# Evidence of cure for extranodal nasal-type natural killer/T-cell lymphoma with current treatment: an analysis of the CLCG database

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# Supplementary data

### **Methods**

#### Definitions of primary site and primary tumor invasion

Primary site is defined according to primary symptoms and radiologic findings. As described in detail in our previous study, primary disease can be classified by the anatomic site of origin, including the nasal cavity, paranasal sinuses, nasopharynx, oropharynx, hypopharynx, larynx, trachea, oral cavity (gingiva, buccal mucosa, mouth floor) and, more rarely, skin and soft tissues, stomach, small intestine, colon, lung, testis, central nervous system, orbit, or other sites. Accordingly, ENKTCL can be further divided into three distinct clinical subgroups: (i) nasal, (ii) non-nasal-UADT ((i) and (ii) combined are represented as the upper aerodigestive tract [UADT]), and (iii) extra-UADT. Nasal ENKTCL refers to lesions in the nasal cavity and paranasal sinuses. Non-nasal-UADT ENKTCL refers to lesions in the nasopharynx, oropharynx (tonsil, base of tongue, oropharyngeal wall), hypopharynx, oral cavity, larynx, and trachea. Extra-UADT ENKTCL (often termed as "non-nasal" or "extranasal" in the literature) is rare, most commonly reported in cutaneous and soft tissues, gastrointestinal tract, testis, lung, central nervous system, and orbit. In patients with localized disease, it is usually easy to define primary site. In patients with disseminated disease, primary site is usually defined as the extranodal site (e.g., UADT, cutaneous and soft tissues, gastrointestinal tract, testis, lung, central nervous system, orbit, or other sites) according to the first or major clinical symptoms/presentations. <sup>1</sup> If the UADT was involved, patients were designated as nasal or non-nasal UADT ENKTCL (UADT ENKTCL), irrespective of whether other extranodal organs were involved.<sup>2</sup>

Primary tumor invasion (PTI) is defined as the growth of the primary tumor beyond the boundary

of the original extranodal site into a neighboring structure or tissue or as contiguous multisite involvement, regardless of stage. Accordingly, PTI for nasal origin was defined as nasal disease extending to any of the following structures: maxillary sinus, ethmoid sinus, sphenoid sinus, facial skin, orbit, nasopharynx, hard palate or oral cavity. PTI for extra-nasal UADT origin was defined as involvement of multiple sites (nasopharynx, oropharynx, tonsil, tongue base, oral cavity, or larynx) or adjacent structures or tissues (such as the nasal cavity). PTI of extra-UADT origin was defined as primary disease directly extended to adjacent structures or tissues (such as gastric or intestinal lesions to adjacent organs; skin and soft tissue lesions to neurovascular structure or bone; testicular lesions to skin).

#### **Treatment**

A total of 1593 patients received chemotherapy. The chemotherapy regimens were summarized in Supplementary Table 1. Of them, 1296 (81.4%) patients received asparaginase-based regimens. The most commonly used asparaginase-based regimens were CHOP/CHOP-like plus ASP (cyclophosphamide, doxorubicin, vincristine, prednisolone, and L-asparaginase or pegaspargase), followed by PGEMOX (pegaspargase plus gemcitabine and oxaliplatin), GELOX (L-asparaginase plus gemcitabine and oxaliplatin) and LVP (L-asparaginase, vincristine, and prednisolone) regimens. The remaining 297 (18.6%) patients received other non-anthracycline-based regimens, mainly with GDP (gemcitabine, dexamethasone, cisplatin) and DICE (etoposide, cyclophosphamide, cisplatin, dexamethasone) regimens.

Radiotherapy was given according to the principles of involved-site radiation therapy (ISRT) with the primary tumor and adjacent sites encompassed. The median RT dose was 50 Gy.

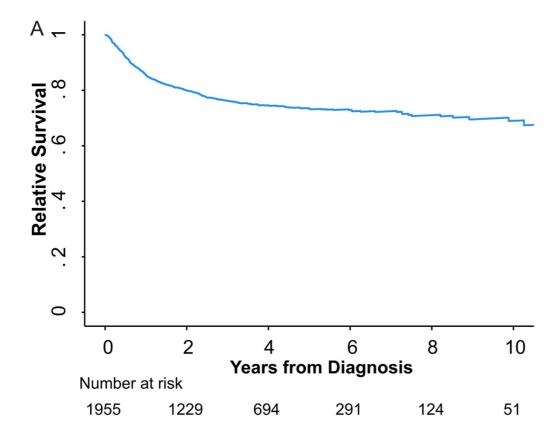
#### Statistical Methods

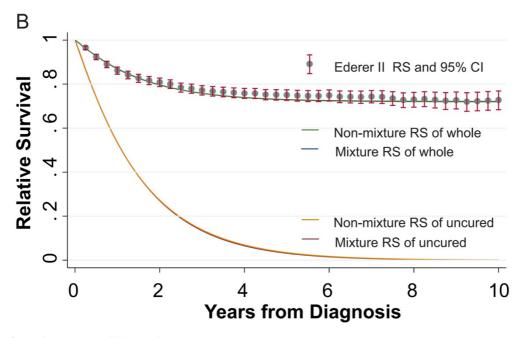
#### Outcome measure

The population mortality files by age and calendar year for male (Supplementary Table 2) and female (Supplementary Table 3) in China were used to calculate the expected survival in an age-, sex-, and calendar year-matched general Chinese population.

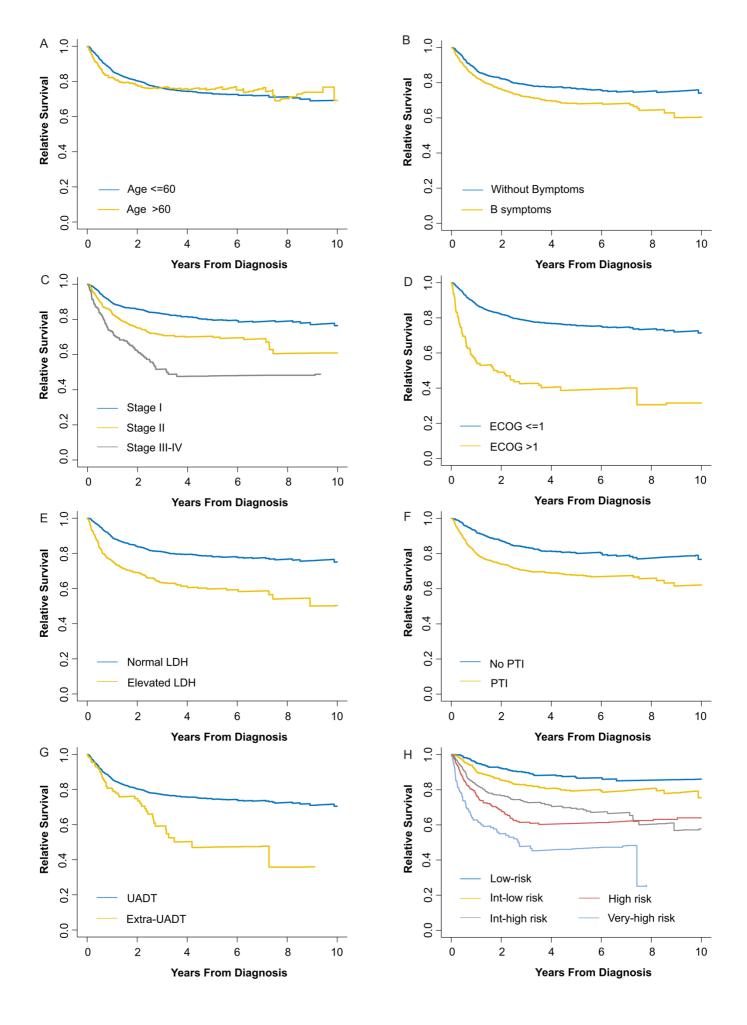
#### Cure model

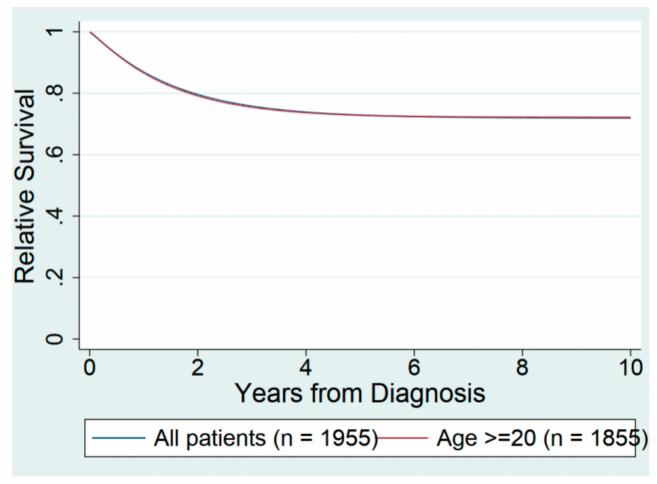
For sensitivity analysis, we fitted mixture and non-mixture cure models. Both models showed good fit on visual examination and gave reasonable estimates of cure fractions, as shown by the closeness of estimated cure fractions to the plateau level of the Ederer II RS curve (Supplementary Fig. 1); however, the non-mixture cure model had a slightly lower Akaike information criterion (Supplementary Table 4).<sup>3</sup> Time to cure was defined as the time between diagnosis and the moment at which excess mortality reaches zero, i.e. the population is considered statistically cured.<sup>4</sup> As Chauvenet et al. proposed, time-to-cure can be estimated as the time at which "almost" all uncured patients would have died.<sup>5</sup> In this study, we estimated cure time as the time at which 95% of the "uncured" patients would have died. This threshold of 95% was considered clinically relevant. Thus, among all the patients who were expected to die from cancer, cure occurred at the moment when 95 patients of 100 had died. From that time, excess mortality attributed to ENKTCL becomes statistically negligible. We restricted follow-up to 10 years, because the number of at-risk patients beyond 10 years was limited, and their inclusion might have led to unreliable estimates.





Supplementary Figure 1





Supplementary Fig. 3

#### Figure legends

Supplementary Fig. 1. **RS** and model fit of non-mixture and mixture cure model. (A) RS estimated using the Ederer II method. (B) Model fit of non-mixture and mixture cure model. Figure 1B shows the Ederer II estimates of RS with 95% confidence intervals, predicted RS curves of the whole cohort from the non-mixture cure model (green line) and mixture cure model (blue line), and predicted RS curves of the uncured patients from the non-mixture cure model (orange line) and mixture cure model (red line). Compared with the Ederer II RS estimates, the predicted RS curves from non-mixture and mixture models fit well and almost overlap. RS, relative survival.

Supplementary Fig. 2. **RS curves by prognostic factors**. RS curves stratified by age (A), B symptoms (B), stage (C), ECOG score (D), LDH (E), PTI (F), primary UADT disease site (G), and NRI-defined risk group (H). RS, relative survival; ECOG, Eastern Cooperative Oncology Group; PTI, primary tumor invasion; LDH, lactate dehydrogenase; UADT, upper aerodigestive tract; NRI, nomogram-revised risk index. Int-low, intermediate-low; Int-high, intermediate-high.

Supplementary Fig. 3. Sensitivity analysis to assess the potential influence of different eligibility criteria. Predicted relative survival curves of the whole cohort of patients with ENKTCL (n = 1955, bule line) and the cohort that excluded patients younger than 20 years old (n = 1855, red line). As is shown, the relative survival curves for the two cohorts almost overlap.

**Supplementary Table 1.** Summary of non-anthracycline-based regimen chemotherapy in 1593 patients who received chemotherapy

| Regimens                    | No. (%)     | Definition and agents  |  |
|-----------------------------|-------------|--|--|
| Asparaginase-based regimens | 1296 (81.4) |  |  |
| CHOP/CHOP-like plus ASP     | 425 (26.7)  | Cyclophosphamide, doxorubicin, vincristine, prednisolone, L-asparaginase or pegaspargase |  |
| PGEMOX                      | 170 (10.7)  | Pegaspargase, gemcitabine, oxaliplatin   |  |
| GELOX                       | 157 (9.9)   | Gemcitabine, oxaliplatin, L-asparaginase   |  |
| LVP (LOD)                   | 153 (9.6)   | L-asparaginase, vincristine, prednisone  |  |
| GDPL                        | 60 (3.8)    | Gemcitabine, dexamethasone, cisplatin, L-asparaginase                                    |  |
| Gem + ASP                   | 53 (3.3)    | Gemcitabine, pegaspargase  |  |
| GAD-M                       | 49 (3.1)    | Gemcitabine, pegaspargase, dexamethasone, methotrexate                                   |  |
| SMILE                       | 29 (1.8)    | L-asparaginase, ifosfamide, methotrexate, etoposide, dexamethasone                       |  |
| AspaMetDex                  | 29 (1.8)    | L-asparaginase, methotrexate, dexamethasone  |  |
| SVILE                       | 28 (1.8)    | Ifosfamide, dexamethasone, pegaspargase, vindesine, and etoposide                        |  |
| DDGP                        | 24 (1.5)    | Dexamethasone, cisplatin, gemcitabine, and pegaspargase                                  |  |
| MESA                        | 22 (1.4)    | Methotrexate, etoposide, dexamethasone, pegaspargase                                     |  |
| IPGDP                       | 21 (1.3)    | Ifosfamide, pegaspargase, gemcitabine, dexamethasone, cisplatin                          |  |
| VIDL                        | 15 (0.9)    | Etoposide, ifosfamide, dexamethasone, L-asparaginase                                     |  |
| Asp alone                   | 9 (0.6)     | L-asparaginase or pegaspargase   |  |
| VDLP                        | 7 (0.4)     | Vincristine, daunorubicin, L-asparaginase, prednisolone                                  |  |
| GDLE                        | 4 (0.3)     | Gemcitabine, dexamethasone, L-asparaginase, etoposide                                    |  |
| DICE-L                      | 3 (0.2)     | Etoposide, cyclophosphamide, cisplatin, dexamethasone, L-asparaginase                    |  |
| IMEP-L                      | 1 (0.1)     | Ifosfamide, methotrexate, etoposide, prednisolone, L-asparaginase                        |  |
| ASP-other                   | 37 (2.3)    | Asparaginase with other combination of cytotoxic drugs                                   |  |
| PLA/Gem/Other regimens      | 297 (18.6)  |  |  |
| GDP                         | 98 (6.2)    | Gemcitabine, dexamethasone, cisplatin  |  |
| DICE                        | 57 (3.6)    | Etoposide, cyclophosphamide, cisplatin, dexamethasone                                    |  |
| ATT                         | 49 (3.1)    | Three regimens combination with ASHAP (doxorubicin, cisplatin, ara-C, solu-medrol), m-   |  |
|                             |             | BACOS (methotrexate, leucovorin, doxorubicin, oncovin, bleomycin, cytoxan, solu-         |  |
|                             |             | medrol), and MINE (mesna, ifosfamide, novantrone, etoposide)                             |  |
| VIDP                        | 17 (1.1)    | Etoposide, ifosfamide, cisplatin, and dexamethasone                                      |  |
| GEMOX                       | 15 (0.9)    | Gemcitabine, oxaliplatin   |  |
| GP                          | 7 (0.4)     | Gemcitabine, cisplatin   |  |
| DIMG                        | 7 (0.4)     | Dexamethasone, ifosfamide, methotrexate, gemcitabine                                     |  |
| DEVIC                       | 4 (0.3)     | Dexamethasone, etoposide, ifosfamide, carboplatin  |  |
| IMEP                        | 2 (0.1)     | Ifosfamide, methotrexate, etoposide, prednisolone  |  |
| Other                       | 41 (2.6)    | Other combination of cytotoxic drugs without asparaginase                                |  |

Abbreviations: ASP, asparaginase (L-asparaginase or pegaspargase); GEM, gemcitabine; MTX, methotrexate; PLA, platinum.

**Supplementary Table 2.** Mortality by age and calendar year for male in China (uploaded as a separated file).

**Supplementary Table 3.** Mortality by age and calendar year for female in China (uploaded as a separated file).

**Supplementary Table 4.** The exploratory model fit analyses of mixture and non-mixture cure model with Weibull distribution for ENKTCL

|  | Mixture Weibull  | Non-mixture Weibull |
|--|------------------|---------------------|
| Cure fraction (%, 95% CI)                    | 72.1 (69.5-74.6) | 71.9 (69.3-74.5)    |
| Median survival time of uncured              | 1.10 (0.96-1.26) | 1.10 (0.96-1.26)    |
| Survival time of 95% probability of cure     | 2.99 (2.43-3.69) | 3.04 (2.46-3.78)    |
| Survival time of 90% probability of cure     | 1.92 (1.56-2.37) | 1.94 (1.56-2.42)    |
| Time to cure (95%)                           | 4.38 (3.61-5.31) | 4.48 (3.66-5.49)    |
| Time to cure (90%)                           | 3.42 (2.86-4.09) | 3.48 (2.89-4.19)    |
| AIC  | 3216.88          | 3214.31             |
| BIC  | 3233.62          | 3231.05             |
| Probability of cure given survival of 5 year | 98.8 (97.3-99.5) | 98.7 (97.1-99.4)    |

ENKTCL, extranodal nasal-type NK/T-cell lymphoma; CI, confidence interval; AIC, Akaike information criterion; BIC, Bayesian information criterion. Time to cure (95%) was estimated as the time at which 95% of the "uncured" patients would have died. Time to cure (90%) was estimated as the time at which 90% of the "uncured" patients would have died.

**Supplementary Table 5.** Median survival time of uncured patients and time to cure by clinical characteristics and risk stratification for extranodal nasal-type NK/T-cell lymphoma

| Variable       | n (%)       | Median survival time of uncured | Time to cure, years (95% CI) |
|----------------|-------------|---------------------------------|------------------------------|
|                | n (%)       | patients, years (95% CI)        | (93% CI)                     |
| Sex            | 1201 (70 () | 1.00 (0.01.1.27)                | 4 44 (2 40 5 (4)             |
| Male           | 1381 (70.6) | 1.08 (0.91-1.27)                | 4.44 (3.49-5.64)             |
| Female         | 574 (29.4)  | 1.16 (0.93-1.45)                | 4.22 (3.12-5.71)             |
| Stage          | 1100 (55.4) | 1 22 (1 25 1 (2)                | 5.00 ( <b>2.5</b> 5 ( 00)    |
| I              | 1123 (57.4) | 1.32 (1.07-1.63)                | 5.08 (3.75-6.88)             |
| II             | 599 (30.6)  | 0.97 (78.3-1.19)                | 3.65 (2.65-5.02)             |
| III-IV         | 233 (11.9)  | 1.01 (0.73-1.40)                | 4.60 (2.77-7.63)             |
| Elevated LDH   |             |                                 |                              |
| No             | 1422 (72.7) | 1.23 (1.06-1.42)                | 3.98 (3.24-4.87)             |
| Yes            | 533 (27.3)  | 0.89 (0.70-1.13)                | 4.49 (3.22-6.27)             |
| Age            |             |                                 |                              |
| ≤60            | 1667 (85.3) | 1.17 (1.02-1.33)                | 4.34 (3.63-5.33)             |
| >60            | 288 (14.7)  | 0.58 (0.39-0.86)                | 2.36 (1.29-4.34)             |
| B symptoms     |             |                                 |                              |
| No             | 1184 (60.6) | 1.12 (0.94-1.33)                | 4.09 (3.19-5.25)             |
| Yes            | 771 (39.4)  | 1.08 (0.86-1.35)                | 4.95 (3.51-6.99)             |
| ECOG score     |             |                                 |                              |
| 0-1            | 1827 (93.5) | 1.21 (1.05-1.39)                | 4.44 (3.64-5.42)             |
| ≥2             | 128 (6.5)   | 0.58 (0.41-0.81)                | 2.87 (1.88-4.39)             |
| PTI            | , ,         |                                 |                              |
| No             | 868 (44.4)  | 1.62 (1.30-2.02)                | 5.19 (3.80-7.10)             |
| Yes            | 1087 (55.6) | 0.88 (0.76-1.02)                | 3.62 (2.95-4.45)             |
| Primary site   | ,           | , ,                             | ,                            |
| UADT           | 1829 (93.6) | 1.05 (0.92-1.21)                | 4.12 (3.40-4.98)             |
| Extra-UADT     | 126 (6.4)   | 2.13 (1.07-4.24)                | 9.11 (3.38-24.6)             |
| NRI            | (0)         | (,                              | 2711 (0.00 = 1.0)            |
| Low risk       | 438 (22.4)  | 1.63 (1.17-2.27)                | 4.45 (2.78-7.13)             |
| Int-low risk   | 564 (28.8)  | 1.43 (1.13-1.82)                | 4.89 (3.16-6.38)             |
| Int-high risk  | 517 (26.4)  | 1.23 (0.91-1.66)                | 5.57 (3.59-8.62)             |
| High risk      | 277 (14.2)  | 0.84 (0.67-1.06)                | 2.95 (2.21-3.95)             |
| Very high risk | 159 (8.1)   | 0.61 (0.43-0.88)                | 3.34 (1.95-5.73)             |

ECOG, Eastern Cooperative Oncology Group; Int, intermediate; LDH, lactate dehydrogenase; NRI, nomogram-revised risk index; UADT, upper aerodigestive tract; PTI, primary tumor invasion. Int-low, intermediate-low; Int-high, intermediate-high. Time to cure was estimated as the time at which 95% of the "uncured" patients would have died.

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