

Prephase rituximab/prednisone therapy and aging-related, proinflammatory cytokine milieu in older, vulnerable patients with newly diagnosed diffuse large B-cell lymphoma

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Supplemental Table 1. Toxicity events by stage

Characteristic	N	Stage I/II, N = 13	Stage III/IV, N = 20	p-value	q-value
Toxic events	33	8 (62%)	14 (70%)	0.7	>0.9
Severe toxic events	33	4 (31%)	8 (40%)	0.7	>0.9
G3+ nonheme toxicity / G4+ heme toxicity	33			>0.9	>0.9
At least 1 G3+ nonheme / G4+ heme toxicity		6 (46%)	10 (50%)		
No G3+ nonheme / G4+ heme toxicities		7 (54%)	10 (50%)		
G3+ nonheme toxicities	33			>0.9	>0.9
At least 1 G3+ nonheme toxicity		6 (46%)	10 (50%)		
No G3+ nonheme toxicities		7 (54%)	10 (50%)		
G3+ heme toxicities	33			0.026	0.2
At least 1 G3+ heme toxicity		6 (46%)	17 (85%)		
No G3+ heme toxicities		7 (54%)	3 (15%)		
G4+ heme toxicities	33			>0.9	>0.9
At least 1 G4+ heme toxicity		2 (15%)	4 (20%)		
No G4+ heme toxicities		11 (85%)	16 (80%)		

Abbreviations: G3, grade 3; G4, grade 4. Bold/Italic indicates significant p-value.

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Supplemental Table 2. Correlations of baseline GA measures with baseline pro-inflammatory cytokine levels

GA measures	Cytokine levels	Spearman's Rho	p-value	q-value
TUG	TNF-alpha	0.65	5.92e-05	0.005
IADL	TNF-alpha	-0.48	8.49e-04	0.024
CARG score	IL-6	0.50	9.27e-04	0.024
% risk	TNF-alpha	0.43	1.44e-03	0.024
pKPS	IL-10	-0.48	1.68e-03	0.024
TUG	IL-2	0.54	1.78e-03	0.024
CARG score	TNF-alpha	0.45	1.90e-03	0.024
ADL	IL-10	-0.42	2.40e-03	0.027
pKPS	TNF-alpha	-0.46	2.68e-03	0.027
pKPS	IL-2	-0.58	3.36e-03	0.029
cKPS	IL-6	-0.45	3.91e-03	0.029
LDH (log level)	IL-10	0.40	4.27e-03	0.029
IADL	IL-10	-0.38	4.45e-03	0.029
LDH (log level)	TNF-alpha	0.42	4.58e-03	0.029
IADL	IL-2	-0.54	5.60e-03	0.033
cKPS	IL-2	-0.55	6.32e-03	0.033

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ADL	TNF-alpha	-0.36	6.56e-03	0.033
% risk	IL-6	0.41	6.73e-03	0.033
Activity limitation	IL-6	-0.60	6.90e-03	0.033
IADL	IL-6	-0.42	7.68e-03	0.034
cKPS	TNF-alpha	-0.33	8.00e-03	0.034
LDH (log level)	IL-2	0.48	9.30e-03	0.038
TUG	IL-10	0.41	1.05e-02	0.041
CARG score	IL-10	0.41	1.32e-02	0.049

Abbreviations: GA, geriatric assessment; LDH, lactate dehydrogenase; cKPS, clinician-rated karnofsky performance scale; pKPS, patient-rated karnofsky performance scale; ADL, activities of daily living; IADL, instrumental activities of daily living; TUG, timed-get-up and go; CARG, cancer and aging group; IFN, interferon; IL, interleukin; TNF, tumor necrosis factor. Bold indicates significant q-value.

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Supplemental Table 3. Baseline and change in GA measures by stage

GA measures	N	Stage I/II, N = 13 (median, IQR)	Stage III/IV, N = 20 (median, IQR)	p-value	q-value
Baseline LOG LDH	33	5.48 (5.26, 5.58)	5.77 (5.54, 6.02)	0.010	0.089
Baseline cKPS	33	90 (70, 90)	70 (60, 90)	0.11	0.2
Baseline pKPS	33	90 (80, 100)	80 (60, 90)	0.037	0.13
Baseline ADL	33	85 (45, 100)	42 (18, 80)	0.021	0.094
Baseline IADL	33	14.00 (13.00, 14.00)	12.00 (7.75, 14.00)	0.068	0.2
Baseline Activity limitation	33	56 (50, 62)	44 (41, 62)	0.6	0.6
Baseline TUG	31	9 (9, 11)	14 (11, 20)	0.009	0.089
Baseline CARG score	33	9.00 (7.00, 11.00)	10.50 (10.00, 13.25)	0.045	0.14
Baseline % risk	33	54 (54, 77)	54 (54, 89)	0.3	0.4
Δ in LOG LDH	30	0.00 (-0.20, 0.24)	0.01 (-0.25, 0.10)	0.3	0.4
Δ in cKPS	32	0 (0, 0)	10 (0, 15)	0.3	0.4
Δ in pKPS	32	0 (0, 10)	0 (-10, 10)	0.3	0.4
Δ in ADL	30	2 (-5, 6)	0 (-12, 10)	0.6	0.6
Δ in IADL	31	0.00 (-0.25, 0.00)	0.00 (-0.50, 0.50)	0.6	0.6
Δ in Activity limitation	31	0 (-12, 6)	0 (-17, 0)	0.2	0.3

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Δ in TUG	29	-1.0 (-1.2, 1.0)	-0.7 (-3.8, 1.0)	0.6	0.6
Δ in CARG score	32	0.00 (0.00, 2.00)	-1.00 (-3.00, 0.00)	0.016	0.094
Δ in % risk	32	0 (0, 0)	0 (-8, 0)	0.2	0.3

Abbreviations: IQR, interquartile range; LDH, lactate dehydrogenase; cKPS, clinician-rated karnofsky performance scale; pKPS, patient-rated karnofsky performance scale; ADL, activities of daily living; IADL, instrumental activities of daily living; TUG, timed-get-up and go; CARG, cancer and aging group. Bold/Italic indicates significant p-value.

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Supplemental Table 4. Baseline and change in cytokine levels by stage

Cytokine levels	N	Stage I/II, N = 13 (median, IQR)	Stage III/IV, N = 20 (median, IQR)	p-value	q-value
Baseline LOG IFN-gamma	32	1.81 (1.35, 2.31)	2.93 (2.14, 3.82)	0.010	0.2
Baseline LOG IL-10	32	-0.29 (-0.81, 1.41)	0.09 (-0.32, 2.65)	0.12	0.5
Baseline LOG IL-12	32	-1.81 (-2.39, -1.32)	-1.86 (-2.50, -1.36)	0.8	0.8
Baseline LOG IL-13	32	-1.11 (-4.41, -0.88)	-0.81 (-1.65, -0.37)	0.2	0.5
Baseline LOG IL-1-beta	32	-5.30 (-5.30, -5.30)	-5.30 (-5.30, -5.30)	0.6	0.7
Baseline LOG IL-2	31	-5.30 (-5.30, -1.95)	-2.43 (-4.51, -1.03)	0.057	0.4
Baseline LOG IL-4	32	-3.35 (-3.58, -3.04)	-3.15 (-4.17, -2.59)	0.8	0.8
Baseline LOG IL-6	32	0.73 (-0.32, 1.18)	1.07 (0.59, 1.81)	0.2	0.5
Baseline LOG IL-8	30	2.44 (2.20, 2.94)	2.81 (2.50, 3.05)	0.5	0.7
Baseline LOG TNF-alpha	32	0.87 (0.56, 1.69)	1.77 (1.53, 2.29)	0.020	0.2
Δ in LOG IFN-gamma	30	0.51 (-0.55, 1.30)	-0.88 (-1.72, 0.50)	0.2	0.5
Δ in LOG IL-10	30	-0.33 (-1.00, 0.10)	-0.72 (-2.19, 0.20)	0.6	0.7
Δ in LOG IL-12	30	-0.12 (-0.47, 0.11)	-0.03 (-0.30, 0.36)	0.4	0.7
Δ in LOG IL-13	30	0.62 (-1.09, 2.74)	-0.06 (-1.12, 0.49)	0.5	0.7
Δ in LOG IL-1-beta	30	0.03 (0.00, 2.74)	0.00 (0.00, 0.00)	0.11	0.5

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Δ in LOG IL-2	29	0.67 (-0.08, 1.92)	0.00 (-0.82, 1.28)	0.2	0.5
Δ in LOG IL-4	30	-0.19 (-0.52, 0.19)	-0.34 (-0.99, 0.10)	0.9	0.9
Δ in LOG IL-6	30	-0.22 (-1.29, 0.29)	-1.02 (-1.77, -0.35)	0.2	0.5
Δ in LOG IL-8	28	0.27 (-1.75, 1.69)	-0.38 (-0.75, 0.24)	0.5	0.7
Δ in LOG TNF-alpha	30	-1.18 (-1.55, 0.02)	-0.35 (-1.01, -0.03)	0.6	0.7

Abbreviations: IQR, interquartile range; IFN, interferon; IL, interleukin; TNF, tumor necrosis factor. Bold/Italic indicates significant p-value.

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Supplemental Table 5. Association of baseline and change in cytokine levels with toxicity events

Characteristic (IQR)	N	At least 1 toxic event, N = 22	No toxic events, N = 11	p-value	q-value
Baseline LOG IFN-GAMMA	32	2.42 (1.81, 3.39)	2.28 (1.43, 2.99)	0.3	0.7
Baseline LOG IL10	32	1.50 (-0.38, 2.29)	-0.29 (-0.87, 0.00)	0.012	0.12
Baseline LOG IL12	32	-1.58 (-2.35, -1.33)	-2.28 (-2.51, -1.74)	0.059	0.3
Baseline LOG IL13	32	-0.72 (-1.11, -0.34)	-3.47 (-5.30, -0.95)	0.006	0.12
Baseline LOG IL1-BETA	32	-5.30 (-5.30, -5.30)	-5.30 (-5.30, -5.30)	0.4	0.8
Baseline LOG IL2	31	-2.64 (-5.30, -1.52)	-4.07 (-5.30, -2.25)	0.2	0.7
Baseline LOG IL4	32	-3.32 (-3.54, -2.41)	-3.54 (-4.40, -3.08)	0.068	0.3
Baseline LOG IL6	32	1.17 (0.37, 1.81)	0.66 (-0.31, 1.02)	0.09	0.3
Baseline LOG IL8	30	2.80 (2.29, 2.95)	2.75 (2.44, 3.25)	0.6	0.8
Baseline LOG TNF-ALPHA	32	1.80 (0.87, 2.31)	1.45 (0.82, 1.65)	0.067	0.3
Δ in LOG IFN-GAMMA	30	-15 (-75, 77)	4 (-35, 32)	0.6	0.8
Δ in LOG IL10	30	-51 (-96, -1)	-39 (-792, 38)	0.9	0.9
Δ in LOG IL12	30	1 (-37, 17)	-10 (-15, 9)	0.5	0.8
Δ in LOG IL13	30	-24 (-214, 78)	0 (-235, 53)	0.7	0.8
Δ in LOG IL1-BETA	30	0 (0, 37)	0 (0, 1)	0.5	0.8
Δ in LOG IL2	29	20 (-32, 40)	0 (-9, 35)	0.3	0.7
Δ in LOG IL4	30	-9 (-27, -0)	-7 (-32, 9)	0.7	0.8
Δ in LOG IL6	30	-46 (-118, -12)	-55 (-173, 23)	0.6	0.8
Δ in LOG IL8	28	-13 (-30, 24)	-4 (-32, 47)	0.5	0.8
Δ in LOG TNF-ALPHA	30	-40 (-95, 18)	-34 (-56, -17)	0.8	0.8

Abbreviations: IQR, interquartile range; IFN, interferon; IL, interleukin; TNF, tumor necrosis factor.

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Figure Legend.

Supplemental Figure 1: Changes in individual pro-inflammatory cytokine levels (log-transformed) pre- and post-prephase therapy with q-value indicated. **Abbreviations:** IL, interleukin; TNF, tumor necrosis factor.

Supplemental Figure 1. Changes in cytokine levels from pre- to post-prephase therapy

