


```

                                         Aat2
                                         |
TAGCGGTTTACTCACGGGGATTTCCAAGTCTCCACCCATTGACGTCAATGGGAGTTTGTGTTTGGCACCAAAATCAACGGGACTTTCCAAAATGTCGTA
401 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 500
ATCGCCAAACTGAGTGCCCTAAAGGTTCCAGAGGTGGGGTAACTGCAGTTACCCTCAAACAAAACCGTGGTTTTAGTTGCCCTGAAAGTTTTACAGCAT
                                         Eci1
                                         |

ACA ACTCCGCCCATTTGACGCAAATGGGCGGTAGGCGTGTACGGTGGGAGGTCTATATAAGCAGAGCTGGTTTTAGTGAACCGTCAGATCCGCTAGCGCTA
501 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 600
TGTTGAGGCGGGTAACTGCGTTTACCCGCCATCCGCACATGCCACCCTCCAGATATATTCGTCTCGACCAAATCACTTGGCAGTCTAGGCGATCGCGAT
                                         Nhe1 Afel
                                         |
                                         |

SpDon
Ale1 |
Age1 |
Nco1 ||
                                         BsrG1
                                         |
                                         ApaL1
                                         |
CCGGTCCGCCACCATGGTGAGCGAGCTGATTAAGGAGAACATGCACATGAAGCTGTACATGGAGGGCACCGTGAACAACCACCACTTCAAGTGCACATCCG
601 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 700
GGCCAGCGGTGGTACCACTCGCTCGACTAATTCCCTCTTGTACGTGTACTTCGACATGTACCTCCCGTGGCACTTGTGGTGGTGAAGTTCACGTGTAGGC
mKate2 > M V S E L I K E N M H M K L Y M E G T V N N H H F K C T S E

SpAcc
|
AGGGCGAAGGCAAGCCCTACGAGGGCACCCAGACCATGAGAATCAAGGCGGTTCGAGGGCGGCCCTCTCCCCTTCGCCTTCGACATCCTGGCTACCAGCTT
701 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 800
TCCCCTTCGCTTCGGGATGCTCCCGTGGGTCTGGTACTCTTAGTTCCGCCAGCTCCCGCCGGGAGAGGGGAAGCGGAAGCTGTAGGACCGATGGTTCGAA
mKate2 > G E G K P Y E G T Q T M R I K A V E G G P L P F A F D I L A T S F

SpDon
|
CATGTACGGCAGCAAACCTTCATCAACCACACCCAGGGCATCCCCGACTTCTTTAAGCAGTCCCTTCCCCGAGGGCTTCACATGGGAGAGAGTCACCACA
801 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 900
GTACATGCCGTCGTTTTGGAAGTAGTTGGTGTGGGTCCCGTAGGGGCTGAAGAAATTCGTCAGGAAGGGGCTCCCGAAGTGTACCCTCTCTCAGTGGTGT
mKate2 > M Y G S K T F I N H T Q G I P D F F K Q S F P E G F T W E R V T T
```



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                                Sma1
                                Bgl1
                                Bbs1|
                                ||
                                Nar1
                                Kas1|
                                ||
1401  CTTTCGCGGGCGACGATGCCCCCGGGCCGCTCTTCCCCTCCATCGTGGGGCGCCCCAGGCACCAGGGCGTGATGGTGGGCATGGGTTCAGAAGGATTCCCTAT
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 1500
    GAAGCGCCCGCTGCTACGGGGGGCCCGCAGAAGGGGAGGTAGCACCCCGGGGTCCGTTGGTCCCCTACCACCCGTACCCAGTCTTCCCTAAGGATA
Actin  > F A G D D A P R A V F P S I V G R P R H Q G V M V G M G Q K D S Y

                                Eco57
                                Bmr1
1501  GTGGGCGACGAGGCCAGAGCAAGAGAGGCATCCTCACCTGAAGTACCCCATCGAGCACGGCATCGTCACCAACTGGGACGACATGGAGAAAATCTGGC
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 1600
    CACCCGCTGCTCCGGGTCTCGTTCCTCCTCCGTAGGAGTGGGACTTCATGGGGTAGCTCGTGCCGTAGCAGTGGTTGACCCTGCTGTACCTCTTTTAGACCG
Actin  > V G D E A Q S K R G I L T L K Y P I E H G I V T N W D D M E K I W H

                                BseR1
                                Dra3
                                BstAP
                                |
                                |
1601  ACCACACCTTCTACAATGAGCTGCGTGTGGCTCCCGAGGAGCACCCCGTGCTGCTGACCGAGGCCCCCCTGAACCCCAAGGCCAACCGCGAGAAGATGAC
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 1700
    TGGTGTGGAAGATGTTACTCGACGCACACCGAGGGCTCCTCGTGGGGCACGACGACTGGCTCCGGGGGACTTGGGGTTCCGGTTGGCGCTCTTCTACTG
Actin  > H T F Y N E L R V A P E E H P V L L T E A P L N P K A N R E K M T

                                Bsa1
                                BseY1
1701  CCAGATCATGTTTGAGACCTTCAACACCCCCAGCCATGTACGTTGCTATCCAGGCTGTGCTATCCCTGTACGCCTCTGGCCGTACCACTGGCATCGTGATG
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 1800
    GGTCTAGTACAAACTCTGGAAGTTGTGGGGTCCGTACATGCAACGATAGGTCCGACACGATAGGGACATGCGGAGACCGGCATGGTGACCGTAGCACTAC
Actin  > Q I M F E T F N T P A M Y V A I Q A V L S L Y A S G R T T G I V M

                                BstE2
                                Bcg1a
                                PflF1|
                                Dra3|
                                SpAcc
                                Bcg1b
                                Xcm1
1801  GACTCCGGTGACGGGGTCACCCACACTGTGCCCATCTACGAGGGGTATGCCCTCCCCCATGCCATCCTGCGTCTGGACCTGGCTGGCCGGGACCTGACTG
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 1900
    CTGAGGCCACTGCCCCAGTGGGTGTGACACGGGTAGATGCTCCCCATACGGGAGGGGGTACGGTAGGACGCAGACCTGGACCGACCGGCCCTGGACTGAC
Actin  > D S G D G V T H T V P I Y E G Y A L P H A I L R L D L A G R D L T D

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          BsaB1                               BamH1      Xba1      Bcl1BsaB1
          |                                   |             |             |             |
CAGATGTGGATCAGCAAGCAGGAGTATGACGAGTCCGGCCCTCCATCGTCCACCGCAAATGCTTCTAGGGATCCACCGGATCTAGATAACTGATCATAA
2401 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 2500
GTCTACACCTAGTCGTTTCGTCCTCATACTGCTCAGGCCGGGGAGGTAGCAGGTGGCGTTTACGAAGATCCCTAGGTGGCCTAGATCTATTGACTAGTATT
Actin > Q M W I S K Q E Y D E S G P S I V H R K C F *

                      SpDon   Dra1                               Bsm1      Hpa1
                      |       |                               ||         |
TCAGCCATACCACATTTGTAGAGGTTTTACTTGCTTTAAAAAACCTCCACACCTCCCCTGAACCTGAAACATAAAATGAATGCAATTGTTGTTGTTAA
2501 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 2600
AGTCGGTATGGTGTAACATCTCCAAAATGAACGAAATTTTTTGAGGGGTGTGGAGGGGGACTTGGACTTTGTATTTTACTTACGTTAACAACAACAATT

          polyA   Psil       polyA                               polyA       Bts1 Bsm1
          |       |         |                                   |             |             |
CTTGTTTATTGTCAGCTTATAATGGTTACAAAATAAAGCAATAGCATCACAAATTTTCACAAAATAAAGCATTTTTTTTTCACTGCATTCTAGTTGTGGTTTGTCC
2601 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 2700
GAACAAAATAACGTCGAATATTACCAATGTTTATTTTCGTTATCGTAGTGTTTAAAGTGTTTATTTTCGTAAAAAAGTACGTAAGATCAACACCAAACAGG

                      Mlu1       SpDon       Ssp1
                      |         |         |
AAACTCATCAATGTATCTTAACGCGTAAATTGTAAGCGTTAATATTTTGTAAAATTTCGCGTTAAATTTTTGTAAATCAGCTCATTTTTTAACCAATAG
2701 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 2800
TTTGAGTAGTTACATAGAATTGCGCATTTAACATTTCGCAATTATAAAAACAATTTTAAGCGCAATTTAAAAACAATTTAGTCGAGTAAAAAATTGGTTATC

                      Psil                                               BsaXb
                      |                                                           |
GCCGAAATCGGCCAAAATCCCTTATAAAATCAAAAAGAATAGACCGAGATAGGGTTGAGTGTGTTCCAGTTTGGAAACAAGAGTCCACTATTAAGAACGTGG
2801 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 2900
CGGCTTTAGCCGTTTTAGGGAATATTTAGTTTTCTTATCTGGCTCTATCCCAACTCACAACAAGGTCAAACCTTGTTCTCAGGTGATAATTTCTTGCAACC

                      Drd1   BsaXa                               BtgZ1   Dra3
                      |     |                                   |     |
ACTCCAACGTCAAAGGGCGAAAAACCGTCTATCAGGGCGATGGCCCACTACGTGAACCATCACCCCTAATCAAGTTTTTTTGGGGTTCGAGGTGCCGTAAAGC
2901 |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 3000
TGAGGTTGCAGTTTCCCGCTTTTTGGCAGATAGTCCCGCTACCGGGTGATGCACTTGGTAGTGGGATTAGTTCAAAAAACCCAGCTCCACGGCATTTCG
```



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                Eci1              Eci1      Eci1    Eci1        Nco1           polyA
                |                |          ||       |          |           |
AGTCCC GCCCTAACTCCGCCC ATCCGCCCCTAACTCCGCCCAGTTCGCCCATTCTCCGCCCCATGGCTGACTAATTTTTTTTTTTATTTATGCAGAGGCC
3501 |-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----* 3600
TCAGGGCGGGATTGAGGCGGGTAGGGCGGGATTGAGGCGGGTCAAGGCGGGTAAGAGGCGGGGTACCGACTGATTAATAAATAAATAAATAACGTCTCCGG

                                  Avr2
                                  |
                                Stu1|
                                ||
          Sfi1                    SpDon                    BseR1    Cla1                    BsaB1
          |                        |                         |       |                      |
GAGGCCGCCTCGGCCTCTGAGCTATTCCAGAAGTAGTGAGGAGGCTTTTTTGGAGGCCTAGGCTTTTGCAAAGATCGATCAAGAGACAGGATGAGGATCG
3601 |-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----* 3700
CTCCGGCGGAGCCGAGACTCGATAAGGTCTTCATCACTCCTCCGAAAAACCTCCGGATCCGAAAACGTTTCTAGCTAGTTCTCTGTCTACTCTCTAGC

                                  BspM1
                                  |
                                BfuA1                     Eag1                               Bmr1
                                  |                        |                               |
TTTCGCATGATTGAACAAGATGGATTGCACGCAGGTTCTCCGGCCGCTTGGGTGGAGAGGCTATTCCGGCTATGACTGGGCACAACAGACAATCGGCTGCT
3701 |-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----* 3800
AAAGCGTACTAACTTGTCTACCTAACGTGCGTCCAAGAGGCCGGCGAACCCACCTCTCCGATAAGCCGATACTGACCCGTGTTGTCTGTTAGCCGACGA

                                  Nar1
                                  |
                                Kas1|                                   Drd1
                                  ||                                 |
CTGATGCCCGCGTGTTCCGGCTGTCACGCAGGGGCGCCCGGTTCTTTTTGTCAAGACCGACCTGTCCGGTGCCCTGAATGAACTGCAAGACGAGGCAGC
3801 |-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----* 3900
GACTACGGCGGCACAAGGCCGACAGTCGCGTCCCCGCGGGCCAAGAAAAACAGTTCTGGCTGGACAGGCCACGGGACTTACTTGACGTTCTGCTCCGTCG

                                  Pvu2
                                  |
                                Msc1                    Fsp1                    PflF1                    Eco57
                                |                        |                         |                      |
GCGGCTATCGTGGCTGGCCACGACGGGCGTTCCTTGCAGACTGTGCTCGACGTTGTCAAGCGGGAAGGGACTGGCTGCTATTGGGCGAAGTGGCG
3901 |-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----*-----* 4000
CGCCGATAGCACCGACCGGTGCTGCCCGCAAGGAACGCGTCGACACGAGCTGCAACAGTGACTTCGCCCTTCCCTGACCGACGATAACCCGCTTCACGGC

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                                     BssS1
                                     BspM1
                                     BfuA1
BsrB1      BstB1                                     SpAcc
|           |                                     |
AGCGGGACTCTGGGGTTCGAAATGACCGACCAAGCGACGCCCAACCTGCCATCACGAGATTTTCGATTCCACCGCCGCTTCTATGAAAGGTTGGGCTTCG
4501  |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 4600
TCGCCCTGAGACCCCAAGCTTTACTGGCTGGTTCGCTGCGGGTTGGACGGTAGTGCTCTAAAGCTAAGGTGGCGGCGGAAGATACTTTCCAACCCGAAGC

                                     NaeI
                                     NgoM4 |
                                     Bpm1 |
                                     | |
GAATCGTTTTCCGGGACGCCGGCTGGATGATCCTCCAGCGCGGGGATCTCATGCTGGAGTTCTTCGCCACCCTAGGGGGAGGCTAACTGAAACACGGAA
4601  |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 4700
CTTAGCAAAAGGCCCTGCGGCCGACCTACTAGGAGGTCGCGCCCCTAGAGTACGACCTCAAGAAGCGGGTGGGATCCCCCTCCGATTGACTTTGTGCCTT

                                     polyA
                                     |
                                     polyA
                                     |
GGAGACAATACCGGAAGGAACCCGCGCTATGACGGCAATAAAAAGACAGAATAAAACGCACGGTGTGGGTTCGTTTGTTCATAAACCGGGGGTTCGGTCC
4701  |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 4800
CCTCTGTTATGGCCTTCTTGGGCGGATACTGCCGTTATTTTTCTGTCTTATTTTTCGCTGCCACAACCCAGCAAACAAGTATTTGCGCCCCAAGCCAGG

                                     Bsa1
                                     |
CAGGGCTGGCACTCTGTTCGATAACCCACCGAGACCCCATTTGGGGCCAATACGCCCGCGTTTCTTCTTTTCCCCACCCCAAGTTCGGGTGAAG
4801  |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 4900
GTCCCGACCGTGAGACAGCTATGGGGTGGCTCTGGGGTAACCCCGGTTATGCGGGCGCAAAGAAGGAAAAGGGGTGGGGTGGGGGGTTCAAGCCACTTC

                                     BstAP
                                     AlwN1 |
                                     Bsu36 |
GCCAGGGCTCGCAGCCAACGTCGGGGCGGCAGGCCCTGCCATAGCCTCAGGTTACTCATATATACTTTAGATTGATTTAAAACCTTCATTTTTAATTTAA
4901  |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 5000
CGGGTCCCAGCGTCGGTTGCAGCCCCCGCGTCCGGGACGGTATCGGAGTCCAATGAGTATATATGAAATCTAACTAAATTTTGAAGTAAAATTTAAATT
```

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                                 BspH1
                                 |
5001 AAGGATCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTTCGTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAA
      |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 5100
      TTCCTAGATCCACTTCTAGGAAAAACTATTAGAGTACTGGTTTTAGGGAATTGCACTCAAAAAGCAAGGTGACTCGCAGTCTGGGGCATCTTTTCTAGTTT

                                 BpuE1
                                 |
5101 GGATCTTCTTGAGATCCTTTTTTCTGC GCGTAATCTGCTGCTTGCAAACAAAAAACCACCGCTACCAGCGGTGGTTTGTTTGCCGGATCAAGAGCTAC
      |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 5200
      CCTAGAAGAACTCTAGGAAAAAAGACGCGCATTAGACGACGAACGTTTGTTTTTTGGTGGCGATGGTCGCCACCAAAACAAACGGCCTAGTTCTCGATG

      Eco57                               SpAcc
      |                                   |
5201 CAACTCTTTTCCGAAGGTAAGTGGCTTCAGCAGAGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAAGAAGTCTGT
      |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 5300
      GTTGAGAAAAAGGCTTCCATTGACCGAAGTCGTCTCGCGTCTATGGTTTATGACAGGAAGATCACATCGGCATCAATCCGGTGGTGAAGTTCTTGAGACA

                                 AlwN1                               BpuE1
                                 |                                   |
5301 AGCACCGCTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGCTGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTA
      |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 5400
      TCGTGGCGGATGTATGGAGCGAGACGATTAGGACAATGGTCACCGACGACGGTCACCGCTATTGACGACAGAATGGCCCAACCTGAGTTCTGCTATCAAT

                                 ApaL1       BseY1                               SpAcc
                                 |           |                                   |
5401 CCGGATAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGGAGCGAACGACCTACACCGAACTGAGATACCTACAGCGTGAGC
      |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 5500
      GGCTATTCGCGTGCAGCCGACTTGCCCCCAAGCACGTGTGTCGGGTCGAACCTCGCTTGCTGGATGTGGCTTGACTCTATGGATGTCGCACTCG

                                 BciV1
                                 |     |
                                 Eci1   BssS1
                                 |     |
5501 TATGAGAAAGCGCCACGCTTCCCGAAGGGAGAAAGCGGCAGGATACCGGTAAGCGGCAGGGTCGGAACAGGAGAGCGCACGAGGGAGCTTCCAGGGGG
      |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 5600
      ATACTCTTTCGCGGTGCGAAGGGCTTCCCTCTTTCGCTGTCCATAGGCCATTGCGCGTCCCAGCCTTGTCTCTCGCGTGCCTCGAAGGTCCCCC
```

```
                SpAcc                Drd1                BpuE1                SpAcc                Eci1
                |                |                |                |                |
5601 AAACGCCTGGTATCTTTATAGTCCTGTCGGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGGCGGAGCCTATGGAAAAAC
   |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 5700
   TTTGCGGACCATAGAAATATCAGGACAGCCCAAAGCGGTGGAGACTGAACTCGCAGCTAAAAAACTACGAGCAGTCCCCCGCCTCGGATACCTTTTTG
```

```
                SpDon
                BspLU|
                ||
5701 GCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGGCCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTA
   |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 5800
   CGGTCGTTGCGCCGAAAAATGCCAAGGACCGAAAAACGACCGGAAAACGAGTGTACAAGAAAGGACGCAATAGGGGACTAAGACACCTATTGGCATAAT
```

```
                Nsi1
                BfrB1 |
                | |
5801 CCGCCATGCAT
   |||||*| 5811
   GGCGGTACGTA
```

Found:

Aat2	Afe1	Age1	Ale1	AlwN1	ApaL1	Ase1	Avr2	BamH1	Bbs1	Bcg1a	Bcg1b	BciV1	Bcl1
BfrB1	BfuA1	Bgl1	Bgl2	BmgB1	Bmr1	Bpm1	Bpu10	BpuE1	Bsa1	BsaB1	BsaXa	BsaXb	BseR1
BseY1	Bsm1	BspE1	BspH1	BspLU	BspM1	BsrB1	BsrD1	BsrG1	BssS1	BstAP	BstB1	BstE2	Bsu36
BtgZ1	Bts1	Clal	Dra1	Dra3	Drd1	Eag1	Ear1	Eci1	Eco57	EcoRV	Fsp1	Hpa1	Kas1
Mfe1	Mlu1	Msc1	Nae1	Nar1	Nco1	Nde1	NgoM4	Nhe1	Nsi1	PflF1	polyA	PshA1	Psi1
Pvu2	Rsr2	Sall	Sap1	Sca1	SexA1	Sfi1	Sma1	SnaB1	SpAcc	SpDon	Sph1	Ssp1	Stu1
Xba1	Xcm1	Xho1											

Unique:

Afe1	Age1	Ase1	BamH1	Bcg1a	Bcg1b	Bcl1	Bgl2	BmgB1	Bpu10	BspE1	BspLU	BsrG1	BstB1
BstE2	Clal	EcoRV	Fsp1	Hpa1	Mfe1	Mlu1	Nde1	Nhe1	PshA1	Rsr2	Sall	Sca1	SexA1
Sfi1	Sma1	SnaB1	Xba1	Xho1									

Not found:

Aar1	Acc65	Acl1	Afl2	Ahd1	Apa1	Asc1	AsiS1	Baela	Baelb	BbvC1	Blp1	Bsg1	BsiW1
BsmB1	BssH2	BstX1	BstZ1	BxatB	BxatL	BxatR	BxatP	_Chi	Ecl2	EcoK	EcoN1	EcoR1	FCatB
FCatL	FCatR	FCatP	ScFRT	Fse1	FspA1	Hind3	I_Ceu	Kpn1	loxP	Not1	Nru1	Pac1	PflM1
Pme1	Pml1	PspOM	Pst1	Pvu1	R4atB	R4atL	R4atP	R4atR	Sac1	Sac2	SanD1	Sbf1	Sgf1
SgrA1	Spe1	Srf1	Swal	T3RNA	T7RNA	T7Ter	PISce	Xmn1					

Excluded by site complexity:

Acc1	Acil	Afl3	Alu1	Alw1	Apo1	Ava1	Ava2	Ban1	Ban2	Bbv1	BceA1	Bfa1	Bme15
BsaA1	BsaH1	BsaJ1	BsaW1	BseM2	BsiE1	BsiH1	Bs11	BsmA1	BsmF1	Bsp12	BspCa	BspCb	Bsr1
BsrF1	BssK1	BstF5	BstN1	BstU1	BstY1	Btg1	Cac8	CviJ1	Dde1	Eae1	EcoO1	Fau1	Fnu4H
Fok1	Hae2	Hae3	Hga1	Hha1	Hinc2	Hinf1	HinP1	Hpa2	Hph1	Hpy99	Hpy1	Hpy3	HpyC3
HpyC4	HpyC5	Mae3	Mbo2	Mnl1	Mse1	Msl1	MspA1	Mwo1	Nci1	Nla3	Nla4	Nsp1	Ple1
PpuM1	Rsa1	Sau3A	Sau96	SfaN1	Sfc1	Sml1	Sty1	Taq1	Tat1	Tfi1	Tse1	Tsp45	Tsp50
TspR1													