



Esterase-5 gene sequences from the *Drosophila buzzatii* cluster species.

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Introduction

The autosomal gene Esterase-5 (Est-5) was described as a possible member of the carboxyl/cholinesterases gene family (Cygler *et al.*, 1992; Hemilä *et al.*, 1994; Oakeshott *et al.*, 1999, 2005). The members of this gene family have high evolutionary rates and low amino acid identity probably due to their diversification in DNA sequence and biochemical function, relative to the ecological and behavioral changes of the hosts species (Oakeshott *et al.*, 1999, 2005). In *Drosophila* spp two groups of carboxyl/cholinesterases were described: α -esterases and β -esterases, associated to the hydrolyse of α and β -naftil-acetate, respectively. The α -esterase group was characterized from *Drosophila melanogaster* (Robin *et al.*, 2000). In *D. buzzatii* species the Esterase-5 gene encodes possibly the Esterase-2 protein that is homologous to Esterase-9 from *D. melanogaster* (Oakeshott *et al.*, 1993; East *et al.*, 1990). That protein is expressed in digestive tract and Malpighi tubes in *D. buzzatii*, indicating a possible relationship with alimentary behavior (East *et al.*, 1982); however, the true function is still unknown (East *et al.*, 1990).

The *D. buzzatii* cluster (*D. repleta* group: *D. buzzatii* complex) is a monophyletic group (Ruiz *et al.*, 1993; Manfrin *et al.*, 2001), composed by seven species: *D. buzzatii*, *D. serido*, *D. antonietae*, *D. seriema*, *D. gouveai*, *D. borborema*, and *D. koepferae*, and the main diagnostic characteristic is the aedagus morphology (Silva and Sene, 1991). They are endemic of the South American continent, with the exception of *D. buzzatii* that was introduced in other continents (Barker *et al.*, 1985). The species of this cluster are cryptic and cactophilic, whose larvae feed exclusively on decayed tissues of cacti (Pereira *et al.*, 1983). The distribution of the *D. buzzatii* cluster species ranges from Northeast region of Brazil to Chaco Domain in Argentina and is associated with different species of cacti (Manfrin and Sene, 2006). The cladogenetic events in this cluster are associated to pre-Quaternary vicariant events (Manfrin *et al.*, 2001; De Brito *et al.*, 2002), and the population structure and diversification are associated to the retraction and expansion events of the dry vegetation (caatinga and savanna like vegetations) and moist vegetation (rainforest like vegetations) of the South America, during the paleoclimatic changes, as the Quaternary glaciations (Sene *et al.*, 1988).

Historical events, ecological interactions, and genetic characteristics of the species are fundamental to understanding the current patterns of distribution, species diversification, and population variation. For the *Drosophila buzzatii* cluster a lot of historical information is available; however, little information exists regarding the genes associated with ecological interactions, such as the adaptation to the use of decaying tissues of cactus to feed their larvae. Thus, this study aimed to evaluate information in sequences of the Esterase-5 gene in species of the *D. buzzatii* cluster for future works.

Material and Methods

We extracted DNA from 18 individuals of natural populations of the species of the *D. buzzatii* cluster: *D. buzzatii* (2 individuals), *D. koepferae* (3 individuals), *D. borborema* (1 individual), *D.*

serido (3 individuals), *D. seriema* (3 individuals), *D. antonietae* (3 individuals), and *D. gouveae* (3 individuals). For DNA extraction, we used the Puregene kit (GENTRA Systems). We performed the PCR reaction using three sets of primers (Table 1), designed from *D. borborema* sequence from GeneBank (DQ453800) (Figure 1). The PCR reaction conditions were: 1 µl DNA template, 5 µl Buffer, Magnesium Chloride free, 3 µl Magnesium Chloride (2mM/ µl), 4 µl dNTP (4 pmol/µl), 2 µl of each primer (5 pmol/ µl), 0.2 µl Taq polymerase (5 U/ µl) and 32.8 µl of water in a final volume of 50 µl per reaction. The thermocycler program used was one cycle of 94°C, 4 minutes; 35 cycles of 94°C, 40 seconds, 53°C, 40 seconds, 72°C, one minute, and 72°C 3 minutes, and a final extension of 4°C without limit of time. The sequencing reaction was performed directly from the PCR fragments following the protocol in Piccinalli *et al.* (2007). The sequences were checked by eye using the program Chromas lite v 2.0 (Free Software from: http://www.techneysisium.com.au/chromas_lite.html, access: 18/12/2009) and aligned with Piccinalli *et al.* (2007) sequences and *D. virilis* (CH940652) and *D. mojavensis* (CH933806) from Gene Bank using Bioedit program v. 7.0.9.0 (Hall, 1999). To check for possible phylogenetic information, we performed a Maximum Parsimony test (10000 replications) in PAUP v 4.0 program (Swofford, 2001).

Table 1. Characteristics of the primers used to amplify the fragments of Esterase-5 gene.

Name of the primer	Sequence	Size (bp)	Molecular Weight	Annealing temperature °C
Ester-2F	TGG ACT GAA GGA CCA GGT TT	20	6197.1	57.3
Ester-2R	CTT GTC GTA CTC GGG CAG AT	20	6124.1	59.4
Ester-CH1 F	GGG CAC CTT GCA CAA CTA TT	20	6058.0	57.3
Ester-CH1R	AGC CAT GCC AGA AGA TCC TA	20	6076.0	57.3
Ester-CH2F	ATA TG TGT GCA CGG AGC AG	20	6187.0	57.3
Ester-CH3R	CTT CAT CTT CTT GCT GGA CT	20	6006.0	55.3

Results and Discussion

We tested all sets of primers (Table 1) and the combinations of these primers to produce DNA sequences that cover at least the exon 4 and 5 from the Esterase-5 gene. In the first test we used the set of primers Ester-2F and Ester-2R (Figure 1) that amplified a DNA fragment with approximately 1158 bp in all seven species. The set Ester-2F and Ester-2R cover a big portion of the exon 4 and 5 from the Esterase-5 gene. However, the sequences from the species *D. antonietae* and *D. gouveae* showed low quality of the sequences and some samples of the species *D. seriema*, *D. antonietae*, *D. gouveae*, *D. koepferae*, and *D. buzzatii* do not show PCR products. A second test was performed to produce an internal fragment with a better quality of the sequences. We used the set Ester-2F and Ester-CH1R that showed amplified products of 781 bp (Figure 1). However, these products showed low concentration of the PCR products in the species *D. antonietae*, *D. gouveae*, and *D. buzzatii*, and we made a re-amplification to produce PCR products in high concentration to perform the DNA sequence reaction. The re-amplified products do not show ghost bands, and the concentration was enough to perform sequencing reactions in all species. In a third test we used the set Ester-CH2F and Ester-2R to cover the final portion of the Esterase-5 gene and the exon 5. This set of primers produced fragments with 1039 bp, but can only be obtained from the species *D. borborema*, *D. seriema*, *D. koepferae*, and *D. buzzatii*. Even performing re-amplification and changing the PCR conditions, DNA sequences from the species *D. antonietae* and *D. gouveae* could not be obtained. The other combinations of the primers (Ester-2F and Ester-CH3R, Ester-CH2F and Ester-CH1R,

Ester-CH2F and Ester-CH3R, Ester-CH1F and Ester-2R, Ester-CH1F and Ester-CH1R, Ester-CH1F and Ester-CH3R) did not produce DNA fragments even with modifications in the PCR conditions. The consensus sequence sizes varied from 614 bp in the *D. antonietae*, 740 bp from *D. gouveae*, 888 bp from *D. borborema*, 1032 bp from *D. serido*, 1062 bp from *D. kopferae*, 1085 bp from *D. seriema*, and 1091 from *D. buzzatii* (see Appendix 1).

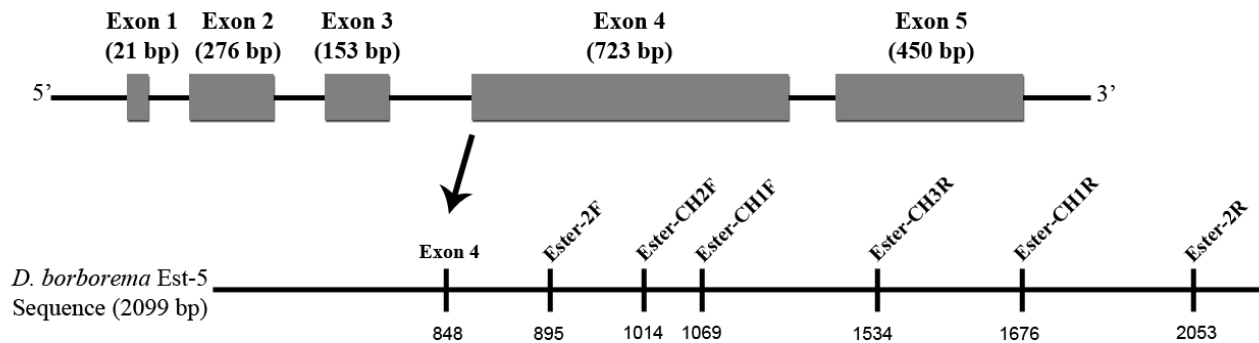


Figure 1. Location of the primers, based on the *D. borborema* sequence from GeneBank, used in this work. In the top of the figure, representation of Esterase-5 gene and exons positions. In the low position of the figure, primer location, the numbers represent the position of the primers in base pairs, related of the *D. borborema* from GeneBank.

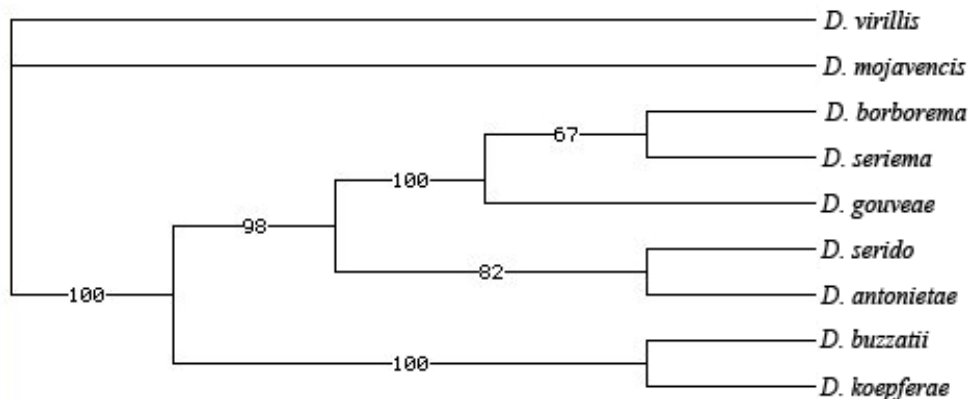


Figure 2. Maximum Parsimony tree based on the consensus sequence of the species used in this work. *D. virillis* and *D. mojavensis* was used as a outgroup. Values of bootstrap calculated based on 10000 replications.

The Maximum Parsimony analyses showed that the species of the *D. buzzatii* form a clade (bootstrap values of 100%), separated from the external species *D. virillis* and *D. mojavensis* (Figure 2). The species in the cluster formed two clades. One encompassed *D. buzzatii* and *D. kopferae* (bootstrap values of 100%). The second one encompassed the other five species with bootstrap values of 98%. This second clade is divided in two groups: one with *D. serido* and *D. antonietae* species (bootstrap values of 82%) and the second group with the species *D. borborema*, *D. seriema*, and *D. gouveae* (bootstrap values of 100%). This phylogenetic hypothesis using the Esterase-5 gene

information is very similar with the ones elaborated using mitochondrial DNA (Manfrin *et al.*, 2001) and chromosomal inversions (Ruiz and Wasserman, 1993; Ruiz *et al.*, 1996). The similarity between the phylogenies obtained with the Esterase-5 gene and the other phylogenies (Manfrin *et al.*, 2001; Ruiz and Wasserman, 1993; Ruiz *et al.*, 1996) seems to be related to the historical relationship with the species of the *D. buzzatii* cluster; even the Esterase-5 showed many indications that it is a gene related to adaptative functions (East *et al.*, 1982). In this work we generated sequences from the Esterase-5 gene from the seven *D. buzzatii* cluster species, and it is a first step to perform future work. The genetic sequences are the base to develop new specific primer sets to be used in molecular systematics, phylogeography, adaptation, and ecological works and more.

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References: Barker, J.S.F., F.M. Sene, P.D. East, and M.A.Q.R. Pereira 1985, *Genética* 67: 161-170; Cygler, M., J. Schrag, *et al.* 1992, *Biotechnology & Genetic Engineering Reviews* 10: 143-184; De Brito, R.A., M.H. Manfrin, and F.M. Sene 2002, *Genetics and Molecular Biology* 25(2): 161-171; East, P., 1982, *Ecological Genetics and Evolution*: 323-338; East, P., A. Graham, and G. Whittington 1990, *Ecological and Evolutionary Genetics of Drosophila*: 389-406; Hall, T., 1999, BioEdit: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT; Hemilä, H., T.T. Koivula, and L. Palva 1994, *Biochimica et Biophysica Acta* 1210(2): 249; Manfrin, M.H., R.A. De Brito, and F.M. Sene 2001, *Annals of the Entomological Society of America* 94(3): 333-346; Manfrin, M.H., and F.M. Sene 2006, *Genetica* 126(1-2): 57-75; Oakeshott, J.G., C. Claudianos, P.M. Campbell, R.D. Newcomb, and R.J. Russell 2005, *Comprehensive Molecular Insect Science* 5: 309-381; Oakeshott, J.G., C. Claudianos, R.J. Russell, and G.C. Robin 1999, *Bioessays* 21(12): 1031-1042; Oakeshott, J.G., E.A. Van Papenrecht, T.M. Boyce, M.J. Healy, and R.J. Russell 1993, *Genética* 90(2): 239-268; Pereira, M.A.Q.R., C.R. Vilela, and F.M. Sene 1983, *Ciência e Cultura* 35: 1313-1319; Piccinali, R.V., L.J. Mascord, J.S.F. Barker, J.G. Oakeshott, and E. Hasson 2007, *Journal of Molecular Evolution* 64(2): 158-170; Robin, G.C.Q., C. Claudianos, R.J. Russell, and J.G. Oakeshott 2000, *Journal of Molecular Evolution* 51(2): 149-160; Ruiz, A., A.M. Cansian, M.R. Alvez, and F.M. Sene 1996, *Revista Brasileira de Genética* 19: 269; Ruiz, A., and M. Wasserman 1993, *Heredity* 70: 582-596; Sene, F.M., M.A.Q.R. Pereira, and C.R. Vilela 1988, *Pacific Science* 42(1-2): 81-88; Silva, A.F.G., and F.M. Sene 1991, *Revista Brasileira de Entomologia* 35: 455-468; Swofford, D., 2001, PAUP* 4.0 beta 5: *Phylogenetic Analysis Using Parsimony (and other methods)*, Sinauer Associates Sunderland, MA.

Appendix 1. Sequences aligned by the program Bioedit v 7.0.9.0 (Hall, 1999) using Blossum 62. * DNA Sequence of *D. borborema* from GeneBank (DQ453800). ** DNA Sequence from Robin (2000).

	10	20	30	40	50
D. bor *		ATGTCAGAAAGGTGTTTGT	TTTCAATAAAAA	CATACATT	TAAATTCCATT
D. bor	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	-----	-----	-----	-----
D. buzz **	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	-----	-----	-----	-----
D. buzz	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	-----	-----	-----	-----
D. koep	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	-----	-----	-----	-----
D. anto	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	-----	-----	-----	-----
D. gouv	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	-----	-----	-----	-----
D. serido	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	-----	-----	-----	-----
D. serie	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	-----	-----	-----	-----

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                60      70      80      90      100
      D. bor *  ....|....|....|....|....|....|....|....|....|....|
      D. bor    ATCATAAATTAATGTGAATTTAATAATAATTTTCATTTTATTTAGTTTGGA
      D. buzz ** .....
      D. buzz    .....
      D. koep     .....
      D. anto     .....
      D. gouv     .....
      D. serido   .....
      D. serie    .....

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                110     120     130     140     150
      D. bor *  ....|....|....|....|....|....|....|....|....|....|
      D. bor    GAGCTGTGAAATATCATTGCCCGTGGGCCAAATCCGGGGCGTTAAGCGCC
      D. buzz ** .....
      D. buzz    .....
      D. koep     .....
      D. anto     .....
      D. gouv     .....
      D. serido   .....
      D. serie    .....

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                160     170     180     190     200
      D. bor *  ....|....|....|....|....|....|....|....|....|....|
      D. bor    GCAGCCTTTACGATGATGATTACTATAGCTTCGAGCGTTTGCCCTTTGGC
      D. buzz ** .....
      D. buzz    .....
      D. koep     .....
      D. anto     .....
      D. gouv     .....
      D. serido   .....
      D. serie    .....

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                210     220     230     240     250
      D. bor *  ....|....|....|....|....|....|....|....|....|....|
      D. bor    AAGCCGCCAGTGGGTGAACTACGCTTTAAGGCACCTGTGCCAGTCGAGCC
      D. buzz ** .....TACACTGCGCTTTAAGGCACCTGTGCCGGTTGAGCC
      D. buzz    .....
      D. koep     .....
      D. anto     .....
      D. gouv     .....
      D. serido   .....
      D. serie    .....

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                260     270     280     290     300
      D. bor *  ....|....|....|....|....|....|....|....|....|....|
      D. bor    CTGGTCCGGTGTGCTGGACTGCACCCACTTTGCCGAAAAGCCCGTGCAGA
      D. buzz ** CTGGTCCGGTGTGCTGGACTGCACCCACTTTGCAAATAAGCCGGTGCAGA
      D. buzz    .....
      D. koep     .....
      D. anto     .....
      D. gouv     .....
      D. serido   .....
      D. serie    .....

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          310      320      330      340      350
D. bor *  ....|....|....|....|....|....|....|....|....|....|
D. bor    AGGGCTTGTTAACAGGCATCATAGAGGGCAGCGAGGACTGCCTTTATTTG
D. buzz ** .....
D. buzz    AGGGCTTGCTAACGGGCATCATAGAAGGCAGCGAGGACTGCCTTTATCTG
D. koep    .....
D. anto    .....
D. gouv    .....
D. serido  .....
D. serie   .....

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          360      370      380      390      400
D. bor *  ....|....|....|....|....|....|....|....|....|....|
D. bor    AATGTATACGCCAAACAGGTGACTTGCTAAAAAATAATATATCACTTTGA
D. buzz ** .....
D. buzz    AATGTATACGCCAAACAGGTGACTTGCTAAAAAATAATATATCACTTTGA
D. koep    .....
D. anto    .....
D. gouv    .....
D. serido  .....
D. serie   .....

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          410      420      430      440      450
D. bor *  ....|....|....|....|....|....|....|....|....|....|
D. bor    TGGATTGTTCAAATGTTCAAGTGATCTCACTGATAACAGCTGAAGAGCGC
D. buzz ** .....
D. buzz    TGGATTGTTCAAATTTCCAAGTGATTTCACTGATAACAGCTGAAGAGCGC
D. koep    .....
D. anto    .....
D. gouv    .....
D. serido  .....
D. serie   .....

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          460      470      480      490      500
D. bor *  ....|....|....|....|....|....|....|....|....|....|
D. bor    AAAACCCCTACCTGTTATGGTCTATATATACGGAGGCGCTTTCTCAATAG
D. buzz ** .....
D. buzz    GAAACCATTACCTGTTATGGTCTATATATACGGAGGCGCTTTCTCCATAG
D. koep    .....
D. anto    .....
D. gouv    .....
D. serido  .....
D. serie   .....

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          510      520      530      540      550
D. bor *  ....|....|....|....|....|....|....|....|....|....|
D. bor    GCGAGGCCACTCGGGATATCTACTCGCCCGATTATTTTATGGCTAAAGAT
D. buzz ** .....
D. buzz    GCGAGGCCACTCGGGAGATCTACTCGCCCGATTACTTTTATGGCTAAAGAT
D. koep    .....
D. anto    .....
D. gouv    .....
D. serido  .....
D. serie   .....

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          560      570      580      590      600
      ....|....|....|....|....|....|....|....|....|....|
D. bor *   GTGGTTCTAGTCACCCTCAACTATCGCGTTGATTGTCTAGGTAAGTGATT
D. bor     -----
D. buzz ** GTGGTTCTAGTCACCCTCAATTATCGCGTTGATTGCCTAGGTAAGTGATT
D. buzz     -----
D. koep     -----
D. anto     -----
D. gouv     -----
D. serido   -----
D. serie    -----

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          610      620      630      640      650
      ....|....|....|....|....|....|....|....|....|....|
D. bor *   GTAAACTTATCGTTTGATTATCTCTTAAATAAGTTAGTTATATTAGTTGT
D. bor     -----
D. buzz ** GTAAACTTATCGTTTGATTATCTCTTAAAGAAGTTTGTATATTAGTTGT
D. buzz     -----
D. koep     -----
D. anto     -----
D. gouv     -----
D. serido   -----
D. serie    -----

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          660      670      680      690      700
      ....|....|....|....|....|....|....|....|....|....|
D. bor *   TCAGCTAGTTTCTCTTATCAATTTGGGTATAAAGTTCGGCCTAGCTTTAA
D. bor     -----
D. buzz ** TTAGCTAGTTTCTCTTATCAATGTGGGTATAAAGTTCGGCTAGCTTTAA
D. buzz     -----
D. koep     -----
D. anto     -----
D. gouv     -----
D. serido   -----
D. serie    -----

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          710      720      730      740      750
      ....|....|....|....|....|....|....|....|....|....|
D. bor *   ATTATTTTTT-TGGTAAAAGCTTTTTCTTTGCTCCCCTAGCTAAGCTTT
D. bor     -----
D. buzz ** GTTATTTTTTATGGCAAAAGCTTTTTCTTTGCTCCCCTAGCTAAGCTTT
D. buzz     -----
D. koep     -----
D. anto     -----
D. gouv     -----
D. serido   -----
D. serie    -----

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          760      770      780      790      800
      ....|....|....|....|....|....|....|....|....|....|
D. bor *   CGCACTCAGCGTACATTGGAAGTCTGCTATTTTCACTCGTTTTCTCTCTCT
D. bor     -----
D. buzz ** CGCACTCAGCGTACATTGGAAGTCTGCTGTTTTCACTCTCTCTCTCTCTCT
D. buzz     -----
D. koep     -----
D. anto     -----
D. gouv     -----
D. serido   -----
D. serie    -----

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      810      820      830      840      850
D. bor *   ....|....|....|....|....|....|....|....|....|....|
D. bor     CTCTCTTTCTCTTTTCTCTTTTCTATTTCCTTTTCTT-----TTTCTT
D. buzz ** CTCTTTACCGTTTCCCCCTCTTTCTTTCTCCTTTTATTCCATTCTCTT
D. buzz     -----
D. koep     -----
D. anto     -----
D. gouv     -----
D. serido   -----
D. serie    -----

      860      870      880      890      900
D. bor *   ....|....|....|....|....|....|....|....|....|....|
D. bor     TAAGGTTTCCTTAGTCTAAAAGACCCAGTTTGAGGTGCCCGGAAATGC
D. buzz ** TAAGGTTTCCTTAGTCTAAAAGACCCAGCTGTGAGGTGCCGGGAAATGC
D. buzz     -----
D. koep     -----
D. anto     -----
D. gouv     -----
D. serido   -----
D. serie    -----

      910      920      930      940      950
D. bor *   ....|....|....|....|....|....|....|....|....|....|
D. bor     TGGACTGAAGGACCAGGTTTTAGCTATCAAATGGGTAAATCAATATATTT
D. buzz ** TGGGCTAAAGGATCAGGTGTTAGCTATCAAATGGGTAAATCAATATATTT
D. buzz     -----CAGGTTTTAGCTATCAAATGGGTAAATCAATATATTT
D. koep     -----GACCAGGTTTTAGCTATCAAATGGGTAAATCAATATATTT
D. anto     -----
D. gouv     -----
D. serido   -----
D. serie    -----TTTTAGCTATCAAATGGGTAAATCAATATATTT

      960      970      980      990      1000
D. bor *   ....|....|....|....|....|....|....|....|....|....|
D. bor     CCTATTTCAATGGTGATGTGAACAACATAACGGTGTTTGGTGAAAGCGCC
D. buzz ** CCTATTTCAATGGTGCTGTGAACAACATAACGGTGTTTCGGTGAAAGCGCC
D. buzz     CCTATTTCAATGGTGATGTGAACAACATAACGGTGTTTCGGTGAAAGCGCC
D. koep     CCTATTTCAATGGTGATGTGAACAACATAACGGTATTCGGTGAAAGCGCC
D. anto     -----
D. gouv     -----GTTTGGTGAAAGCGCC
D. serido   -----GGTGATGTGAAGAACATAACGGTGTTTGGTGAAAGCGCC
D. serie    CCTATTTCAATGGTGATGTGAACAACATAACGGTGTTTGGTGAAAGCGCC

      1010     1020     1030     1040     1050
D. bor *   ....|....|....|....|....|....|....|....|....|....|
D. bor     GGCGGCTGCTCCACCCACTATATGATGTGCACGGAGCAGACGCGGGACT
D. buzz ** GGCGGCTGCTCCACGCACTATATGATGTGCACGGAGCAGACGCGTGGACT
D. buzz     GGCGGCTGCTCCACGCACTATATGATGTGCACGGAGCAGACGCGTGGACT
D. koep     GGCGGCTGCTCCACGCACTATATGATGTGCACGGAGCAGACGCGTGGACT
D. anto     -----ACGGAGCAGACGCGTGGACT
D. gouv     GGCGGCTGCTCCACCCACTATATGATGTGCACGGAGCAGACGCGTGGACT
D. serido   GGCGGCTGTTCCACCCACTATATGATGTGCACAGAGCAGACGCGTGGACT
D. serie    GGCGGCTGCTCCACCCACTATATGATGTGCACGGAGCAGACGCGTGGACT

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	1060	1070	1080	1090	1100
					
D. bor *	CTTCCACAAGGCCATTCCCATGTCGGGCACCTTGCACAAC	ATTGGTCCA			
D. bor	CTTCCACAAGGCCATTCCCATGTCGGGCACCTTGCACAAC	ATTGGTCCA			
D. buzz **	CTTCCACAAGGCCATTCCCATGTCGGGCACCTTGCACAAC	ATTGGTCTA			
D. buzz	CTTCCACAAGGCCATTCCCATGTCGGGCACCTTGCACAAC	ATTGGTCTA			
D. koep	CTTCCACAAGGCCATTCCCATGTCGGGCACCTTGCACAAC	ATTGGTCTA			
D. anto	CTTCCACAAGGCCATTCCCATGTCGGGCACCTTGCACAAC	ATTGGTCCA			
D. gouv	CTTCCACAAGGCCATTCCCATGTCGGGCACCTTGCACAAC	ATTGGTCCA			
D. serido	CTTCCACAAGGCCATTCCCATGTCGGGCACCTTGCACAAC	ATTGGTCCA			
D. serie	CTTCCACAAGGCCATTCCCATGTCGGGCACCTTGCACAAC	ATTGGTCCA			

	1110	1120	1130	1140	1150
					
D. bor *	ATACC-CCTCCCGCAGACTTTGCCTATCGACTGGCCAA	-GGGGAATGGGT			
D. bor	ATACG-TCGCCCCGAGACTTTGCCTATCGTCTGGCCAA	-GTTGAATGGGT			
D. buzz **	ATACA-CCGCCCCGAGACTTTGCCTATCGCCTGGCCAA	-GTTGAATGGGT			
D. buzz	ATACA-CCGCCCCGAGACTTTGCCTATCGCCTGGCCAA	-GTTGAATGGGT			
D. koep	ATACA-CCGCCCCGAGACTTTGCCTATCGCCTGGCCAA	-GTTGAATGGGT			
D. anto	ACACGGCCNNCCGAGACTTTGCCTATCGTCTGGCCAA	AGGTGAATGGGT			
D. gouv	ATACC-CCGCCCCGAGACTTTGCCTATCGACTGGCCAA	-GGTGAATGGGT			
D. serido	ACACGGC-GCCCCGAGACTTTGCCTATCGTCTGGCCAA	-GGTGAATGGGT			
D. serie	ATACC-CCTCCCGCAGACTTTGCCTATCGTCTGGCCAA	-GGTGAATGGGT			

	1160	1170	1180	1190	1200
					
D. bor *	ATGAGGGC-GAAAACAATGATCGCCAGGTGCTGGACTATTTGCGCACTGT				
D. bor	ATGAGGGC-GAAAACAATGATCGCCAGGTGCTGGACTATTTGCGCACTGT				
D. buzz **	TTGAAGGC-GAAAACAATGATCGCCAGGTGCTGGACTATTTGCGTACTGT				
D. buzz	TTGAAGGC-GAAAACAATGATCGCCAGGTGCTGGACTATTTGCGTACTGT				
D. koep	TTGAGGGC-GAAAACAATGATCGCCAGGTGCTGGACTATTTGCGTACTGT				
D. anto	ATNAGGGCTGAAAACAATGATCGCCAGGTGCTGNACTATTTGCGCACTGT				
D. gouv	ATGAGGGC-GAAAACAATGATCGCCAGGTGCTGGACTATTTGCGCACTGT				
D. serido	ATGAGGGC-GAAAACAATGATCGCCAGGTGCTGGACTATTTGCGCACTGT				
D. serie	ATGAGGGC-GAAAACAATGATCGCCAGGTGCTGGACTATTTGCGCACTGT				

	1210	1220	1230	1240	1250
					
D. bor *	CCCAGCAGAGCAGTTGGTTAACCACAGCCTGCTCACTCCT	-GAGGATCGG			
D. bor	GCCGGCAGAGCAGTTGGTTAACCACAGCCTGCTCACTCCT	TGAGGATCGG			
D. buzz **	GCCGGCAGAACAGCTGGTTAACCACAGTCTGCTCACCCCT	-GAAGATCGG			
D. buzz	GCCGGCAGAACAGCTGGTTAACCACAGTCTGCTCACCCCT	-GAGGATCGG			
D. koep	GCCGGCAGAACAGCTGGTTAACCACAGTCTGCTCACCCCT	-GAGGATCGG			
D. anto	GCCGGCAGAGCAGCTGGTTAGCCACACCCTGCTCACCCCT	-GAGGATCGG			
D. gouv	ACCAGCAGAGCAGCTGGTTAACCACAGCCTGCTCACACCT	-GAGGATCGG			
D. serido	GCCGGCAGAGCAGCTGGTTAGCCACAGCCTGCTCACCCCT	-GAGGATCGG			
D. serie	GCCGGCAGAGCAGTTGGTTAACCACAGCCTGCTCACCCCT	-GAGGATCGG			

	1260	1270	1280	1290	1300
					
D. bor *	CGCACCGGACTTATCTATGCCTTTGGCCCCACCGGGAGCCATACGTGAT				
D. bor	CGCAACGGACTTATCTATGCCTTTGGCCCCACCGTGGAGCCATACGTGAT				
D. buzz **	CGCAACGGACTCATCTATGCCTTTGGCCCCACCGTGGAGCCATACGTGAT				
D. buzz	CGCAACGGACTCATCTATGCCTTTGGCCCCACCGTGGAGCCATACGTGAT				
D. koep	CGCAACGGACTCATCTATGCCTTTGGCCCCACCGTGGAGCCATACGTGAT				
D. anto	CGNAACGGACTCATCTATGCNTTTGGCCCCACCGTGGAGCCATACGTGAT				
D. gouv	CGCAACGGACTTATCTATGCCTTTGGCCCCACCGTGGAGCCATACGTGAT				
D. serido	CGCAACGGACTCATCTATGCCTTTGGCCCCACCGTGGAGCCATACGTGAT				
D. serie	CGCAACGGACTTATCTATGCCTTTGGCCCCACCGTGGAGCCATACGTGAT				

	1310	1320	1330	1340	1350
					
D. bor *	GGTAGACTGTGTGGCTCCAAAGCCACAGTTGGAAATGGTGCGCAATGCTT				
D. bor	GGTAGACTGTGTGGCTCCAAAGCCACAGCTGGAGATGGTGCGCGATGCTT				
D. buzz **	GACAGACTGTGTTACTCCAAAGCCACAGCTGGAGATGGTGCGCAAAGCTT				
D. buzz	GGCAGACTGTGTAGCTCCAAAGCCACAGCTGGAGATGGTGCGCGAAGCTT				
D. koep	GGCAGACTGTGTAGCTCCAAAGCCACAGCTGGAGATGGTGCGCGAAGCTT				
D. anto	GGTAGACTGTGTGGCTCCAAAGCCACAGCTGGAGATGGTGCGCGATGCTT				
D. gouv	GGTAGACTGTGTGGCTCCAAAACCCACAGCTGGAGATGGTGCGCGATGCTT				
D. serido	GGTAGACTGTGTGGCTCCAAAGCCACAGCTAGAGATGGTGCGCGATGCTT				
D. serie	GATAGACTGTGTGGCTCCAAAGCCACAGCTGGAGATGGTGCGCGATGCTT				

	1360	1370	1380	1390	1400
					
D. bor *	GGAGCAATAAGTTACCAGCCATGCTGGGCGGCACCTCTTTTGAAGGTCTC				
D. bor	GGAGCAATAAGTTACCAGTCATGCTGGGCGGCACCTCTTTTGAAGGTCTC				
D. buzz **	GGAGCAATAAGTTACCAGTCATGCTGGGCGGTACTTCTTTTGAAGGTCTC				
D. buzz	GGAGCAATAAGTTACCAGTCATGCTGGGCGGTACTTCTTTTGAAGGTCTC				
D. koep	GGAGCAATAAGTTACCAGTCATGCTGGGCGGTACTTCTTTTGAAGGTCTC				
D. anto	GGAGCAATAAGTTACCAGTCATGCTGGGCGGTACTTCTTTTGAAGGTCTC				
D. gouv	GGAGCAATAAGTTACCAGCCATGCTGGGCGGCACCTCTTTTGAAGGTCTC				
D. serido	GGAGCAATAAGTTACCAGTCATGCTGGGCGGTACTTCTTTTGAAGGTCTC				
D. serie	GGAGCAATAAGTTACCAGTCATGCTGGGCGGCACCTCTTTTGAAGGTCTC				

	1410	1420	1430	1440	1450
					
D. bor *	TTCATGTATCCCGCCTTGAAGGCTAATCCTAAGGGCATGGATAGCCTGCC				
D. bor	TTCATGTATCCCGCCTTAAAGGCCAATCCTAAGGGCATGGATAGCCTGCC				
D. buzz **	TTCATGTATCCCGCCTTGAAGGCTAATCCTAAGGGCATGGATAGCCTGCC				
D. buzz	TTCATGTATCCCGCCTTGAAGGCTAATCCTAAGGGCATGGATAGCCTGCC				
D. koep	TTCATGTATCCCGCCTTGAAGGCTAATCCTAAGGGCATGGATAGCCTGCC				
D. anto	TTCATGTATCCCGCCTTGAAGGCTAATCCTAAGGGCATGGATAGCCTGCC				
D. gouv	TTCATGTATCCCGCCTTAAAGGCCAATCCTAAGGGCATGGATAGCCTGCC				
D. serido	TTCATGTATCCCGCCTTGAAGGCTAATGCTAAGGGCATGGATAGCCTGCC				
D. serie	TTCATGTATCCCGCCTTAAAGGCCAATCCTAAGGGCATGGATAGCCTGCC				

	1460	1470	1480	1490	1500
					
D. bor *	GCAGGATCTGCTGCGATTGACGCCCTTATGAAGTGCGTGTGCTCAATACGG				
D. bor	GCAGGATCTGTTGCGATTGACGCCCTTATGAAGTGCGTGTGCTCAATACGG				
D. buzz **	GCAGGATCTGTTGCGATTGACGCCCATGAAGTGCGTGTGCTCAATACGG				
D. buzz	GCAGGATCTCTTGCGATTGACGCCCATGAAGTGCGTGTGCTCAATACGG				
D. koep	GCAGGATCTGTTGCGATTGACGCCCATGAAGTGCGTGTGCTCAATACGG				
D. anto	GCAGGATCTGTTGCGATTGACGCCCTTATGAAGTGCGTGTGCTCAATAAGG				
D. gouv	GCAGGATCTGTTGCGATTGACGCCCTTATGAAGTGCGTGTGCTCAATACGG				
D. serido	GCAGGATCTGTTGCGATTGACGCCCATGAAGTGCGTGTGCTCAATACGG				
D. serie	GCAGGATCTGTTGCGATTGACGCCCTTATGAAGTGCGTGTGCTCAATACGG				

	1510	1520	1530	1540	1550
					
D. bor *	AGCAACAGAATTTAGAGTCCAGCAAGAAGATGAAGCAATTGTACTTTGGC				
D. bor	AGCAACAGAATCTAGAGTCCAGCAAGAAGATGAAGCAATTGTACTTTGGC				
D. buzz **	AGCAACAGAATTTAGAGTCCAGCAAGAAGATGAAGCAATTGTACTTTGGC				
D. buzz	AGCAACAGAATTTAGAGTCCAGCAAGAAGATGAAGCAATTGTACTTTGGC				
D. koep	AGCAACAGAATTTAGAGTCCAGCAAGAAGATGAAGCAATTGTACTTTGGC				
D. anto	AACAACAGAATCTAGAGTCCAGCAAGAAGATGAAGCAATTGTACTTTGGC				
D. gouv	AGCAACAGAATCTAGAGTCCAGCAAGAAGATGAAGCAATTGTACTTTGGC				
D. serido	AACAACAGAATCTAGAGTCCAGTCCAGCAAGAAGATGAAGCAATTGTACTTTGGC				
D. serie	AGCAACAGAATCTAGAGTCCAGCAAGAAGATGAAGCAATTGTACTTTGGC				

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                1560      1570      1580      1590      1600
D. bor *      ....|....|....|....|....|....|....|....|....|....|
D. bor        GATGACACGCCCAGCTCCAAGCTCATCATGAACTTCATGGACGTGAGTT-
D. buzz **    GATGCCACGCCCAGCTCCAAGTTCATCATGAACTTCATGGACGTGAGCTG
D. buzz       GATGCTACACCCAGCTCCAAGCTCATCATGAACTTCATGGACGTGAGCT-
D. koep       GATGCTACACCCAGCTCCAAGCTCATCATGAACTTCATGGACGTGAGCT-
D. anto       GATGCCACGCCCAGCTCCAAGCTCATCATGAACTTTATGGACGNNNTN-
D. gouv       GATGACACGCCCAGCTCCAAGCTCATCATGAACTTCATGGACGTGAGCT-
D. serido     GATGCCACGCCCAGCTCCAAGCTCATCACGAACTTCATGGACGTGAGTT-
D. serie      GATGCCACCCCAGCTCCAAGTTCATCATGAACTTCATGGACGTGAGTT-

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                1610      1620      1630      1640      1650
D. bor *      ....|....|....|....|....|....|....|....|....|....|
D. bor        CGAATCACTTAAGTTTAACTATATAAC--TACTCTA-ATTCTTTAT----
D. buzz **    CGAATCACTTAAGTTTAACTATATAAC--TACTCTA-ATTCTTTGT----
D. buzz       CGAATCACTTAAGTTTAACTATATAAC--TACTCTA-ATTCTTTGT----
D. koep       CGAATCACTTAAGTTTAACTATATAAC--TACTCTA-ATTCTTTGT----
D. anto       NNNATCACTTAAGTTTAACTATATACT-GTACTCTATATACTGTACT---
D. gouv       CGAATCACTTAAGTTTAACTATATACTGTACTCTA-ATTCTTTAT----
D. serido     CGAATCACTTAAGTTTAACTATATATT-GTACTCTATATACTGTACTCTA
D. serie      CGAATCACT---TTTAACTATATACTGTACTCTA-ATTCTTTAT----

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                1660      1670      1680      1690      1700
D. bor *      ....|....|....|....|....|....|....|....|....|....|
D. bor        -----ATTTCTACTTTTAGTACTATTCTTATAGGATCTTCTGGCAT
D. buzz **    -----ATTTCTACTTTTAGTACTATTCTTATAGGATCTTCTGGCAT
D. buzz       -----ATTTCTACTTTTAGTACTATTCTTATAGGATCTTCTGGCAT
D. koep       -----ATTTCTACTTTTAGTACTATTCTTATAGGATCTTCTGGCAT
D. anto       -----ATTTCTACTTTTAGTACTATTCTTATAGGATCTTCTGGCAT
D. gouv       -----ATTTCTACTTTTAGTACTATTCTTATAGGATCTTCTGGCAT
D. serido     ATTTCTTTATATTTCTACTTTTAGTACTATTCTTATAGGATTTTCTGGCAT
D. serie      -----ATTTCTACTTTTAGTACTATTCTTATAGGATCTTCTGGCAT

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                1710      1720      1730      1740      1750
D. bor *      ....|....|....|....|....|....|....|....|....|....|
D. bor        GGCTTCCATCGCACTTTGCAAGCTCGCTGGCATAACGCCACGGCCCCCAC
D. buzz **    GGCTTCCATCGCACTTTGCAAGCTCGCTGGCATAACGCCACGGCTCCAC
D. buzz       GGCTTCCATCGCACTTTGCAAGCTCGCTGGCATAACGCCACGGCTCCAC
D. koep       GGCTTCCATCGCACTTTGCAAGCTCGCTGGCATAACGCCACGGCTCCAC
D. anto       -----
D. gouv       GGCTTCCATCGCACTTTGCAAGCTCGCTGGCATAACGCCACGG-----
D. serido     GGCTTCCATCGTACTTTGCAAGCTCGCTGGCATAACGCCACGGCTCCAC
D. serie      GGCTTCCATCGCACTTTGCAAGCTCGCTGGCATAACGCCACGGCTCCAC

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                1760      1770      1780      1790      1800
D. bor *      ....|....|....|....|....|....|....|....|....|....|
D. bor        TTACTTCTATCGCTTTGATTTTGATTACCTGGTTTCAATTTCTACCGTA
D. buzz **    ATACTTCTATCGCTTTGATTTTGATTACCTGATTTCAATTTCTACCGTA
D. buzz       ATACTTCTATCGCTTTGATTTTGATTACCTGATTTCAATTTCTACCGTA
D. koep       ATACTTCTATCGCTTTGATTTTGATTACCTGATTTCAATTTCTACCGTA
D. anto       -----
D. gouv       -----
D. serido     TTACTTCTATCGCTTTGATTTTGATTACCTGATTTCAATTTCTTCCGTA
D. serie      TTACTTCTATCGCTTTGATTTTGATTACCTGATTTCAATTTCTACCGTA

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	1810	1820	1830	1840	1850
					
D. bor *	AGAAATTCTG	TGGGGATGACATCAAGCAGGGCGTTGCCCATGCCGATGA			
D. bor	AGAAATTCTG	TGGGGATGACATCAAGCAGGGCGTTGCCCATGCCGATGA			
D. buzz **	AAAAATTCTG	TGGCGATGACATCAAGCAGGGCGTTGCCCATGCCGATGA			
D. buzz	AAAAATTCTG	TGGCGATGACATCAAGCAGGGCGTTGCCCATGCCGATGA			
D. koep	AAAAATTCTG	TGGCGATGACATCAAGCAGGGCGTTGCCCATGCCGATGA			
D. anto	-----	-----	-----	-----	-----
D. gouv	-----	-----	-----	-----	-----
D. serido	AGAAATTCTG	TGGCGATGACATTAAGCAGGGCGTTGCCCATGCCGATGA			
D. serie	AGAAATTCTG	TGGGGATGACATCAAGCAGGGCGTTGCCCATGCCGATGA			

	1860	1870	1880	1890	1900
					
D. bor *	GTTGAGCTATTTGTTCCGCAACTCGGAGTCGTGGAAACTGGACAAAGCAT				
D. bor	GTTGAGCTATTTGTTCCGCAACTCGGAGTCGTGGAAACTGGACAAAGCAT				
D. buzz **	GTTGAGCTATTTGTTCCGTAACCTCGGAGTCGTGGAAACTGGACAAAGCCT				
D. buzz	GTTGAGCTATTTGTTCCGTAACCTCGGAGTCGTGGAAACTGGACAAAGCCT				
D. koep	GTTGAGCTATTTGTTCCGTAACCTCGGAGTCGTGGAAACTGGACAAAGCCT				
D. anto	-----	-----	-----	-----	-----
D. gouv	-----	-----	-----	-----	-----
D. serido	GTTGAGCTATTTGTTCCGCAACTCGGAGTCGTGGAAACTAGACAAAGCCT				
D. serie	GTTGAGCTATTTGTTCCGCAACTCGGAGTCGTGGAAACTGGACAAAGCCT				

	1910	1920	1930	1940	1950
					
D. bor *	CCGCTGAATATCGCACCATTCAACGGCTGGTCGACATTTGGACATCCTTT				
D. bor	CCGCTGAATATCGCACCATTCAACGGCTGGTCGACA-----				
D. buzz **	CTGCTGAATATCGCACCATTCAACGGCTGGTCGACATTTGGACATCCTTT				
D. buzz	CTGCTGAATATCGCACCATTCAACGGCTGGTCGACATTTGGACATCCTTT				
D. koep	CTGCTGAATATCGCACCATTCAACGGCTGGTCGACATTTGGACATCCTTT				
D. anto	-----	-----	-----	-----	-----
D. gouv	-----	-----	-----	-----	-----
D. serido	CCGCTGAATATCGCACCATTCAACGGCTGGTCGACATTTGGACATCCTTT				
D. serie	CCGATGAATATCGCACCATTCAACGTCTGGTCGACATTTGGACATCCTTT				

	1960	1970	1980	1990	2000
					
D. bor *	GCAGCTACTTCCAATCCCAACTGCGCTGAGACAGCGCATCTCGATTGGCA				
D. bor	-----	-----	-----	-----	-----
D. buzz **	GCAGCTACTTCCAATCCCAACTGCGCT-----				
D. buzz	GCAGCTACTTCCAATCCCAACTGCGCTGAGACTGCGCATCTCGATTGGCA				
D. koep	GCAGCTACTTCCAATCCCAACTGCGCTGAGACTGCGCATCTCGA-----				
D. anto	-----	-----	-----	-----	-----
D. gouv	-----	-----	-----	-----	-----
D. serido	GCAGCTACTTCCAATCCCAACTGCGCTGAGACAGCGCATCTCGATTGGC-				
D. serie	GCAGCTACTTCCAATCCCAACTGCGCTGAGACAGCGCATCTCGATTGGCA				

	2010	2020	2030	2040	2050
					
D. bor *	GCCAGCTAAGCAGGAGCAGCCTCAGCGTGTGCTCAACATTGGCAAAGAGG				
D. bor	-----	-----	-----	-----	-----
D. buzz **	-----	-----	-----	-----	-----
D. buzz	GCCAGCGAAGCAGGAGCAGCCTCAGC-----				
D. koep	-----	-----	-----	-----	-----
D. anto	-----	-----	-----	-----	-----
D. gouv	-----	-----	-----	-----	-----
D. serido	-----	-----	-----	-----	-----
D. serie	GCCAGCTAAGCAGGAGCAGCCTCAGC-----				

	2060	2070	2080	2090	2100
D. bor *					
D. bor	TGGAGCTCATCGATCTGCCCGAGTACGACAAGCTGCTCGTCTGGGATAGT				
D. buzz **	----- ----- ----- ----- ----- ----- ----- ----- ----- -----				
D. buzz	----- ----- ----- ----- ----- ----- ----- ----- ----- -----				
D. koep	----- ----- ----- ----- ----- ----- ----- ----- ----- -----				
D. anto	----- ----- ----- ----- ----- ----- ----- ----- ----- -----				
D. gouv	----- ----- ----- ----- ----- ----- ----- ----- ----- -----				
D. serido	----- ----- ----- ----- ----- ----- ----- ----- ----- -----				
D. serie	----- ----- ----- ----- ----- ----- ----- ----- ----- -----				

	2110	2120
D. bor *		
D. bor	CTATATAAGAAGGAACACTTATTTTAG	
D. buzz **	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	
D. buzz	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	
D. koep	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	
D. anto	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	
D. gouv	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	
D. serido	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	
D. serie	----- ----- ----- ----- ----- ----- ----- ----- ----- -----	

Guide to Authors

Drosophila Information Service prints short research, technique, and teaching articles, descriptions of new mutations, and other material of general interest to *Drosophila* researchers. The current publication schedule for regular issues is annually, with the official publication date being December. The annual issue will include material submitted during the calendar year. To help us meet this target date, we request that submissions be sent by 15 December, but articles are accepted at any time. A receipt deadline of 31 December is a firm deadline, due to printer submission schedules. Electronic submissions are encouraged, and may be required for lengthy or complex articles.

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Waters, R.L., J.T. Smith, and R.R. Brown 1990, *J. Genet.* 47: 123-134.

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