Thank you very much for your kind considerations and constructive comments. Following your advices, we immediately reanalyzed the data excluding those of TNM stage 4 and new 46 cases enrollment .It was still found that the expression level of VGLL3 was correlated with N stage, TNM stage and HER2 status. COX univariate and multivariate analyses show that high levels of VGLL3 expression was independent poor prognostic factors.

The original table 1 ,2 and figure 3 was corrected.

The language was corrected again

Thanks again for waiting for a long time reply.