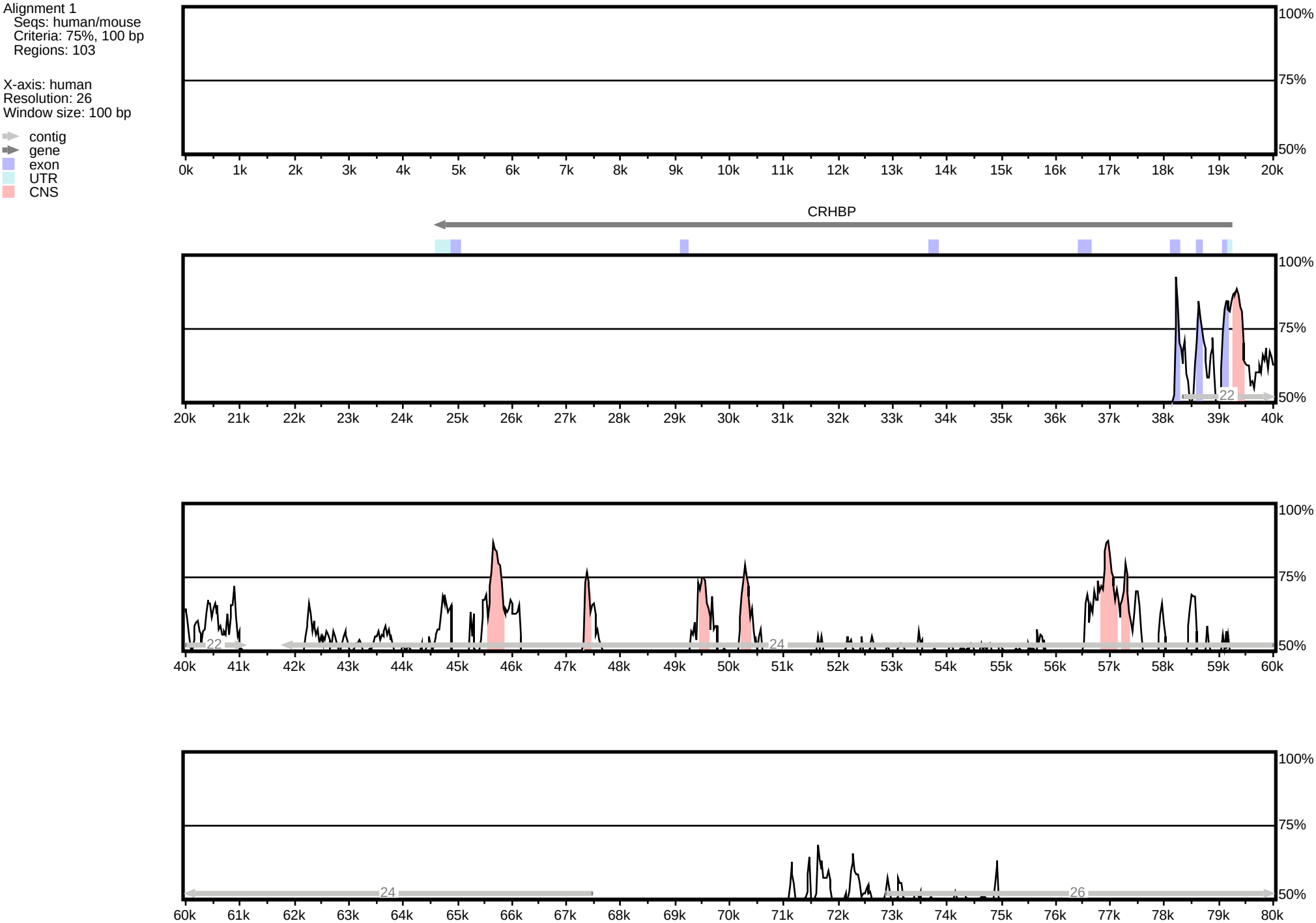
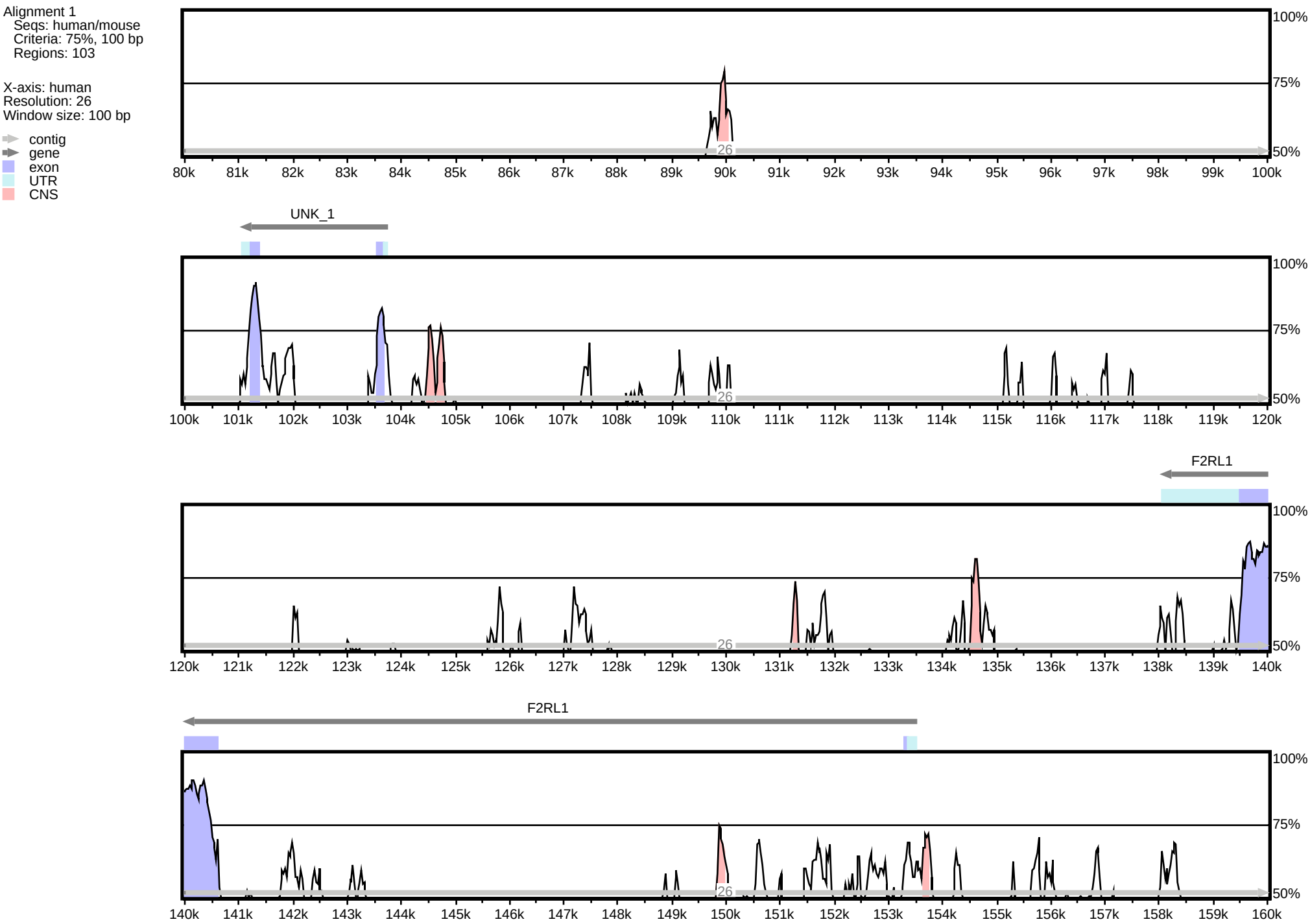
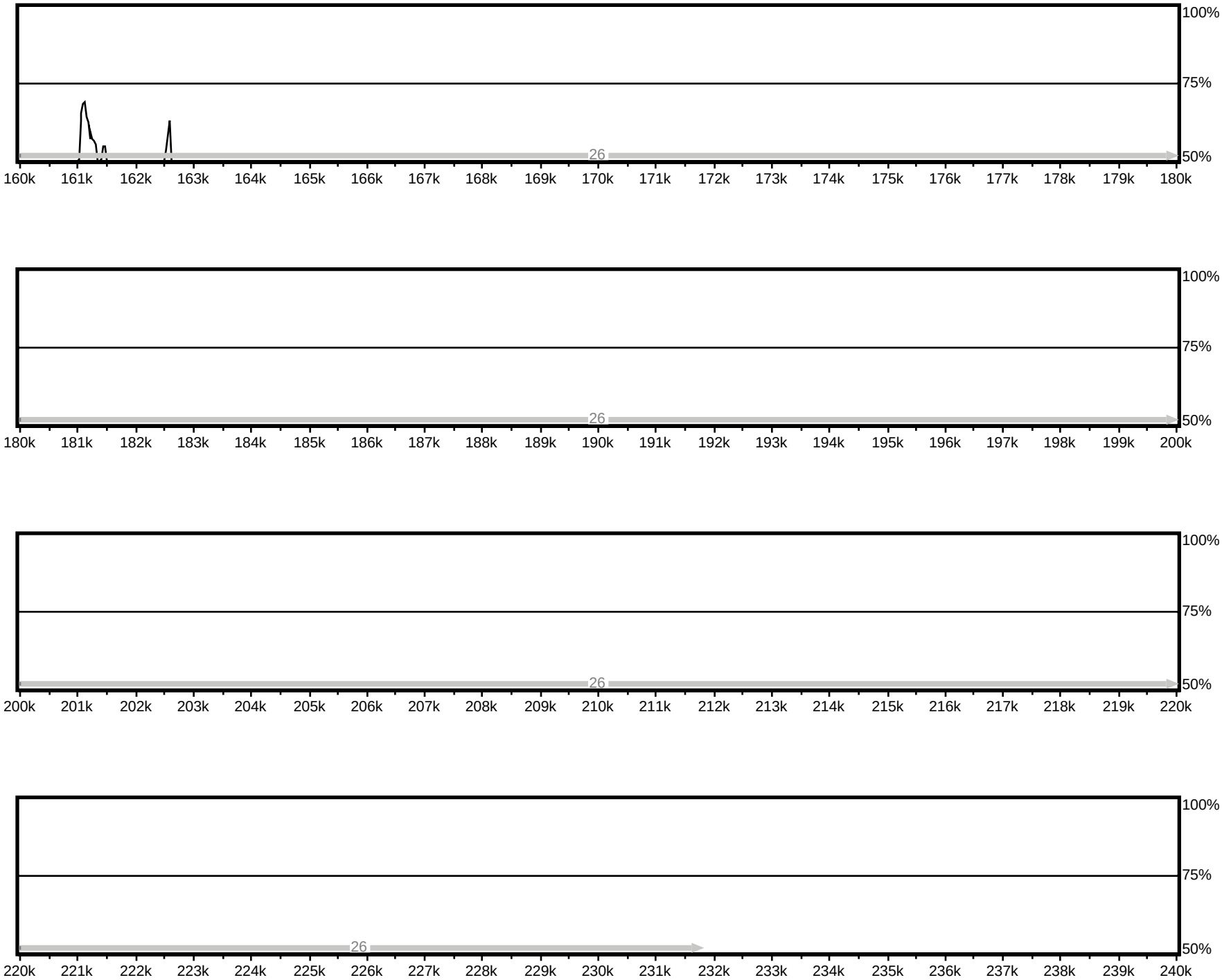

Alignment 1
Seqs: human/mouse
Criteria: 75%, 100 bp
Regions: 103
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
100%
75%
50%
0k 1k 2k 3k 4k 5k 6k 7k 8k 9k 10k 11k 12k 13k 14k 15k 16k 17k 18k 19k 20k
CRHBP
20k 21k 22k 23k 24k 25k 26k 27k 28k 29k 30k 31k 32k 33k 34k 35k 36k 37k 38k 39k 40k
22
40k 41k 42k 43k 44k 45k 46k 47k 48k 49k 50k 51k 52k 53k 54k 55k 56k 57k 58k 59k 60k
22
24
60k 61k 62k 63k 64k 65k 66k 67k 68k 69k 70k 71k 72k 73k 74k 75k 76k 77k 78k 79k 80k
24
26

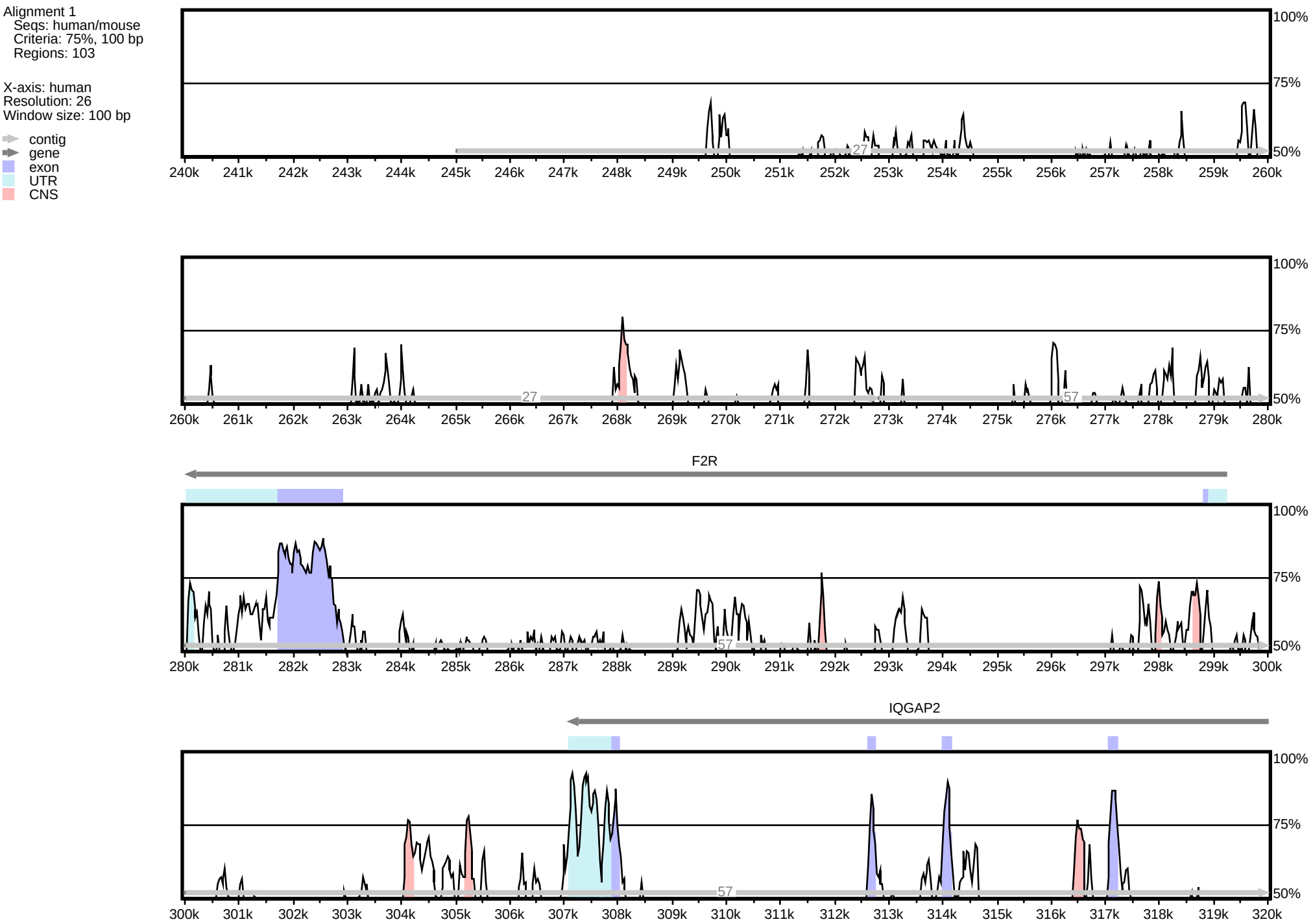

Alignment 1
Seqs: human/mouse
Criteria: 75%, 100 bp
Regions: 103
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
100%
75%
50%
26
80k 81k 82k 83k 84k 85k 86k 87k 88k 89k 90k 91k 92k 93k 94k 95k 96k 97k 98k 99k 100k
UNK_1
100k 101k 102k 103k 104k 105k 106k 107k 108k 109k 110k 111k 112k 113k 114k 115k 116k 117k 118k 119k 120k
F2RL1
120k 121k 122k 123k 124k 125k 126k 127k 128k 129k 130k 131k 132k 133k 134k 135k 136k 137k 138k 139k 140k
F2RL1
140k 141k 142k 143k 144k 145k 146k 147k 148k 149k 150k 151k 152k 153k 154k 155k 156k 157k 158k 159k 160k

Alignment 1
Seqs: human/mouse
Criteria: 75%, 100 bp
Regions: 103

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

contig
gene
exon
UTR
CNS




Alignment 1
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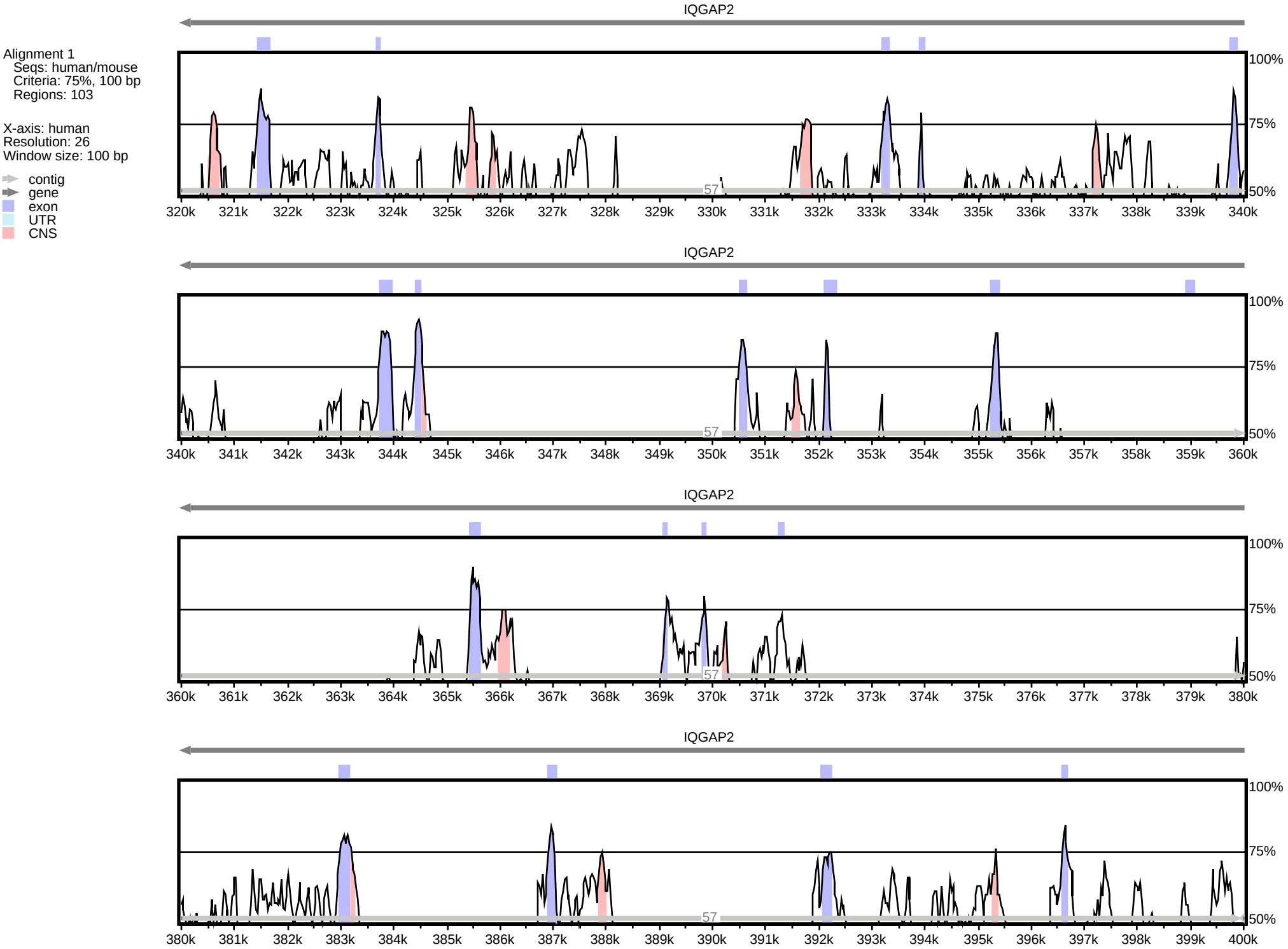
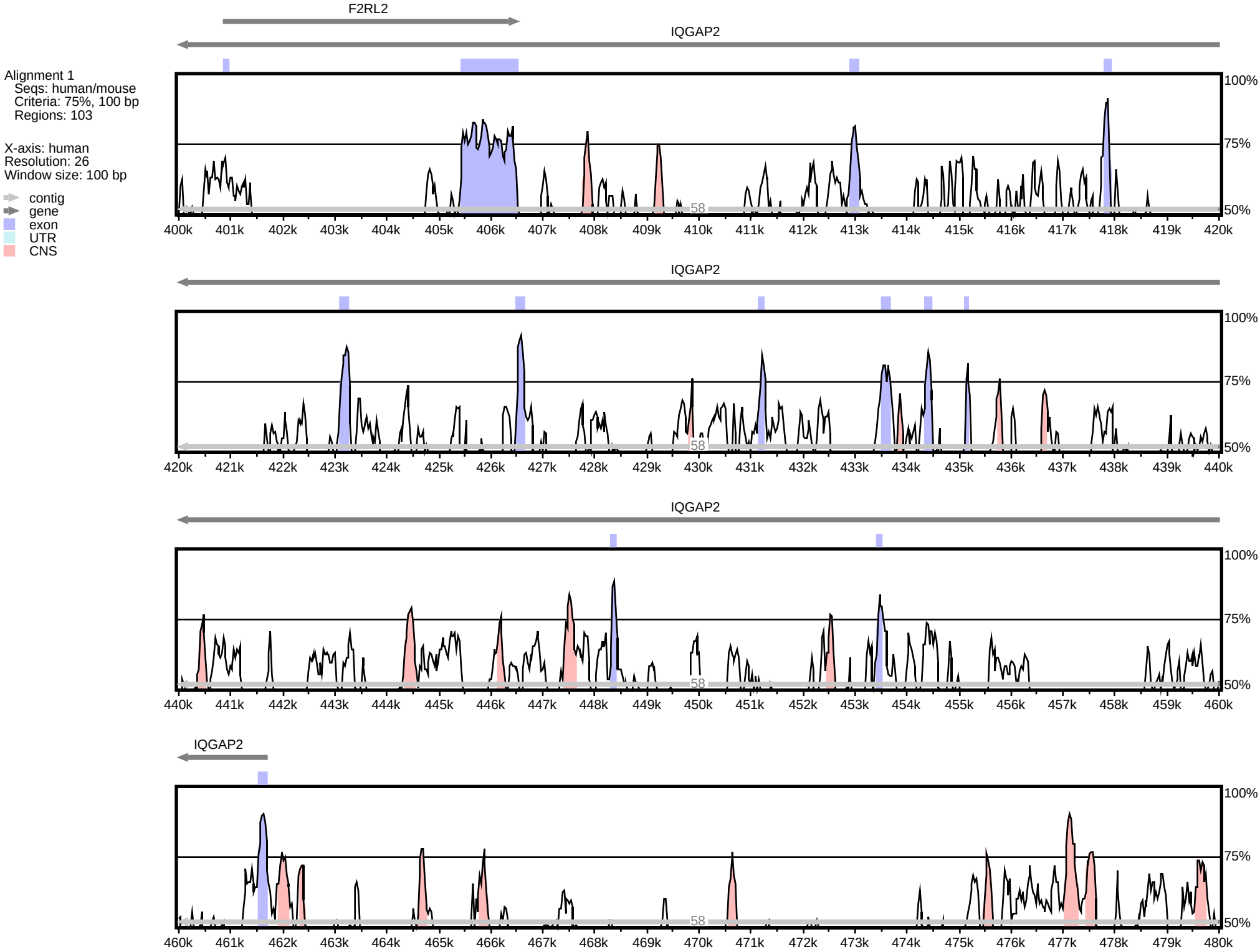
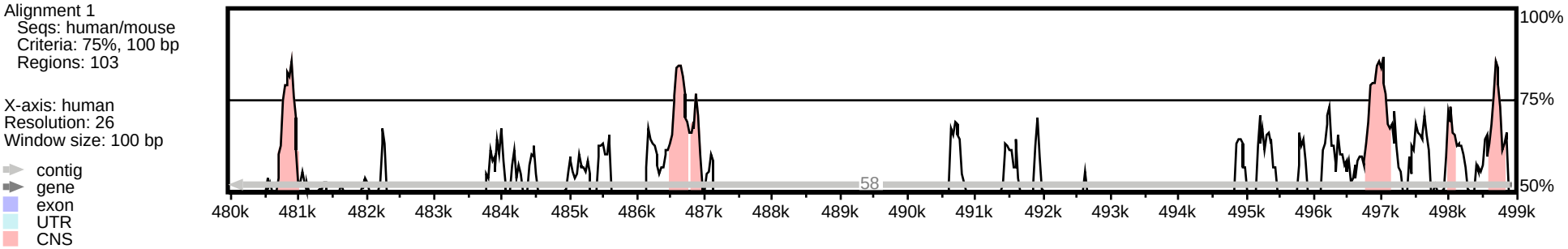
Criteria: 75%, 100 bp
Regions: 103
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
100%
75%
50%
27
240k 241k 242k 243k 244k 245k 246k 247k 248k 249k 250k 251k 252k 253k 254k 255k 256k 257k 258k 259k 260k
260k 261k 262k 263k 264k 265k 266k 267k 268k 269k 270k 271k 272k 273k 274k 275k 276k 277k 278k 279k 280k
F2R
57
280k 281k 282k 283k 284k 285k 286k 287k 288k 289k 290k 291k 292k 293k 294k 295k 296k 297k 298k 299k 300k
IQGAP2
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: 103
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
IQGAP2
100%
75%
50%
57
320k 321k 322k 323k 324k 325k 326k 327k 328k 329k 330k 331k 332k 333k 334k 335k 336k 337k 338k 339k 340k
IQGAP2
340k 341k 342k 343k 344k 345k 346k 347k 348k 349k 350k 351k 352k 353k 354k 355k 356k 357k 358k 359k 360k
IQGAP2
360k 361k 362k 363k 364k 365k 366k 367k 368k 369k 370k 371k 372k 373k 374k 375k 376k 377k 378k 379k 380k
IQGAP2
380k 381k 382k 383k 384k 385k 386k 387k 388k 389k 390k 391k 392k 393k 394k 395k 396k 397k 398k 399k 400k


F2RL2
IQGAP2
Alignment 1
Seqs: human/mouse
Criteria: 75%, 100 bp
Regions: 103
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
100%
75%
50%
58
400k 401k 402k 403k 404k 405k 406k 407k 408k 409k 410k 411k 412k 413k 414k 415k 416k 417k 418k 419k 420k
IQGAP2
420k 421k 422k 423k 424k 425k 426k 427k 428k 429k 430k 431k 432k 433k 434k 435k 436k 437k 438k 439k 440k
IQGAP2
440k 441k 442k 443k 444k 445k 446k 447k 448k 449k 450k 451k 452k 453k 454k 455k 456k 457k 458k 459k 460k
IQGAP2
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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Criteria: 75%, 100 bp
Regions: 103
X-axis: human
Resolution: 26
Window size: 100 bp
contig
gene
exon
UTR
CNS
100%
75%
50%
58
480k
481k
482k
483k
484k
485k
486k
487k
488k
489k
490k
491k
492k
493k
494k
495k
496k
497k
498k
499k