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Alphabetized List of Recognition Specificities

All restriction endonuclease recognition specificities available from New England Biolabs are listed below. For enzymes that recognize non-palindromic sequences, the complementary sequence of each strand is listed. For example, CCTC(7/6) and (6/7)GAGG both represent an MnII (NEB #R0163) site.

All recognition sequences are written 5' to 3' using the single letter code nomenclature with the point of cleavage indicated by a "/".

Numbers in parentheses indicate point of cleaveage for non-palindromic enzymes.

For example, GGTCTC(1/5) indicates cleavage at:

5' ...GGTCTCN'...3'

3' ...CCAGAGNNNNV...5'

Recognition Sequence	Enzyme
AVCGTT	Acil
A/AGCTT	HindIII HindIII-HF®
AAT/ATT	Sspl Sspl-HF®
/AATT	MuCl Tsp509l
A/CATGT	Pcil
ACCGGT	Agel Agel-HF Agel-HF® RE-Mx®
ACCTGC(4/8)	BspM BfuAl
ACCWGGT	SexAl
A/CGCGT	Mul
ACGGC(12/14)	BceAl
ACGT	HpyCH4IV
ACN/GT	HpyCH4III
(10/15)ACNNNNGTAYC(12/7)	Bael
(9/12)ACNNNNNCTCC(10/7)	BsaXI
ACRYGT	Afilli
ACTAGT	Spel Spel-HF® Spel RE-Mx®
ACTGG(1/-1)	Bsrl
ACTGGG(5/4)	Bmrl
A/GATCT	BgIII
AGC/GCT	Afel
AG/CT	Alul
AGG/CCT	Stul
AGT/ACT	Scal Scal-HF® Scal-HF® RE-Mx®
AT/CGAT	Clal BspDI
ATCTATGTCGGGTGCGGAGAAAGAGGTAAT(-15/-19)	PI-Scel
ATGCAT	Nsil

AT/TAAT	Asel
ATTT/AAAT	Swal
(11/13)CAANNNNGTGG(12/10)	CspCl
C/AATTG	Miel Miel-HF® Miel-HF® RE-Mix®
CACGAG(-5/-1)	BssSI
CACGTC(-3/-3)	BmgBl
CAC/GTG	PmII
CACNCAC	Dulli
CACNNN/GTG	Dralli Dralli-HF®
CACNN/NNGTG	Alel
CAGCAG(25/27)	E∞P15I
CAG/CTG	Pull Pull-HF®
CAGNNN/CTG	AwNI
CAGTG(2/0)	BtsIMuti
NNCASTGNN/	TspRI
CATATG	Ndel
CATG	NIaIII
C/ATG	CviAll
/CATG	Fatl
CAYNN/NNRTG	MsII
CC(12/16)	FspEl
CCANNNN/NNNTGG	Xcml
CCANNNN/NTGG	BstXI
CCANNN/NTGG	PfIM
CCATC(4/5)	Bccl
C/CATGG	Ncol Ncol-HF® Ncol-HF® RE-Mix®
CCCAGC(-5/-1)	BseYl
CCCGC(4/6)	Faul
CCC/GGG	Smal
C/CCGGG	Xmal TspM
(0/-1)CCD	NtCviPII
CCDG(10/14)	LpnPl
CCGC(-3/-1)	Acil
CCGC/GG	Sacll
CCGCTC(-3/-3)	BsrBl
C/CGG	Mspl Hpall
CC/NGG	SaFl
/CCNGG	BssKl StyD4I
C/CNNGG	BsaJI
CCNNNNN/NNGG	BsII
C/CRYGG	Btgl
CC/SGG	Nail
C/CTAGG	Avril
CCTC(7/6)	MnII

CCTCAGC(-5/-2)	BbvCl
CCTCAGC CCTCAGC	Nb.BbvCl
CCTCAGC(-5/-7)	Nt.BbvCl
CCTGCA/GG	Sbfl Sbfl-HF®
CCTNAGC(-5/-2)	Bpu10I
CC/TNAGG	Bsu36l
CCTNN/NNNAGG	EcoNI
	HpyAV
CCTTC(6/5)	
CC/WGG	BstNI
/CCWGG	PspGI
C/CWWGG	Styl Styl-HF®
(10/12)CGANNNNNNTGC(12/10)	Bogl
CGAT/CG	Pwl Pwl-HF®
CG/CG	BstUl
C/GGCCG	Eagl Eagl-HF®
CG/GWCCG	Rsrll
CGRY/CG	BsiEl
C/GTACG	BsiW
CGTCTC(1/5)	BsmBl
CGWCG/	Hpy991
CMG/CKG	MspA1I
CNNR(9/13)	MspJI
CR/CCGGYG	SgrAl
C/TAG	Bfal
CTCAG(9/7)	BspCNI
C/TCGAG	Xhol PaeR7I Tlil Xhol RE-Mix®
CTCTTC(1/4)	Earl
CTGAAG(16/14)	Acul
CTGCA/G	Pstl Pstl-HF®
CTGGAG(16/14)	Bpml
C/TNAG	Ddel
C/TRYAG	Sfcl
C/TTAAG	Afili
CTTGAG(16/14)	BpuEl
C/TYRAG	Smll
CYCGRG	Aval BsoBl
GAAGA(8/7)	Mooll
GAAGAC(2/6)	BbsI
GAANN/NNTTC	Xmnl
GAATGC(1/-1)	Bsml
GAATGC	Nb.Bsml
G/AATTC	EcoRI EcoRI-HF® EcoRI-HF® RE-Mx®
GACGC(5/10)	Hgal
GACGT/C	Aatll

GACN/NNGTC	Tth111I PfiFI
GACNN/NNGTC	PshAl
GACNNVNNGTC	Ahdl
GACNNN/NNGTC	Drdl
GAG/CTC	Eco53kl
GAGCT/C	SacI SacI-HF®
GAGGAG(10/8)	BseRI
GAGTC(4/5)	Plel
GAGTC(4/-5)	Nt.BstNBI
GAGTC(5/5)	Myl
G/ANTC	Hinfl
GAT/ATC	ECRVECRV-HF® ECRV-HF® RE-Mx®
/GATC	Mbol Sau3Al Dpnll BfuCl
GATC	Dpnl
GATNN/NNATC	BsaBl
G/AWTC	Tfil
GCAATG(2/0)	BsrDI
GCAATG	Nb.BsrDl
GCAGC(8/12)	Bbvl
GCAGTG(2/0)	BtsI
GCAGTG	Nb.Bts1
GCANNNN/NTGC	BstAPI
GCATC(5/9)	SfaNI
GCATG/C	Sphl Sphl-HF®
GCCGAG(21/19)	NmeAll
GCC/GGC	Nael
G/CCGGC	NgoMV
GCCNNNN/NGGC	BgII
GCGAT/CGC	AsiSI
GCGATG(10/14)	BtgZI
G/CGC	HinP1I
GCG/C	Hhal
G/CGCGC	BssHII
GC/GGCCGC	Notl Notl-HF® Notl-HF® RE-Mx®
GC/NGC	Fnu4HI
GCN/NGC	Cac8I
GCNNNN/NNGC	Mwol
G/CTAGC	Nhel Nhel-HF® Nhel-HF® RE-Mx®
GCTAG/C	Bmtl Bmtl-HF®
GCTCTTC(1/4)	Sapi BspQi
GCTCTTC(1/4) GCTCTTC(1/-7)	
	Sapl BspQl
GCTCTTC(1/-7)	Sapl BspQl Nt.BspQl

GGATC(4/5)	AW
GGATC(4/-5)	NtAwl
G/GATCC	BamHI BamHI-HF®
	Fokl
GGATG(9/13)	BtsCl
GGATG(2/0)	
GG/CC	Haelli Phoi
GGCCGG/CC	Fsel
GGCCNNNN/NGGCC	Sfil
GG/CGCC	Narl
G/GCGCC	Kasl
GGC/GCC	Sfol
GGCGC/C	PluTI
GG/CGCGCC	Ascl Ascl RE-Mx®
GGCGGA(11/9)	Edi
GGGAC(10/14)	BsmFl
GGGCC/C	Apal
G/GGCCC	PspOM
G/GNCC	Sau96l
GGN/NCC	NIaIV
GGTAC/C	Kpnl Kpnl-HF® Kpnl-HF® RE-Mix®
G/GTACC	Acc65I
GGTCTC(1/5)	Bsal Bsal-HF®
GGTGA(8/7)	HphI
G/GTNACC	BstEll BstEll-HF® BstEll-HF® RE-Mx®
G/GWCC	Avail
G/GYRCC	Banl
GKGCMC	BaeGl
GR/CGYC	BsaHl
GRGCY/C	Banll
GT/AC	Rsal
G/TAC	CviQl
GTATAC	BstZ17I
GTATCC(6/5)	BciVI
G/TCGAC	Sall-HF® Sall-HF® RE-Mx®
GTCTC(1/-5)	NtBsmAl
GTCTC(1/5)	BsmAl BcoDl
G/TGCAC	ApaLl
GTGCAG(16/14)	Bsgl
GT/MKAC	Accl
GTN/NAC	Hpy166II
/GTSAC	Tsp45l
GTT/AAC	Hpal
GTT/AAC	Hpal

RCATGYY Nagel RCCGGYY Burf RCGATCY Beht RCGCOYY Hoell RGCCY Child-1 RGCMCCY Ecol109 RGGMCCY Pp.M TACATATAGGTCCTANGGTACCGAY(-91-13) Foell TACAGTA SnaB TACAGTA BepHI TACAGA BepHI TACAGA BupE TACAGA Tonga TACAGA Mul TACAGA Hoyl88I TACAGA Hoyl88I TACAGA Hoyl88II TACAGA Hoyl88II TACAGA Hoyl88II TACAGA Hoyl88II TACAGA Hoyl88II TACAGA Hoyl88II TACAGA HoylANA TACAGA HoylANA <	R/AATTY	Apol
RICCGEY BerII RICCY BerII RICCCY CURLI RICCY CURLI RICCY ECOTORI RICCKCY ECOTORI RICCKCY PUMI TACCITANCAGGTCTMAGGTAGCGAM.(4/13) I-Ceul TACIGTA Small TACGGATACAGGGTAMT.(4/13) I-Seel TACAGGAT BepBI TCCCGGA BepBI TCCGAA TacA TCCGAA Mmel TCCGAA Mul TCCGAA Hy/188I TCCNICA Hy/188II TCCNICA Hy/188II TCCNICAA Hy/188II TCCTAGA Hy/188II TCCTAGA Hy/188II TCCTAGA Hy/188II TCCTAGA Hy/18VI TGGCA Hy/18VI TGGCA Hy/18VI TGGCAA Hy/18VI TGGCAA Hy/18VI TGGCAA Hy/18VI TGGCAA Hy/18VI TGGCAA	RCATG/Y	Nspl
RGGCYY Habil RGCY CviK-1 RGGNCCY Ecoc1091 RGGNCCY PpuM TAACTATIAACGGTCCTAAGGTAGC9A4-91-13) I-Ceul TACKGTA SnaBI TACKGTA BapHI TCCACGA BapHI TCCCGA BapHI TCCACQA BapHI TCCACQA TaqA TCCACQC18) Mmel TCCACQCA Nul TCCACQCA Nul TCCACQCA Hay188I TCNIGA Hay188II TCNINGA Hay188III TCCACACA Bell TCGCAA Hay168III TGCACA HpDCH4V TGCGCA FapI TGCGCA FapI TGCGCA Mel TGCACACCACTATTATGGGT(-134-17) Mel TGTACA Mel TTAA Mel TTAA Mel TTATAA Pacl Pacl RE-Mx89 TTATAA Pacl Pacl RE-Mx89 TTATAA	R/CCGGY	BsrFI
RGCY CVIK-1 RGCNCCY Ecoc1091 RGCNCCY PpuM TAACTATAACGGTCCTAAGGTAGCGAA(-91-13) I-Coul TACIGTA SnaBI TACIGTA SnaBI TACIGTA BapHI TCCAGCA BapEI TCCCGA BapEI TCCRAC(2018) MmeI TCCAA TaqA TCCAGCA Nul TCNIGA Hoy188I TCNINGA Hoy188II TCTAGA Mola RE-Mid0 TGCACA Bell TGCACA HoyCH4V TGCACA HoyCH4V TGCGCA FapI TGGCCA McI TGGCCA McI TGGCCA McI TGGCCA McI TGGCCA McI TGGCCA McI TGTACA BsiG TTAA McI TTATA McI TTATAA PsiI TTCGAA BsiBI TTCGAA <td>R/GATCY</td> <td>BstYl</td>	R/GATCY	BstYl
RGGNCCY Eco01091 RGGNCCY PpuM TACTATACGGTCCTAGGGTAGCGA(-91-13) I-Ceul TAGGTA SnaBI TAGGGA BspHI TCATCA BspHI TCCGGA BspEI TCCGAC(20/18) Mmel TCCGAC TagA TCGACA Ntul TCONIGA Hpy188I TCTACA Xbal Xbal RE-Mx@ TGATCA Bell TGACA HpyCH4V TGCA FspI TGCACACCACCTATTATGGGTATTATGGGT(-134-17) PI-PspI TGGCA Med TGTACA Med TTATA Med TTATA Med TTATA PscI Pack RE-Mx@ TTATA PscI Pack RE-Mx@ TTATATA PscI Pack RE-Mx@ TTATATA PscI Pack RE-Mx@ TTATATA PscI Pack RE-Mx@ TTATAA PscI Pack RE-Mx@ TTATAA PscI Pack RE-Mx@ TTATAA PscI Pack RE-Mx@ TTATAA	RGCGC/Y	Haell
RGGMCCY PpuM TACTATAMCGGTCCTANGGTACCGAN(-9/-13) FCeul TACGTA SnaBl TAGGGATAMCAGGGTAAT(-9/-13) FScol TCATGA BspHI TCCCGA BspEI TCCCGAC BspEI TCCGAC(20/18) Mmel TCCGAC Taqfq TCGAC NnII TCONGA Hpy188I TCNINGA Hpy188III TCTACA Abal Xbal RE-Mx0 TGCA HpyCH4V TGCA HpyCH4V TGCA FspI TGGCAA MscI TGGCA MscI TGGCA MscI TGTACA BsrGI TGGCA MscI TGGCA MscI TTAA MscI TTAA MscI TTAA MscI TTATAA Pacl Pacl RE-Mx0 TTAGAA Pacl Pacl RE-Mx0 TTAGAA Pacl Pacl RE-Mx0 TTAGAA Pacl Pacl RE-Mx0 TTAGAA	RG/CY	CviKI-1
TACCTATACCGGTCCTAGGGTAGCGAA(-9i-13) I-Ceul TACGGTA ShaBl TAGGGATACCAGGGTAAT(-9i-13) I-Scol TICATCA BapHI TCCGGA BapEl TCCGAC(20i18) Mmel TCGGA TaqP TCGICGA Nrul TCNIGA Hpy188II TCNINGA Hpy188III TICACA Xbal Xbal RE-Mx® TIGATCA Bell TGCICA HpyCH4V TGCICA FapI TGCICAA Med TGCICAA Med TGCICAA Med TGCICAA Med TITAA Med TITAA Med TTAA Med TTAATITAA Ped Pacl RE-Mx® TITATAA Pel Pacl RE-Mx® TITAGAA BelBI TTICAAA BelBI TTICAAA Pel Pacl RE-Mx® TTICAAA Pel Pacl RE-Mx® TTICAAA Pel Pacl RE-Mx® TTICAAA Pel Pacl RE-Mx® <td>RG/GNCCY</td> <td>E∞O109I</td>	RG/GNCCY	E∞O109I
TACIGTA SnaBl TACIGTA I-Scal TICATGA BspHI TICATGA BspEI TCCGGA BspEI TCCRAC(20/18) Mmel TCCGA TaqP TCGGA Ntul TCNIGA Hpy188I TCNINGA Hpy188II TICTAGA Xbal Xbal RE-Mx® TIGATCA Bell TGCA HpyCH4V TGCGCA FspI TGCCAA Msd TGCACA Msd TGCAAAACACTATTATGGGTATTATGGGT(-13'-17') PHPspI TGCAA Msd TITAA Msd TITAA Msel TTAATIAA Ped Pacl RE-Mx® TITATAA Psil TITCGAA BstBI TITCGAA BstBI TITCGAA BstBI TITCAAA PspI WCCTCGACB PspX WCCGGW BsaM YACGTR BsaM	RG/GWCCY	PpuM
TAGGGATAACAGGGTAAT(-9I-13) I-Soel TICATGA BspHI TICCGGA BspEI TCCRAC(20/18) Mmel TCGA TaqPI TCGGA Nrul TCNICA Hpy188II TCAINGA Hpy188III TICAGA Xbal Xbal RE-Mw9 TIGATCA BdI TGCA HpyCH4V TGCGA FspI TGCGAACAGCTAITATGGGT(-13I-17) PI-PspI TGCCAA MscI TIGTACA BsrGI TITAA MseI TTAATITAA Pad Pacl RE-Mw9 TTAATATAA PsiI TTICAAA BstBI TITI/AAA Dral VOTCGABB PspXI WCCGGW BsaMI VACIGTR BsaAI	TAACTATAACGGTCCTAAGGTAGCGAA(-9/-13)	I-Ceul
TICATGA BspHI TICCGGA BspEI TICCRAC(20/18) Mmel TICGA TaqA TICGICGA Nul TCNICA Hpy188II TICAGA Xbal Xbal RE-Mx® TICATCA BdI TIGCA HpyCH4V TIGCA FspI TIGCACACAGCTAITATGGGT(-13-17) PI-PspI TIGCACA MscI TIGTACA BsrGI TITAA MscI TITAA MscI TITATAA Pad Pacl RE-Mx® TITICGAA BstBI TITICAAA BstBI TITICAAA PspXI WICCGGW BsaWI YACIGTR BsaAI	TAC/GTA	SnaBl
TCCGGGA BspEI TCCRAC(20/18) MmeI TCGA TaqP TCGGGA NruI TCNIGA Hpy188I TCNINGA Hpy188III TCTAGA Xbal Xbal RE-Mx® TIGATCA BdI TGCA HpyCH4V TGCIGCA FspI TGCCAACACACTATTATGCGT(-13/-17) Pl-PspI TGCIGCA MscI TIGTACA BsrGI TITAA MseI TTAATITAA PacI PacI RE-Mx® TITATAA PsiI TTICGAA BstBI TITI/GAA BstBI TITI/AAA DraI VC/TCGACB PspX WICCGGW BsaWI YACIGTR BsaAI	TAGGGATAACAGGGTAAT(-9/-13)	I-Scel
TCCRAC(20/18) Mmel TICGA Taq ⁹ TCGCGA Nrul TCNIGA Hpy1881 TCNINGA Hpy188II TICATAGA Xball Xball RE-Mx® TIGATCA Bdl TGCA HpyCH4V TGCIGCA Fspl TGCCAAACACCTATTATGGGT(-13/-17) PI-Pspl TGGCCA Mscl TIGTACA BsrGl TITAATTAA Msel TTAATTTAA Psil TTCGAA BstBl TITICGAA BstBl TITI/AAA Dral VC/TCGAGB PspXl WCCCGGW BsaWl YAC/GTR BsaAl	T/CATGA	BspHI
T/CGA Taq ^A TCG/CGA Nrul TCN/GA Hpy188I TC/NINGA Hpy188III T/CATGA Xbal Xbal RE-Mx® T/GATCA Bd TGCA HpyCH4V TGC/CGA Fspl TGC/CCA Fspl TGC/CCA Mscl TGC/CCA Mscl TGTACA BsrGl T/TAA Mscl TITATYTAA Pacl Pacl RE-Mx® TTATYTAA Psil TTICGAA BstBl TTICAA Drai VC/TCGAGB PspX WCCGGW BsaM YAC/GTR BsaM	T/CCGGA	BspEl
TCG/CGA Nrul TCN/CGA Hpy1881 TCN/NGA Hpy188III TICATAGA Xbal Xbal RE-Mx® TICATCA BdI TGCA HpyCH4V TGCCA Fspl TGCCAACCAGCTATTATGGGT(-13/-17) PI-Pspl TGGCCA Mscl TIGTACA BsrGI TITAAT Msel TTAATITAA Pacl Pacl RE-Mx® TTAYTAA Psil TTCGAA BsiBl TTCGAA BsiBl TTTI/AAA Dral VCTCGAGB PspXI WCCGGW BsaMI YAC/GTR BsaAI	TCCRAC(20/18)	Mmel
TCN/GA Hpy188II TC/NNGA Hpy188III T/CAGA Xbal Xbal RE-Mx® T/GATCA BdI TGCA HpyCH4V TGC/GCA Fspl TGC/CCA Mcd T/GATCA BsrGI T/GACA BsrGI T/TAA Mcel TTAAT/TAA Pacl Pacl RE-Mx® TT/CGAA BstBI TT//AAA Dral VC/TCGAGB PspM WICCGGW BsaMI YAC/GTR BsaA	T/CGA	Taq ^q
TCINNGA Hpy188III TICTAGA Xbal Xbal RE-Mx® TIGATCA BdI TIGCA HpyCH4V TGCGCA Fspl TGGCAAACAGCTAITTATGGGTAITTATGGGT(-13I-17) PI-Pspl TGGCCA Mcd TIGATCA BsrGI TITAA Mel TTAATITAA Pacl Pacl RE-Mx® TTAATAA PsiI TTI/CGAA BstBI TTI/AAA Dral VC/TCGAGB PspXI WCCGGW BsaWI VACIGTR BsaAI	TCG/CGA	Nrul
T/CTAGA Xbal Xbal RE-Mx® T/GATCA Bdl TG/CA HpyCH4V TGC/GCA Fspl TGGCAAACAGCTATTATGGGT(-13/-17) Pl-Pspl TGGCCA Mscl T/GTACA BsrGl T/TAA Msel TTAAT/TAA Pacl Pacl RE-Mx® TTAYTAA Psil TT/CGAA BstBl TT//AAA Dral VC/TCGAGB PspX WCCGGW BsaWl YAC/GTR BsaAl	TCN/GA	Hpy188I
T/GATCA Bell TG/CA HpyCH4V TGC/GCA Fspl TGCCAAACACCTATTATCGGTA(TAJ-17) PI-Pspl TGGCCA Mscl T/GTACA BsrGl T/TAAA Msel TTAAT/TAA Pacl Pacl RE-Mx® TTA/TAA Psil TT/CGAA BstBl TT/CGAA BstBl TT/CGAA BstBl TT/CAAA BstBl TC/CCAGB PspX WCCCGGW BsaWI VAC/GTR	TC/NNGA	Hpy188III
TG/CA HpyCH4V TGC/GCA Fspl TGGCAAACAGCTATTATGGGTATTATGGGT(-13/-17) PI-Pspl TGGCCA Mscl T/GTACA BsrGl T/TAA Msel TTAAT/TAA Pacl Pacl RE-Mx® TTA/TAA Psil TT/CGAA BstBl TT/CGAA BstBl VC/TCGAGB PspXl WCCGGW BsaWl YAC/GTR BsaA	T/CTAGA	Xbal Xbal RE-Mx®
TGC/GCA Fspl TGGCAAACAGCTATTATGGGT(-13/-17) PI-Pspl TGG/CCA Mscl T/GTACA BsrGl T/TAA Msel TTAAT/TAA Pacl Pacl RE-Mx® TTA/TAA Psil TT/CGAA BstBl TTT//AAA Dral VC/TCGAGB PspXl WCCGGW BsaWl YAC/GTR BsaAl	T/GATCA	BdI
TGGCAMACAGCTATTATGGGTATTATGGGT(-13/-17) PI-Pspl TGG/CCA Mscl T/GTACA BsrGl T/TAA Msel TTAAT/TAA Pacl Pacl RE-Mx® TTATAA Psil TT/CGAA BstBl TTT//AAA Dral VC/TCGAGB PspXl W/CCGGW BsaWl YAC/GTR BsaAl	TG/CA	HpyCH4V
TGG/CCA Mscl T/GTACA BsrGl T/TAA Msel TTAAT/TAA Pacl Pacl RE-Mx® TTA/TAA Psil TT/CGAA BstBl TTT/AAA Dral VC/TCGAGB PspXl W/CCGGW BsaWl YAC/GTR BsaAl	TGC/GCA	Fspl
T/GTACA BsrGl T/TAA Msel TTAAT/TAA Pacl Pacl RE-Mx® TTA/TAA Psil TT/CGAA BstBl TTT//AAA Dral VC/TCGAGB PspXl W/CCGGW BsaWl YAC/GTR BsaAl	TGGCAAACAGCTATTATGGGTATTATGGGT(-13/-17)	PI-PspI
T/TAA Msel TTAAT/TAA Pacl Pacl RE-Mx® TTATAA Psil TT/CGAA BstBl TTT/AAA Dral VC/TCGAGB PspXl W/CCGGW BsaAl	TGG/CCA	Mscl
TTAAT/TAA Pacl Pacl RE-Mx® TTA/TAA Psil TT/CGAA BstBl TTT/AAA Dral VC/TCGAGB PspXl W/CCGGW BsaWl YAC/GTR BsaAl	T/GTACA	BsrGl
TTATAA Psil TT/CGAA BstBl TTT/AAA Dral VC/TCGAGB PspXl W/CCGGW BsaWl YAC/GTR BsaAl	Т/ТАА	Msel
TTI/CGAA BstBl TTTI/AAA Dral VC/TCGAGB PspXl W/CCGGW BsaWl YAC/GTR BsaAl	TTAAT/TAA	Pacl Pacl RE-Mx®
TTT/AAA Dral VC/TCGAGB PspXl W/CCGGW BsaWl YAC/GTR BsaAl	ТТАТАА	Psil
VC/TCGAGB PspXl W/CCGGW BsaWl YAC/GTR BsaAl	TT/CGAA	BstBl
W/CCGGW BsaWl YAC/GTR BsaAl	TTT/AAA	Dral
YAC/GTR BsaAl	VC/TCGAGB	PspXI
	W/CCGGW	BsaWl
Y/GGCCR Eael	YAC/GTR	BsaAl
	Y/GGCCR	Eael