Niraparib Maintenance Treatment Improves Time Without Symptoms or Toxicity (TWiST) Versus Routine Surveillance in Recurrent Ovarian Cancer: A TWiST Analysis of the ENGOT-OV16/NOVA Trial

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PURPOSE This study estimated time without symptoms or toxicity (TWiST) with niraparib compared with routine surveillance (RS) in the maintenance treatment of patients with recurrent ovarian cancer.

PATIENTS AND METHODS Mean progression-free survival (PFS) was estimated for niraparib and RS by fitting parametric survival distributions to Kaplan-Meier data for 553 patients with recurrent ovarian cancer who were enrolled in the phase III ENGOT-OV16/NOVA trial. Patients were categorized according to the presence or absence of a germline BRCA mutation—gBRCAmut and non-gBRCAmut cohorts. Mean time with toxicity was estimated based on the area under the Kaplan-Meier curve for symptomatic grade 2 or greater fatigue, nausea, and vomiting adverse events (AEs). Time with toxicity was the number of days a patient experienced an AE post-random assignment and before disease progression. TWiST was estimated as the difference between mean PFS and time with toxicity. Uncertainty was explored using alternative PFS estimates and considering all symptomatic grade 2 or greater AEs.

RESULTS In the gBRCAmut and non-gBRCAmut cohorts, niraparib treatment resulted in a mean PFS benefit of 3.23 years and 1.44 years, respectively, and a mean time with toxicity of 0.28 years and 0.10 years, respectively, compared with RS. Hence, niraparib treatment resulted in a mean TWiST benefit of 2.95 years and 1.34 years, respectively, compared with RS, which is equivalent to more than four-fold and two-fold increases in mean TWiST between niraparib and RS in the gBRCAmut and non-gBRCAmut cohorts, respectively. This TWiST benefit was consistent across all sensitivity analyses, including modeling PFS over 5-, 10-, and 15-year time horizons.

CONCLUSION Patients who were treated with niraparib compared with RS experienced increased mean TWiST. Thus, patients who were treated with niraparib in the ENGOT-OV16/NOVA trial experienced more time without symptoms or symptomatic toxicities compared with control.

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INTRODUCTION Ovarian cancer (OC) was diagnosed in 240,000 women worldwide in 2012.1 Most women are diagnosed with advanced cancer and prognosis is poor, which is reflected by OC being one of the most deadly forms of women’s cancer, with a 5-year survival rate of 47.4% and approximately 151,917 deaths in 2012.1,2 The standard of care for first-line treatment is platinum- and taxane-based chemotherapy, which results in high initial response rates; however, relapse rates for women who are diagnosed with advanced OC can be up to 95%.3,4 Given the lack of a cure for recurrent OC (ROC) and the negative impact on patient quality of life (QoL), there is a need for effective maintenance treatments that delay disease progression without the toxic effects that offset this benefit and affect patient QoL. The poly(ADP-ribose) polymerase inhibitor (PARPi) niraparib is approved by the US Food and Drug Administration and European Medicines Agency for use as maintenance treatment in
patients with recurrent epithelial ovarian, fallopian tube, or primary peritoneal cancer who have achieved complete or partial response to platinum-based chemotherapy. In the randomized controlled phase III ENGOT-OV16/NOVA trial of niraparib, patients were categorized according to the presence or absence of a germline BRCA mutation—gBRCA mut and non-gBRCA mut cohorts. Results demonstrated that niraparib treatment significantly extended median progression-free survival (PFS) in the gBRCA mut (21.0 months vs 5.5 months; P < .001) and non-gBRCA mut cohorts (9.3 months vs 3.9 months; P < .001).

The ENGOT-OV16/NOVA trial demonstrated that patient QoL remained stable during treatment and the pre-progression period, with no significant differences observed between treatment arms. Therefore, niraparib treatment prolongs PFS compared with placebo and maintains QoL. To explore this benefit further, this analysis estimated the time without symptoms or toxicity (TWiST) between niraparib and routine surveillance (RS) in patients with gBRCA mut and non-gBRCA mut tumors who were enrolled in ENGOT-OV16/NOVA. TWiST is an established methodology that partitions PFS into two health states and evaluates the tradeoff between treatment toxicity and time to progression; therefore, TWiST provides an estimate of time when a patient is free from progression and toxicity and thus likely to maintain a good QoL.

PATIENTS AND METHODS

Data Source

This analysis used data from the gBRCA mut and non-gBRCA mut cohorts of ENGOT-OV16/NOVA (ClinicalTrials.gov identifier: NCT01847274). Eligible patients were age 18 years or older; had histologically diagnosed ovarian, fallopian tube, or primary peritoneal cancer with predominantly high-grade serous histologic features; and received two or more platinum-based chemotherapy regimens. Patients must have had platinum-sensitive disease—complete or partial response and disease progression—more than 6 months after their penultimate platinum-based chemotherapy. Patients were randomly assigned 2:1 to receive once-daily niraparib 300 mg or placebo. Comparison with placebo was chosen to reflect the impact of niraparib on clinical practice. As such, placebo is referred to as RS from here on. The ENGOT-OV16/NOVA primary end point was PFS, which was defined as the time from treatment randomization to the earliest date of disease progression or death from any cause. Secondary end points included patient-reported outcomes, chemotherapy-free interval, PFS on the next line of therapy, time to first and second subsequent therapy, and overall survival. Efficacy data were analyzed in the intention-to-treat population, which was defined as all patients who underwent random assignment in each cohort. Safety data were analyzed in the safety population, defined as all patients who received one or more doses of treatment.

Study Participants

This analysis was performed on the intention-to-treat population of the gBRCA mut (niraparib, n = 138; RS, n = 65) and non-gBRCA mut cohorts (niraparib, n = 234; RS, n = 116) from ENGOT-OV16/NOVA. Patient baseline characteristics were balanced across cohorts (Data Supplement Table S1). One patient in the gBRCA mut cohort was excluded from the niraparib data set as the patient had only received first-line chemotherapy before initiating niraparib. One patient from the non-gBRCA mut cohort was excluded from the RS data set as no prior therapy line was recorded.

Statistical Analysis

Statistical analysis was performed using RStudio, version 1.1.38 (Boston, MA).

PFS. Patient-level data (database lock: June 20, 2016) from ENGOT-OV16/NOVA were used to derive Kaplan-Meier PFS curves for niraparib and RS for the gBRCA mut and non-gBRCA mut cohorts.

The time horizon for PFS extrapolation was based on expert clinical opinion such that patients could be progression free for up to 20 years. In addition, a recent National Institute for Health and Care Excellence (NICE) appraisal reported that it is biologically plausible for patients receiving niraparib to be progression free more than 10 years. Therefore, the base case analysis assumed that patients could be progression free for up to 20 years. In addition, the assumption that patients could be progression free for up to 5, 10, and 15 years was explored in sensitivity analyses.

PFS extrapolation was performed over a 20-year time horizon by fitting standard parametric survival distributions to niraparib and RS Kaplan-Meier PFS data. NICE Decision Support Unit guidelines were followed. The best-fitting distribution between treatment arms for each cohort, referred to as the base case, was selected on the basis of clinical plausibility and statistical and visual fit. The determination of the best-fitting distribution is further reported in the Data Supplement.

The generalized gamma distribution was identified as the statistically best-fitting distribution across treatment arms for both cohorts. Visual inspection confirmed that the generalized gamma distribution fit the data well (Fig 1). Clinical plausibility was assessed by comparing 5-year survival estimates with long-term survival available in platinum-sensitive ROC (PSROC) for PARPi. Study 19 olaparib data demonstrated that approximately 16% and 14% of patients with BRCA mut and BRCA wild-type tumors, respectively, remained on treatment—that is, progression free—at 5 years. Of the six standard parametric survival distributions, the generalized gamma distribution most closely modeled long-term survival from Study 19, with 22% and 13% of patients progression free at 5 years in the gBRCA mut and non-gBRCA mut cohorts, respectively.
Thus, this analysis used the generalized gamma distribution for the base case PFS extrapolation.

Mean PFS for niraparib and RS was calculated as the area under the curve (Data Supplement, Equation S1).

**Toxicity and TWiST.** In ENGOT-OV16/NOVA, safety was assessed by adverse event (AE) monitoring, laboratory tests, vital signs, and physical examinations. AEs were collected from the time of consent through treatment discontinuation. New serious AEs were collected for 30 days after treatment discontinuation. Few patients had no significant impact on QoL. The impact of hematologic AEs—for example, anemia, neutropenia, and thrombocytopenia—was not assessed, as a previous analysis concluded that they had no significant effect on QoL in ENGOT-OV16/NOVA. Time with toxicity for each patient was calculated as the number of days a patient experienced grade 2 or greater fatigue, nausea, or vomiting after random assignment and before disease progression. The number of days a patient experienced one of these AEs was calculated by subtracting the AE end date from the start date (Eq 1).

\[
AE\ duration = end\ date - start\ date. \quad (1)
\]

Multiple occurrences of these AEs experienced by a patient over the same time period were not counted twice. As such, the earliest start date and the latest end date were taken. In cases in which only the month and year were reported for the AE start or end date, data were imputed as the earliest start date and the latest end date were taken. In cases in which only the month and year were reported for the AE start or end date, data were imputed as the earliest start date and the latest end date were taken.

**FIG 1.** Kaplan-Meier (KM) curves for progression-free survival (PFS) and generalized gamma distributions over a 20-year time horizon for the (A) gBRCA mut and (B) non-gBRCA mut cohorts, and KM curves for time with toxicity (TOX) for the (C) gBRCA mut and (D) non-gBRCA mut cohorts. gBRCA mut, germline BRCA mutation; RS, routine surveillance.
progression. Time with toxicity was calculated for 202 patients (niraparib, n = 137; RS, n = 65) and 349 patients (niraparib, n = 234; RS, n = 115) in the gBRCAmut and non-gBRCAmut cohorts, respectively.

Kaplan-Meier time with toxicity curves were estimated for niraparib and RS for both cohorts (Fig 1). Mean time with toxicity was calculated as the restricted Kaplan-Meier mean (Data Supplement, Equation S1).

Mean TWiST was estimated as the difference between mean PFS and mean time with toxicity for each arm (Eq 2). The mean TWiST difference between niraparib and RS was calculated as the difference between mean TWiST for niraparib and RS.

\[ \text{Mean TWiST} = \text{mean PFS} - \text{mean time with toxicity}. \]  

**Sensitivity Analyses**

Sensitivity analyses assessed the impact of alternative PFS distributions, PFS time horizon, and the inclusion of all symptomatic grade 2 or greater AEs on the mean TWiST difference between niraparib and RS. To assess the impact of different PFS distributions, alternative distributions were selected by considering statistical and visual fit and clinical plausibility. Two additional standard parametric survival distributions—log-normal and log-logistic—were explored as they closely modeled 5-year survival data available from Study 19 (log-normal: gBRCAmut, 11.46%; non-gBRCAmut, 2.91%; log-logistic: gBRCAmut, 11.03%; non-gBRCAmut, 3.98%).

In addition, both distributions were a good statistical and visual fit to the observed data for both cohorts (Data Supplement, Tables S2–S3, Fig. S1–S4). Flexible PFS distributions were also explored per NICE Decision Support Unit guidelines and published literature. Two distributions—normal k = 1 and odds k = 3—were explored as they closely modeled 5-year survival data available from Study 19 for both cohorts (normal k = 1: gBRCAmut, 17.75%; non-gBRCAmut, 9.38%; odds k = 3: gBRCAmut, 20.54%; non-gBRCAmut, 9.21%). Both distributions were a good statistical fit to the observed data for both cohorts (Data Supplement Tables S4–S5). The Data Supplement (Fig S5–S8) shows that distributions fit the observed data well.

In addition, sensitivity analyses were performed on the PFS time horizon, where all patients were assumed to have experienced disease progression by 5, 10, or 15 years.

Finally, the inclusion of all symptomatic grade 2 or greater AEs experienced by more than 10% of patients in either treatment arm of ENGOT-OV16/NOVA (Table 1) was explored.

**Validation**

The TWiST methodology underwent internal and external validation. PFS extrapolation estimates were validated against published long-term survival data in PSROC for PARPi. In addition, modeled outcomes were validated using median survival time to align with the reporting of ENGOT-OV16/NOVA, which is standard practice in clinical trials. The analysis was conducted internally by two independent analysts (L.W. and H.G.), and the inputs were checked for accuracy by TESARO: A GSK Company (Waltham, MA). An external statistician reviewed the approach and provided suggestions for improvement, which included Kaplan-Meier toxicity curves and flexible PFS methodology. All feedback obtained was incorporated into the final analysis and this publication.

**RESULTS**

**Base Case Results**

Base case PFS extrapolation distributions—generalized gamma over 20 years—and Kaplan-Meier time with toxicity curves are presented in Figure 2 for niraparib and RS. Table 2 lists mean PFS, mean time with toxicity, and the estimated mean TWiST difference between niraparib and RS for both cohorts.

For the gBRCAmut cohort, niraparib was associated with a mean PFS benefit of 4.14 years and a mean time with toxicity of 0.31 years, which is equivalent to a mean TWiST of 3.83 years. In addition, RS was associated with mean PFS and mean time with toxicity of 0.91 and 0.03 years, respectively, which is equivalent to a mean TWiST of 0.88 years. Hence, niraparib treatment resulted in a mean TWiST benefit of 2.95 years compared with RS.

For the non-gBRCAmut cohort, niraparib was associated with a mean PFS of 2.59 years and a mean time with toxicity of 0.13 years, which is equivalent to a mean TWiST of 2.46 years. In addition, RS was associated with mean PFS and mean time with toxicity of 0.14 and 0.02 years, respectively, which is equivalent to a mean TWiST of 1.12 years. Hence, niraparib treatment resulted in a mean TWiST benefit of 1.34 years compared with RS.

**Sensitivity Analyses**

For all sensitivity analyses, niraparib demonstrated a consistent mean TWiST benefit compared with RS when examining PFS at 5, 10, and 15 years (Table 2 and Data Supplement Fig S9–S25). When extrapolated PFS was modeled with more conservative estimates, mean TWiST benefit ranged from 1.15 to 3.65 years and 0.52 to 1.23 years for the gBRCAmut and non-gBRCAmut cohorts, respectively. When all symptomatic grade 2 or greater AEs were included, mean TWiST benefit was 3.01 and 1.22 for the gBRCAmut and non-gBRCAmut cohorts, respectively.

**Valueation**

Median modeled and observed PFS from ENGOT-OV16/NOVA were similar for both niraparib and RS for gBRCAmut (niraparib: modeled v observed, 21.0 v 21.0 months; RS: modeled v observed, 5.52 v 5.5 months) and non-gBRCAmut
DISCUSSION

This analysis sought to evaluate the difference in TWiST between niraparib and RS in the maintenance treatment of patients with PSROC. TWiST is the amount of time—measured in years—during which patients are progression and toxicity free and thus likely to maintain good QoL. Mean PFS extrapolated from ENGOT-OV16/NOVA over a 20-year time horizon demonstrated that patients receiving niraparib, compared with RS, experienced increased mean PFS in both cohorts. Sensitivity analyses were performed at additional clinically relevant 5-, 10-, and 15-year PFS timeframes, demonstrating improved TWiST in niraparib-treated patients. Furthermore, subtracting the mean time with toxicity for grade 2 or greater fatigue, nausea, and vomiting from mean PFS resulted in an increased mean TWiST. Base case results showed that niraparib was associated with a mean TWiST benefit compared with RS. These TWiST benefits demonstrate that patients receiving niraparib will maintain good QoL. Specifically, compared with RS, patients with gBRCAmut and non-gBRCAmut tumors who receive niraparib will gain 2.95 years and 1.34 years, respectively, during which they are progression and toxicity free. In addition, results demonstrate that the mean TWiST for patients receiving RS in the gBRCAmut and non-gBRCAmut cohorts, respectively. Robustness of the niraparib TWiST benefit compared with RS was confirmed through consistency across all sensitivity analyses in both cohorts. The greatest mean TWiST benefit observed in these sensitivity analyses for niraparib, compared with RS, was 3.65 years and 1.23 years for the gBRCAmut and non-gBRCAmut cohorts, respectively. When alternative PFS estimates were considered, niraparib mean TWiST was up to five-fold and two-fold greater than the mean TWiST for patients receiving RS in the gBRCAmut and non-gBRCAmut cohorts, respectively.

### TABLE 1. Incidence of All Symptomatic Grade ≥ 2 Adverse Events Experienced by ≥ 10% of Patients in Either Treatment Arm by Cohort of ENGOT-OV16/NOVA

<table>
<thead>
<tr>
<th>Adverse Event</th>
<th>gBRCAmut Cohort</th>
<th>Non-gBRCAmut Cohort</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Niraparib (n = 136)</td>
<td>RS (n = 65)</td>
</tr>
<tr>
<td>Fatigue</td>
<td>43 (31.6)</td>
<td>8 (12.3)</td>
</tr>
<tr>
<td>Nausea</td>
<td>30 (22.1)</td>
<td>6 (9.2)</td>
</tr>
<tr>
<td>Hypertension</td>
<td>26 (19.1)</td>
<td>4 (6.2)</td>
</tr>
<tr>
<td>Constipation</td>
<td>17 (12.5)</td>
<td>2 (3.1)</td>
</tr>
<tr>
<td>Headache</td>
<td>15 (11.0)</td>
<td>2 (3.1)</td>
</tr>
<tr>
<td>Vomiting</td>
<td>14 (10.3)</td>
<td>1 (1.5)</td>
</tr>
<tr>
<td>Abdominal pain</td>
<td>10 (7.4)</td>
<td>5 (7.7)</td>
</tr>
<tr>
<td>Urinary tract infection</td>
<td>8 (5.9)</td>
<td>4 (6.2)</td>
</tr>
<tr>
<td>Dyspepsia</td>
<td>7 (5.1)</td>
<td>1 (1.5)</td>
</tr>
<tr>
<td>Back pain</td>
<td>7 (5.1)</td>
<td>2 (3.1)</td>
</tr>
<tr>
<td>Dyspnea</td>
<td>6 (4.4)</td>
<td>0</td>
</tr>
<tr>
<td>Decreased appetite</td>
<td>6 (4.4)</td>
<td>1 (1.5)</td>
</tr>
<tr>
<td>Diarrhea</td>
<td>5 (3.7)</td>
<td>2 (3.1)</td>
</tr>
<tr>
<td>Arthralgia</td>
<td>5 (3.7)</td>
<td>2 (3.1)</td>
</tr>
<tr>
<td>Insomnia</td>
<td>5 (3.7)</td>
<td>0</td>
</tr>
<tr>
<td>Nasopharyngitis</td>
<td>4 (2.9)</td>
<td>1 (1.5)</td>
</tr>
<tr>
<td>Abdominal distension</td>
<td>2 (1.5)</td>
<td>1 (1.5)</td>
</tr>
<tr>
<td>Dizziness</td>
<td>2 (1.5)</td>
<td>1 (1.5)</td>
</tr>
<tr>
<td>Cough</td>
<td>2 (1.5)</td>
<td>0</td>
</tr>
<tr>
<td>Myalgia</td>
<td>1 (0.7)</td>
<td>1 (1.5)</td>
</tr>
<tr>
<td>Palpitations</td>
<td>1 (0.7)</td>
<td>0</td>
</tr>
</tbody>
</table>

NOTE. Data provided as No. (%) unless otherwise noted. Abbreviations: gBRCAmut, germline BRCA mutation; RS, routine surveillance.
patients receiving niraparib spend more time progression and toxicity free with respect to grade 2 or greater fatigue, nausea, and vomiting than do patients receiving RS. This analysis demonstrates that niraparib treatment maintained QoL among progression-free patients, which is consistent with a previous analysis that showed that patients receiving niraparib had stable QoL during the preprogression period. Furthermore, all symptoms as measured by the Functional Assessment of Cancer Therapy-Ovarian Symptom Index, with the exception of nausea, remained stable or improved over time with niraparib treatment.\textsuperscript{8} ROC has been associated with a negative psychological and physical impact on patients, particularly as patients receive additional lines of subsequent chemotherapy.\textsuperscript{16} Therefore, as PFS decreases with each subsequent line of chemotherapy,\textsuperscript{17} and subsequent chemotherapy is associated with burdensome AEs and a negative impact on QoL, there is a need for an alternative to RS for the maintenance treatment of patients with PSROC. Effective PARPi, such as niraparib, which can extend PFS without negatively affecting QoL, can significantly improve the prognosis of PSROC compared with RS.

This publication reports one of the first TWiST analyses conducted for a PARPi for the maintenance treatment of patients with PSROC and is the first TWiST analysis that evaluates both gBRCA\textsuperscript{mut} and non-gBRCA\textsuperscript{mut} populations. The only other published TWiST analysis—by Friedlander et al\textsuperscript{10}—evaluated the TWiST difference between olaparib and placebo in the SOLO-2 trial; however, these results are not comparable with those presented here as a result of a different methodology used. Friedlander et al\textsuperscript{10} used PFS trial data only, whereas our study extrapolated PFS. Furthermore, the TWiST difference between olaparib and placebo was limited to gBRCA\textsuperscript{mut} patients only because of trial eligibility.\textsuperscript{10}

FIG 2. Kaplan-Meier curves for time with toxicity (TOX), mean progression-free survival (PFS) generalized gamma distributions, and mean time without symptoms or toxicity (TWiST) for (A) niraparib and (B) routine surveillance (RS) in the gBRCA\textsuperscript{mut} cohort and for (C) niraparib and (D) RS in the non-gBRCA\textsuperscript{mut} cohort over a 20-year time horizon. gBRCA\textsuperscript{mut}, germline BRCA mutation.
TABLE 2. Base Case and Sensitivity Analysis Results for Niraparib Versus Routine Surveillance

<table>
<thead>
<tr>
<th>Scenario</th>
<th>Mean PFS, Years</th>
<th>Mean Time With Toxicity, Years</th>
<th>Mean TWiST, Years</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>gBRCAmut cohort</strong></td>
<td></td>
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<tr>
<td>Base case</td>
<td></td>
<td></td>
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<tr>
<td>Generalized gamma PFS distribution (20-year time horizon)</td>
<td>4.14 (2.20 to 7.04; 1.25)</td>
<td>0.91 (0.51 to 1.90; 0.35)</td>
<td>3.23</td>
</tr>
<tr>
<td>Sensitivity analysis</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Generalized gamma PFS distribution (5-year time horizon)</td>
<td>2.21 (1.73 to 2.81; 0.25)</td>
<td>0.78 (0.52 to 1.22; 0.16)</td>
<td>1.43</td>
</tr>
<tr>
<td>Generalized gamma PFS distribution (10-year time horizon)</td>
<td>3.09 (2.09 to 4.50; 0.60)</td>
<td>0.85 (0.51 to 1.50; 0.24)</td>
<td>2.23</td>
</tr>
<tr>
<td>Generalized gamma PFS distribution (15-year time horizon)</td>
<td>3.68 (2.15 to 5.76; 0.94)</td>
<td>0.89 (0.50 to 1.74; 0.30)</td>
<td>2.79</td>
</tr>
<tr>
<td>Log-normal PFS distribution</td>
<td>2.53 (1.72 to 3.67; 0.47)</td>
<td>0.63 (0.47 to 0.86; 0.09)</td>
<td>1.90</td>
</tr>
<tr>
<td>Log-logistic PFS distribution</td>
<td>2.57 (1.80 to 3.51; 0.42)</td>
<td>0.65 (0.48 to 0.91; 0.10)</td>
<td>1.92</td>
</tr>
<tr>
<td>Normal, k = 1 PFS distribution</td>
<td>3.66 (1.92 to 9.79; 1.97)</td>
<td>0.72 (0.50 to 1.98; 0.41)</td>
<td>2.94</td>
</tr>
<tr>
<td>Odds, k = 3 PFS distribution</td>
<td>4.80 (2.05 to 17.41; 4.02)</td>
<td>0.87 (0.49 to 6.38; 2.04)</td>
<td>3.93</td>
</tr>
<tr>
<td>Inclusion of all symptomatic grade ≥ 2 AEs</td>
<td>4.14 (2.20 to 7.04; 1.25)</td>
<td>0.91 (0.51 to 1.90; 0.35)</td>
<td>3.23</td>
</tr>
<tr>
<td><strong>Non-gBRCAmut cohort</strong></td>
<td></td>
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<tr>
<td>Base case</td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>Generalized gamma PFS distribution (20-year time horizon)</td>
<td>2.59 (1.60 to 3.82; 0.56)</td>
<td>1.14 (0.65 to 1.98; 0.33)</td>
<td>1.44</td>
</tr>
<tr>
<td>Sensitivity analysis</td>
<td></td>
<td></td>
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</tr>
<tr>
<td>Generalized gamma PFS distribution (5-year time horizon)</td>
<td>1.51 (1.19 to 1.86; 0.15)</td>
<td>0.89 (0.52 to 1.11; 0.15)</td>
<td>0.63</td>
</tr>
<tr>
<td>Generalized gamma PFS distribution (10-year time horizon)</td>
<td>2.00 (1.39 to 2.66; 0.31)</td>
<td>1.09 (0.59 to 1.49; 0.22)</td>
<td>0.91</td>
</tr>
<tr>
<td>Generalized gamma PFS distribution (15-year time horizon)</td>
<td>2.33 (1.50 to 3.27; 0.44)</td>
<td>1.13 (0.68 to 1.80; 0.28)</td>
<td>1.20</td>
</tr>
<tr>
<td>Log-normal PFS distribution</td>
<td>1.32 (1.05 to 1.72; 0.16)</td>
<td>0.98 (0.47 to 0.75; 0.06)</td>
<td>0.73</td>
</tr>
<tr>
<td>Log-logistic PFS distribution</td>
<td>1.43 (1.12 to 1.87; 0.17)</td>
<td>0.60 (0.47 to 0.78; 0.07)</td>
<td>0.83</td>
</tr>
<tr>
<td>Normal, k = 1 PFS distribution</td>
<td>2.23 (1.30 to 5.22; 0.99)</td>
<td>0.90 (0.55 to 2.77; 0.60)</td>
<td>1.33</td>
</tr>
<tr>
<td>Odds, k = 3 PFS distribution</td>
<td>2.41 (1.36 to 6.77; 1.38)</td>
<td>1.37 (0.61 to 5.22; 1.24)</td>
<td>1.04</td>
</tr>
<tr>
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<td>1.14 (0.65 to 1.80; 0.33)</td>
<td>1.44</td>
</tr>
</tbody>
</table>

Abbreviations: AE, adverse event; gBRCAmut, germline BRCA mutation; k, knotts; PFS, progression-free survival; RS, routine surveillance; TWIST, time without symptoms or toxicity.
extrapolation of ENGOT-OV16/NOVA Kaplan-Meier PFS data, as opposed to comparing mean time with toxicity with restricted mean PFS only, thereby estimating the complete TWiST benefit expected for patients. Finally, this analysis has been demonstrated to be robust, as a TWiST benefit with niraparib was consistent across all sensitivity analyses performed.

This analysis has some limitations. The base case PFS was extrapolated over 20 years. Although it is unrealistic that long-term PFS would align with this cutoff point, this was based on expert clinical opinion and was accepted in two previous health technology appraisal submissions.18,19 Furthermore, the impact of alternative PFS estimates was explored through sensitivity analyses. Results indicated that patients who were treated with niraparib still experienced a TWiST benefit compared with RS even when more conservative PFS estimates were modeled. An additional limitation is that only the most common symptomatic non-hematologic grade 2 or greater AEs were included in the base case analysis. However, all symptomatic grade 2 or greater AEs experienced by more than 10% of ENGOT-OV16/NOVA patients was explored in a sensitivity analysis. These results demonstrated that niraparib-treated patients continue to experience an increased mean TWiST compared with RS, with the gBRCAmut mean TWiST increasing with respect to the base case analysis because of a smaller proportional increase in niraparib toxicity compared with RS. This is in agreement with results of a previous analysis in which a higher proportion of patients receiving RS reported pain compared with niraparib-treated patients.8

Future analyses could include a quality-adjusted TWiST analysis when mature overall survival data are available from ENGOT-OV16/NOVA. Moreover, an analysis of the safety profile of niraparib in ENGOT-OV16/NOVA recommended that patients with a baseline body weight of less than 77 kg or platelet count less than 150,000/μL may benefit from a 200-mg per day starting dose.20 Subsequent investigations could focus on the TWiST difference with this alternative starting dose.

TWiST estimates from ENGOT-OV16/NOVA indicate that niraparib-treated patients experienced increased mean TWiST compared with patients receiving RS. Therefore, patients receiving niraparib compared with control in ENGOT-OV16/NOVA experienced more time without symptoms or symptomatic toxicities and thus have the potential to experience longer undiminished QoL.
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**AUTHORS’ DISCLOSURES OF POTENTIAL CONFLICTS OF INTEREST**

Niraparib Maintenance Treatment Improves Time Without Symptoms or Toxicity (TWiST) Versus Routine Surveillance in Recurrent Ovarian Cancer: A TWiST Analysis of the ENGOT-OV16/NOVA Trial

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