

Cancer Res Treat. 2019;51(1):413-414

https://doi.org/10.4143/crt.2018.335

Correspondence

Open Access

TRIUMPH Trial: One Small Step Could Become One Giant Leap for Precision Oncology in Head and Neck Cancer

Bhumsuk Keam, MD, PhD¹, **Hye Ryun Kim,** MD, PhD², **Hwan Jung Yun,** MD, PhD³

¹Department of Internal Medicine, Seoul National University Hospital, Seoul, ²Division of Medical Oncology, Department of Internal Medicine, Yonsei Cancer Center, Yonsei University College of Medicine, Seoul, ³Department of Internal Medicine, Chungnam National University Hospital, Daejeon, Korea

Recent remarkable progress in the fields of cancer genomics, computational analysis and drug discovery have changed the whole paradigm in cancer research. So called precision oncology, defined as molecular profiling of tumors to identify druggable alterations, is rapidly developing and waiting for entering the mainstream of cancer research as well as practice [1]. In the era of precision oncology, traditional classification based on organ or pathology do not have clinical meaning anymore. Molecular subtype base on next-generation sequencing (NGS) will lead us to appropriate molecular targeted agents.

Head and neck squamous cell carcinoma (HNSCC) is not a specific disease entity, but rather a broad category of diverse tumor types arising from various anatomic structures including oral cavity, oropharynx, hypopharynx, larynx and paranasal sinus. HNSCC is highly heterogeneous group of disease arises from the mucosal lining of the upper aerodigestive tract, demonstrates squamous differentiation, and involves older men with a long history of smoking. The prognosis by anatomic subsite quite differ. Human papilloma virus positive oropharyngeal cancer showed very good prognosis while oral cavity cancer had worst prognosis. In the era of NGS, HNSCC become more heterogeneous by mutational status [2-4].

Traditional design of clinical trials has faced big challenge for these heterogeneity and rarity. One solution is umbrella trial. In an umbrella trial, patients with specified cancer type are centrally screened and assigned to one of several molecularly defined subtrials where they receive matched targeted agents [5]. Some novel umbrella trials such as BATTLE trial [6] or MOSCATO trial [7] suggest that such a biomarker driven clinical trial can give appropriate benefit to the patients. To perform an umbrella trial, precise NGS based molecular phenotyping should be practical and working in the level of clinic. To date, there was surprisingly few study deals with the feasibility of molecular phenotyping for umbrella trial. We should answer the questions whether precision oncology is just a theory or whether it is realistically feasible. We should answer how we implement the precision oncology into the clinic and prove the patients' benefit.

In this issue of Cancer Research and Treatment, Lim et al. [8] reported the feasibility of targeted NGS to guide the treatment of HNSCC. The authors tested the feasibility from tissue sample process to analysis of NGS and mRNA expression at the practical level. Mutation profiles were similar with prior reports [2-4] and the authors found several targetable alterations such as PIK3CA, CDKN2A and CCND1.

Based on this success, KCSG (Korean Cancer Study Group) Head and Neck Cancer and Esophageal Cancer Committee launched novel umbrella trial: Translational blomarker Driven UMbrella Project for Head and Neck (TRIUMPH, NCT 03292250), which is the first umbrella trial for HNSCC in the world. TRIUMPH trial is for recurred/metastatic HNSCC, consisting of 5 targeted therapies including phosphoinositide 3-kinase inhibitor BYL719, pan-HER inhibitor poziotinib, fibroblast growth factor receptor inhibitor nintedanib, CDK4/6 inhibitor abemaciclib, andd immune checkpoint inhibitor durvalumab+/tremelimumab. TRIUMPH trial is investigator-initiated trial, and Korean Cancer Study Group affiliated 37 institutes participate. We think TRIUMPH trial is one small step for head and neck cancer, but hope to be one giant leap for precision oncology in HNSCC.

Correspondence: Hwan Jung Yun, MD, PhD

Department of Internal Medicine, Chungnam National University Hospital, 282 Munhwa-ro, Jung-gu, Daejeon 35015, Korea Tel: 82-42-280-7157, Fax: 82-42-257-5753, E-mail: hjyun@cnuh.co.kr

Received June 4, 2018 Accepted July 27, 2018 Published Online July 31, 2018

References

- 1. Sheikine Y, Kuo FC, Lindeman NI. Clinical and technical aspects of genomic diagnostics for precision oncology. J Clin Oncol. 2017;35:929-33.
- 2. Agrawal N, Frederick MJ, Pickering CR, Bettegowda C, Chang K, Li RJ, et al. Exome sequencing of head and neck squamous cell carcinoma reveals inactivating mutations in NOTCH1. Science. 2011;333:1154-7.
- 3. Stransky N, Egloff AM, Tward AD, Kostic AD, Cibulskis K, Sivachenko A, et al. The mutational landscape of head and neck squamous cell carcinoma. Science. 2011;333:1157-60.
- 4. Cancer Genome Atlas Network. Comprehensive genomic characterization of head and neck squamous cell carcinomas. Nature. 2015:517:576-82.
- 5. Renfro LA, Sargent DJ. Statistical controversies in clinical research: basket trials, umbrella trials, and other master protocols: a review and examples. Ann Oncol. 2017;28:34-43.
- 6. Kim ES, Herbst RS, Wistuba II, Lee JJ, Blumenschein GR Jr, Tsao A, et al. The BATTLE trial: personalizing therapy for lung cancer. Cancer Discov. 2011;1:44-53.
- 7. Massard C, Michiels S, Ferte C, Le Deley MC, Lacroix L, Hollebecque A, et al. High-throughput genomics and clinical outcome in hard-to-treat advanced cancers: results of the MOSCATO 01 Trial. Cancer Discov. 2017;7:586-95.
- 8. Lim SM, Cho SH, Hwang IG, Choi JW, Chang H, Ahn MJ, et al. Implementation the feasibility of targeted next-generation sequencing to guide the treatment of head and neck squamous cell carcinoma. Cancer Res Treat. 2019;51:300-12.