

Table SIII. Patient characteristics of 955 NSCLC samples downloaded from The Cancer Genome Atlas.

Subset	Group	Total (n=955)		<75 (n=776)		≥75 (n=179)		P-values
		Median (range)	n (%)	Median (range)	n (%)	Median (range)	n (%)	
Age, years		67 (38-90)	-	65 (38-74)	-	78 (75-90)	-	<0.0001
Sex	Male		569 (59.6)		470 (60.6)		99 (55.3)	0.2
	Female		386 (40.4)		306 (39.4)		80 (44.7)	
Histology	Ad		488 (51.1)		397 (51.2)		91 (50.8)	0.6
	Sq		467 (48.9)		379 (48.8)		88 (49.2)	
Race	Asian		17 (1.8)		12 (1.5)		5 (2.8)	-
	Not hispanic or latino		675 (70.7)		553 (71.3)		122 (68.2)	
	Hispanic or latino		15 (1.6)		14 (1.8)		1 (0.6)	
	African American		80 (8.4)		68 (8.8)		12 (6.7)	
	American Indian or Alaskan Native		1 (0.1)		1 (0.1)		0 (0.0)	
	Unknown		167 (17.5)		128 (16.5)		39 (21.8)	
EGFR mutation	Mutant		77 (8.1)		64 (8.2)		13 (7.3)	0.8
	Wild type ^a		878 (91.9)		712 (91.8)		166 (92.7)	

^aIncluding a case without information of EGFR mutational status. PS, performance status; Ad, adenocarcinoma; Sq, squamous cell carcinoma.