iRecSpot-EF: Effective Sequence Based Features for Recombination Hotspot Prediction

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Supporting Information (#2):

List of features selected by using the AdaBoost feature selection technique. AdaBoost has selected features based on relation probability. Among 40,679 features only 425 features are essential for this dataset. However, The number 425 is not fixed, it depends on the dataset.

The table below contains feature ranking, feature number, feature structure (looking view), and feature importance (probability) it comes from.

Table 1: Features observation

Feature Ranking	Feature Number	Feature Structure	Feature Importance
1	9954	z-axis = $(A+T)$ - $(C+G)$	0.008
2	30769	AGGCT	0.008
3	30710	AGAGC	0.008
4	12413	ATA_TC	0.006
5	13005	CTT_CC	0.006
6	40130	GTGTT	0.006
7	35526	CTTAC	0.006
8	4949	CGGG	0.006
9	1610	GCC	0.006
10	18664	CAAAG	0.006
11	20716	CAACA	0.004
12	813	CCG	0.004
13	33333	CGGGC	0.004
14	38464	TGGTG	0.004
15	22625	AGTTT	0.004
16	20755	CACGA	0.004
17	40214	TACGC	0.004
18	35206	CCCAC	0.004
19	7711	TATT	0.004
20	32559	AAGCG	0.004
21	2535	GTA_T	0.004
22	38322	TCGCT	0.004
23	9325	TTCG	0.004
24	21984	TCTTG	0.004
25	2985	GCC_A	0.004
26	7903	AGTT	0.004
27	6863	AGCT	0.004
			Continued on next page

Feature Ranking	Feature Number	ued from previous page Feature Structure	Feature Importance
28	32069	TATAC	0.004
$\begin{vmatrix} 26 \\ 29 \end{vmatrix}$	32588	AATCA	0.004
$\begin{vmatrix} 29 \\ 30 \end{vmatrix}$		CCCGG	0.004
31	18839 30459	AAATA	0.004
32		AGTA	0.004
1	5848	1	
33	8421	ATAG	0.004
34	13508	GTT_AA	0.004
35	37409	CGCTT	0.004
36	34394	TGTGT	0.004
37	24020	TCTGA	0.004
38	33987	GTTAA	0.004
39	27838	GTGTC	0.004
40	11912	TTC_AG	0.004
41	37378	CGATT	0.004
42	1486	GCT	0.004
43	21097	CTAAT	0.004
44	12169	ACC_AT	0.004
45	1993	GCA	0.004
46	31502	GACCC	0.004
47	32006	TACAC	0.004
48	32638	ACATC	0.004
49	4697	TGTA	0.004
50	23060	CGCGA	0.004
51	23479	GCGGG	0.004
52	20219	AAATA	0.004
53	20483	AGCAA	0.004
54	11249	GGA_CT	0.004
55	24933	CCAAC	0.004
56	22411	ACCCA	0.004
57	37332	CCTGA	0.004
58	4021	GCGG	0.004
59	18452	AGCGA	0.004
60	20211	AAAGA	0.004
61	6387	ATGC	0.004
62	25548	GCTCA	0.004
63	13650	TAT_CT	0.002
64	7367	AGAT	0.002
65	11380	GTA_GA	0.002
66	16047	TTGCG	0.002
67	18128	TTTCG	0.002
68	13069	GAC_CC	0.002
69	23433	GCCAT	0.002
70	29403	CTTTA	0.002
71	40324	TCCAA	0.002
$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	10603	CCA_CA	0.002
73	9261	TCCG	0.002
74	14066	AAACT	0.002
75	7131	GGTC	0.002
		CGA	
76	1712	GCCA	0.002
77	17305		0.002
78	35091	CACGA	0.002
79	31996	TAATA	0.002
80	7139	GTAC	0.002
81	3231	AATT	0.002
			Continued on next page

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Feature Ranking	Feature Number	Feature Structure	Feature Importance
82	13060	GAC_AA	0.002
83	2414	C_TCT	0.002
84	3659	TGCC	0.002
85	25992	TCCAG	0.002
86	1701	CAG	0.002
87	17295	GCCCG	0.002
88	2405	CTA_G	0.002
89	8612	GCAG	0.002
90	23847	TAGAG	0.002
91	18410	AGAAT	0.002
92	22611	AGTGA	0.002
93	33263	CGACG	0.002
94	29993	TAGAT	0.002
95	15424	GGGTG	0.002
96	36977	ATACT	0.002
97	25151	CGGTG	0.002
98	19953	TGACT	0.002
99	7868	ACTG	0.002
100	10503	CAC_AG	0.002
100	3273	AGCA	0.002
101	31944	GTTAG	0.002
102	27362	CTTTT	0.002
103	24247	TTGGG	0.002
104	1266	TGC	0.002
1			
106	4053	GGGG	0.002
107	24520	ACTAG	0.002
108	24757	ATGGC	0.002
109	10901	CTC_GC	0.002
110	38219	TATCA	0.002
111	9012	CCGG	0.002
112	15383	GGCGG	0.002
113	37173	CAGGC	0.002
114	34539	AAACA	0.002
115	12218	ACG_GT	0.002
116	32396	TTCCA	0.002
117	32760	AGAGG	0.002
118	8414	AGTT	0.002
119	4042	GGCC	0.002
120	2498	G_GAC	0.002
121	29431	GAAGG	0.002
122	2477	GCC_G	0.002
123	36947	AGTGA	0.002
124	36948	AGTGA	0.002
125	12550	CAC_AC	0.002
126	1228	GCG	0.002
127	15439	GGTCG	0.002
128	32703	ACGTG	0.002
129	19031	CGTGG	0.002
130	14059	AAACA	0.002
131	27281	CTCCT	0.002
132	27279	CTCCG	0.002
133	32711	ACTAG	0.002
134	33722	GCGGT	0.002
135	5586	GGGC	0.002
199	9900	GGGO	
			Continued on next page

Feature Ranking	Feature Number	ed from previous page Feature Structure	Feature Importance
136	27544	GCCGG	0.002
137	40354	TCCTT	0.002
138	6337	AGAA	0.002
139	14282	ACTAT	0.002
140	21480	GGAAG	0.002
140	14527	ATGTG	$\begin{vmatrix} 0.002 \\ 0.002 \end{vmatrix}$
141	26020	TCGAA	$\begin{vmatrix} 0.002 \\ 0.002 \end{vmatrix}$
143	1252	TAG	$\begin{vmatrix} 0.002 \\ 0.002 \end{vmatrix}$
143	29682	GGACT	
			0.002
145	11792	TGC_CG TTCGC	0.002
146	26262		0.002
147	21203	CTTGA	0.002
148	30633	ACGAT	0.002
149	31328	CGTTG	0.002
150	2360	C_CTA	0.002
151	21414	GCGAC	0.002
152	14606	CACCC	0.002
153	9672	GGCA	0.002
154	38379	TGACA	0.002
155	28653	AGACC	0.002
156	9671	GGAT	0.002
157	19731	TACGA	0.002
158	8484	CCAG	0.002
159	11910	TTC_AC	0.002
160	22009	$TGA_{}GT$	0.002
161	9223	TAAT	0.002
162	9666	GGAC	0.002
163	3123	TCGC	0.002
164	13154	GATTT	0.002
165	16136	AACAG	0.002
166	22442	ACGAT	0.002
167	6219	TGCC	0.002
168	40070	GTCAC	0.002
169	40167	TAAAG	0.002
170	13547	TAACA	0.002
171	25846	TAAGC	0.002
172	5673	TCCA	0.002
173	14700	CCACA	0.002
174	26959	CATCG	0.002
175	31432	CTTAG	0.002
176	1039	ACT	0.002
177	9767	TCAT	0.002
178	38867	ACTGA	0.002
179	12362	AGT_AT	0.002
180	6199	TCGT	0.002
181	37097	CAAAT	0.002
182	28228	TGTAA	0.002
183	33411	CTCAA	0.002
184	1813	AGG	0.002
185	14636	CAGCA	0.002
186	33791	GGATG	0.002
187	1069	CCG	0.002
188	4239	AACT	0.002
189	24897	CAGTT	0.002
100	_ 1001		Continued on next page
			Continued on next page

190	D 1		red from previous page	
191	Feature Ranking	Feature Number	Feature Structure	Feature Importance
192				
193				1
194				1
195	I .			1
196	I .			1
197	195			0.002
198	196	1076	CGG	0.002
199	197	10745	CGA_GT	0.002
200 33785 GGA GT 0.002 201 5741 TTC G 0.002 202 18473 AGG AT 0.002 203 38859 ACT CA 0.002 204 26343 AAA AG 0.002 205 4733 TTT G 0.002 206 22528 AGA TG 0.002 207 23798 TAA GC 0.002 208 29736 GGG AG 0.002 209 10802 CGG AG 0.002 210 40233 TAG AT 0.002 211 8566 C TGT 0.002 212 3067 GTT C 0.002 213 30767 AGG CG 0.002 214 23420 GCA TA 0.002 215 19098 CTC GT 0.002 <td< td=""><td>198</td><td>23764</td><td>GTTGA</td><td>0.002</td></td<>	198	23764	GTTGA	0.002
201 5741 TTC _ G 0.002 202 18473 AGG _ AT _ O.002 203 38859 ACT _ CA _ O.002 204 26343 AAA _ AG _ O.002 205 4733 TTT _ G _ O.002 206 22528 AGA _ TG _ O.002 207 23798 TAA _ GC _ O.002 208 29736 _ GGG _ AG _ O.002 209 10802 _ CGG _ CT _ O.002 210 40233 _ TAG _ AT _ O.002 211 _ 8566 _ C _ TGT _ O.002 212 _ 3067 _ GTT _ C _ O.002 213 _ 30767 _ AGG _ CG _ O.002 214 _ 23420 _ GCA _ TA _ O.002 215 _ 19998 _ CTC _ GT _ O.002 216 _ 13095 _ GAG _ AG _ O.002 217 _ 11039 _ GAC _ TG _ O.002 218 _ 3214 _ A_ ACT _ O.002 219 _ 31478 _ GAA _ GC _ O.002 220 _ 10630 _ CCC _ AC _ O.002 221 _ 2930 _ C _ TGC _ O.002 222 _ 14582 _ CAA _ GC _ O.002 223 _ 20657 _ ATG _ CT _ O.002 224 _ 6270 _ T _ TTT _ O.002 225 _ 33860 _ GGT _ AA _ O.002 228 _ 20649 _ ATG _ AT _ O.002 228 _ 20649 _ ATG _ AT _ O	199	18025	$TTA_{}AT$	0.002
202 18473 AGG_AT 0.002 203 38859 ACTCA 0.002 204 26343 AAAAG 0.002 205 4733 TTTG 0.002 206 22528 AGA_TG 0.002 207 23798 TAA_GC 0.002 208 29736 GGG_AG 0.002 209 10802 CGG.CT 0.002 210 40233 TAG_AT 0.002 211 8566 CTGT 0.002 212 3067 GTT_C 0.002 213 30767 AGG_CG O.002 214 23420 GCA_TA 0.002 215 19098 CTC_GT 0.002 216 13095 GAG_AG 0.002 217 11039 GAC_TG 0.002 218 3214 A_ACT 0.002 219 31478 GAA_GC 0.002 220 10630 <td< td=""><td>200</td><td>33785</td><td>$GGA_{}GT$</td><td>0.002</td></td<>	200	33785	$GGA_{}GT$	0.002
203 38859 ACT CA 0.002 204 26343 AAA AG 0.002 205 4733 TTT G 0.002 206 22528 AGA TG 0.002 207 23798 TAA GC 0.002 208 29736 GGG AG 0.002 209 10802 CGG.CT 0.002 210 40233 TAG AT 0.002 211 8566 C TGT 0.002 212 3067 GTT C 0.002 213 30767 AGG CG 0.002 214 23420 GCA TA 0.002 215 19998 CTC GT 0.002 216 13095 GAG AG 0.002 217 11039 GAC TG 0.002 218 3214 A ACT 0.002 221	201	5741	TTCG	0.002
204 26343 AAA AG 0.002 205 4733 TTT G 0.002 206 22528 AGA TG 0.002 207 23798 TAA GC 0.002 208 29736 GGG AG 0.002 209 10802 CGG.CT 0.002 210 40233 TAG AT 0.002 211 8566 C TGT 0.002 212 3067 GTT_C 0.002 213 30767 AGG CG 0.002 214 23420 GCA TA 0.002 215 19098 CTC GT 0.002 216 13095 GAG AG 0.002 217 11039 GAC.TG 0.002 218 3214 A_ACT 0.002 219 31478 GAA GC 0.002 221 2930 C_TGC 0	202	18473	AGGAT	0.002
204 26343 AAA AG 0.002 205 4733 TTT G 0.002 206 22528 AGA TG 0.002 207 23798 TAA GC 0.002 208 29736 GGG AG 0.002 209 10802 CGG.CT 0.002 210 40233 TAG AT 0.002 211 8566 C TGT 0.002 212 3067 GTT_C 0.002 213 30767 AGG CG 0.002 214 23420 GCA TA 0.002 215 19098 CTC GT 0.002 216 13095 GAG AG 0.002 217 11039 GAC.TG 0.002 218 3214 A_ACT 0.002 219 31478 GAA GC 0.002 221 2930 C_TGC 0	203	38859	ACTCA	0.002
205 4733 TTTG 0.002 206 22528 AGATG 0.002 207 23798 TAAGC 0.002 208 29736 GGGAG 0.002 209 10802 CGG_CT 0.002 210 40233 TAGAT 0.002 211 8566 CTGT 0.002 212 3067 GTT_C 0.002 213 30767 AGGCG 0.002 214 23420 GCATA 0.002 215 19098 CTC_GT 0.002 216 13095 GAG_AG 0.002 217 11039 GAC_TG 0.002 218 3214 A_ACT 0.002 219 31478 GAAGC 0.002 220 10630 CCC_AC 0.002 221 2930 C_TGC 0.002 222 14582 CAA_GC 0.002 223 20657 <td< td=""><td>1</td><td></td><td></td><td>1</td></td<>	1			1
206 22528 AGA	1			
207 23798 TAAGC 0.002 208 29736 GGGAG 0.002 209 10802 CGG_CT 0.002 210 40233 TAGAT 0.002 211 8566 CTGT 0.002 212 3067 GTT_C 0.002 213 30767 AGGC 0.002 214 23420 GCATA 0.002 215 19098 CTCGT 0.002 216 13095 GAG_AG 0.002 217 11039 GAC_TG 0.002 218 3214 A_ACT 0.002 219 31478 GAAGC 0.002 220 10630 CCC_AC 0.002 221 2930 C_TGC 0.002 222 14582 CAAGC 0.002 223 20657 ATGCT 0.002 224 6270 TTTTT 0.002 225 33860	1			1
208 29736 GGGAG 0.002 209 10802 CGG_CT 0.002 210 40233 TAGAT 0.002 211 8566 CTGT 0.002 212 3067 GTT_C 0.002 213 30767 AGGCG 0.002 214 23420 GCATA 0.002 215 19098 CTCGT 0.002 216 13095 GAG_AG 0.002 217 11039 GAC_TG 0.002 218 3214 A_ACT 0.002 219 31478 GAAGC 0.002 220 10630 CCC_AC 0.002 221 2930 C_TGC 0.002 222 14582 CAAGC 0.002 222 14582 CAAGC 0.002 223 20657 ATGTTT 0.002 224 6270 TTTTT 0.002 225 33860	1			1
209 10802 CGG.CT 0.002 210 40233 TAG	1	l .		1
210 40233 TAG	1			1
211 8566 CTGT 0.002 212 3067 GTT_C 0.002 213 30767 AGGCG 0.002 214 23420 GCA_TA 0.002 215 19098 CTCGT 0.002 216 13095 GAG_AG 0.002 217 11039 GAC_TG 0.002 218 3214 A_ACT 0.002 219 31478 GAA_GC 0.002 220 10630 CCC_AC 0.002 221 2930 C_TGC 0.002 222 14582 CAA_GC 0.002 222 14582 CAA_GC 0.002 223 20657 ATG_CT 0.002 224 6270 TTTT 0.002 225 33860 GGT_AA 0.002 226 34838 AGC_GC 0.002 227 31486 GAA_TC 0.002 228 20649 AT	1	l .		1
212 3067 GTT_C 0.002 213 30767 AGGCG 0.002 214 23420 GCATA 0.002 215 19098 CTCGT 0.002 216 13095 GAG_AG 0.002 217 11039 GAC_TG 0.002 218 3214 A_ACT 0.002 219 31478 GAAGC 0.002 220 10630 CCC_AC 0.002 221 2930 C_TGC 0.002 221 2930 C_TGC 0.002 222 14582 CAA_GC 0.002 223 20657 ATG_CT 0.002 224 6270 T_TTT 0.002 225 33860 GGT_AA 0.002 226 34838 AGC_GC 0.002 227 31486 GAA_TC 0.002 228 20649 ATG_AT 0.002 230 34833 AGC_	I .			1
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214 23420 GCATA 0.002 215 19098 CTCGT 0.002 216 13095 GAG_AG 0.002 217 11039 GAC_TG 0.002 218 3214 A_ACT 0.002 219 31478 GAAGC 0.002 220 10630 CCC_AC 0.002 221 2930 C_TGC 0.002 222 14582 CAAGC 0.002 223 20657 ATGCT 0.002 224 6270 TTTT 0.002 225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 <td></td> <td></td> <td></td> <td>1</td>				1
215 19098 CTCGT 0.002 216 13095 GAG_AG 0.002 217 11039 GAC_TG 0.002 218 3214 A_ACT 0.002 219 31478 GAAGC 0.002 220 10630 CCC_AC 0.002 221 2930 C_TGC 0.002 222 14582 CAA_GC 0.002 223 20657 ATGCT 0.002 224 6270 TTTT 0.002 225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584				1
216 13095 GAG_AG 0.002 217 11039 GAC_TG 0.002 218 3214 A_ACT 0.002 219 31478 GAAGC 0.002 220 10630 CCC_AC 0.002 221 2930 C_TGC 0.002 222 14582 CAA_GC 0.002 223 20657 ATGCT 0.002 224 6270 TTTT 0.002 225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 229 12280 AGA_GG 0.002 231 26443 AATCA 0.002 231 26443 AATAA 0.002 233 32645 ACCAC 0.002 234 8584 GAC_AC 0.002 235 20680	I .			1
217 11039 GAC_TG 0.002 218 3214 A_ACT 0.002 219 31478 GAAGC 0.002 220 10630 CCC_AC 0.002 221 2930 C_TGC 0.002 222 14582 CAA_GC 0.002 223 20657 ATGCT 0.002 224 6270 TTTT 0.002 225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 229 12280 AGA_GG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GAACA 0.002 237 9125		l .		1
218 3214 AACT 0.002 219 31478 GAAGC 0.002 220 10630 CCC.AC 0.002 221 2930 CTGC 0.002 222 14582 CAAGC 0.002 223 20657 ATGCT 0.002 224 6270 TTTT 0.002 225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 229 12280 AGAGG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 231 26443 AATACA 0.002 233 32645 ACCATG 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 238 24848 CACCG				1
219 31478 GAA		l .		1
220 10630 CCC_AC 0.002 221 2930 C_TGC 0.002 222 14582 CAA_GC 0.002 223 20657 ATGCT 0.002 224 6270 TTTT 0.002 225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 229 12280 AGA_GG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 238 24848 CACCG 0.002 239 36377				1
221 2930 C_TGC 0.002 222 14582 CAA_GC 0.002 223 20657 ATGCT 0.002 224 6270 TTTT 0.002 225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAA_TC 0.002 228 20649 ATG_AT 0.002 229 12280 AGA_GG 0.002 230 34833 AGCCT 0.002 231 26443 AAT_CA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATT_AG 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGG 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002				1
222 14582 CAAGC 0.002 223 20657 ATGCT 0.002 224 6270 TTTT 0.002 225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATG_AT 0.002 229 12280 AGA_GG 0.002 230 34833 AGCCT 0.002 231 26443 AATAT 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.00	I .			1
223 20657 ATGCT 0.002 224 6270 TTTT 0.002 225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 229 12280 AGA_GG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.		2930	$C_{}TGC$	0.002
224 6270 TTTT 0.002 225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 229 12280 AGA_GG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 237 9125 GCAG 0.002 238 24848 CACCG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002	222	14582	CAAGC	0.002
225 33860 GGTAA 0.002 226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 229 12280 AGA_GG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCATG 0.002	223	20657	ATGCT	0.002
226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 229 12280 AGAGG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCATG 0.002	224	6270	$T_{}TTT$	0.002
226 34838 AGCGC 0.002 227 31486 GAATC 0.002 228 20649 ATGAT 0.002 229 12280 AGA_GG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002	225	33860	GGTAA	0.002
227 31486 GAATC 0.002 228 20649 ATGAT 0.002 229 12280 AGAGG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 AGGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002			AGCGC	1
228 20649 ATGAT 0.002 229 12280 AGAGG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACCG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002				1
229 12280 AGA_GG 0.002 230 34833 AGCCT 0.002 231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002				1
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231 26443 AATCA 0.002 232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002	I .			1
232 7964 CATG 0.002 233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002				1
233 32645 ACCAC 0.002 234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002				1
234 8584 GACA 0.002 235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002				1
235 20680 ATTAG 0.002 236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002				1
236 6255 TTCT 0.002 237 9125 GCAG 0.002 238 24848 CACCG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCATG 0.002				1
237 9125 GCAG 0.002 238 24848 CACGG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002				1
238 24848 CACCG 0.002 239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002				1
239 36377 TGCGT 0.002 240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002				1
240 18435 AGCAA 0.002 241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002				1
241 4788 ACGG 0.002 242 38338 TCGTT 0.002 243 14719 CCA_TG 0.002	I .			1
242 38338 TCGTT 0.002 243 14719 CCATG 0.002				1
243 14719 CCATG 0.002				1
				1
Continued on next page	243	14719	CCATG	0.002
				Continued on next page

Table 1 Continued from previous page			
Feature Ranking	Feature Number	Feature Structure	Feature Importance
244	20709	CAAAC	0.002
245	19777	TAGTT	0.002
246	16163	AAGAA	0.002
247	34985	ATGAT	0.002
248	11207	GCT_AG	0.002
249	25819	GTTTA	0.002
250	19122	CTGCT	0.002
251	11966	TTG_TC	0.002
252	9138	GCGC	0.002
253	25945	TATGT	0.002
254	452	GA_G	0.002
255	4166	TGAT	0.002
256	18428	AGATA	0.002
257	8557	CTCG	0.002
I .		TCCAG	0.002
258	32135		1
259	14235	ACCTA	0.002
260	7998	CCTT	0.002
261	14130	AAGCT	0.002
262	19628	GTGCA	0.002
263	14508	ATGCA	0.002
264	4704	TTAA	0.002
265	25365	GACGC	0.002
266	18690	CAATT	0.002
267	5269	AAGG	0.002
268	2584	$T_{-}ATA$	0.002
269	2585	TAT_A	0.002
270	12905	CTA_AT	0.002
271	37277	CCCTC	0.002
272	19426	GCTTT	0.002
273	13955	TTC_AA	0.002
274	31089	CCACT	0.002
275	10184	ACT_AG	0.002
276	24607	AGCTG	0.002
277	25497	GCCGT	0.002
278	28480	AAGTG	0.002
279	15042	CTGTT	0.002
280			0.002
1	20403	ACGGA	1
281	12652	CCA_CA	0.002
282	16393	AGCAT	0.002
283	6051	GCAC	0.002
284	3425	CTAA	0.002
285	19366	GCGAC	0.002
286	35374	CGGCC	0.002
287	26674	AGGCT	0.002
288	22175	TTCTG	0.002
289	28421	AACAC	0.002
290	27090	CCTCT	0.002
291	7664	GTGA	0.002
292	5134	TACT	0.002
293	22785	CAATT	0.002
294	22155	TTCCA	0.002
295	24710	ATCAC	0.002
296	27721	GGTAT	0.002
297	31719	GGAAG	0.002
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Feature Ranking	Feature Number	Feature Structure	Feature Importance
298	23959	TCCGG	0.002
299	23076	CGGAA	0.002
300	11498	TAA_AT	0.002
301	6512	CTGA	0.002
302	30304	TGTTG	0.002
303	30386	TTGCT	0.002
304	21575	GGTAG	0.002
305	13340	GGC_TA	0.002
306	2022	TAT	0.002
307	17703	TAGAG	0.002
308	18213	AAGAC	0.002
309	5104	GTGA	0.002
310	16504	ATAGG	0.002
311	6581	GCGG	0.002
312	14332	AGATA	0.002
313	30407	TTTAG	0.002
314	30597	ACCAC	0.002
315	778	ACC	0.002
316	22177	TTCTT	0.002
317	31679	GCGTG	0.002
318	32236	TGACA	0.002
319	36690	AATCT	0.002
320	35516	CTGTA	0.002
321		GATGG	0.002
321	15191		
I .	11694	TCG_CC	0.002
323	20308	AATGA	0.002
324	5331	AGGC	0.002
325	4385	CCAA	0.002
326	4490	GACC	0.002
327	18897	CCTCT	0.002
328	16372	AGAGA	0.002
329	24162	TGTTT	0.002
330	27904	TAATG	0.002
331	39092	ATGGA	0.002
332	20296	AATAG	0.002
333	17698	TACTT	0.002
334	10090	ACA_AT	0.002
335	5177	TCTA	0.002
336	12661	CCA_GC	0.002
337	28442	AACGT	0.002
338	35473	CTCCT	0.002
339	30487	AACGG	0.002
340	1568	CAA	0.002
341	21795	TAGAA	0.002
342	15966	TGTTC	0.002
343	31202	CCTTT	0.002
344	27671	GGCGG	0.002
345	32258	TGATT	0.002
346	7786	TTCC	0.002
347	8708	TAAG	0.002
348	7487	CCTT	0.002
349	12196	ACG_AA	0.002
350	5442	CGAC	0.002
351	22213	TTTAC	0.002
301	22210	111110	Continued on next page
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Feature Ranking	Feature Number	ned from previous page Feature Structure	Feature Importance
352	40435	TGAGA	0.002
353	14301	ACTTC	0.002
354	30121	TCGAT	0.002
355	8681	GTCA	0.002
356	14930	CGTCT	0.002
357	7701	TAGG	0.002
358	16313	ACGGT	0.002
359	3727	AACT	0.002
360	6997	CGGG	0.002
		l .	1
361	4427	CGCC	0.002
362	23555	GGCAA	0.002
363	33156	CCCAA	0.002
364	40660	TTTGA	0.002
365	34745	ACGGT	0.002
366	3298	ATAC	0.002
367	22737	ATTCT	0.002
368	23669	GTAGC	0.002
369	20229	AACAC	0.002
370	6675	TAGC	0.002
371	7437	CACG	0.002
372	11622	TCA_AC	0.002
373	13406	GGT_TC	0.002
374	36304	TCTCG	0.002
375	36607	AAATG	0.002
376	9313	TTAA	0.002
377	3924	CGGG	0.002
378	3856	CAGA	0.002
379	7831	AAGT	0.002
380	27979	TATCA	0.002
381	8886	ACGT	0.002
382	22730	ATTAT	0.002
383	26695	AGTAG	0.002
384	14005	TTGGC	0.002
385	21851	TATTA	0.002
386	32781	AGCCC	0.002
387	32969	ATTAT	0.002
388	6552	GATA	0.002
389	20976	CGACG	0.002
390	31159	CCGGG	0.002
391	8769	TGAA	0.002
392	2010	GTC	0.002
393	32955	ATGTA	0.002
394	23097	CGGGT	0.002
395	36106	TACAT	0.002
396	24003	TCTAA	0.002
397	4533	GCGG	0.002
398	7796	TTGG	0.002
399	11050	GAG_AT	0.002
400	33012	CAAGA	0.002
401	6648	GTTA	0.002
402	21559	GGGGG	0.002
403	22209	TTGTT	0.002
404	38571	TTGCA	0.002
405	24011	TCTCA	0.002
100		0	Continued on next page
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Feature Ranking	Feature Number	Feature Structure	Feature Importance
406	3888	CCGA	0.002
407	6596	GGAG	0.002
408	31631	GCCCG	0.002
409	20325	ACAAC	0.002
410	8233	TCCA	0.002
411	14395	AGGTA	0.002
412	3849	CACA	0.002
413	18661	CAAAC	0.002
414	2543	GTC ₋ T	0.002
415	39207	CAGAG	0.002
416	5210	TGTC	0.002
417	16933	CGGAC	0.002
418	39209	CAGAT	0.002
419	27834	GTGGT	0.002
420	31828	GGTGA	0.002
421	1451	CCC	0.002
422	33693	GCCTC	0.002
423	25587	GGAGA	0.002
424	12616	CAT_AG	0.002
425	34628	AATAA	0.002

Table 1 shows, feature ranking, feature number/index (0-base indexing), feature structure, and feature importance. From the above table $`_'$ represents 1-gap, $`__'$ represents 2-gap, $`__'$ represents 3-gap and so on.

Table 2: Probability calculation among features

Ranks of feature	Probability per feature	Group Probability
1-3	0.008	$0.008 \times 3 = 0.024$
4-10	0.006	$0.006 \times 7 = 0.042$
11-62	0.004	$0.004 \times 52 = 0.208$
63-425	0.002	$0.002 \times 363 = 0.726$

Table 2 shows, the important number of features according to their probability. The overall probability is 1.00.