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Real-World Testing Patterns for Risk Assessment and Implications on the Adoption of Novel Therapeutics in Chronic Lymphocytic Leukemia: IgHV Mutation Status, FISH Cytogenetic, and Immunophenotyping

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Disclosures

- Consulting: BeiGene
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Introduction

- Prognostic testing, including immunoglobulin heavy-chain variable region gene (IgHV) mutation status, cytogenetic abnormalities by fluorescence in situ hybridization (FISH), and immunophenotyping, has been recommended in all newly diagnosed patients with chronic lymphocytic leukemia/small lymphocytic lymphoma (CLL/SLL) prior to treatment initiation, and even in previously treated patients in some settings
- Recent data have shown that disease with high-risk genetic features is better managed with novel agents than traditional chemoimmunotherapy. As such, the need for testing has become more relevant for disease management
- However, there is limited recent data on real-world patterns of testing for risk factor assessment and in-turn, pattern of evidence-based treatment selection



Objectives

To examine:

- Frequency and results of testing
- Timing of testing by line of therapy
- Factors associated with the receipt of testing



Methods

- **Study design:** Retrospective, observational study
- **Data source:** Flatiron Health EHR-derived database
- **Study period:** January 2014 to May 2021
- **Study population:**
 - Adults who were newly diagnosed with CLL/SLL
 - Index date: the first CLL/SLL diagnosis date during the identification period (July 2014 - February 2021)

Inclusion criteria

- Aged ≥ 18 years at index date
- Continuous enrollment of 6 months pre- and 3 months post-index date
- Patients who died within 3 months post-index date should be retained



Methods: Study Outcomes and Analysis Plan

- **Study outcomes:**
 - Frequency, results and timing of the following tests:
 - IgHV
 - FISH cytogenetic:
 - 11q deletion [del(11q)]
 - 13q deletion [del(13q)]
 - 17p deletion [del(17p)]
 - Trisomy 12 [+12])
 - Other biomarkers (including CD38 and ZAP-70) by immunophenotyping
- **Statistical analysis:**
 - Descriptive analyses: to examine the frequency and results in the overall population and compared by patient characteristics and across sociodemographic groups
 - Multivariable logistic regression: to examine factors associated with the likelihood of receiving testing
 - Statistical significance: p-value <0.05



Results: Patient Characteristics

- A total of 3,037 CLL patients were included
- Most patients were elderly (median age=73), male (62.3%), and white (74.6%)
- Most patients (92%) received treatment in community practices, with 54.1% commercially-insured

Table 1. Demographic and Clinical Characteristics of CLL Patient Population

| | CLL/SLL Patients (N=3,037) |
|--------------------------------|----------------------------|
| Age 65+ years, n (%) | 2,38 (78.4%) |
| Male, n (%) | 1,892 (62.3%) |
| Whites, n (%) | 2,265 (74.6%) |
| Hispanics, n (%) | 94 (3.1%) |
| Region, n (%) | |
| Midwest | 367 (12.1%) |
| Northeast | 515 (17.0%) |
| South | 1,217 (40.1%) |
| West | 680 (22.4%) |
| Other/missing | 258 (8.5%) |
| Health insurance, n (%) | |
| Commercial | 1,643 (54.1%) |
| Government | 1,120 (36.9%) |
| Other | 274 (9.0%) |

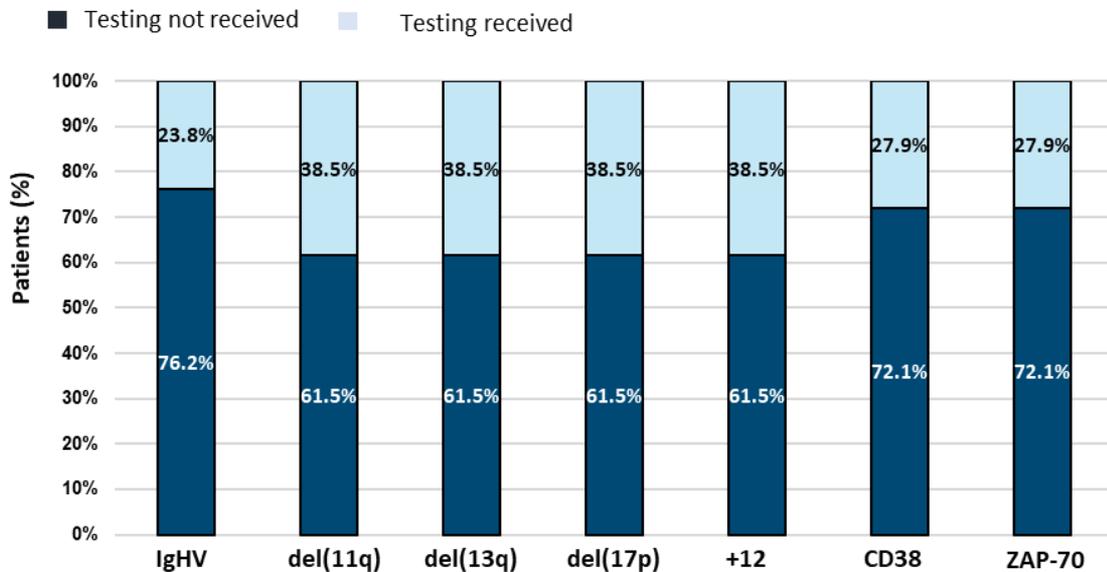
| | CLL/SLL Patients (N=3,037) |
|--|----------------------------|
| Community center, n (%) | 2,794 (92.0%) |
| BMI category at index | |
| Underweight (BMI < 18.5) | 30 (1.0%) |
| Normal Weight (18.5 <= BMI <25) | 620 (20.4%) |
| Overweight (25 <= BMI < 30) | 839 (27.6%) |
| Obese (>=30) | 801 (26.4%) |
| Unknown | 747 (24.6%) |
| Stage at index | |
| Stage I-II | 299 (9.9%) |
| Stage III | 106 (3.5%) |
| Stage IV | 193 (6.4%) |
| Stage Missing | 2,193 (72.2%) |
| ECOG status at index (Categorical), n (%) | |
| 0 | 719 (23.7%) |
| 1 | 464 (15.3%) |
| 2 | 95 (3.1%) |
| >=3 | 21 (0.7%) |
| Missing | 1,738 (57.2%) |



Results: Frequency of Risk Factor Testing

- Testing pattern
 - Over half of CLL patients did not receive risk factor testing: IgHV mutation analyses (76.2%, n=2,315), FISH (61.5%, n=1,868) and immunophenotyping (72.1%, n=2,190)
 - Of those who had testing, the majority (99%) had it done once prior to starting first-line of therapy

Figure 1. Real-world frequency of risk assessment testing



Results: Risk Factor Testing - Subgroup Analyses

- Significant differences in the receipt of testing were observed between different age, gender, race/ethnicity, and regional subgroups
- Among patients who received testing, the presence of high-risk biomarkers was as follows: unmutated IgHV (56.1%), del(17p) present (14.4%), del(11q) present (16.9%), and CD38 present (30.8%)
- Compared to patients <65 years, testing results in elderly patients ≥65 years showed a lower presence of unmutated IgHV (53.8%) and del(11q) (15.7%) while higher del(17p) (14.7%) and +12 (28.1%)

Table 2. Disparity in risk factor testing evaluation among various subgroups of patients with CLL

| | Age (<65 vs. 65+) (%) | Sex (M vs. F) (%) | Race (White vs. Non-White) (%) | Hispanic (Yes/No) (%) | Practice type (academic vs. community) (%) | Insurance (commercial vs. government) (%) |
|---------------|-----------------------------|-------------------------|--------------------------------------|-----------------------------|--|---|
| IgHV | 32.5, 21.4* | 25.4, 21* | 24.7, 21* | 16, 24.0 | 30, 23* | 25.2, 22.1 |
| FISH | 42.7, 37.3* | 40.0, 36.0* | 39.0, 37.0 | 35.1, 38.6 | 39.5, 38.4 | 38.1, 39.1 |
| CD38 or ZAP70 | 29.0, 26.1 | 29.8, 27.4 | 28.2, 26.9 | 19.1, 28.2 | 20.6, 28.5* | 28.2, 27.8 |

* p<0.05

Note: FISH cytogenetic tests include 11q deletion [del(11q)], 13q deletion [del(13q)], 17p deletion [del(17p)], Trisomy 12 [+12]

Abbreviations: CLL/SLL, Chronic lymphocytic leukemia /small lymphocytic leukemia



Results: Risk Factor Testing - Subgroup Analyses

- No significant disparity was observed in white vs. non-white patients except for a lower incidence of mutated IgHV and del(13q) presence
- Compared to tested men, tested women had a lower presence of unmutated IgHV (53.9%), del(11q) (11.4%) and CD38+ (25.8%) while higher del(17p) (18.2%)
- The impact of risk testing on therapy selection was investigated: patients with del(17p) had a higher likelihood than those who tested negative (73.6% vs. 48.4%) of being treated with novel agents (ibrutinib, acalabrutinib, or venetoclax)
- In contrast, 26.4% of those who tested del(17p) present and 39.8% among those who did not get tested received chemotherapy

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Results: Factors Associated with the Receipt of Testing

- 3 multivariable logistic regressions were performed to identify factors associated with receipt of IgHV, FISH cytogenetic, and immunophenotyping tests
- Patients who were older, female, or those living in the west of US were significantly less likely to receive IgHV testing
- Similar results were observed in the receipt of FISH cytogenetic testing: patients who were older, female or those living in the west of US were significantly less likely to receive FISH testing
- Multivariable analysis shows patients who live in the northeast or west were less likely to receive immunophenotyping tests

Table 3. Multivariate logistic regression: Factors/predictors associated with CLL patients receiving testing

| Effect | Testing (Outcome variable) | | | | | | | | |
|--|----------------------------|--------------|--------------|---------------|--------------|--------------|---------------|--------------|--------------|
| | IgHV | | | FISH | | | CD38 or ZAP70 | | |
| | Odds Ratio | Lower CL | Upper CL | Odds Ratio | Lower CL | Upper CL | Odds Ratio | Lower CL | Upper CL |
| Age group: 65+ vs <65 | 0.572* | 0.466 | 0.702 | 0.786* | 0.652 | 0.947 | 0.891 | 0.728 | 1.092 |
| Gender: Female vs Male | 0.815* | 0.682 | 0.974 | 0.857* | 0.736 | 0.999 | 0.868 | 0.734 | 1.026 |
| Race Non-white vs White | 0.854 | 0.694 | 1.051 | 0.958 | 0.804 | 1.142 | 0.974 | 0.804 | 1.179 |
| Ethnicity: Hispanic or Latino vs Unknown | 0.614 | 0.344 | 1.096 | 0.873 | 0.559 | 1.365 | 0.605 | 0.353 | 1.038 |
| Region (Reference: South) | | | | | | | | | |
| Mid West | 0.64 | 0.479 | 0.855 | 0.956 | 0.751 | 1.215 | 0.834 | 0.646 | 1.077 |
| Northeast | 0.868 | 0.683 | 1.104 | 1.01 | 0.817 | 1.249 | 0.758* | 0.603 | 0.953 |
| Other/Missing | 0.759 | 0.206 | 2.793 | 1.014 | 0.352 | 2.921 | 1.207 | 0.399 | 3.651 |
| West | 0.549* | 0.431 | 0.698 | 0.716* | 0.587 | 0.874 | 0.520* | 0.416 | 0.650 |
| Payer type (Reference: Commercial) | | | | | | | | | |
| Government | 0.970 | 0.801 | 1.174 | 1.11 | 0.942 | 1.307 | 1.030 | 0.862 | 1.230 |
| Other | 0.846 | 0.617 | 1.16 | 1.011 | 0.774 | 1.322 | 0.988 | 0.736 | 1.326 |
| Practice type: Academic vs Community | 1.412 | 0.375 | 5.32 | 0.92 | 0.311 | 2.716 | 0.424 | 0.135 | 1.331 |

* p<0.05

Note: FISH cytogenetic tests include 11q deletion [del(11q)], 13q deletion [del(13q)], 17p deletion [del(17p)], Trisomy 12 [+12]

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Discussion

- The NCCN guidelines recommend novel agents for patients with high-risk CLL/SLL. Thus, all patients are advised to complete risk-factor testing for both prognostication and selection of optimal, evidence-based therapy
- Despite the recommendations, there remains a significant number of patients who do not undergo FISH and/or IgHV mutation status testing prior to therapy
- Health disparities, across age, gender, race/ethnicity, regional subgroups, and insurance status, in testing are identified



Conclusions

- This real-world data highlights not only a significant gap in testing, but that this suboptimal testing is more common in vulnerable populations
- Despite identification of del(17), a quarter of CLL patients failed to receive novel agents in the frontline setting
- There is an unmet need for further education and refinement of clinical practice
- This is necessary to achieve the best clinical outcome in CLL patients through robust risk-assessment testing and optimal therapeutic triaging.

