Figure S3. PCA plots. A) Basal-like *BRCA1* (*n* = 20) and sporadic (*n* = 10) tumors visualized using the 110 reporter genes included in the basal *BRCA1* signature. B) Luminal B *BRCA2* (*n* = 16) and sporadic (*n* = 48) tumors visualized using the 100 reporter genes included in the *BRCA2* signature.