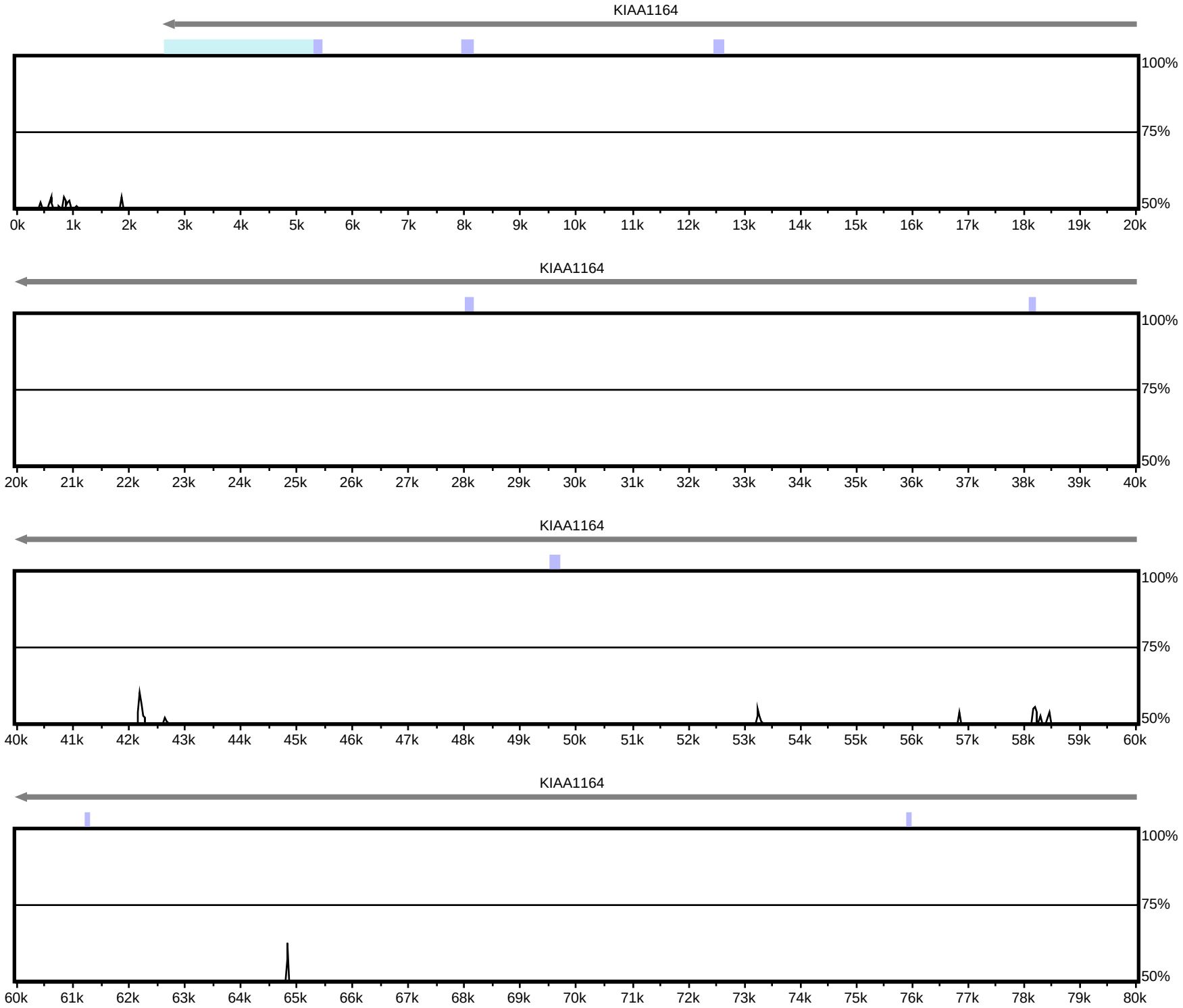
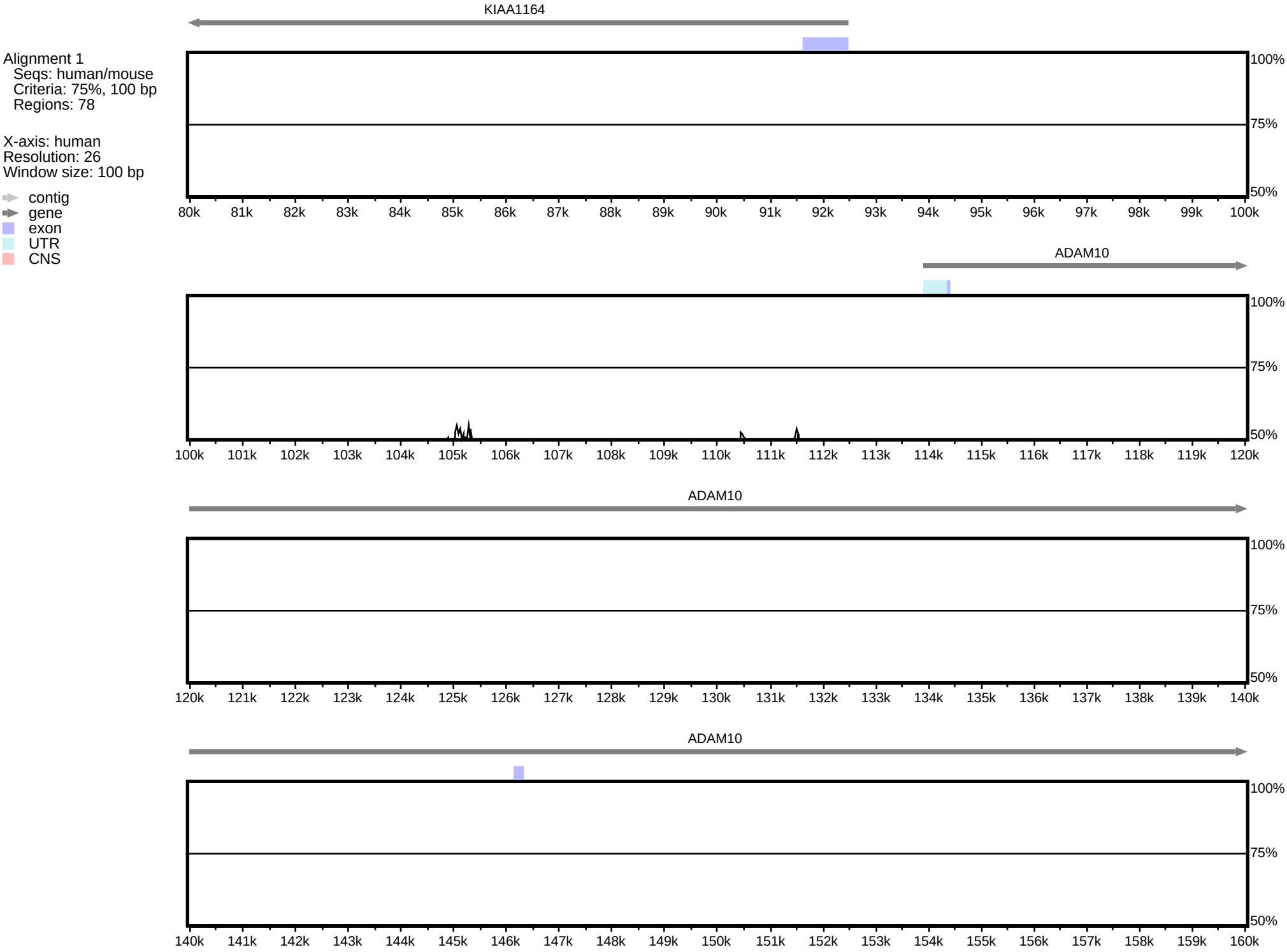
Alignment 1
Seqs: human/mouse
Criteria: 75%, 100 bp
Regions: 78

X-axis: human
Resolution: 26
Window size: 100 bp

contig
gene
exon
UTR
CNS




Alignment 1
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Criteria: 75%, 100 bp
Regions: 78
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
KIAA1164
ADAM10
ADAM10
ADAM10
100%
75%
50%
80k 81k 82k 83k 84k 85k 86k 87k 88k 89k 90k 91k 92k 93k 94k 95k 96k 97k 98k 99k 100k
100k 101k 102k 103k 104k 105k 106k 107k 108k 109k 110k 111k 112k 113k 114k 115k 116k 117k 118k 119k 120k
120k 121k 122k 123k 124k 125k 126k 127k 128k 129k 130k 131k 132k 133k 134k 135k 136k 137k 138k 139k 140k
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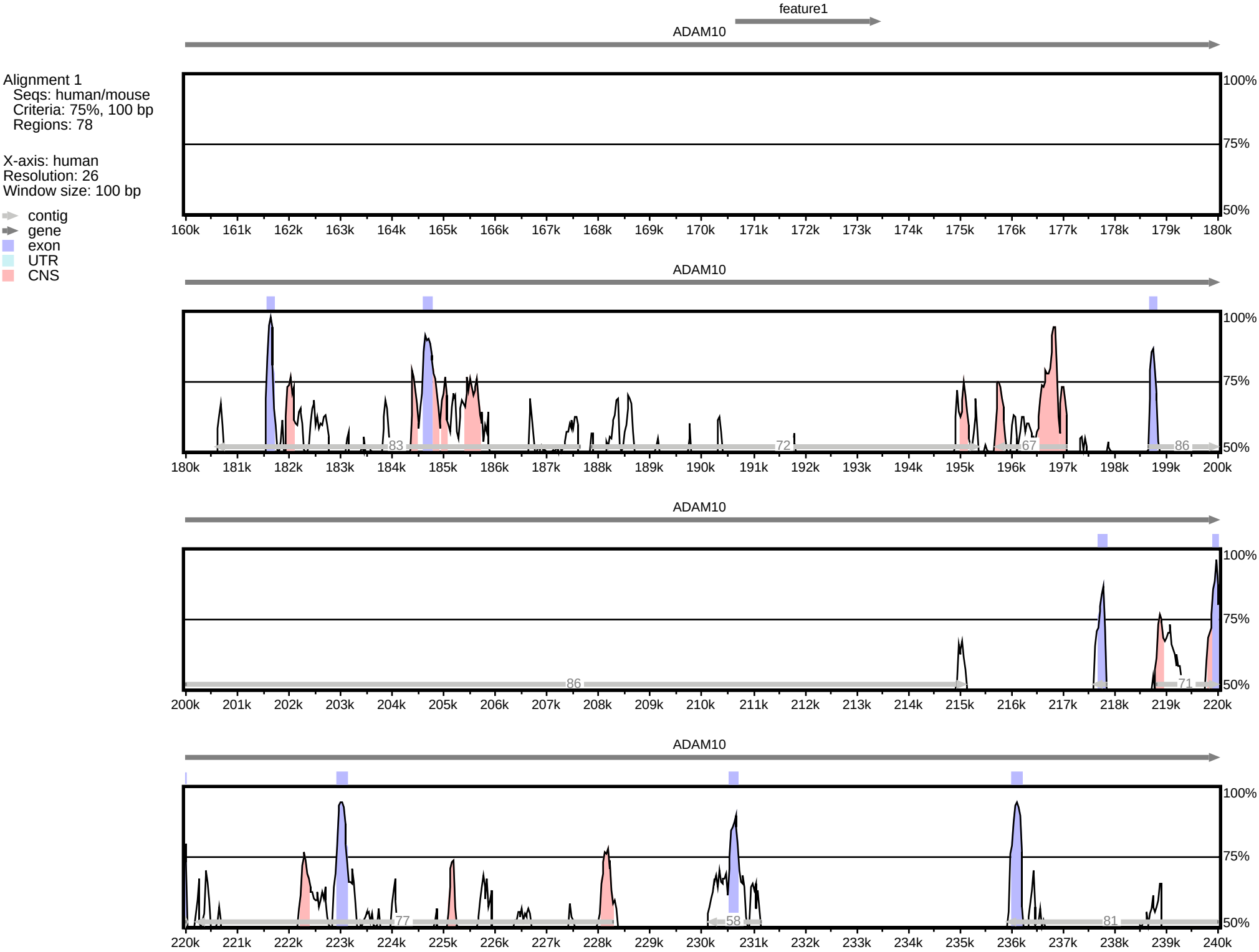
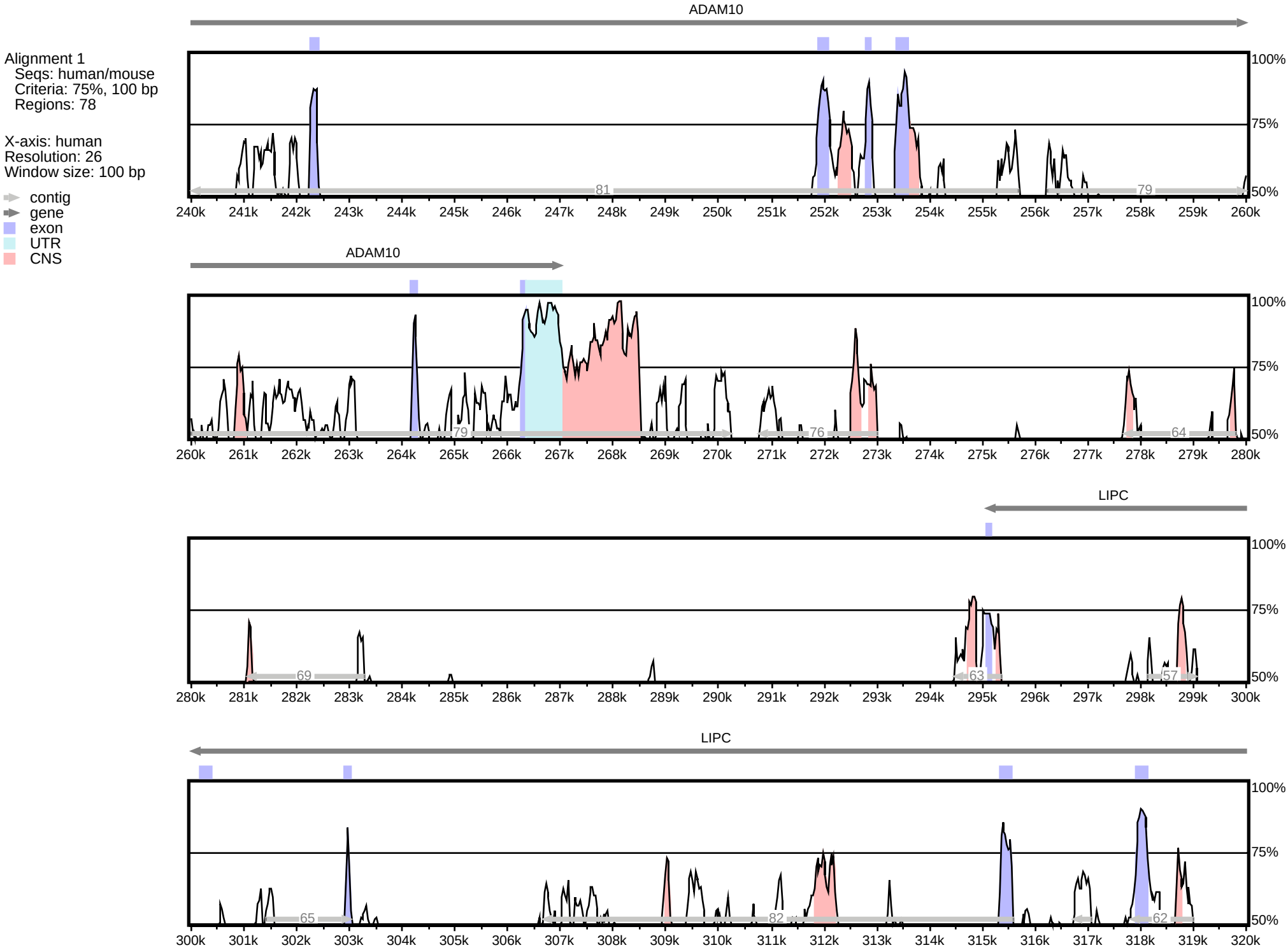
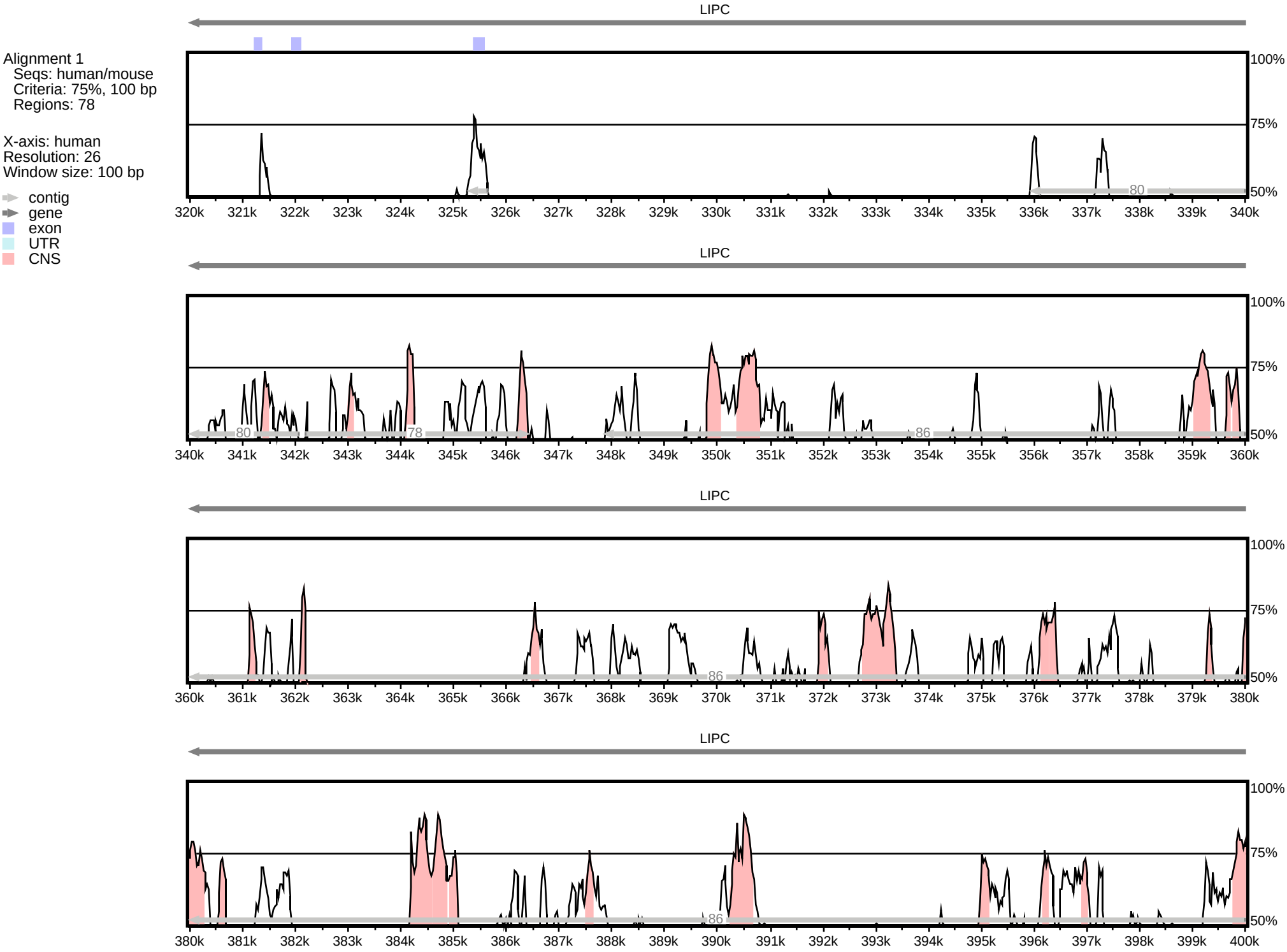

feature1
ADAM10
Alignment 1
Seqs: human/mouse
Criteria: 75%, 100 bp
Regions: 78
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
100%
75%
50%
160k 161k 162k 163k 164k 165k 166k 167k 168k 169k 170k 171k 172k 173k 174k 175k 176k 177k 178k 179k 180k
83
72
67
86
180k 181k 182k 183k 184k 185k 186k 187k 188k 189k 190k 191k 192k 193k 194k 195k 196k 197k 198k 199k 200k
86
71
200k 201k 202k 203k 204k 205k 206k 207k 208k 209k 210k 211k 212k 213k 214k 215k 216k 217k 218k 219k 220k
77
58
81
220k 221k 222k 223k 224k 225k 226k 227k 228k 229k 230k 231k 232k 233k 234k 235k 236k 237k 238k 239k 240k


Alignment 1
Seqs: human/mouse
Criteria: 75%, 100 bp
Regions: 78
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
ADAM10
81
79
100%
75%
50%
240k 241k 242k 243k 244k 245k 246k 247k 248k 249k 250k 251k 252k 253k 254k 255k 256k 257k 258k 259k 260k
ADAM10
79
76
64
260k 261k 262k 263k 264k 265k 266k 267k 268k 269k 270k 271k 272k 273k 274k 275k 276k 277k 278k 279k 280k
LIPC
69
63
57
280k 281k 282k 283k 284k 285k 286k 287k 288k 289k 290k 291k 292k 293k 294k 295k 296k 297k 298k 299k 300k
LIPC
65
82
62
300k 301k 302k 303k 304k 305k 306k 307k 308k 309k 310k 311k 312k 313k 314k 315k 316k 317k 318k 319k 320k


Alignment 1
Seqs: human/mouse
Criteria: 75%, 100 bp
Regions: 78
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
LIPC
100%
75%
50%
80
320k 321k 322k 323k 324k 325k 326k 327k 328k 329k 330k 331k 332k 333k 334k 335k 336k 337k 338k 339k 340k
LIPC
100%
75%
50%
80
78
86
340k 341k 342k 343k 344k 345k 346k 347k 348k 349k 350k 351k 352k 353k 354k 355k 356k 357k 358k 359k 360k
LIPC
100%
75%
50%
86
360k 361k 362k 363k 364k 365k 366k 367k 368k 369k 370k 371k 372k 373k 374k 375k 376k 377k 378k 379k 380k
LIPC
100%
75%
50%
86
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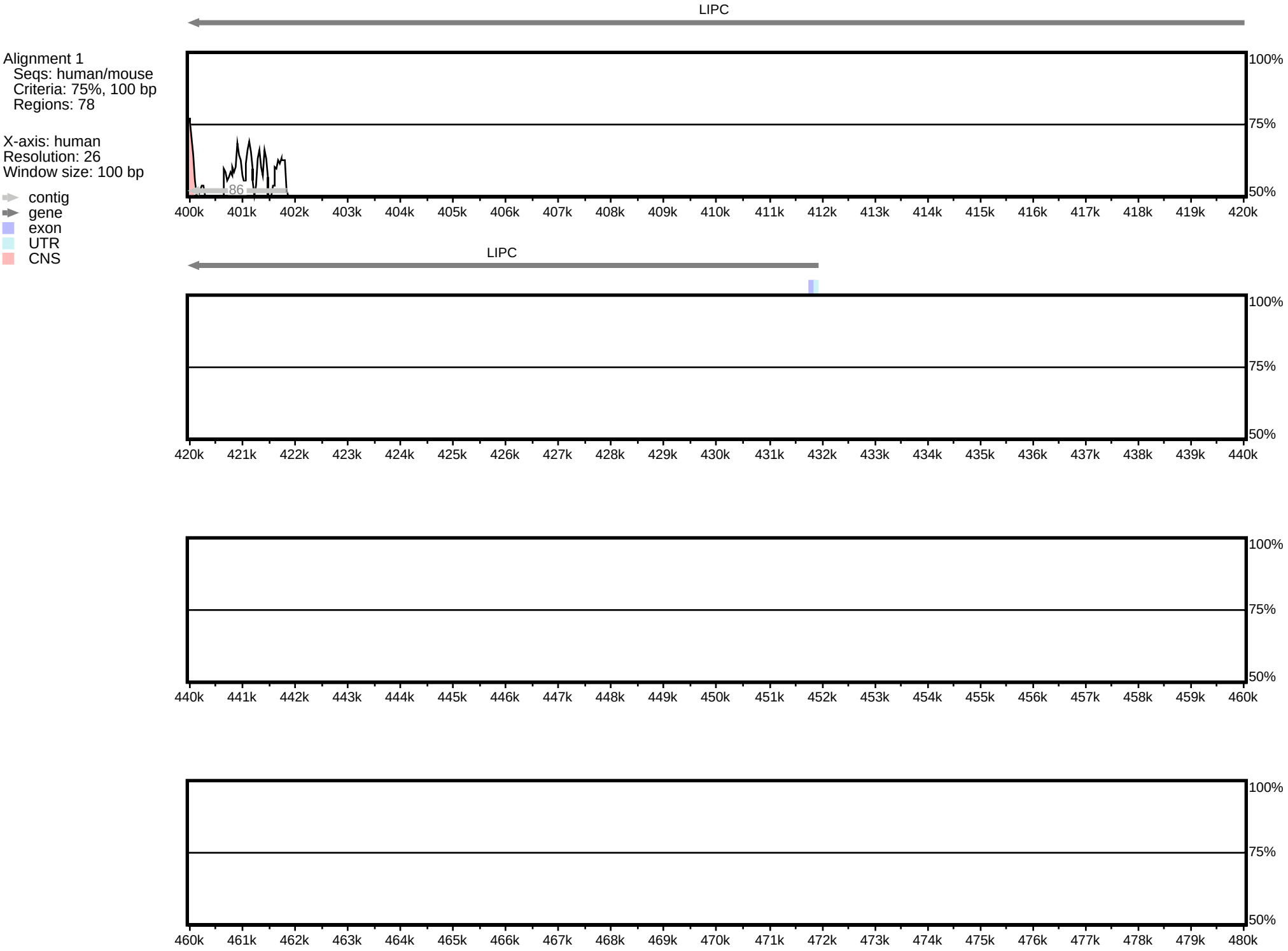
LIPC
Alignment 1
Seqs: human/mouse
Criteria: 75%, 100 bp
Regions: 78
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
86
100%
75%
50%
400k 401k 402k 403k 404k 405k 406k 407k 408k 409k 410k 411k 412k 413k 414k 415k 416k 417k 418k 419k 420k
LIPC
420k 421k 422k 423k 424k 425k 426k 427k 428k 429k 430k 431k 432k 433k 434k 435k 436k 437k 438k 439k 440k
440k 441k 442k 443k 444k 445k 446k 447k 448k 449k 450k 451k 452k 453k 454k 455k 456k 457k 458k 459k 460k
460k 461k 462k 463k 464k 465k 466k 467k 468k 469k 470k 471k 472k 473k 474k 475k 476k 477k 478k 479k 480k

Alignment 1
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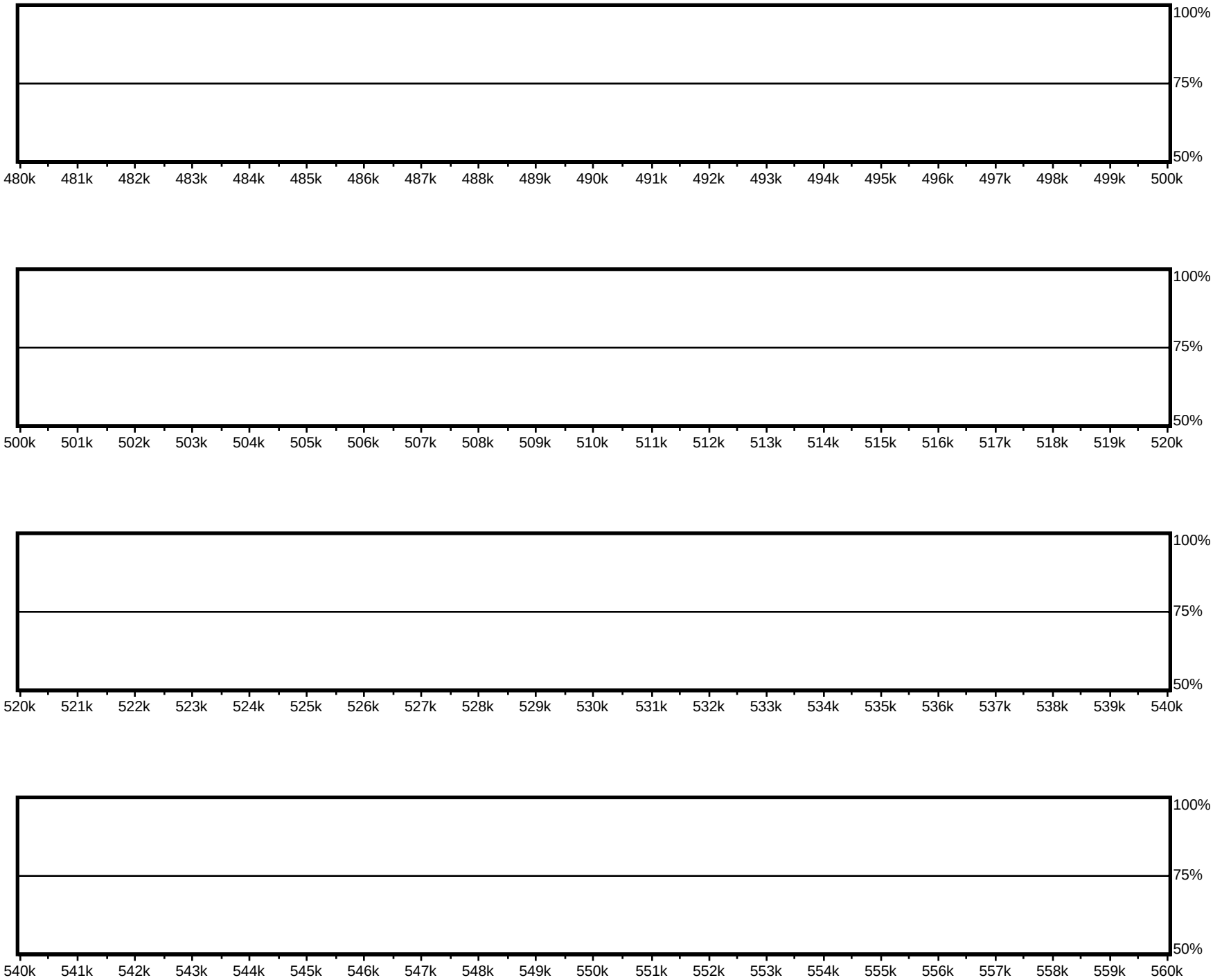
Window size: 100 bp

contig
gene
exon
UTR
CNS



Alignment 1
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Regions: 78

X-axis: human
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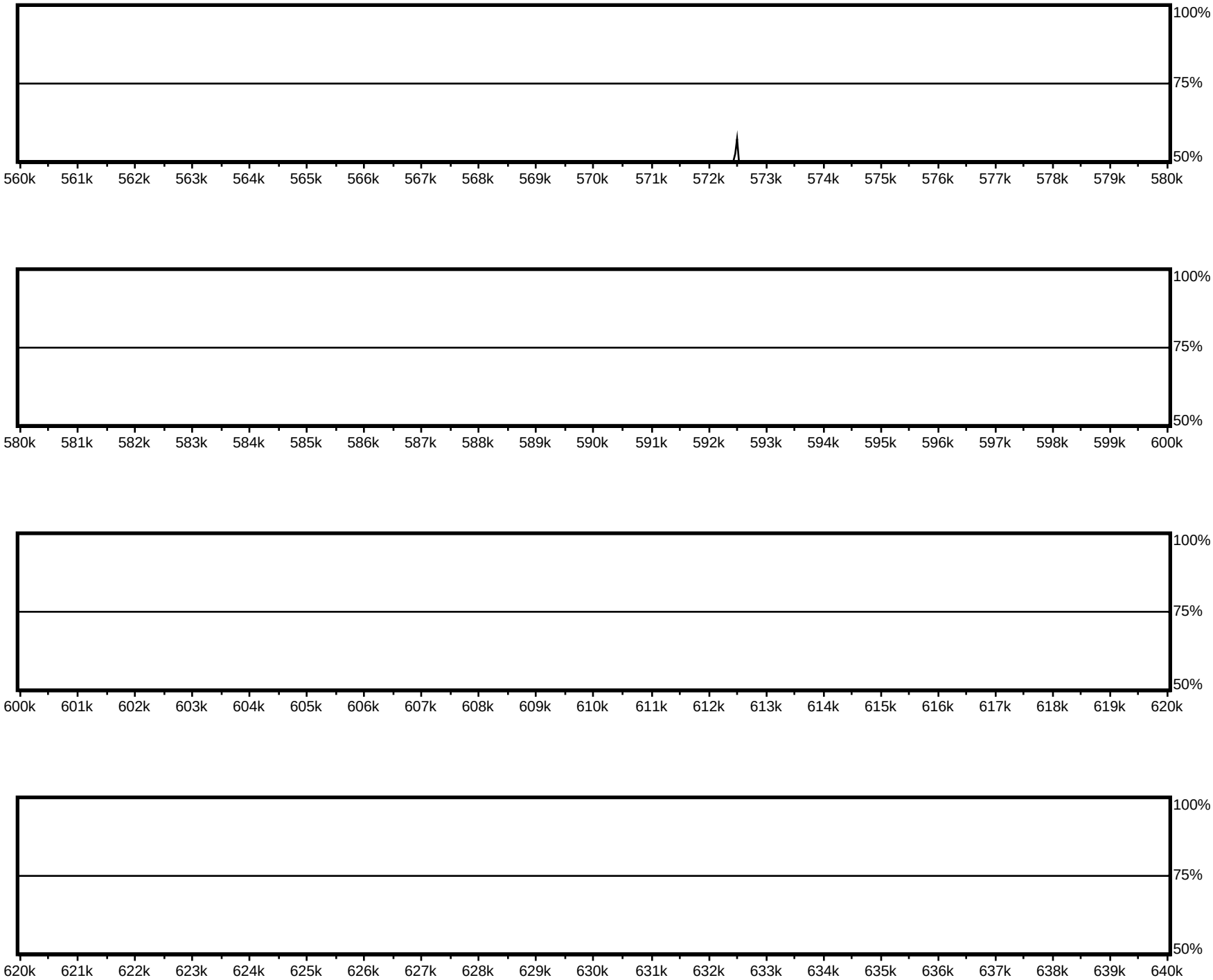
Window size: 100 bp

contig
gene
exon
UTR
CNS



Alignment 1
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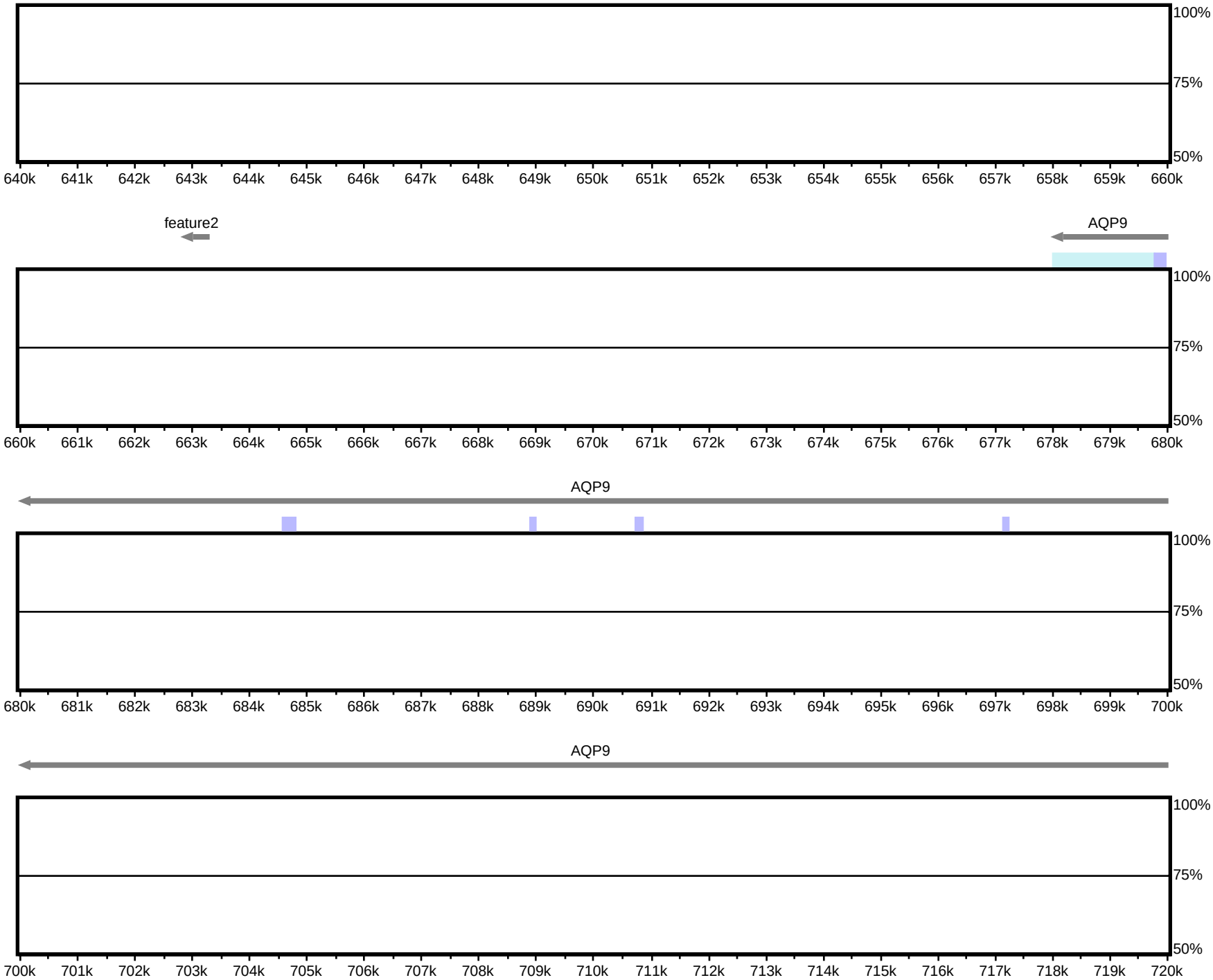
contig
gene
exon
UTR
CNS



Alignment 1
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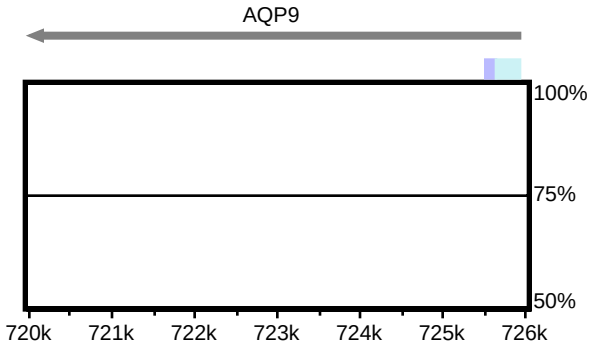
Regions: 78

X-axis: human
Resolution: 26
Window size: 100 bp

contig
gene
exon
UTR
CNS