Supplementary Figure S27. Stratification of BC patients using the epigenetic dormancy signature

а

Genes included in the

Expression

low

high

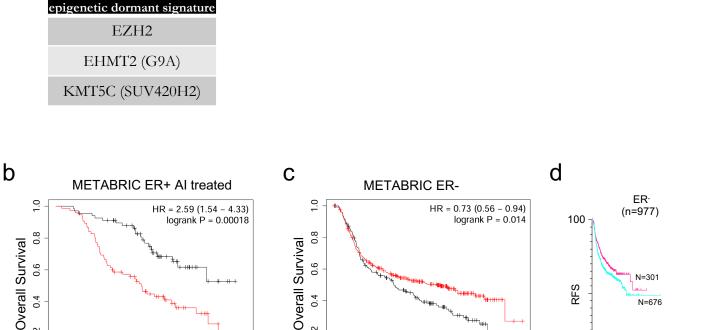
50

100

OS (updated) time (months)

150

200



N=676

High

Low

100 150 200

Months

HR 0.813

0.63 - 1

0 50

P=0.056

0.4

0.2

Expression

low

high

100

150

OS (updated) time (months)

200

Supplementary Figure S27. Stratification of BC patients using the epigenetic dormancy signature. a) Genes included in the epigenetic dormancy signature b) Overall Survival (OS) curves for ER-positive (ER+) patients stratified based on the expression of the epigenetic dormancy signature (high vs low G9a/EZH2/KMT5C expression) c) Overall Survival (OS) curves for ER-negative (ER-) patients treated with Aromatase Inhibitors (AI) stratified based on the expression of the epigenetic dormancy signature d) Relapse free survival (RFS) curves for ER-negative (ER-) patients stratified based on the expression of the epigenetic dormancy signature.