

## **Supplemental Tables and Figures - MS#ADV-2021-006923R3**

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**Supplemental Figure 3.** Kaplan-Meier estimates of survival (progression-free and overall) according to risk groups based on the risk factors TMTV >220 cm<sup>3</sup> and ECOG-PS ≥2 in the PETAL (A, B), GOYA (C,D) and real-world (E,F) series in patients aged <60 years.

**Supplemental Figure 4.** Kaplan-Meier estimates of survival (progression-free and overall) according to risk groups based on the risk factors TMTV >220 cm<sup>3</sup> and ECOG-PS ≥2 in the PETAL (A, B), GOYA (C,D) and real-world (E,F) series in patients aged ≥60 years.

**Supplemental Table 1. Main characteristics of the two clinical trials (GOYA, PETAL) and the real-world series**

<b>Study name</b>	<b>Clinical trial identifier Phase N patients</b>	<b>Inclusion period</b>	<b>N centers and location</b>	<b>Patient characteristics</b>	<b>Treatment</b>
<b>GOYA</b>	NCT01287741 Phase III n=1315	July 2011 to June 2014	207 centers in 29 countries	Previously untreated DLBCL, ECOG-PS 0-2, and either IPI $\geq 2$ or IPI = 1 with or without bulky disease (defined as one lesion $\geq 7.5$ cm), or with an IPI = 0 and bulky disease, aged $\geq 18$ years	Obinutuzumab (Gazyva/Gazyvaro) G-CHOP versus R-CHOP
<b>PETAL</b>	NCT00554164 Phase III n= 510	November 2007 to December 2012	23 German centers	Untreated aggressive CD20-positive lymphoma with ECOG-PS 0-3  (patients with T-cell lymphoma were excluded)	After 2 cycles of R-CHOP, PET-positive patients with untreated aggressive CD20-positive lymphoma with ECOG-PS $\leq 3$ were randomly assigned to treatment, 4 cycles of R-CHOP $\pm 2$ additional doses of rituximab
<b>Real world series</b>	n=349	2004 to 2016	4 European centers	Patients aged $\geq 18$ years, with DLBCL	R-CHOP, 6 or 8 cycles

**Supplemental Table 2. Clinical characteristics of patients stratified by age (<60 years and ≥60 years) and by series**

Characteristic, n (%)	Age <60 years			Age ≥60 years		
	PETAL (n = 229)	GOYA (n = 557)	Real-world series (n = 94)	PETAL (n=280)	GOYA (n=758)	Real-world series (n=234)
<b>Median age, years (range)</b>	50 (18-59)	49 (18-59)	48.5 (17-59)	69 (60-80)	68 (60-86)	69 (60-80)
<b>Sex</b>						
<b>Male</b>	126 (55.0)	245 (44.0)	53 (56.4)	153 (54.6)	386 (50.9)	109 (46.6)
<b>Female</b>	103 (45.0)	312 (56.0)	41 (43.6)	127 (45.4)	372 (49.1)	125 (53.4)
<b>Histology</b>						
<b>DLBCL NOS</b>	198 (86.5)	508 (91.2)	16 (84.2)	280 (100)	678 (89.4)	155 (92.3)
<b>FL grade 3B</b>	14 (6.1)	0 (0)	0 (0)	0 (0)	0 (0)	0 (0)
<b>De novo transformed</b>	17 (7.4)	0 (0)	0 (0)	0 (0)	0 (0)	10 (6.0)
<b>Other</b>	0 (0)	49 (8.8)	3 (15.8)	0 (0)	80 (10.6)	3 (1.8)
<b>Missing</b>	0	0	75	0	0	66
<b>ECOG-PS</b>						
<b>0–1</b>	211 (93.4)	490 (88.0)	79 (84.0)	240 (86.0)	662 (87.3)	167 (71.4)
<b>≥ 2</b>	15 (6.6)	67 (12.0)	15 (16.0)	39 (14.0)	96 (12.7)	67 (28.6)
<b>Missing</b>	3	0	0	0	0	0
<b>Ann Arbor stage</b>						
<b>I-II</b>	101 (44.3)	135 (24.2)	39 (41.5)	113 (40.4)	179 (23.6)	49 (20.9)
<b>III-IV</b>	127 (55.7)	422 (75.8)	55 (58.5)	167 (59.6)	579 (76.4)	185 (79.1)
<b>Missing</b>	1	0	0	0	0	0
<b>Extranodal sites</b>						
<b>&lt; 2</b>	156 (68.4)	353 (63.4)	61 (64.9)	193 (68.9)	501 (66.1)	138 (62.2)
<b>≥ 2</b>	72 (31.6)	204 (36.6)	33 (35.1)	87 (31.1)	257 (33.9)	84 (37.8)
<b>Missing</b>	0	0	1	0	0	12
<b>Elevated LDH (&gt; ULN)</b>						

Characteristic, n (%)	Age <60 years			Age ≥60 years		
	PETAL (n = 229)	GOYA (n = 557)	Real-world series (n = 94)	PETAL (n=280)	GOYA (n=758)	Real-world series (n=234)
<b>No</b>	101 (44.3)	235 (42.2)	40 (43.0)	121 (43.2)	323 (42.6)	82 (35.0)
<b>Yes</b>	127 (55.7)	322 (57.8)	53 (57.0)	159 (56.8)	431 (56.9)	152 (65.0)
<b>Missing</b>	1	0	1	0	4	0
<b>aalPI</b>						
<b>0–1</b>	137 (60.4)	293 (52.6)	50 (53.2)	153 (54.6)	393 (51.8)	89 (38.0)
<b>2–3</b>	90 (39.7)	264 (47.4)	44 (46.8)	127 (45.4)	365 (48.2)	145 (62.0)
<b>IPI</b>						
<b>0-2</b>	176 (77.2)	417 (74.9)	60 (63.8)	138 (49.3)	319 (42.1)	76 (32.5)
<b>3-5</b>	52 (22.8)	140 (25.1)	34 (36.2)	142 (50.7)	439 (57.9)	158 (67.5)
<b>Missing</b>	1	0	0	0	0	0
<b>Treatment</b>						
<b>R-CHOP</b>	222 (96.9)	274 (49.2)	94 (100)	263 (93.9)	380 (50.1)	234 (100)
<b>G-CHOP</b>	-	283 (50.8)	-	-	378 (49.9)	-
<b>Intensified CHOP / CHOP</b>	7 (3.1)	-	-	17 (6.1)	-	-
<b>TMTV</b>						
<b>Median cm<sup>3</sup> (range)</b>	178 (1-6750)	311.7 (2-8113)	153 (0-2416)	173 (1-8896)	250.8 (1.7-5334)	207.2 (0-3764)
<b>TMTV &gt;220 cm<sup>3</sup></b>	109 (47.6)	333 (59.8)	34 (36.2)	121 (43.2)	392 (51.7)	114 (48.7)
<b>TMTV and ECOG-PS risk factors</b>						
<b>0 factor</b>	119 (52.0)	207 (37.2)	56 (59.6)	151 (54.1)	346 (45.6)	98 (41.9)
<b>1 factor</b>	96 (41.9)	300 (53.9)	27 (28.7)	96 (34.4)	336 (44.3)	91 (38.9)
<b>2 factors</b>	14 (6.1)	50 (9.0)	11 (11.7)	32 (11.5)	76 (10.0)	45 (19.2)

\* TMTV > 220 cm<sup>3</sup> and ECOG-PS ≥2

### Supplemental Table 3 .Calibration methodology

Initially proposed for microarrays and then validated for several imaging biomarkers in PET (Orlhac et al J Nucl Med 2018), CT (Orlhac et al Radiology 2019) and MR (Orlhac et al Eur Rad 2021), the ComBat approach estimates the location (mean) and/or scale (variance) (L/S) adjustments needed to align the statistical distributions of a given biomarker measured under different conditions. Here, we applied this approach to all patients from the phase III GOYA trial (NCT01287741) with available TMTV to a cohort within the phase III REMARC trial (NCT01122472).

Following the language as described for microarrays in ComBat, assuming  $TMTV_{ij}$  represents the total metabolic tumor volume for sample  $j$  from clinical trial  $i$ . We define a L/S model that assumes:

$$TMTV_{ij} = \alpha + X\beta + \gamma_i + \delta_i e_{ij}$$

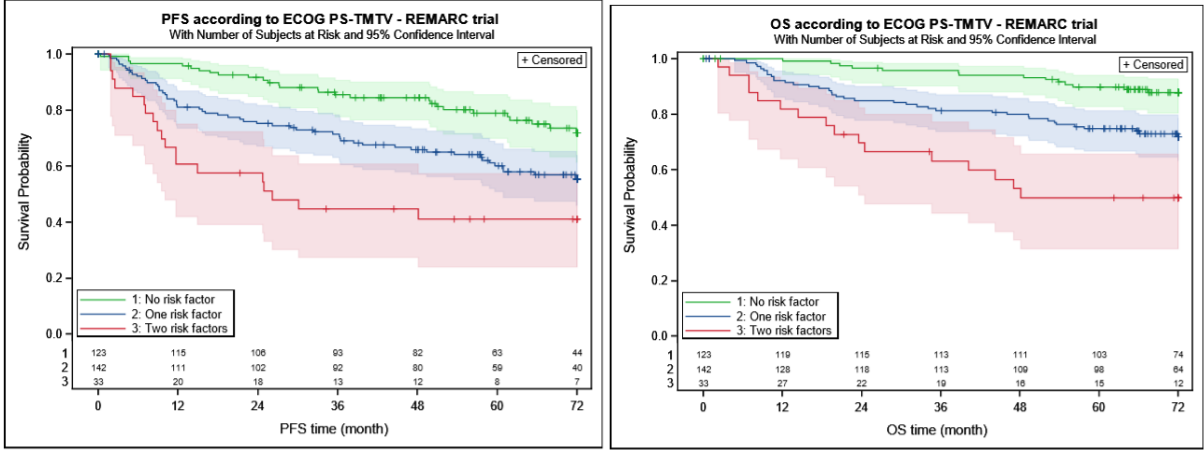
where  $TMTV_{ij}$  is the overall TMTV expression,  $X$  is a design matrix for sample conditions, and  $\beta$  is the vector of regression coefficients corresponding to  $X$ . The error terms,  $e_{ij}$  can be assumed to follow a Normal distribution with expected value of zero and variance  $\sigma^2$ . The  $i$  and  $j$  represent the additive and multiplicative trial effects for trial  $i$ . The trial-adjusted data,  $TMTV_{ij}^*$ , are given by

$$TMTV_{ij}^* = \frac{TMTV_{ij} - \hat{\alpha} + X\hat{\beta} - \hat{\gamma}_i}{\hat{\delta}_i} + \hat{\alpha} + X\hat{\beta}$$

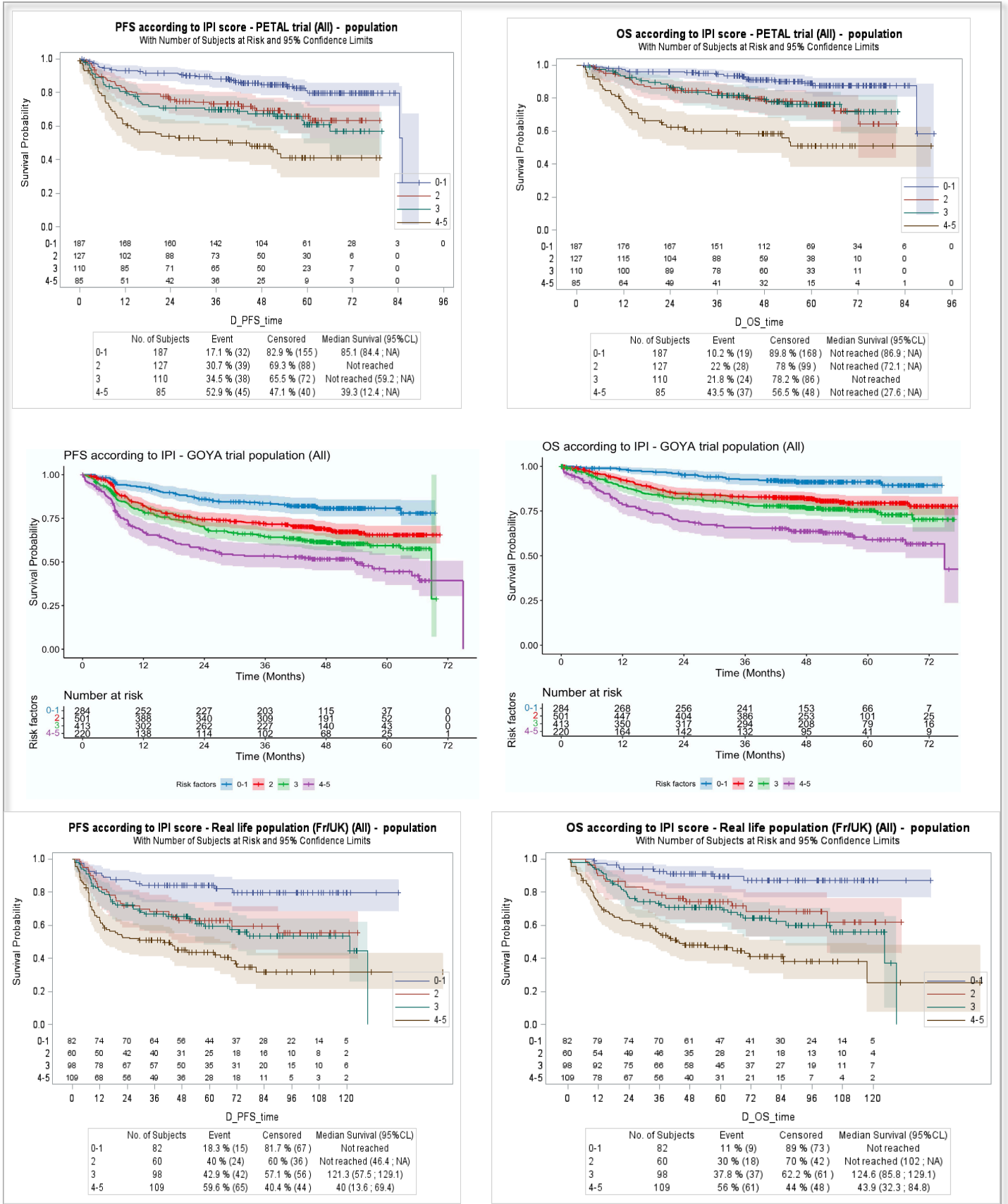
Where  $\hat{\alpha}$ ,  $\hat{\beta}$ ,  $\hat{\gamma}_i$ , and  $\hat{\delta}_i$  are the estimators for respective parameters  $\alpha$ ,  $\beta$ ,  $\gamma_i$ , and  $\delta_i$ .

We applied the ComBat L/S adjustment to 1315 patients with available TMTV measurements from the phase III GOYA trial and leveraged 301 patients from the phase III REMARC trial as a reference using study and age  $\geq 60$  years as an indicator variable in our model matrix.

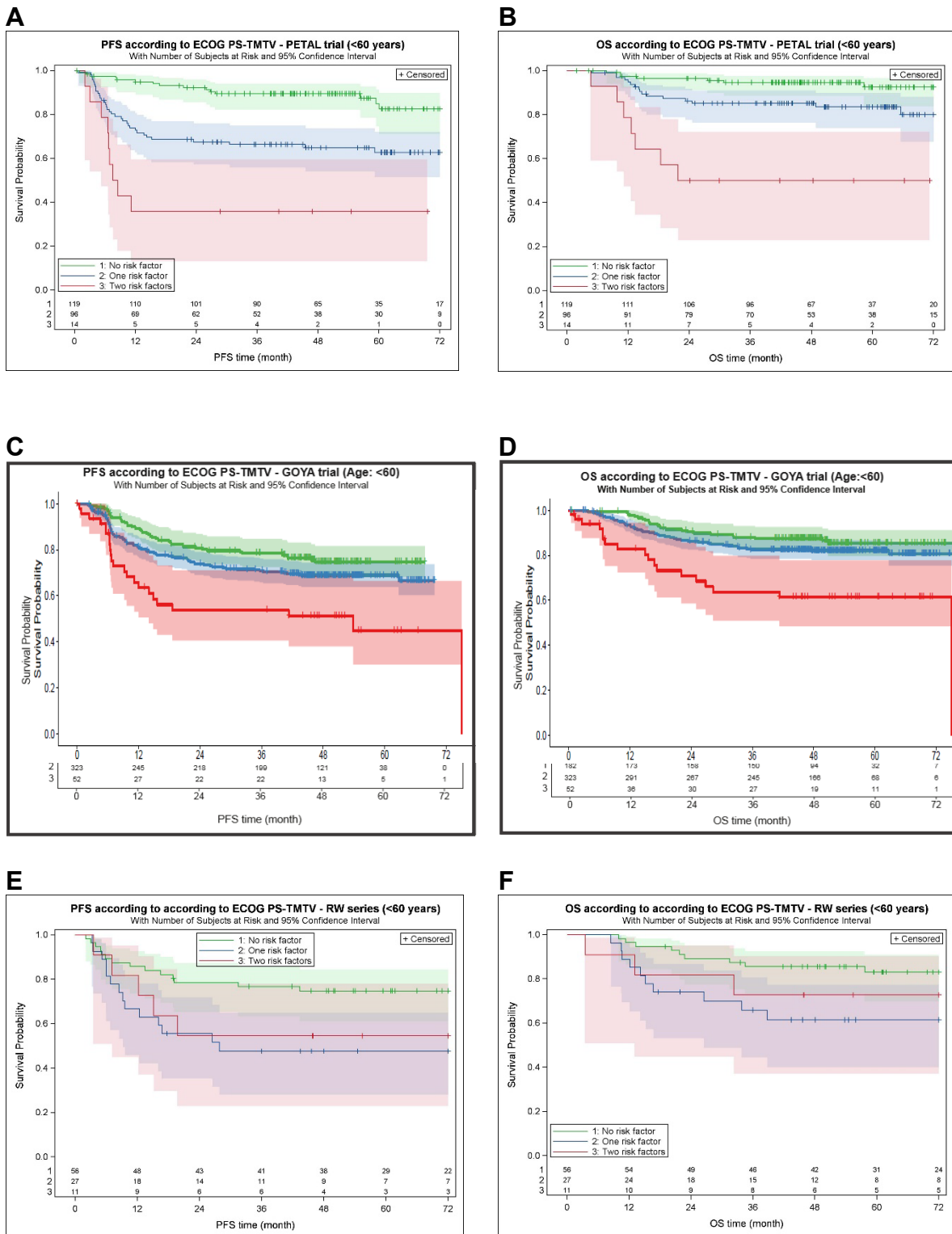
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**Supplemental Figure 4.** Kaplan-Meier estimates of survival (progression-free and overall) according to risk groups based on the risk factors TMTV >220 cm<sup>3</sup> and ECOG-PS ≥2 in the PETAL (A, B), GOYA (C,D) and real-world (E,F) series in patients aged ≥60 years.

