

# Supplementary Materials: Targeted Next-Generation Sequencing of 117 Routine Clinical Samples Provides Further Insights into the Molecular Landscape of Uveal Melanoma.

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Table S1. Concordance Data

Table S2. NGS Quality Comparison

Table S3. Clinical, Molecular and Histopathological data 117 patients

Enrichment Method	Reads Mapped %	Reads Mapped % after UMI Removed	Number of Bases with Coverage More Than 0 (%)	Mean Depth of Coverage for the Bases with Coverage More Than 0
PCR	92.06	16.33	92.99	302x
Hybrid Capture	97.68	74.37	99.99	497.25x

Table S4. Monosomy 3 SF3B1 Cohort Data.



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