                      | 130 (39%)     | 145 (40%)       | 113 (34%)       | 50 (34%)     |         |
| Missing                       | 260           | 348             | 416             | 288          |         |
| IGHV                          |               |                 |                 |              |         |
| Mutated                       | 150 (51%)     | 172 (54%)       | 170 (59%)       | 64 (61%)     | 0.166   |
| Unmutated                     | 144 (49%)     | 144 (46%)       | 119 (41%)       | 41 (39%)     |         |
| Missing                       | 299           | 397             | 459             | 328          |         |
| FISH (prior to treatment)     |               |                 |                 |              |         |
| 13q-                          | 118 (44%)     | 136 (43%)       | 126 (41%)       | 50 (34%)     | *       |
| Normal                        | 71 (27%)      | 76 (24%)        | 80 (26%)        | 41 (28%)     |         |
| +12                           | 43 (16%)      | 62 (20%)        | 58 (19%)        | 33 (23%)     |         |

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| 110-                         | -         | 0         | Age 05-74 N=740 | Age 10 IN-HOL | r value |
|------------------------------|-----------|-----------|-----------------|---------------|---------|
| h                            | 22 (8%)   | 26 (8%)   | 23 (7%)         | 16 (11%)      |         |
| 17p-                         | 10(4%)    | 11 (3%)   | 16 (5%)         | 6 (4%)        |         |
| other                        | 3 (1%)    | 5 (2%)    | 4 (1%)          | 1 (1%)        |         |
| Missing                      | 326       | 397       | 441             | 286           |         |
| FISH groupings               |           |           |                 |               |         |
| Normal, 13q, trisomy 12      | 232 (88%) | 274 (88%) | 264 (87%)       | 124 (85%)     | 662.0   |
| 11q, 17p                     | 32 (12%)  | 37 (12%)  | 39 (13%)        | 22 (15%)      |         |
| Treated                      | 268 (45%) | 282 (40%) | 296 (40%)       | 108 (25%)     | <0.001  |
| Median TFT Rai 0 patients    | 5.9       | 8.8       | 8.6             | not reached   | 0.047   |
| Median TFT Rai I-II patients | 1.9       | 2.5       | 2.5             | 3.4           | 0.216   |
| Dead                         | 112 (19%) | 171 (24%) | 254 (34%)       | 199 (46%)     | <0.001  |
| Median OS (yrs)              | 11.8      | 10.9      | 9.0             | 6.4           | <0.001  |

p-value not provided because chi-square test is not appropriate and calculation of Fisher's exact p-value was not feasible (too many FISH categories).

## Table 2

Prognostic Parameters, Time to First Treatment (TFT), and Overall Survival (OS)

|                                     | Median TFT (years) | P value | Median OS (years) | P value |
|-------------------------------------|--------------------|---------|-------------------|---------|
| Rai Risk at diagnosis (N=2397)      |                    |         |                   |         |
| Low (Rai stage 0)                   | 8.0                | < 0.001 | 11.44             | < 0.001 |
| Intermediate (Rai stage I-II)       | 2.4                |         | 8.8               |         |
| High (Rai stage III-IV)             | 0.1                |         | 5.2               |         |
| ALC (× 10 <sup>9</sup> /L) (N=2450) |                    |         |                   |         |
| 30                                  | 5.6                | < 0.001 | 10.2              | 0.003   |
| >30                                 | 2.4                |         | 8.7               | 1       |
| CD38 (N=1828)                       |                    |         |                   |         |
| Negative                            | 8.0                | < 0.001 | 11.9              | < 0.001 |
| Positive                            | 2.8                |         | 8.5               | 1       |
| ZAP-70 (N=1175)                     |                    |         |                   |         |
| Negative                            | 9.3                | < 0.001 | Not reached       | < 0.001 |
| Positive                            | 3.2                |         | 10.8              | 1       |
| IgVH Mutation (N=1004)              |                    |         |                   |         |
| Mutated                             | 11.0               | < 0.001 | Not reached       | < 0.001 |
| Unmutated                           | 2.8                |         | 9.7               | 1       |
| FISH (prior to treatment) (N=715)   |                    |         |                   |         |
| 13q-                                | Not reached        | < 0.001 | Not reached       | < 0.001 |
| Normal                              | 8.7                |         | Not reached       | 1       |
| +12                                 | 5.4                |         | 10.9              | 1       |
| 11q-                                | 2.4                |         | 8.4               |         |
| 17p-                                | 5.2                |         | 7.6               |         |
| FISH groupings                      |                    |         |                   |         |
| Normal, 13q, trisomy 12             | 8.7                | < 0.001 | Not reached       | < 0.001 |
| 11q, 17p                            | 3.7                |         | 7.6               |         |

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Table 3

Multivariate Models

| Prognostic Factor                          |      |                  |       |           |      |           |      |         |
|--|------|------------------|-------|-----------|------|-----------|------|---------|
|  | Ϋ́   | Age <55          | Age   | Age 55-64 | Age  | Age 65-74 | Age  | e 75    |
|  | HR   | p-value          | HR    | p-value   | HR   | p-value   | HR   | p-value |
| Rai Stage: High vs Low                     | 7.7  | <0.001           | 9.0   | <0.001    | 6.7  | <0.001    | 10.6 | <0.001  |
| Rai Stage: Intermediate vs Low             | 2.5  | <0.001           | 2.4   | <0.001    | 2.8  | <0.001    | 4.3  | <0.001  |
| ALC (>30 vs 30)                            | 1.3  | 0.070            | 1.8   | <0.001    | 2.2  | <0.001    | 2.9  | <0.001  |
| Rai Stage: High vs Low                     | 14.6 | <0.001           | 16.0  | <0.001    | 6.9  | <0.001    | 11.3 | <0.001  |
| Rai Stage: Intermediate vs Low             | 2.4  | <0.001           | 2.4   | <0.001    | 2.6  | <0.001    | 2.6  | <0.001  |
| CD38 (Positive vs negative)                | 2.3  | <0.001           | 2.8   | <0.001    | 1.7  | <0.001    | 1.7  | 0.037   |
| Rai Stage: High vs Low                     | 19.7 | <0.001           | 9.7   | <0.001    | 15.0 | <0.001    | 6.8  | 0.001   |
| Rai Stage: Intermediate vs Low             | 2.3  | <0.001           | 2.2   | <0.001    | 3.6  | < 0.001   | 1.1  | 0.802   |
| ZAP-70 (Positive vs negative)              | 2.0  | <0.001           | 1.9   | <0.001    | 2.8  | <0.001    | 5.6  | <0.001  |
| Rai Stage: High vs Low                     | 16.1 | <0.001           | 8.6   | <0.001    | 30.0 | < 0.001   | 11.7 | <0.001  |
| Rai Stage: Intermediate vs Low             | 2.7  | <0.001           | 2.1   | <0.001    | 2.5  | <0.001    | 1.3  | 0.587   |
| IgVH Mutation (UM vs. M)                   | 3.8  | <0.001           | 3.6   | <0.001    | 4.7  | <0.001    | 5.4  | 0.002   |
| Rai Stage: High vs Low                     | 6.2  | <0.001           | 6.2   | <0.001    | 10.0 | < 0.001   | 13.2 | <0.001  |
| Rai Stage: Intermediate vs Low             | 2.8  | <0.001           | 2.1   | 0.002     | 4.9  | <0.001    | 3.2  | 0.003   |
| FISH (17p, 11q vs normal, 13q, trisomy 12) | 2.7  | 0.001            | 3.0   | <0.001    | 2.8  | <0.001    | 2.0  | 0.078   |
|  | OVE  | OVERALL SURVIVAL | IAVIV | ,         |      |           |      |         |
| Prognostic Factor                          | Αg   | Age <55          | Ag(   | Age 55-64 | Age  | Age 65-74 | Age  | e 75    |
|  | HR   | p-value          | HR    | p-value   | HR   | p-value   | HR   | p-value |
| Rai Stage: High vs Low                     | 3.6  | 0.002            | 5.0   | <0.001    | 2.6  | <0.001    | 3.8  | <0.001  |
| Rai Stage: Intermediate vs Low             | 2.2  | <0.001           | 1.9   | <0.001    | 1.9  | <0.001    | 1.8  | <0.001  |
| ALC  | 1.4  | 0.126            | 1.5   | 0.027     | 1.7  | 0.001     | 1.5  | 0.052   |
| Rai Stage: High vs Low                     | 11.8 | <0.001           | 4.4   | <0.001    | 2.1  | 0.047     | 3.2  | <0.001  |
| Rai Stage: Intermediate vs Low             | 2.3  | 0.008            | 1.7   | 0.021     | 1.9  | 0.001     | 2.0  | 0.003   |
| CD38 (Positive vs negative)                | 1.6  | 0.077            | 2.2   | <0.001    | 1.3  | 0.138     | 1.5  | 0.038   |
| Rai Stage: High vs Low                     | 22.8 | 0.005            | 9.1   | <0.001    | 1.0  | 0.982     | 9.1  | <0.001  |

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| II   | IME TO | TIME TO FIRST TREATMENT | REATM | IENT      |     |           |     |         |
|--|--------|-------------------------|-------|-----------|-----|-----------|-----|---------|
| Prognostic Factor                          | Чg     | Age <55                 | Age   | Age 55-64 | уgA | Age 65-74 | Age | ge 75   |
|  | HR     | p-value                 | HR    | p-value   | HR  | p-value   | HR  | p-value |
| Rai Stage: Intermediate vs Low             | 2.8    | 0.013                   | 1.4   | 0.333     | 2.1 | 0.024     | 4.1 | <0.001  |
| ZAP-70 (Positive vs negative)              | 2.6    | 0.011                   | 2.2   | 0.025     | 1.2 | 0.578     | 0.7 | 0.426   |
| Rai Stage: High vs Low                     | 34.0   | <0.001                  | 1.7   | 0.475     | 2.1 | 0.319     | 2.7 | 0.358   |
| Rai Stage: Intermediate vs Low             | 2.2    | 0.074                   | 1.5   | 0.225     | 2.0 | 0.019     | 1.8 | 0.120   |
| IgVH Mutation (UM vs. M)                   | 6.2    | <0.001                  | 2.8   | 0.004     | 3.2 | <0.001    | 2.0 | 0.073   |
| Rai Stage: High vs Low                     | *      |                         | 4.5   | 0.007     | 7.4 | <0.001    | 3.7 | 0.002   |
| Rai Stage: Intermediate vs Low             | 1.7    | 0.294                   | 1.0   | 0.936     | 2.3 | 0.007     | 2.7 | 0.008   |
| FISH (17p, 11q vs normal, 13q, trisomy 12) | 5.2    | 0.001                   | 4.6   | 0.001     | 3.0 | 0.001     | 1.4 | 0.372   |
|  |        |                         |       |           |     |           |     |         |

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Too few events to estimate.