

Huynh K, Quoc G V, Thanh T N C, Loc H N, Thanh V C. Whole-genome sequencing of three local rice varieties (*Oryza sativa* L.) in Vietnam. *Plant Science Today*. 2021;8(3):437-444. <https://doi.org/10.14719/pst.2021.8.3.1047>

Supplementary Tables

Table S1. Total numbers of SNP and InDel in this study

Samples	SNPs	InDels
NTCD-T	1,959,489	174,506
Tai Nguyen	1,859,736	157,693
NTCD-TN	2,040,301	184,111

Table S2. Number of SNPs and InDel effects by impact

Samples	SNPs				InDels		
	High	Low	Moderate	Modifier	High	Moderate	Modifier
NTCD-T	1,829	57,925	59,671	7,363,679	2,013	3,027	801,689
Tai Nguyen	1,758	56,417	58,098	7,021,002	1,786	2,940	724,851
NTCD-TN	1,869	59,222	61,112	7,607,886	2,071	3,158	842,807

Table S3. Number of SNPs effects by functional class

Samples	Missense	Nonsense	Silent
NTCD-T	60,174	895	53,135
Tai Nguyen	58,598	845	51,85
NTCD-TN	61,635	915	54,313

Table S4. Non-synonymous SNPs and high-effect InDels

Samples	Non-synonymous SNPs (%)	High-effect InDels (%)
NTCD-T	0.79	0.25
Tai Nguyen	0.81	0.24
NTCD-TN	0.79	0.24