Large registry analysis to accurately define second malignancy rates and risks in a well-characterized cohort of 744 consecutive multiple myeloma patients followed-up for 25 years

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Contributions: ME, RW designed the research; ME, GI, MK, OL, MP, HR, JW, AMM, MS collected data; ME, MK, RW, GI, OL, MS analyzed the data; all authors interpreted the data and made critical contributions to the manuscript. ME wrote the manuscript.