**Draft genome of *Cicer reticulatum* L., the wild progenitor of chickpea provides insights into the domestication process**

SUPPORTING TEXT

**Text S1.** CDS and Protein sequences of 4 RGH obtained by mapping the identified RGF in the advanced draft genome of chickpea ICC4958, onto the *C. reticulatum* genome.

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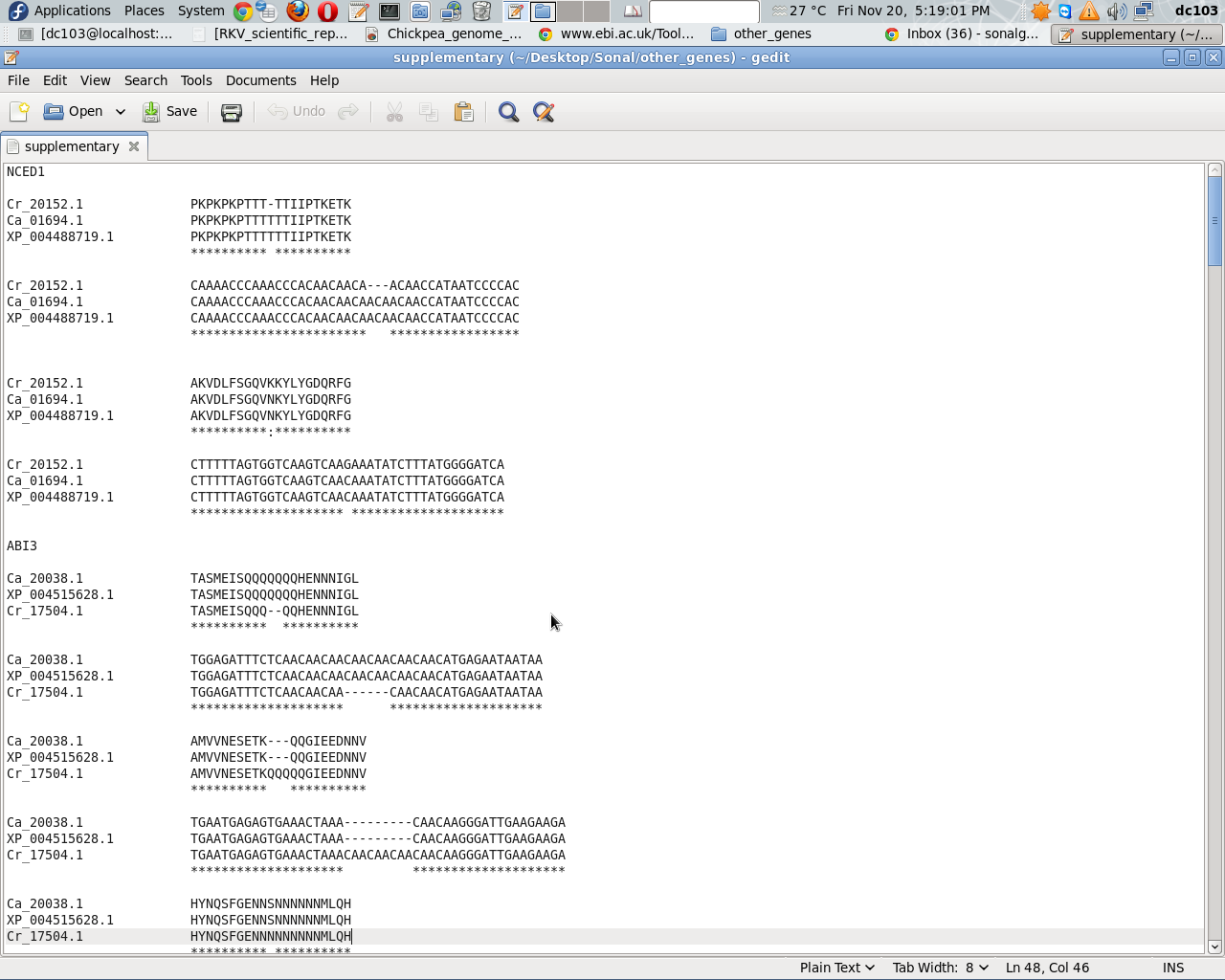
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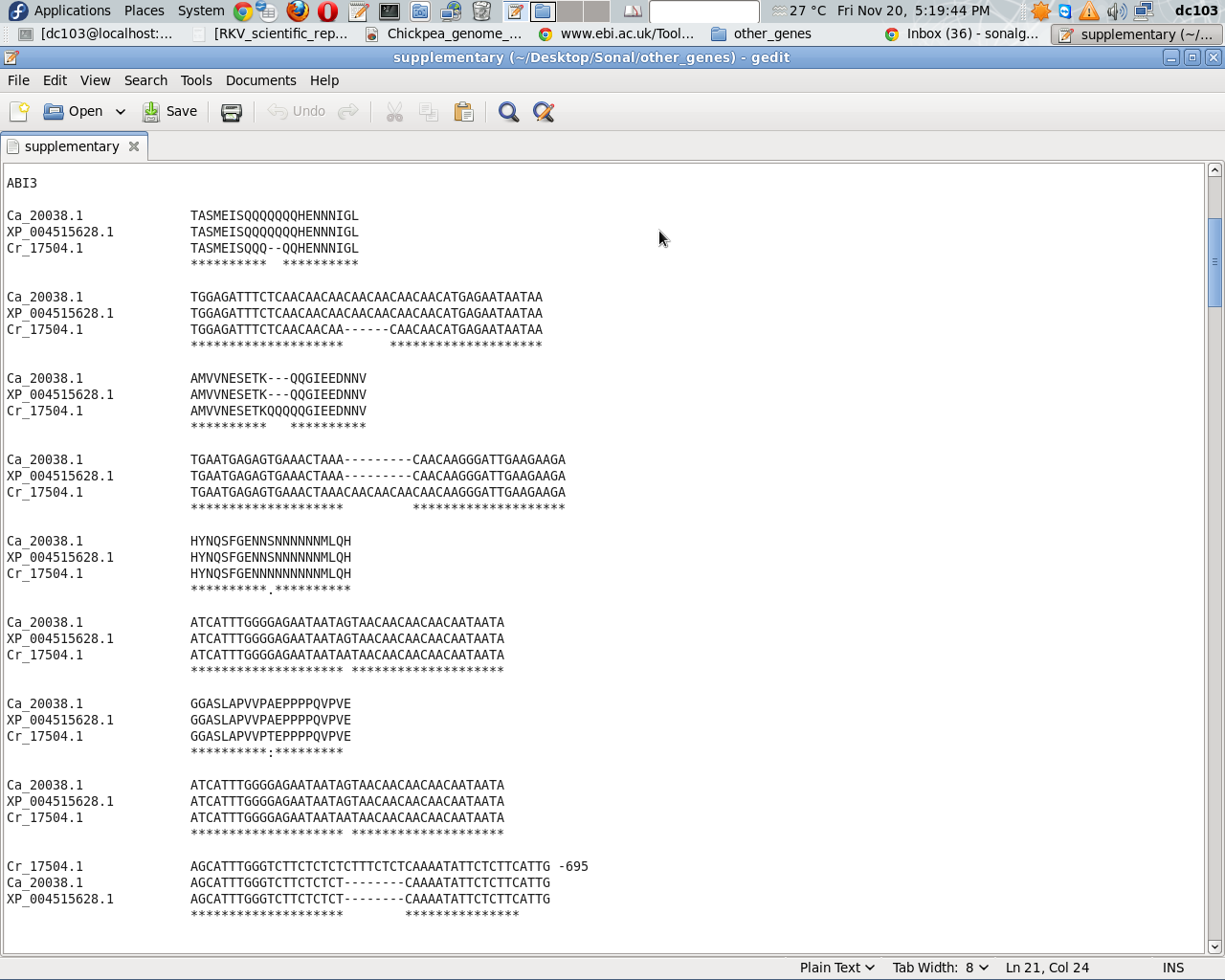
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**Text S2.** Sequence alignment of domestication-associated candidate gene orthologues in cultivated and wild chickpea. Pepetide and CDS sequences were aligned using ClustalW. Functional annotations are mentioned at the top of the box. Cr represents *C. reticulatum*, Ca represents C. arietinum ICC4958 (*desi*). GenBank accession numbers (XP) are used to denote genes/peptides of chickpea CDC Frontier (*kabuli*).

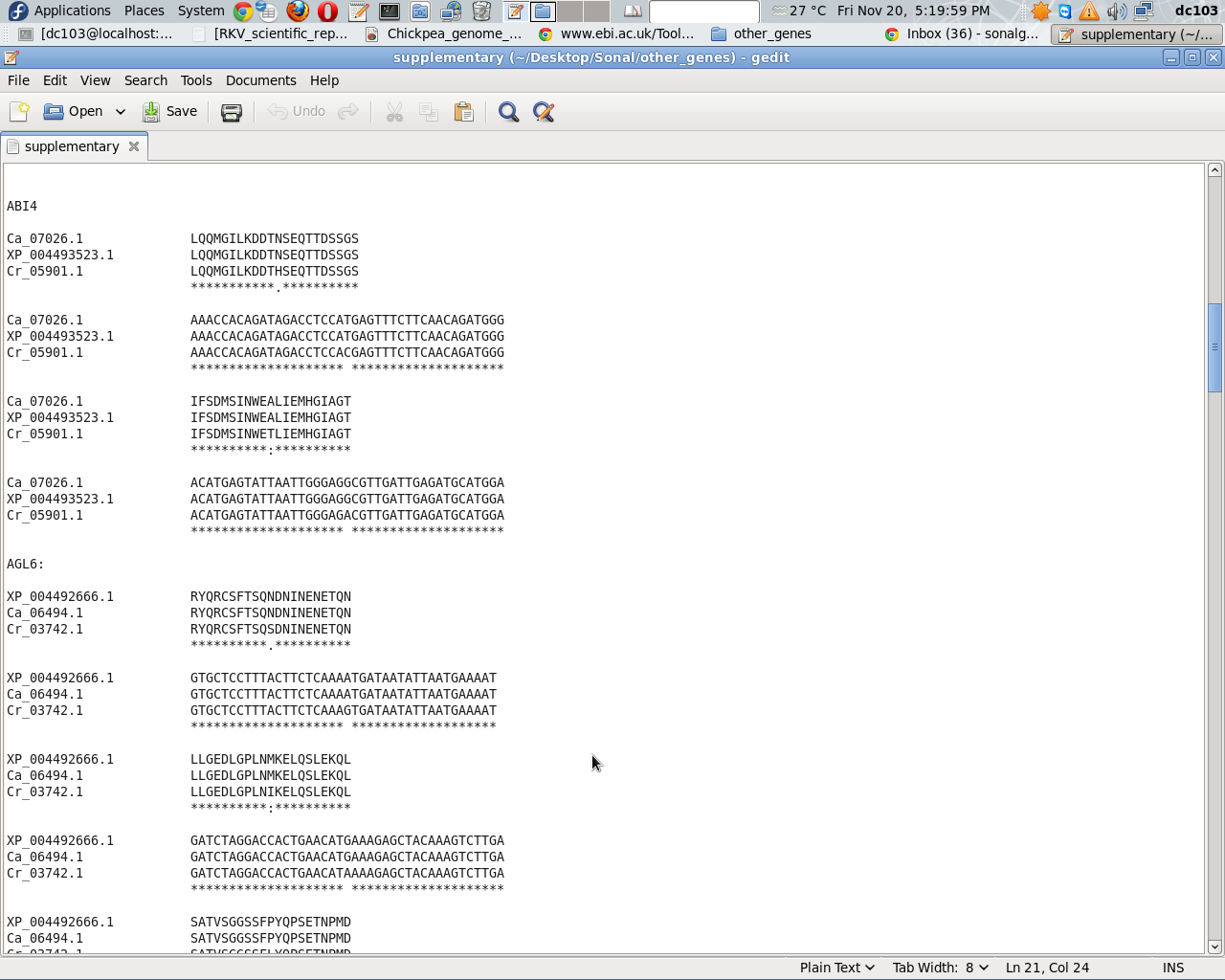
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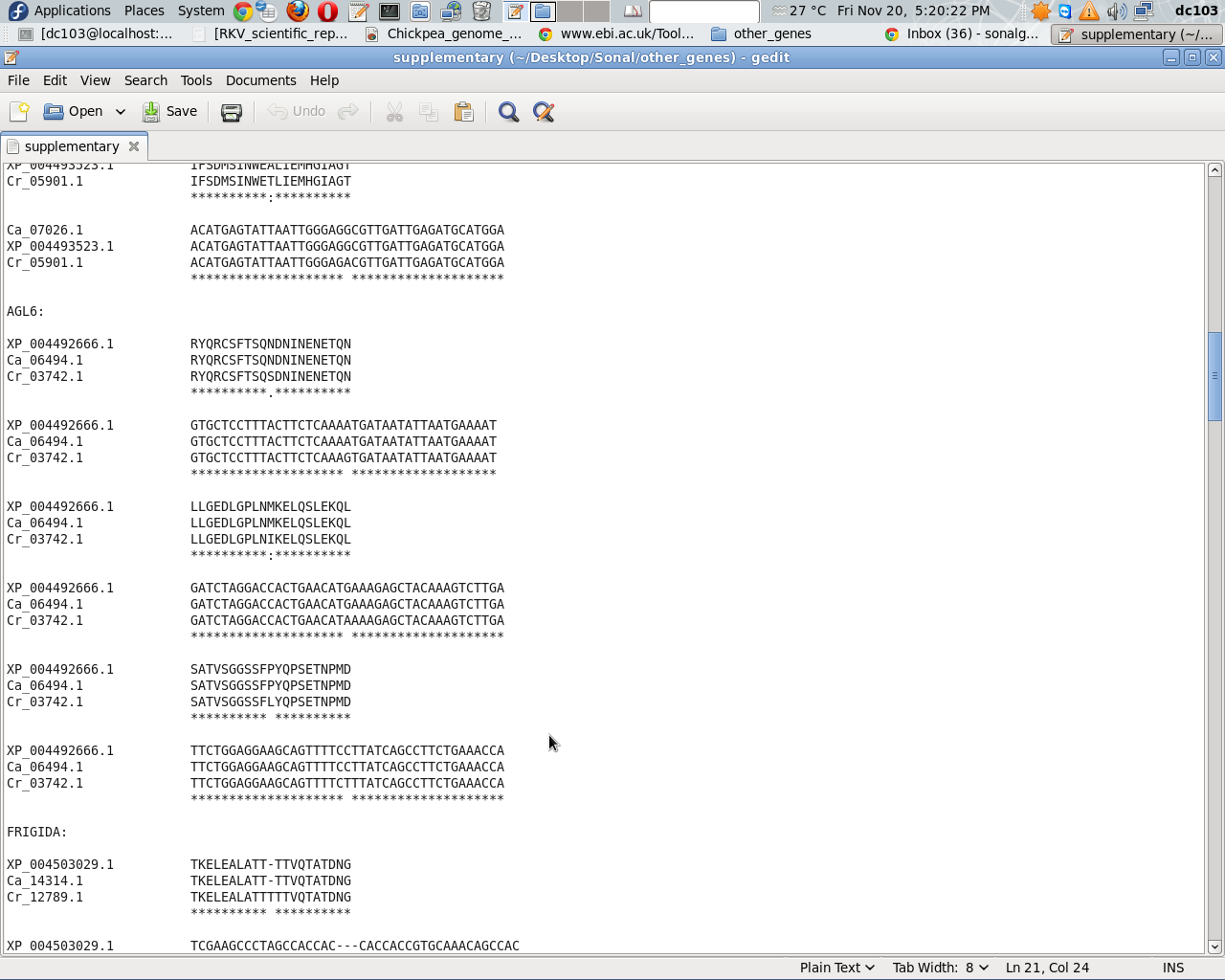
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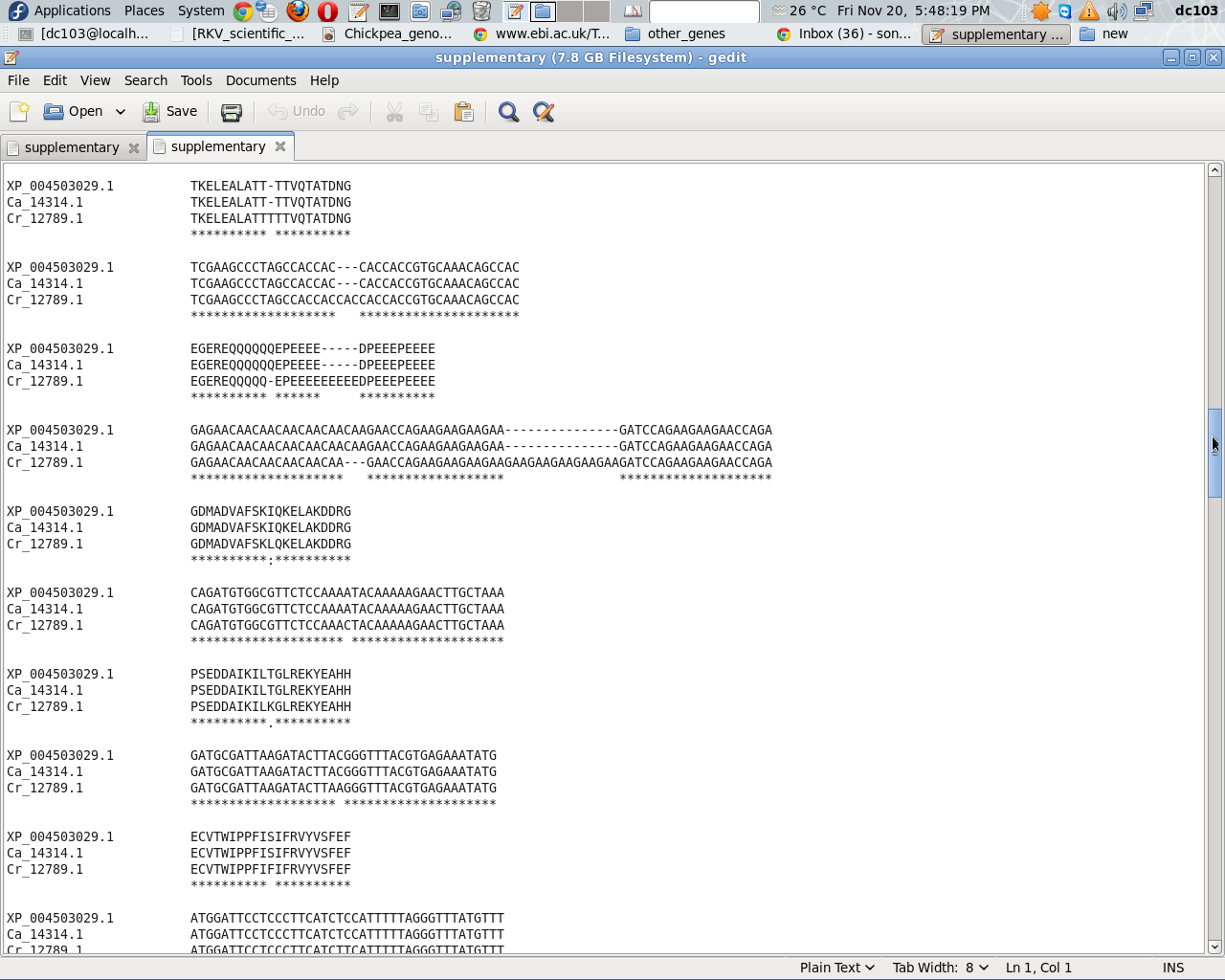
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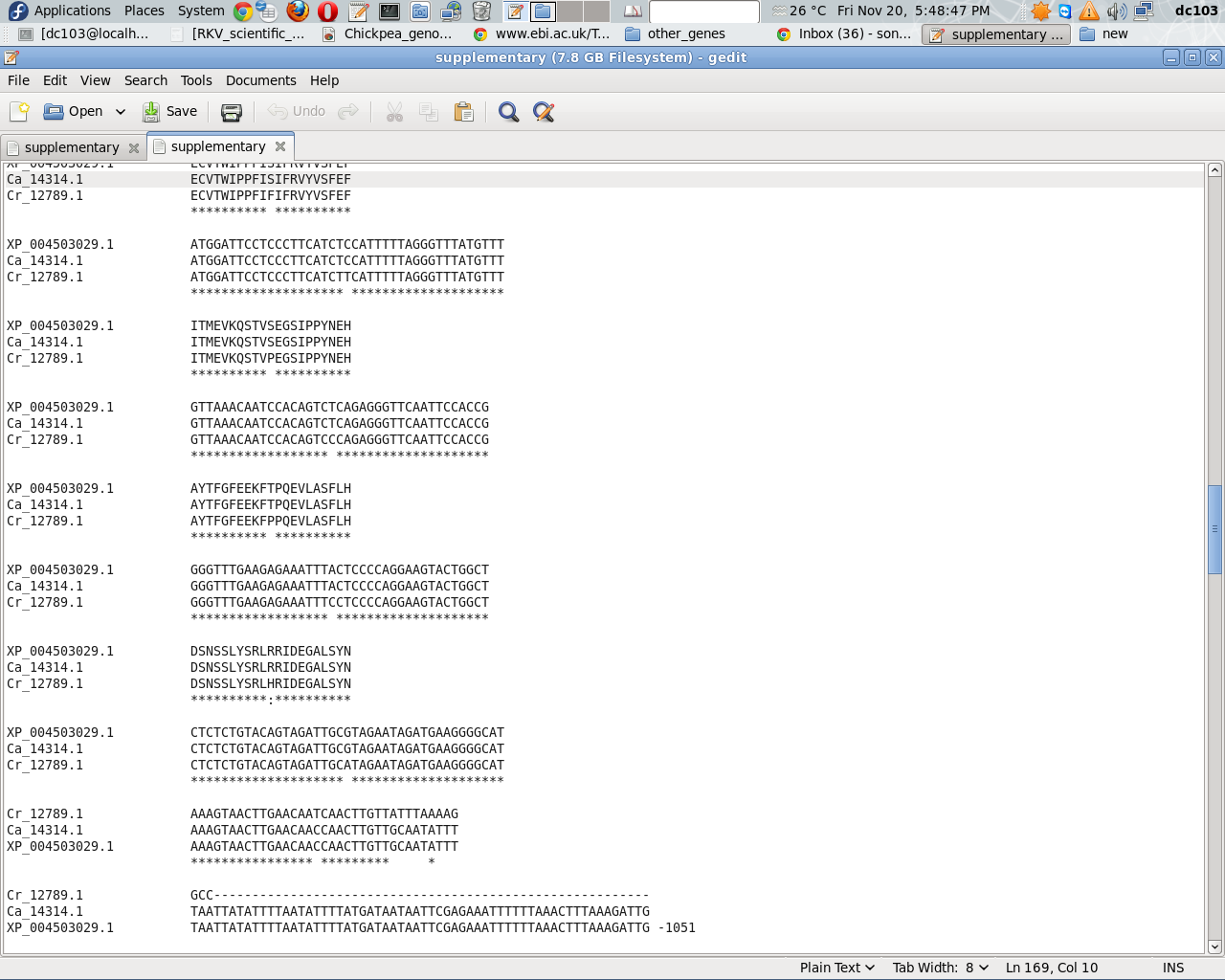


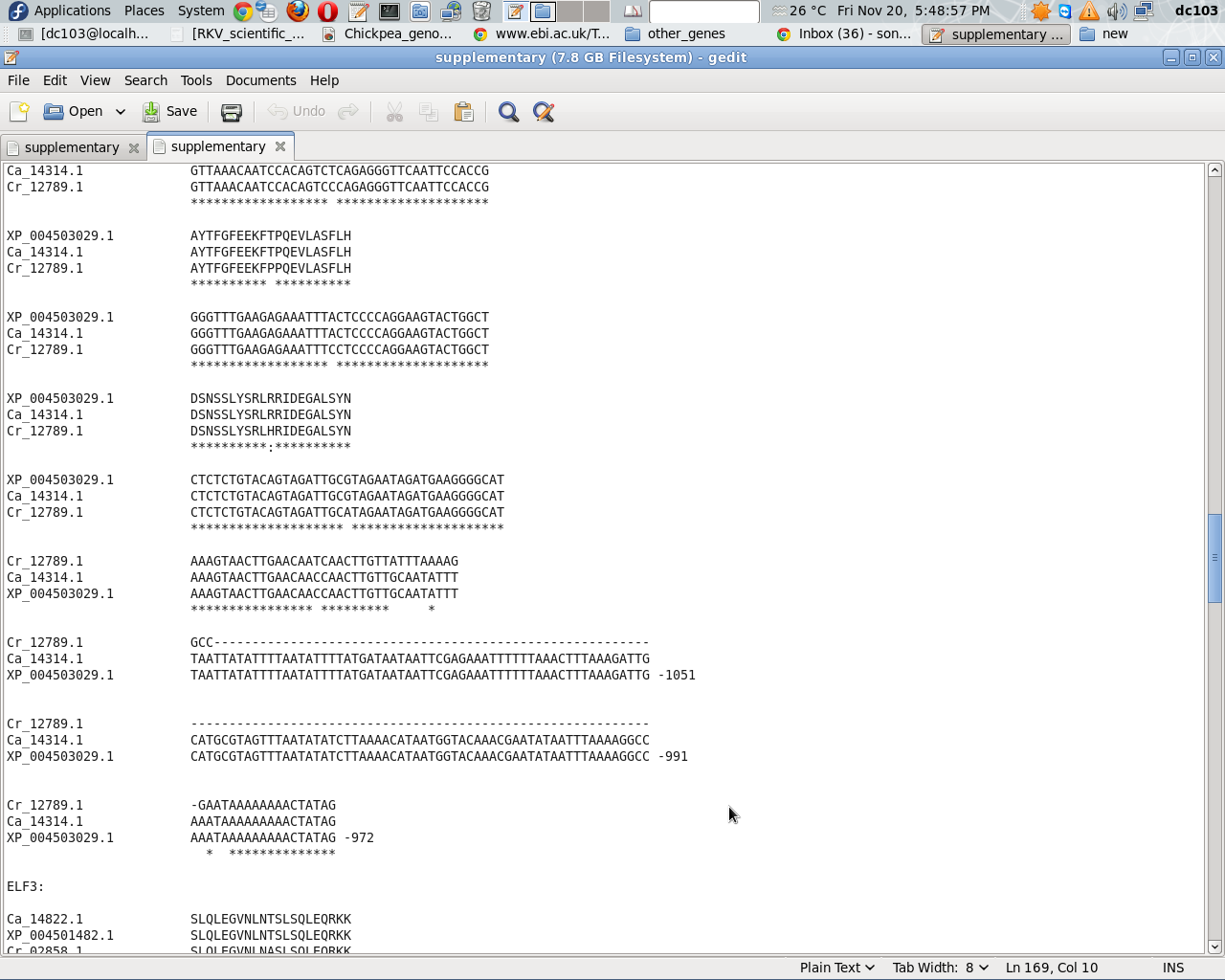
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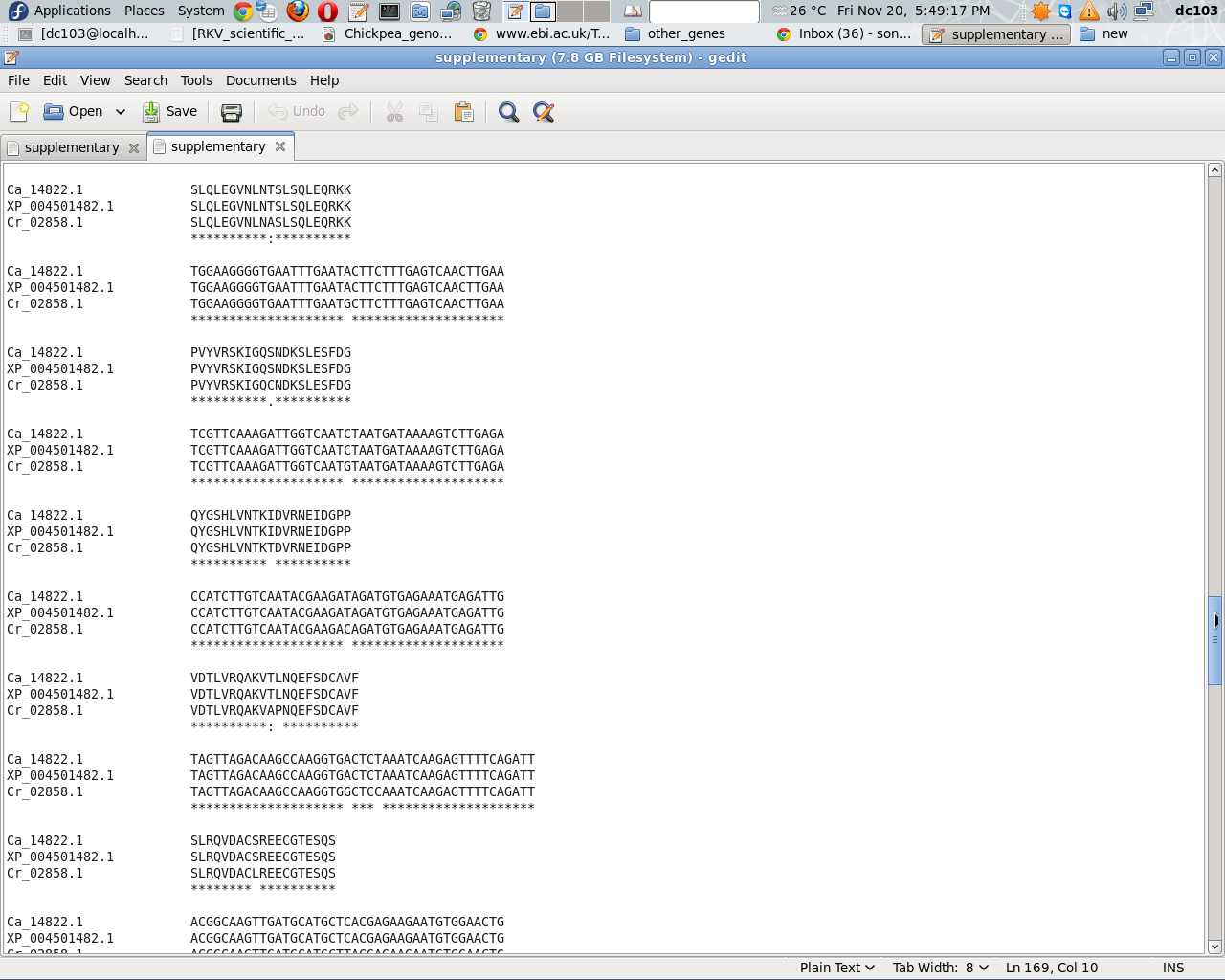
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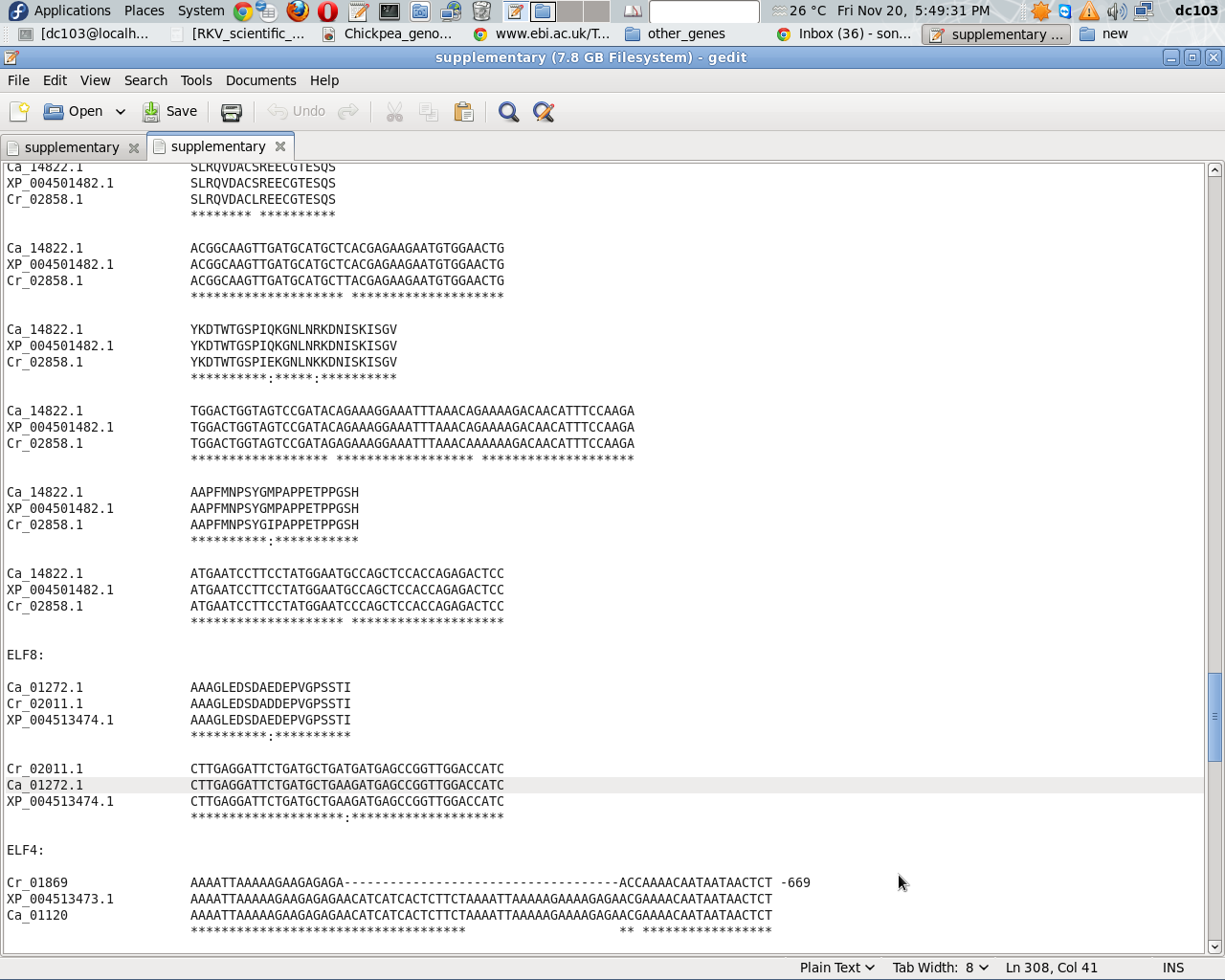




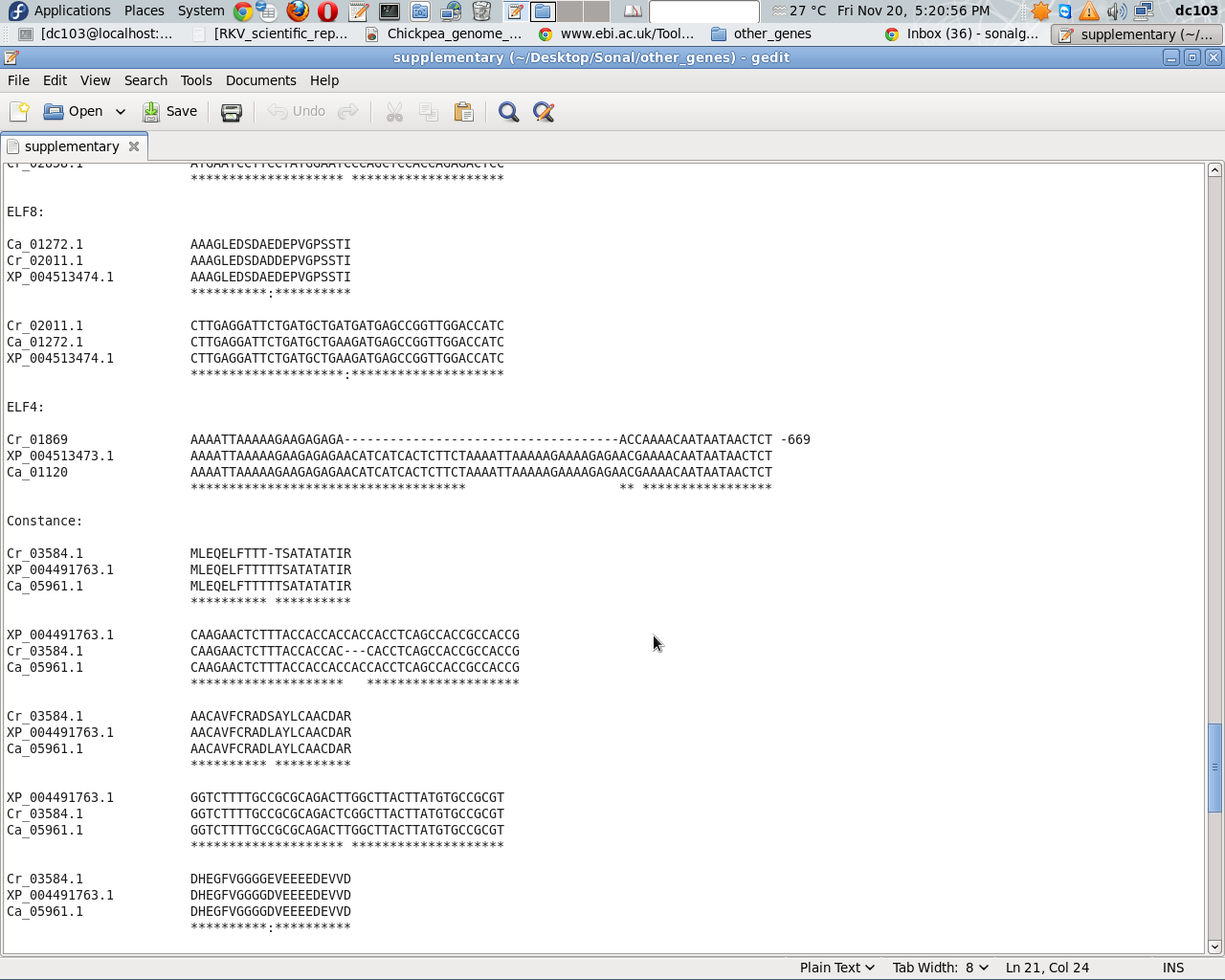


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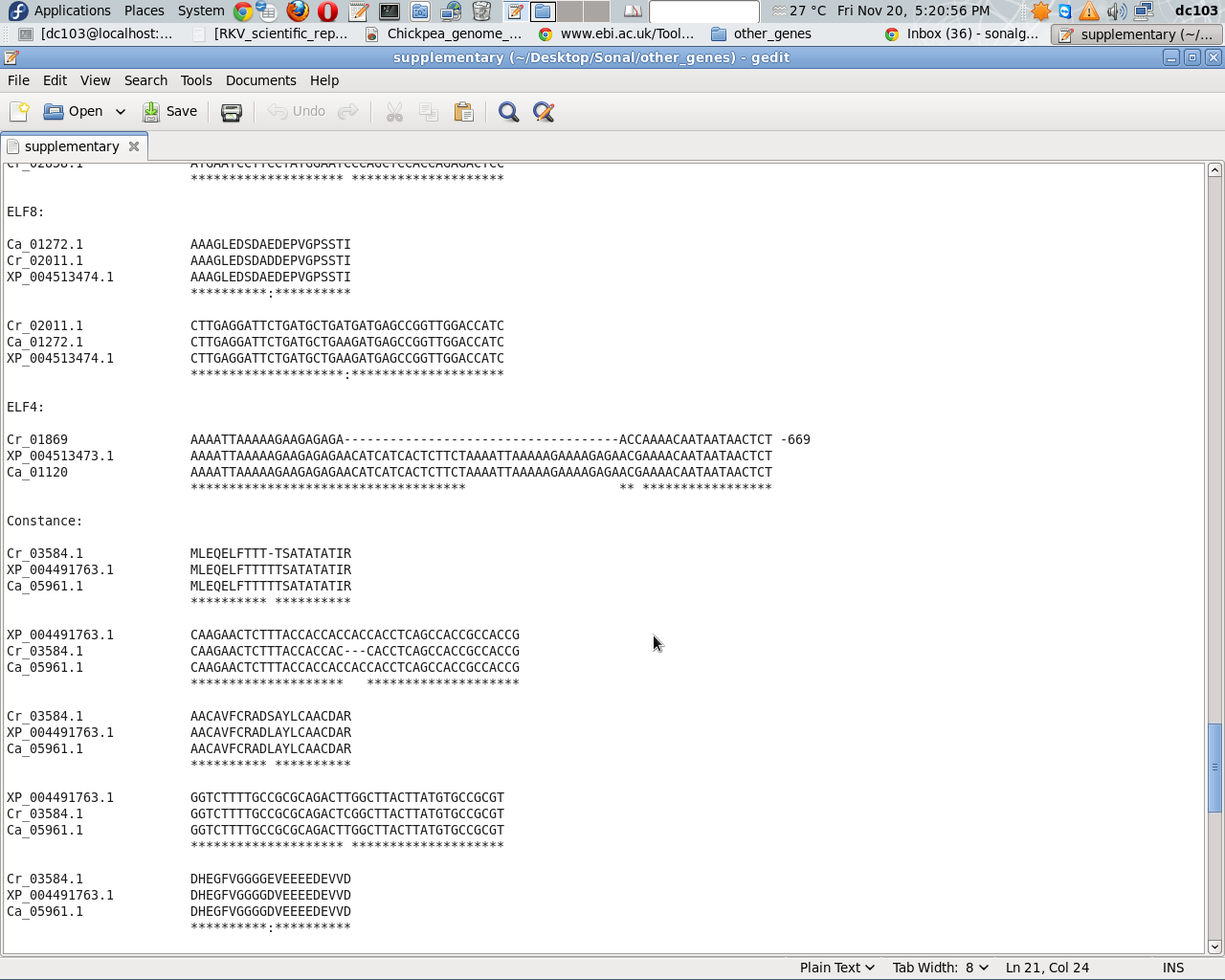




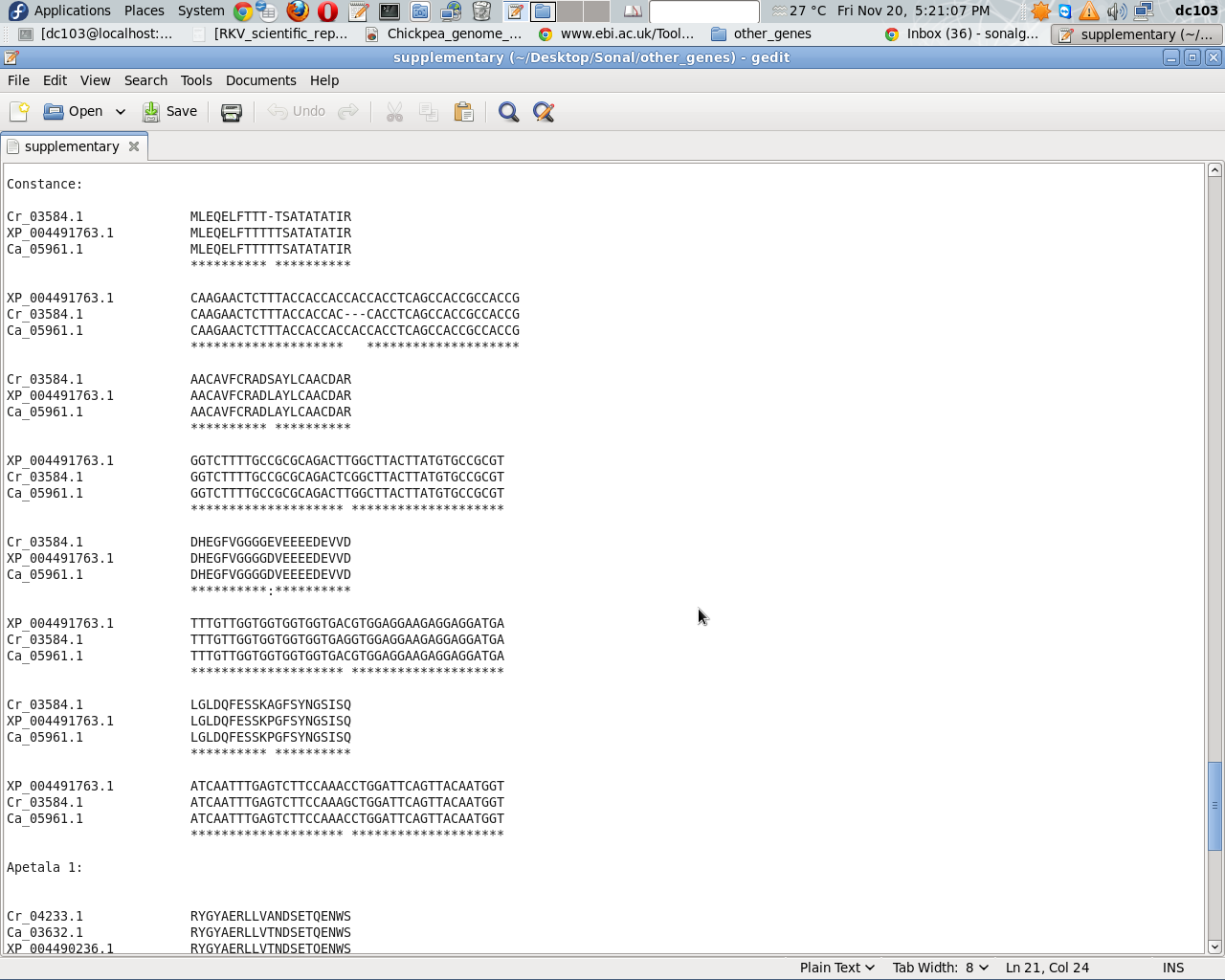
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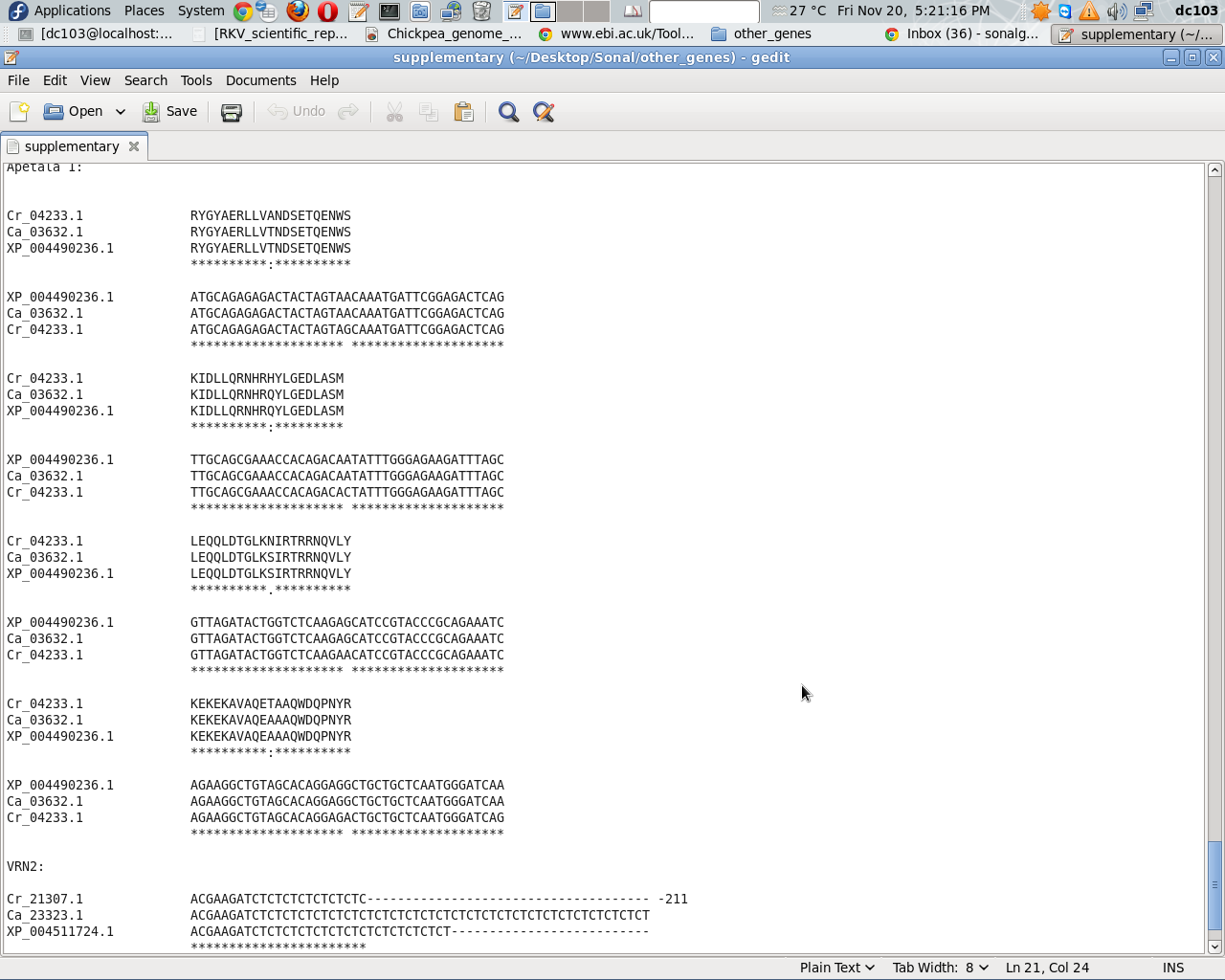
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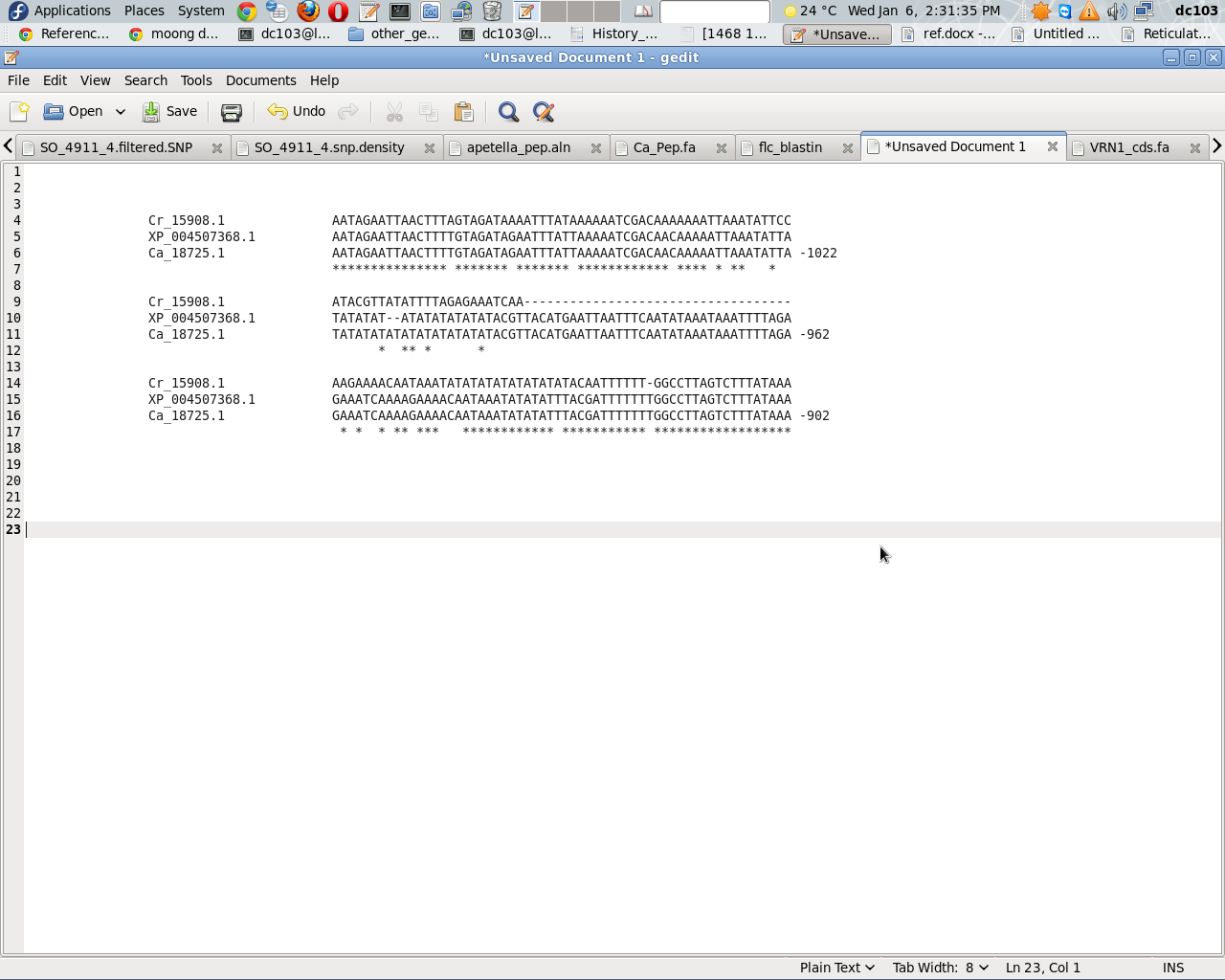
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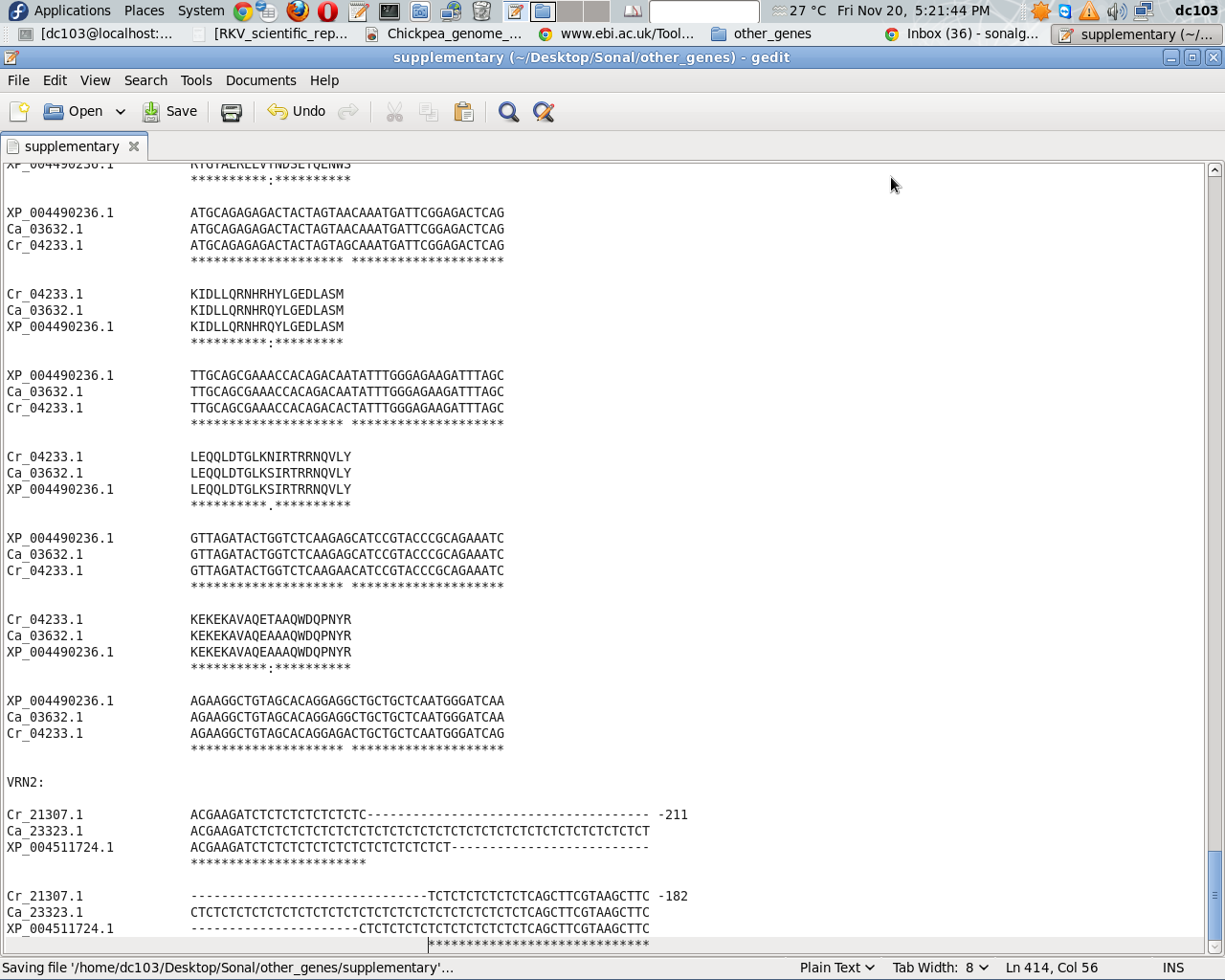
Apetala 1



VRN1



VRN2



BB

