

A real-world cohort study of EGFR TKIs in patients with NSCLC with uncommon *EGFR* mutations (UpSwinG)

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Presenter DISCLOSURES

Ineligible Company (formerly: Commercial Interest)	Relationship(s)
Boehringer Ingelheim Inc.	Speaker bureau fees/Expert testimony
MSD Inc.	Speaker bureau fees/Expert testimony
Elli Lilly Japan	Speaker bureau fees/Expert testimony
Ono Pharma. Inc.	Speaker bureau fees/Expert testimony
Chugai Pharma Inc.	Speaker bureau fees/Expert testimony

Introduction

While most patients with *EGFR*^{m+} NSCLC harbor 'common' mutations (Del19/L858R), uncommon mutations are highly heterogeneous; these can be categorized into four groups:

Exon 18	Exon 19	Exon 20	Exon 21
E709X	Del19	Ex20ins	L858R
G719X	Ex19ins	S768I	L861Q
	L747P/S	T790M	

7–23% of *EGFR* mutations are 'uncommon' mutations¹

- Common (sensitive to all TKIs; afatinib approved in this setting)
- 'Major' uncommon (sensitive to TKIs; afatinib approved in this setting)
- Ex20ins (considered resistant to TKIs but highly heterogeneous)
- Others (little data on TKI sensitivity; highly heterogeneous)
- T790M (resistant to 1st- and 2nd-gen TKIs)

- Up to 25% of *EGFR*^{m+} tumors harbor compound mutations¹
- Increased use of sensitive sequencing-based detection methods and liquid biopsy will increase the frequency of uncommon mutations detected in real-world clinical practice²

EGFR^{m+}, *EGFR* mutation-positive; gen, generation; TKI, tyrosine kinase inhibitor.

1. Yang JC, et al. J Thorac Oncol 2020;15:803–15; 2. Kobayashi Y & Mitsudomi T. Cancer Sci 2016;107:1179–86



Methods

UpSwinG (NCT04179890): non-interventional global study (nine countries*) that assessed medical records of consecutive patients with uncommon *EGFR* mutations treated in real-world practice (max 15 per site)

Aims (uncommon mutations cohort)

- 1) Investigate the treatment of patients with uncommon *EGFR* mutations
- 2) Assess the efficacy of EGFR TKIs in each uncommon mutation category
- 3) Assess how *EGFR* mutations are detected in real-world practice

Patients (n=246)

- All had at least one uncommon mutation
- All received an EGFR TKI (afatinib, gefitinib, erlotinib, or osimertinib) in 1st- or 2nd-line
- *EGFR* mutation detection undertaken locally using different methodologies

Key exclusion criteria

- Treated in a clinical trial
- Active brain metastases
- Patients with acquired T790M only and treated with osimertinib

Primary objective: TTF

Secondary objectives: ORR, OS, DoR

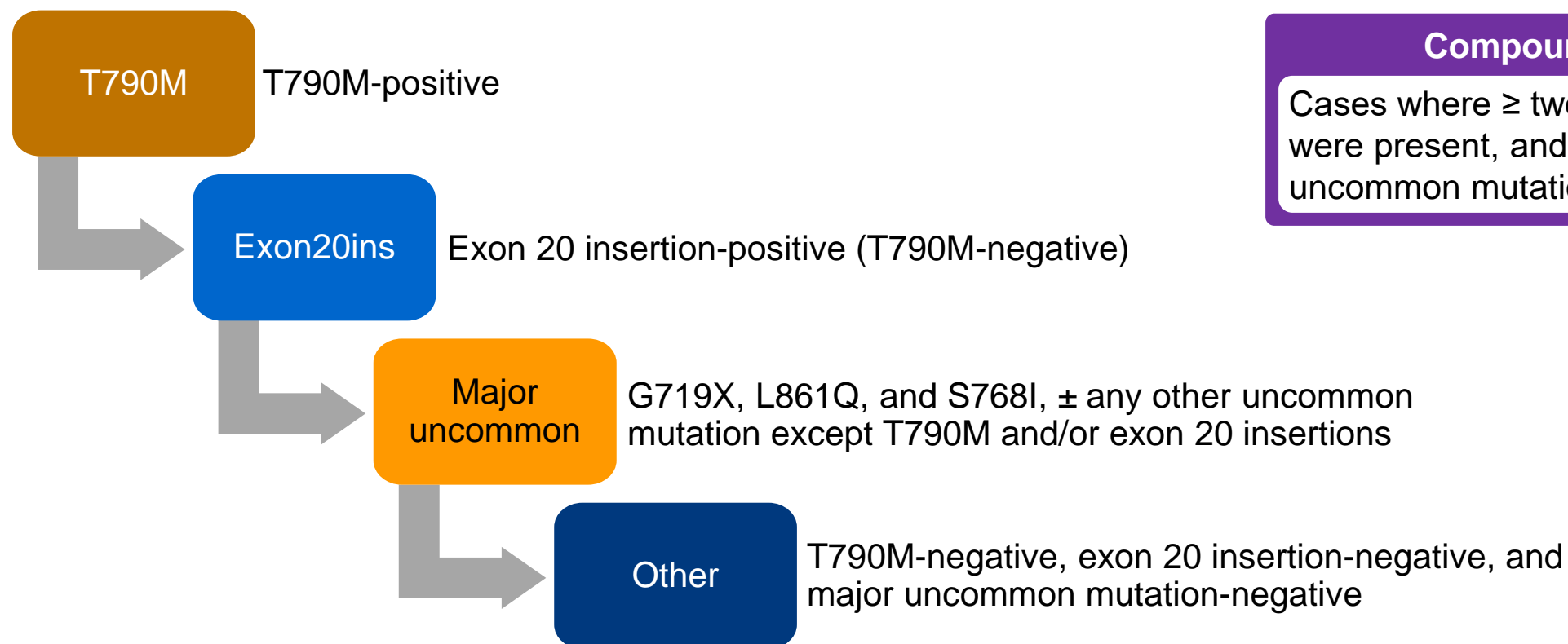
DoR, duration of response; ORR, overall response rate; OS, overall survival; TTF, time-to-treatment failure.

*Austria, France, Germany, Italy, Japan, South Korea, Spain, Taiwan, United Kingdom



Methods (continued)

Patients were categorized hierarchically according to tumor mutation



Compound mutations

Cases where \geq two *EGFR* mutations were present, and \geq one was an uncommon mutation

Results

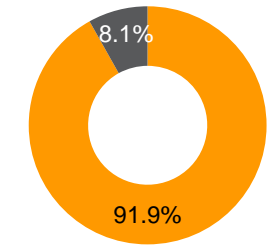
Patient characteristics were generally similar regardless of which EGFR TKI was received as index therapy

		All (N=246)	1 st -gen EGFR TKIs (n=106*)	Afatinib (n=132)
	Female, n (%)	138 (56.1)	66 (62.3)	67 (50.8)
	Median age, years (range)	69.5 (27.0–93.0)	70.5 (42.0–91.0)	68.5 (27.0–93.0)
	Never-smoker, n (%)	129 (52.4)	64 (60.4)	62 (47.0)
BMI, n (%)	Normal	116 (47.2)	42 (39.6)	68 (51.5)
	Underweight	17 (6.9)	9 (8.5)	7 (5.3)
	Overweight/obese	49 (19.9)	21 (19.8)	27 (20.5)
Ethnicity, n (%)	Caucasian	23 (9.3)	8 (7.5)	13 (9.8)
	Asian	206 (83.7)	87 (82.1)	114 (86.4)
	Unknown/not collected	17 (6.9)	11 (10.4)	5 (3.8)
Stage/histology, n (%)	Stage IIIB/C	37 (15.0)	14 (13.2)	22 (16.7)
	Stage IV	209 (85.0)	92 (86.8)	110 (83.3)
	Adenocarcinoma	239 (97.2)	102 (96.2)	129 (97.7)
	Brain metastases	17 (6.9)	5 (4.7)	12 (9.1)
ECOG PS, n (%)	0	35 (14.2)	9 (8.5)	25 (18.9)
	1	124 (50.4)	61 (57.5)	59 (44.7)
	2–4	31 (12.6)	14 (13.2)	17 (12.9)
	Unknown	56 (22.8)	22 (20.8)	31 (23.5)

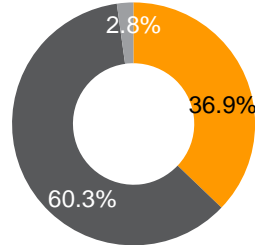
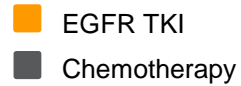
BMI, body mass index; ECOG PS, Eastern Cooperative Oncology Group performance status

Results (continued)

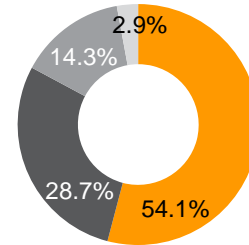
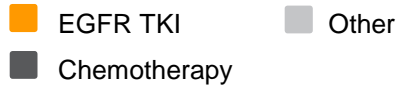
EGFR TKIs were generally the 1st-line treatment of choice for uncommon mutations



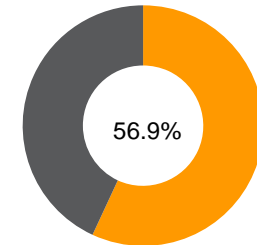
1st-line treatment*



2nd-line treatment*

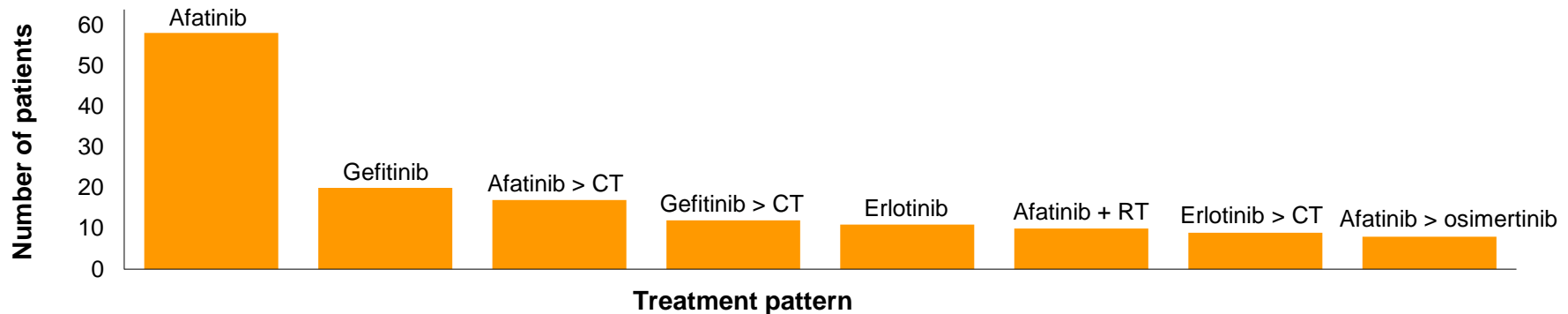


Index therapy^{†,‡}



Subsequent therapy

Of patients received > one line of therapy



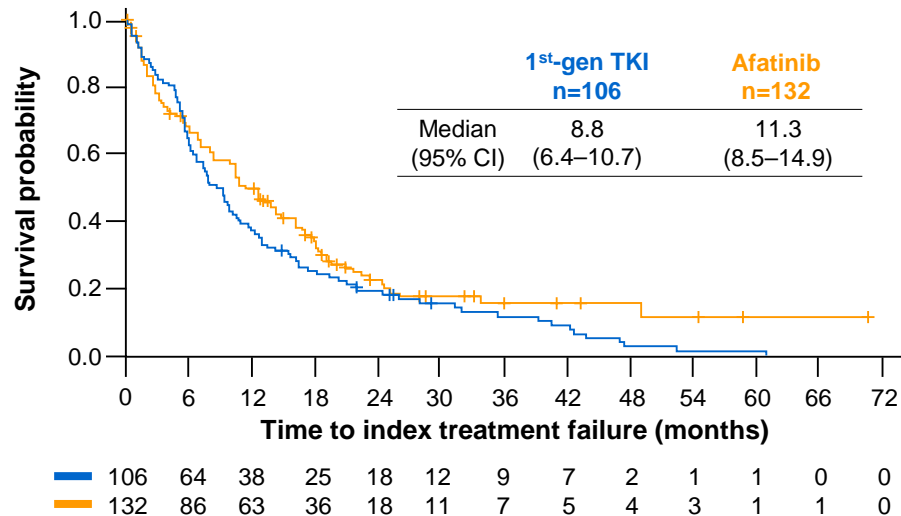
CT, chemotherapy; RT, radiotherapy.

*An additional patient was treated with chemotherapy plus bevacizumab; [†]Includes one patient treated with gefitinib/erlotinib; [‡]Includes one patient treated with afatinib/gefitinib

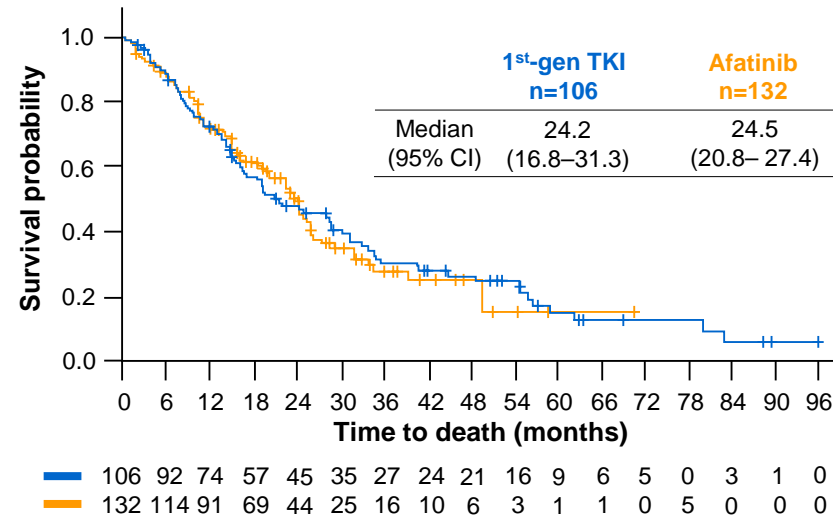
Results (continued)

EGFR TKIs conferred encouraging TTF, ORR and OS; the OS was likely confounded by the high uptake of subsequent treatments and less censoring for the 1st-generation EGFR TKIs due to their longer availability*

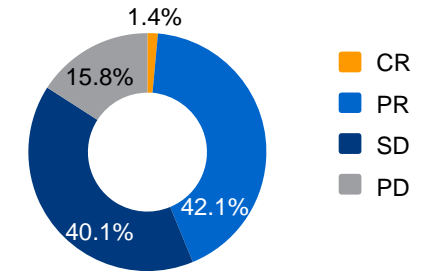
TTF



OS



Response to index EGFR TKI therapy†



CI, confidence interval; CR, complete response; PD, progressive disease; PR, partial response; SD, stable disease.

*Results from patients treated with osimertinib not shown due to small sample size; †Evaluable patients

Results (continued)

Clinical outcomes varied according to mutation category in patients treated with EGFR TKIs*

	Any TKI (n=246)				1 st -gen EGFR TKIs (n=106)				Afatinib (n=132)			
	TTF, mos	OS, mos	ORR [†] , %	DoR [†] , mos	TTF, mos	OS, mos	ORR [†] , %	DoR [†] , mos	TTF, mos	OS, mos	ORR [†] , %	DoR [†] , mos
All patients	9.9	24.4	43.4	10.0	8.8	24.2	44.1	6.0	11.3	24.5	43.8	12.0
■ Major uncommon	11.3	25.7	49.1	10.0	9.8	28.5	47.3	6.5	14.3	24.5	50.6	12.0
■ Exon 20 insertion	5.5	22.5	17.4	19.3	5.2	21.0	16.7	33.0	8.3	22.5	18.8	5.5
■ T790M	2.8	32.7	20.0	6.0	2.1	14.2	0	-	5.7	-	33.3	6.0
■ Other	7.4	13.4	43.8	7.5	7.3	12.8	55.6	4.5	10.8	20.2	28.6	10.5
■ Compound	12.3	28.7	48.6	10.0	12.4	31.3	48.3	6.0	12.6	23.4	52.5	10.0

Mos, months.

*Results from patients treated with osimertinib not shown due to small sample size; [†]Evaluable patients

Results (continued)

Most patients (76.8%) received the approved starting dose of EGFR TKIs (gefitinib 250 mg; erlotinib 150 mg; afatinib 40 mg); outcomes were not affected by starting dose*

	Any TKI (n=189)				1 st -gen EGFR TKIs (n=91)				Afatinib (n=93)			
	TTF, mos	OS, mos	ORR [†] , %	DoR [†] , mos	TTF, mos	OS, mos	ORR [†] , %	DoR [†] , mos	TTF, mos	OS, mos	ORR [†] , %	DoR [†] , mos
All patients	10.7	25.6	45.4	10.0	9.3	28.2	47.6	6.0	12.8	24.8	44.2	12.0
■ Major uncommon	11.9	28.5	50.0	10.0	10.4	30.2	51.5	6.5	15.7	24.5	49.2	12.0
■ Exon 20 insertion	10.6	22.5	16.7	-	6.0	21.0	0	-	15.5	22.5	22.2	-
■ T790M	2.8	23.0	21.4	6.0	2.1	14.2	0	-	5.7	-	33.3	6.0
■ Other	11.5	20.2	50.0	6.0	7.4	13.4	55.6	4.5	13.3	24.8	33.3	12.0
■ Compound	12.6	30.2	50.0	10.0	14.0	33.0	52.0	4.5	12.6	23.4	51.7	10.0

*Results from patients treated with osimertinib not shown due to small sample size; [†]Evaluable patients



Key findings and conclusions

- EGFR TKIs are 1st-line treatment of choice for uncommon mutations in everyday clinical practice
- Afatinib was the most commonly used EGFR TKI
- Overall, median TTF was 9.9 months and median OS was 24.4 months
- Strongest results were seen in major uncommon and compound mutations
- Treatment with an EGFR TKI should be considered for most patients with uncommon mutations

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