

# Description of Additional Supplementary File

## Discovering the drivers of clonal hematopoiesis

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**Supplementary Data file 1. Mutational profiles of signatures active across blood samples in the metastasis cohort.**

**Supplementary Data file 2. Compendium of clonal hematopoiesis driver genes.**

**Supplementary Data file 3. Non-coding elements significant by the analysis of three driver discovery methods.**

**Supplementary Data file 4. Examples of mutations potentially affecting transcription factor binding sites in enhancer elements.** P-values associated with the score of Transcription Factor binding to wild-type sequences within regulatory elements were computed using FIMO (see main Methods.)