












CORRECTION **OPEN**

Correction: Tracking daratumumab clearance using mass spectrometry: implications on M protein monitoring and reusing daratumumab

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We noted a mistake in the Introduction section in the following sentence: “and ability to differentiate therapeutic monoclonal antibodies from endogenous M proteins based on unique retention time and mass”. The use of mass and retention time is actually not true for the MALDI method. We would like to correct this sentence and replace it with: “with ability to aid in differentiating therapeutic monoclonal antibodies from endogenous M proteins on the basis of mass to charge”.



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