# THE LANCET Oncology

## Supplementary appendix

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Supplement to: Rajan A, Carter CA, Berman A, et al. Cixutumumab for patients with recurrent or refractory advanced thymic epithelial tumours: a multicentre, open-label, phase 2 trial. *Lancet Oncol* 2014; published online January 15. http://dx.doi.org/10.1016/S1470-2045(13)70596-5.

## Supplementary appendix

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#### Additional methods

#### **Patients**

Adequate organ and bone marrow function was defined as: leucocytes  $>3\times10^9$  cells per L, absolute neutrophil count  $>1\cdot5\times10^9$  cells per L, haemoglobin >90 g/L, platelets  $>100\times10^9$  cells per L, total bilirubin  $\le1\cdot5$  times institutional upper limit of normal (ULN), aspartate aminotransferase or alanine aminotransferase <3 times institutional ULN (5 times if liver function test elevations are due to liver metastases), and creatinine  $\le1\cdot5$  times institutional ULN (or creatinine clearance >1 mL per s/m² for patients with creatinine levels above institutional normal).

Patients with symptomatic brain metastases, fasting blood glucose  $\geq 6.7$  mmol/L or above the institutional ULN, major surgery, radiation therapy, chemotherapy, biological therapy, or hormonal therapy within 4 weeks of enrolling, and unresolved residual toxic effects related to previous treatment were ineligible for this study. Concurrent steroids for myasthenia gravis or other thymic epithelial tumour-related paraneoplastic conditions were permitted provided the dose was stable for  $\geq 8$  weeks prior to enrolment.

#### **Procedures**

Baseline evaluation consisted of a complete history and physical examination, tumour imaging with a CT scan of the chest, abdomen and pelvis and PET scan, and laboratory assessment including a complete blood count, prothrombin time, partial thromboplastin time, serum chemistries, lipid profile, amylase, lipase, HbA1c, IGF-1 levels, quantitative immunoglobulin assessment, urinalysis, and electrocardiogram.

Although pharmacokinetic analysis was proposed initially, it was not done after completion of the study since pharmacokinetic data were available for every 3-week dosing of cixutumumab by the time enrolment was completed to this trial.

#### Pharmacodynamic analysis

For multiparameter flow cytometric analysis, mononuclear cells were obtained by centrifugation and the cells were viably frozen until analysis. For assessment of interferon  $\gamma$  expression in CD4+ T cells, mononuclear cells were incubated with phorbol myristate acetate, ionomycin, and monensin for 4 h, stained for CD4 and interferon  $\gamma$ , and analysed in a MACSQuant flow cytometer (Miltenyi Biotec, Bergisch Gladbach, Germany). Peripheral blood populations were defined as central memory CD4 T cells (CD4+ CCR7+, CD27+, CD45RA-), naive CD4 T cells (CD4+ CCR7+, CD27+, CD45RA+), regulatory T cells (Tregs; CD4+CD25hiFoxp3+), and analysed in an LSR II (BD Biosciences). Circulating endothelial cells (CECs; CD45–, CD31+, CD146+ CD133–) and circulating endothelial progenitor cells (CEPs; CD45–, CD31+, CD146– CD133+) were analysed in a MACSQuant flow cytometer. All flow cytometric data were analysed using FlowJo software (Tree Star, Ashland, OR, USA).

#### Statistical analysis

The association between a variety of clinical and laboratory parameters and response and survival was determined using the Kaplan-Meier method and a log-rank test. As a total of 44 survival analyses were performed, p values for these analyses need to be interpreted in that context. For parameters obtained at multiple timepoints, the association with survival according to values at cycle 1, day 1 (CID1), cycle 2, day 1 (C2D1), and the difference between the two timepoints were all evaluated. Paired differences in total and free IGF-1, and in IGF-1R, between baseline and values at later timepoints were evaluated for significance using a Wilcoxon signed rank test.

Separately by histology, responses were divided into three categories as partial response, stable disease, and progressive disease, or into two response groups: partial response and no response. A Jonckheere-Terpstra test for trend was used to assess the significance of the trends in the changes in CEPs and CECs over the three response categories, whereas a Wilcoxon rank sum test was used to compare the changes in CEPs and CECs when only two response groups were compared.

All p values are two-tailed and presented without any adjustment for multiple comparisons.

	Ethnic origin	WHO subtype	Cycles (n)	New or worsening* autoimmune disease during treatment	Previous regimens of systemic treatment (n)	Previous radiation therapy	Previous surgery	Presence of extra- thoracic metastases	Time to response (months)
57-year-old man	White	В3	46	No	3	Yes	Yes	No	19.5
52-year-old man	White	B2	18	No	4	No	Yes	Yes	8.5
39-year-old woman	White	B1	6	Yes	3	Yes	No	Yes	3
42-year-old man	Black	AB	24	Yes	2	No	Yes	No	4.5
59-year-old woman	White	B2	15	Yes	2	No	Yes	No	7.5

\*Details of worsening autoimmune disease during treatment are provided in table 4. Supplemental Table 1: Details of partial responders

	Response or disease stabilisation	Duration of response or disease stabilisation (months)
Patient description		
57-year-old man, B3 subtype		
PAC	No	NA
Ifosfamide	Yes	4
Belinostat*	Yes	11
52-year-old man, B2 subtype		
Saracatinib*	No	NA
Belinostat*	No	NA
Paclitaxel	Yes	4
39-year-old woman, B1 subtype		
ADOC	Yes	35
Pemetrexed	Yes	9
Pemetrexed	Yes	7
42-year-old man, AB subtype		
Pemetrexed	Yes	23
59-year-old woman, B2 subtype		
ADOC	Yes	16
Carboplatin plus paclitaxel	Yes	4

NA=not applicable; PAC=cisplatin, doxorubicin, cyclophosphamide. ADOC=cisplatin, doxorubicin, cyclophosphamide, vincristine. \*Clinical trial. Supplemental Table 2: Previous systemic treatment for recurrent disease in patients responding to cixutumumab

	Patients (n)	Median (IQR) concentration	p (paired)*
Free IGF-1 in serum (nmol/L)			
Baseline, day 1	24	0.1 (0.02–0.11)	NA
Cycle 1, day 3	23	0.3 (0.22–0.41)	<0.0001
Cycle 1, day 7	22	0.5 (0.32–0.63)	<0.0001
Cycle 1, day 14	21	0.4 (0.25–0.57)	<0.0001
Total IGF-1 in serum (nmol/L)			
Baseline, day 1	24	7.4 (2.3–13.0)	NA
Cycle 1, day 3	23	13.0 (6.5–20.5)	0.0001
Cycle 1, day 7	22	16.1 (11.0–28.6)	<0.0001
Cycle 1, day 14	21	18.3 (10.3–29.0)	<0.0001
IGF-1R in PBMC (pmol/g)			
Baseline, day 1	13	8-1 (6-2–13-0)	NA
Cycle 1, day 3	13	6.3 (4.9–9.1)	0.006
Cycle 1, day 7	13	6.3 (3.9–8.6)	0.006
Cycle 2, day 1	13	5.0 (3.5–7.0)	0.001

NA=not applicable. PBMC=peripheral blood mononuclear cells. \*Wilcoxon signed rank test.

Supplemental Table 3: Concentrations of IGF-1 in serum and IGF-1R in PBMCs before and after treatment with cixutumumab

	Patients (n)	Median (IQR) amount of phospho-AKT in PBMCs (RU)	p (paired)*
Baseline, day 1	13	8.1 (5.9–12.3)	NA
Cycle 1, day 3	13	7.0 (6.3–10.4)	0.69
Cycle 1, day 7	13	8-4 (6-1–13-4)	0.60
Cycle 2, day 1	13	8.3 (5.9–14.1)	0.61

NA=not applicable. PBMCs=peripheral blood mononuclear cells. RU=relative unit. \*Student's t test.

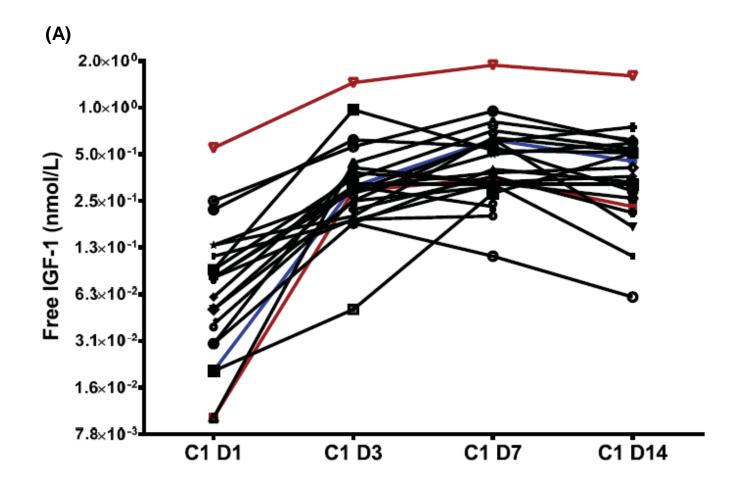
Supplemental Table 4: Concentrations of phospho-AKT, normalised against total protein in PBMCs before and after treatment with cixutumumab

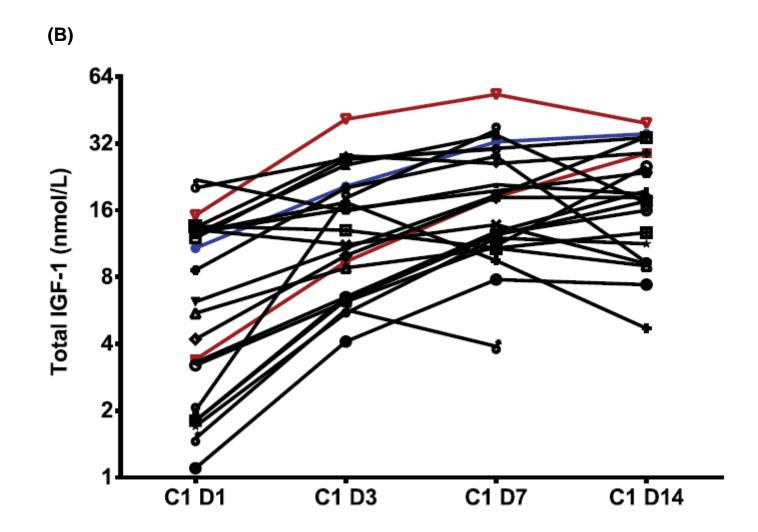
	Timepoint	Samples analysed (n)	Median (range) value	р
Interferon γ-expressing CD4+ T cells		••		0.0017
Partial response	C2D1 to C1D1	5	1·31 (0·32 to 2·04)	
Stable disease	C2D1 to C1D1	31	1·11 (0·42 to 2·21)	
Progressive disease	C2D1 to C1D1	11	0.93 (0.48 to 1.16)	
Circulating endothelial progenitors				0.049
Responders (thymoma)	C2D1 to C1D1	5	-8·13 (-30·97 to 119·28)	
Non-responders (thymoma)	C2D1 to C1D1	31	-63.29 (-1604.22  to  259.69)	
Circulating endothelial cells				0.29
Responders (thymoma)	C2D1 to C1D1	5	$-28 \cdot 11 \ (-180 \cdot 18 \text{ to } 12 \cdot 73)$	
Non-responders (thymoma)	C2D1 to C1D1	31	10.91 (-533.46 to 451.20)	
Anti-interferon α antibody				0.021
Responders	Pretreatment	5	609 (60·50 to 10 044)	
Non-responders	Pretreatment	30	15 356 (7 to 22 381)	
Anti-interleukin 22 antibody				0.0048
Responders	Pretreatment	5	100 (32 to 309)	
Non-responders	Pretreatment	30	516·75 (16 to 10 097)	
Anti-GMCSF antibody		••		0.014
Responders	Pretreatment	5	212·50 (43·50 to 526)	
Non-responders	Pretreatment	29	28 (19 to 247·50)	
Anti-interferon γ antibody				0.0091
New or worse autoimmune condition	Pretreatment	8	108 (46 to 405·50)	
No new or worse autoimmune condition	Pretreatment	38	49.75 (3 to 1414)	
Interferon γ-expressing CD4+ T cells				0.011
New or worse autoimmune condition	Pretreatment	8	4·80 (1·21 to 24·60)	
No new or worse autoimmune condition	Pretreatment	39	14·20 (1·40 to 48·60)	
Interleukin 17-expressing CD4+ T cells				0.033
New or worse autoimmune condition	Pretreatment	8	0·14 (0·06 to 1·15)	
No new or worse autoimmune condition	Pretreatment	39	0.46 (0.04 to 2.51)	
Interleukin 4-expressing CD4+ T cells				0.021
New or worse autoimmune condition	Pretreatment	8	0.55 (0.18 to 2.61)	
No new or worse autoimmune condition	Pretreatment	39	2·34 (0·14 to 17·30)	
Interleukin 17-expressing CD4+ T cells				0.0058
New or worse autoimmune condition	C2D1 to C1D1	8	0·13 (0·06 to 0·58)	
No new or worse autoimmune condition	C2D1 to C1D1	39	0.56 (0.04  to  3.13)	

Interferon γ-expressing CD4+ T cells				0.016
New or worse autoimmune condition	C2D1 to C1D1	8	4·34 (1·49 to 24·50)	
No new or worse autoimmune condition	C2D1 to C1D1	39	14·80 (1·62 to 54)	
Interleukin 4-expressing CD4+ T cells				0.043
New or worse autoimmune condition	C2D1 to C1D1	8	0.63 (0.23 to 4.74)	
No new or worse autoimmune condition	C2D1 to C1D1	39	2·38 (0·16 to 11·50)	

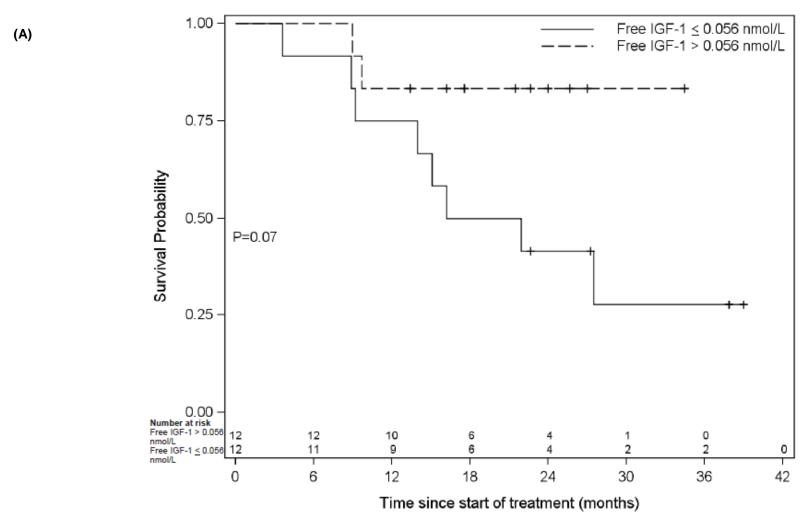
C2D1=cycle 2, day 1. C1D1=cycle 1, day 1. GMCSF=granulocyte macrophage colony-stimulating factor. Supplemental Table 5: Results of selected immune correlative studies

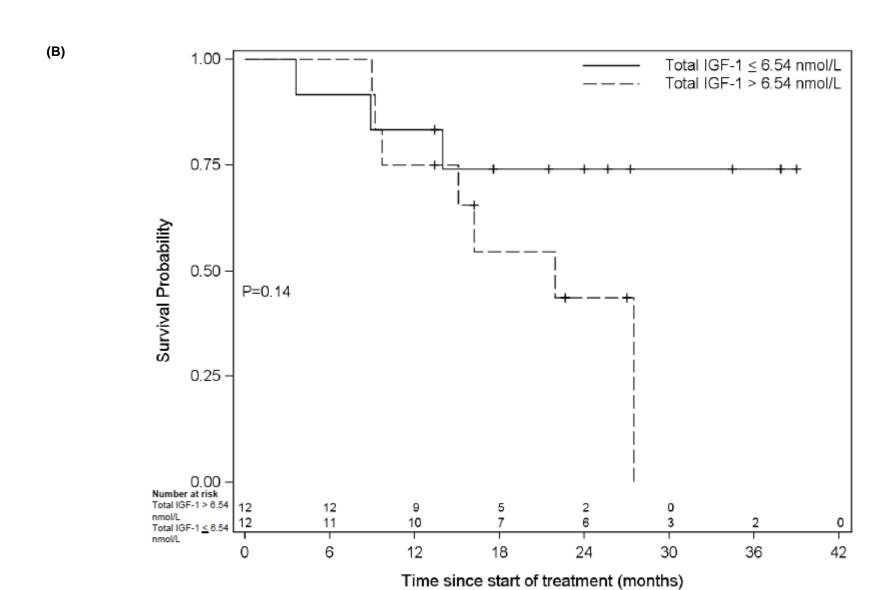
Supplemental Figure 1: Change in amounts of (A) free IGF-1 and (B) total IGF-1 after treatment with cixutumumab C1D1=cycle 1, day 1. C1D3=cycle 1, day 3. C1D7=cycle 1, day 7. C1D14=cycle 1, day 14.





Supplemental Figure 2: Kaplan-Meier curves for overall survival in patients with thymoma according to amounts of (A) free IGF-1 ( $\leq$ 0.056 nmol/L vs >0.056 nmol/L) and (B) total IGF-1 ( $\leq$ 6.54 nmol/L vs >6.54 nmol/L) before treatment





### Supplemental Figure 3: Association between amounts of IGF-1 and pre-existing or new or worsening autoimmune disease

- (A) Free IGF-1 and pre-existing autoimmune disease (p=0.51). (B) Free IGF-1 and new or worsening autoimmune disease (p=0.94).
- (C) Total IGF-1 and pre-existing autoimmune disease (p=0.68). (D) Total IGF-1 and new or worsening autoimmune disease (p=0.82).

