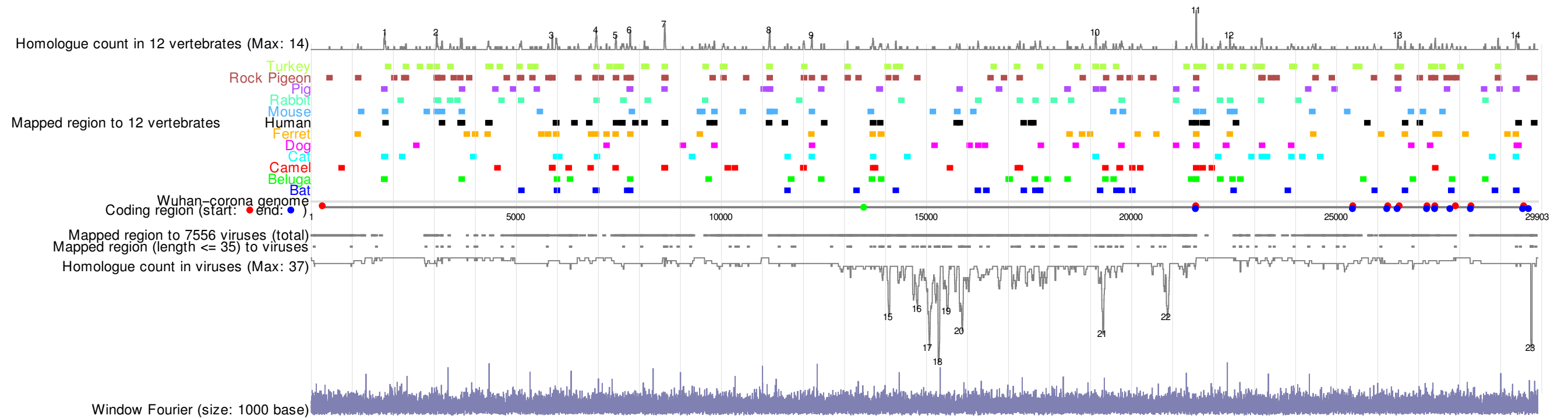


Homologue mapping



Frequent homologues (Vertebrates)

No.	homologue	start	end	count	in CDS
1	AATTTTA	1787	1793	6	Yes
2	GATGAGGATGAAGAAGAAGGT	3047	3067	6	Yes
3	CTACAAAGAAAACAGTTAC	5866	5884	5	Yes
4	ATAAATATTATAATTTG	6944	6960	7	Yes
5	TTGCAT	7425	7430	5	Yes
6	CCATCCATCTTTACTTTTGATAAA	7764	7786	7	Yes
7	TTTTTGTTGCTGCTATTTTCTATTTA	8607	8632	9	Yes
8	ATTTCTCTGTTTGTTTTGT	11167	11186	7	Yes
9	AAAAGT	12198	12203	5	Yes
10	ATAAAATAGAAGAA	19121	19134	6	Yes
11	TGTTTGTTTTCTTGTTTTATTGC	21564	21587	14	Yes
12	TATTAAAATATAATGAA	22389	22405	5	Yes
13	TTTTCT	26495	26500	5	No
14	CCTAA	29375	29379	5	Yes

Frequent homologues (Viruses)

No.	homologue	start	end	count	in CDS
15	CAAGATCTCAATGGTAACTGGTATGATTTCCGGTGATT	14068	14104	21	Yes
16	TGGTAATGCTGC	14775	14786	18	Yes
17	ACAAAACGTAATGTCATCCCTACTATAACTCAAATGAATCTTAAGTATGCCATTAGTGCAAAGAATAGAGCTCGCACCGTAGCTGGTGTCTCTAT	15031	15125	32	Yes
18	CTTATGGGTTGGGATTATCCTAAATGTGATAGAGCCATGCCTAA	15280	15323	37	Yes
19	GATGCCACAACCTGCTTATGCTAATAGTGTTTTTAACAT	15490	15527	19	Yes
20	CAAAACAATGTTTTATGTCTGAAGCAAAATGTTGGACTGAGACTGACCTTACTAAAGGACCTCATGAATTTTGCTCTCAACATACA	15805	15891	26	Yes
21	GGTTGTGATGGTGGCAGTTTGTATGTAAATAAACATGCATTCCACACACC	19276	19325	27	Yes
22	TATGAGAGTTATACATTTTGGTGCTGGTTCTGATAAAGGAGTTGCACCAGGTAC	20850	20903	21	Yes
23	CCGAGGCCACGCGGAGTACGATCGAGTGTACAG	29732	29764	32	No