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INTRODUCTION

The development of new disease biomarkers is essential for the progress of personalized medicine. This clinical approach, allows a great number of patients to access more efficient and safer therapeutic protocols, based on molecular findings in tissue samples obtained from patients for diagnostic or therapeutic purposes.

Biobanks, as repositories of human biological samples, have the responsibility to provide researchers with samples of known quality and an estimated potential for the analysis of DNA, RNA, proteins and antigenicity. A key element in this context is the knowledge, control and registration of all pre-analytical factors related to the final analytical performance of samples.

This issue is particularly complex when dealing with samples from a wide range of tissues, obtained through various procedures (biopsy, resection or post-mortem) and potentially tested with different analytical procedures.

OPTIMARK PROJECT

The purpose of the project, that is currently being carried out by 17 centres of the Spanish Network of Biobanks, is to select and validate the essential preanalytical variables relevant in analytical testing and high predictive value for the applicability of research samples of new tissue biomarkers.

The project will allow a retrospective analysis of the potential of tissue samples stored in the Network, and the effectiveness of the current codification systems that include pre-analytical variables. Additionally, the project aims at developing optimized protocols for obtaining samples of various tissue origins, controlling key pre-analytical factors (Fig. 1), and the evaluation (algorithms) of analytes (generic and specific) related to biomarkers (Fig. 2).

In the first phase, 374 patients between 30 and 65 years (in the case of brain samples up to 75 years without vascular pathology)

EXPECTED IMPACT OF PRE-ANALYTICAL FACTORS ON TISSUE QUALITY

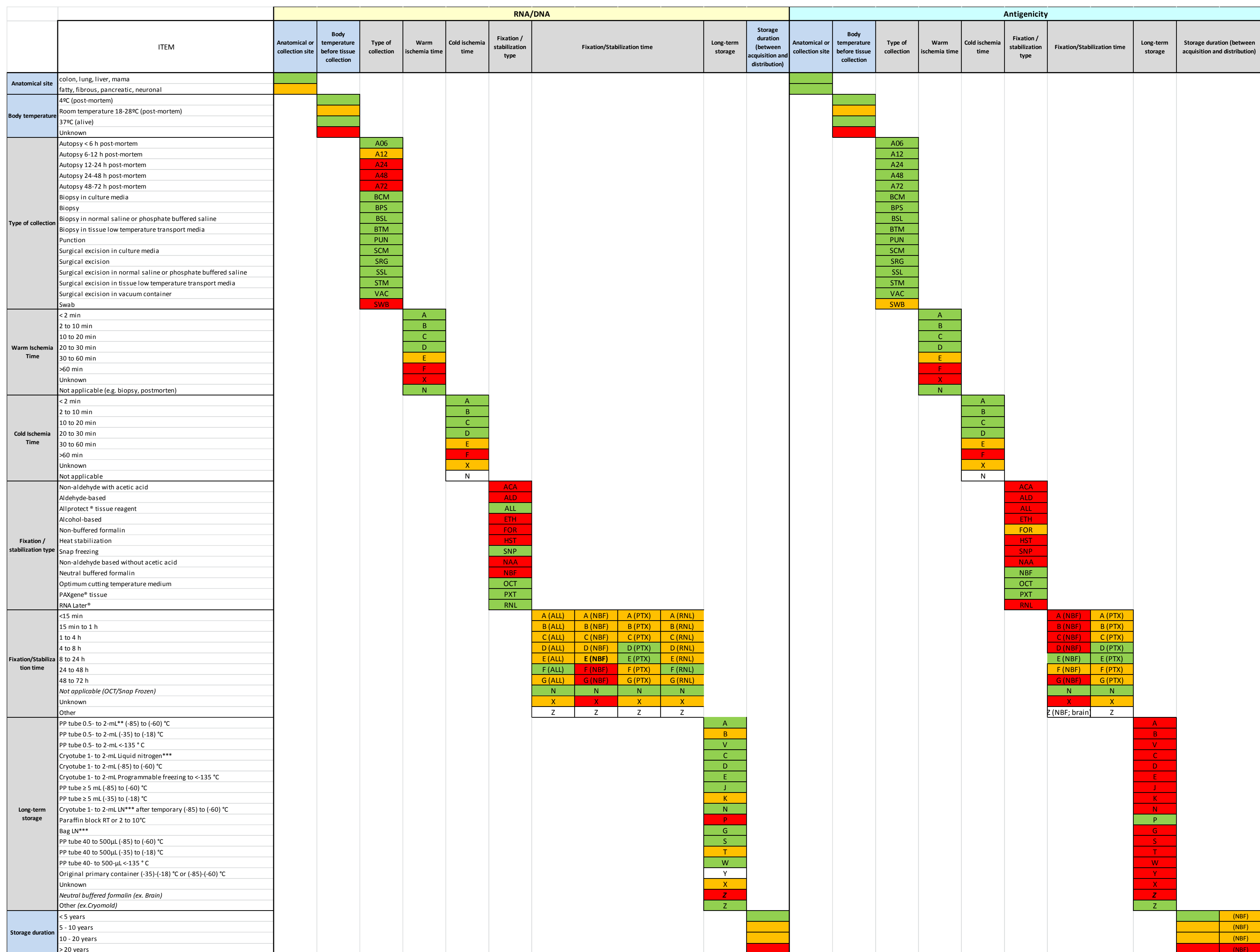


Figure 1. Expected impact by colours of selected SPREC v.2.0 (gray cell) and BRISQ (blue cell) preanalytical factors on tissue quality, over the nucleic acids and antigenicity. Green: optimal quality or not effect; yellow: moderate quality or unknown effect; red: sub optimal quality.

have been selected to test their tissue samples in order to evaluate long term storage based on RNA and antigenicity quality. A representative number of frozen and formalin-fixed paraffin-embedded tissue samples stored less than 1 year, 1-5 years, 5-10 years and more than 20 years have been selected from colon (n=104), brain (57), lung (76), breast (85), stomach (17) and endometrium (35).

Correlations will help to analyse the impact of other key preanalytical factors.

PROPOSALS AND CONCLUSIONS

1. The SPREC coding system and a subset of BRISQ variables have been used to estimate the expected quality for each analyte group (Fig. 1).
2. The Research Group has addressed strategies independently, and a decision tree has been developed to assess the quality of each analyte group (Fig. 2).
3. Selected retrospective samples are being analysed in order to validate the proposed procedure, to evaluate the effect of several preanalytical factors, as well as the cost-effectiveness in tissue biomarkers research.
4. A prospective tissue sampling phase shall be required to validate the gaps on available information.

ALGORITHMS PROPOSED TO TEST QUALITY

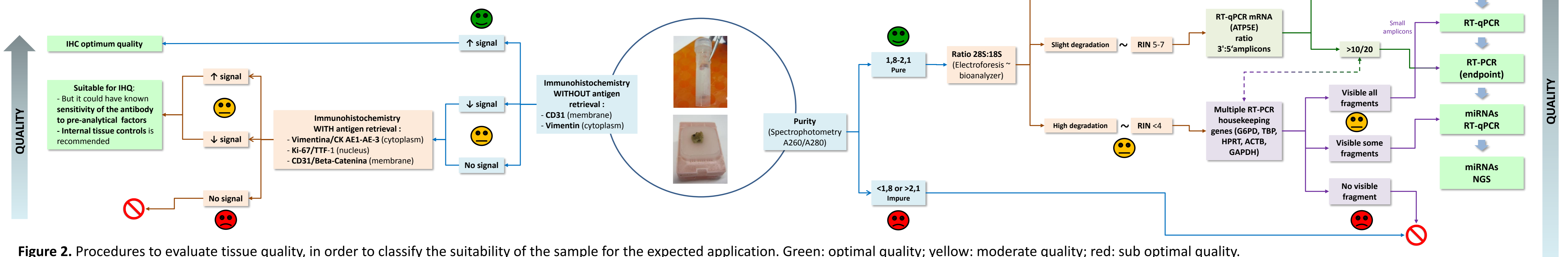


Figure 2. Procedures to evaluate tissue quality, in order to classify the suitability of the sample for the expected application. Green: optimal quality; yellow: moderate quality; red: sub optimal quality.

INSTITUTIONS