

Table 1, Patient clinicopathological characteristics and TS polymorphism variant analysis in 140 NSCLC patients harboring driver EGFR mutations.

Characteristics		TS polymorphism,n,(%)			
		Wildtype	3R	3R+6bp-	6bp-
Sex,n,(%)		68(49)	6(4)	25(18)	41(29)
	Male	27(19)	4(3)	5(4)	14(10)
	Female	41(29)	2(1)	20(14)	27(19)
Age,years,n,(%)	< 65	46(33)	5(4)	18(13)	22(16)
	≥65	21(15)	1(1)	7(5)	20(14)
Stage,n,(%)	I	18(13)	3(2)	6(4)	9(6)
	II	5(4)	0(0)	1(1)	3(2)
	III	3(2)	11(8)	2(1)	5(4)
	IV	39(28)	2(1)	15(11)	19(14)
	NA	2(1)	0(0)	1(1)	6(4)
EGFR mutation, n(%)	19DEL	25(18)	2(1)	11(8)	16(11)
	21L858R	33(24)	3(2)	11(8)	22(16)
	19DEL and 21L858R	0(0)	0(0)	0(0)	1(1)
	Rare mutations	8(6)	1(1)	3(2)	4(3)