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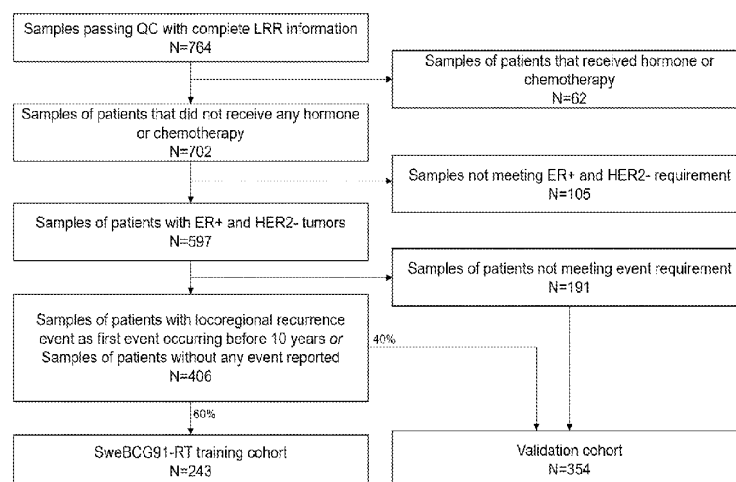
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(54) Title: METHODS AND GENOMIC CLASSIFIERS FOR PROGNOSIS OF BREAST CANCER AND IDENTIFYING SUBJECTS NOT LIKELY TO BENEFIT FROM RADIOTHERAPY

FIG. 1



(57) Abstract: The present disclosure relates to systems and methods for providing individualized prognostic assessments of breast cancer recurrence (e.g., locoregional recurrence). The systems and methods involve measuring gene expression from a patient sample to create a gene expression signature which identifies subjects who are not likely to benefit from radiotherapy following breast cancer surgery.

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