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COMPOUND 1		COMPOUND 2		COMPOUND 3	
COMPOUND 4		COMPOUND 5		COMPOUND 6	
COMPOUND 7		COMPOUND 8		COMPOUND 9	

FIG. 1

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COMPOUND 10	COMPOUND 11	COMPOUND 12
COMPOUND 13	COMPOUND 14	COMPOUND 15
COMPOUND 16	COMPOUND 17	COMPOUND 18

FIG. 1
(CONT.-1)

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COMPOUND 19	COMPOUND 20	COMPOUND 21
COMPOUND 22	COMPOUND 23	COMPOUND 24
COMPOUND 25	COMPOUND 26	COMPOUND 27

FIG. 1
(CONT.-2)

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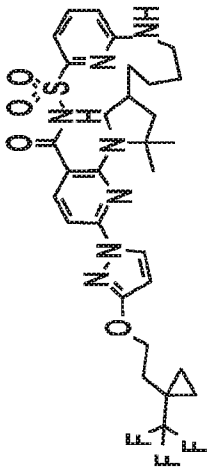
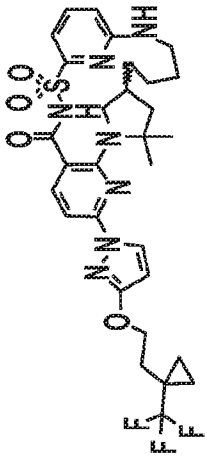
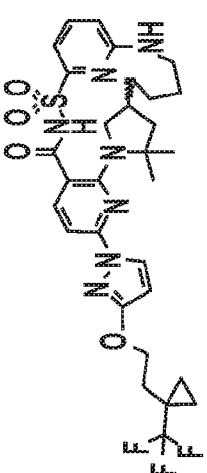
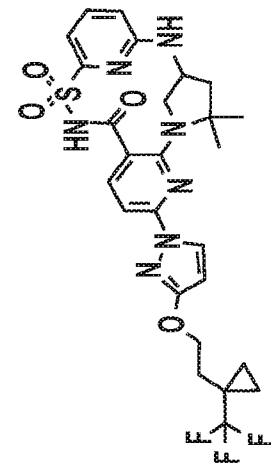
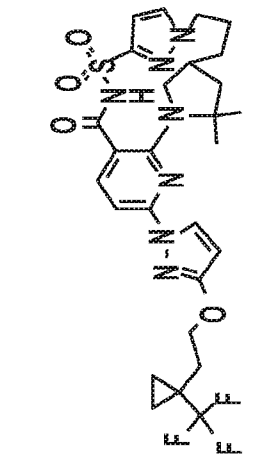
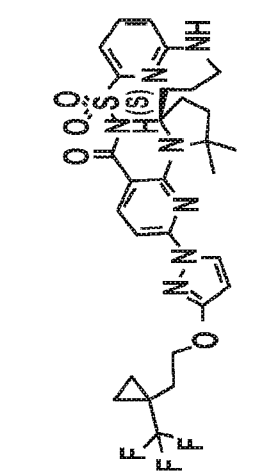
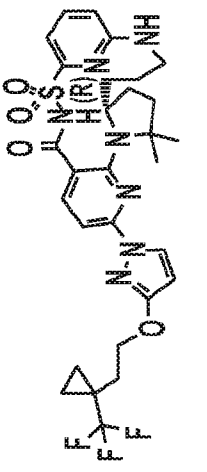
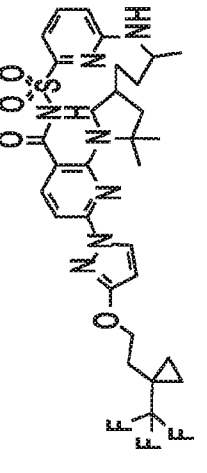
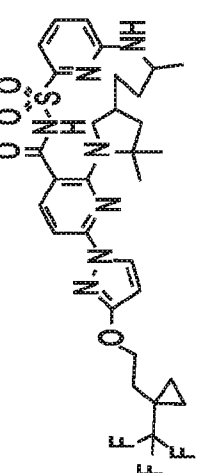
COMPOUND 28		COMPOUND 29		COMPOUND 30	
COMPOUND 31		COMPOUND 32		COMPOUND 33	
COMPOUND 34		COMPOUND 35		COMPOUND 36	

FIG. 1
(CONT.-3)

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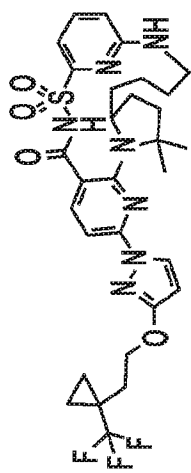
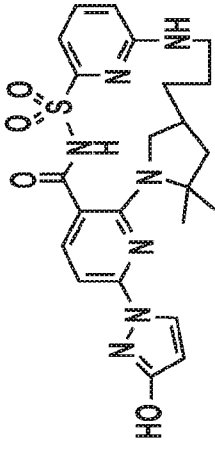
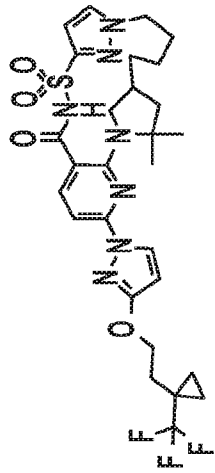
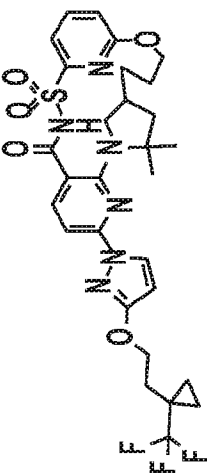
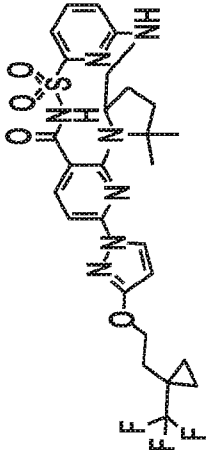
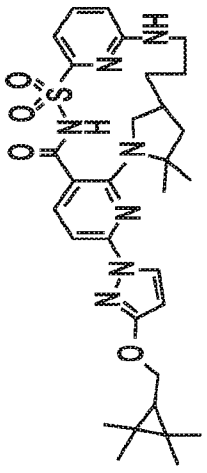
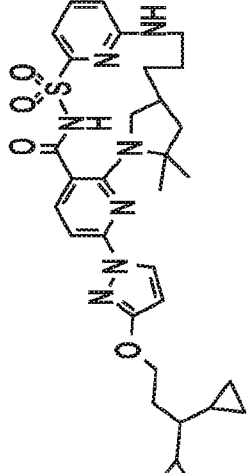
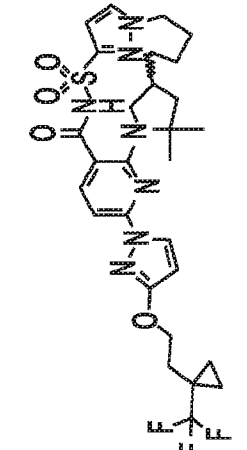
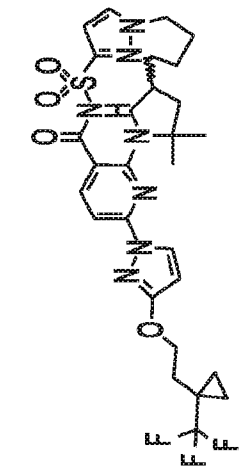
COMPOUND 37		COMPOUND 38		COMPOUND 39	
COMPOUND 40		COMPOUND 41		COMPOUND 42	
COMPOUND 43		COMPOUND 44		COMPOUND 45	

FIG. 1
(CONT.-4)

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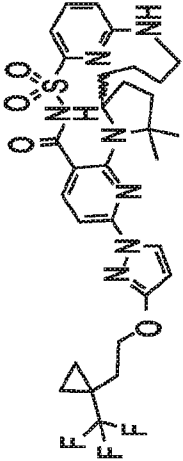
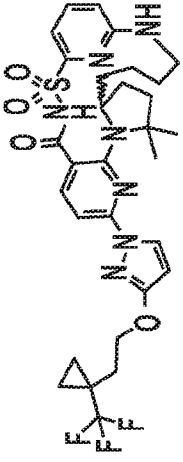
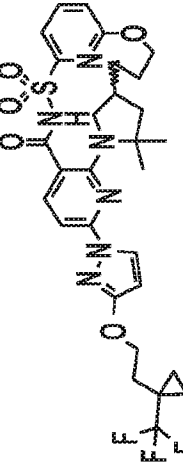
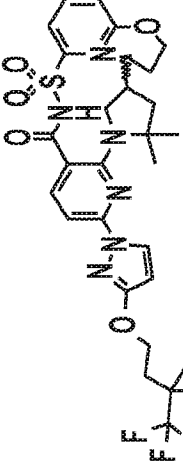
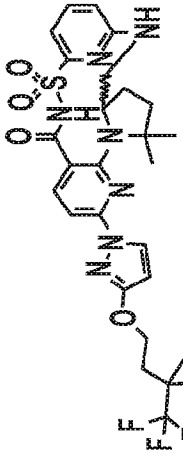
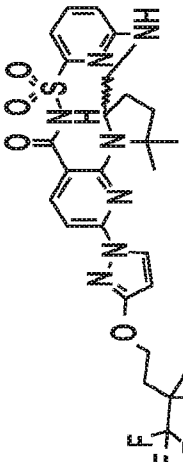
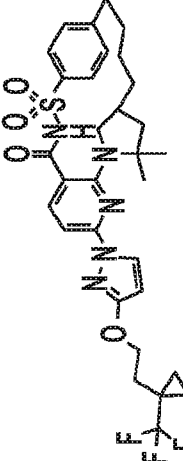
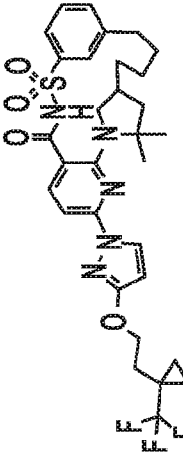
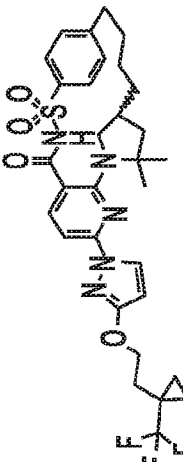
COMPOUND 46	COMPOUND 47	COMPOUND 48
		
COMPOUND 49	COMPOUND 50	COMPOUND 51
		
COMPOUND 52	COMPOUND 53	COMPOUND 54
		

FIG. 1
(CONT.-5)

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COMPOUND 55	COMPOUND 56	COMPOUND 57
COMPOUND 58	COMPOUND 59	COMPOUND 60
COMPOUND 61	COMPOUND 62	COMPOUND 63

FIG. 1
(CONT.-6)

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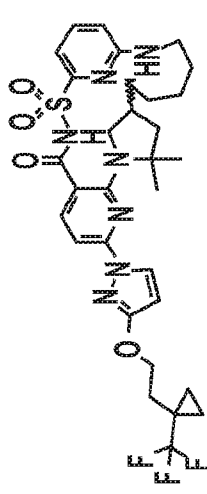
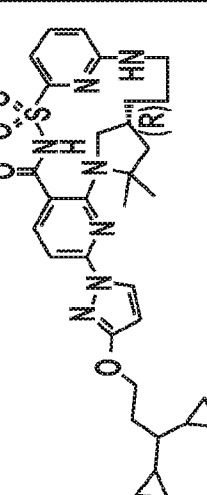
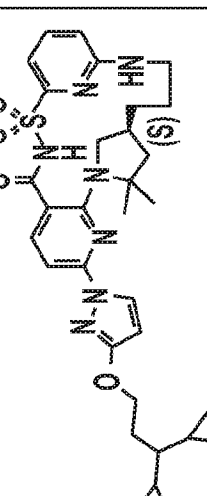
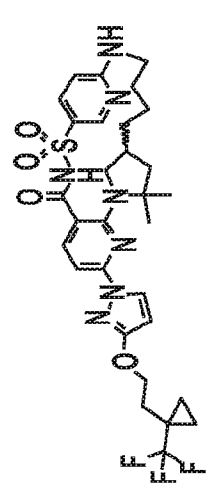
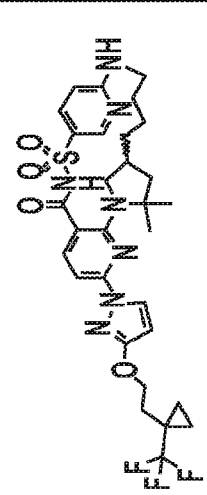
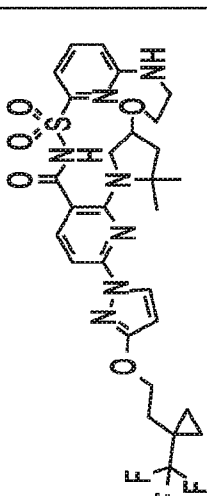
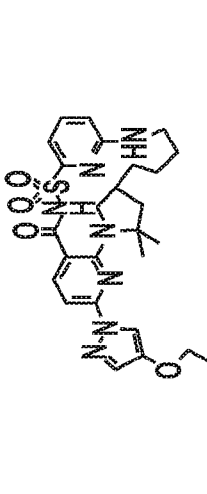
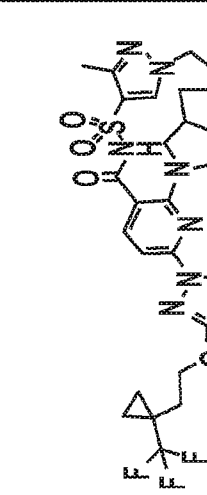
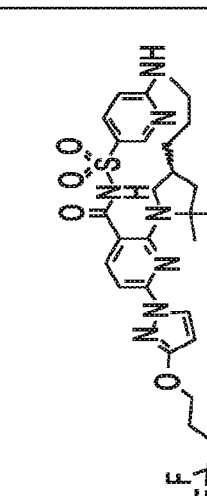
COMPOUND 64	
COMPOUND 65	
COMPOUND 66	
COMPOUND 67	
COMPOUND 68	
COMPOUND 69	
COMPOUND 70	
COMPOUND 71	
COMPOUND 72	

FIG. 1
(CONT.-7)

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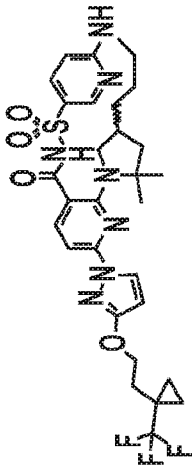
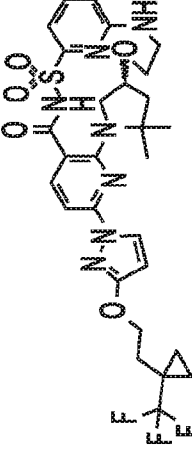
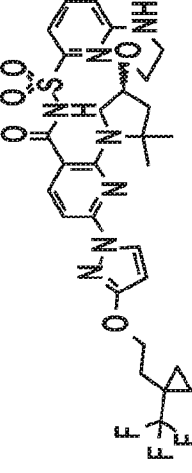
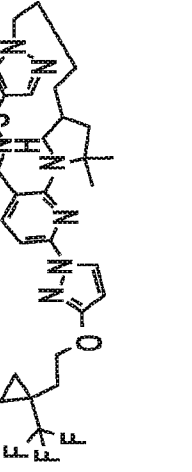
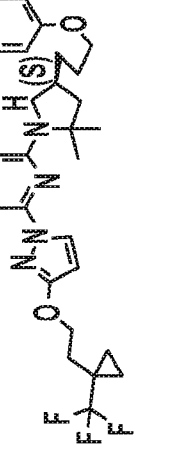
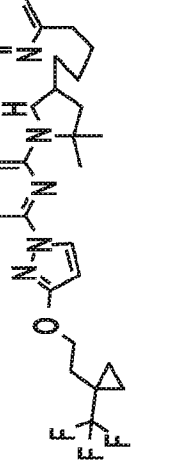
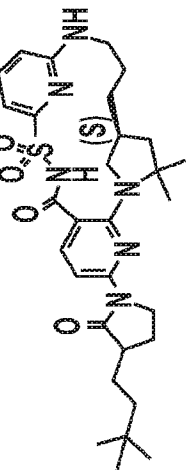
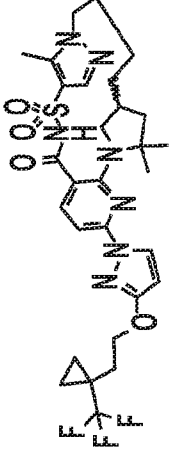
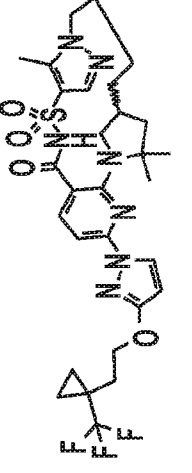
COMPOUND 73	COMPOUND 74	COMPOUND 75
		
COMPOUND 76	COMPOUND 77	COMPOUND 78
		
COMPOUND 79	COMPOUND 80	COMPOUND 81
		

FIG. 1
(CONT.-8)

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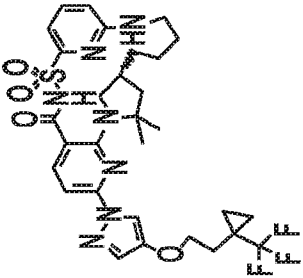
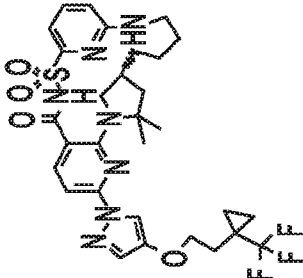
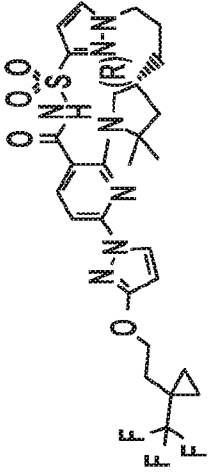
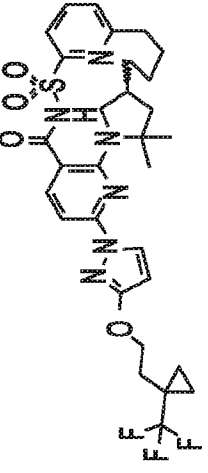
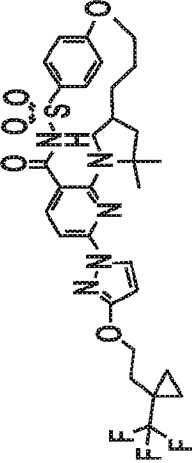
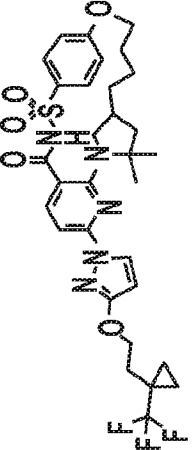
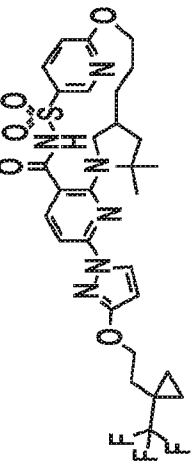
COMPOUND 82	COMPOUND 83	COMPOUND 84
		
COMPOUND 85	COMPOUND 86	COMPOUND 87
		
COMPOUND 88	COMPOUND 89	COMPOUND 90
		

FIG. 1
(CONT.-9)

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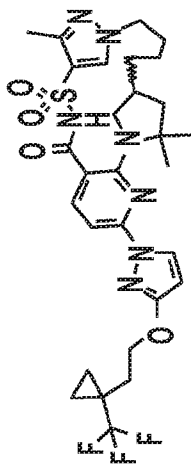
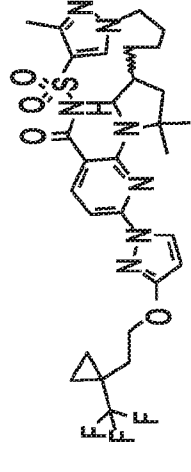
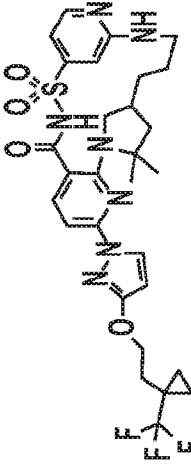
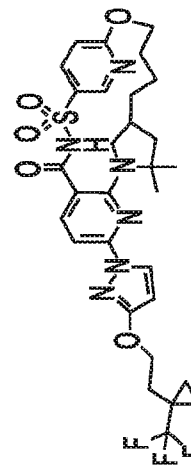
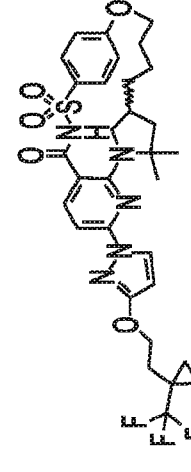
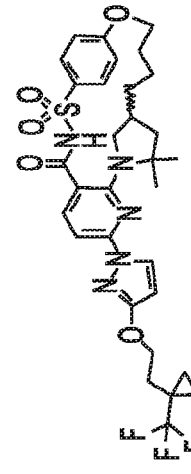
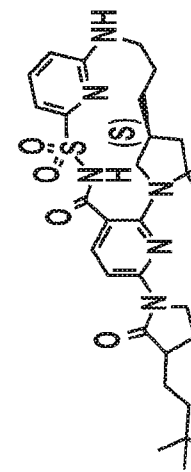
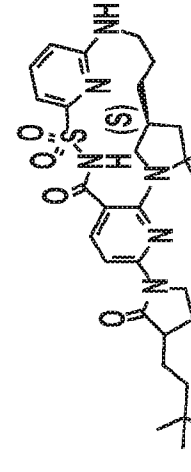
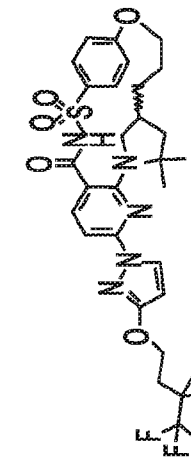
COMPOUND 91	
COMPOUND 92	
COMPOUND 93	
COMPOUND 94	
COMPOUND 95	
COMPOUND 96	
COMPOUND 97	
COMPOUND 98	
COMPOUND 99	

FIG. 1
(CONT.-10)

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COMPOUND 100	COMPOUND 101	COMPOUND 102
COMPOUND 103	COMPOUND 104	COMPOUND 105
COMPOUND 106	COMPOUND 107	COMPOUND 108

FIG. 1
(CONT.-11)

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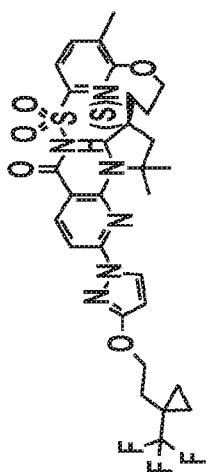
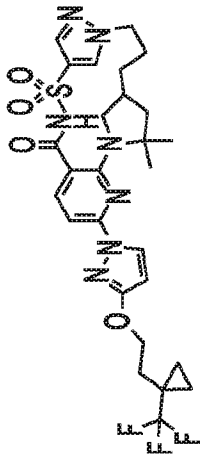
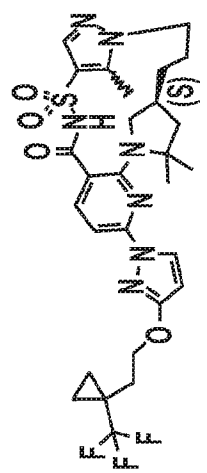
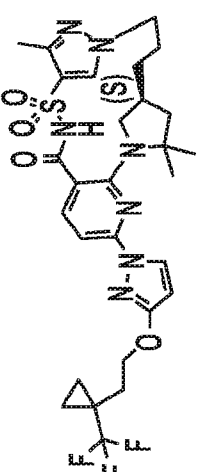
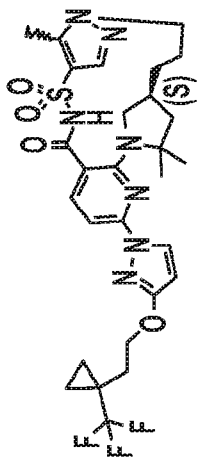
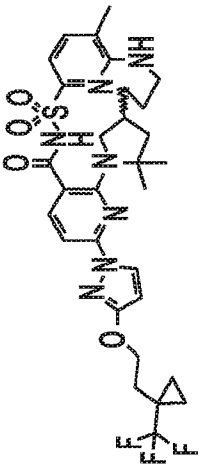
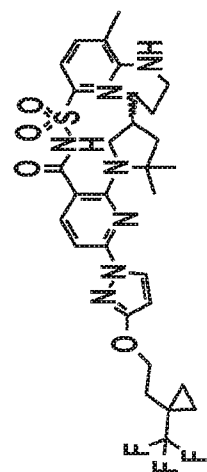
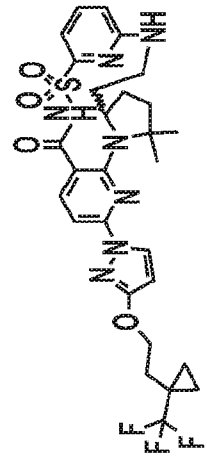
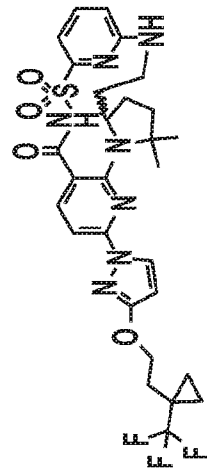
COMPOUND 109	COMPOUND 110	COMPOUND 111
		
COMPOUND 112	COMPOUND 113	COMPOUND 114
		
COMPOUND 115	COMPOUND 116	COMPOUND 117
		

FIG. 1
(CONT.-12)

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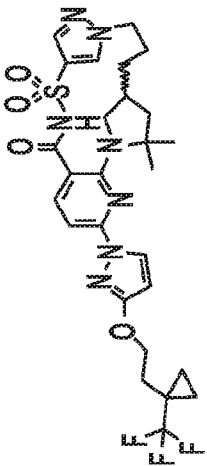
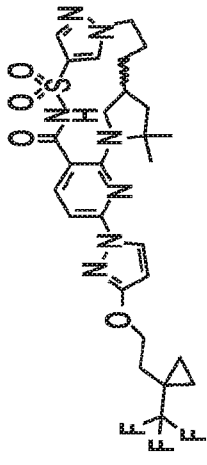
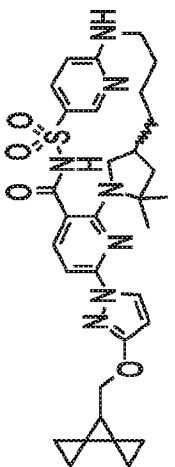
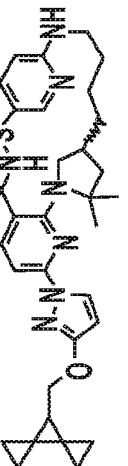
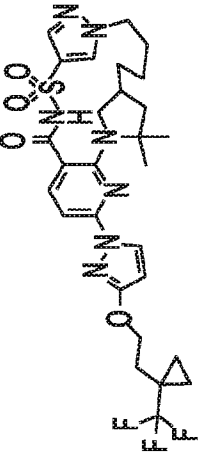
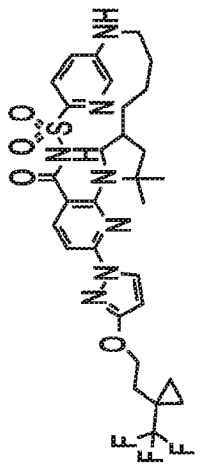
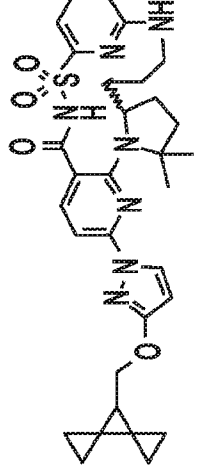
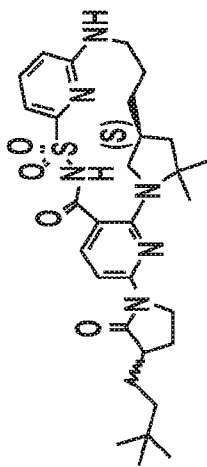
COMPOUND 118	COMPOUND 119	COMPOUND 120
		
COMPOUND 121	COMPOUND 122	COMPOUND 123
		
COMPOUND 124	COMPOUND 125	COMPOUND 126
		

FIG. 1
(CONT.-13)

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COMPOUND 127	COMPOUND 128	COMPOUND 129
COMPOUND 130	COMPOUND 131	COMPOUND 132
COMPOUND 133	COMPOUND 134	COMPOUND 135

FIG. 1
(CONT.-14)

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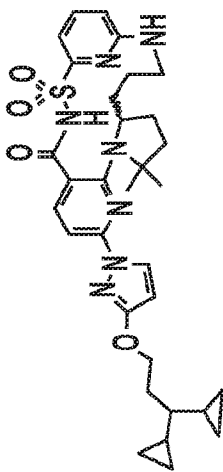
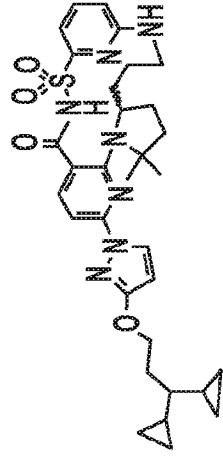
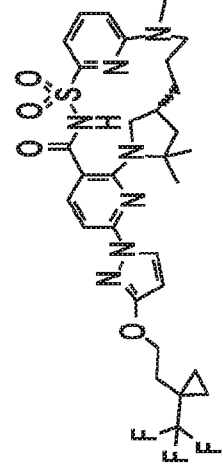
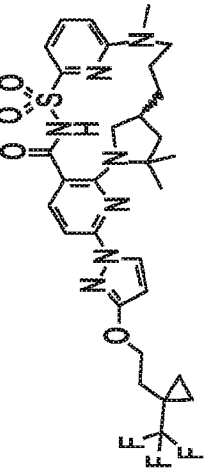
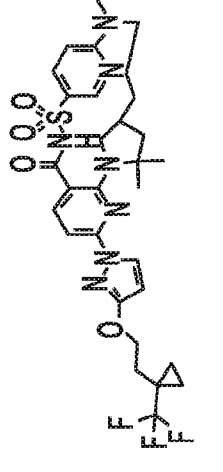
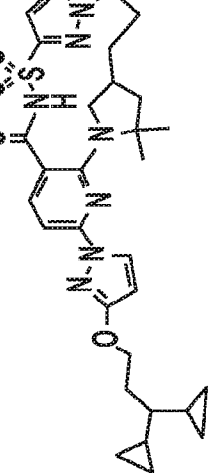
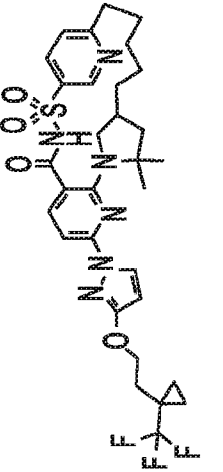
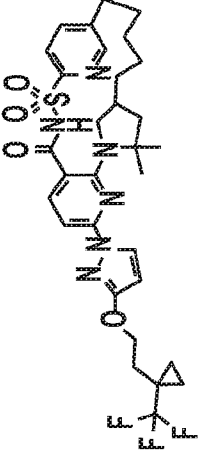
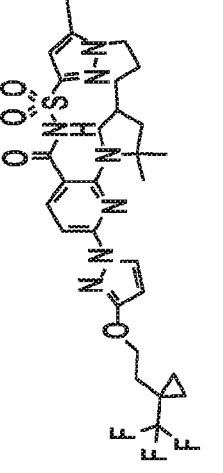
COMPOUND 136	COMPOUND 137	COMPOUND 138
		
COMPOUND 139	COMPOUND 140	COMPOUND 141
		
COMPOUND 142	COMPOUND 143	COMPOUND 144
		

FIG. 1
(CONT.-15)

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COMPOUND 145	COMPOUND 146	COMPOUND 147
COMPOUND 148	COMPOUND 149	COMPOUND 150
COMPOUND 151	COMPOUND 152	COMPOUND 153

FIG. 1
(CONT.-16)

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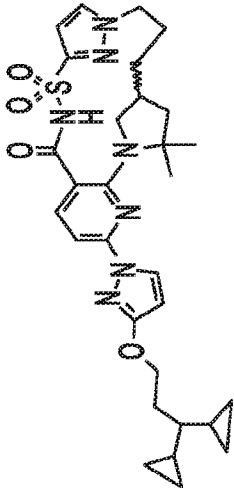
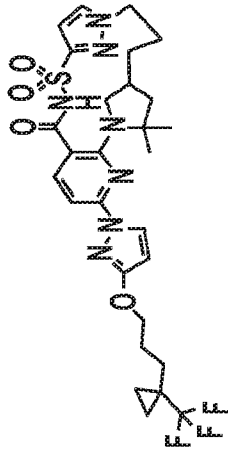
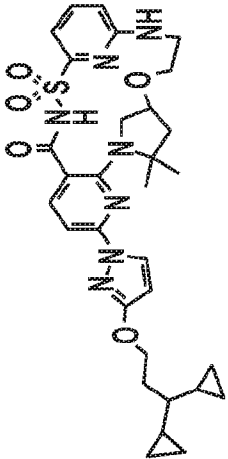
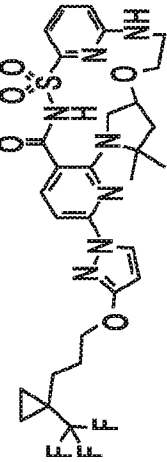
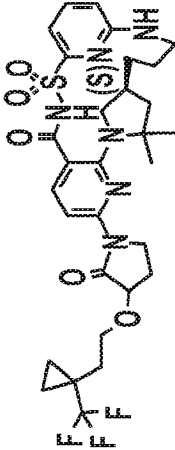
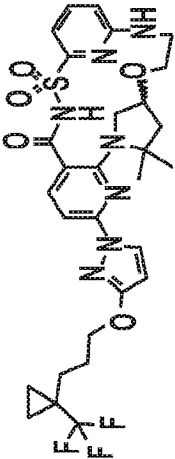
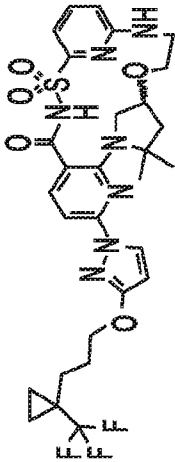
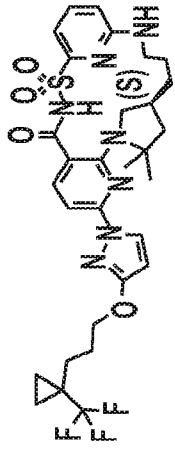
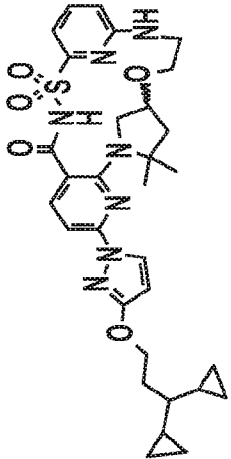
COMPOUND 154	COMPOUND 155	COMPOUND 156
		
COMPOUND 157	COMPOUND 158	COMPOUND 159
		
COMPOUND 160	COMPOUND 161	COMPOUND 162
		

FIG. 1
(CONT.-17)

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COMPOUND 163	COMPOUND 164	COMPOUND 165
COMPOUND 166	COMPOUND 167	COMPOUND 168
COMPOUND 169	COMPOUND 170	COMPOUND 171

FIG. 1
(CONT.-18)

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COMPOUND 172	COMPOUND 173	COMPOUND 174
COMPOUND 175	COMPOUND 176	COMPOUND 177
COMPOUND 178	COMPOUND 179	COMPOUND 180

FIG. 1
(CONT.-19)

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COMPOUND 181	COMPOUND 182	COMPOUND 183
COMPOUND 184	COMPOUND 185	COMPOUND 186
COMPOUND 187	COMPOUND 188	COMPOUND 189

FIG. 1
(CONT.-20)

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COMPOUND 190	COMPOUND 191	COMPOUND 192
COMPOUND 193	COMPOUND 194	COMPOUND 195
COMPOUND 196	COMPOUND 197	COMPOUND 198

FIG. 1
(CONT.-21)

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COMPOUND 199	COMPOUND 200	COMPOUND 201
COMPOUND 202	COMPOUND 203	COMPOUND 204
COMPOUND 205	COMPOUND 206	COMPOUND 207

FIG. 1
(CONT.-22)

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COMPOUND 208	COMPOUND 209	COMPOUND 210
COMPOUND 211	COMPOUND 212	COMPOUND 213
COMPOUND 214	COMPOUND 215	COMPOUND 216

FIG. 1
(CONT.-23)

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COMPOUND 217	COMPOUND 218	COMPOUND 219
COMPOUND 220	COMPOUND 221	COMPOUND 222
COMPOUND 223	COMPOUND 224	COMPOUND 225

FIG. 1
(CONT.-24)

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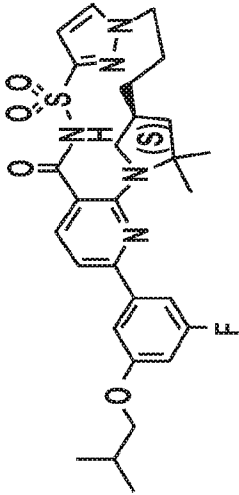
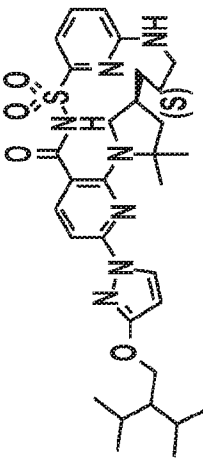
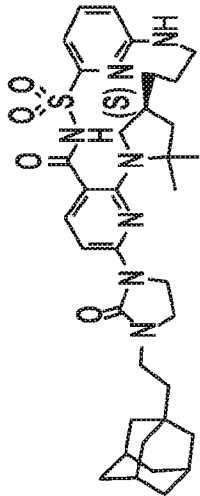
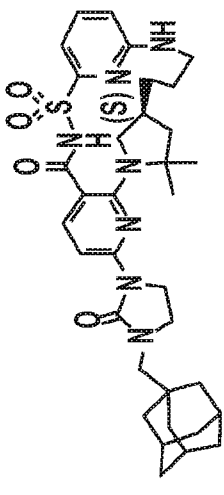
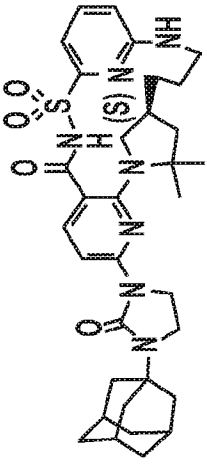
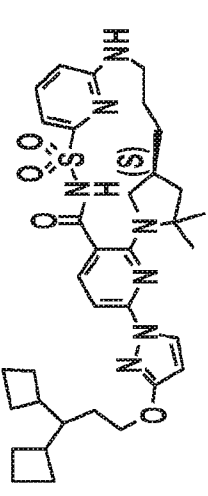
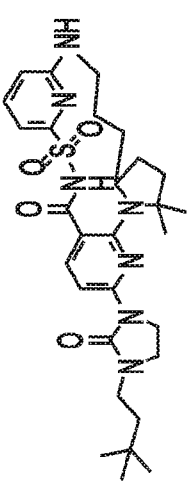
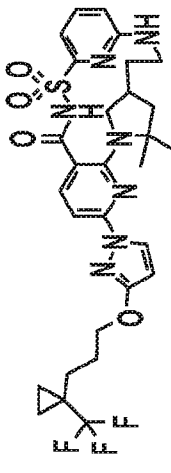
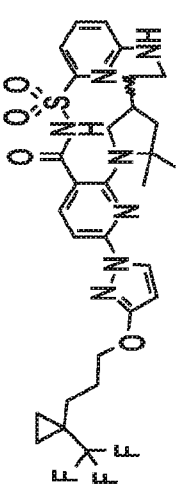
COMPOUND 226	COMPOUND 227	COMPOUND 228
		
COMPOUND 229	COMPOUND 230	COMPOUND 231
		
COMPOUND 232	COMPOUND 233	COMPOUND 234
		

FIG. 1
(CONT.-25)

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COMPOUND 235	COMPOUND 236	COMPOUND 237
COMPOUND 238	COMPOUND 239	COMPOUND 240
COMPOUND 241	COMPOUND 242	COMPOUND 243

FIG. 1
(CONT.-26)

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COMPOUND 244		COMPOUND 245		COMPOUND 246	
COMPOUND 247		COMPOUND 248		COMPOUND 249	
COMPOUND 250		COMPOUND 251		COMPOUND 252	

FIG. 1
(CONT.-27)

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COMPOUND 253	COMPOUND 254	COMPOUND 255
COMPOUND 256	COMPOUND 257	COMPOUND 258
COMPOUND 259	COMPOUND 260	COMPOUND 261

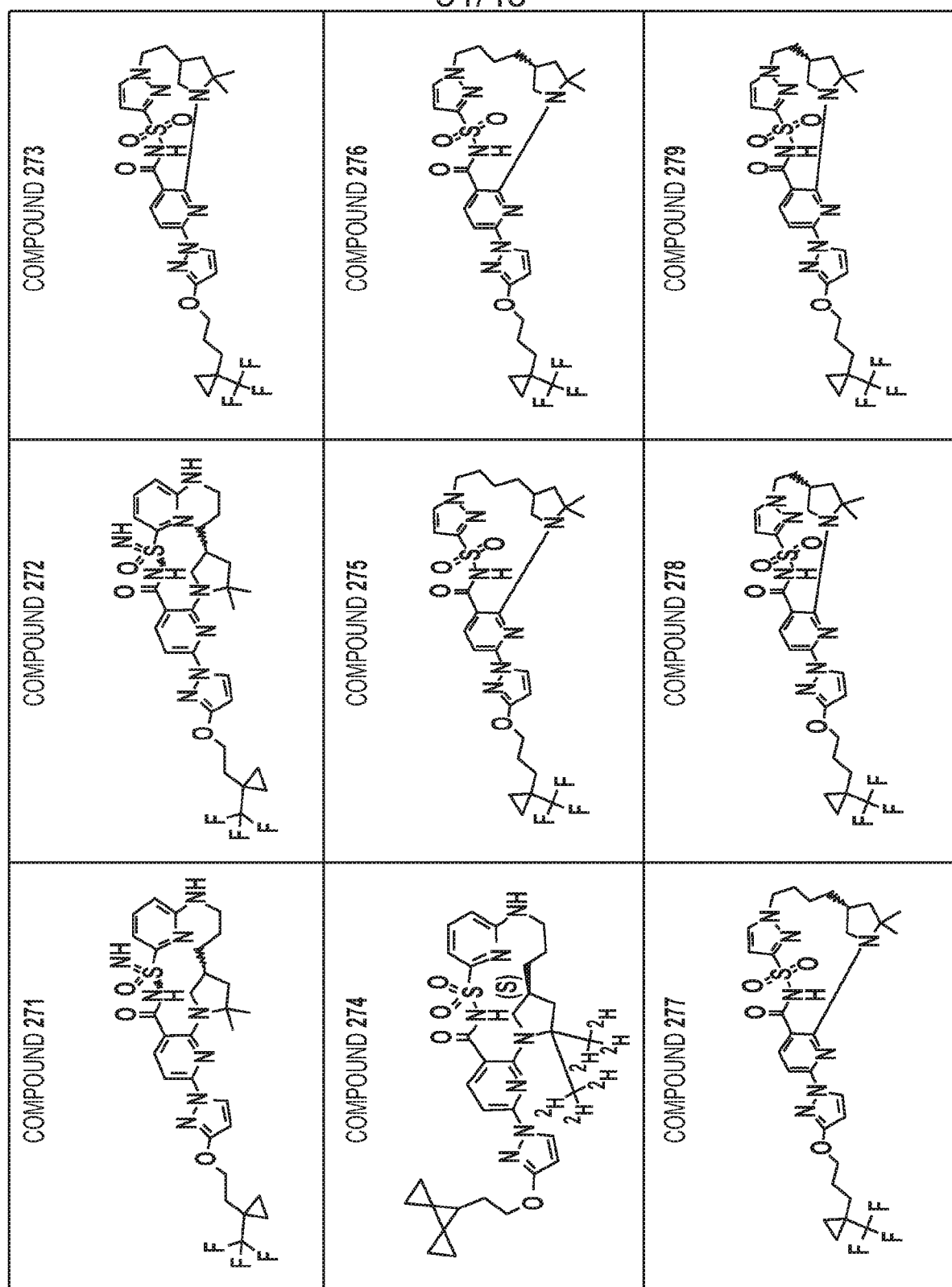
FIG. 1
(CONT.-28)

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COMPOUND 262	COMPOUND 263	COMPOUND 264
COMPOUND 265	COMPOUND 266	COMPOUND 267
COMPOUND 268	COMPOUND 269	COMPOUND 270

FIG. 1
(CONT.-29)

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**FIG. 1**
(CONT.-30)

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COMPOUND 280	COMPOUND 281	COMPOUND 282
COMPOUND 283	COMPOUND 284	COMPOUND 285
COMPOUND 286	COMPOUND 287	COMPOUND 288

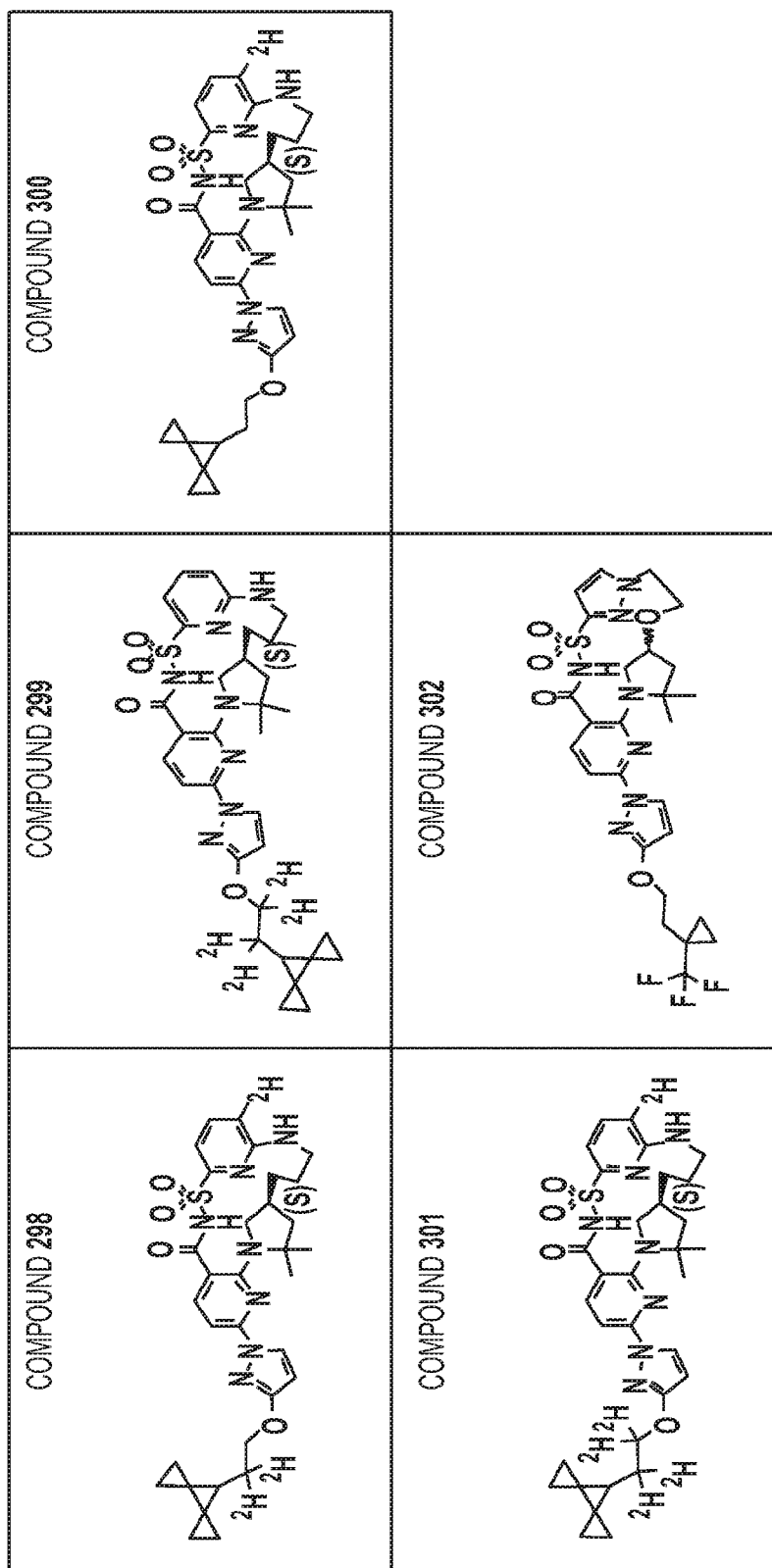
FIG. 1
(CONT.-31)

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COMPOUND 289	COMPOUND 290	COMPOUND 291
COMPOUND 292	COMPOUND 293	COMPOUND 294
COMPOUND 295	COMPOUND 296	COMPOUND 297

FIG. 1
(CONT.-32)

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**FIG. 1**
(CONT.-33)

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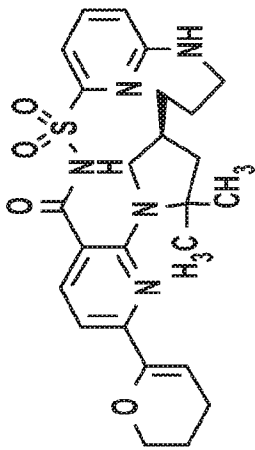
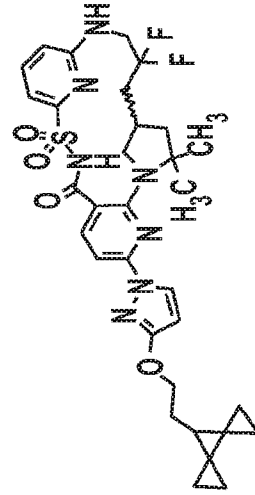
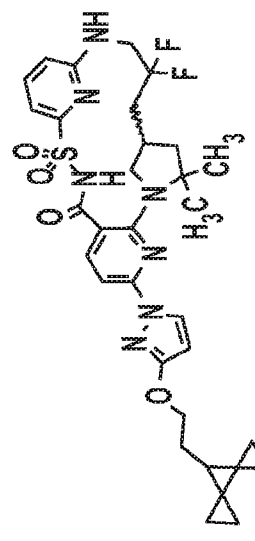
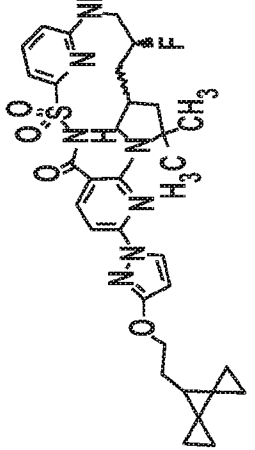
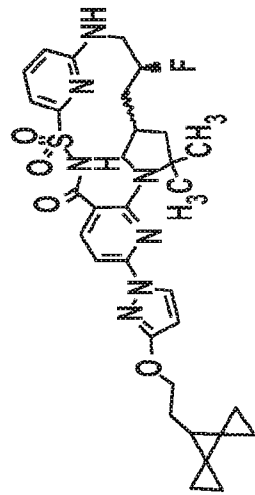
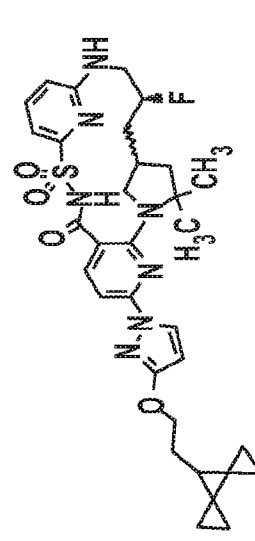
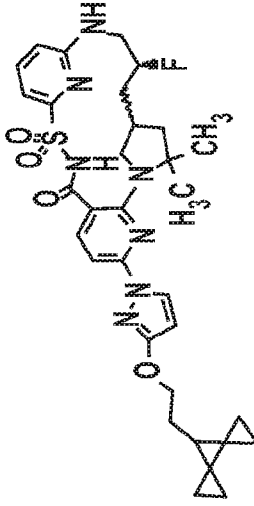
COMPOUND 303	
COMPOUND 304	
COMPOUND 305	
COMPOUND 306	
COMPOUND 307	
COMPOUND 308	
COMPOUND 309	

FIG. 1
(CONT.-34)

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MUTATION cDNA NAME	MUTATION PROTEIN NAME	MUTATION LEGACY NAME
c1A>G		M1V
c.54- 5940_273+10250de l21kb	pSer18ArgfsX16	CFTRdele2,3
c.91C>T	p.Arg31Cys	R31C
c.115C>T	p.Gln39X	Q39X
c.137C>A	p.Ala46Asp	A46D
c.165-1G>A	No protein name	297-1G->A
c.166G>A	p.Glu56Lys	E56K
c.174_175insA	p.Arg59LysfsX10	306insA
c.178G>T	p.Glu60X	E60X
c.200C>T	p.Pro67Leu	P67L
c.220C>T	p.Arg74Trp	R74W
c.223C>T	p.Arg75X	R75X
c.224G>A	p.Arg75Gln	R75Q
c.254G>A	p.Gly85Glu	G85E
c.262_263delTT	p.Leu88IlefsX22	394delTT
c.273+1G>A	No protein name	405+1G->A
c.274-1G>A	No protein name	406-1G->A
c.274G>A	p.Glu92Lys	E92K
c.274G>T	p.Glu92X	E92X
c.292C>T	p.Gln98X	Q98X
c.313delA	p.Ile105SerfsX2	444delA
c.325_327delTATins G	p.Tyr109GlyfsX4	457TAT->G
c.328G>C	p.Asp110His	D110H
c.349C>T	p.Arg117Cys	R117C
c.350G>A	p.Arg117His	R117H
c.366T>A	p.Tyr122X	Y122X
c.442delA	p.Ile148LeufsX5	574delA
c.443T>C	p.Ile148Thr	I148T

FIG. 2

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MUTATION cDNA NAME	MUTATION PROTEIN NAME	MUTATION LEGACY NAME
c.489+1G>T	No protein name	621+1G->T
c.531delT	p.Ile177MetfsX12	663delT
c.532G>A	p.Gly178Glu	G178R
c.543_546delTAGT	p.Leu183PhefsX5	675del4
c.579+1G>T	No protein name	711+1G->T
c.579+3A>G	No protein name	711+3A->G
c.579+5G>A	No protein name	711+5G->A
c.580-1G>T	No protein name	712-1G->T
C.59SOT	p.His199Tyr	H199Y
C.613CM	p.Pro205Ser	P205S
c.617T>G	p.Leu20GTrp	L206W
C.6580T	p.Gln220X	Q220X
c.580T>G	p.Leu227Arg	L227R
c.720_741delAGGG AGAATGATGATGAA GTAC	p.Gly241GlufsX13	852del22
c.828C>A	p.Cys276X	C276X
c.948delT	p.Phe316LeufsX1 2	1078delT
c.988G>T	p.Gly330X	G330X
c.1000C>T	p.Arg334Trp	R334W
c.1007T>A	p.Ile336Lys	I336K
c.1013C>T	p.Thr338Ele	T338I
c.1021T>C	p.Ser341Pro	S341P
c.1022_1023insTC	p.Phe342HisfsX2 8	1154insTC
c.1040G>A	p.Arg347His	R347H
c.1040G>C	p.Arg347Pro	R347P
c.1055G>A	p.Arg352Gln	R352Q
c.[1075C>A; 10799C> A]	p.[Gln359Lys;Thr 360Lys]	Q359K/T360K
c.1081delT	p.Trp361GlyfsX8	1213delT

FIG. 2 (CONT.-1)

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MUTATION cDNA NAME	MUTATION PROTEIN NAME	MUTATION LEGACY NAME
c.1116+1G>A	No protein name	1248+1G->A
c.1127_1128insA	p.Gln378AlafsX4	1259insA
c.1153_1154insAT	p.Asn386lfeX3	1288insTA
c.1202G>A or c.1203G>A	p.Trp401X	W401X
c.1209+1G>A	No protein name	1341+1G->A
c.1210-12[5]	No protein name	5T
c.1210-12[7]	No protein name	7T
c.1240C>T	p.Gln414X	Q414X
c.1329_1330insAGA T	p.Ile444ArgfsX3	1461ins4
c.1340delA	p.Lys447ArgfsX2	1471delA
c.1364C>A	p.Ala455Glu	A455E
c.1393-1G>A	No protein name	1525-1G->A
c.1397C>A or c.1397C>G	p.Ser466X	S466X
c.1400T>C	p.Leu467Pro	L467P
c.1408A>G	p.Met470Val	M470V
c.1418delG	p.Gly473GlufsX54	1548delG
c.1466C>A	p.Ser489X	S489X
c.1475C>T	p.Ser492Phe	S492F
c.1477C>T	p.Gln493X	Q493X
c.1519_1521delATC	p.Ile507del	I507del
c.1521_1523delCTT	p.Phe508del	F508del
c.1545_1546delTA	p.Tyr515X	1677delTA
c.1558G>T	p.Val520Phe	V520F
c.1573C>T	p.Gln525X	Q525X
c.1585-8G>A	No protein name	1717-8G->A
c.1585-1G>A	No protein name	1717-1G->A
c.1624G>T	p.Gly542X	G542X

FIG. 2 (CONT.-2)

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MUTATION cDNA NAME	MUTATION PROTEIN NAME	MUTATION LEGACY NAME
c.1645A>C or c.1G47T>G	p.Ser549Arg	S549R
c.1646G>A	p.Ser549Asn	S549N
c.1650delA	p.Gly551ValfsX8	1782delA
c.1651G>A	p.Gly551Ser	G551S
c.1652G>A	p.Gly551Asp	G551D
c.1654C>T	p.Gln552X	Q552X
c.1657C>T	p.Arg553X	R553X
c.1673T>C	p.Leu558Ser	L558S
c.1675G>A	p.Ala559Thr	A559T
c.1679G>A	p.Arg560Lys	R560K
c.1679G>C	p.Arg560Thr	R560T
c.1679+1G>C	No protein name	1811+1G->C
c.1679+1.6kbA>G	No protein name	1811+1.6kbA->G
c.1680-1G>A	No protein name	1812-1G->A
c.1682C>A	p.Ala561Glu	A561E
c.1692delA	p.Asp565MetfsX7	1824delA
c.1705T>G	p.Tyr569Asp	Y569D
c.1727G>C	p.Gly576Ala	G576A
c.1736A>G	p.Asp579Gly	D579G
c.1753G>T	p.Glu585X	E585X
c.1766+1G>A	No protein name	1898+1G->A
c.1766+1G>C	No protein name	1898+1G->C
c.1766+3A>G	No protein name	1898+3A->G
c.1841A>G	p.Asp614Gly	D614G
c.1923_1931del9ins	p.Ser641ArgfsX5	2055del9->A
c.1973_1985del13ins sAGAAA	p.Arg658 LysfsX4	2105- 2117del13insA GAAA

FIG. 2 (CONT.-3)

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MUTATION cDNA NAME	MUTATION PROTEIN NAME	MUTATION LEGACY NAME
c.1986_1989delAAC T	p.Thr663ArgfsX8	2118del4
c.2002C>T	p.Arg668Cys	R668C
c.2012delT	p.Leu671X	2143delT
c.2051_2052delAAi nsG	p.Lys684SerfsX38	2183AA->G+
c.2051_2052delAAi nsGnsG	p.Lys684SerfsX38	2183delAA->G#
c.2052_2053insA	p.Gln685ThrfsX4	2184insA
c.2052delA	p.Lys684AsnfsX38	2184delA
c.21250T	p.Arg709X	R709X
c.2128A>T	p.Lys710X	K710X
c.2175_2176insA	p.Glu726ArgfsX4	2307insA
c.2195T>G	p.Leu732X	L732X
c.2215delG	p.Val739TyrfsX16	2347delG
c.2260G>A	p.Val754Met	V754M
c.2290C>T	p.Arg764X	R764X
c.2353C>T	p.Arg785X	R785X
c.2374C>T	p.Arg792X	R792X
c.2424_2425insAT	p.Ser809IlefsX13	2556insAT
c.2453delT	p.Leu818TrpfsX3	2585delT
c.2462_2463delGT	p.Ser821ArgfsX4	No legacy name
c.2464G>T	p.Glu822X	E822X
c.2490+1G>A	No protein name	2622+1G->A
c.2491G>T	p.Glu831X	E831X
c.2537G>A or c.2538G>A	p.Trp846X	W846X
c.25470A	p.Tyr849X	Y849X
c.2551C>T	p.Arg851X	R851X
c.2583delT	p.Phe861LeufsX3	2711delT
c.2657+2_2657+3in sA	No protein name	2789+2insA

FIG. 2 (CONT.-4)

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MUTATION cDNA NAME	MUTATION PROTEIN NAME	MUTATION LEGACY NAME
c.2657+5G>A	No protein name	2789+5G->A
c.2658-IG>C	No protein name	2790-IG->C
c.2668C>T	p.Gln890X	Q890X
c.2735C>A	p.Ser912X	S912X
c.2737_2738insG		2869insG
c.2739T>A	p.Tyr913X	Y913X
c.2764_2765insAG	p.Val922GlufsX2	2896insAG
c.2780T>C	p.Leu927Pro	L927P
c.2834C>T	p.Ser945Leu	S945L
c.2875delG	p.Ala959HisfsX9	3007delG
c.2908G>C	p.Gly970Arg	G970R
c.2930C>T	p.Ser977Phe	S977F
c.2988G>A	No protein name	3120G->A
c.2988+1G>A	No protein name	3120+1G->A
c.2989- 977_3367+248del	No protein name	3121- 977_3499+248 del2515
c.2989-1G>A	No protein name	3121-1G->A
c.2991G>C	p.Leu997Phe	L997F
c.3002_3003delTG	p.Val1001AspfsX4 5	3132delTG
c.3080T>C	p.Ile1027Thr	I1027T
c.3140-26A>G	No protein name	3272-26A->G
c.3154T>G	p.Phe1052Val	F1052V
c.3160C>G	p.His1054Asp	H1054D
c.3181G>C	p.Gly1061Arg	G1061R
c.3194T>C	p.Leu1065Pro	L1065P
c.3196C>T	p.Arg1066Cys	R1066C
c.3197G>A	p.Arg1066His	R1066H
c.3205G>A	p.Gly1069Arg	G1069R
c.3208C>T	p.Arg1070Trp	R1070W

FIG. 2 (CONT.-5)

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MUTATION cDNA NAME	MUTATION PROTEIN NAME	MUTATION LEGACY NAME
c.3209G>A	p.Arg1070Gln	R1070Q
c.3222T>A	p.Phe1074Leu	F1074L
c.3230T>C	p.Leu1077Pro	L1077P
c.3266G>A	p.Trp1089X	W1089X
c.3276C>A or c.3276C>G	p.Tyr1092X	Y1092X
c.3302T>A	p.Met1101Lys	M1101K
c.3310G>T	p.Glu1104X	E1104X
c.3454G>C	p.Asp1152His	D1152H
c.3472C>T	p.Arg1158X	R1158X
c.3484C>T	p.Arg1162X	R1162X
c.3485G>T	p.Arg1162Leu	R1162L
c.3528delC	p.Lys1177SerfsX1 5	3659delC
c.3535_3536insTCA A	p.Thr1179IlefsX1 7	3667ins4
c.3587C>G	p.Ser1196X	S1196X
c.3605delA	p.Asp1202AlafsX 9	3737delA
c.3611G>A or c.3612G>A	p.Trp1204X	W1204X
c.3659delC	p.Thr1220LysfsX8	3791delC
c.3691delT	p.Ser1231ProfsX4	3821delT
c.3700A>G	p.Ile1234Val	I1234V
c.3705T>G	p.Ser1235Arg	S1235R
c.3717+12191C>T	No protein name	3849+10kbC->T
c.371S-1G>A	No protein name	3850-1G->A
c.3731G>A	p.Gly1244Glu	G1244E
c.3744delA	p. Lys1250ArgfsX9	3876delA
c.3752G>A	p.Ser1251Asn	S1251N
c.3763T>C	p.Ser1255Pro	S1255P

FIG. 2 (CONT.-6)

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MUTATION cDNA NAME	MUTATION PROTEIN NAME	MUTATION LEGACY NAME
c.37640A	p.Ser1255X	S1255X
c.3773_3774insT	p.Leu1258PhefsX 7	3905insT
c.3808G>A	p.Asp1270Asn	D1270N
c.3346G>A	p.Trp12S2X	W1282X
c.3873+IG>A	No protein name	4005+IG->A
c.3883delA	p.Ile1295PhefsX3 3	4015delA
c.3884_3885insT	p.Ser1297PhefsX 5	4016insT
c.3909C>G	p.Asn1303Lys	N1303K
c.3937C>T	p.Gln1313X	Q1313X
c.3964- 78_4242+577del	NULL	CFTRdele22,23
c.4046G>A	p.Gly1349Asp	G1349D
c.4077_4080delTGT TinsAA	No protein name	4209TGTT->AA
c.4111G>T	p.Glu1371X	E1371X
c.4196_4197delTC	p.Cys1400X	4326delTC
c.4234C>T	p.Gln1412X	Q1412X
c.4242+IG>T	No protein name	4374+IG->T
c.4251delA	p.Glu1418ArgfsX 14	4382delA
c.4296_4297insGA	p.Ser1435GlyfsX 4	4428insGA

FIG. 2 (CONT.-7)