Author's response to reviews

Title: Screening of exon 11 for BRCA1 using the High Resolution melting for diagnostic in Moroccan breast cancer patients

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Reviewer: Emilia Wiechec

Minor points:

Question 1 The authors state that the method is highly sensitive but do not support this statement; what was the lowest amount of DNA necessary for the analysis?

➔ We have now specified the lowest amount of DNA necessary for the analysis (20 ng).

Question 3 How many patients out of 71 screened showed mutations? only 2?

➔ Yes, only two.

Question 4 The authors state that the results were validated with SNaPshot analysis, which is not mentioned anywhere apart from the abstract and no results is shown.

➔ The software used in the analysis was “Applied Biosystems SeqScape Software v2.5 SNaPshot analysis”. We have now included this information in the new version of the manuscript.

Question 5 It would be good to have more information about the patients chosen for the study

➔ This is a good point. We indeed have now added the clinical data of the patients showing mutations.

Question 6 English grammar should be corrected in few places.

➔ We have now corrected this in the new version of the manuscript.