



# Baseline Amide Proton Transfer Imaging at 3T Fails to Predict Early Response to Induction Chemotherapy in Nasopharyngeal Carcinoma

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**Background:** Early identification of nasopharyngeal carcinoma (NPC) patients with high risk of failure to induction chemotherapy (IC) would facilitate prompt individualized treatment decisions and thus reduce toxicity and improve overall survival rate. This study aims to investigate the value of amide proton transfer (APT) imaging in predicting short-term response of NPC to IC and its potential correlation with well-established prognosis-related clinical characteristics.

**Methods and Materials:** A total of 80 pathologically confirmed NPC patients receiving pre-treatment APT imaging at 3T were retrospectively enrolled. Using asymmetry analysis, APT maps were calculated with mean (APT<sub>mean</sub>), 90<sup>th</sup> percentile (APT<sub>90</sub>) of APT signals in manually segmented NPC measured. APT values were compared among groups with different histopathological subtypes, clinical stages (namely, T, M, N, and overall stages), EBV-related indices (EBV-DNA), or responses to induction chemotherapy, using Mann-Whitney U test or Kruskal–Wallis H test.

**Results:** NPC showed significantly higher APT<sub>mean</sub> than normal nasopharyngeal tissues (1.81  $\pm$  0.62% vs.1.32  $\pm$  0.56%, *P* <0.001). APT signals showed no significant difference between undifferentiated and differentiated NPC subtypes groups, different EBV-DNA groups, or among T, N, M stages and overall clinical stages of II, III, IVA and IVB (all *P* >0.05). Similarly, baseline APT-related parameters did not differ significantly among

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different treatment response groups after IC, no matter if evaluated with RECIST criteria or sum volumetric regression ratio (SVRR) (all P > 0.05).

**Conclusion:** NPC showed significantly stronger APT effect than normal nasopharyngeal tissue, facilitating NPC lesion detection and early identification. However, stationary baseline APT values exhibited no significant correlation with histologic subtypes, clinical stages and EBV-related indices, and showed limited value to predict short-term treatment response to IC.

Keywords: amide proton transfer (APT), nasopharyngeal carcinoma (NPC), Epstein–Barr virus (EBV), clinical stage, induction chemotherapy

#### INTRODUCTION

The number of the newly diagnosed nasopharyngeal carcinoma (NPC) patients has been reported to be over 129,000 worldwide in 2018, among which 70% is aggregated in East Asia and Southeast Asia (1). In particular, the incidence is as high as 9.69/100,000 in Southeastern China (2). Currently, induction chemotherapy (IC) followed by concurrent chemoradiotherapy (CCRT) has become the standard treatment for locoregionally NPC for its better prognosis than CCRT alone (3, 4). However, around 10% patients still do not respond well to IC. Identification of those non-responders in advance would facilitate more prompt personalized treatment decisions, avoiding chemotherapy toxicity and unnecessary costs. In addition, tumor response to IC has been identified as an independent prognostic factor for long-term survival (5, 6). Hence, pretreatment prediction of short-term response to IC is essential for better individualized management of NPC patients.

Due to substantial biological heterogeneity, NPCs with the same clinical stage might have completely different responses to IC. Identification of biomarkers that could reflect tumor microenvironment characteristics may benefit the prediction of treatment response. With the advancement of morphological and function imaging techniques, MRI has been widely used for NPC evaluation by providing information at both macroscopic and microscopic levels (7-10). Recently, APT-weighted (APT<sub>w</sub>) imaging has emerged as a powerful tool to probe the chemical exchange saturation transfer (CEST) effect between endogenous proteins/peptide amide protons and bulk water (11). Because of overexpressed proteins in tumors, APT<sub>w</sub> imaging has been increasingly used in tumor detection, grading, and treatment response evaluation of various types of cancer (12–26). Although APT<sub>w</sub> imaging has shown promising efficacy in predicting middleterm outcome after complete standard treatment (induction

chemotherapy followed by concurrent chemoradiotherapy) (26) or 2-year long-term outcome prediction in NPC patients (27), little is known about whether APT signal could predict short-term treatment response to IC only.

Besides, NPC can be classified into distinct subtypes by wellestablished prognosis-related characteristics, e.g. Epstein–Barr virus (EBV) infection status (28), different histopathological subtypes (29) and clinical stages, etc. It remains unknown whether these distinct NPC subtypes with different prognosis have different protein metabolism levels, which potentially could be quantified by  $APT_w$  imaging. Therefore, if there is any association between baseline APT measurements and wellestablished prognosis-related clinical characteristics of NPC, such as histopathological subtypes, clinical stages, and status of Epstein–Barr (EB) virus infection, the value of APT in predicting prognosis would be further strengthened.

In this study, we aimed to investigate the potential value of  $APT_w$  imaging in predicting short-term response to IC and its potential association with prognosis-related histopathological subtypes, clinical stages, and EBV-related indices.

#### MATERIALS AND METHODS

#### **Patient Enrollment**

This retrospective study was approved by the local ethics committee of the Cancer Hospital & Shenzhen Hospital, Chinese Academy of Medical Sciences, and informed consent was waived. From December 2019 to August 2021, 114 patients with suspected NPC who had undergone baseline APT-weighed imaging were enrolled for further selection. The inclusion and exclusion criteria were as follows: 1) biopsy confirmed NPC; 2) the maximal diameter of the primary nasopharyngeal lesion was larger than 1 cm; 3) conventional MRI and APT-weighted imaging were performed before treatment; 4) each patient had received at least one examination for clinical staging work-up, such as chest X-ray, abdominal ultrasonography, chest and abdomen CT, bone SPECT scan, or whole-body PET-CT scan. Clinical staging for each NPC patient was performed based on the 8th edition of the American Joint Committee on Cancer (AJCC) (30); APT-weighted images were reviewed by a boardcertified radiologist (ZL with more than 10 years of experience in head and neck imaging) and those with degraded image quality,

**Abbreviations:** NPC, nasopharyngeal carcinoma; APT, amide proton transfer; CEST, chemical exchange saturation transfer; EBV, Epstein–Barr virus; AJCC, American Joint Committee on Cancer; WASSR, water saturation shift referencing; APT<sub>mean</sub>, Mean value of APT asymmetry; APT<sub>90</sub>, 90th percentile of APT asymmetry; RECIST, Response Evaluation Criteria in Solid Tumors; SVRR, sum volumetric regression ratio; EBER, EBV-encoded small RNA; ISH, *in situ* hybridization; EBV-DNA, Epstein–Barr deoxyribonucleic acid; EA-IgA, early antigen antibody immunoglobulin A; VCA-IgA, viral capsid antibody immunoglobulin A; MTR, magnetization transfer ratio.

namely, overt signal lost, motion artifacts, geometric distortion, were excluded from analysis.

# **Treatment Strategy**

All NPC patients received standard treatment. For NPC patients with clinical stages of I and II, concurrent chemoradiotherapy or radiotherapy alone was given, whereas IC followed by concurrent chemoradiotherapy were prescribed to patients with clinical stages of III and IV. The brief treatment procedure was: 1) IC (GP regimen): Gemcitabine  $(1 \text{ g/m}^2, \text{ Days } 1 \text{ and } 8) + \text{cisplatin}$  $(75 \text{ mg/m}^2, \text{ Day 1});$  or Paclitaxel  $(135-175 \text{ mg/m}^2, \text{ Day 1}) +$ cisplatin (75 mg/m<sup>2</sup>, Day 1), 3 weeks per cycle  $\times$  2–3 cycles; 2) Concurrent chemoradiotherapy: cisplatin (30 mg/m<sup>2</sup>, Day 1, once a week for 7 times, iv) and concurrent radiation therapy (in a total of 6.6-7.6 weeks, accumulated dose of 66-78 Gy for primary lesion, while 60-70 Gy for regional lymph nodes). With intensity-modulated radiation therapy being used, conventional 6MV-X fractionated radiotherapy (200 cGy each time × 1 time per day  $\times$  5 days per week) was performed, which covers the primary NPC lesion and bilateral cervical lymphatics at risk.

# Follow-Up and Study Endpoint

In this study, the endpoint was set between after IC completion and before next-step concurrent chemoradiotherapy. Standard clinical MRI follow-up was performed at the endpoint for each subject.

# **MRI Study**

MRI study was performed on a 3T MR scanner (Discovery MR 750w, General Electric) using a 32-channel head and neck joint coil. Local shimming was placed to cover the whole nasopharyngeal carcinoma to reduce static magnetic field inhomogeneity. Referencing to T2-weighted images, a singleslice axial plane covering the largest primary nasopharyngeal carcinoma was selected for APT imaging with chemical shiftselective fat suppression and single-shot fast spin echo (FSE) readout (4 RF saturation pulses with duration of 500 ms for each pulse and without interpulse delay, RF saturation power B<sub>1</sub> of 2  $\mu$ T, TR = 6.5 s with minimum TE, slice thickness = 4 mm, field of view =  $22 \times 22$  cm<sup>2</sup>, matrix =  $128 \times 128$ ). In additional to an unsaturated scan, 27 CEST-weighted images with frequency offsets of 0,  $\pm$  0.5,  $\pm$  1.0,  $\pm$  1.5,  $\pm$  2.0,  $\pm$  2.5,  $\pm$  3.0,  $\pm$  3.5,  $\pm$  $4.5, \pm 5.0, \pm 5.5, \pm 6.0, \pm 6.5$ , and  $\pm 7.0$  ppm were acquired. The total acquisition time is approximately 4.5 min. A water saturation shift referencing (WASSR) map was collected with  $B_1$  of 0.5  $\mu$ T (frequency offsets between ±240 Hz with intervals of 48 Hz) to correct B<sub>0</sub> inhomogeneity.

# **Data Analysis**

APT data was analyzed in MATLAB (MathWorks). Z-spectra (S) was normalized by the unsaturated scan (S<sub>0</sub>), interpolated by smoothing splines, and corrected the field inhomogeneity with WASSR (31, 32). Pixel-wise APT-weighted value was calculated using asymmetry analysis of APT<sub>w</sub> (APT-weighted) =  $S(-3.5 \text{ ppm})/S_0 - S(3.5 \text{ ppm})/S_0$ .

Region of interest (ROI) of primary NPC was manually delineated on the unsaturated image by a radiologist (LZ with

6 years of experience in head and neck imaging) using opensource ITK-SNAP software (http://itksnap.org) with referencing conventional anatomical images, namely, T1-weighted, T2weighted, and contrast enhanced T1-weighted images. Overt hemorrhage, air space, necrosis and vessels were excluded. For patients with tumor confined in one side of nasopharynx without involving the midline of posterior wall of nasopharynx, the normal nasopharyngeal tissue was delineated as well. ROI delineation was then reviewed and approved by a senior radiologist (ZL with more than 10 years of experience in head and neck imaging). APT<sub>w</sub> signals, namely, the average APT (APT<sub>mean</sub>) and 90<sup>th</sup> percentile (APT<sub>90</sub>), were obtained from each ROI.

# **Treatment Response Evaluation**

The largest diameter of NPC was manually measured on the slice with the largest area of lesion on axial T<sub>1</sub>-weighted contrast enhanced images by a board-certified radiologist (LZ with 6 years of experience in head and neck imaging) on the vendor workstation. The measurement was reviewed by a senior radiologist (ZL), with any discrepancies resolved through consensus discussion. Then, the treatment response of primary tumor was determined by calculating the relative change percentage of tumor maximal diameter between pre- and posttreatment, as  $\Delta D = (D_{pre-treatment} - D_{post-treatment})/D_{pre-treatment}$ and classified as complete response (CR,  $\Delta D = 100\%$ ), partial response (PR,  $\Delta D \ge 30\%$ ), stable disease (SD,  $-20\% < \Delta D < 30\%$ ), and progression disease (PD,  $\Delta D \le -20\%$ ) according to the RECIST criteria version 1.1 (33).

Tumor volume was measured by manually delineating and summing NPC lesion from consecutive 2D slices on contrastenhanced  $T_1$ -weighted image by LZ and then reviewed by a senior radiologist (ZL). The sum volumetric regression ratio (SVRR) was defined as the tumor volume reduction percentage before and after treatment, and 50% reduction in SVRR was arbitrarily chosen as the cut-off value.

# Histopathology and Epstein–Barr (EB) Virus-Related Indices

The following pathological information were collected: 1) histology of nasopharyngeal carcinoma based on WHO classification, namely, nonkeratinizing squamous cell carcinoma (differentiated subtype and undifferentiated subtype), keratinizing squamous cell carcinoma and basaloid squamous cell carcinoma (34); 2) EBV-encoded small RNA (EBER) status using *in situ* hybridization (ISH). Serum test reports before treatment were also examined to obtain the following information: 1) copy numbers of viral Epstein–Barr deoxyribonucleic acid (EBV-DNA) in plasma samples before treatment, whereas copy numbers of 400 as threshold; 2) serum level of early antigen antibody (EA-IgA); and 3) serum level of viral capsid antibody (VCA-IgA).

# **Statistical Analysis**

All the statistical analyses were performed using SPSS software v. 26.0 (IBM, Armonk, NY). Paired Student's t-test was performed to evaluate the difference in  $APT_w$  signals between normal and

cancerous nasopharyngeal tissues. Continuous variables were displayed as median (25th–75th percentile). Mann–Whitney U test was used between the two histopathological subtypes. Kruskal–Wallis H test was conducted to assess any significant differences among groups with different clinical stages or treatment responses. Receiver operating characteristic (ROC) analysis was performed and the area under the curve (AUC) was calculated. The optimal cutoff values were determined by maximizing the sum of sensitivity and specificity using YOUDEN index. Pearson's correlation analysis was used to evaluate the relationship between APT<sub>w</sub> measurements and tumor regression rate.

# RESULTS

A total of 80 patients were included in the analysis (**Figure 1**) with clinical characteristics summarized in **Table 1**.

#### **Tissue Characterization**

A total of 17 patients showed NPC confined in one side and not involving the midline of nasopharynx. **Figures 2A, B** illustrate APT<sub>w</sub> maps and histograms in NPC and normal nasopharyngeal tissue regions in a 66-year-old male patient. The APT<sub>mean</sub> of the NPC were significantly higher than that of corresponding normal nasopharyngeal tissue ( $1.81 \pm 0.62\% vs.1.32 \pm 0.56\%$ , *P* <0.001) (**Figure 2C**). However, APT<sub>90</sub> showed no significant difference between NPC and corresponding normal nasopharyngeal tissue ( $3.68 \pm 0.84\% vs.3.25 \pm 1.07\%$ , *P* = 0.115). The optimal cutoff value of APT<sub>mean</sub> in discriminating NPC with contralateral nasopharyngeal tissue was 1.66% with a sensitivity of 0.765, a specificity of 0.765, and an AUC of 0.727 (**Figure 2D**).

Among the enrolled 80 patients, 66 had undifferentiated carcinoma, 14 had differentiated carcinoma, and no patients had nonkeratinizing squamous cell carcinoma. All  $APT_w$  signals

did not differ significantly between undifferentiated and differentiated NPC groups (all P > 0.05) (**Figure 3**), as summarized in **Table 2**.

#### **TNM Stage**

No significant difference was found in  $APT_w$  signals among primary tumor stages of T1 to T4, among lymph node stages of N1 to N3 (**Figure 4**), between M0 and M1 stages, among overall clinical stages of II, III, IVA, and IVB (all *P* >0.05) (**Table 2**).

#### **EBV-Related Indices**

The 67 patients with available EBER status all showed positive EBER on the specimen using *in situ* hybridization in all cases. For the 76 cases with available serum EBV-DNA quantification, APT<sub>w</sub> signals was found to be comparable between EBV-DNA positive group and EBV-DNA negative group (**Figure 5**). Similarly, there was no significant difference in APT<sub>w</sub> signals between EA-IgA positive group (N = 18) and EA-IgA negative group (N = 16), and between VCA-IgA positive group (N = 27) and VCA-IgA negative group (N = 7) (all *P* >0.05) for the 34 NPC patients with available serum expression of EBV antibody (**Table 3**).

#### **Treatment Response**

Among 66 patients with available treatment response evaluation after overall treatment, 53 patients had received induction chemotherapy (**Table 4**). For patients receiving overall treatment, no significant correlation was found neither between APT<sub>mean</sub> and SVRR ( $R^2 = -0.014$ , P = 0.738) (**Figure 6A**) nor between APT<sub>90</sub> and SVRR ( $R^2 = -0.001$ , P =0.339) (**Figure 6B**). Similarly, for patients who received induction chemotherapy, SVRR did not significantly correlate with neither APT<sub>mean</sub> ( $R^2 = -0.019$ , P = 0.883) (**Figure 6C**), nor APT<sub>90</sub> ( $R^2 = 0.008$ , P = 0.238) (**Figure 6D**). Baseline APT-related parameters did not differ significantly among different treatment



**TABLE 1** | Clinical characteristics of the enrolled 80 patients.

Clinical characteristics	Values/No. of patients
Age (years)	41.5
	(34-51.25)
Gender	
Male	55
remaie	25
Histology	0
Nonkeratinizing squamous cell carcinoma	0
Undifferentiated carcinoma	66
Differentiated carcinoma	14
Primary tumor stage	10
11	16
12	8
13	41
14	15
Nodal stage	
NO	1
N1	25
N2	25
N3	29
Metastatic stage	
MO	66
M1	11
Mx	3
Clinical stage	
1	0
ll	5
III	33
IVA	27
IVB	12
Not available	3
Treatment strategy	
Radiotherapy only	1
Concurrent chemoradiotherapy	15
Induction chemotherapy and concurrent chemoradiotherapy	58
Treatment strategy not available	6
MRI scan	
Baseline MR scan	80
Follow-up MR scan after induction chemotherapy	53
Follow-up MR scan after radiotherapy only	1
Follow-up MR scan after concurrent	12
chemoradiotherapy	
No follow-up MB scan	14

response groups after overall treatment or after induction chemotherapy, no matter evaluated with the RECIST criteria or SVRR (all P > 0.05). Besides, no significant difference in tumor regression rate was found between EBV-DNA (+) (n = 16) and EBV-DNA (-) (n = 37) group who received induction chemotherapy, as evaluated with the RECIST criteria (52.0 (38.3–57.2)% vs.58.0 (35.0–100.0)%, P = 0.407) and SVRR (67.0 (52.5–88)% vs. 79.0 (56.0–100.0)%, P = 0.441), respectively.

### DISCUSSION

In this study, we investigated the value of pre-treatment  $APT_w$  signal in predicting short-term response after IC in NPC patients and its potential association with prognosis-related clinical

characteristics, namely, histopathological subtypes, clinical stages, and EBV-related indices. Unfortunately, based on current result, the stationary baseline APT-weighted imaging showed limited value to predict response to IC and no significant correlation with different histopathological subtypes, EBV infection statuses and clinical stages.

The nasopharynx APT MRI is challenging due to susceptibility artifacts, and swallowing and jaw motion (22, 35). Our image quality was visually comparable to those shown in previous APT studies (22, 35) without apparent signal loss, geometric distortion, or any artifacts. In this study, significantly higher mean APT value in NPC tissue was observed than that in normal nasopharyngeal tissue, which could facilitate NPC lesion detection and early identification. The stronger APT effect may reflect rapidly proliferating tumor cells and abnormally active proteo-synthesis in cancerous tissues, as postulated previously (36).

Although APT-weighted imaging has been proven to be useful in differentiating head and neck malignant and benign tumors (22, 23), it was reported to be difficult in distinguishing malignant tumor types in head and neck regions (23). Our study showed that  $APT_w$  signals failed to discriminate between undifferentiated and differentiated NPC histology subtypes, which may imply similar protein synthesis status of these two histology subtypes before treatment. As for clinical stages, no significant difference of  $APT_w$  signals was shown among T, N, and M stages, and overall clinical stages.

EB-virus infection was recognized as the most common etiology of NPC, especially for nasopharyngeal non-keratinizing squamous cell carcinoma in endemic area (37). Consistently, all cases with available EBER in situ hybridization were tested EBER positive in our study. The serum EBV-DNA copy number has already been widely used to screen population with the high risk of developing NPC in endemic areas (38) and may serve as a biomarker to monitor and predict treatment response (39). However, pretreatment EBV-DNA status does not show apparent relationship with tumor regression rate after IC in this study. In contrast, a multi-center phase II randomized controlled study showed that an early decrease of EBV-DNA copy number indicates better treatment response (40). This might imply that monitoring the dynamic change of EBV-infection status throughout the treatment course using EBV-DNA copy number might be more effective than merely detecting a stationary pretreatment status in predicting treatment response. However, EBV-DNA copy number quantification data were not available after treatment in this retrospective study. Besides, although it seems no significant correlation of APT signals with EBV-related indices was presented, the correlation between APT<sub>w</sub> signals change and EBV-DNA copy number alteration should be investigated in the future.

This study focused on the investigation of the value of APT imaging in predicting short-term survival of NPC patients. Although Qamar et al. (26, 27) demonstrated that the mean APT value change between pretreatment and intra-treatment 2 weeks later indeed could predict the middle-term outcome six months after the completion of standard treatment, both pretreatment and intra-treatment mean APT value did not



FIGURE 2 | Representative ROI delineation on unsaturated S0 images, APT<sub>w</sub> maps, and histograms in NPC (A) and normal nasopharyngeal tissue (B) regions in a 66-year-old male patient. Comparison of APT<sub>w</sub> signals, including APT<sub>mean</sub>, APT<sub>90</sub> (C) between NPC and corresponding normal nasopharynx tissues; ROC analyses of APT<sub>mean</sub> in discriminating NPC and normal nasopharynx tissue (D).



#### TABLE 2 | APT-related parameters in different histology and TNM stage groups.

	No.	APT <sub>mean</sub> (%)	APT <sub>90</sub> (%)
Nonkeratinizing squamous cell carcinoma	0	_	_
Undifferentiated carcinoma	66	1.72 (1.36-2.10)	3.54 (3.05-4.04)
Differentiated carcinoma	14	1.77 (1.45-1.99)	3.20 (3.14–3.77)
P-value	_	0.810	0.695
Primary tumor stage			
T1	16	1.81 (1.60-2.07)	3.70 (3.46–4.11)
T2	8	2.01 (1.72-2.42)	3.67 (3.23-4.52)
ТЗ	41	1.64 (1.32-1.96)	3.39 (2.86-3.74)
T4	15	1.80 (1.31-2.26)	3.22 (3.03-4.16)
P-value	_	0.330	0.103
Nodal stage			
NO	1	-	_
N1	25	1.76 (1.48–2.00)	3.44 (3.14-4.09)
N2	25	1.86 (1.48-2.14)	3.59 (3.06-4.28)
N3	29	1.72 (1.32-2.00)	3.45 (3.07-3.82)
P-value	_	0.644	0.867
Metastatic stage			
MO	66	1.74 (1.48-2.01)	3.42 (3.05-4.03)
M1	11	1.53 (1.02-2.21)	3.74 (3.64-4.05)
Mx <sup>a</sup>	3	_	_
P-value	_	0.432	0.048*
Overall tumor stage			
1	0	_	_
11	5	1.76 (1.62–1.80)	3.25 (2.89-4.04)
III	33	1.86 (1.58–2.14)	3.56 (3.14-4.09)
IVA	27	1.59 (1.36–1.91)	3.16 (2.95–3.58)
IVB	12	1.62 (1.08–2.18)	3.75 (3.67-4.10)
Not available	3		
P-value	_	0.343	0.068

\*P < 0.05 indicates significant difference.  $Mx^a$ , metastatic status undetermined.



male patient with N3 stage NPC (C).



TABLE 3 | APT-related parameters in groups with different EBV-related indices.

	No.	APT <sub>mean</sub> (%)	APT <sub>90</sub> (%)
EBV-DNA			
+	23	1.72 (1.48–2.04)	3.44 (3.12-4.04)
-	53	1.69 (1.28-2.06)	3.45 (3.05–3.78)
Not available	4	_	_
P-value	_	0.995	0.730
EA-IgA			
+	18	1.70 (1.27–1.78)	3.14 (2.94-3.60)
-	16	1.84 (1.49–2.20)	3.54 (3.16-4.15)
Not available	46		
P-value	_	0.164	0.126
EB-VCA-IgA			
+	27	1.72 (1.59–1.74)	3.05 (2.92-3.38)
-	7	1.80 (1.59–2.00)	3.45 (2.92-4.01)
Not available	46		
P-value	_	0.531	0.206
EBER			
+	67	1.73 (1.40-2.08)	3.45 (3.05-4.03)
-	0		· _ /
Not available	13	-	-

show significant difference between responders and nonresponders. Such observations were consistent with our results that no significant difference in pretreatment  $APT_w$  signals was presented among different treatment response groups. This might also imply that monitoring the dynamic protein change throughout the treatment course using  $APT_w$  imaging may be more effective than detecting a stationary baseline pretreatment status in predicting treatment response.

Aside from the relatively limited sample size and imbalanced distribution in some groups, several limitations should be noted. First, conventional MTR asymmetry analysis was used to quantify APT

effect, which would be contaminated with magnetization transfer and nuclear overhauser enhancement effects. However, the method has been histologically demonstrated to be applicable in tumor characterization (41). In addition, the APT<sub>w</sub> signal in tumor has proved mostly attributed to the APT contribution under the currently utilized RF power of 2  $\mu$ T (42). Therefore, the findings regarding APT are valid. Second, this study employed single-slice APT imaging to shorten the scan time, and the limited spatial coverage may be inadequate to characterize intra-tumoral heterogeneity. 3D APT scanning integrated with fast data sampling strategies would be desired for whole-tumor characterization in the future.

TABLE 4 | Baseline APT-related parameters for different treatment response group.

	No.	APT <sub>mean</sub> (%)	APT <sub>90</sub> (%)
Overall treatment response			
PD	0	_	-
SD	8	2.11 (1.59–2.36)	3.52 (2.76-4.16)
PR	41	1.77 (1.50–2.04)	3.39 (3.13–3.74)
CR	17	1.72 (1.32–2.18)	3.65 (2.95-4.10)
P-value	_	0.562	0.880
Overall treatment response based on SVRR			
<50%	13	1.62 (1.38–2.11)	3.21 (3.05–3.56)
>50%	53	1.80 (1.48–2.16)	3.45 (3.13–4.04)
P-value	_	0.514	0.306
Treatment response after induction chemotherapy			
PD	0	-	-
SD	6	2.11 (1.33–2.46)	3.52 (2.65–3.92)
PR	35	1.77 (1.45–2.01)	3.25 (3.10–3.78)
CR	12	1.72 (1.53–2.22)	3.89 (3.15-4.20)
P-value	_	0.838	0.590
Treatment response after induction chemotherapy based	d on SVRR		
<50%	11	1.53 (1.33–2.01)	3.21 (3.01–3.57)
>50%	42	1.80 (1.49–2.17)	3.49 (3.13–4.04)
P-value	-	0.380	0.369





#### Conclusion

In conclusion, NPC showed significantly stronger APT effect than normal nasopharyngeal tissue, facilitating NPC lesion detection and early identification. However, baseline APT values showed limited value in predicting short-term response to IC and no significant correlation with different histopathological subtypes, EBV infection status, and clinical stages.

# DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/supplementary material. Further inquiries can be directed to the corresponding authors.

# ETHICS STATEMENT

The studies involving human participants were reviewed and approved by the Cancer Hospital & Shenzhen Hospital, Chinese Academy of Medical Sciences. Written informed consent for

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participation was not required for this study in accordance with the national legislation and the institutional requirements.

# **AUTHOR CONTRIBUTIONS**

DL and YW contributed to project idea and supervision and to manuscript revision, and maintained integrity of this manuscript. ZL and LZ implemented the whole study, analyzed the data, and drafted the manuscript. QY and LQ contributed to statistical analysis and reviewed the manuscript. TL, HL, and CC collected the raw MRI data. YL collected the pathological data. PC and CQ collected clinical information. XL provided technical support. All authors listed have made a substantial, direct, and intellectual contribution to the work and approved it for publication.

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Conflict of Interest: Author LQ was employed by GE Healthcare.

The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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