NRG Oncology/RTOG 0436 Data Dictionary Dictionary for NCT00655876-D1-Dataset.csv

Note #1: Local failure was estimated by the Kaplan-Meier method (per Section 13.4.1.2 in protocol) and not the cumulative incidence method, which was erroneously printed in the manuscript.

Note #2: Adverse events (AEs) were graded with the Common Terminology Criteria for Adverse Events version 3.0 (CTCAE v3.0).

Note #3: Data can be used to approximate published study findings, but exact reproduction of previous manuscripts may not be possible in some cases (e.g., when data must be modified for de-identification purposes or have undergone further data cleaning).

See also the companion dataset / data dictionary:

NCT00655876-D2-Dataset.csv / NCT00655876-D2-Data-Dictionary.pdf

#	Variable	Description	Coding for categorical/Min-max for continuous
1	study no	Study number	Character length 9 [RTOG-0436]
2	cn_deidentified	De-identified	Length 8
		patient ID	
3	rx	Assigned treatment	1 = RT + Chemo + Cetuximab [radiation therapy + paclitaxel +
			cisplatin + cetuximab]
			2 = RT + Chemo [radiation therapy + paclitaxel + cisplatin]
4	case_status	Case Status	1 = Eligible
			2 = No protocol treatment
			3 = Ineligible
5	reason_for_exclusion	Reason for	Character length 100
		Exclusion	Reason is not blank (n=16) [analysis_flag_1=1]
			Reason is blank (n=328) [analysis_flag_1=2]
6	analysis_flag_1	Eligible and	1=No (n=16)
		evaluable for	2=Yes (n=328)
		efficacy endpoints	
7	analysis_flag_2	Eligible and	1=No (n=25)
		evaluable for	2=Yes (n=319)
		adverse events	
8	reason_no_treatment	Reason patient did	Character length 100
		not receive	Reason is not blank (n=9) [case_status=2]
		protocol treatment	Reason is blank (n=335) [case_status=1,3]
9	age	Age (years)	Continuous
			Min-max = 32-87
			. = Not applicable (n=16; analysis_flag_1=1)
10	agefm	Age (years)-	1 = <u>≤</u> 49
		Categorized	2 = 50-59
			3 = 60-69
			4 = 70-79
			$5 = \ge 80$
<u> </u>			. = Not applicable (n=16; analysis_flag_1=1)
11	sex	Sex	1 = Male
			2 = Female
			. = Not applicable (n=16; analysis_flag_1=1)
12	race	Race	1 = American Indian or Alaskan Native
			2 = Asian
			3 = Black or African American
			4 = Native Hawaiian or Other Pacific Islander
			5 = White
			6 = More than 1 race
			9 = Unknown
			. = Not applicable (n=16; analysis_flag_1=1)

#	Variable	Description	Coding for categorical/Min-max for continuous
13	ethnicity	Ethnicity	1 = Hispanic or Latino
			2 = Not Hispanic or Latino
			9 = Unknown
			. = Not applicable (n=16; analysis_flag_1=1)
14	zubrod	Zubrod	Values = 0, 1, 2
		performance status	. = Not applicable (n=16; analysis_flag_1=1)
15	tstage	T stage (clinical)	1 = T1
			2 = T2
			3 = T3
			4 = T4
			. = Not applicable (n=16; analysis flag 1=1)
16	nstage	N stage (clinical)	1 = N0
10	listage	it stage (enniour)	2 = N1
			$=$ Not applicable (n=16; analysis_flag_1=1)
17	mstage	M stage (clinical)	1 = M0
17	instage	wi stuge (chinear)	2 = M1a
			. = Not applicable (n=16; analysis_flag_1=1)
18	lesion size	Cancer lesion size,	$1 = -100$ applicable (n=10, analysis_nag_1=1)
10	lesion_size	· · · · · · · · · · · · · · · · · · ·	$2 = \ge 5 \text{ cm}$
		cm	
10	histolo are	Histologia findinga	. = Not applicable (n=16; analysis_flag_1=1) 1 = Adenocarcinoma
19	histology	Histologic findings	
			2 = Squamous cell
20	1'		. = Not applicable (n=16; analysis_flag_1=1)
20	celiac	Celiac nodes status	1 = Present
			2 = Absent
			. = Not applicable (n=16; analysis_flag_1=1)
21	endoscopy_results	Results of the	1 = Clinical complete response (cCR) (biopsy not required)
		Endoscopic	2 = Residual disease or suspicion of residual disease and negative
		Assessment	biopsy
			3 = Residual disease or suspicion of residual disease at the time of
			endoscopy and positive biopsy
			4 = Residual disease or suspicion of residual disease at the time of
			endoscopy and no biopsy performed to pathologically prove disease
			5 = Endoscopy not performed
			. = Not applicable (n=16; analysis_flag_1=1)
22	ccr	cCR Status	0= Yes, cCR (endoscopy_results = 1)
			1= No, residual disease (endoscopy_results = 2,3,4)
			999 = Endoscopy not performed (endoscopy_results = 5)
			. = Not applicable (n=16; analysis_flag_1=1)
23	interim_adeno_flag	Included in the	1=No (n=194)
		protocol-specified	2=Yes (n=150*)
		interim cCR	
		analysis of	*Note: One patient was deemed to be ineligible after this interim
		adenocarcinoma	analysis so they have analysis_flag_1=1.
		patients?	
24	ccr_adeno	cCR Status at the	0 = Yes, cCR
		interim analysis for	1= No, residual disease
		adenocarcinoma	. = Not applicable (n=194; interim_adeno_flag =1)
		patients	
25	interim_scc_flag	Included in the	1=No (n=231)
		protocol-specified	2=Yes (n=113)
		interim cCR	
		analysis of	
	1	Squamous cell	

#	Variable	Description	Coding for categorical/Min-max for continuous
		patients?	
26	ccr_scc	cCR Status at the	0= Yes, cCR
		interim analysis for	1= No, residual disease
		squamous cell	. = Not applicable (n=231; interim_scc_flag =1)
		patients	
27	surgery_followup	Surgery in follow-	1=No
		up for recurrent or	2=Yes
		residual disease?	. = Not applicable (n=16; analysis_flag_1=1)
28	surgery_followup_reason	Reason for surgery	1 = Residual disease (n=26)
		in follow-up	2 = Recurrent disease (n = 18)
			999 = Surgery not performed (n=284; surgery_followup=1)
			. = Not applicable (n=16; analysis_flag_1=1)
29	tvscore	Tumor volume	1 = Per protocol
		contouring score	2 = Acceptable variation
			3 = Unacceptable variation
			9 = Not evaluable
			. = Not applicable (n=16; analysis_flag_1=1)
30	oarscore	Organs at risk	1 = Per protocol
		contouring score	2 = Acceptable variation
			3 = Unacceptable variation
			9 = Not evaluable
21	, 1	T 1	. = Not applicable (n=16; analysis_flag_1=1)
31	tvscore_dva	Tumor volume	1 = Per protocol
		dose volume	2 = Acceptable variation
		analysis score	3 = Unacceptable variation 9 = Not evaluable
32	oarscore dva	Organs at risk dose	. = Not applicable (n=16; analysis_flag_1=1) 1 = Per protocol
32	oarscore_dva	volume analysis	2 = Acceptable variation
		score	9 = Not evaluable
		score	. = Not evaluation (n=16; analysis_flag_1=1)
33	chemo_review	Overall	1 = Per protocol
55	enemo_review	Chemotherapy/	2 = Not per protocol
		Targeted Agent	9 = Not evaluable
		Review Score	. = Not applicable (n=16; analysis_flag_1=1)
34	chemo review details	Details of Overall	1 = Per protocol, no modifications or delays
		Chemotherapy/	2 = Per protocol, modifications and/or delays
		Targeted Agent	$3 =$ Not per protocol, modifications and/or delays with $\ge 80\%$ of
		Review Score	protocol dose given
			4 = Not per protocol, modifications and/or delays with $< 80%$ of
			protocol dose given
			9 = Not evaluable
			. = Not applicable (n=16; analysis_flag_1=1)
35	chemo_modification_ae	Chemotherapy/	1 = No
	_	Targeted Agent	2 = Yes
		dose	. = Not applicable (n=16; analysis_flag_1=1)
		modified/delayed	
		due to AEs	Note: This is based on the site's reporting on whether the
			chemotherapy/targeted agent dose was modified due to AEs.
			This is not based on the chemotherapy/targeted agent review score.
36	chemo_termination_ae	Chemotherapy/	1 = No
		Targeted Agent	2 = Yes
		treatment	. = Not applicable (n=16; analysis_flag_1=1)
		terminated due to	

#	Variable	Description	Coding for categorical/Min-max for continuous
		AEs	Note: This is based on the site's reporting on whether the
			chemotherapy/targeted agent treatment was terminated due to AEs.
			This is not based on the chemotherapy/targeted agent review score.
37	local_failure	Local failure status	0 = censored
			1 = event
			2 = competing event (death)
			. = Not applicable (n=16; analysis_flag_1=1)
38	local_failure_months	Local failure time	. = Not applicable (n=16; analysis_flag_1=1)
		(months)	Notes Time from the date of non-deniention to the date of local
			<i>Note: Time from the date of randomization to the date of local</i>
20		Overall survival	failure or last follow-up/death.
39	survival		0 = censored
		status	1 = event (death)
40	and in a south a	Overall survival	. = Not applicable (n=16; analysis_flag_1=1)
40	survival_months	time (months)	. = Not applicable (n=16; analysis_flag_1=1)
			Note: Time from the date of randomization to the date of death or
			last follow-up.
41	max_overall_grade	Overall highest	0 = grade 0
		treatment-related	1 = grade 1
		grade per patient	2 = grade 2
			3 = grade 3
			4 = grade 4
			5 = grade 5
			. = Not applicable (n=25; analysis_flag_2=1)
42	max_gi_grade	Overall highest	0 = grade 0
		grade for	1 = grade 1
		treatment-related	2 = grade 2
		gastrointestinal	3 = grade 3
		adverse events	4 = grade 4
			. = Not applicable (n=25; analysis_flag_2=1)
43	max_hem_grade	Overall highest	0 = grade 0
		grade for	1 = grade 1
		treatment-related	2 = grade 2
		hematologic	3 = grade 3
		adverse events	4 = grade 4
4.4		O	. = Not applicable (n=25; analysis_flag_2=1)
44	max_acne_grade	Overall highest	0 = grade 0
		grade for	1 = grade 1
		treatment-related	$2 = \operatorname{grade} 2$
		acniform rash	3 = grade 3
15	man nanham 1- 001	adverse events	. = Not applicable (n=25; analysis_flag_2=1)
45	max_nonhem_grade_90days	Overall highest	0 = grade 0
		grade for treatment-related	1 = grade 1
			2 = grade 2 3 = grade 3
		non-hematologic adverse events ≤	3 = grade 3 4 = grade 4
		90 days from	4 - grade 4 5 = grade 5
		-	
		treatment start	. = Not applicable (n=25; analysis_flag_2=1)