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Management of NK cell malignancies

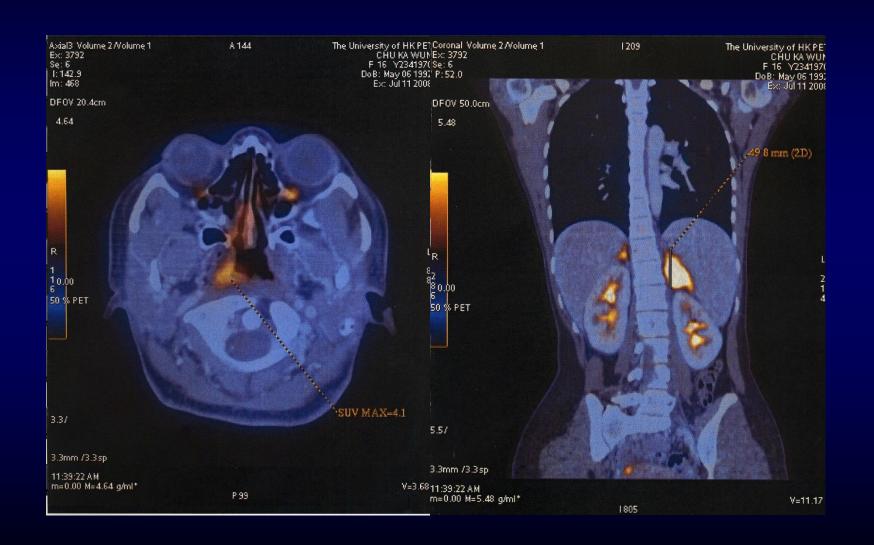
Principles

- 1. Accurate staging is needed for NK/T-cell at all anatomical locations
- 2. "Non-nasal" NK/T-cell lymphomas may have occult nasal involvement, and hence are disseminated "nasal" lymphomas
- 3. NK/T-cell lymphomas of all stages require chemotherapy
- 4. Radiotherapy alone should not be used

Cutaneous NK cell lymphoma, nasal type



Cutaneous NK/T cell lymphoma, nasal type Disseminated NK cell lymphoma



Management of NK cell malignancies

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L-asparaginase containing regimens

Table 1. Novel chemotherapeutic regimens for NK/T-cell lymphomas

Regimen	Protocol	Reference
AspaMetDex	Escherichia coli L-asparaginase: 6000 U/m² IM, days 2, 4, 6, and 8 Methotrexate: 3000 mg/m² IV, day 1 Dexamethasone: 40 mg orally, days 1-4	16
LVP	L-asparaginase: 6000 IU/m² IV, days 1-5 Vincristine: 1.4/m² IV, day 1 Prednisolone: 100 mg orally, days 1-5	59
GELOX	Gemcitabine: 1000 mg/m ² IV, days 1 and 8 E. coli L-asparaginase: 6000 units/m ² IM, days 1-7 Oxaliplatin: 130 mg/m ² IV, day 1	60
SMILE	Dexamethasone: 40 mg IV or orally, days 2-4 Methotrexate: 2000 mg/m² IV, day 1 Ifosfamide: 1500 mg/m² IV, days 2-4 E. coli L-asparaginase: 6000 U/m² IV, days 8, 10, 12, 14, 16, 18, and 20 Etoposide: 100 mg/m² IV, days 2-4	62

IM, intramuscularly.

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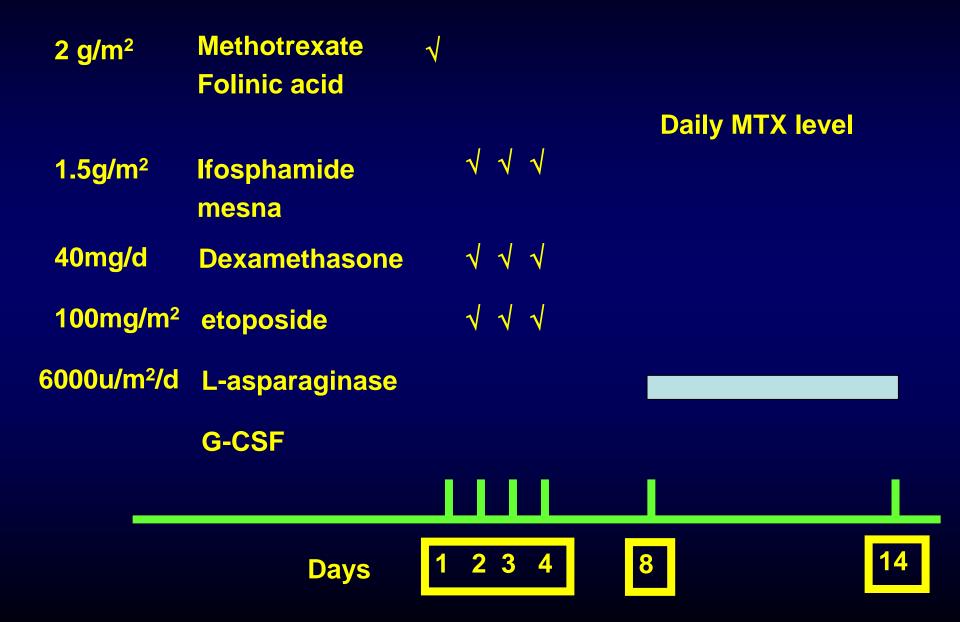
IM, intramuscularly.

Rationale for the SMILE protocol

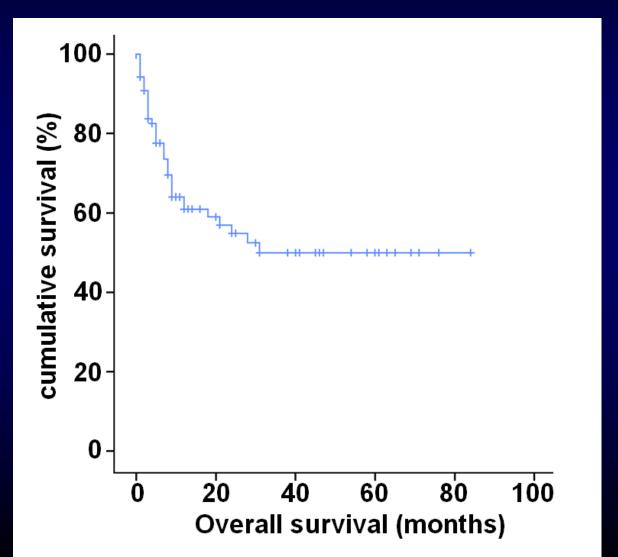
SMILE

- 1. Non-MDR dependent drugs: dexamethasone, ifosfamide, methotrexate
- 2. L-asparaginase: effective as a single agent
- 3. Etoposide (MDR-dependent): effective for CAEBV and other EBV related lymphoproliferation

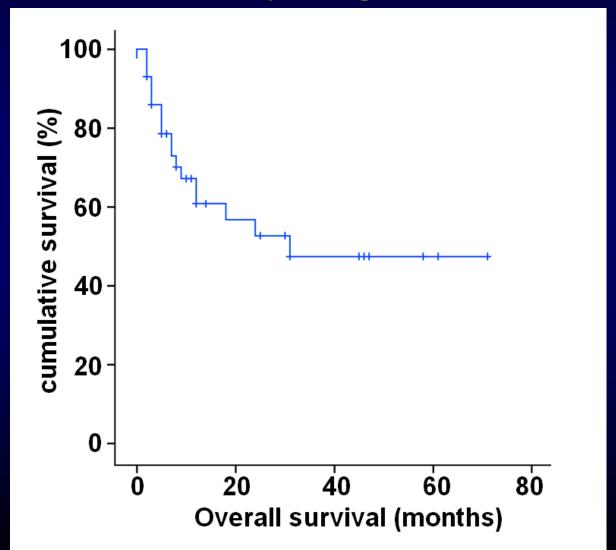
SMILE protocol for NK cell lymphomas (K. Oshimi)



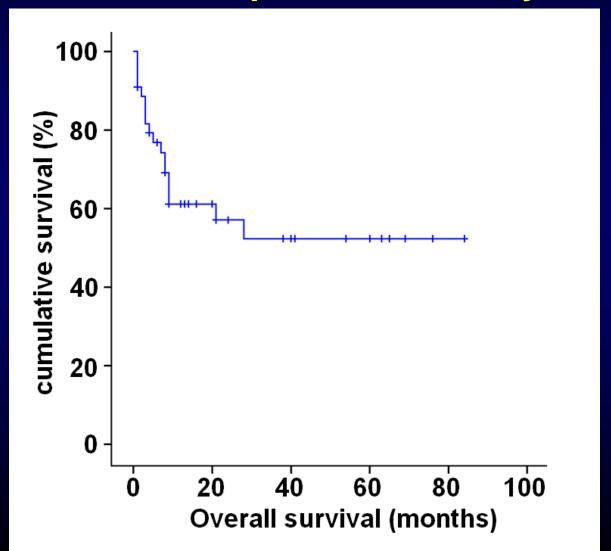
Overall survival



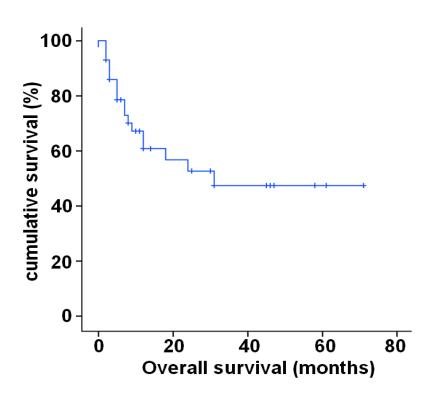
Overall survival: newly-diagnosed



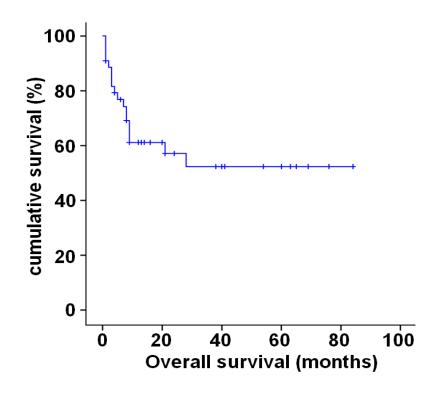
Overall survival: relapsed / refractory



Similar OS for newly-diagnosed and relapsed / refractory cases

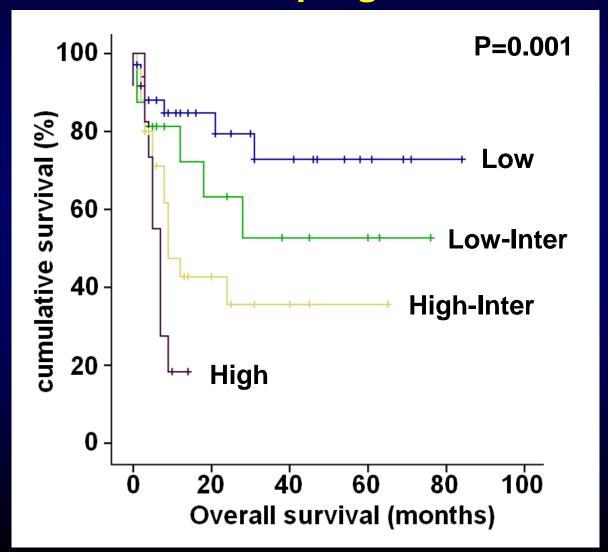


Newly diagnosed



Relapsed / refractory

Overall survival: IPI for prognostication



Non MDR related chemotherapy

Promace-CytaBOM (III)
DHAP (I)

SMILE (II)



Non MDR related chemotherapy

Promace-CytaBOM (III) + RT

SMILE (II)



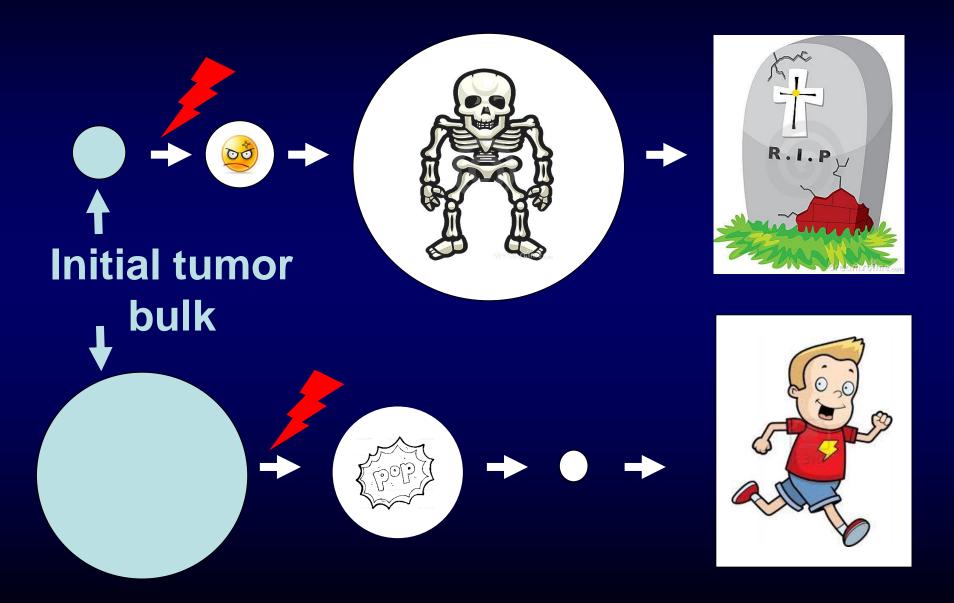
Prognostication of NK/T-cell lymphoma treated by SMILE

Current prognostic models rely on constitutional features of the patients and lymphoma load

Hypothesis

Sensitivity of lymphoma cells to chemotherapy is a more important indicator of outcome

Lymphoma sensitivity to treatment



NK/T-cell lymphoma Monitoring of response to SMILE

- 1. EBV DNA
- 2. PET/CT scan

NK/T-cell lymphoma Monitoring of response to SMILE

1. EBV DNA

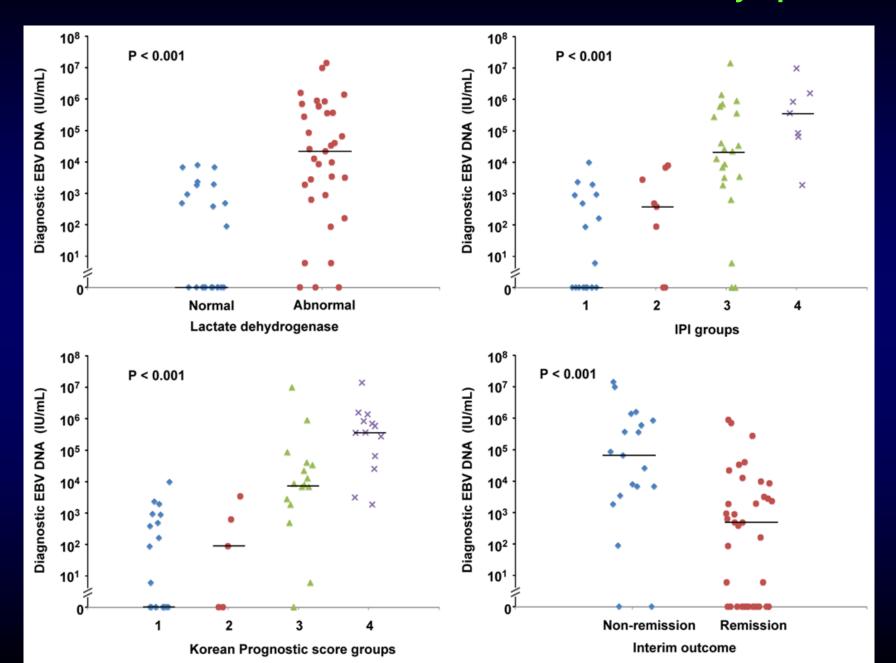
2. PET/CT scan

Prognostic factors for SMILE

Prospective evaluation of

- 1. Presentation EBV DNA
- 2. Non-detectable EBV DNA after SMILE
- 3. Dynamic patterns
 - A. persistent non-detectable
 - B. detectable, but < presentation
 - C. detectable, but > presentation

Presentation EBV DNA for SMILE treated NK/T-cell lymphoma



Presentation plasma EBV DNA

- 1. Correlated very well with conventional parameters of tumor load, including LDH, IPI, KIPI, and remission
- 2. Did not correlate with overall survival or disease free survival
- 3. Implications: EBV DNA reflects tumor load, but tumor load is not the only factor that impacts on survival

Dynamic changes of EBV DNA

1. Reduction of EBV DNA after SMILE

- 2. Levels of EBV DNA during SMILE therapy
 - A. non-detectable
 - B. detectable, but < presentation
 - C. detectable, but > presentation

Dynamic changes reflect tumor sensitivity to SMILE therapy

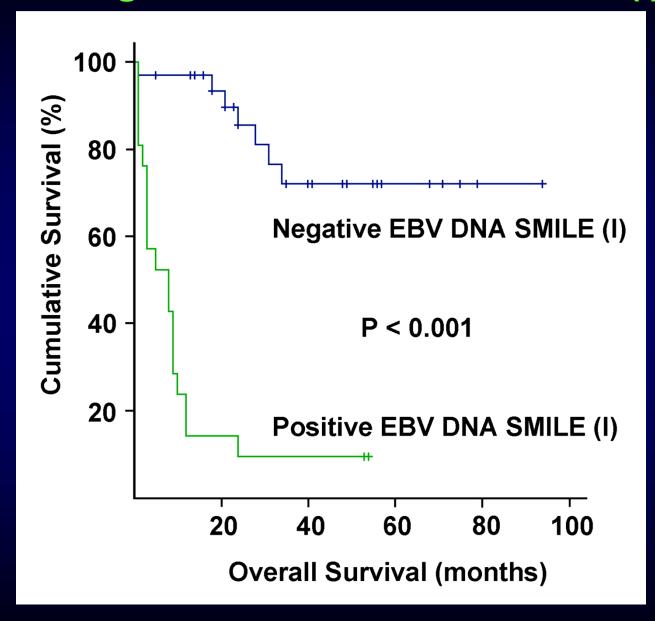
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Impact of negative EBV DNA after SMILE (I) on OS



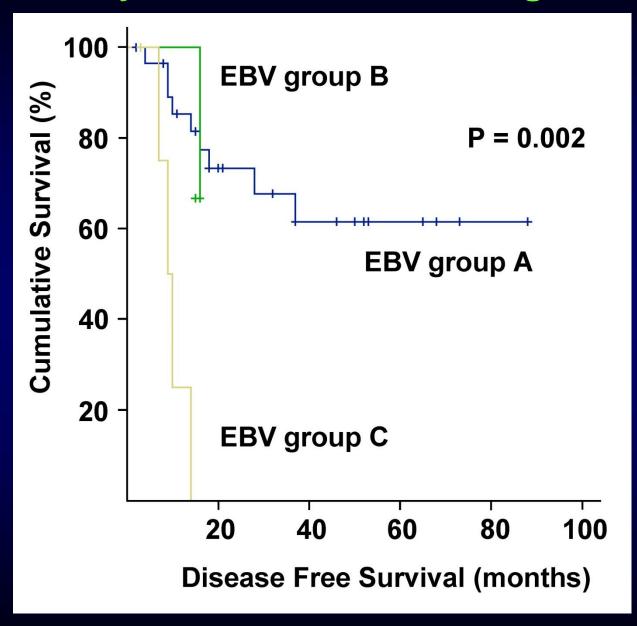
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Dynamic changes reflect tumor sensitivity to SMILE therapy

Impact of dynamic EBV DNA changes on DFS



Prognostic factors for SMILE treated NK/T-cell lymphoma

Table 4. Multivariate analysis of prognostic factors for survivals after SMILE therapy						
Significant factors	P value	Hazard ratio	95% confidence interval			
Whole cohort						
Overall survival						
Albumin Negative EBV DNA after SMILE (I)	0.005 < 0.001	3.589 12.883	1.458 - 8.831 4.759 - 34.876			
Disease free survival						
Age Dynamic EBV change grouping	0.015 0.002	1.078 4.072	1.014 – 1.145 1.676 – 9.890			
Patients with quantifiable EBV DNA at presentation						
Overall survival						
Albumin Negative EBV DNA after SMILE (I)	0.002 < 0.001	5.764 19.887	1.896 – 17.516 5.331 – 74.104			
Disease free survival						
Age Dynamic EBV change grouping	0.047 0.036	1.067 2.849	1.001 - 1.137 $1.070 - 7.587$			

Conclusions

- 1. Presentation EBV DNA reflects tumor load, but does not impact on survivals
- 2. Negative EBV DNA after SMILE (I) reflects superior response to chemotherapy, and therefore impacts on OS
- 3. For patients already in CR, non-detectable EBV DNA means optimal suppression of tumor cells, and therefore impacts on DFS

NK/T-cell lymphoma Monitoring of response to SMILE

- EBV DNA
- 2. PET/CT scan

Role of interim PET/CT scan in SMILE

Design

- 1. Baseline PET/CT
- 2. Interim PET/CT after 2 3 cycles
- 3. End-treatment PET/CT

Parameters
SUVmax, ∆SUVmax, tumor volume,
5-point Deauville score

Role of interim PET/CT in SMILE therapy

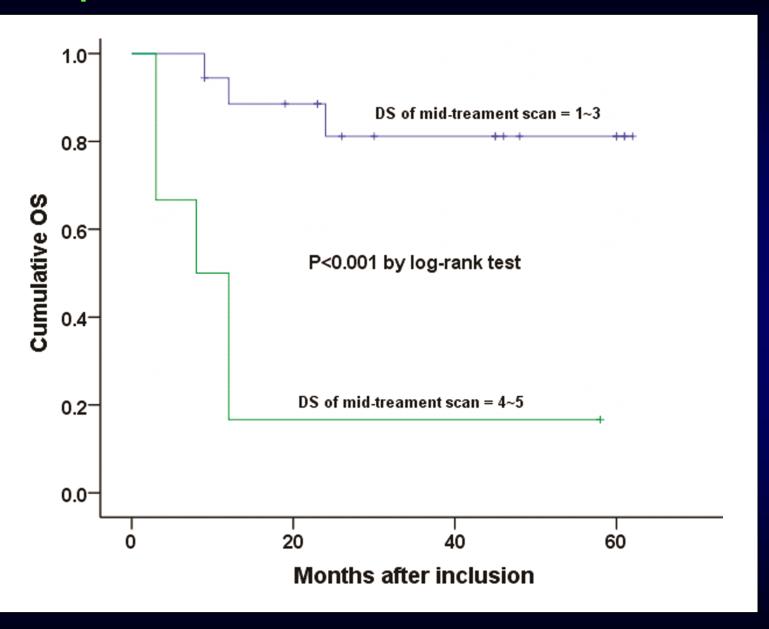
Multivariate analysis OS

Deauville 5-point score P < 0.001

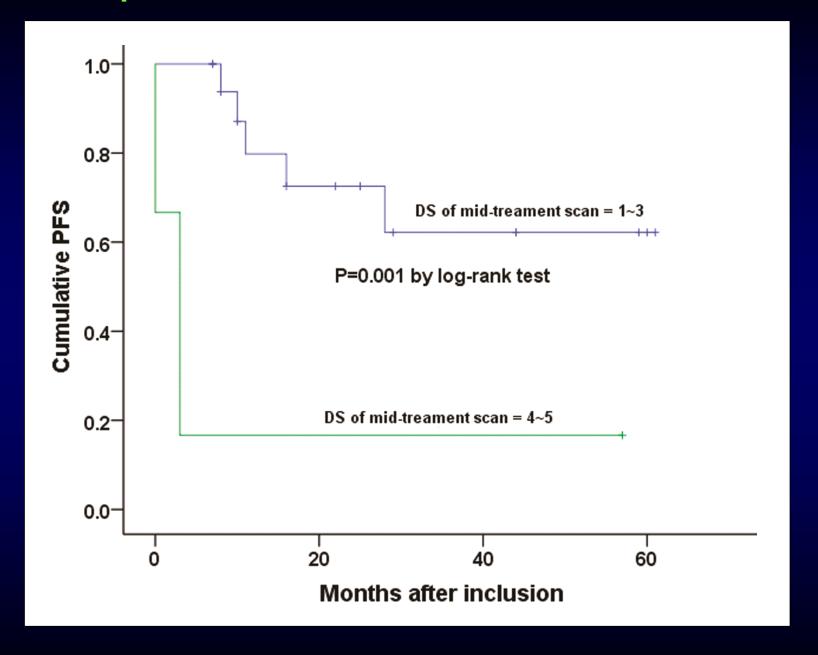
Multivariate analysis PFS

Deauville 5-point score P < 0.001

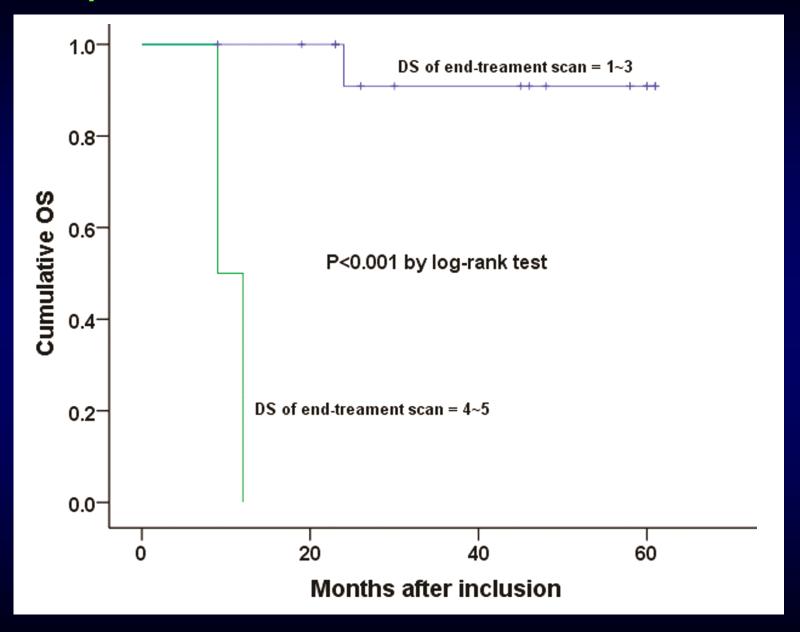
Impact of mid-treatment PET/CT on OS



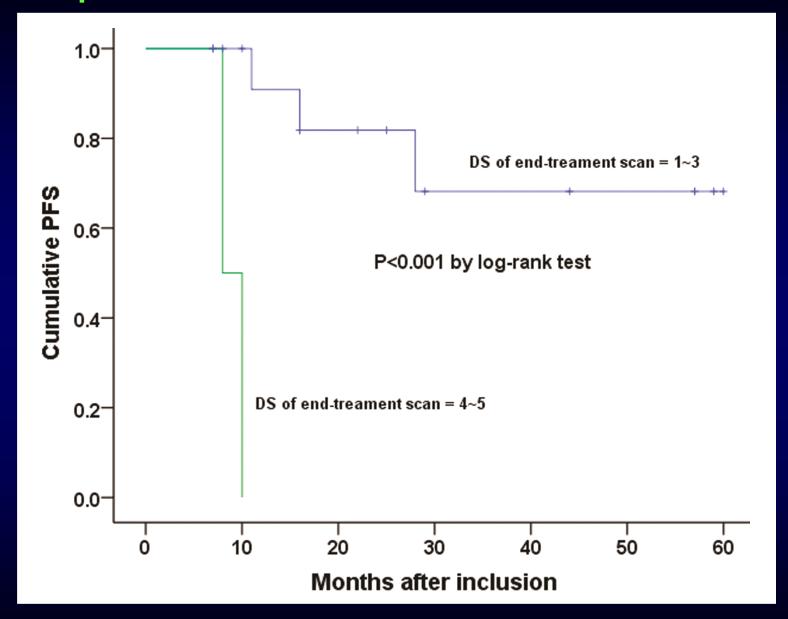
Impact of mid-treatment PET/CT on PFS



Impact of end-of-treatment PET/CT on OS



Impact of end-of-treatment PET/CT on PFS



Role of interim PET/CT in SMILE therapy

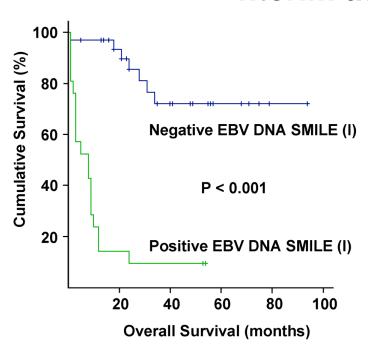
Should aim at achieving a negative PET/CT scan after 2 – 3 cycles of SMILE

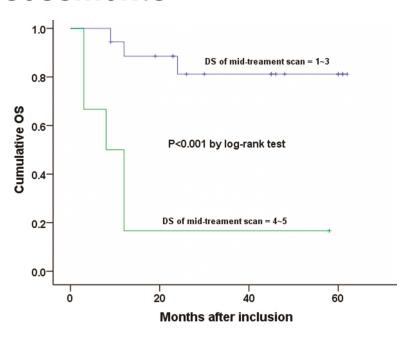
Patients with a Deauville score of 4 – 5 at interim scan will not do well, and may require additional therapy to achieve durable remission

Which is more prognostic

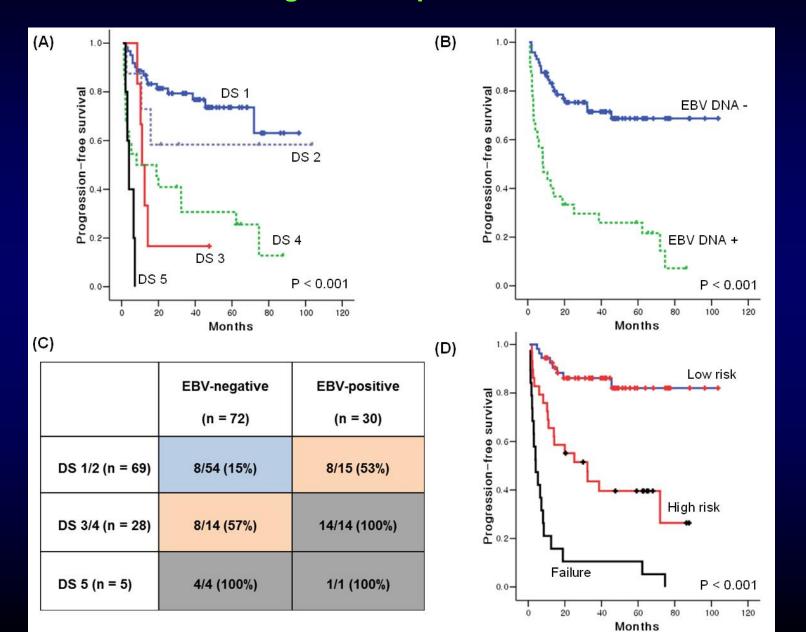
EBV DNA or PET/CT scan

Interim assessments

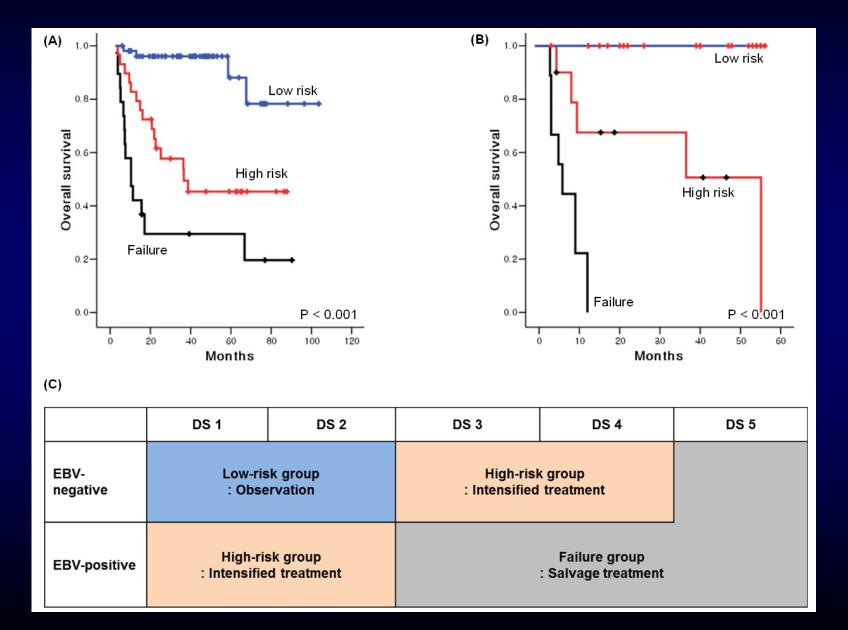




Combined EBV DNA and PET/CT at end of treatment Prognostic impact



Combined EBV DNA and PET/CT at end of treatment Prognostic impact



Incorporation of presentation EBV DNA in prognostication

Age
Stage
Nasal / non-nasal
Nodal involvement
EBV DNA (detectable / non-detectable)

Other L-asparaginase containing regimens

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