



**Supplementary Figure S7: Relationship of the 26 probeset signature to the 16 metagenes and seven known prognostic signatures in TNBC**

The 394 TNBC samples were analyzed for the expression of 16 metagenes and seven previously published prognostic signatures as described in Figure 3 and hierarchical clustered using Pearson correlation as distance metric.

Abbreviations:

SAMmean = 26 probeset signature

wound.score\$score = Wound response signature

rs.394\$score = recurrence score

ggi.score\$score = genomic grade index

gene70.score\$score = 70-gene signature

sabatier.score\$score = medullary like signature

Tesch7.score\$score = 7-gene immune response module

sdpp.sore\$score = stroma derived prognostic predictor