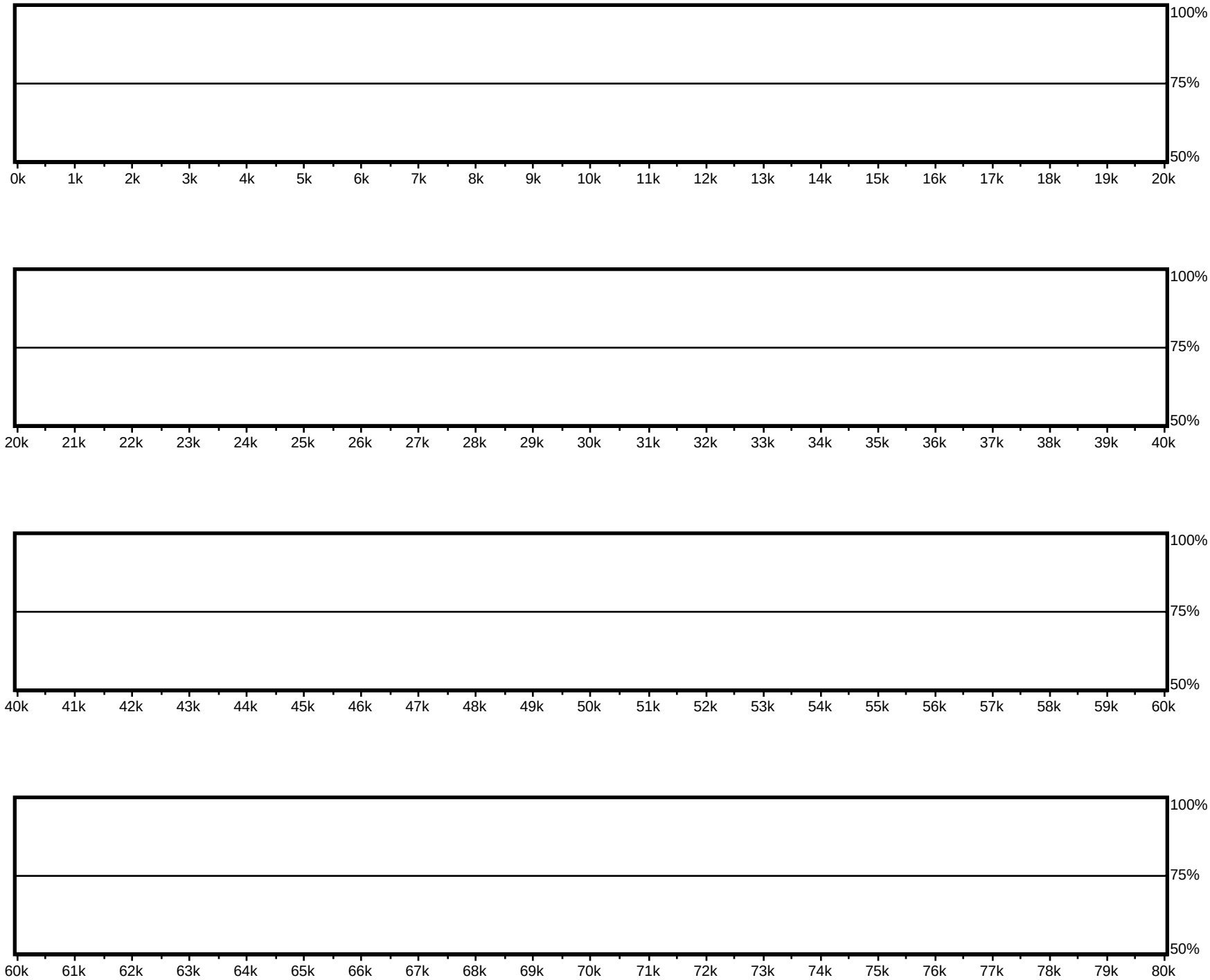
Alignment 1
Seqs: human/lemur
Criteria: 75%, 100 bp
Regions: 388

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

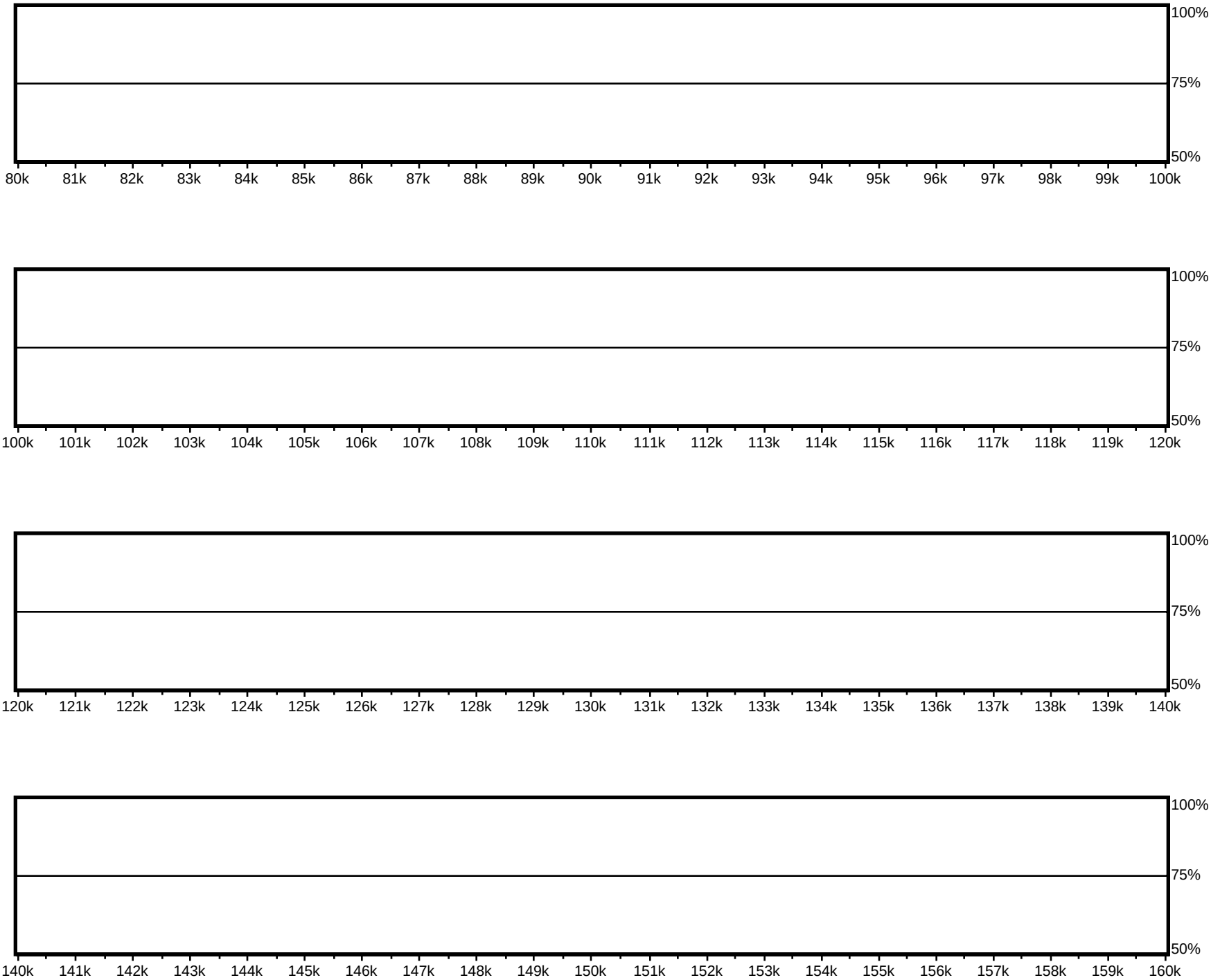
contig
gene
exon
UTR
CNS

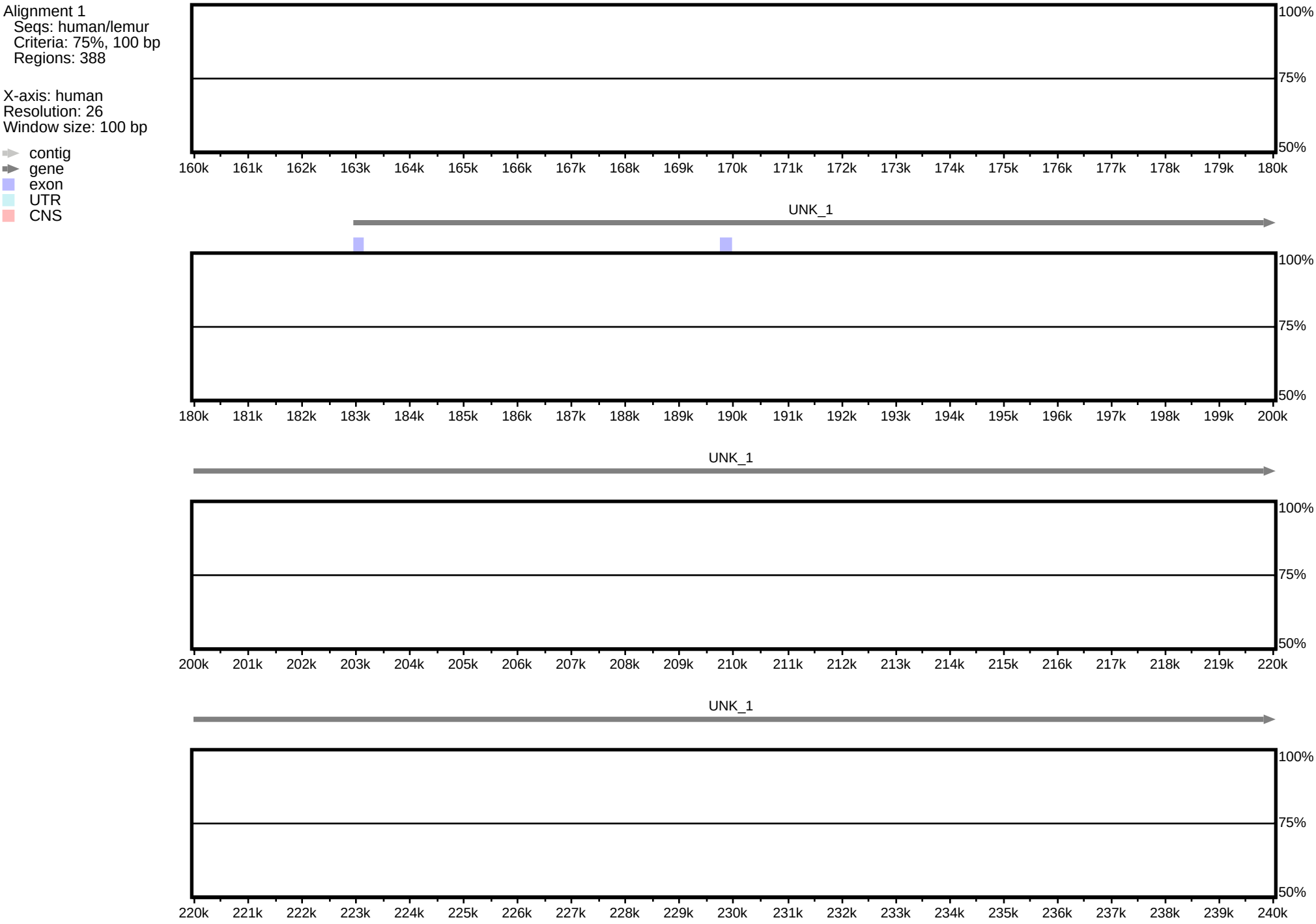


Alignment 1
Seqs: human/lemur
Criteria: 75%, 100 bp
Regions: 388

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

contig
gene
exon
UTR
CNS

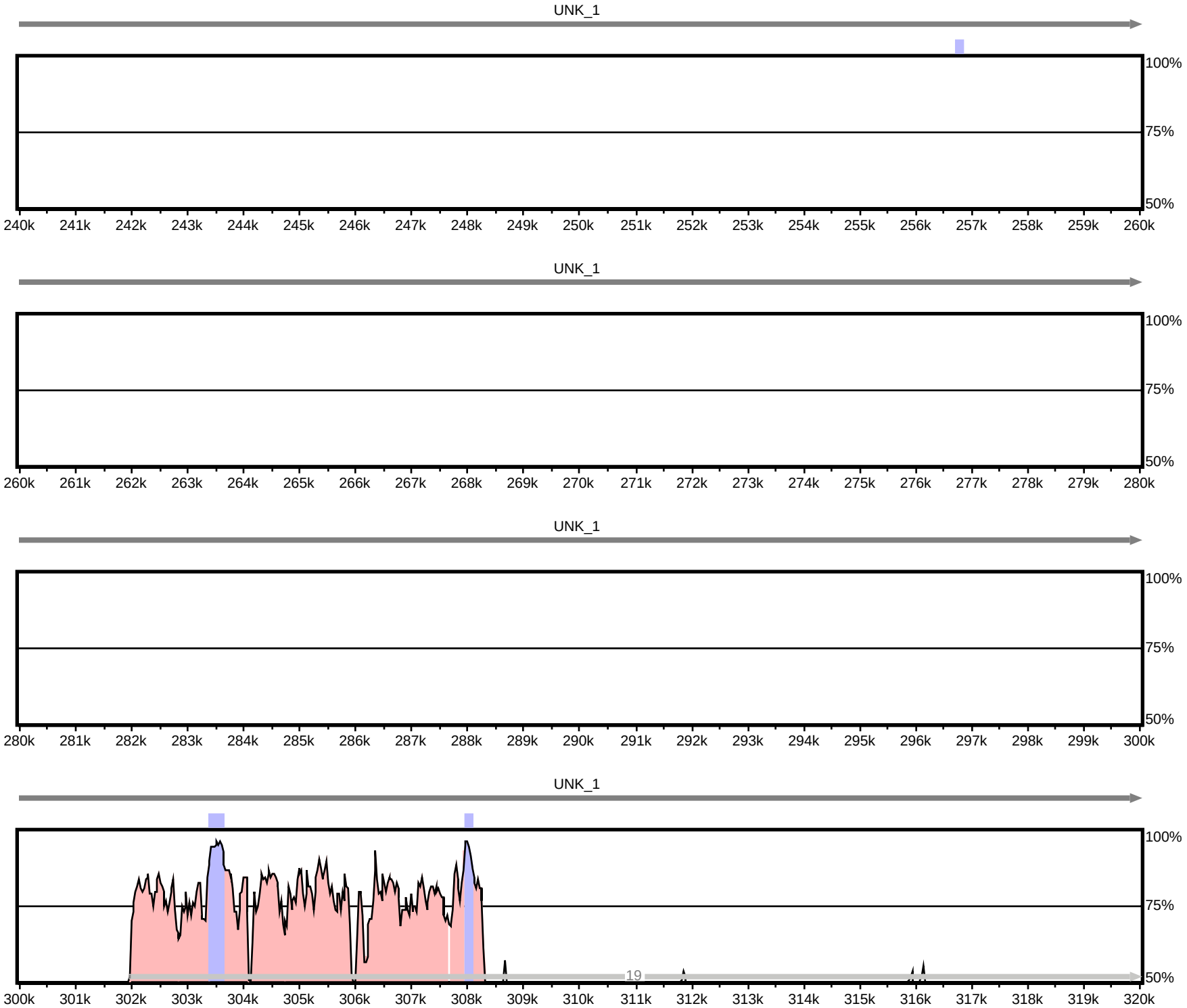



Alignment 1
Seqs: human/lemur
Criteria: 75%, 100 bp
Regions: 388
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
100%
75%
50%
160k 161k 162k 163k 164k 165k 166k 167k 168k 169k 170k 171k 172k 173k 174k 175k 176k 177k 178k 179k 180k
UNK_1
180k 181k 182k 183k 184k 185k 186k 187k 188k 189k 190k 191k 192k 193k 194k 195k 196k 197k 198k 199k 200k
UNK_1
200k 201k 202k 203k 204k 205k 206k 207k 208k 209k 210k 211k 212k 213k 214k 215k 216k 217k 218k 219k 220k
UNK_1
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/lemur
Criteria: 75%, 100 bp
Regions: 388

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

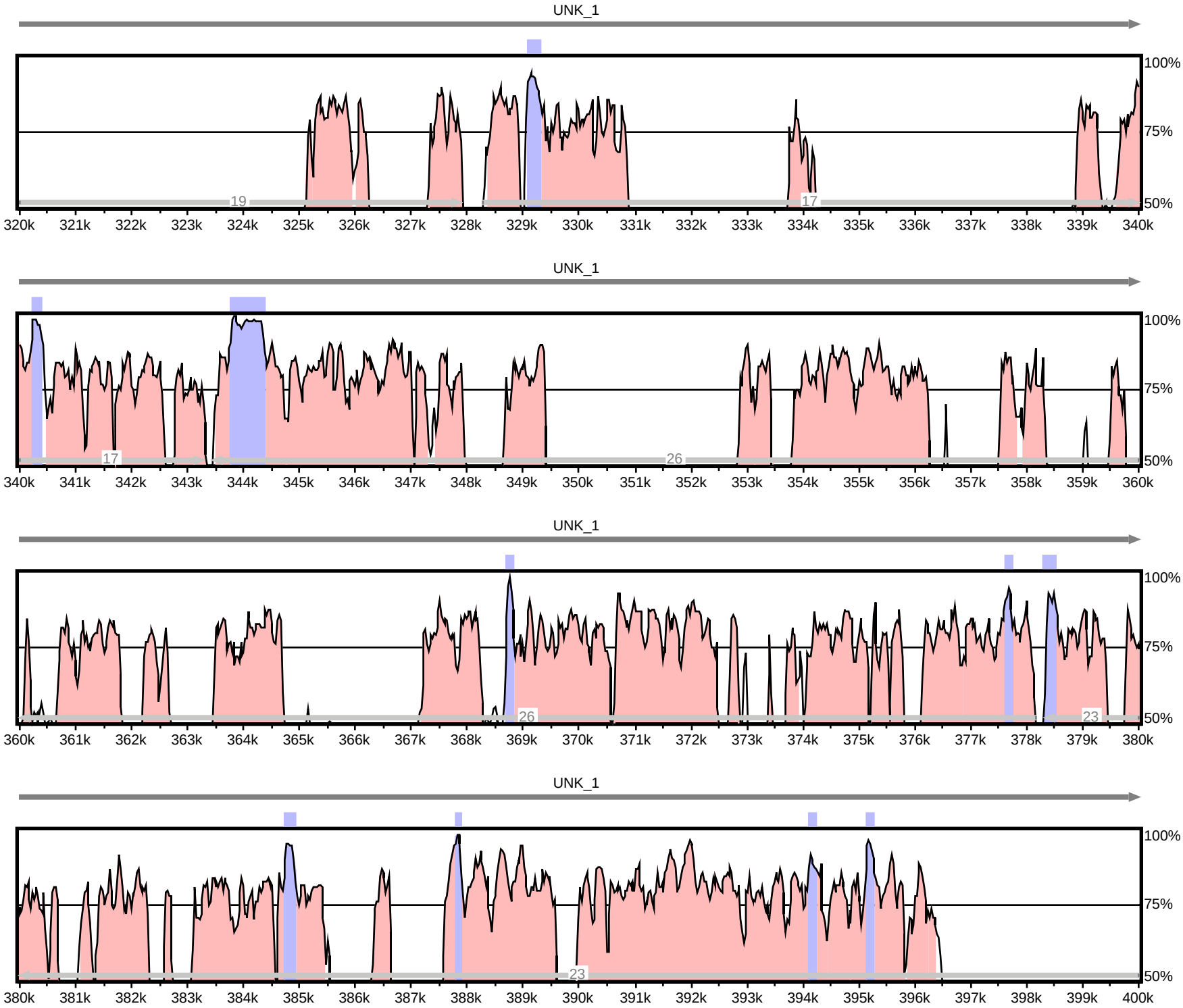
contig
gene
exon
UTR
CNS

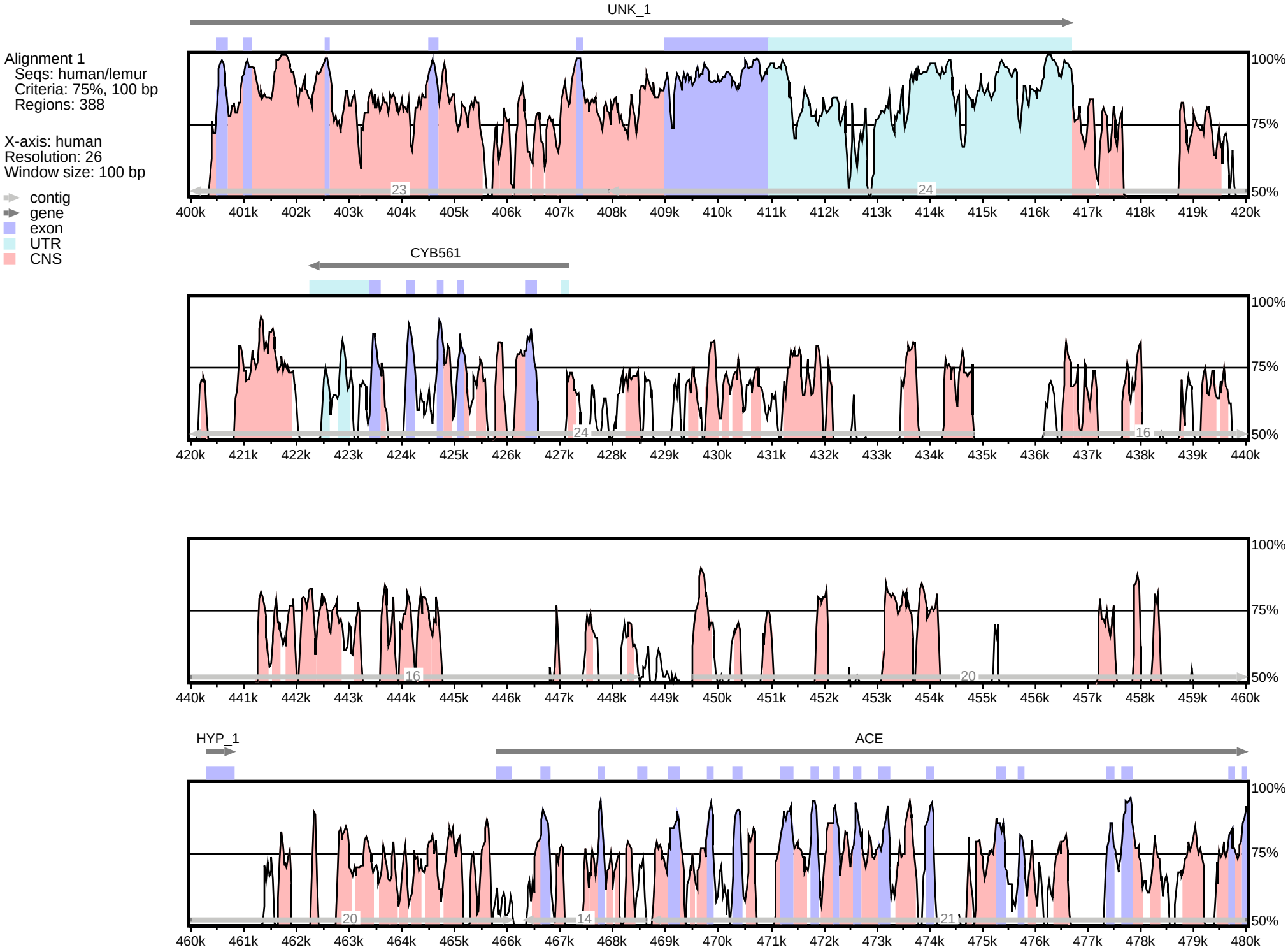


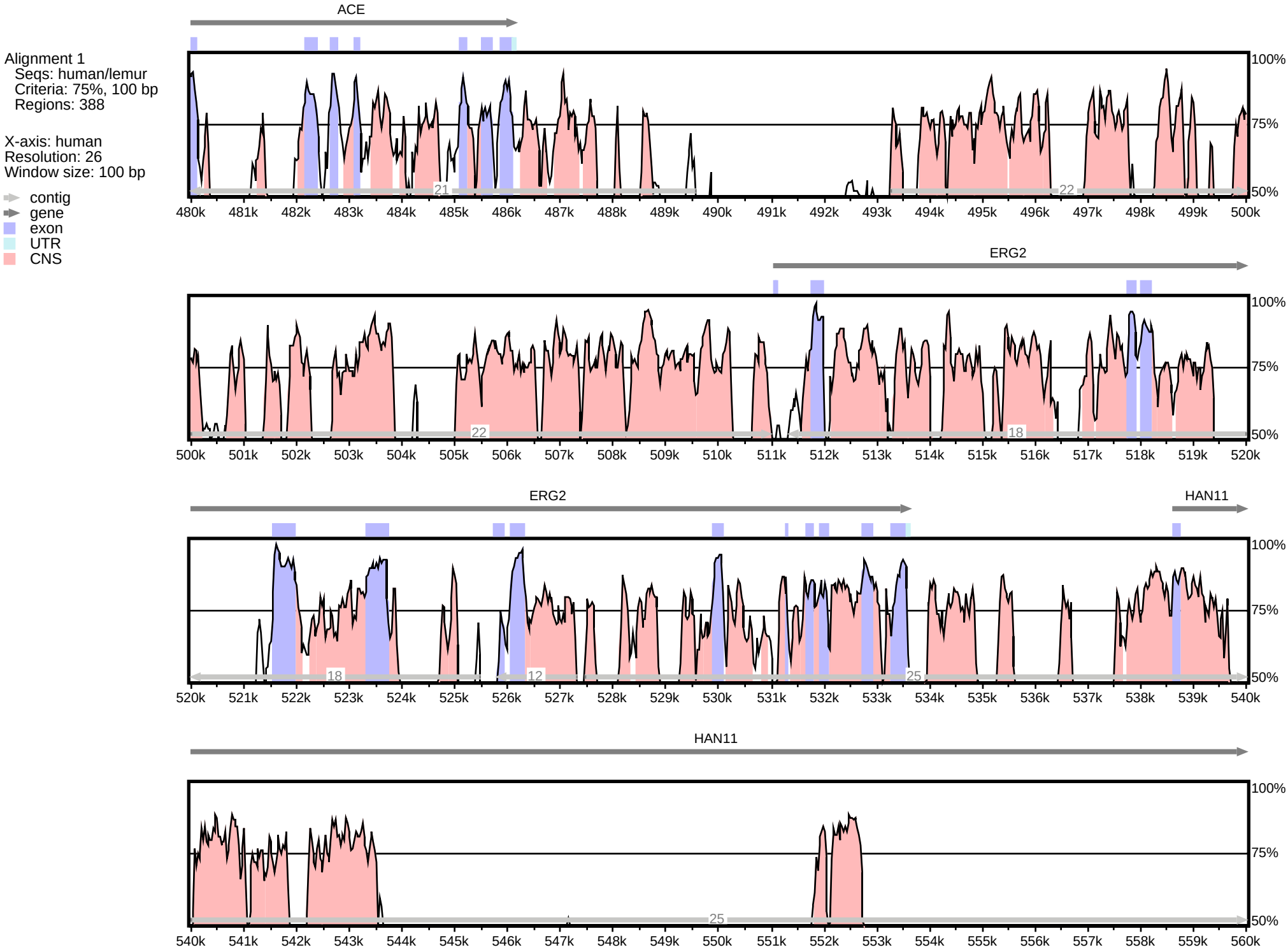
Alignment 1
Seqs: human/lemur
Criteria: 75%, 100 bp
Regions: 388

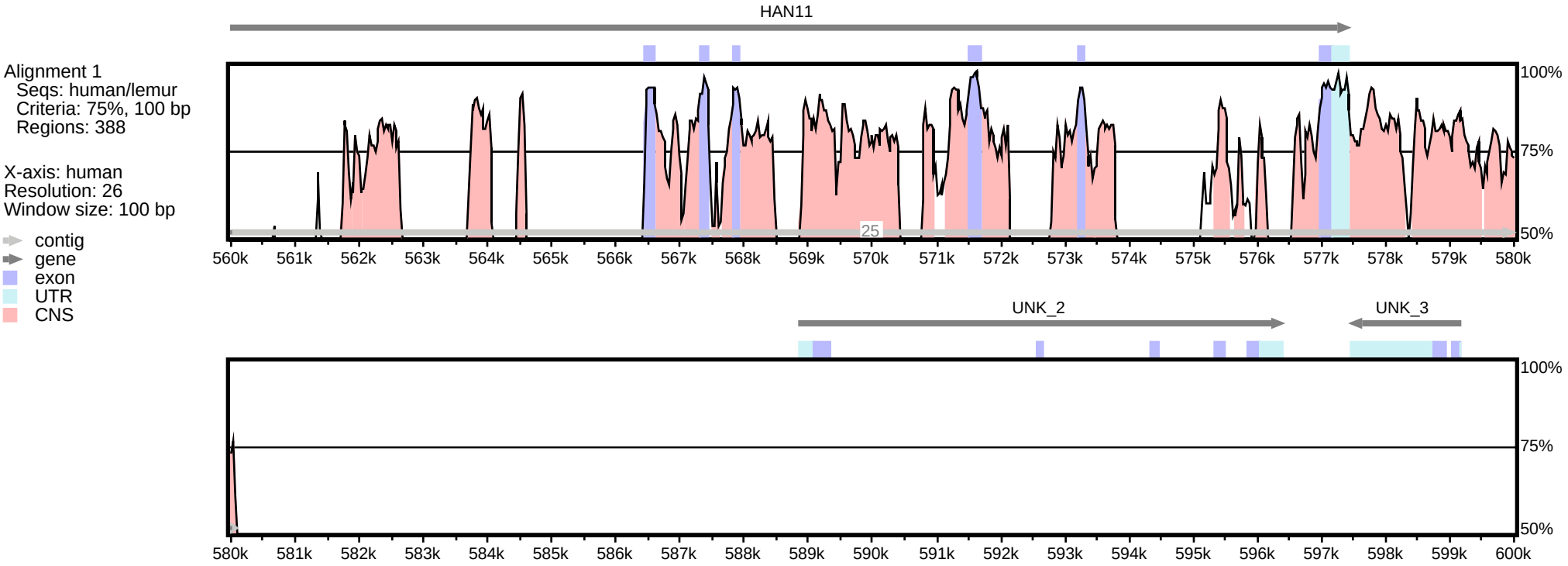
X-axis: human
Resolution: 26
Window size: 100 bp

contig
gene
exon
UTR
CNS




Alignment 1
Seqs: human/lemur
Criteria: 75%, 100 bp
Regions: 388
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
UNK_1
CYB561
HYP_1
ACE
100%
75%
50%
400k 401k 402k 403k 404k 405k 406k 407k 408k 409k 410k 411k 412k 413k 414k 415k 416k 417k 418k 419k 420k
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
Seqs: human/lemur
Criteria: 75%, 100 bp
Regions: 388
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
ACE
ERG2
HAN11
100%
75%
50%
480k 481k 482k 483k 484k 485k 486k 487k 488k 489k 490k 491k 492k 493k 494k 495k 496k 497k 498k 499k 500k
500k 501k 502k 503k 504k 505k 506k 507k 508k 509k 510k 511k 512k 513k 514k 515k 516k 517k 518k 519k 520k
520k 521k 522k 523k 524k 525k 526k 527k 528k 529k 530k 531k 532k 533k 534k 535k 536k 537k 538k 539k 540k
540k 541k 542k 543k 544k 545k 546k 547k 548k 549k 550k 551k 552k 553k 554k 555k 556k 557k 558k 559k 560k


HAN11
Alignment 1
Seqs: human/lemur
Criteria: 75%, 100 bp
Regions: 388
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
100%
75%
50%
25
560k
561k
562k
563k
564k
565k
566k
567k
568k
569k
570k
571k
572k
573k
574k
575k
576k
577k
578k
579k
580k
UNK_2
UNK_3
100%
75%
50%
580k
581k
582k
583k
584k
585k
586k
587k
588k
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