

Analysis of Nasopharyngeal Carcinoma using Exome Sequencing

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Abstract

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Acknowledgments

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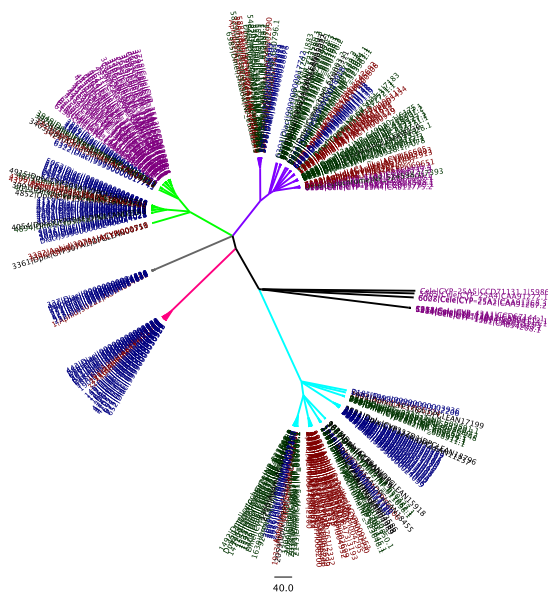


Figure 1. Foobar. Rest of figure 2 caption. Caption should be left justified, as specified by the options to the caption package.

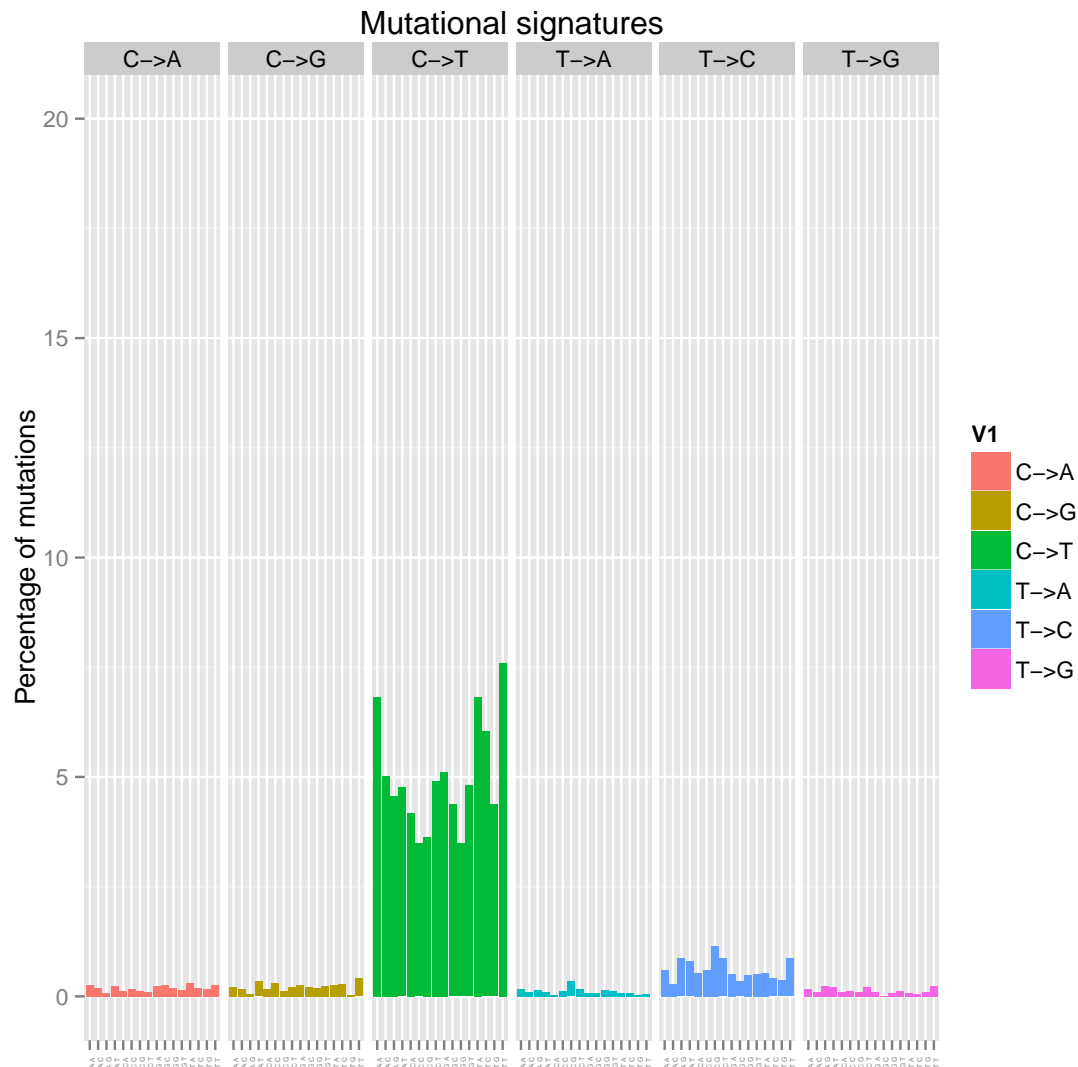


Figure 2. Mutational signature of somatic mutations generated using NPC tumor/normal pair. Rest of figure 2 caption. Caption should be left justified, as specified by the options to the caption package.