A1. Feature Selection Results

TABLE I. GAIN RATIO WITH RANKER SEARCH

Average Merit	Average Rank	Attribute									
0.171 +- 0.003	1 +- 0	31 Protein_Family									
0.157 +- 0.002	2 +- 0	32 Protein_Group									
0.112 +- 0.003	3 +- 0	33 Sub_domain									
0.093 +- 0.031	5.1 +- 3.3	7 Blosum62									
0.086 +- 0.008	5.1 +- 0.54	30 Natural_Freqs_Diff									
0.079 +- 0.004	5.7 +- 0.46	29 Hydropathy_Index_Diff									
0.045 +- 0.002	7.4 +- 0.66	27 Side_Chain_Polarity_Diff									
0.044 +- 0.008	8.3 +- 1.73	1 Original_Amino_Acid									
0.038 +- 0.003	9.2 +- 0.87	5 Conservation_Consensus_Family									
	10.5 +-	2 SNP_Amino_Acid									
0.033 +- 0.006	1.69										
	12.5 +-	4 Conservation_Wild_Allkinase									
0.011 +- 0.001	1.12										
	13.5 +-	12 binding_site									
0.008 +- 0.001	1.12										
	13.5 +-	20 Side_Chain_Polarity_Wild									
0.027 +- 0.014	6.26										
	14.5 +-	26 Van_der_Waals_Volume_Wild									
0.024 +- 0.016	5.55										
	15.1 +-	13 Side_Chain_Polarity									
0.001 +- 0	1.45										
0.001 +- 0.002	15.5 +- 1.2	24 Avg_Mass_Wild									
	15.9 +-	9 modification									
0 +- 0	1.37										
	17.8 +-	11 is_active_site									
0.005 + -0.014	3.71										

TABLE II. CORRELATION BASED FEATURE SELECTION

Number of Folds (%)	Attribute								
10(100 %)	7 Blosum62								
10(100 %)	31 Protein_Family								
10(100 %)	32 Protein_Group								
8(80%)	12 binding_site								
2(20 %)	4 Conservation_Wild_Allkinase								
1(10%)	3 Conservation_Consensus_Allkinase								

TABLE III. RELIEF BASED FEATURE SELECTION

Average Merit	Average	Attribute
Average Ment	Rank	Attribute
0.25 +- 0.005	1 +- 0	31 Protein_Family
0.206 +- 0.005	2 +- 0	32 Protein_Group
0.087 +- 0.006	3.2 +- 0.4	1 Original_Amino_Acid
0.082 +- 0.005	3.8 +- 0.4	7 Blosum62
0.055 +- 0.005	5.4 +- 0.49	33 Sub_domain
0.057 +- 0.008	5.6 +- 0.49	2 SNP_Amino_Acid
0.037 +- 0.005	7 +- 0	27 Side_Chain_Polarity_Diff
0.027 +- 0.003	8.4 +- 0.8	5 Conservation_Consensus_Family
0.026 +- 0.001	9 +- 0.63	26 Van_der_Waals_Volume_Wild
	10.9 +-	
0.022 +- 0.001	0.94	23 Natural_Freqs_Wild
0.022 +- 0.003	11 +- 1.48	13 Side_Chain_Polarity
0.022 +- 0.001	11 +- 0.89	30 Natural_Freqs_Diff
	12.7 +-	
0.019 +- 0.001	0.64	24 Avg_Mass_Wild
	15.2 +-	
0.014 +- 0.002	1.25	22 Hydropathy_Index_Wild
0.014 +- 0.002	15.3 +- 0.9	20 Side_Chain_Polarity_Wild
	15.8 +-	
0.014 +- 0.002	1.66	19 Van_der_Waals_Volume
	16.6 +-	
0.013 +- 0.002	0.66	29 Hydropathy_Index_Diff
	17.7 +-	
0.012 +- 0.002	1.62	16 Natural_Freqs
	18.9 +-	
0.011 +- 0.002	1.14	17 Avg_Mass
	19.5 +-	
0.01 +- 0.002	0.67	15 Hydropathy_Index
	22.1 +-	
0.007 +- 0.002	1.58	4 Conservation_Wild_Allkinase
0.006 +- 0.002	22.3 +- 1.1	25 Isoelectric_Point_Wild
	23.1 +-	
0.006 +- 0.002	1.51	18 Isoelectric_Point
	23.9 +-	
0.005 +- 0.002	1.22	14 Side_Chain_Charge
0.005 +- 0.001	24.4 +- 0.8	28 Side_Chain_Charge_Diff
	25.3 +-	
0.004 +- 0.002	1.49	21 Side_Chain_Charge_Wild

TABLE IV. WRAPPER: SMO POLY KERNEL AND BEST FIRST SEARCH

Number of Folds (%)	Attribute
10(100 %)	7 Blosum62
10(100 %)	31 Protein_Family
7(70%)	23 Natural_Freqs_Wild
6(60%)	15 Hydropathy_Index
4(40%)	9 modification
3(30%)	27 Side_Chain_Polarity_Diff
3(30%)	29 Hydropathy_Index_Diff
2(20%)	16 Natural_Freqs
2(20%)	22 Hydropathy_Index_Wild
1(10%)	12 binding_site
1(10%)	13 Side_Chain_Polarity
1(10%)	14 Side_Chain_Charge
1(10%)	20 Side_Chain_Polarity_Wild
1(10%)	21 Side_Chain_Charge_Wild
1(10%)	26 Van_der_Waals_Volume_Wild

TABLE V. WRAPPER: SMO POLY KERNEL AND GENETIC SEARCH

Number of Folds (%)	Attribute
10(100 %)	7 Blosum62
10(100 %)	31 Protein_Family
8(80%)	16 Natural_Freqs
8(80%)	21 Side_Chain_Charge_Wild
7(70%)	6 Conservation_Consensus_Group
7(70%)	13 Side_Chain_Polarity
6(60%)	15 Hydropathy_Index
6(60%)	23 Natural_Freqs_Wild
6(60%)	25 Isoelectric_Point_Wild
5(50%)	3 Conservation_Consensus_Allkinase
5(50%)	9 modification
5(50%)	30 Natural_Freqs_Diff
4(40%)	12 binding_site
4(40%)	29 Hydropathy_Index_Diff
3(30%)	18 Isoelectric_Point
2(20%)	8 is_signal_peptide
2(20%)	10 is_pk_domain
2(20%)	22 Hydropathy_Index_Wild
2(20%)	27 Side_Chain_Polarity_Diff
2(20%)	28 Side_Chain_Charge_Diff
1(10%)	4 Conservation_Wild_Allkinase
1(10%)	5 Conservation_Consensus_Family
1(10%)	11 is_active_site
1(10%)	14 Side_Chain_Charge
1(10%)	17 Avg_Mass
1(10%)	24 Avg_Mass_Wild
1(10%)	26 Van_der_Waals_Volume_Wild
1(10%)	33 Sub_domain

A2. List of Prioritization of EGFR Mutations

ID	Gene	MutationPos		MutantTyne	Disease_In_Cosmic J4	A F	RandomForest NBTree	FT	DecisionTable	DTNR	I W/I	BayesNet	NaiveRaves	SVM	NeuralNetwork	Score	Avg_Weighted_Score	Positive_Weighted_Score
	EGFR	713	К	F		s ye			yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	715	ı	<u>.</u> S		s ye		•	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	719	G	<u>A</u>	· · · · · · · · · · · · · · · · · · ·	s ye	ľ	•	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	719		V	· · · · · · · · · · · · · · · · · · ·	s ye	ľ		yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	720		F	' '	s ye		_	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	720		P		s ye		yes	•	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	721		A	' I'	s ye		yes	•	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
_	EGFR	721		D	· · · · · · · · · · · · · · · · · · ·	s ye	ľ		yes	yes	yes	ves	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	721		S	· · · · · · · · · · · · · · · · · · ·	s ye	ľ		yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	723		S		s ye			yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	724		S	' I'	s ye			yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
12	EGFR	725	Т	M	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
13	EGFR	727	Υ	С	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
14	EGFR	729	G	E	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
15	EGFR	729	G	R	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
16	EGFR	730	L	F	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
17	EGFR	731	W	R	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
18	EGFR	733	Р	S	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
19	EGFR	733	P	Т	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
20	EGFR	735	G	S	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
21	EGFR	741	Р	L	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
22	EGFR	743		Р	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
23	EGFR	743	A	S	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	743	Α	Т	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
_	EGFR	747	L	P		s ye		yes		yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	747	L	S	' I'	s ye		•	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	751		I	' I'	s ye		•	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
_	EGFR	753		F		s ye			yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
_	EGFR	753		<u>S</u>	· · · · · · · · · · · · · · · · · · ·	s ye	ľ		yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	755		D		s ye			yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
_	EGFR	758		G		s ye		yes		yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	761		G	' '	s ye		_	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000 1.0000
	EGFR EGFR	761 766		N T	' I'	s ye		•	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000 1.0000	1.0000
	EGFR	770		N	· · · · · · · · · · · · · · · · · · ·	s ye	ľ		yes	yes	yes yes	yes yes	yes yes	yes	yes	1.0000	1.0000	1.0000
_	EGFR	773		1		s ye		_	yes yes	yes yes	yes	yes	yes	yes yes	yes	1.0000	1.0000	1.0000
	EGFR	773		Υ		s ye		yes	•	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	776		C		s ye		yes	•	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	776		G	' I'	s ye		•	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	777		Q	<i>'</i>	s ye	· · · · · · · · · · · · · · · · · · ·		yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	779		C		s ye			yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	779		F	· · · · · · · · · · · · · · · · · · ·	s ye	ľ		yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
43	EGFR	779	G	S		s ye		_	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
44	EGFR	784	S	F	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
45	EGFR	796	G	A	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
46	EGFR	796	G	S	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	798	L	F	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
48	EGFR	799	L	R		s ye		yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
49	EGFR	810	G	D	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	810		S	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	814	_	Р		s ye		yes		yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	829		Q		s ye		yes		yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
	EGFR	832		С		s ye		_	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
54	EGFR	832	R	L	yes ye	s ye	es yes	yes	yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000

						1	1	1						
55 EGFR	833 L	V yes	yes ye		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
56 EGFR	835 H	L yes	yes ye		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
57 EGFR	838 L	P yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
58 EGFR	838 L	V yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
59 EGFR	839 A	T yes	yes ye	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
60 EGFR	842 N	D yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
61 EGFR	847 T	A yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
62 EGFR	847 T	l yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
63 EGFR	849 Q	L yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
64 EGFR	850 H	D yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
65 EGFR	850 H	N yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
66 EGFR	853 I	T yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
67 EGFR	854 T	A yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
68 EGFR	855 D	G yes	yes yes	es yes	yes yes	yes	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
69 EGFR	855 D	N yes	yes yes	es yes	yes yes	ves	yes	yes	yes	yes	yes	1.0000	1.0000	1.0000
70 EGFR	856 F	L yes	yes yes	- '	yes yes	yes	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
71 EGFR	856 F	S yes	yes yes		yes yes	ves	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
72 EGFR	857 G	E yes	yes yes		yes yes	yes	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
73 EGFR	858 L	A yes	yes yes		yes yes	yes	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
74 EGFR	858 L	G yes	yes yes		yes yes	ves	yes	yes		yes	yes	1.0000	1.0000	1.0000
75 EGFR	858 L	K yes	yes yes		yes yes	yes	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
76 EGFR	858 L	M yes	yes yes		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
77 EGFR	858 L	Q yes	yes yes		yes yes	ves	yes	yes		ves	yes	1.0000	1.0000	1.0000
78 EGFR	858 L	W yes	yes yes	- '	yes yes	yes	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
79 EGFR	860 K		yes yes		yes yes	yes	_	yes				1.0000	1.0000	1.0000
80 EGFR	861 L	F yes	yes yes	· ·	yes yes	·	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
81 EGFR	861 L		yes yes		· '	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
-	861 L	P yes R yes			yes yes	yes	yes	-		yes	yes	1.0000	1.0000	1.0000
82 EGFR		765	yes yes		yes yes	yes	yes	yes	·	yes	yes			
83 EGFR	861 L	V yes	yes yes		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
84 EGFR	862 L	P yes	yes yes	- '	yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
85 EGFR	863 G	D yes	yes yes		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
86 EGFR	863 G	S yes	yes ye:		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
87 EGFR	864 A	E yes	yes ye		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
88 EGFR	864 A	T yes	yes ye		yes yes	yes	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
89 EGFR	866 E	K yes	yes ye	· ·	yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
90 EGFR	868 E	G yes	yes ye		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
91 EGFR	869 Y	C yes	yes yes	· ·	yes yes	yes	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
92 EGFR	871 A	G yes	yes ye	· ·	yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
93 EGFR	871 A	T yes	yes ye		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
94 EGFR	873 G	E yes	yes ye	· ·	yes yes	yes	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
95 EGFR	874 G	S yes	yes ye		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
96 EGFR	875 K	R yes	yes ye	· ·	yes yes	yes	yes	yes	·	yes	yes	1.0000	1.0000	1.0000
97 EGFR	884 E	K yes	yes ye		yes yes	yes	yes	yes		yes	yes	1.0000	1.0000	1.0000
98 EGFR	728 K	R yes	yes no		yes yes	yes	yes	yes	·	yes	yes	0.9091	0.9083	0.9101
99 EGFR	733 P	L yes	yes ye	es no	yes yes	yes	yes	yes	yes	yes	yes	0.9091	0.9092	0.9090
100 EGFR	745 K	R yes	yes no		yes yes	yes	yes	yes	yes	yes	yes	0.9091	0.9083	0.9101
101 EGFR	746 E	K yes	yes yes		yes yes	yes	yes	yes	yes	yes	yes	0.9091	0.9092	0.9090
102 EGFR	746 E	V yes	yes no		yes yes	yes	yes	yes	yes	yes	yes	0.9091	0.9083	0.9101
103 EGFR	752 S	P yes	yes yes	es no	yes yes	yes	yes	yes	yes v	yes	yes	0.9091	0.9092	0.9090
104 EGFR	752 S	Y yes	yes ye	es no	yes yes	yes	yes	yes	yes	yes	yes	0.9091	0.9092	0.9090
105 EGFR	754 K	R yes	yes no	yes	yes yes	yes	yes	yes	yes	yes	yes	0.9091	0.9083	0.9101
106 EGFR	758 E	D yes	yes ye:	es no	yes yes	yes	yes	yes	yes	yes	yes	0.9091	0.9092	0.9090
107 EGFR	767 A	V yes	yes ye:	es no	yes yes	yes	yes	yes	yes	yes	yes	0.9091	0.9092	0.9090
108 EGFR	768 S	C yes	yes yes	es no	yes yes	yes	yes	yes	yes	yes	yes	0.9091	0.9092	0.9090
109 EGFR	768 S	N yes	yes yes	es no	yes yes	yes	yes	yes	yes	yes	yes	0.9091	0.9092	0.9090
110 EGFR	783 T	l yes	yes yes		yes yes	yes	yes				yes	0.9091	0.9092	0.9090
		1'												

111 EGFR	784 S	D	yes	yes	VAS	no	yes	VAS	yes	yes	yes	yes	yes	yes	0.9091	0.9092	0.9090
112 EGFR	784 S	v	yes	yes	•	no	yes		yes	yes	,	•	yes	yes	0.9091	0.9092	0.9090
113 EGFR	785 T	- i	yes	yes	•	no	yes		yes	yes	,	•	yes	yes	0.9091	0.9092	0.9090
114 EGFR	785 T	N	yes	yes		no	yes		yes	yes		•	yes	yes	0.9091	0.9092	0.9090
115 EGFR	792 L	D	yes	yes	•	no	yes	,	yes	yes		•	yes	yes	0.9091	0.9092	0.9090
116 EGFR	803 R	ı	yes	yes	*	no	yes	,	yes	-	yes		yes	yes	0.9091	0.9092	0.9090
117 EGFR	804 E	G	1				_		1	yes	·	•		,	0.9091		0.9090
118 EGFR	806 K	4	yes	yes	,	no	yes	,	yes	yes			yes	yes	0.9091	0.9092 0.9083	0.9101
	806 K	A F	yes	yes		yes	yes	,	yes	yes		•	yes	yes	0.9091	0.9083	0.9101
119 EGFR 120 EGFR	811 S		yes	yes		yes	yes	,	yes	yes		•	yes	yes	0.9091	0.9083	0.9090
		P.	yes	yes	yes	no	yes		yes	yes	·	•	yes	yes			
121 EGFR	812 Q	K	yes	yes		yes	yes	,	yes	yes	-	•	yes	yes	0.9091	0.9083	0.9101
122 EGFR	818 C	Y -	yes	yes	*	no	yes	,	yes	yes		•	yes	yes	0.9091	0.9092	0.9090
123 EGFR	821 I	1	yes	yes		yes	yes	,	yes	yes		•	yes	yes	0.9091	0.9083	0.9101
124 EGFR	836 R	L	yes	yes	•	no	yes		yes	yes	·	•	yes	yes	0.9091	0.9092	0.9090
125 EGFR	841 R	K -	yes	yes	yes	no	yes	,	yes	yes		•	yes	yes	0.9091	0.9092	0.9090
126 EGFR	846 K	R	yes	yes		yes	yes	,	yes	yes		•	yes	yes	0.9091	0.9083	0.9101
127 EGFR	860 K	E	yes	yes		yes	yes	•	yes	yes	·	•	yes	yes	0.9091	0.9083	0.9101
128 EGFR	870 H	N	yes	yes		no	yes	,	yes	yes	yes	•	yes	yes	0.9091	0.9092	0.9090
129 EGFR	870 H	R	yes	yes		no	yes		yes	yes	-	•	yes	yes	0.9091	0.9092	0.9090
130 EGFR	715 I	L	yes	yes		no	yes		yes	yes	·	•	yes	yes	0.8182	0.8174	0.8190
131 EGFR	744 I	М	yes	yes		no	yes	,	yes	yes			yes	yes	0.8182	0.8174	0.8190
132 EGFR	744 I	V	yes	yes		no	yes	,	yes	yes		•	yes	yes	0.8182	0.8174	0.8190
133 EGFR	746 E	I	yes	yes		no	yes	,	yes	yes		•	yes	yes	0.8182	0.8174	0.8190
134 EGFR	748 R	K	yes	yes	no	no	yes	0.8182	0.8174	0.8190							
135 EGFR	759 I	N	yes	yes	no	no	yes	,	yes	yes	yes	yes	yes	yes	0.8182	0.8174	0.8190
136 EGFR	759 I	V	yes	yes	no	no	yes	0.8182	0.8174	0.8190							
137 EGFR	773 H	R	yes	yes	no	no	yes	0.8182	0.8174	0.8190							
138 EGFR	787 Q	R	yes	yes	no	no	yes	0.8182	0.8174	0.8190							
139 EGFR	795 F	S	yes	yes	no	no	yes	0.8182	0.8174	0.8190							
140 EGFR	805 H	I	yes	yes	no	no	yes	0.8182	0.8174	0.8190							
141 EGFR	807 D	N	yes	yes	no	no	yes	0.8182	0.8174	0.8190							
142 EGFR	921 S	R	yes	yes	no	no	yes	0.8182	0.8174	0.8190							
143 EGFR	712 F	S	yes	yes	yes	yes	no	0.2727	0.2755	0.2751							
144 EGFR	798 L	Н	yes	yes	yes	yes	no	0.2727	0.2755	0.2751							
145 EGFR	826 N	S	yes	yes	yes	yes	no	0.2727	0.2755	0.2751							
146 EGFR	834 V	Α	yes	yes	yes	yes	no	0.2727	0.2755	0.2751							
147 EGFR	845 V	Α	yes	yes	yes	yes	no	0.2727	0.2755	0.2751							
148 EGFR	851 V	Α	yes	yes	yes	yes	no	0.2727	0.2755	0.2751							
149 EGFR	712 F	L	yes	yes	no	yes	no	0.1818	0.1837	0.1852							
150 EGFR	717 V	A	yes	yes	no	yes	no	0.1818	0.1837	0.1852							
151 EGFR	726 V	М	yes	yes	no	yes	no	0.1818	0.1837	0.1852							
152 EGFR	727 Y	Н	yes	yes	no	yes	no	0.1818	0.1837	0.1852							
153 EGFR	741 P	Н	yes	yes	yes	no	0.1818	0.1846	0.1841								
154 EGFR	742 V	A	yes	yes	no	yes	no	0.1818	0.1837	0.1852							
155 EGFR	750 A	Р	yes	yes	yes	no	0.1818	0.1846	0.1841								
156 EGFR	769 V	L	yes	yes	no	yes	no	0.1818	0.1837	0.1852							
157 EGFR	775 C	Υ	yes	yes	yes	no	0.1818	0.1846	0.1841								
158 EGFR	801 Y	Н	yes	yes	no	yes	no	0.1818	0.1837	0.1852							
159 EGFR	802 V	F	yes	yes		yes	no	0.1818	0.1837	0.1852							
160 EGFR	802 V	I	yes	yes			no	0.1818	0.1837	0.1852							
161 EGFR	819 V	А	yes	yes			no		no	no			no	no	0.1818		0.1852
162 EGFR	834 V	L	yes	yes			no		no	no			no	no	0.1818		0.1852
163 EGFR	834 V	М	yes	yes		•	no		no	no			no	no	0.1818		0.1852
164 EGFR	843 V	l I	yes	yes			no		no	no			no	no	0.1818		0.1852
165 EGFR	851 V	l I	yes	yes			no		no	no			no	no	0.1818		0.1852
166 EGFR	897 V	lı .	yes	yes			no		no	no			no	no	0.1818		0.1852
,	1-	1.	17	,,	-	.,				1		-			,	2.100,	3.1002

| 16 | 7 EGFR | 738 | V | G | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |
|----|--------|-----|---|---|-----|-----|----|----|----|----|----|----|----|----|----|----|--------|--------|--------|
| 16 | B EGFR | 765 | V | M | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |
| 16 | 9 EGFR | 769 | V | M | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |
| 17 | EGFR | 774 | V | L | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |
| 17 | 1 EGFR | 774 | V | M | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |
| 17 | 2 EGFR | 776 | R | Н | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |
| 17 | B EGFR | 786 | V | M | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |
| 17 | 4 EGFR | 813 | Υ | С | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |
| 17 | 5 EGFR | 831 | R | Н | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |
| 17 | 6 EGFR | 832 | R | Н | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |
| 17 | 7 EGFR | 845 | V | М | yes | yes | no | 0.0909 | 0.0929 | 0.0942 |